

**Supplementary Data 6. Novel alternative spliced transcripts annotated in AceView.**

<b>Gene</b>	<b>mRNA novel (not in RefSeq)</b>	<b>NCBI gene ID (if any)</b>	<b>Transcript extent on genome (nt)</b>	<b>Transcript length (nt)</b>	<b>Number of exons</b>	<b>CDS length (aa)</b>	<b>Gene descriptor</b>
<a href="#">4HBT.0</a>	<a href="#">4HBT.0.aSep08</a>		35437	721	6	240	thioesterase adipose associated BFIT1 (4HBT.0) alternative variant aSep08, mRNA.
<a href="#">4HBT.0</a>	<a href="#">4HBT.0.bSep08</a>		2496	399	3	132	thioesterase adipose associated BFIT1 (4HBT.0) alternative variant bSep08, mRNA.
<a href="#">4HBT.0</a>	<a href="#">4HBT.0.cSep08</a>		28355	342	3	90	thioesterase adipose associated BFIT1 (4HBT.0) alternative variant cSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.aSep08</a>		83571	5087	18	593	5'-nucleotidase cytosolic II (5_nucleotid.0) alternative variant aSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.bSep08</a>		71057	1793	8	580	5'-nucleotidase cytosolic II (5_nucleotid.0) alternative variant bSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.cSep08</a>		64390	740	9	229	5'-nucleotidase cytosolic II (5_nucleotid.0) alternative variant cSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.dSep08</a>		11283	927	10	178	cytosolic II (5_nucleotid.0) alternative variant dSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.eSep08</a>		3405	416	6	116	5'-Nucleotidase cytosolic II (5_nucleotid.0) alternative variant eSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.fSep08</a>		90074	623	3	60	putative protein (5_nucleotid.0) alternative variant fSep08, mRNA.
<a href="#">5_nucleotid.0</a>	<a href="#">5_nucleotid.0.gSep08</a>		1072	831	3	54	5'-nucleotidase cytosolic II CRA b (5_nucleotid.0) alternative variant gSep08, mRNA.
<a href="#">6PGD.0</a>	<a href="#">6PGD.0.aSep08</a>		16205	2132	13	514	phosphogluconate dehydrogenase (6PGD.0) alternative variant aSep08, mRNA.
<a href="#">6PGD.0</a>	<a href="#">6PGD.0.bSep08</a>		4258	976	4	158	phosphogluconate dehydrogenase (6PGD.0) alternative variant bSep08, mRNA.
<a href="#">6PGD.0</a>	<a href="#">6PGD.0.cSep08</a>		2911	875	4	144	phosphogluconate dehydrogenase (16.2 kD) (6PGD.0) alternative variant cSep08, mRNA.
<a href="#">6PGD.0</a>	<a href="#">6PGD.0.dSep08</a>		1203	391	2	123	phosphogluconate dehydrogenase (6PGD.0) alternative variant dSep08, mRNA.
<a href="#">6PGD.0</a>	<a href="#">6PGD.0.eSep08</a>		842	350	2	77	CRA d like (6PGD.0) alternative variant eSep08, mRNA.
<a href="#">7a5</a>	<a href="#">7a5.aSep08</a>	<a href="#">314539</a>	11886	524		95	putative binding protein 7a5 (7a5) mRNA.
<a href="#">7tm_2.0</a>	<a href="#">7tm_2.0.aSep08</a>		17118	742		224	brain-specific angiogenesis inhibitor (7tm_2.0) mRNA.
<a href="#">2331ex4-5</a>	<a href="#">2331ex4-5.aSep08</a>	<a href="#">415058</a>	2317	356		61	class I gene fragment 2331 (2331ex4-5) mRNA.
<a href="#">2747a1a2</a>	<a href="#">2747a1a2.aSep08</a>	<a href="#">414861</a>	17174	658		87	2747a1a2 pseudogene (9.2 kD) (2747a1a2) mRNA.
<a href="#">A</a>	<a href="#">A.bSep08</a>	<a href="#">24152</a>	91664	637		103	agouti (11.3 kD) (A) alternative variant bSep08, mRNA.
<a href="#">A1cf</a>	<a href="#">A1cf.bSep08</a>	<a href="#">170912</a>	12864	388	2	82	putative protein (A1cf) alternative variant bSep08, mRNA.
<a href="#">A2bp1</a>	<a href="#">A2bp1.bSep08</a>	<a href="#">302920</a>	58346	3167	6	132	ataxin 2 binding protein 1 (A2bp1) alternative variant bSep08, mRNA.
<a href="#">A2bp1</a>	<a href="#">A2bp1.cSep08</a>	<a href="#">302920</a>	203638	809	2	113	ataxin 2 binding protein 1 (A2bp1) alternative variant cSep08, mRNA.
<a href="#">A2bp1</a>	<a href="#">A2bp1.eSep08</a>	<a href="#">302920</a>	37031	493	2	46	ataxin 2 binding protein 1 (A2bp1) alternative variant eSep08, mRNA.

<a href="#">A2m</a>	<a href="#">A2m.cSep08</a>	<a href="#">24153</a>	98876	1036	7	345	murinoglobulin (A2m) alternative variant cSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.cSep08</a>	<a href="#">297568</a>	98876	1036	7	345	murinoglobulin (A2m) alternative variant cSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.dSep08</a>	<a href="#">24153</a>	7540	1513	8	205	CRA b (22.7 kD) (A2m) alternative variant dSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.dSep08</a>	<a href="#">297568</a>	7540	1513	8	205	CRA b (22.7 kD) (A2m) alternative variant dSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.eSep08</a>	<a href="#">24153</a>	4166	494	4	145	alpha-2-macroglobulin CRA b (A2m) alternative variant eSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.eSep08</a>	<a href="#">297568</a>	4166	494	4	145	alpha-2-macroglobulin CRA b (A2m) alternative variant eSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.fSep08</a>	<a href="#">24153</a>	2801	1336	3	56	CRA b (A2m) alternative variant fSep08, mRNA.
<a href="#">A2m</a>	<a href="#">A2m.fSep08</a>	<a href="#">297568</a>	2801	1336	3	56	CRA b (A2m) alternative variant fSep08, mRNA.
<a href="#">A2M.1</a>	<a href="#">A2M.1.aSep08</a>		2852	490		163	alpha-2-macroglobulin (A2M.1) mRNA.
<a href="#">A2M.2</a>	<a href="#">A2M.2.aSep08</a>		10711	318		105	complement component (A2M.2) mRNA.
<a href="#">A2M_recep.0</a>	<a href="#">A2M_recep.0.aSep08</a>		5160	513		127	murinoglobulin (A2M_recep.0) mRNA.
<a href="#">A2M_recep.1</a>	<a href="#">A2M_recep.1.aSep08</a>		4325	506		125	murinoglobulin (A2M_recep.1) mRNA.
<a href="#">A3galt2</a>	<a href="#">A3galt2.bSep08</a>	<a href="#">171553</a>	8487	613	4	111	alpha 1,3-galactosyltransferase 2 (12.9 kD) (A3galt2) alternative variant bSep08, mRNA.
<a href="#">A4galt</a>	<a href="#">A4galt.bSep08</a>	<a href="#">63888</a>	1889	406	1	135	alpha 1,4-galactosyltransferase (A4galt) alternative variant bSep08, mRNA.
<a href="#">A4_EXTRA.0</a>	<a href="#">A4_EXTRA.0.aSep08</a>		33327	385		128	amyloid -like protein 2 (A4_EXTRA.0) mRNA.
<a href="#">AA926063</a>	<a href="#">AA926063.aSep08</a>	<a href="#">294284</a>	8208	815	1	44	AA926063gene (AA926063) alternative variant aSep08, mRNA.
<a href="#">AA926063</a>	<a href="#">AA926063.bSep08</a>	<a href="#">294284</a>	12519	416	1	97	AA926063gene (10.3 kD) (AA926063) alternative variant bSep08, mRNA.
<a href="#">Aaas</a>	<a href="#">Aaas.aSep08</a>	<a href="#">300259</a>	9396	1790	10	516	achalasia adrenocortical insufficiency alacrimia (Aaas) alternative variant aSep08, mRNA.
<a href="#">Aaas</a>	<a href="#">Aaas.cSep08</a>	<a href="#">300259</a>	9189	1459	13	275	achalasia adrenocortical insufficiency alacrimia (Aaas) alternative variant cSep08, mRNA.
<a href="#">Aaas</a>	<a href="#">Aaas.dSep08</a>	<a href="#">300259</a>	1000	704	3	137	achalasia adrenocortical insufficiency alacrimia (14.7 kD) (Aaas) alternative variant dSep08, mRNA.
<a href="#">Aaas</a>	<a href="#">Aaas.eSep08</a>	<a href="#">300259</a>	10864	688	7	115	achalasia adrenocortical insufficiency alacrimia (Aaas) alternative variant eSep08, mRNA.
<a href="#">AAA_5.0</a>	<a href="#">AAA_5.0.aSep08</a>		6002	731		235	axonemal dynein heavy Dnahc8 (AAA_5.0) mRNA.
<a href="#">AAA_5.1</a>	<a href="#">AAA_5.1.aSep08</a>		80616	1584		442	ATPase associated with various cellular activities, AAA-5 (AAA_5.1) mRNA.
<a href="#">Aadat</a>	<a href="#">Aadat.bSep08</a>	<a href="#">29416</a>	40594	1798	7	387	aminoadipate aminotransferase (Aadat) alternative variant bSep08, complete mRNA.
<a href="#">Aak1</a>	<a href="#">Aak1.aSep08</a>	<a href="#">500244</a>	97336	2011	2	533	AP2 associated kinase 1 (Aak1) alternative variant aSep08, mRNA.
<a href="#">Aak1</a>	<a href="#">Aak1.bSep08</a>	<a href="#">500244</a>	18543	1167	5	388	AP2 associated kinase 1 (Aak1) alternative variant bSep08, mRNA.
<a href="#">Aak1</a>	<a href="#">Aak1.cSep08</a>	<a href="#">500244</a>	96627	1442	3	341	AP2 associated kinase 1 (38.1 kD) (Aak1) alternative variant cSep08, complete mRNA.
<a href="#">Aak1</a>	<a href="#">Aak1.dSep08</a>	<a href="#">500244</a>	8426	347	3	115	AP2 associated kinase 1 (Aak1) alternative variant dSep08, mRNA.

<a href="#">Aamp</a>	<a href="#">Aamp.bSep08</a>	<a href="#">301512</a>	4108	1662	7	266	angio-associated, migratory cell protein (28.8 kD) (Aamp) alternative variant bSep08, mRNA.
<a href="#">AARP2CN.0</a>	<a href="#">AARP2CN.0.aSep08</a>		4695	462		153	ribosome assembly protein CRA b (AARP2CN.0) mRNA.
<a href="#">Aars</a>	<a href="#">Aars.aSep08</a>	<a href="#">292023</a>	21985	3460	22	968	alanyl-tRNA synthetase (106.8 kD) (Aars) alternative variant aSep08, mRNA.
<a href="#">Aars</a>	<a href="#">Aars.bSep08</a>	<a href="#">292023</a>	2270	598	5	193	alanyl-tRNA synthetase (Aars) alternative variant bSep08, mRNA.
<a href="#">Aars</a>	<a href="#">Aars.cSep08</a>	<a href="#">292023</a>	3735	769	4	171	alanyl-tRNA synthetase (Aars) alternative variant cSep08, mRNA.
<a href="#">Aars</a>	<a href="#">Aars.dSep08</a>	<a href="#">292023</a>	6390	408	4	135	alanyl-tRNA synthetase (Aars) alternative variant dSep08, mRNA.
<a href="#">Aars</a>	<a href="#">Aars.fSep08</a>	<a href="#">292023</a>	2912	433	2	76	alanyl-tRNA synthetase (Aars) alternative variant fSep08, mRNA.
<a href="#">Aarsd1</a>	<a href="#">Aarsd1.bSep08</a>	<a href="#">619440</a>	3981	716	5	87	putative cytoplasmic protein of metazoan origin (9.2 kD) (Aarsd1) alternative variant bSep08, mRNA.
<a href="#">Aarsd1</a>	<a href="#">Aarsd1.cSep08</a>	<a href="#">619440</a>	614	307	3	65	threonyl/alanyl tRNA synthetase, SAD (Aarsd1) alternative variant cSep08, mRNA.
<a href="#">Aasdh</a>	<a href="#">Aasdh.aSep08</a>	<a href="#">364136</a>	2635	707		205	aminoadipate-semialdehyde dehydrogenase (Aasdh) mRNA.
<a href="#">Aasdhppt</a>	<a href="#">Aasdhppt.bSep08</a>	<a href="#">300328</a>	11150	1781	3	252	aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase (29.4 kD) (Aasdhppt) alternative variant bSep08, mRNA.
<a href="#">Aasdhppt</a>	<a href="#">Aasdhppt.cSep08</a>	<a href="#">300328</a>	10971	789	2	119	aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase (14.0 kD) (Aasdhppt) alternative variant cSep08, mRNA.
<a href="#">Aass</a>	<a href="#">Aass.aSep08</a>	<a href="#">296925</a>	19610	1950		500	aminoadipate-semialdehyde synthase (Aass) alternative variant aSep08, mRNA.
<a href="#">Aass</a>	<a href="#">Aass.cSep08</a>	<a href="#">296925</a>	21082	734		229	aminoadipate-semialdehyde synthase (Aass) alternative variant cSep08, mRNA.
<a href="#">Aass</a>	<a href="#">Aass.dSep08</a>	<a href="#">296925</a>	16921	871		221	aminoadipate-semialdehyde synthase (Aass) alternative variant dSep08, mRNA.
<a href="#">Aass</a>	<a href="#">Aass.eSep08</a>	<a href="#">296925</a>	1479	260		86	aminoadipate-semialdehyde synthase (Aass) alternative variant eSep08, mRNA.
<a href="#">Abat</a>	<a href="#">Abat.bSep08</a>	<a href="#">81632</a>	36734	792	1	264	4-aminobutyrate aminotransferase (Abat) alternative variant bSep08, mRNA.
<a href="#">Abba-1</a>	<a href="#">Abba-1.aSep08</a>	<a href="#">307845</a>	10440	598	6	199	actin-bundling protein with BAIAP2 homology (Abba-1) alternative variant aSep08, mRNA.
<a href="#">Abba-1</a>	<a href="#">Abba-1.bSep08</a>	<a href="#">307845</a>	2985	410	3	136	actin-bundling protein with BAIAP2 homology (Abba-1) alternative variant bSep08, mRNA.
<a href="#">Abba-1</a>	<a href="#">Abba-1.cSep08</a>	<a href="#">307845</a>	754	401	2	133	actin-bundling protein with BAIAP2 homology (Abba-1) alternative variant cSep08, mRNA.
<a href="#">Abba-1</a>	<a href="#">Abba-1.dSep08</a>	<a href="#">307845</a>	10109	396	4	132	actin-bundling protein with BAIAP2 homology (Abba-1) alternative variant dSep08, mRNA.
<a href="#">ABC2_membran e.0</a>	<a href="#">ABC2_membrane.0.aSe p08</a>		10293	3151	9	401	WHITE (ABC2_membrane.0) alternative variant aSep08, mRNA.
<a href="#">ABC2_membran e.0</a>	<a href="#">ABC2_membrane.0.bSe p08</a>		3515	590	3	146	WHITE (ABC2_membrane.0) alternative variant bSep08, mRNA.

<a href="#">ABC2_membran e.0</a>	<a href="#">ABC2_membrane.0.cSe p08</a>		9818	504	3	25	putative protein (ABC2_membrane.0) alternative variant cSep08, mRNA.
<a href="#">Abca1</a>	<a href="#">Abca1.bSep08</a>	<a href="#">313210</a>	5724	2107	6	264	ATP-binding cassette, sub-family A (ABC1), member 1 (Abca1) alternative variant bSep08, mRNA.
<a href="#">Abca2</a>	<a href="#">Abca2.bSep08</a>	<a href="#">79248</a>	1287	856	6	238	ATP-binding cassette sub-family A member 2 like (Abca2) alternative variant bSep08, mRNA.
<a href="#">Abca2</a>	<a href="#">Abca2.cSep08</a>	<a href="#">79248</a>	649	535	2	117	ATP-binding cassette sub-family A member 2 like (12.8 kD) (Abca2) alternative variant cSep08, mRNA.
<a href="#">Abca2</a>	<a href="#">Abca2.dSep08</a>	<a href="#">79248</a>	1293	744	4	100	ATP-binding cassette sub-family A member 2 like (11.1 kD) (Abca2) alternative variant dSep08, mRNA.
<a href="#">Abca3</a>	<a href="#">Abca3.aSep08</a>	<a href="#">302973</a>	13174	3456	14	815	ATP-binding cassette, sub-family A (ABC1), member 3 (Abca3) alternative variant aSep08, mRNA.
<a href="#">Abca4</a>	<a href="#">Abca4.bSep08</a>	<a href="#">310836</a>	11110	1778	7	524	ATP-binding cassette, sub-family A (ABC1), member 4 (Abca4) alternative variant bSep08, mRNA.
<a href="#">Abca4</a>	<a href="#">Abca4.cSep08</a>	<a href="#">310836</a>	6552	705	4	234	ATP-binding cassette, sub-family A (ABC1), member 4 (Abca4) alternative variant cSep08, mRNA.
<a href="#">Abca4</a>	<a href="#">Abca4.dSep08</a>	<a href="#">310836</a>	11587	615	5	205	ATP-binding cassette, sub-family A (ABC1), member 4 (Abca4) alternative variant dSep08, mRNA.
<a href="#">Abca4</a>	<a href="#">Abca4.eSep08</a>	<a href="#">310836</a>	2821	364	3	115	ATP-binding cassette, sub-family A (ABC1), member 4 (Abca4) alternative variant eSep08, mRNA.
<a href="#">Abca4</a>	<a href="#">Abca4.fSep08</a>	<a href="#">310836</a>	1650	365	2	52	ATP-binding cassette, sub-family A (ABC1), member 4 (Abca4) alternative variant fSep08, mRNA.
<a href="#">Abca5</a>	<a href="#">Abca5.bSep08</a>	<a href="#">286970</a>	738	623	2	32	ATP-binding cassette, sub-family A (ABC1), member 5 (3.8 kD) (Abca5) alternative variant bSep08, mRNA.
<a href="#">Abca7</a>	<a href="#">Abca7.aSep08</a>	<a href="#">299609</a>	7584	2758	20	891	ATP-binding cassette, sub-family A (ABC1), member 7 (Abca7) alternative variant aSep08, mRNA.
<a href="#">Abca7</a>	<a href="#">Abca7.bSep08</a>	<a href="#">299609</a>	3261	1347	7	198	ATP-binding cassette, sub-family A (ABC1), member 7 (Abca7) alternative variant bSep08, mRNA.
<a href="#">Abca7</a>	<a href="#">Abca7.cSep08</a>	<a href="#">299609</a>	677	342	3	113	ATP-binding cassette, sub-family A (ABC1), member 7 (Abca7) alternative variant cSep08, mRNA.
<a href="#">Abca7</a>	<a href="#">Abca7.dSep08</a>	<a href="#">299609</a>	1023	770	2	98	ATP-binding cassette, sub-family A (ABC1), member 7 (10.5 kD) (Abca7) alternative variant dSep08, mRNA.
<a href="#">Abca8</a>	<a href="#">Abca8.aSep08</a>	<a href="#">303637</a>	16174	2111		629	ATP-binding cassette, sub-family A (ABC1), member 8 (Abca8) alternative variant aSep08, mRNA.
<a href="#">Abca8</a>	<a href="#">Abca8.bSep08</a>	<a href="#">303637</a>	7001	739		171	ATP-binding cassette, sub-family A (ABC1), member 8 (Abca8) alternative variant bSep08, mRNA.
<a href="#">Abca8a</a>	<a href="#">Abca8a.aSep08</a>	<a href="#">303638</a>	34455	2819	20	726	ATP-binding cassette, sub-family A (ABC1), member 8a (Abca8a) alternative variant aSep08, mRNA.
<a href="#">Abca9</a>	<a href="#">Abca9.aSep08</a>	<a href="#">287788</a>	3696	378		125	ATP-binding cassette, sub-family A (ABC1), member 9 (Abca9) mRNA.
<a href="#">Abca16</a>	<a href="#">Abca16.aSep08</a>	<a href="#">293444</a>	40703	408		125	ATP-binding cassette, sub-family A (ABC1), member 16 (Abca16) mRNA.
<a href="#">Abcb1</a>	<a href="#">Abcb1.bSep08</a>	<a href="#">170913</a>	25433	458	4	97	ATP-binding cassette, sub-family B (MDR/TAP), member 1 (Abcb1) alternative variant bSep08, mRNA.
<a href="#">Abcb6</a>	<a href="#">Abcb6.bSep08</a>	<a href="#">140669</a>	5010	2336	2	341	ATP-binding cassette, sub-family B (MDR/TAP), member 6 (38.2 kD) (Abcb6) alternative variant bSep08, mRNA.

<a href="#">Abcb7</a>	<a href="#">Abcb7.bSep08</a>	<a href="#">302395</a>	34626	1785	7	384	ATP-binding cassette, sub-family B (MDR/TAP), member 7 (Abcb7) alternative variant bSep08, mRNA.
<a href="#">Abcb7</a>	<a href="#">Abcb7.cSep08</a>	<a href="#">302395</a>	7179	373	2	93	ATP-binding cassette, sub-family B (MDR/TAP), member 7 (Abcb7) alternative variant cSep08, mRNA.
<a href="#">Abcb8</a>	<a href="#">Abcb8.bSep08</a>	<a href="#">362302</a>	6491	901	6	224	ATP-binding cassette, sub-family B (MDR/TAP), member 8 (24.9 kD) (Abcb8) alternative variant bSep08, mRNA.
<a href="#">Abcb8</a>	<a href="#">Abcb8.cSep08</a>	<a href="#">362302</a>	5620	187	2	40	ATP-binding cassette, sub-family B (MDR/TAP), member 8 (Abcb8) alternative variant cSep08, mRNA.
<a href="#">Abcb9</a>	<a href="#">Abcb9.bSep08</a>	<a href="#">63886</a>	2338	790	2	123	ATP-binding cassette, sub-family B (MDR/TAP), member 9 (Abcb9) alternative variant bSep08, mRNA.
<a href="#">Abcc4</a>	<a href="#">Abcc4.bSep08</a>	<a href="#">170924</a>	43580	1787		568	ATP-binding cassette, sub-family C (CFTR/MRP), member 4 (Abcc4) alternative variant bSep08, mRNA.
<a href="#">Abcc4</a>	<a href="#">Abcc4.cSep08</a>	<a href="#">170924</a>	56766	1197		399	ATP-binding cassette, sub-family C (CFTR/MRP), member 4 (Abcc4) alternative variant cSep08, mRNA.
<a href="#">Abcc4</a>	<a href="#">Abcc4.dSep08</a>	<a href="#">170924</a>	9600	748		68	ATP-binding cassette, sub-family C (CFTR/MRP), member 4 (Abcc4) alternative variant dSep08, mRNA.
<a href="#">Abcc5</a>	<a href="#">Abcc5.bSep08</a>	<a href="#">116721</a>	34336	2183	6	223	ATP-binding cassette, sub-family C (CFTR/MRP), member 5 (25.7 kD) (Abcc5) alternative variant bSep08, mRNA.
<a href="#">Abcc5</a>	<a href="#">Abcc5.cSep08</a>	<a href="#">116721</a>	31246	1259	5	204	ATP-binding cassette, sub-family C (CFTR/MRP), member 5 (23.4 kD) (Abcc5) alternative variant cSep08, mRNA.
<a href="#">Abcc8</a>	<a href="#">Abcc8.bSep08</a>	<a href="#">25559</a>	926	605	2	190	atp-binding cassette sub-family c member 8 like (Abcc8) alternative variant bSep08, mRNA.
<a href="#">Abcc8</a>	<a href="#">Abcc8.dSep08</a>	<a href="#">25559</a>	1212	308	3	102	ATP-binding cassette C member 8 like (Abcc8) alternative variant dSep08, mRNA.
<a href="#">Abcc8</a>	<a href="#">Abcc8.eSep08</a>	<a href="#">25559</a>	2329	480	5	91	ATP-binding cassette C member 8 like (Abcc8) alternative variant eSep08, mRNA.
<a href="#">Abcc10</a>	<a href="#">Abcc10.cSep08</a>	<a href="#">316231</a>	506	287	2	95	ATP-binding cassette, sub-family C (CFTR/MRP), member 10 (Abcc10) alternative variant cSep08, mRNA.
<a href="#">Abcd1</a>	<a href="#">Abcd1.bSep08</a>	<a href="#">363516</a>	815	733	2	99	ATP-binding cassette, sub-family D (ALD), member 1 (Abcd1) alternative variant bSep08, mRNA.
<a href="#">Abcd1</a>	<a href="#">Abcd1.cSep08</a>	<a href="#">363516</a>	7361	1508	4	66	ATP-binding cassette, sub-family D (ALD), member 1 (Abcd1) alternative variant cSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.aSep08</a>	<a href="#">299196</a>	15890	1194	14	397	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant aSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.bSep08</a>	<a href="#">299196</a>	11232	769	5	255	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant bSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.cSep08</a>	<a href="#">299196</a>	5368	774	6	209	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant cSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.dSep08</a>	<a href="#">299196</a>	4034	581	4	143	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant dSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.eSep08</a>	<a href="#">299196</a>	2093	336	3	95	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant eSep08, mRNA.
<a href="#">Abcd4</a>	<a href="#">Abcd4.fSep08</a>	<a href="#">299196</a>	1420	595	4	67	ATP-binding cassette, sub-family D (ALD), member 4 (Abcd4) alternative variant fSep08, mRNA.
<a href="#">Abce1</a>	<a href="#">Abce1.aSep08</a>	<a href="#">361390</a>	18206	1247		364	ATP-binding cassette, sub-family E (OABP), member 1 (Abce1) mRNA.

<a href="#">Abcf1</a>	<a href="#">Abcf1.bSep08</a>	<a href="#">85493</a>	2209	1205	5	113	ATP-binding cassette, sub-family F (GCN20), member 1 (12.6 kD) (Abcf1) alternative variant bSep08, mRNA.
<a href="#">Abcf1</a>	<a href="#">Abcf1.dSep08</a>	<a href="#">85493</a>	739	414	4	94	ATP-binding cassette, sub-family F (GCN20), member 1 (Abcf1) alternative variant dSep08, mRNA.
<a href="#">Abcf2</a>	<a href="#">Abcf2.aSep08</a>	<a href="#">311959</a>	12670	2458	15	624	ATP-binding cassette, sub-family F (GCN20), member 2 (71.4 kD) (Abcf2) alternative variant aSep08, complete mRNA.
<a href="#">Abcf2</a>	<a href="#">Abcf2.bSep08</a>	<a href="#">311959</a>	4160	806	5	244	ATP-binding cassette, sub-family F (GCN20), member 2 (Abcf2) alternative variant bSep08, mRNA.
<a href="#">Abcf2</a>	<a href="#">Abcf2.cSep08</a>	<a href="#">311959</a>	2017	722	4	162	ATP-binding cassette, sub-family F (GCN20), member 2 (Abcf2) alternative variant cSep08, mRNA.
<a href="#">Abcf3</a>	<a href="#">Abcf3.bSep08</a>	<a href="#">287982</a>	6774	806	5	198	ATP-binding cassette sub-family F member 3 like (Abcf3) alternative variant bSep08, mRNA.
<a href="#">Abcf3</a>	<a href="#">Abcf3.cSep08</a>	<a href="#">287982</a>	7330	705	6	147	ATP-binding cassette sub-family F member 3 like (Abcf3) alternative variant cSep08, mRNA.
<a href="#">Abcf3</a>	<a href="#">Abcf3.dSep08</a>	<a href="#">287982</a>	1443	668	4	128	ATP-binding cassette sub-family F member 3 like (14.1 kD) (Abcf3) alternative variant dSep08, mRNA.
<a href="#">Abcg1</a>	<a href="#">Abcg1.aSep08</a>	<a href="#">85264</a>	59333	914	7	304	ATP-binding cassette, sub-family G (WHITE), member 1 (Abcg1) alternative variant aSep08, mRNA.
<a href="#">Abcg2</a>	<a href="#">Abcg2.bSep08</a>	<a href="#">312382</a>	100846	686	5	165	ATP-binding cassette, sub-family G (WHITE), member 2 (Abcg2) alternative variant bSep08, mRNA.
<a href="#">Abcg2</a>	<a href="#">Abcg2.cSep08</a>	<a href="#">312382</a>	24813	423	4	111	ATP-binding cassette, sub-family G (WHITE), member 2 (Abcg2) alternative variant cSep08, mRNA.
<a href="#">Abcg3</a>	<a href="#">Abcg3.bSep08</a>	<a href="#">498327</a>	29874	765	6	254	ATP-binding cassette, sub-family G (WHITE), member 3 (Abcg3) alternative variant bSep08, mRNA.
<a href="#">Abcg3</a>	<a href="#">Abcg3.cSep08</a>	<a href="#">498327</a>	10876	722	3	88	ATP-binding cassette, sub-family G (WHITE), member 3 (9.7 kD) (Abcg3) alternative variant cSep08, mRNA.
<a href="#">Abcg3l1</a>	<a href="#">Abcg3l1.bSep08</a>	<a href="#">289453</a>	10183	765	4	141	ATP-binding cassette, sub-family G (WHITE), member 3-like 1 (15.8 kD) (Abcg3l1) alternative variant bSep08, mRNA.
<a href="#">Abcg3l1</a>	<a href="#">Abcg3l1.cSep08</a>	<a href="#">289453</a>	1089	390	2	58	ATP-binding cassette, sub-family G (WHITE), member 3-like 1 (Abcg3l1) alternative variant cSep08, mRNA.
<a href="#">Abcg3l1</a>	<a href="#">Abcg3l1.dSep08</a>	<a href="#">289453</a>	4702	1442	3	31	ATP-binding cassette, sub-family G (WHITE), member 3-like 1 (3.6 kD) (Abcg3l1) alternative variant dSep08, mRNA.
<a href="#">Abcg3l2</a>	<a href="#">Abcg3l2.bSep08</a>	<a href="#">360910</a>	5415	403	1	89	ATP-binding cassette sub-family G member 3 family like (9.6 kD) (Abcg3l2) alternative variant bSep08, mRNA.
<a href="#">Abcg3l2</a>	<a href="#">Abcg3l2.bSep08</a>	<a href="#">360997</a>	5415	403	1	89	ATP-binding cassette sub-family G member 3 family like (9.6 kD) (Abcg3l2) alternative variant bSep08, mRNA.
<a href="#">Abcg4</a>	<a href="#">Abcg4.bSep08</a>	<a href="#">300664</a>	7807	1299	4	357	ATP-binding cassette, sub-family G (WHITE), member 4 (Abcg4) alternative variant bSep08, mRNA.
<a href="#">ABC_tran.0</a>	<a href="#">ABC_tran.0.bSep08</a>		18903	665	1	133	ATP-binding cassette member like (ABC_tran.0) alternative variant bSep08, mRNA.
<a href="#">ABC_tran.1</a>	<a href="#">ABC_tran.1.aSep08</a>		2713	979		273	ATP-binding cassette sub-family A member 7 like (ABC_tran.1) mRNA.
<a href="#">Abhd3</a>	<a href="#">Abhd3.bSep08</a>	<a href="#">291793</a>	11070	1277	5	214	putative protein of ancient origin (Abhd3) alternative variant bSep08, mRNA.

<a href="#">Abhd4</a>	<a href="#">Abhd4.bSep08</a>	<a href="#">364380</a>	14578	980	4	209	putative protein of ancient origin (Abhd4) alternative variant bSep08, mRNA.
<a href="#">Abhd4</a>	<a href="#">Abhd4.cSep08</a>	<a href="#">364380</a>	6277	693	3	173	putative protein of ancient origin (Abhd4) alternative variant cSep08, mRNA.
<a href="#">Abhd4</a>	<a href="#">Abhd4.dSep08</a>	<a href="#">364380</a>	11463	696	4	142	putative protein of eukaryotic origin (16.6 kD) (Abhd4) alternative variant dSep08, mRNA.
<a href="#">Abhd5</a>	<a href="#">Abhd5.bSep08</a>	<a href="#">316122</a>	20074	717	1	207	putative protein of ancient origin (Abhd5) alternative variant bSep08, mRNA.
<a href="#">Abhd6</a>	<a href="#">Abhd6.bSep08</a>	<a href="#">305795</a>	17835	354	3	63	putative protein of vertebrate origin (Abhd6) alternative variant bSep08, mRNA.
<a href="#">Abhd6</a>	<a href="#">Abhd6.cSep08</a>	<a href="#">305795</a>	4534	729	2	35	putative protein (Abhd6) alternative variant cSep08, mRNA.
<a href="#">Abhd7</a>	<a href="#">Abhd7.bSep08</a>	<a href="#">289440</a>	12871	725	3	137	putative protein of ancient origin (15.7 kD) (Abhd7) alternative variant bSep08, mRNA.
<a href="#">Abhd8</a>	<a href="#">Abhd8.bSep08</a>	<a href="#">306338</a>	1056	938	1	114	putative protein of metazoan origin (Abhd8) alternative variant bSep08, mRNA.
<a href="#">Abhd10</a>	<a href="#">Abhd10.bSep08</a>	<a href="#">303953</a>	5563	840	1	60	putative protein of mammalian origin (Abhd10) alternative variant bSep08, mRNA.
<a href="#">Abhd11</a>	<a href="#">Abhd11.aSep08</a>	<a href="#">360831</a>	2729	1159	6	321	PGAP1-like and alpha/beta hydrolase fold-1 (Abhd11) alternative variant aSep08, mRNA.
<a href="#">Abhd11</a>	<a href="#">Abhd11.bSep08</a>	<a href="#">360831</a>	1697	663	3	157	putative protein of ancient origin (Abhd11) alternative variant bSep08, mRNA.
<a href="#">Abhd11</a>	<a href="#">Abhd11.dSep08</a>	<a href="#">360831</a>	2784	994	5	127	putative protein of ancient origin (13.5 kD) (Abhd11) alternative variant dSep08, mRNA.
<a href="#">Abhd11</a>	<a href="#">Abhd11.eSep08</a>	<a href="#">360831</a>	2002	1421	3	39	putative protein (Abhd11) alternative variant eSep08, mRNA.
<a href="#">Abhd12</a>	<a href="#">Abhd12.bSep08</a>	<a href="#">499913</a>	55382	981	8	266	putative protein of ancient origin (29.7 kD) (Abhd12) alternative variant bSep08, mRNA.
<a href="#">Abhd12</a>	<a href="#">Abhd12.cSep08</a>	<a href="#">499913</a>	24114	1027	10	237	putative protein of ancient origin (26.9 kD) (Abhd12) alternative variant cSep08, mRNA.
<a href="#">Abhd13</a>	<a href="#">Abhd13.bSep08</a>	<a href="#">306630</a>	9926	375	1	93	putative protein (Abhd13) alternative variant bSep08, mRNA.
<a href="#">Abhd14a</a>	<a href="#">Abhd14a.aSep08</a>	<a href="#">300982</a>	4640	1186	5	303	putative protein of ancient origin (Abhd14a) alternative variant aSep08, mRNA.
<a href="#">Abhd14a</a>	<a href="#">Abhd14a.cSep08</a>	<a href="#">300982</a>	888	705	2	170	putative secreted or extracellular protein precursor of ancient origin (18.9 kD) (Abhd14a) alternative variant cSep08, mRNA.
<a href="#">Abhd14a</a>	<a href="#">Abhd14a.dSep08</a>	<a href="#">300982</a>	8261	910	5	166	putative protein of ancient origin (Abhd14a) alternative variant dSep08, mRNA.
<a href="#">Abhd14a</a>	<a href="#">Abhd14a.fSep08</a>	<a href="#">300982</a>	4032	974	4	111	putative protein of ancient origin (12.3 kD) (Abhd14a) alternative variant fSep08, mRNA.
<a href="#">Abhd14b</a>	<a href="#">Abhd14b.bSep08</a>	<a href="#">300983</a>	3058	639	4	186	putative protein of ancient origin (Abhd14b) alternative variant bSep08, mRNA.
<a href="#">Abhd14b</a>	<a href="#">Abhd14b.cSep08</a>	<a href="#">300983</a>	4035	1344	3	158	putative protein of ancient origin (Abhd14b) alternative variant cSep08, mRNA.
<a href="#">Abhd14b</a>	<a href="#">Abhd14b.eSep08</a>	<a href="#">300983</a>	4391	853	3	96	putative cytoplasmic protein of ancient origin (10.7 kD) (Abhd14b) alternative variant eSep08, mRNA.
<a href="#">Abi1</a>	<a href="#">Abi1.bSep08</a>	<a href="#">79249</a>	32763	594	5	197	abl-interactor 1 (Abi1) alternative variant bSep08, mRNA.

<a href="#">Abi2</a>	<a href="#">Abi2.aSep08</a>	<a href="#">286928</a>	217692	1836	10	469	abl-interactor 2 (Abi2) alternative variant aSep08, complete mRNA.
<a href="#">Abi2</a>	<a href="#">Abi2.bSep08</a>	<a href="#">286928</a>	120653	1801	4	436	abl-interactor 2 (Abi2) alternative variant bSep08, mRNA.
<a href="#">Abi2</a>	<a href="#">Abi2.dSep08</a>	<a href="#">286928</a>	18292	598	5	199	abl-interactor 2 (Abi2) alternative variant dSep08, mRNA.
<a href="#">Abi2</a>	<a href="#">Abi2.eSep08</a>	<a href="#">286928</a>	28101	2825	3	127	abl-interactor 2 (Abi2) alternative variant eSep08, mRNA.
<a href="#">Abi2</a>	<a href="#">Abi2.fSep08</a>	<a href="#">286928</a>	4481	445	2	96	abl-interactor 2 (Abi2) alternative variant fSep08, mRNA.
<a href="#">Abi3</a>	<a href="#">Abi3.bSep08</a>	<a href="#">303476</a>	1824	744	4	196	ABI gene family, member 3 (Abi3) alternative variant bSep08, mRNA.
<a href="#">Abi3</a>	<a href="#">Abi3.cSep08</a>	<a href="#">303476</a>	2508	783	4	152	ABI gene family, member 3 (Abi3) alternative variant cSep08, mRNA.
<a href="#">Abi3</a>	<a href="#">Abi3.dSep08</a>	<a href="#">303476</a>	2155	579	2	144	ABI gene family, member 3 (Abi3) alternative variant dSep08, mRNA.
<a href="#">Abl1</a>	<a href="#">Abl1.aSep08</a>	<a href="#">311860</a>	8353	3502	4	677	c-abl oncogene 1, receptor tyrosine kinase (Abl1) alternative variant aSep08, mRNA.
<a href="#">Abl1</a>	<a href="#">Abl1.bSep08</a>	<a href="#">311860</a>	24705	1310	7	330	c-abl oncogene 1, receptor tyrosine kinase (Abl1) alternative variant bSep08, mRNA.
<a href="#">Abl2</a>	<a href="#">Abl2.bSep08</a>	<a href="#">304883</a>	1207	812	1	82	v-abl Abelson murine leukemia viral oncogene homolog 2 (arg, Abelson-related gene) (Abl2) alternative variant bSep08, mRNA.
<a href="#">Ablim1</a>	<a href="#">Ablim1.bSep08</a>	<a href="#">307989</a>	10306	1252		236	actin-binding LIM protein 1 (Ablim1) alternative variant bSep08, mRNA.
<a href="#">Ablim3</a>	<a href="#">Ablim3.aSep08</a>	<a href="#">307395</a>	14819	2544		225	actin binding LIM protein family, member 3 (Ablim3) mRNA.
<a href="#">Abp1</a>	<a href="#">Abp1.bSep08</a>	<a href="#">65029</a>	1290	758	3	252	amiloride binding protein 1 (amine oxidase, copper-containing) (Abp1) alternative variant bSep08, mRNA.
<a href="#">Abp1</a>	<a href="#">Abp1.cSep08</a>	<a href="#">65029</a>	13291	841	3	53	amiloride binding protein 1 (amine oxidase, copper-containing) (6.1 kD) (Abp1) alternative variant cSep08, mRNA.
<a href="#">Abpa</a>	<a href="#">Abpa.aSep08</a>	<a href="#">361551</a>	1266	401	3	92	androgen binding protein, alpha (Abpa) alternative variant aSep08, mRNA.
<a href="#">Abr</a>	<a href="#">Abr.aSep08</a>	<a href="#">287537</a>	39644	3635	9	429	active BCR-related gene (Abr) alternative variant aSep08, mRNA.
<a href="#">Abr</a>	<a href="#">Abr.bSep08</a>	<a href="#">287537</a>	14712	511	4	169	active BCR-related gene (Abr) alternative variant bSep08, mRNA.
<a href="#">Abr</a>	<a href="#">Abr.cSep08</a>	<a href="#">287537</a>	1170	382	1	78	active BCR-related gene (Abr) alternative variant cSep08, mRNA.
<a href="#">Abt1</a>	<a href="#">Abt1.bSep08</a>	<a href="#">306960</a>	1314	311	2	103	activator of basal transcription 1 (Abt1) alternative variant bSep08, mRNA.
<a href="#">Abtb1</a>	<a href="#">Abtb1.bSep08</a>	<a href="#">297432</a>	628	509	2	132	putative protein of eukaryotic origin (Abtb1) alternative variant bSep08, mRNA.
<a href="#">Abtb1</a>	<a href="#">Abtb1.cSep08</a>	<a href="#">297432</a>	3739	734	7	95	putative protein of metazoan origin (Abtb1) alternative variant cSep08, mRNA.
<a href="#">Acaa1</a>	<a href="#">Acaa1.bSep08</a>	<a href="#">24157</a>	5491	555	6	184	acetyl-Coenzyme A acyltransferase 1 (Acaa1) alternative variant bSep08, mRNA.
<a href="#">Acaa1</a>	<a href="#">Acaa1.cSep08</a>	<a href="#">24157</a>	3680	696	5	163	acetyl-Coenzyme A acyltransferase 1 (Acaa1) alternative variant cSep08, mRNA.
<a href="#">Acaa1</a>	<a href="#">Acaa1.dSep08</a>	<a href="#">24157</a>	3300	605	6	158	acetyl-Coenzyme A acyltransferase 1 (Acaa1) alternative variant dSep08, mRNA.



<a href="#">Acaa1</a>	<a href="#">Acaa1.eSep08</a>	<a href="#">24157</a>	756	326	3	70	acetyl-Coenzyme A acyltransferase 1 (Acaa1) alternative variant eSep08, mRNA.
<a href="#">Acaa2</a>	<a href="#">Acaa2.bSep08</a>	<a href="#">170465</a>	45017	1281	10	397	acetyl-Coenzyme A acyltransferase 2 (41.9 kD) (Acaa2) alternative variant bSep08, mRNA.
<a href="#">Acaa2</a>	<a href="#">Acaa2.cSep08</a>	<a href="#">170465</a>	5105	511	3	114	acetyl-Coenzyme A acyltransferase 2 (Acaa2) alternative variant cSep08, mRNA.
<a href="#">Acaa2</a>	<a href="#">Acaa2.dSep08</a>	<a href="#">170465</a>	2486	392	2	22	acetyl-Coenzyme A acyltransferase 2 (Acaa2) alternative variant dSep08, mRNA.
<a href="#">Acacb</a>	<a href="#">Acacb.aSep08</a>	<a href="#">116719</a>	4467	976		261	acetyl-Coenzyme A carboxylase beta (Acacb) mRNA.
<a href="#">Acad9</a>	<a href="#">Acad9.bSep08</a>	<a href="#">294973</a>	1987	749	4	109	acyl-Coenzyme A dehydrogenase family, member 9 (Acad9) alternative variant bSep08, mRNA.
<a href="#">Acads</a>	<a href="#">Acads.bSep08</a>	<a href="#">64304</a>	862	532	3	90	acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (Acads) alternative variant bSep08, mRNA.
<a href="#">Acadvl</a>	<a href="#">Acadvl.bSep08</a>	<a href="#">25363</a>	1782	699	6	233	acyl-Coenzyme A dehydrogenase very long chain (Acadvl) alternative variant bSep08, mRNA.
<a href="#">Acadvl</a>	<a href="#">Acadvl.cSep08</a>	<a href="#">25363</a>	1326	748	6	169	acyl-Coenzyme A dehydrogenase very long chain (Acadvl) alternative variant cSep08, mRNA.
<a href="#">Acadvl</a>	<a href="#">Acadvl.dSep08</a>	<a href="#">25363</a>	1398	1229	3	105	acyl-Coenzyme A dehydrogenase very long chain (Acadvl) alternative variant dSep08, mRNA.
<a href="#">Acadvl</a>	<a href="#">Acadvl.eSep08</a>	<a href="#">25363</a>	708	534	3	82	acyl-Coenzyme A dehydrogenase very long chain (Acadvl) alternative variant eSep08, mRNA.
<a href="#">Acadvl</a>	<a href="#">Acadvl.fSep08</a>	<a href="#">25363</a>	721	625	2	46	acyl-Coenzyme A dehydrogenase very long chain (5.2 kD) (Acadvl) alternative variant fSep08, mRNA.
<a href="#">Acan</a>	<a href="#">Acan.bSep08</a>	<a href="#">58968</a>	15963	3978	6	1040	aggrecan (Acan) alternative variant bSep08, mRNA.
<a href="#">Acan</a>	<a href="#">Acan.cSep08</a>	<a href="#">58968</a>	17423	770	3	246	aggrecan (Acan) alternative variant cSep08, mRNA.
<a href="#">Acan</a>	<a href="#">Acan.dSep08</a>	<a href="#">58968</a>	19999	642	4	150	aggrecan (Acan) alternative variant dSep08, mRNA.
<a href="#">Acat1</a>	<a href="#">Acat1.bSep08</a>	<a href="#">25014</a>	12108	1457	5	120	acetyl-coenzyme A acetyltransferase 1 (Acat1) alternative variant bSep08, mRNA.
<a href="#">Acat1</a>	<a href="#">Acat1.cSep08</a>	<a href="#">25014</a>	1576	814	2	79	mitochondrial acetoacetyl-coa thiolase (8.2 kD) (Acat1) alternative variant cSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.bSep08</a>	<a href="#">308100</a>	34558	1379	8	368	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant bSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.cSep08</a>	<a href="#">308100</a>	27216	1000	7	326	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant cSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.dSep08</a>	<a href="#">308100</a>	13322	742	6	244	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant dSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.eSep08</a>	<a href="#">308100</a>	4658	510	2	169	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant eSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.fSep08</a>	<a href="#">308100</a>	6654	628	3	122	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant fSep08, mRNA.
<a href="#">Acat2</a>	<a href="#">Acat2.gSep08</a>	<a href="#">308100</a>	6650	513	2	90	acetyl-Coenzyme A acetyltransferase 2 (Acat2) alternative variant gSep08, mRNA.
<a href="#">Acbd4</a>	<a href="#">Acbd4.aSep08</a>	<a href="#">303577</a>	10469	1874	11	328	acyl-coA-binding protein, ACBP (37.5 kD) (Acbd4) alternative variant aSep08, mRNA.
<a href="#">Acbd4</a>	<a href="#">Acbd4.cSep08</a>	<a href="#">303577</a>	7911	830	8	276	acyl-coA-binding protein, ACBP (Acbd4) alternative variant cSep08, mRNA.

<a href="#">Acbd4</a>	<a href="#">Acbd4.eSep08</a>	<a href="#">303577</a>	2742	412	3	99	putative protein of mammalian origin (Acbd4) alternative variant eSep08, mRNA.
<a href="#">Acbd4</a>	<a href="#">Acbd4.fSep08</a>	<a href="#">303577</a>	1768	545	6	77	putative protein (8.1 kD) (Acbd4) alternative variant fSep08, mRNA.
<a href="#">Acbd4</a>	<a href="#">Acbd4.gSep08</a>	<a href="#">303577</a>	1395	517	4	49	putative protein (Acbd4) alternative variant gSep08, mRNA.
<a href="#">Acbd4</a>	<a href="#">Acbd4.hSep08</a>	<a href="#">303577</a>	1230	779	3	50	putative protein of vertebrate origin (Acbd4) alternative variant hSep08, mRNA.
<a href="#">Acbd5</a>	<a href="#">Acbd5.bSep08</a>	<a href="#">307170</a>	29590	3012	9	352	putative protein, with a coiled coil domain, of metazoan origin (Acbd5) alternative variant bSep08, mRNA.
<a href="#">Acbd6</a>	<a href="#">Acbd6.bSep08</a>	<a href="#">289125</a>	14533	268	1	82	ankyrin (Acbd6) alternative variant bSep08, mRNA.
<a href="#">Accn2</a>	<a href="#">Accn2.bSep08</a>	<a href="#">79123</a>	2996	1137	7	264	amiloride-sensitive cation channel 2 neuronal CRA c (Accn2) alternative variant bSep08, mRNA.
<a href="#">Accn2</a>	<a href="#">Accn2.cSep08</a>	<a href="#">79123</a>	23959	655	6	170	amiloride-sensitive cation channel 2 neuronal CRA b (Accn2) alternative variant cSep08, mRNA.
<a href="#">Accn2</a>	<a href="#">Accn2.dSep08</a>	<a href="#">79123</a>	1407	521	2	99	putative protein (Accn2) alternative variant dSep08, mRNA.
<a href="#">Accn3</a>	<a href="#">Accn3.bSep08</a>	<a href="#">286920</a>	915	505	1	138	amiloride-sensitive cation channel 3 (Accn3) alternative variant bSep08, mRNA.
<a href="#">Accn4</a>	<a href="#">Accn4.bSep08</a>	<a href="#">63882</a>	741	413	1	93	amiloride-sensitive cation channel 4, pituitary (10.3 kD) (Accn4) alternative variant bSep08, mRNA.
<a href="#">Accs</a>	<a href="#">Accs.aSep08</a>	<a href="#">311218</a>	7402	1794	3	543	1-aminocyclopropane-1-carboxylate synthase homolog (Arabidopsis)(non-functional) (Accs) alternative variant aSep08, mRNA.
<a href="#">Accs</a>	<a href="#">Accs.bSep08</a>	<a href="#">311218</a>	13900	1908	10	523	1-aminocyclopropane-1-carboxylate synthase homolog (Arabidopsis)(non-functional) (Accs) alternative variant bSep08, mRNA.
<a href="#">Accs</a>	<a href="#">Accs.cSep08</a>	<a href="#">311218</a>	3480	403	1	87	1-aminocyclopropane-1-carboxylate synthase homolog (Arabidopsis)(non-functional) (Accs) alternative variant cSep08, mRNA.
<a href="#">Acd</a>	<a href="#">Acd.bSep08</a>	<a href="#">307798</a>	2727	1378	9	183	adrenocortical dysplasia (19.9 kD) (Acd) alternative variant bSep08, mRNA.
<a href="#">Acd</a>	<a href="#">Acd.cSep08</a>	<a href="#">307798</a>	1772	1465	4	164	adrenocortical dysplasia (18.1 kD) (Acd) alternative variant cSep08, mRNA.
<a href="#">Acd</a>	<a href="#">Acd.dSep08</a>	<a href="#">307798</a>	1811	1493	4	161	adrenocortical dysplasia (17.4 kD) (Acd) alternative variant dSep08, mRNA.
<a href="#">Acd</a>	<a href="#">Acd.eSep08</a>	<a href="#">307798</a>	2739	1173	6	135	adrenocortical dysplasia (14.8 kD) (Acd) alternative variant eSep08, mRNA.
<a href="#">Ace</a>	<a href="#">Ace.bSep08</a>	<a href="#">24310</a>	4279	816	5	271	angiotensin I converting enzyme (peptidyl-dipeptidase A) 1 (Ace) alternative variant bSep08, mRNA.
<a href="#">Ace2</a>	<a href="#">Ace2.bSep08</a>	<a href="#">302668</a>	2902	1053	1	82	angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 (Ace2) alternative variant bSep08, mRNA.
<a href="#">Ace3</a>	<a href="#">Ace3.aSep08</a>	<a href="#">498012</a>	7972	1039		217	angiotensin I converting enzyme (peptidyl-dipeptidase A) 3 (Ace3) mRNA.
<a href="#">Acin1</a>	<a href="#">Acin1.aSep08</a>	<a href="#">305884</a>	12088	2422	12	565	apoptotic chromatin condensation inducer 1 (65.3 kD) (Acin1) alternative variant aSep08, mRNA.
<a href="#">Acin1</a>	<a href="#">Acin1.bSep08</a>	<a href="#">305884</a>	7875	1540	9	237	apoptotic chromatin condensation inducer 1 (Acin1) alternative variant bSep08, mRNA.

<a href="#">Acin1</a>	<a href="#">Acin1.cSep08</a>	<a href="#">305884</a>	4016	1621	4	154	apoptotic chromatin condensation inducer 1 (Acin1) alternative variant cSep08, mRNA.
<a href="#">Acin1</a>	<a href="#">Acin1.fSep08</a>	<a href="#">305884</a>	469	370	2	93	apoptotic chromatin condensation inducer 1 (Acin1) alternative variant fSep08, mRNA.
<a href="#">Acin1</a>	<a href="#">Acin1.gSep08</a>	<a href="#">305884</a>	1149	501	2	82	apoptotic chromatin condensation inducer 1 (Acin1) alternative variant gSep08, mRNA.
<a href="#">Acly</a>	<a href="#">Acly.cSep08</a>	<a href="#">24159</a>	5605	607	5	169	ATP citrate lyase (Acly) alternative variant cSep08, mRNA.
<a href="#">Acly</a>	<a href="#">Acly.dSep08</a>	<a href="#">24159</a>	1342	412	3	117	ATP citrate lyase (Acly) alternative variant dSep08, mRNA.
<a href="#">Acn9</a>	<a href="#">Acn9.aSep08</a>	<a href="#">362323</a>	65028	695	2	123	ACN9 homolog (S. cerevisiae) (Acn9) alternative variant aSep08, mRNA.
<a href="#">Acn9</a>	<a href="#">Acn9.cSep08</a>	<a href="#">362323</a>	41029	441	2	58	ACN9 homolog (S. cerevisiae) (6.9 kD) (Acn9) alternative variant cSep08, complete mRNA.
<a href="#">Aco1</a>	<a href="#">Aco1.bSep08</a>	<a href="#">50655</a>	9447	411	3	67	aconitase 1 (Aco1) alternative variant bSep08, mRNA.
<a href="#">Aco2</a>	<a href="#">Aco2.bSep08</a>	<a href="#">79250</a>	31752	755	5	251	aconitase 2, mitochondrial (Aco2) alternative variant bSep08, mRNA.
<a href="#">Aco2</a>	<a href="#">Aco2.cSep08</a>	<a href="#">79250</a>	17662	948	2	58	aconitase 2, mitochondrial (6.8 kD) (Aco2) alternative variant cSep08, mRNA.
<a href="#">Aco2</a>	<a href="#">Aco2.dSep08</a>	<a href="#">79250</a>	42612	417	3	48	aconitase 2, mitochondrial (5.5 kD) (Aco2) alternative variant dSep08, mRNA.
<a href="#">Acot3</a>	<a href="#">Acot3.bSep08</a>	<a href="#">314304</a>	4845	1031	3	336	acyl-CoA thioesterase 3 (Acot3) alternative variant bSep08, mRNA.
<a href="#">Acot7</a>	<a href="#">Acot7.bSep08</a>	<a href="#">26759</a>	35070	706	4	182	acyl-CoA thioesterase 7 (Acot7) alternative variant bSep08, mRNA.
<a href="#">Acot7</a>	<a href="#">Acot7.cSep08</a>	<a href="#">26759</a>	30849	371	1	123	acyl-CoA thioesterase 7 (Acot7) alternative variant cSep08, mRNA.
<a href="#">Acot8</a>	<a href="#">Acot8.bSep08</a>	<a href="#">170588</a>	11270	726	4	200	acyl-CoA thioesterase 8 (Acot8) alternative variant bSep08, mRNA.
<a href="#">Acot8</a>	<a href="#">Acot8.cSep08</a>	<a href="#">170588</a>	4604	743	4	113	acyl-CoA thioesterase 8 (Acot8) alternative variant cSep08, mRNA.
<a href="#">Acot8</a>	<a href="#">Acot8.dSep08</a>	<a href="#">170588</a>	8612	502	3	106	acyl-CoA thioesterase 8 (Acot8) alternative variant dSep08, mRNA.
<a href="#">Acot8</a>	<a href="#">Acot8.fSep08</a>	<a href="#">170588</a>	3067	695	3	86	acyl-CoA thioesterase 8 (Acot8) alternative variant fSep08, mRNA.
<a href="#">Acot12</a>	<a href="#">Acot12.bSep08</a>	<a href="#">170570</a>	1552	336	1	80	acyl-CoA thioesterase 12 (Acot12) alternative variant bSep08, mRNA.
<a href="#">Acox1</a>	<a href="#">Acox1.aSep08</a>	<a href="#">50681</a>	8246	2970	9	424	acyl-Coenzyme A oxidase 1, palmitoyl (Acox1) alternative variant aSep08, mRNA.
<a href="#">Acox2</a>	<a href="#">Acox2.bSep08</a>	<a href="#">252898</a>	3353	578	1	132	acyl-Coenzyme A oxidase 2, branched chain (Acox2) alternative variant bSep08, mRNA.
<a href="#">Acox3</a>	<a href="#">Acox3.bSep08</a>	<a href="#">83522</a>	45377	549	4	114	acyl-Coenzyme A oxidase 3, pristanoyl (Acox3) alternative variant bSep08, mRNA.
<a href="#">Acp2</a>	<a href="#">Acp2.bSep08</a>	<a href="#">24162</a>	9066	2669	11	215	acid phosphatase 2, lysosomal (24.9 kD) (Acp2) alternative variant bSep08, complete mRNA.
<a href="#">Acpp</a>	<a href="#">Acpp.aSep08</a>	<a href="#">56780</a>	45917	1295	1	430	acid phosphatase, prostate (Acpp) alternative variant aSep08, mRNA.
<a href="#">Acpt</a>	<a href="#">Acpt.bSep08</a>	<a href="#">308569</a>	4215	686	2	133	acid phosphatase, testicular (Acpt) alternative variant bSep08, mRNA.

<a href="#">Acpt</a>	<a href="#">Acpt.cSep08</a>	<a href="#">308569</a>	4859	737	3	114	acid phosphatase, testicular (Acpt) alternative variant cSep08, mRNA.
<a href="#">Acpt</a>	<a href="#">Acpt.dSep08</a>	<a href="#">308569</a>	4215	761	2	88	acid phosphatase, testicular (Acpt) alternative variant dSep08, mRNA.
<a href="#">Acr</a>	<a href="#">Acr.bSep08</a>	<a href="#">24163</a>	5908	1254	1	381	acrosin (42.7 kD) (Acr) alternative variant bSep08, mRNA.
<a href="#">Acsbg1</a>	<a href="#">Acsbg1.bSep08</a>	<a href="#">171410</a>	6947	614	4	204	acyl-CoA synthetase bubblegum family member 1 (Acsbg1) alternative variant bSep08, mRNA.
<a href="#">Acsbg1</a>	<a href="#">Acsbg1.cSep08</a>	<a href="#">171410</a>	6747	799	3	86	acyl-CoA synthetase bubblegum family member 1 (Acsbg1) alternative variant cSep08, mRNA.
<a href="#">Acsf2</a>	<a href="#">Acsf2.bSep08</a>	<a href="#">619561</a>	11147	729	7	243	acyl-CoA synthetase family member 2 (Acsf2) alternative variant bSep08, mRNA.
<a href="#">Acsf2</a>	<a href="#">Acsf2.cSep08</a>	<a href="#">619561</a>	5724	609	3	203	acyl-CoA synthetase family member 2 (Acsf2) alternative variant cSep08, mRNA.
<a href="#">Acsf3</a>	<a href="#">Acsf3.aSep08</a>	<a href="#">498962</a>	40393	2282	7	629	acyl-CoA synthetase family member 3 (Acsf3) alternative variant aSep08, mRNA.
<a href="#">Acsf3</a>	<a href="#">Acsf3.bSep08</a>	<a href="#">498962</a>	8422	427	3	142	acyl-CoA synthetase family member 3 (Acsf3) alternative variant bSep08, mRNA.
<a href="#">Acsf3</a>	<a href="#">Acsf3.cSep08</a>	<a href="#">498962</a>	8956	344	1	25	acyl-CoA synthetase family member 3 (Acsf3) alternative variant cSep08, mRNA.
<a href="#">Acsl1</a>	<a href="#">Acsl1.bSep08</a>	<a href="#">25288</a>	41288	1500	5	208	acyl-CoA synthetase long-chain family member 1 (24.0 kD) (Acsl1) alternative variant bSep08, mRNA.
<a href="#">Acsl3</a>	<a href="#">Acsl3.bSep08</a>	<a href="#">114024</a>	4744	495	1	151	acyl-CoA synthetase long-chain family member 3 (Acsl3) alternative variant bSep08, mRNA.
<a href="#">Acsl4</a>	<a href="#">Acsl4.bSep08</a>	<a href="#">113976</a>	35957	622	3	162	acyl-CoA synthetase long-chain family member 4 (Acsl4) alternative variant bSep08, mRNA.
<a href="#">Acsl4</a>	<a href="#">Acsl4.cSep08</a>	<a href="#">113976</a>	33856	398	1	76	acyl-CoA synthetase long-chain family member 4 (Acsl4) alternative variant cSep08, mRNA.
<a href="#">Acsl5</a>	<a href="#">Acsl5.aSep08</a>	<a href="#">94340</a>	11065	1387	10	368	acyl-CoA synthetase long-chain family member 5 (Acsl5) alternative variant aSep08, mRNA.
<a href="#">Acsl5</a>	<a href="#">Acsl5.bSep08</a>	<a href="#">94340</a>	2085	748	2	78	acyl-CoA synthetase long-chain family member 5 (8.9 kD) (Acsl5) alternative variant bSep08, mRNA.
<a href="#">Acsl5</a>	<a href="#">Acsl5.cSep08</a>	<a href="#">94340</a>	1458	594	1	10	acyl-CoA synthetase long-chain family member 5 (1.1 kD) (Acsl5) alternative variant cSep08, mRNA.
<a href="#">Acsl6</a>	<a href="#">Acsl6.bSep08</a>	<a href="#">117243</a>	30872	1350	10	383	acyl-CoA synthetase long-chain family member 6 (Acsl6) alternative variant bSep08, mRNA.
<a href="#">Acsl6</a>	<a href="#">Acsl6.cSep08</a>	<a href="#">117243</a>	5323	440	5	146	acyl-CoA synthetase long-chain family member 6 (Acsl6) alternative variant cSep08, mRNA.
<a href="#">Acsl6</a>	<a href="#">Acsl6.dSep08</a>	<a href="#">117243</a>	5628	722	2	108	acyl-CoA synthetase long-chain family member 6 (Acsl6) alternative variant dSep08, mRNA.
<a href="#">Acsl6</a>	<a href="#">Acsl6.eSep08</a>	<a href="#">117243</a>	19613	480	4	87	acyl-CoA synthetase long-chain family member 6 (Acsl6) alternative variant eSep08, mRNA.
<a href="#">Acsl6</a>	<a href="#">Acsl6.fSep08</a>	<a href="#">117243</a>	9562	1244	2	71	acyl-CoA synthetase long-chain family member 6 (Acsl6) alternative variant fSep08, mRNA.
<a href="#">Acsm2</a>	<a href="#">Acsm2.bSep08</a>	<a href="#">246263</a>	15462	801	5	246	acyl-CoA synthetase medium-chain family member 2 (Acsm2) alternative variant bSep08, mRNA.
<a href="#">Acsm2</a>	<a href="#">Acsm2.cSep08</a>	<a href="#">246263</a>	5618	765	4	114	acyl-CoA synthetase medium-chain family member 2 (12.8 kD) (Acsm2) alternative variant cSep08, mRNA.

<a href="#">Acsm2</a>	<a href="#">Acsm2.dSep08</a>	<a href="#">246263</a>	1327	305	3	67	acyl-CoA synthetase medium-chain family member 2 (Acsm2) alternative variant dSep08, mRNA.
<a href="#">Acsm3</a>	<a href="#">Acsm3.bSep08</a>	<a href="#">24763</a>	994	649		38	acyl-CoA synthetase medium-chain family member 3 (Acsm3) alternative variant bSep08, mRNA.
<a href="#">Acss1</a>	<a href="#">Acss1.cSep08</a>	<a href="#">296259</a>	8725	288	2	88	acyl-CoA synthetase short-chain family member 1 (Acss1) alternative variant cSep08, mRNA.
<a href="#">Acss2</a>	<a href="#">Acss2.bSep08</a>	<a href="#">311569</a>	6385	1286	1	239	acyl-CoA synthetase short-chain family member 2 (Acss2) alternative variant bSep08, mRNA.
<a href="#">Acta1</a>	<a href="#">Acta1.bSep08</a>	<a href="#">29437</a>	667	242		80	actin, alpha 1, skeletal muscle (Acta1) alternative variant bSep08, mRNA.
<a href="#">Actb</a>	<a href="#">Actb.bSep08</a>	<a href="#">81822</a>	2543	1900	4	336	beta actin (37.2 kD) (Actb) alternative variant bSep08, mRNA.
<a href="#">Actb</a>	<a href="#">Actb.cSep08</a>	<a href="#">81822</a>	937	847	2	205	beta-actin (Actb) alternative variant cSep08, mRNA.
<a href="#">Actb</a>	<a href="#">Actb.eSep08</a>	<a href="#">81822</a>	1239	749	4	125	actin beta (Actb) alternative variant eSep08, mRNA.
<a href="#">Actc1</a>	<a href="#">Actc1.bSep08</a>	<a href="#">29275</a>	5829	1383	6	328	actin, alpha, cardiac muscle 1 (36.5 kD) (Actc1) alternative variant bSep08, complete mRNA.
<a href="#">Actg1</a>	<a href="#">Actg1.aSep08</a>	<a href="#">287876</a>	2866	1923	6	404	actin, gamma 1 (Actg1) alternative variant aSep08, mRNA.
<a href="#">Actg1</a>	<a href="#">Actg1.bSep08</a>	<a href="#">287876</a>	2443	1713	6	375	actin, gamma 1 (41.8 kD) (Actg1) alternative variant bSep08, mRNA.
<a href="#">Actg1</a>	<a href="#">Actg1.dSep08</a>	<a href="#">287876</a>	1729	1179	4	132	actin, gamma 1 (14.6 kD) (Actg1) alternative variant dSep08, mRNA.
<a href="#">Actl6a</a>	<a href="#">Actl6a.bSep08</a>	<a href="#">361925</a>	5141	742	7	246	actin-like 6A (Actl6a) alternative variant bSep08, mRNA.
<a href="#">Actl6a</a>	<a href="#">Actl6a.cSep08</a>	<a href="#">361925</a>	8152	595	6	198	actin-like 6A (Actl6a) alternative variant cSep08, mRNA.
<a href="#">Actl6a</a>	<a href="#">Actl6a.dSep08</a>	<a href="#">361925</a>	8252	736	7	159	actin-like 6A (Actl6a) alternative variant dSep08, mRNA.
<a href="#">Actl6b</a>	<a href="#">Actl6b.cSep08</a>	<a href="#">288563</a>	1398	664	2	81	actin-like 6B (Actl6b) alternative variant cSep08, mRNA.
<a href="#">Actn1</a>	<a href="#">Actn1.bSep08</a>	<a href="#">81634</a>	4610	841	5	200	actinin, alpha 1 (Actn1) alternative variant bSep08, mRNA.
<a href="#">Actn1</a>	<a href="#">Actn1.cSep08</a>	<a href="#">81634</a>	2859	1635	2	82	actinin, alpha 1 (Actn1) alternative variant cSep08, mRNA.
<a href="#">Actn2</a>	<a href="#">Actn2.aSep08</a>	<a href="#">291245</a>	68109	2950	21	959	actinin alpha 2 (Actn2) alternative variant aSep08, mRNA.
<a href="#">Actn2</a>	<a href="#">Actn2.bSep08</a>	<a href="#">291245</a>	5922	773	6	241	actinin alpha 2 (Actn2) alternative variant bSep08, mRNA.
<a href="#">Actn4</a>	<a href="#">Actn4.bSep08</a>	<a href="#">63836</a>	2579	823	5	243	actinin alpha 4 (Actn4) alternative variant bSep08, mRNA.
<a href="#">Actn4</a>	<a href="#">Actn4.cSep08</a>	<a href="#">63836</a>	1507	917	2	176	actinin alpha 4 (18.9 kD) (Actn4) alternative variant cSep08, mRNA.
<a href="#">Actn4</a>	<a href="#">Actn4.dSep08</a>	<a href="#">63836</a>	8164	507	5	169	actinin alpha 4 (Actn4) alternative variant dSep08, mRNA.
<a href="#">Actn4</a>	<a href="#">Actn4.eSep08</a>	<a href="#">63836</a>	964	406	2	135	actinin alpha 4 (Actn4) alternative variant eSep08, mRNA.
<a href="#">Actr1a</a>	<a href="#">Actr1a.bSep08</a>	<a href="#">294010</a>	17099	1399	10	365	ARP1 actin-related protein 1 homolog A (yeast) (Actr1a) alternative variant bSep08, mRNA.
<a href="#">Actr1a</a>	<a href="#">Actr1a.cSep08</a>	<a href="#">294010</a>	2603	600	4	62	ARP1 actin-related protein 1 homolog A (yeast) (Actr1a) alternative variant cSep08, mRNA.
<a href="#">Actr1b</a>	<a href="#">Actr1b.bSep08</a>	<a href="#">316333</a>	11403	739	6	169	ARP1 actin-related protein 1 homolog B (yeast) (Actr1b) alternative variant bSep08, mRNA.
<a href="#">Actr1b</a>	<a href="#">Actr1b.cSep08</a>	<a href="#">316333</a>	1104	754	3	167	ARP1 actin-related protein 1 homolog B (yeast) (18.6 kD) (Actr1b) alternative variant cSep08, mRNA.
<a href="#">Actr1b</a>	<a href="#">Actr1b.dSep08</a>	<a href="#">316333</a>	11875	750	6	142	ARP1 actin-related protein 1 homolog B (yeast) (Actr1b) alternative variant dSep08, mRNA.

<a href="#">Actr1b</a>	<a href="#">Actr1b.eSep08</a>	<a href="#">316333</a>	4301	580	5	141	ARP1 actin-related protein 1 homolog B (yeast) (Actr1b) alternative variant eSep08, mRNA.
<a href="#">Actr2</a>	<a href="#">Actr2.bSep08</a>	<a href="#">289820</a>	6414	1679	2	86	ARP2 actin-related protein 2 homolog (yeast) (13.5 kD) (Actr2) alternative variant bSep08, mRNA.
<a href="#">Actr3</a>	<a href="#">Actr3.bSep08</a>	<a href="#">81732</a>	18772	424	4	141	ARP3 actin-related protein 3 homolog (yeast) (Actr3) alternative variant bSep08, mRNA.
<a href="#">Actr3b</a>	<a href="#">Actr3b.aSep08</a>	<a href="#">362298</a>	93498	1849	6	356	ARP3 actin-related protein 3 homolog B (yeast) (Actr3b) alternative variant aSep08, mRNA.
<a href="#">Actr3b</a>	<a href="#">Actr3b.bSep08</a>	<a href="#">362298</a>	65189	524	2	174	ARP3 actin-related protein 3 homolog B (yeast) (Actr3b) alternative variant bSep08, mRNA.
<a href="#">Actr5</a>	<a href="#">Actr5.aSep08</a>	<a href="#">362258</a>	8076	1406	5	468	ARP5 actin-related protein 5 homolog (yeast) (Actr5) alternative variant aSep08, mRNA.
<a href="#">Actr5</a>	<a href="#">Actr5.cSep08</a>	<a href="#">362258</a>	4009	1047	4	166	ARP5 actin-related protein 5 homolog (yeast) (Actr5) alternative variant cSep08, mRNA.
<a href="#">Actr8</a>	<a href="#">Actr8.aSep08</a>	<a href="#">361107</a>	15369	2876	13	624	ARP8 actin-related protein 8 homolog (S. cerevisiae) (70.5 kD) (Actr8) alternative variant aSep08, complete mRNA.
<a href="#">Actr8</a>	<a href="#">Actr8.bSep08</a>	<a href="#">361107</a>	2276	1006	3	83	ARP8 actin-related protein 8 homolog (S. cerevisiae) (Actr8) alternative variant bSep08, mRNA.
<a href="#">Actr8</a>	<a href="#">Actr8.cSep08</a>	<a href="#">361107</a>	792	533	2	78	ARP8 actin-related protein 8 homolog (S. cerevisiae) (Actr8) alternative variant cSep08, mRNA.
<a href="#">Actr8</a>	<a href="#">Actr8.dSep08</a>	<a href="#">361107</a>	2615	543	3	62	ARP8 actin-related protein 8 homolog (S. cerevisiae) (Actr8) alternative variant dSep08, mRNA.
<a href="#">Actr10</a>	<a href="#">Actr10.bSep08</a>	<a href="#">299121</a>	26871	2817	5	290	actin-related protein 10 homolog (S. cerevisiae) (32.2 kD) (Actr10) alternative variant bSep08, complete mRNA.
<a href="#">Actr10</a>	<a href="#">Actr10.cSep08</a>	<a href="#">299121</a>	20236	715	1	236	actin-related protein 10 homolog (S. cerevisiae) (Actr10) alternative variant cSep08, mRNA.
<a href="#">Acvr1</a>	<a href="#">Acvr1.bSep08</a>	<a href="#">79558</a>	14806	792	4	232	activin A receptor, type 1 (Acvr1) alternative variant bSep08, mRNA.
<a href="#">Acvr2b</a>	<a href="#">Acvr2b.bSep08</a>	<a href="#">25366</a>	2613	711	1	236	activin receptor IIB (Acvr2b) alternative variant bSep08, mRNA.
<a href="#">Acvr11</a>	<a href="#">Acvr11.bSep08</a>	<a href="#">25237</a>	7768	826	5	184	activin A receptor type II-like 1 (Acvr11) alternative variant bSep08, mRNA.
<a href="#">Acvr11</a>	<a href="#">Acvr11.cSep08</a>	<a href="#">25237</a>	1440	744	2	128	activin A receptor type II-like 1 (Acvr11) alternative variant cSep08, mRNA.
<a href="#">Acvr11</a>	<a href="#">Acvr11.dSep08</a>	<a href="#">25237</a>	7159	728	3	79	activin A receptor type II-like 1 (Acvr11) alternative variant dSep08, mRNA.
<a href="#">Acy1</a>	<a href="#">Acy1.bSep08</a>	<a href="#">300981</a>	4252	1644	12	313	aminoacylase 1 (35.4 kD) (Acy1) alternative variant bSep08, mRNA.
<a href="#">Acy1</a>	<a href="#">Acy1.cSep08</a>	<a href="#">300981</a>	495	409	2	105	aminoacylase 1 (Acy1) alternative variant cSep08, mRNA.
<a href="#">Acy1</a>	<a href="#">Acy1.eSep08</a>	<a href="#">300981</a>	1800	437	4	95	aminoacylase 1 (10.4 kD) (Acy1) alternative variant eSep08, mRNA.
<a href="#">Acy3</a>	<a href="#">Acy3.bSep08</a>	<a href="#">293653</a>	2199	784	2	261	aspartoacylase (aminoacylase) 3 (Acy3) alternative variant bSep08, mRNA.
<a href="#">Acyp1</a>	<a href="#">Acyp1.aSep08</a>	<a href="#">299203</a>	6945	587	2	129	acylphosphatase 1, erythrocyte (common) type (Acyp1) alternative variant aSep08, mRNA.
<a href="#">Acyp1</a>	<a href="#">Acyp1.bSep08</a>	<a href="#">299203</a>	6885	606	2	99	acylphosphatase 1, erythrocyte (common) type (11.3 kD) (Acyp1) alternative variant bSep08, mRNA.

<a href="#">Ada</a>	<a href="#">Ada.bSep08</a>	<a href="#">24165</a>	1002	388	1	64	adenosine deaminase (Ada) alternative variant bSep08, mRNA.
<a href="#">Adam2</a>	<a href="#">Adam2.bSep08</a>	<a href="#">56806</a>	6515	900	3	245	a disintegrin metalloprotease domain 2 (Adam2) alternative variant bSep08, mRNA.
<a href="#">Adam2</a>	<a href="#">Adam2.cSep08</a>	<a href="#">56806</a>	16246	747	9	124	fertilin beta (Adam2) alternative variant cSep08, mRNA.
<a href="#">Adam3</a>	<a href="#">Adam3.bSep08</a>	<a href="#">57021</a>	7496	266	4	74	a disintegrin and metalloproteinase domain 3 (cyritestin) (Adam3) alternative variant bSep08, mRNA.
<a href="#">Adam3</a>	<a href="#">Adam3.cSep08</a>	<a href="#">57021</a>	7086	391	4	64	a disintegrin and metalloproteinase domain 3 (cyritestin) (7.1 kD) (Adam3) alternative variant cSep08, mRNA.
<a href="#">Adam5</a>	<a href="#">Adam5.bSep08</a>	<a href="#">498654</a>	8464	751	9	232	a disintegrin and metalloproteinase domain 5 (Adam5) alternative variant bSep08, mRNA.
<a href="#">Adam5</a>	<a href="#">Adam5.cSep08</a>	<a href="#">498654</a>	19089	688	6	229	a disintegrin and metalloproteinase domain 5 (Adam5) alternative variant cSep08, mRNA.
<a href="#">Adam5</a>	<a href="#">Adam5.dSep08</a>	<a href="#">498654</a>	12189	317	3	44	a disintegrin and metalloproteinase domain 5 (Adam5) alternative variant dSep08, mRNA.
<a href="#">Adam9</a>	<a href="#">Adam9.bSep08</a>	<a href="#">290834</a>	35911	481	2	55	a disintegrin and metalloproteinase domain 9 (meltrin gamma) (Adam9) alternative variant bSep08, mRNA.
<a href="#">Adam10</a>	<a href="#">Adam10.aSep08</a>	<a href="#">29650</a>	34125	1374	5	457	a disintegrin and metalloproteinase domain 10 (Adam10) alternative variant aSep08, mRNA.
<a href="#">Adam10</a>	<a href="#">Adam10.bSep08</a>	<a href="#">29650</a>	20858	591	1	49	a disintegrin and metalloproteinase domain 10 (5.8 kD) (Adam10) alternative variant bSep08, mRNA.
<a href="#">Adam11</a>	<a href="#">Adam11.bSep08</a>	<a href="#">360638</a>	1035	347	5	70	a disintegrin and metalloproteinase domain 11 (Adam11) alternative variant bSep08, mRNA.
<a href="#">Adam12</a>	<a href="#">Adam12.aSep08</a>	<a href="#">679837</a>	91722	2928		699	a disintegrin and metalloproteinase domain 12 (meltrin alpha) (77.6 kD) (Adam12) mRNA.
<a href="#">Adam17</a>	<a href="#">Adam17.bSep08</a>	<a href="#">57027</a>	4448	852	4	202	a disintegrin and metalloproteinase domain 17 (Adam17) alternative variant bSep08, mRNA.
<a href="#">Adam17</a>	<a href="#">Adam17.cSep08</a>	<a href="#">57027</a>	1174	774	1	113	a disintegrin and metalloproteinase domain 17 (3.6 kD) (Adam17) alternative variant cSep08, mRNA.
<a href="#">Adam18</a>	<a href="#">Adam18.aSep08</a>	<a href="#">57029</a>	15673	279		42	a disintegrin and metalloproteinase domain 18 (4.8 kD) (Adam18) mRNA.
<a href="#">Adam19</a>	<a href="#">Adam19.aSep08</a>	<a href="#">303068</a>	68212	1776		561	a disintegrin and metalloproteinase domain 19 (meltrin beta) (Adam19) alternative variant aSep08, mRNA.
<a href="#">Adam19</a>	<a href="#">Adam19.bSep08</a>	<a href="#">303068</a>	45740	492		163	a disintegrin and metalloproteinase domain 19 (meltrin beta) (Adam19) alternative variant bSep08, mRNA.
<a href="#">Adam23</a>	<a href="#">Adam23.bSep08</a>	<a href="#">301460</a>	3105	502	2	133	a disintegrin and metalloproteinase domain 23 (Adam23) alternative variant bSep08, mRNA.
<a href="#">Adam23</a>	<a href="#">Adam23.cSep08</a>	<a href="#">301460</a>	10852	3586	2	57	a disintegrin and metalloproteinase domain 23 (Adam23) alternative variant cSep08, mRNA.
<a href="#">Adam32</a>	<a href="#">Adam32.aSep08</a>	<a href="#">361170</a>	66006	1480	9	439	a disintegrin and metalloproteinase domain 32 (Adam32) alternative variant aSep08, mRNA.
<a href="#">Adam32</a>	<a href="#">Adam32.bSep08</a>	<a href="#">361170</a>	27139	769	9	256	a disintegrin and metalloproteinase domain 32 (Adam32) alternative variant bSep08, mRNA.
<a href="#">Adam32</a>	<a href="#">Adam32.cSep08</a>	<a href="#">361170</a>	27162	727	8	241	a disintegrin and metalloproteinase domain 32 (Adam32) alternative variant cSep08, mRNA.
<a href="#">Adam32</a>	<a href="#">Adam32.dSep08</a>	<a href="#">361170</a>	32091	745	1	204	a disintegrin and metalloproteinase domain 32 (Adam32) alternative variant dSep08, mRNA.

<a href="#">Adam32</a>	<a href="#">Adam32.eSep08</a>	<a href="#">361170</a>	10204	604	1	132	a disintegrin and metallopeptidase domain 32 (Adam32) alternative variant eSep08, mRNA.
<a href="#">Adam33</a>	<a href="#">Adam33.bSep08</a>	<a href="#">311425</a>	1810	754	7	171	a disintegrin and metallopeptidase domain 33 (Adam33) alternative variant bSep08, mRNA.
<a href="#">Adam33</a>	<a href="#">Adam33.cSep08</a>	<a href="#">311425</a>	2057	471	4	79	a disintegrin and metallopeptidase domain 33 (Adam33) alternative variant cSep08, mRNA.
<a href="#">Adam33</a>	<a href="#">Adam33.dSep08</a>	<a href="#">311425</a>	2027	657	3	69	a disintegrin and metallopeptidase domain 33 (Adam33) alternative variant dSep08, mRNA.
<a href="#">Adamdec1</a>	<a href="#">Adamdec1.bSep08</a>	<a href="#">290338</a>	5722	432	1	144	ADAM-like, decysin 1 (Adamdec1) alternative variant bSep08, mRNA.
<a href="#">Adamts2</a>	<a href="#">Adamts2.aSep08</a>	<a href="#">287899</a>	11780	1278	6	425	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 2 (Adamts2) alternative variant aSep08, mRNA.
<a href="#">Adamts2</a>	<a href="#">Adamts2.bSep08</a>	<a href="#">287899</a>	3252	792		172	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 2 (Adamts2) alternative variant bSep08, mRNA.
<a href="#">Adamts2</a>	<a href="#">Adamts2.cSep08</a>	<a href="#">287899</a>	959	214	2	71	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 2 (Adamts2) alternative variant cSep08, mRNA.
<a href="#">Adamts4</a>	<a href="#">Adamts4.aSep08</a>	<a href="#">66015</a>	2922	1254		417	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 4 (Adamts4) mRNA.
<a href="#">Adamts6</a>	<a href="#">Adamts6.bSep08</a>	<a href="#">361886</a>	62411	659	7	202	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 (Adamts6) alternative variant bSep08, mRNA.
<a href="#">Adamts7</a>	<a href="#">Adamts7.bSep08</a>	<a href="#">315879</a>	7693	2431	7	736	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 7 (Adamts7) alternative variant bSep08, mRNA.
<a href="#">Adamts7</a>	<a href="#">Adamts7.cSep08</a>	<a href="#">315879</a>	8808	397	2	33	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 7 (Adamts7) alternative variant cSep08, mRNA.
<a href="#">Adamts9</a>	<a href="#">Adamts9.bSep08</a>	<a href="#">312566</a>	12235	398	4	132	a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 9 (Adamts9) alternative variant bSep08, mRNA.
<a href="#">Adamts13</a>	<a href="#">Adamts13.aSep08</a>	<a href="#">362091</a>	3224	752		250	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 13 (Adamts13) mRNA.
<a href="#">Adamts17</a>	<a href="#">Adamts17.aSep08</a>	<a href="#">293004</a>	22257	373		123	ADAM metallopeptidase with thrombospondin type 1 motif, 17 (Adamts17) mRNA.
<a href="#">Adamts18</a>	<a href="#">Adamts18.aSep08</a>	<a href="#">361412</a>	71985	1704		498	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 18 (Adamts18) alternative variant aSep08, mRNA.
<a href="#">Adamts18</a>	<a href="#">Adamts18.bSep08</a>	<a href="#">361412</a>	6211	550		110	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 18 (Adamts18) alternative variant bSep08, mRNA.
<a href="#">Adamts19</a>	<a href="#">Adamts19.bSep08</a>	<a href="#">361332</a>	34964	1390	2	295	a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 19 (Adamts19) alternative variant bSep08, mRNA.
<a href="#">Adamts1</a>	<a href="#">Adamts1.aSep08</a>	<a href="#">362539</a>	99305	493		111	ADAMTS-like 1 (Adamts1) mRNA.



<a href="#">Adamtsl2</a>	<a href="#">Adamtsl2.aSep08</a>	<a href="#">311827</a>	7141	1677		360	ADAMTS-like 2 (Adamtsl2) mRNA.
<a href="#">Adamtsl3</a>	<a href="#">Adamtsl3.bSep08</a>	<a href="#">308787</a>	118115	905	5	125	ADAMTS-like 3 (Adamtsl3) alternative variant bSep08, mRNA.
<a href="#">Adamtsl4</a>	<a href="#">Adamtsl4.bSep08</a>	<a href="#">310670</a>	4337	757	5	252	ADAMTS-like 4 (Adamtsl4) alternative variant bSep08, mRNA.
<a href="#">Adamtsl4</a>	<a href="#">Adamtsl4.cSep08</a>	<a href="#">310670</a>	1221	728	3	242	ADAMTS-like 4 (Adamtsl4) alternative variant cSep08, mRNA.
<a href="#">Adamtsl4</a>	<a href="#">Adamtsl4.eSep08</a>	<a href="#">310670</a>	1329	946	3	140	ADAMTS-like 4 (Adamtsl4) alternative variant eSep08, mRNA.
<a href="#">Adamtsl4</a>	<a href="#">Adamtsl4.fSep08</a>	<a href="#">310670</a>	2416	346	4	115	ADAMTS-like 4 (Adamtsl4) alternative variant fSep08, mRNA.
<a href="#">Adamtsl4</a>	<a href="#">Adamtsl4.gSep08</a>	<a href="#">310670</a>	3678	514	4	113	ADAMTS-like 4 (Adamtsl4) alternative variant gSep08, mRNA.
<a href="#">Adamtsl5</a>	<a href="#">Adamtsl5.aSep08</a>	<a href="#">314626</a>	1640	1437	3	399	ADAMTS-like 5 (Adamtsl5) alternative variant aSep08, mRNA.
<a href="#">Adamtsl5</a>	<a href="#">Adamtsl5.bSep08</a>	<a href="#">314626</a>	601	517	1	171	ADAMTS-like 5 (Adamtsl5) alternative variant bSep08, mRNA.
<a href="#">ADAM_CR.1</a>	<a href="#">ADAM_CR.1.aSep08</a>		1692	308		102	domain 19 (ADAM_CR.1) mRNA.
<a href="#">ADAM_CR.2</a>	<a href="#">ADAM_CR.2.aSep08</a>		1121	686		117	A disintegrin metallopeptidase domain 8 (ADAM_CR.2) mRNA.
<a href="#">Adaptin_N.0</a>	<a href="#">Adaptin_N.0.aSep08</a>		89654	1393		463	adaptor-related protein complex 3 beta CRA a (Adaptin_N.0) mRNA.
<a href="#">Adar</a>	<a href="#">Adar.bSep08</a>	<a href="#">81635</a>	961	712	2	81	adenosine deaminase, RNA-specific (Adar) alternative variant bSep08, mRNA.
<a href="#">Adarb1</a>	<a href="#">Adarb1.dSep08</a>	<a href="#">25367</a>	89753	550	4	50	adenosine deaminase, RNA-specific, B1 (5.8 kD) (Adarb1) alternative variant dSep08, complete mRNA.
<a href="#">Adck1</a>	<a href="#">Adck1.bSep08</a>	<a href="#">366698</a>	8332	371	1	109	putative protein (Adck1) alternative variant bSep08, mRNA.
<a href="#">Adck2</a>	<a href="#">Adck2.bSep08</a>	<a href="#">312258</a>	12821	2296	6	567	uncharacterized aarF domain-containing protein kinase 2 (62.9 kD) (Adck2) alternative variant bSep08, mRNA.
<a href="#">Adck2</a>	<a href="#">Adck2.cSep08</a>	<a href="#">312258</a>	4176	1599	2	120	uncharacterized aarF domain-containing protein kinase 2 like (13.3 kD) (Adck2) alternative variant cSep08, mRNA.
<a href="#">Adck4</a>	<a href="#">Adck4.aSep08</a>	<a href="#">308453</a>	22872	1677	15	524	ABC-1 (Adck4) alternative variant aSep08, mRNA.
<a href="#">Adck4</a>	<a href="#">Adck4.bSep08</a>	<a href="#">308453</a>	10845	1403	10	294	ABC-1 (32.5 kD) (Adck4) alternative variant bSep08, mRNA.
<a href="#">Adck4</a>	<a href="#">Adck4.cSep08</a>	<a href="#">308453</a>	6117	761	6	183	uncharacterized aarF domain-containing protein kinase 4 (Adck4) alternative variant cSep08, mRNA.
<a href="#">Adck4</a>	<a href="#">Adck4.dSep08</a>	<a href="#">308453</a>	13617	1389	8	144	putative protein of ancient origin (Adck4) alternative variant dSep08, mRNA.
<a href="#">Adck4</a>	<a href="#">Adck4.eSep08</a>	<a href="#">308453</a>	2340	793	3	72	putative protein of ancient origin (Adck4) alternative variant eSep08, mRNA.
<a href="#">Adck5</a>	<a href="#">Adck5.aSep08</a>	<a href="#">362943</a>	17808	1980	15	602	ABC-1 (Adck5) alternative variant aSep08, mRNA.
<a href="#">Adck5</a>	<a href="#">Adck5.bSep08</a>	<a href="#">362943</a>	1248	873	6	243	putative protein of ancient origin (Adck5) alternative variant bSep08, mRNA.
<a href="#">Adck5</a>	<a href="#">Adck5.cSep08</a>	<a href="#">362943</a>	1382	698	5	138	ABC-1 (Adck5) alternative variant cSep08, mRNA.
<a href="#">Adck5</a>	<a href="#">Adck5.dSep08</a>	<a href="#">362943</a>	15832	734	6	129	uncharacterized aarF domain-containing protein kinase 5 like (15.0 kD) (Adck5) alternative variant dSep08, mRNA.

<a href="#">Adck5</a>	<a href="#">Adck5.fSep08</a>	<a href="#">362943</a>	474	388	2	67	putative protein of vertebrate origin (Adck5) alternative variant fSep08, mRNA.
<a href="#">Adcy1</a>	<a href="#">Adcy1.aSep08</a>	<a href="#">305509</a>	10406	409		136	adenylate cyclase 1 (Adcy1) mRNA.
<a href="#">Adcy2</a>	<a href="#">Adcy2.bSep08</a>	<a href="#">81636</a>	161235	3238	21	854	adenylate cyclase 2 (Adcy2) alternative variant bSep08, mRNA.
<a href="#">Adcy2</a>	<a href="#">Adcy2.cSep08</a>	<a href="#">81636</a>	50556	486	6	161	adenylate cyclase 2 (Adcy2) alternative variant cSep08, mRNA.
<a href="#">Adcy2</a>	<a href="#">Adcy2.dSep08</a>	<a href="#">81636</a>	2910	518	2	63	adenylate cyclase 2 (Adcy2) alternative variant dSep08, mRNA.
<a href="#">Adcy3</a>	<a href="#">Adcy3.bSep08</a>	<a href="#">64508</a>	3617	748	6	142	adenylate cyclase 3 (Adcy3) alternative variant bSep08, mRNA.
<a href="#">Adcy3</a>	<a href="#">Adcy3.dSep08</a>	<a href="#">64508</a>	727	389	2	38	adenylate cyclase 3 (Adcy3) alternative variant dSep08, mRNA.
<a href="#">Adcy4</a>	<a href="#">Adcy4.bSep08</a>	<a href="#">54223</a>	5971	970	8	303	adenylate cyclase 4 (34.2 kD) (Adcy4) alternative variant bSep08, mRNA.
<a href="#">Adcy4</a>	<a href="#">Adcy4.dSep08</a>	<a href="#">54223</a>	515	434	2	101	adenylate cyclase 4 CRA c (10.8 kD) (Adcy4) alternative variant dSep08, mRNA.
<a href="#">Adcy7</a>	<a href="#">Adcy7.aSep08</a>	<a href="#">84420</a>	5999	879	8	293	adenylate cyclase 7 (Adcy7) alternative variant aSep08, mRNA.
<a href="#">Adcy9</a>	<a href="#">Adcy9.aSep08</a>	<a href="#">302950</a>	12316	4390		461	adenylate cyclase 9 (Adcy9) mRNA.
<a href="#">Adcy10</a>	<a href="#">Adcy10.aSep08</a>	<a href="#">59320</a>	6903	758		252	adenylate cyclase 10 (Adcy10) mRNA.
<a href="#">Adcyap1</a>	<a href="#">Adcyap1.bSep08</a>	<a href="#">24166</a>	2413	977	3	113	adenylate cyclase activating polypeptide 1 (Adcyap1) alternative variant bSep08, mRNA.
<a href="#">Adcyap1</a>	<a href="#">Adcyap1.cSep08</a>	<a href="#">24166</a>	1047	258	3	67	adenylate cyclase activating polypeptide 1 (Adcyap1) alternative variant cSep08, mRNA.
<a href="#">Adcyap1r1</a>	<a href="#">Adcyap1r1.bSep08</a>	<a href="#">24167</a>	3466	349	2	115	adenylate cyclase activating polypeptide 1 receptor 1 (Adcyap1r1) alternative variant bSep08, mRNA.
<a href="#">Add1</a>	<a href="#">Add1.bSep08</a>	<a href="#">24170</a>	14273	1034	7	241	adducin 1 (alpha) (Add1) alternative variant bSep08, mRNA.
<a href="#">Add1</a>	<a href="#">Add1.cSep08</a>	<a href="#">24170</a>	1544	617	2	97	adducin 1 (alpha) (Add1) alternative variant cSep08, mRNA.
<a href="#">Add1</a>	<a href="#">Add1.dSep08</a>	<a href="#">24170</a>	1234	650	2	32	adducin 1 (alpha) (Add1) alternative variant dSep08, mRNA.
<a href="#">Add1</a>	<a href="#">Add1.fSep08</a>	<a href="#">24170</a>	31433	410	4	82	adducin 1 (alpha) (Add1) alternative variant fSep08, mRNA.
<a href="#">Add2</a>	<a href="#">Add2.cSep08</a>	<a href="#">24171</a>	8744	619	4	206	adducin 2 (beta) (Add2) alternative variant cSep08, mRNA.
<a href="#">Add2</a>	<a href="#">Add2.dSep08</a>	<a href="#">24171</a>	11554	725	6	162	adducin 2 (beta) (Add2) alternative variant dSep08, mRNA.
<a href="#">Add2</a>	<a href="#">Add2.eSep08</a>	<a href="#">24171</a>	21312	543	3	91	adducin 2 (beta) (10.6 kD) (Add2) alternative variant eSep08, mRNA.
<a href="#">Add3</a>	<a href="#">Add3.bSep08</a>	<a href="#">25230</a>	90494	1196	7	273	adducin 3 (gamma) (Add3) alternative variant bSep08, mRNA.
<a href="#">Add3</a>	<a href="#">Add3.cSep08</a>	<a href="#">25230</a>	9042	728	6	242	adducin 3 (gamma) (Add3) alternative variant cSep08, mRNA.
<a href="#">Add3</a>	<a href="#">Add3.eSep08</a>	<a href="#">25230</a>	74609	437	2	67	adducin 3 (gamma) (7.9 kD) (Add3) alternative variant eSep08, mRNA.

<a href="#">Adfp</a>	<a href="#">Adfp.bSep08</a>	<a href="#">298199</a>	11200	1144	2	381	adipose differentiation related protein (Adfp) alternative variant bSep08, mRNA.
<a href="#">Adfp</a>	<a href="#">Adfp.cSep08</a>	<a href="#">298199</a>	27494	1593	5	378	adipose differentiation related protein (Adfp) alternative variant cSep08, mRNA.
<a href="#">Adh1</a>	<a href="#">Adh1.bSep08</a>	<a href="#">24172</a>	3679	1121	3	261	alcohol dehydrogenase 1 (class I) (Adh1) alternative variant bSep08, mRNA.
<a href="#">Adh1</a>	<a href="#">Adh1.dSep08</a>	<a href="#">24172</a>	300	208		69	alcohol dehydrogenase 1 (class I) (Adh1) alternative variant dSep08, mRNA.
<a href="#">Adh1</a>	<a href="#">Adh1.eSep08</a>	<a href="#">24172</a>	299	207	2	68	alcohol dehydrogenase 1 (class I) (Adh1) alternative variant eSep08, mRNA.
<a href="#">Adh1</a>	<a href="#">Adh1.fSep08</a>	<a href="#">24172</a>	28416	795	5	11	alcohol dehydrogenase 1 (class I) (1.1 kD) (Adh1) alternative variant fSep08, mRNA.
<a href="#">Adh5</a>	<a href="#">Adh5.bSep08</a>	<a href="#">100145871</a>	8196	623	5	194	alcohol dehydrogenase 5 (Adh5) alternative variant bSep08, mRNA.
<a href="#">Adh6</a>	<a href="#">Adh6.aSep08</a>	<a href="#">310903</a>	30523	1680	2	390	alcohol dehydrogenase 6 (class V) (Adh6) alternative variant aSep08, mRNA.
<a href="#">Adh7</a>	<a href="#">Adh7.bSep08</a>	<a href="#">171178</a>	7048	754	1	166	alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide (18.3 kD) (Adh7) alternative variant bSep08, mRNA.
<a href="#">Adhfe1</a>	<a href="#">Adhfe1.bSep08</a>	<a href="#">362474</a>	10688	725	7	241	alcohol dehydrogenase, iron containing, 1 (Adhfe1) alternative variant bSep08, mRNA.
<a href="#">Adhfe1</a>	<a href="#">Adhfe1.cSep08</a>	<a href="#">362474</a>	9863	752	9	178	alcohol dehydrogenase, iron containing, 1 (Adhfe1) alternative variant cSep08, mRNA.
<a href="#">Adhfe1</a>	<a href="#">Adhfe1.dSep08</a>	<a href="#">362474</a>	6066	906	4	68	alcohol dehydrogenase, iron containing, 1 (Adhfe1) alternative variant dSep08, mRNA.
<a href="#">Adhfe1</a>	<a href="#">Adhfe1.eSep08</a>	<a href="#">362474</a>	5427	739	4	36	alcohol dehydrogenase, iron containing, 1 (4.4 kD) (Adhfe1) alternative variant eSep08, mRNA.
<a href="#">ADH_N.0</a>	<a href="#">ADH_N.0.aSep08</a>		33226	591		129	vesicle amine transport protein 1 homolog-like (ADH_N.0) mRNA.
<a href="#">adh_short.0</a>	<a href="#">adh_short.0.aSep08</a>		2425	1037	3	307	carbonyl reductase (adh_short.0) alternative variant aSep08, mRNA.
<a href="#">adh_short.1</a>	<a href="#">adh_short.1.aSep08</a>		6167	1086		227	retinol dehydrogenase (adh_short.1) mRNA.
<a href="#">Adi1</a>	<a href="#">Adi1.bSep08</a>	<a href="#">298934</a>	6042	621	4	151	acireductone dioxygenase 1 (18.0 kD) (Adi1) alternative variant bSep08, mRNA.
<a href="#">Adipor1</a>	<a href="#">Adipor1.bSep08</a>	<a href="#">289036</a>	14959	829	5	199	adiponectin receptor 1 (Adipor1) alternative variant bSep08, mRNA.
<a href="#">Adipor1</a>	<a href="#">Adipor1.cSep08</a>	<a href="#">289036</a>	16877	808	5	96	adiponectin receptor 1 (Adipor1) alternative variant cSep08, mRNA.
<a href="#">Adipor2</a>	<a href="#">Adipor2.bSep08</a>	<a href="#">312670</a>	35833	454	3	37	adiponectin receptor 2 (Adipor2) alternative variant bSep08, mRNA.
<a href="#">Adk</a>	<a href="#">Adk.bSep08</a>	<a href="#">25368</a>	283840	836	8	278	adenosine kinase (Adk) alternative variant bSep08, mRNA.
<a href="#">Adk</a>	<a href="#">Adk.cSep08</a>	<a href="#">25368</a>	87096	422	3	69	adenosine kinase (Adk) alternative variant cSep08, mRNA.
<a href="#">Adk</a>	<a href="#">Adk.dSep08</a>	<a href="#">25368</a>	1363	311	2	31	adenosine kinase (3.3 kD) (Adk) alternative variant dSep08, mRNA.
<a href="#">Adnp2</a>	<a href="#">Adnp2.bSep08</a>	<a href="#">307236</a>	20606	785	4	261	ADNP homeobox 2 (Adnp2) alternative variant bSep08, mRNA.

<a href="#">AdoHcyase_NA D.0</a>	<a href="#">AdoHcyase_NAD.0.aSep08</a>		16318	495		164	S-adenosylhomocysteine hydrolase-like (AdoHcyase_NAD.0) mRNA.
<a href="#">Adprh</a>	<a href="#">Adprh.bSep08</a>	<a href="#">25371</a>	4612	780	3	149	ADP-ribosylarginine hydrolase (Adprh) alternative variant bSep08, mRNA.
<a href="#">Adprhl2</a>	<a href="#">Adprhl2.bSep08</a>	<a href="#">362600</a>	2148	1315	4	217	ADP-ribosylhydrolase like 2 (23.5 kD) (Adprhl2) alternative variant bSep08, mRNA.
<a href="#">Adprhl2</a>	<a href="#">Adprhl2.cSep08</a>	<a href="#">362600</a>	3834	1037	3	211	ADP-ribosylhydrolase like 2 (21.9 kD) (Adprhl2) alternative variant cSep08, mRNA.
<a href="#">Adrbk1</a>	<a href="#">Adrbk1.bSep08</a>	<a href="#">25238</a>	2741	2089	2	230	adrenergic receptor kinase, beta 1 (26.8 kD) (Adrbk1) alternative variant bSep08, mRNA.
<a href="#">Adrm1</a>	<a href="#">Adrm1.bSep08</a>	<a href="#">65138</a>	695	590	2	88	adhesion regulating molecule 1 (Adrm1) alternative variant bSep08, mRNA.
<a href="#">Adsl</a>	<a href="#">Adsl.bSep08</a>	<a href="#">315150</a>	9136	1927	6	223	adenylosuccinate lyase (24.9 kD) (Adsl) alternative variant bSep08, mRNA.
<a href="#">Adsl</a>	<a href="#">Adsl.cSep08</a>	<a href="#">315150</a>	19093	760	7	122	adenylosuccinate lyase (Adsl) alternative variant cSep08, mRNA.
<a href="#">Adsl</a>	<a href="#">Adsl.dSep08</a>	<a href="#">315150</a>	2690	1084	1	46	adenylosuccinate lyase (Adsl) alternative variant dSep08, mRNA.
<a href="#">Aebp1</a>	<a href="#">Aebp1.bSep08</a>	<a href="#">305494</a>	1985	578	7	192	AE binding protein 1 (Aebp1) alternative variant bSep08, mRNA.
<a href="#">Aebp2</a>	<a href="#">Aebp2.bSep08</a>	<a href="#">297705</a>	13230	5887	4	106	AE binding protein 2 (Aebp2) alternative variant bSep08, mRNA.
<a href="#">Aebp2</a>	<a href="#">Aebp2.cSep08</a>	<a href="#">297705</a>	3663	782	2	33	AE binding protein 2 (Aebp2) alternative variant cSep08, mRNA.
<a href="#">Aer61</a>	<a href="#">Aer61.bSep08</a>	<a href="#">494219</a>	33943	1792	11	466	glycosyltransferase Aer61 (Aer61) alternative variant bSep08, mRNA.
<a href="#">Aer61</a>	<a href="#">Aer61.cSep08</a>	<a href="#">494219</a>	3060	2588	2	72	glycosyltransferase Aer61 (7.9 kD) (Aer61) alternative variant cSep08, mRNA.
<a href="#">Aes</a>	<a href="#">Aes.bSep08</a>	<a href="#">29466</a>	5143	1242	6	187	amino-terminal enhancer of split (Aes) alternative variant bSep08, mRNA.
<a href="#">Aes</a>	<a href="#">Aes.cSep08</a>	<a href="#">29466</a>	3425	352	4	61	amino-terminal enhancer of split (Aes) alternative variant cSep08, mRNA.
<a href="#">Aes</a>	<a href="#">Aes.dSep08</a>	<a href="#">29466</a>	1273	730	3	42	amino-terminal enhancer of split (Aes) alternative variant dSep08, mRNA.
<a href="#">Afap1</a>	<a href="#">Afap1.bSep08</a>	<a href="#">140935</a>	1759	482	1	72	actin filament associated protein 1 (Afap1) alternative variant bSep08, mRNA.
<a href="#">Afap1l1</a>	<a href="#">Afap1l1.bSep08</a>	<a href="#">291565</a>	13236	2899	5	191	actin filament associated protein 1-like 1 (Afap1l1) alternative variant bSep08, mRNA.
<a href="#">Afap1l2</a>	<a href="#">Afap1l2.aSep08</a>	<a href="#">292130</a>	4888	2074		344	actin filament associated protein 1-like 2 (Afap1l2) mRNA.
<a href="#">Aff1</a>	<a href="#">Aff1.bSep08</a>	<a href="#">305152</a>	74363	399	2	100	AF4/FMR2 family, member 1 (Aff1) alternative variant bSep08, mRNA.
<a href="#">Aff2</a>	<a href="#">Aff2.aSep08</a>	<a href="#">293922</a>	16588	468		155	AF4/FMR2 family, member 2 (Aff2) mRNA.
<a href="#">Aff3</a>	<a href="#">Aff3.aSep08</a>	<a href="#">363220</a>	12288	828	5	233	AF4/FMR2 family, member 3 (Aff3) alternative variant aSep08, mRNA.
<a href="#">Aff3</a>	<a href="#">Aff3.bSep08</a>	<a href="#">363220</a>	12185	677	5	182	AF4/FMR2 family, member 3 (Aff3) alternative variant bSep08, mRNA.

<a href="#">Aff3</a>	<a href="#">Aff3.cSep08</a>	<a href="#">363220</a>	2286	637	1	150	AF4/FMR2 family, member 3 (Aff3) alternative variant cSep08, mRNA.
<a href="#">Aff4</a>	<a href="#">Aff4.bSep08</a>	<a href="#">303132</a>	4107	808	5	246	AF4/FMR2 family, member 4 (Aff4) alternative variant bSep08, mRNA.
<a href="#">Aff4</a>	<a href="#">Aff4.cSep08</a>	<a href="#">303132</a>	29551	663	7	221	AF4/FMR2 family, member 4 (Aff4) alternative variant cSep08, mRNA.
<a href="#">Aff4</a>	<a href="#">Aff4.dSep08</a>	<a href="#">303132</a>	9205	607	3	201	AF4/FMR2 family, member 4 (Aff4) alternative variant dSep08, mRNA.
<a href="#">Aff4</a>	<a href="#">Aff4.fSep08</a>	<a href="#">303132</a>	3653	349	2	41	AF4/FMR2 family, member 4 (4.8 kD) (Aff4) alternative variant fSep08, mRNA.
<a href="#">Afg3l1</a>	<a href="#">Afg3l1.bSep08</a>	<a href="#">361436</a>	12792	1675	10	520	AFG3(ATPase family gene 3)-like 1 (S. cerevisiae) (Afg3l1) alternative variant bSep08, mRNA.
<a href="#">Afg3l1</a>	<a href="#">Afg3l1.cSep08</a>	<a href="#">361436</a>	13542	500	6	166	AFG3(ATPase family gene 3)-like 1 (S. cerevisiae) (Afg3l1) alternative variant cSep08, mRNA.
<a href="#">Afg3l2</a>	<a href="#">Afg3l2.bSep08</a>	<a href="#">307350</a>	3525	795	2	105	AFG3(ATPase family gene 3)-like 2 (yeast) (Afg3l2) alternative variant bSep08, mRNA.
<a href="#">Afg3l2</a>	<a href="#">Afg3l2.cSep08</a>	<a href="#">307350</a>	2593	712	2	101	AFG3(ATPase family gene 3)-like 2 (yeast) (Afg3l2) alternative variant cSep08, mRNA.
<a href="#">Afp</a>	<a href="#">Afp.bSep08</a>	<a href="#">24177</a>	7895	1235	6	274	alpha-fetoprotein (Afp) alternative variant bSep08, mRNA.
<a href="#">Afp</a>	<a href="#">Afp.cSep08</a>	<a href="#">24177</a>	6936	764	6	230	alpha-fetoprotein (Afp) alternative variant cSep08, mRNA.
<a href="#">Afp</a>	<a href="#">Afp.dSep08</a>	<a href="#">24177</a>	6817	651	5	211	alpha-fetoprotein precursor (23.5 kD) (Afp) alternative variant dSep08, mRNA.
<a href="#">Afp</a>	<a href="#">Afp.eSep08</a>	<a href="#">24177</a>	4175	812	5	183	alpha-fetoprotein (Afp) alternative variant eSep08, mRNA.
<a href="#">Aftph</a>	<a href="#">Aftph.aSep08</a>	<a href="#">305544</a>	53958	3745	3	905	aftiphilin (98.1 kD) (Aftph) alternative variant aSep08, complete mRNA.
<a href="#">Aftph</a>	<a href="#">Aftph.bSep08</a>	<a href="#">305544</a>	14353	346	3	115	aftiphilin (Aftph) alternative variant bSep08, mRNA.
<a href="#">Aftph</a>	<a href="#">Aftph.cSep08</a>	<a href="#">305544</a>	6842	1708	2	60	aftiphilin (6.5 kD) (Aftph) alternative variant cSep08, mRNA.
<a href="#">Aftph</a>	<a href="#">Aftph.eSep08</a>	<a href="#">305544</a>	9516	404	3	26	aftiphilin (Aftph) alternative variant eSep08, mRNA.
<a href="#">Aga</a>	<a href="#">Aga.bSep08</a>	<a href="#">290923</a>	11789	958	8	249	aspartylglucosaminidase (27.1 kD) (Aga) alternative variant bSep08, mRNA.
<a href="#">Aga</a>	<a href="#">Aga.cSep08</a>	<a href="#">290923</a>	5919	712	6	224	aspartylglucosaminidase (Aga) alternative variant cSep08, mRNA.
<a href="#">Aga</a>	<a href="#">Aga.dSep08</a>	<a href="#">290923</a>	6180	767	5	138	aspartylglucosaminidase (14.7 kD) (Aga) alternative variant dSep08, mRNA.
<a href="#">Aga</a>	<a href="#">Aga.eSep08</a>	<a href="#">290923</a>	1997	419	3	102	aspartylglucosaminidase (Aga) alternative variant eSep08, mRNA.
<a href="#">Aga</a>	<a href="#">Aga.gSep08</a>	<a href="#">290923</a>	2643	492	2	71	aspartylglucosaminidase (8.0 kD) (Aga) alternative variant gSep08, mRNA.
<a href="#">Agbl2</a>	<a href="#">Agbl2.aSep08</a>	<a href="#">366124</a>	11086	562		92	ATP/GTP binding protein-like 2 (Agbl2) mRNA.
<a href="#">Agbl3</a>	<a href="#">Agbl3.aSep08</a>	<a href="#">500076</a>	36603	773	8	257	ATP/GTP binding protein-like 3 (Agbl3) alternative variant aSep08, mRNA.
<a href="#">Agbl3</a>	<a href="#">Agbl3.bSep08</a>	<a href="#">500076</a>	8445	801	1	137	ATP/GTP binding protein-like 3 (16.1 kD) (Agbl3) alternative variant bSep08, mRNA.
<a href="#">Ager</a>	<a href="#">Ager.bSep08</a>	<a href="#">81722</a>	1501	1199	4	147	advanced glycosylation end product-specific receptor (Ager) alternative variant bSep08, mRNA.

<a href="#">Ager</a>	<a href="#">Ager.cSep08</a>	<a href="#">81722</a>	616	413	3	84	advanced glycosylation end product-specific receptor (Ager) alternative variant cSep08, mRNA.
<a href="#">Ager</a>	<a href="#">Ager.dSep08</a>	<a href="#">81722</a>	492	406	2	82	advanced glycosylation end product-specific receptor (Ager) alternative variant dSep08, mRNA.
<a href="#">Ager</a>	<a href="#">Ager.eSep08</a>	<a href="#">81722</a>	951	406	3	81	advanced glycosylation end product-specific receptor (Ager) alternative variant eSep08, mRNA.
<a href="#">Aggf1</a>	<a href="#">Aggf1.aSep08</a>	<a href="#">310005</a>	18992	1757	10	585	angiogenic factor with G patch FHA domains 1 (Aggf1) alternative variant aSep08, mRNA.
<a href="#">Aggf1</a>	<a href="#">Aggf1.bSep08</a>	<a href="#">310005</a>	14943	1789	9	323	angiogenic factor VG5Q (Aggf1) alternative variant bSep08, mRNA.
<a href="#">Aggf1</a>	<a href="#">Aggf1.cSep08</a>	<a href="#">310005</a>	3422	399	2	124	angiogenic factor VG5Q (Aggf1) alternative variant cSep08, mRNA.
<a href="#">Aggf1</a>	<a href="#">Aggf1.dSep08</a>	<a href="#">310005</a>	2158	367	2	81	angiogenic factor VG5Q (Aggf1) alternative variant dSep08, mRNA.
<a href="#">Aggf1</a>	<a href="#">Aggf1.fSep08</a>	<a href="#">310005</a>	1011	278	2	25	putative protein (2.9 kD) (Aggf1) alternative variant fSep08, mRNA.
<a href="#">Agk</a>	<a href="#">Agk.bSep08</a>	<a href="#">502749</a>	51247	616		150	acylglycerol kinase (Agk) alternative variant bSep08, mRNA.
<a href="#">Agl</a>	<a href="#">Agl.bSep08</a>	<a href="#">362029</a>	4138	509	4	169	amylase-1,6-glycosidase, 4-alpha-glucanotransferase (Agl) alternative variant bSep08, mRNA.
<a href="#">Agl</a>	<a href="#">Agl.cSep08</a>	<a href="#">362029</a>	13475	349	3	75	amylase-1,6-glycosidase, 4-alpha-glucanotransferase (Agl) alternative variant cSep08, mRNA.
<a href="#">Agmat</a>	<a href="#">Agmat.bSep08</a>	<a href="#">298607</a>	3141	967	3	113	agmatine ureohydrolase (agmatinase) (12.0 kD) (Agmat) alternative variant bSep08, mRNA.
<a href="#">Agmat</a>	<a href="#">Agmat.cSep08</a>	<a href="#">298607</a>	3088	681	3	101	agmatine ureohydrolase (agmatinase) (10.5 kD) (Agmat) alternative variant cSep08, mRNA.
<a href="#">Ago_hook.0</a>	<a href="#">Ago_hook.0.aSep08</a>		3845	1849	3	616	trinucleotide repeat containing 6a (Ago_hook.0) alternative variant aSep08, mRNA.
<a href="#">Agpat1</a>	<a href="#">Agpat1.bSep08</a>	<a href="#">406165</a>	1030	523	4	173	1-acylglycerol-3-phosphate O-acyltransferase 1 (Agpat1) alternative variant bSep08, mRNA.
<a href="#">Agpat1</a>	<a href="#">Agpat1.cSep08</a>	<a href="#">406165</a>	7855	805	5	171	1-acylglycerol-3-phosphate O-acyltransferase 1 (18.0 kD) (Agpat1) alternative variant cSep08, mRNA.
<a href="#">Agpat1</a>	<a href="#">Agpat1.dSep08</a>	<a href="#">406165</a>	6440	474	4	158	1-acylglycerol-3-phosphate O-acyltransferase 1 (Agpat1) alternative variant dSep08, mRNA.
<a href="#">Agpat1</a>	<a href="#">Agpat1.eSep08</a>	<a href="#">406165</a>	969	768	2	255	1-acylglycerol-3-phosphate O-acyltransferase 1 (Agpat1) alternative variant eSep08, mRNA.
<a href="#">Agpat1</a>	<a href="#">Agpat1.fSep08</a>	<a href="#">406165</a>	6998	828	3	104	1-acylglycerol-3-phosphate O-acyltransferase 1 (Agpat1) alternative variant fSep08, mRNA.
<a href="#">Agpat2</a>	<a href="#">Agpat2.bSep08</a>	<a href="#">311821</a>	10801	885	1	175	1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta) (19.5 kD) (Agpat2) alternative variant bSep08, mRNA.
<a href="#">Agpat3</a>	<a href="#">Agpat3.bSep08</a>	<a href="#">294324</a>	1336	691	2	89	O-acyltransferase 3 (Agpat3) alternative variant bSep08, mRNA.
<a href="#">Agpat3</a>	<a href="#">Agpat3.cSep08</a>	<a href="#">294324</a>	65876	237	3	73	putative protein (Agpat3) alternative variant cSep08, mRNA.
<a href="#">Agpat3</a>	<a href="#">Agpat3.dSep08</a>	<a href="#">294324</a>	65867	349	3	69	putative protein (Agpat3) alternative variant dSep08, mRNA.

<a href="#">Agpat3</a>	<a href="#">Agpat3.eSep08</a>	<a href="#">294324</a>	19692	321	2	39	putative protein (Agpat3) alternative variant eSep08, mRNA.
<a href="#">Agpat4</a>	<a href="#">Agpat4.bSep08</a>	<a href="#">170919</a>	102988	715	3	118	1-acylglycerol-3-phosphate O-acyltransferase 4 (lysophosphatidic acid acyltransferase, delta) (13.0 kD) (Agpat4) alternative variant bSep08, mRNA.
<a href="#">Agpat4</a>	<a href="#">Agpat4.cSep08</a>	<a href="#">170919</a>	33521	359	1	60	1-acylglycerol-3-phosphate O-acyltransferase 4 (lysophosphatidic acid acyltransferase, delta) (Agpat4) alternative variant cSep08, mRNA.
<a href="#">Agpat5</a>	<a href="#">Agpat5.bSep08</a>	<a href="#">306582</a>	30077	488	4	147	1-acylglycerol-3-phosphate O-acyltransferase 5 (lysophosphatidic acid acyltransferase, epsilon) (Agpat5) alternative variant bSep08, mRNA.
<a href="#">Agpat7</a>	<a href="#">Agpat7.bSep08</a>	<a href="#">296048</a>	4249	1099	2	235	1-acylglycerol-3-phosphate O-acyltransferase 7 (lysophosphatidic acid acyltransferase, eta) (25.5 kD) (Agpat7) alternative variant bSep08, mRNA.
<a href="#">Agpat9</a>	<a href="#">Agpat9.bSep08</a>	<a href="#">305166</a>	35122	705	2	234	1-acylglycerol-3-phosphate O-acyltransferase 9 (Agpat9) alternative variant bSep08, mRNA.
<a href="#">Agps</a>	<a href="#">Agps.aSep08</a>	<a href="#">84114</a>	19166	417		136	alkylglycerone phosphate synthase (Agps) mRNA.
<a href="#">Agr2</a>	<a href="#">Agr2.aSep08</a>	<a href="#">298961</a>	10310	701	1	192	anterior gradient 2 ( <i>Xenopus laevis</i> ) (Agr2) alternative variant aSep08, mRNA.
<a href="#">Agrn</a>	<a href="#">Agrn.bSep08</a>	<a href="#">25592</a>	4723	1429	9	320	agrin (Agrn) alternative variant bSep08, mRNA.
<a href="#">Agrn</a>	<a href="#">Agrn.cSep08</a>	<a href="#">25592</a>	3203	797	6	198	agrin (Agrn) alternative variant cSep08, mRNA.
<a href="#">Agrn</a>	<a href="#">Agrn.dSep08</a>	<a href="#">25592</a>	17518	414	4	137	agrin (Agrn) alternative variant dSep08, mRNA.
<a href="#">Agrn</a>	<a href="#">Agrn.eSep08</a>	<a href="#">25592</a>	585	431	2	79	agrin (Agrn) alternative variant eSep08, mRNA.
<a href="#">Agrp</a>	<a href="#">Agrp.aSep08</a>	<a href="#">25582</a>	759	409		116	agouti related protein (Agrp) mRNA.
<a href="#">Agt</a>	<a href="#">Agt.bSep08</a>	<a href="#">24179</a>	2733	795	3	212	angiotensinogen (serpin peptidase inhibitor, clade A, member 8) (Agt) alternative variant bSep08, mRNA.
<a href="#">Agt</a>	<a href="#">Agt.cSep08</a>	<a href="#">24179</a>	1260	636	2	113	angiotensinogen (serpin peptidase inhibitor, clade A, member 8) (Agt) alternative variant cSep08, mRNA.
<a href="#">Agtbbp1</a>	<a href="#">Agtbbp1.bSep08</a>	<a href="#">290986</a>	25600	1063	6	220	ATP/GTP binding protein 1 (Agtbbp1) alternative variant bSep08, mRNA.
<a href="#">Agtbbp1</a>	<a href="#">Agtbbp1.cSep08</a>	<a href="#">290986</a>	25380	768	4	152	ATP/GTP binding protein 1 (Agtbbp1) alternative variant cSep08, mRNA.
<a href="#">Agtbbp1</a>	<a href="#">Agtbbp1.dSep08</a>	<a href="#">290986</a>	15900	402	4	130	ATP/GTP binding protein 1 (Agtbbp1) alternative variant dSep08, mRNA.
<a href="#">Agr2</a>	<a href="#">Agr2.bSep08</a>	<a href="#">24182</a>	3839	2954	2	363	angiotensin II receptor, type 2 (41.3 kD) (Agr2) alternative variant bSep08, mRNA.
<a href="#">Agxt</a>	<a href="#">Agxt.bSep08</a>	<a href="#">24792</a>	5502	934	5	228	alanine-glyoxylate aminotransferase (Agxt) alternative variant bSep08, mRNA.
<a href="#">Agxt</a>	<a href="#">Agxt.cSep08</a>	<a href="#">24792</a>	4007	417	2	111	alanine-glyoxylate aminotransferase (Agxt) alternative variant cSep08, mRNA.
<a href="#">Ahctf1</a>	<a href="#">Ahctf1.aSep08</a>	<a href="#">360886</a>	9518	3180		578	AT hook containing transcription factor 1 (Ahctf1) mRNA.
<a href="#">Ahcy</a>	<a href="#">Ahcy.bSep08</a>	<a href="#">29443</a>	2340	761	1	175	S-adenosylhomocysteine hydrolase (Ahcy) alternative variant bSep08, mRNA.
<a href="#">Ahcy1</a>	<a href="#">Ahcy1.aSep08</a>	<a href="#">362013</a>	49241	2443	17	524	S-adenosylhomocysteine hydrolase-like 1 (58.6 kD) (Ahcy1) alternative variant aSep08, mRNA.

<a href="#">Ahcy1</a>	<a href="#">Ahcy1.cSep08</a>	<a href="#">362013</a>	25485	791	8	190	S-adenosylhomocysteine hydrolase-like 1 (Ahcy1) alternative variant cSep08, mRNA.
<a href="#">Ahcy1</a>	<a href="#">Ahcy1.dSep08</a>	<a href="#">362013</a>	24915	841	6	173	S-adenosylhomocysteine hydrolase-like 1 (Ahcy1) alternative variant dSep08, mRNA.
<a href="#">Ahcy1</a>	<a href="#">Ahcy1.eSep08</a>	<a href="#">362013</a>	6097	2147	8	158	S-adenosylhomocysteine hydrolase-like 1 (Ahcy1) alternative variant eSep08, mRNA.
<a href="#">Ahcy2</a>	<a href="#">Ahcy2.aSep08</a>	<a href="#">312192</a>	125859	1282	10	423	S-adenosylhomocysteine hydrolase-like 2 (Ahcy2) alternative variant aSep08, mRNA.
<a href="#">Ahcy2</a>	<a href="#">Ahcy2.cSep08</a>	<a href="#">312192</a>	16105	536	3	79	S-adenosylhomocysteine hydrolase-like 2 (Ahcy2) alternative variant cSep08, mRNA.
<a href="#">Ahdc1</a>	<a href="#">Ahdc1.aSep08</a>	<a href="#">362617</a>	12553	2503	2	670	AT hook, DNA binding motif, containing 1 (Ahdc1) alternative variant aSep08, mRNA.
<a href="#">Ahi1</a>	<a href="#">Ahi1.bSep08</a>	<a href="#">308923</a>	132267	1887	3	543	abelson helper integration site 1 (Ahi1) alternative variant bSep08, mRNA.
<a href="#">Ahi1</a>	<a href="#">Ahi1.cSep08</a>	<a href="#">308923</a>	72156	2042	7	183	abelson helper integration site 1 (Ahi1) alternative variant cSep08, mRNA.
<a href="#">Ahi1</a>	<a href="#">Ahi1.dSep08</a>	<a href="#">308923</a>	44327	735	7	150	abelson helper integration site 1 (17.3 kD) (Ahi1) alternative variant dSep08, mRNA.
<a href="#">Ahi1</a>	<a href="#">Ahi1.eSep08</a>	<a href="#">308923</a>	12751	695	3	39	abelson helper integration site 1 (4.4 kD) (Ahi1) alternative variant eSep08, mRNA.
<a href="#">Ahr</a>	<a href="#">Ahr.bSep08</a>	<a href="#">25690</a>	35153	2590	2	815	aryl hydrocarbon receptor (92.3 kD) (Ahr) alternative variant bSep08, mRNA.
<a href="#">Ahr</a>	<a href="#">Ahr.cSep08</a>	<a href="#">25690</a>	35153	2690	2	815	aryl hydrocarbon receptor (92.3 kD) (Ahr) alternative variant cSep08, mRNA.
<a href="#">Ahr</a>	<a href="#">Ahr.dSep08</a>	<a href="#">25690</a>	36363	3905	2	815	aryl hydrocarbon receptor (92.3 kD) (Ahr) alternative variant dSep08, complete mRNA.
<a href="#">Ahr</a>	<a href="#">Ahr.eSep08</a>	<a href="#">25690</a>	35153	2432	2	810	aryl hydrocarbon receptor (Ahr) alternative variant eSep08, mRNA.
<a href="#">Ahsa2</a>	<a href="#">Ahsa2.aSep08</a>	<a href="#">305577</a>	7432	702	6	233	AHA1, activator of heat shock protein ATPase homolog 2 (yeast) (Ahsa2) alternative variant aSep08, mRNA.
<a href="#">Ahsa2</a>	<a href="#">Ahsa2.bSep08</a>	<a href="#">305577</a>	5546	520	4	173	AHA1, activator of heat shock protein ATPase homolog 2 (yeast) (Ahsa2) alternative variant bSep08, mRNA.
<a href="#">Ahsg</a>	<a href="#">Ahsg.bSep08</a>	<a href="#">25373</a>	1289	733	2	129	alpha-2-HS-glycoprotein (Ahsg) alternative variant bSep08, mRNA.
<a href="#">Ahsg</a>	<a href="#">Ahsg.cSep08</a>	<a href="#">25373</a>	569	260	2	86	alpha-2-HS-glycoprotein (Ahsg) alternative variant cSep08, mRNA.
<a href="#">Aif1</a>	<a href="#">Aif1.aSep08</a>	<a href="#">29427</a>	741	456	1	151	allograft inflammatory factor 1 (Aif1) alternative variant aSep08, mRNA.
<a href="#">Aifm1</a>	<a href="#">Aifm1.bSep08</a>	<a href="#">83533</a>	6743	484	4	160	apoptosis-inducing factor, mitochondrion-associated 1 (Aifm1) alternative variant bSep08, mRNA.
<a href="#">Aifm1</a>	<a href="#">Aifm1.cSep08</a>	<a href="#">83533</a>	970	625	2	116	apoptosis-inducing factor, mitochondrion-associated 1 (13.4 kD) (Aifm1) alternative variant cSep08, mRNA.
<a href="#">Aifm2</a>	<a href="#">Aifm2.bSep08</a>	<a href="#">361843</a>	6648	898	5	175	apoptosis-inducing factor, mitochondrion-associated 2 (Aifm2) alternative variant bSep08, mRNA.
<a href="#">Aifm2</a>	<a href="#">Aifm2.cSep08</a>	<a href="#">361843</a>	7612	715	2	38	apoptosis-inducing factor, mitochondrion-associated 2 (4.5 kD) (Aifm2) alternative variant cSep08, mRNA.



<a href="#">Aifm2</a>	<a href="#">Aifm2.dSep08</a>	<a href="#">361843</a>	6605	638	2	38	apoptosis-inducing factor, mitochondrion-associated 2 (4.5 kD) (Aifm2) alternative variant dSep08, mRNA.
<a href="#">Aifm3</a>	<a href="#">Aifm3.aSep08</a>	<a href="#">303786</a>	6015	1541	12	177	apoptosis-inducing factor, mitochondrion-associated 3 (19.8 kD) (Aifm3) alternative variant aSep08, mRNA.
<a href="#">Aifm3</a>	<a href="#">Aifm3.bSep08</a>	<a href="#">303786</a>	1258	433	5	144	apoptosis-inducing factor, mitochondrion-associated 3 (Aifm3) alternative variant bSep08, mRNA.
<a href="#">Aifm3</a>	<a href="#">Aifm3.cSep08</a>	<a href="#">303786</a>	2510	621	4	97	apoptosis-inducing factor, mitochondrion-associated 3 (Aifm3) alternative variant cSep08, mRNA.
<a href="#">Aifm3</a>	<a href="#">Aifm3.dSep08</a>	<a href="#">303786</a>	734	384	2	62	apoptosis-inducing factor, mitochondrion-associated 3 (Aifm3) alternative variant dSep08, mRNA.
<a href="#">Aim1l</a>	<a href="#">Aim1l.aSep08</a>	<a href="#">298543</a>	7921	840		280	absent in melanoma 1-like (Aim1l) mRNA.
<a href="#">Aim2</a>	<a href="#">Aim2.aSep08</a>	<a href="#">304987</a>	5808	832		148	absent in melanoma 2 (16.7 kD) (Aim2) mRNA.
<a href="#">Aip</a>	<a href="#">Aip.bSep08</a>	<a href="#">282827</a>	3462	487	1	125	aryl-hydrocarbon receptor-interacting protein (Aip) alternative variant bSep08, mRNA.
<a href="#">Ak7</a>	<a href="#">Ak7.bSep08</a>	<a href="#">314416</a>	32246	1456	8	335	adenylate kinase 7 (Ak7) alternative variant bSep08, mRNA.
<a href="#">Ak7</a>	<a href="#">Ak7.cSep08</a>	<a href="#">314416</a>	23532	735	5	203	adenylate kinase 7 (Ak7) alternative variant cSep08, mRNA.
<a href="#">Ak7</a>	<a href="#">Ak7.dSep08</a>	<a href="#">314416</a>	5379	415	3	104	adenylate kinase 7 (Ak7) alternative variant dSep08, mRNA.
<a href="#">Akap1</a>	<a href="#">Akap1.bSep08</a>	<a href="#">114124</a>	1345	799	2	255	A kinase anchor protein 1 CRA b (Akap1) alternative variant bSep08, mRNA.
<a href="#">Akap1</a>	<a href="#">Akap1.cSep08</a>	<a href="#">114124</a>	761	399	2	83	A-kinase anchor protein (Akap1) alternative variant cSep08, mRNA.
<a href="#">Akap1</a>	<a href="#">Akap1.dSep08</a>	<a href="#">114124</a>	18154	466	2	66	putative protein (Akap1) alternative variant dSep08, mRNA.
<a href="#">Akap1</a>	<a href="#">Akap1.eSep08</a>	<a href="#">114124</a>	11329	780	2	58	putative protein (Akap1) alternative variant eSep08, mRNA.
<a href="#">Akap2</a>	<a href="#">Akap2.bSep08</a>	<a href="#">298024</a>	29517	4463	2	113	A kinase (PRKA) anchor protein 2 (Akap2) alternative variant bSep08, mRNA.
<a href="#">Akap2</a>	<a href="#">Akap2.cSep08</a>	<a href="#">298024</a>	10173	838	3	60	A kinase (PRKA) anchor protein 2 (Akap2) alternative variant cSep08, mRNA.
<a href="#">Akap3</a>	<a href="#">Akap3.bSep08</a>	<a href="#">312720</a>	11908	778	1	207	A kinase (PRKA) anchor protein 3 (Akap3) alternative variant bSep08, mRNA.
<a href="#">Akap4</a>	<a href="#">Akap4.bSep08</a>	<a href="#">79254</a>	4896	317	1	63	A kinase (PRKA) anchor protein 4 (Akap4) alternative variant bSep08, mRNA.
<a href="#">Akap5</a>	<a href="#">Akap5.aSep08</a>	<a href="#">171026</a>	7891	1277	3	212	A kinase (PRKA) anchor protein 5 (23.4 kD) (Akap5) alternative variant aSep08, mRNA.
<a href="#">Akap6</a>	<a href="#">Akap6.bSep08</a>	<a href="#">64553</a>	22161	809	2	77	A kinase (PRKA) anchor protein 6 (Akap6) alternative variant bSep08, mRNA.
<a href="#">Akap7</a>	<a href="#">Akap7.bSep08</a>	<a href="#">361458</a>	51028	626	4	128	A-kinase anchor protein 7 (Akap7) alternative variant bSep08, mRNA.
<a href="#">Akap7</a>	<a href="#">Akap7.cSep08</a>	<a href="#">361458</a>	28414	788	2	81	a-kinase anchor protein 7 (9.2 kD) (Akap7) alternative variant cSep08, mRNA.
<a href="#">Akap8</a>	<a href="#">Akap8.bSep08</a>	<a href="#">116633</a>	6804	1463	6	310	A kinase (PRKA) anchor protein 8 (34.6 kD) (Akap8) alternative variant bSep08, mRNA.
<a href="#">Akap8</a>	<a href="#">Akap8.dSep08</a>	<a href="#">116633</a>	3438	381	5	127	A kinase (PRKA) anchor protein 8 (Akap8) alternative variant dSep08, mRNA.

<a href="#">Akap8</a>	<a href="#">Akap8.eSep08</a>	<a href="#">116633</a>	1242	1070	2	74	A kinase (PRKA) anchor protein 8 (14.9 kD) (Akap8) alternative variant eSep08, mRNA.
<a href="#">Akap8l</a>	<a href="#">Akap8l.bSep08</a>	<a href="#">299569</a>	3426	2847	4	444	a-kinase anchor protein 8-like precursor (47.7 kD) (Akap8l) alternative variant bSep08, mRNA.
<a href="#">Akap8l</a>	<a href="#">Akap8l.cSep08</a>	<a href="#">299569</a>	15484	1149	10	331	A kinase anchor protein 8-like (36.6 kD) (Akap8l) alternative variant cSep08, mRNA.
<a href="#">Akap8l</a>	<a href="#">Akap8l.dSep08</a>	<a href="#">299569</a>	12912	898	5	141	a-kinase anchor protein 8-like (Akap8l) alternative variant dSep08, mRNA.
<a href="#">Akap8l</a>	<a href="#">Akap8l.eSep08</a>	<a href="#">299569</a>	16652	416	2	138	putative protein (Akap8l) alternative variant eSep08, mRNA.
<a href="#">Akap8l</a>	<a href="#">Akap8l.fSep08</a>	<a href="#">299569</a>	13052	977	4	134	A kinase anchor protein 8-like (Akap8l) alternative variant fSep08, mRNA.
<a href="#">Akap9</a>	<a href="#">Akap9.bSep08</a>	<a href="#">246150</a>	102918	730	7	129	A kinase (PRKA) anchor protein (yotiao) 9 (14.8 kD) (Akap9) alternative variant bSep08, mRNA.
<a href="#">Akap10</a>	<a href="#">Akap10.bSep08</a>	<a href="#">360540</a>	26457	2825	1	354	A kinase (PRKA) anchor protein 10 (39.1 kD) (Akap10) alternative variant bSep08, complete mRNA.
<a href="#">Akap11</a>	<a href="#">Akap11.bSep08</a>	<a href="#">498549</a>	11293	3917	1	142	A kinase (PRKA) anchor protein 11 (Akap11) alternative variant bSep08, mRNA.
<a href="#">Akap13</a>	<a href="#">Akap13.bSep08</a>	<a href="#">293024</a>	15828	2136	8	515	A kinase (PRKA) anchor protein 13 (Akap13) alternative variant bSep08, mRNA.
<a href="#">Akap13</a>	<a href="#">Akap13.cSep08</a>	<a href="#">293024</a>	1706	718	2	239	A kinase (PRKA) anchor protein 13 (Akap13) alternative variant cSep08, mRNA.
<a href="#">Akna</a>	<a href="#">Akna.bSep08</a>	<a href="#">362530</a>	5815	707	5	175	AT-hook transcription factor (Akna) alternative variant bSep08, mRNA.
<a href="#">Akr1a1</a>	<a href="#">Akr1a1.bSep08</a>	<a href="#">78959</a>	3661	601	5	168	aldo-keto reductase family 1, member A1 (aldehyde reductase) (Akr1a1) alternative variant bSep08, mRNA.
<a href="#">Akr1b1</a>	<a href="#">Akr1b1.bSep08</a>	<a href="#">24192</a>	3121	573	2	73	aldo-keto reductase family 1, member B1 (aldose reductase) (Akr1b1) alternative variant bSep08, mRNA.
<a href="#">Akr1b7</a>	<a href="#">Akr1b7.bSep08</a>	<a href="#">116463</a>	2005	849	1	125	aldo-keto reductase family 1, member B7 (Akr1b7) alternative variant bSep08, mRNA.
<a href="#">Akr1b8</a>	<a href="#">Akr1b8.bSep08</a>	<a href="#">286921</a>	3531	508	4	70	aldo-keto reductase family 1, member B8 (Akr1b8) alternative variant bSep08, mRNA.
<a href="#">Akr1b10</a>	<a href="#">Akr1b10.bSep08</a>	<a href="#">296972</a>	3949	372	3	82	aldo-keto reductase family 1, member B10 (aldose reductase) (Akr1b10) alternative variant bSep08, mRNA.
<a href="#">Akr1b10</a>	<a href="#">Akr1b10.cSep08</a>	<a href="#">296972</a>	4742	599	4	46	aldo-keto reductase family 1, member B10 (aldose reductase) (Akr1b10) alternative variant cSep08, mRNA.
<a href="#">Akr1c6</a>	<a href="#">Akr1c6.bSep08</a>	<a href="#">307092</a>	7625	763	1	55	aldo-keto reductase family 1, member C6 (Akr1c6) alternative variant bSep08, mRNA.
<a href="#">Akr1c12</a>	<a href="#">Akr1c12.aSep08</a>	<a href="#">361266</a>	33373	1329	11	370	aldo-keto reductase family 1, member C12 (Akr1c12) alternative variant aSep08, mRNA.
<a href="#">Akr1c12</a>	<a href="#">Akr1c12.bSep08</a>	<a href="#">361266</a>	8252	869	7	274	aldo-keto reductase family 1, member C12 (Akr1c12) alternative variant bSep08, mRNA.
<a href="#">Akr1c14</a>	<a href="#">Akr1c14.bSep08</a>	<a href="#">191574</a>	2534	392	3	130	aldo-keto reductase family 1, member C14 (Akr1c14) alternative variant bSep08, mRNA.
<a href="#">Akr1c14</a>	<a href="#">Akr1c14.cSep08</a>	<a href="#">191574</a>	2641	1189	2	34	aldo-keto reductase family 1, member C14 (3.8 kD) (Akr1c14) alternative variant cSep08, mRNA.

<a href="#">Akr1c18</a>	<a href="#">Akr1c18.bSep08</a>	<a href="#">171516</a>	13495	730	3	233	aldo-keto reductase family 1, member C18 (Akr1c18) alternative variant bSep08, mRNA.
<a href="#">Akr1c18</a>	<a href="#">Akr1c18.cSep08</a>	<a href="#">171516</a>	9170	627	2	196	aldo-keto reductase family 1, member C18 (Akr1c18) alternative variant cSep08, mRNA.
<a href="#">Akr1c21</a>	<a href="#">Akr1c21.bSep08</a>	<a href="#">291283</a>	39491	777		190	aldo-keto reductase family 1, member C21 (Akr1c21) alternative variant bSep08, mRNA.
<a href="#">Akr1cl1</a>	<a href="#">Akr1cl1.bSep08</a>	<a href="#">361267</a>	16687	768	6	228	aldo-keto reductase family 1, member C-like 1 (25.9 kD) (Akr1cl1) alternative variant bSep08, complete mRNA.
<a href="#">Akr1cl1</a>	<a href="#">Akr1cl1.cSep08</a>	<a href="#">361267</a>	14597	769	6	201	aldo-keto reductase family 1, member C-like 1 (23.1 kD) (Akr1cl1) alternative variant cSep08, mRNA.
<a href="#">Akr1cl1</a>	<a href="#">Akr1cl1.dSep08</a>	<a href="#">361267</a>	11826	1783	3	59	aldo-keto reductase family 1, member C-like 1 (Akr1cl1) alternative variant dSep08, mRNA.
<a href="#">Akr1e1</a>	<a href="#">Akr1e1.bSep08</a>	<a href="#">307091</a>	13787	651	8	176	aldo-keto reductase family 1, member E1 (Akr1e1) alternative variant bSep08, mRNA.
<a href="#">Akr1e1</a>	<a href="#">Akr1e1.cSep08</a>	<a href="#">307091</a>	4028	360	3	119	aldo-keto reductase family 1, member E1 (Akr1e1) alternative variant cSep08, mRNA.
<a href="#">Akr1e1</a>	<a href="#">Akr1e1.dSep08</a>	<a href="#">307091</a>	2391	660	3	31	aldo-keto reductase family 1, member E1 (3.8 kD) (Akr1e1) alternative variant dSep08, mRNA.
<a href="#">Akr7a3</a>	<a href="#">Akr7a3.bSep08</a>	<a href="#">26760</a>	1840	927	1	104	aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase) (11.0 kD) (Akr7a3) alternative variant bSep08, mRNA.
<a href="#">Akt1</a>	<a href="#">Akt1.bSep08</a>	<a href="#">24185</a>	15053	762	5	133	thymoma viral proto-oncogene 1 (Akt1) alternative variant bSep08, mRNA.
<a href="#">Akt1s1</a>	<a href="#">Akt1s1.bSep08</a>	<a href="#">292887</a>	3857	708	4	221	AKT1 substrate 1 (proline-rich) (Akt1s1) alternative variant bSep08, mRNA.
<a href="#">Akt1s1</a>	<a href="#">Akt1s1.cSep08</a>	<a href="#">292887</a>	3949	507	4	168	AKT1 substrate 1 (proline-rich) (Akt1s1) alternative variant cSep08, mRNA.
<a href="#">Akt2</a>	<a href="#">Akt2.bSep08</a>	<a href="#">25233</a>	42232	649	6	182	thymoma viral proto-oncogene 2 (Akt2) alternative variant bSep08, mRNA.
<a href="#">Akt2</a>	<a href="#">Akt2.cSep08</a>	<a href="#">25233</a>	25543	594	5	110	thymoma viral proto-oncogene 2 (Akt2) alternative variant cSep08, mRNA.
<a href="#">Alad</a>	<a href="#">Alad.bSep08</a>	<a href="#">25374</a>	5426	782	8	204	aminolevulinate, delta-, dehydratase (Alad) alternative variant bSep08, mRNA.
<a href="#">Alad</a>	<a href="#">Alad.cSep08</a>	<a href="#">25374</a>	1623	420	4	112	aminolevulinate, delta-, dehydratase (Alad) alternative variant cSep08, mRNA.
<a href="#">Alad</a>	<a href="#">Alad.dSep08</a>	<a href="#">25374</a>	2112	936	4	81	aminolevulinate, delta-, dehydratase (9.1 kD) (Alad) alternative variant dSep08, mRNA.
<a href="#">Alad</a>	<a href="#">Alad.eSep08</a>	<a href="#">25374</a>	4911	393	2	27	aminolevulinate, delta-, dehydratase (Alad) alternative variant eSep08, mRNA.
<a href="#">Alcam</a>	<a href="#">Alcam.bSep08</a>	<a href="#">79559</a>	33689	3729	10	351	activated leukocyte cell adhesion molecule (Alcam) alternative variant bSep08, mRNA.
<a href="#">Alcam</a>	<a href="#">Alcam.cSep08</a>	<a href="#">79559</a>	14754	1068	7	240	activated leukocyte cell adhesion molecule (26.6 kD) (Alcam) alternative variant cSep08, mRNA.
<a href="#">Alcam</a>	<a href="#">Alcam.dSep08</a>	<a href="#">79559</a>	206151	1289	4	120	activated leukocyte cell adhesion molecule (Alcam) alternative variant dSep08, mRNA.
<a href="#">Aldh1a1</a>	<a href="#">Aldh1a1.bSep08</a>	<a href="#">24188</a>	38926	1403	8	304	aldehyde dehydrogenase family 1, subfamily A1 (32.6 kD) (Aldh1a1) alternative variant bSep08, mRNA.

<a href="#">Aldh1a1</a>	<a href="#">Aldh1a1.cSep08</a>	<a href="#">24188</a>	31859	681	6	81	aldehyde dehydrogenase family 1, subfamily A1 (8.9 kD) (Aldh1a1) alternative variant cSep08, mRNA.
<a href="#">Aldh1a1</a>	<a href="#">Aldh1a1.dSep08</a>	<a href="#">24188</a>	2160	418	1	43	aldehyde dehydrogenase family 1, subfamily A1 (Aldh1a1) alternative variant dSep08, mRNA.
<a href="#">Aldh1a3</a>	<a href="#">Aldh1a3.bSep08</a>	<a href="#">266603</a>	10085	693	6	158	aldehyde dehydrogenase family 1, subfamily A3 (Aldh1a3) alternative variant bSep08, mRNA.
<a href="#">Aldh1a7</a>	<a href="#">Aldh1a7.aSep08</a>	<a href="#">29651</a>	28678	620		137	aldehyde dehydrogenase family 1, subfamily A7 (Aldh1a7) alternative variant aSep08, mRNA.
<a href="#">Aldh1l2</a>	<a href="#">Aldh1l2.aSep08</a>	<a href="#">299699</a>	5595	399	3	133	aldehyde dehydrogenase 1 family, member L2 (Aldh1l2) alternative variant aSep08, mRNA.
<a href="#">Aldh1l2</a>	<a href="#">Aldh1l2.bSep08</a>	<a href="#">299699</a>	3982	1234	2	56	aldehyde dehydrogenase 1 family, member L2 (6.2 kD) (Aldh1l2) alternative variant bSep08, mRNA.
<a href="#">Aldh2</a>	<a href="#">Aldh2.aSep08</a>	<a href="#">29539</a>	33779	2257	7	552	aldehyde dehydrogenase 2, mitochondrial (Aldh2) alternative variant aSep08, mRNA.
<a href="#">Aldh2</a>	<a href="#">Aldh2.cSep08</a>	<a href="#">29539</a>	25872	800	6	253	aldehyde dehydrogenase 2, mitochondrial (Aldh2) alternative variant cSep08, mRNA.
<a href="#">Aldh2</a>	<a href="#">Aldh2.dSep08</a>	<a href="#">29539</a>	4526	492	2	108	aldehyde dehydrogenase 2, mitochondrial (Aldh2) alternative variant dSep08, mRNA.
<a href="#">Aldh3a2</a>	<a href="#">Aldh3a2.bSep08</a>	<a href="#">65183</a>	2182	743	1	168	aldehyde dehydrogenase family 3, subfamily A2 (18.3 kD) (Aldh3a2) alternative variant bSep08, mRNA.
<a href="#">Aldh3b2</a>	<a href="#">Aldh3b2.aSep08</a>	<a href="#">688800</a>	39907	1621	7	479	aldehyde dehydrogenase 3 family, member B2 (52.6 kD) (Aldh3b2) alternative variant aSep08, mRNA.
<a href="#">Aldh3b2</a>	<a href="#">Aldh3b2.bSep08</a>	<a href="#">688800</a>	916	382	1	73	aldehyde dehydrogenase 3 family, member B2 (Aldh3b2) alternative variant bSep08, mRNA.
<a href="#">Aldh5a1</a>	<a href="#">Aldh5a1.aSep08</a>	<a href="#">291133</a>	23422	2520	10	614	aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase) (66.2 kD) (Aldh5a1) alternative variant aSep08, mRNA.
<a href="#">Aldh6a1</a>	<a href="#">Aldh6a1.bSep08</a>	<a href="#">81708</a>	9763	415	5	118	aldehyde dehydrogenase family 6, subfamily A1 (Aldh6a1) alternative variant bSep08, mRNA.
<a href="#">Aldh7a1</a>	<a href="#">Aldh7a1.aSep08</a>	<a href="#">291450</a>	32262	1894	3	539	aldehyde dehydrogenase family 7, member A1 (58.7 kD) (Aldh7a1) alternative variant aSep08, mRNA.
<a href="#">Aldh7a1</a>	<a href="#">Aldh7a1.bSep08</a>	<a href="#">291450</a>	6304	537	1	177	aldehyde dehydrogenase family 7, member A1 (Aldh7a1) alternative variant bSep08, mRNA.
<a href="#">Aldh8a1</a>	<a href="#">Aldh8a1.aSep08</a>	<a href="#">685750</a>	19384	1767		352	aldehyde dehydrogenase 8 family, member A1 (Aldh8a1) mRNA.
<a href="#">Aldh16a1</a>	<a href="#">Aldh16a1.bSep08</a>	<a href="#">361571</a>	7589	528	5	141	aldehyde dehydrogenase 16 family, member A1 (Aldh16a1) alternative variant bSep08, mRNA.
<a href="#">Aldh16a1</a>	<a href="#">Aldh16a1.cSep08</a>	<a href="#">361571</a>	6241	383	3	127	aldehyde dehydrogenase 16 family, member A1 (Aldh16a1) alternative variant cSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.aSep08</a>	<a href="#">24189</a>	3762	1223	9	391	aldolase A, fructose-bisphosphate (Aldoa) alternative variant aSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.cSep08</a>	<a href="#">24189</a>	4770	1077	8	276	aldolase A, fructose-bisphosphate (30.0 kD) (Aldoa) alternative variant cSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.dSep08</a>	<a href="#">24189</a>	4208	941	7	250	aldolase A, fructose-bisphosphate (Aldoa) alternative variant dSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.eSep08</a>	<a href="#">24189</a>	2884	839	7	246	aldolase A, fructose-bisphosphate (Aldoa) alternative variant eSep08, mRNA.

<a href="#">Aldoa</a>	<a href="#">Aldoa.fSep08</a>	<a href="#">24189</a>	3872	1139	6	243	aldolase A, fructose-bisphosphate (26.2 kD) (Aldoa) alternative variant fSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.gSep08</a>	<a href="#">24189</a>	2456	786	7	210	aldolase A, fructose-bisphosphate (Aldoa) alternative variant gSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.hSep08</a>	<a href="#">24189</a>	4247	789	6	208	aldolase A, fructose-bisphosphate (Aldoa) alternative variant hSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.iSep08</a>	<a href="#">24189</a>	2849	742	5	176	aldolase A, fructose-bisphosphate (Aldoa) alternative variant iSep08, mRNA.
<a href="#">Aldoa</a>	<a href="#">Aldoa.jSep08</a>	<a href="#">24189</a>	2035	765	2	83	aldolase A, fructose-bisphosphate (9.3 kD) (Aldoa) alternative variant jSep08, mRNA.
<a href="#">Aldob</a>	<a href="#">Aldob.bSep08</a>	<a href="#">24190</a>	11154	903	8	268	aldolase B fructose-bisphosphate (29.5 kD) (Aldob) alternative variant bSep08, mRNA.
<a href="#">Aldob</a>	<a href="#">Aldob.cSep08</a>	<a href="#">24190</a>	9712	715	6	218	aldolase B fructose-bisphosphate (Aldob) alternative variant cSep08, mRNA.
<a href="#">Aldob</a>	<a href="#">Aldob.dSep08</a>	<a href="#">24190</a>	6709	686	6	126	aldolase B fructose-bisphosphate (Aldob) alternative variant dSep08, mRNA.
<a href="#">Aldob</a>	<a href="#">Aldob.eSep08</a>	<a href="#">24190</a>	5742	1265	2	32	aldolase B fructose-bisphosphate (Aldob) alternative variant eSep08, mRNA.
<a href="#">Aldoc</a>	<a href="#">Aldoc.bSep08</a>	<a href="#">24191</a>	2280	419	3	69	aldolase C, fructose-bisphosphate (Aldoc) alternative variant bSep08, mRNA.
<a href="#">Aldoc</a>	<a href="#">Aldoc.dSep08</a>	<a href="#">24191</a>	1502	352	2	39	aldolase C, fructose-bisphosphate (4.1 kD) (Aldoc) alternative variant dSep08, mRNA.
<a href="#">Aldo_ket_red.0</a>	<a href="#">Aldo_ket_red.0.bSep08</a>		7104	619	1	205	potassium channel beta (Aldo_ket_red.0) alternative variant bSep08, mRNA.
<a href="#">Alg1</a>	<a href="#">Alg1.aSep08</a>	<a href="#">360475</a>	10707	1747	13	462	asparagine-linked glycosylation 1 homolog (yeast, beta-1,4-mannosyltransferase) (Alg1) alternative variant aSep08, mRNA.
<a href="#">Alg1</a>	<a href="#">Alg1.bSep08</a>	<a href="#">360475</a>	5928	709	8	180	asparagine-linked glycosylation 1 homolog (yeast, beta-1,4-mannosyltransferase) (Alg1) alternative variant bSep08, mRNA.
<a href="#">Alg1</a>	<a href="#">Alg1.cSep08</a>	<a href="#">360475</a>	1043	573	2	100	asparagine-linked glycosylation 1 homolog (yeast, beta-1,4-mannosyltransferase) (Alg1) alternative variant cSep08, mRNA.
<a href="#">Alg2</a>	<a href="#">Alg2.aSep08</a>	<a href="#">313231</a>	4567	3032		415	asparagine-linked glycosylation 2 homolog (yeast, alpha-1,3-mannosyltransferase) (47.3 kD) (Alg2) mRNA.
<a href="#">Alg3</a>	<a href="#">Alg3.bSep08</a>	<a href="#">287983</a>	2258	574	1	141	asparagine-linked glycosylation 3 homolog (yeast, alpha-1,3-mannosyltransferase) (Alg3) alternative variant bSep08, mRNA.
<a href="#">Alg5</a>	<a href="#">Alg5.bSep08</a>	<a href="#">295051</a>	8101	822	7	240	asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase) (Alg5) alternative variant bSep08, mRNA.
<a href="#">Alg5</a>	<a href="#">Alg5.cSep08</a>	<a href="#">295051</a>	4942	692	5	164	asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase) (19.0 kD) (Alg5) alternative variant cSep08, mRNA.
<a href="#">Alg5</a>	<a href="#">Alg5.dSep08</a>	<a href="#">295051</a>	3865	497	3	141	asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase) (Alg5) alternative variant dSep08, mRNA.

<a href="#">Alg5</a>	<a href="#">Alg5.fSep08</a>	<a href="#">295051</a>	4162	532	3	81	asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase) (Alg5) alternative variant fSep08, mRNA.
<a href="#">Alg6</a>	<a href="#">Alg6.aSep08</a>	<a href="#">362547</a>	37082	2651	10	227	asparagine-linked glycosylation 6 (25.9 kD) (Alg6) alternative variant aSep08, mRNA.
<a href="#">Alg6</a>	<a href="#">Alg6.bSep08</a>	<a href="#">362547</a>	21030	840	7	203	asparagine-linked glycosylation 6 homolog (Alg6) alternative variant bSep08, mRNA.
<a href="#">Alg6</a>	<a href="#">Alg6.cSep08</a>	<a href="#">362547</a>	10700	1574	2	104	asparagine-linked glycosylation 6 homolog (Alg6) alternative variant cSep08, mRNA.
<a href="#">Alg8</a>	<a href="#">Alg8.bSep08</a>	<a href="#">293129</a>	9239	739	2	119	asparagine-linked glycosylation 8 homolog (yeast, alpha-1,3-glucosyltransferase) (13.0 kD) (Alg8) alternative variant bSep08, mRNA.
<a href="#">Alg9</a>	<a href="#">Alg9.bSep08</a>	<a href="#">367083</a>	36252	756	1	251	asparagine-linked glycosylation 9 homolog (yeast, alpha 1,2-mannosyltransferase) (Alg9) alternative variant bSep08, mRNA.
<a href="#">Alg9</a>	<a href="#">Alg9.cSep08</a>	<a href="#">367083</a>	32159	1184	1	177	asparagine-linked glycosylation 9 homolog (yeast, alpha 1,2-mannosyltransferase) (Alg9) alternative variant cSep08, mRNA.
<a href="#">Alg11</a>	<a href="#">Alg11.bSep08</a>	<a href="#">361174</a>	7255	1729	5	401	asparagine-linked glycosylation 11 homolog (yeast, alpha-1,2-mannosyltransferase) (45.2 kD) (Alg11) alternative variant bSep08, complete mRNA.
<a href="#">Alg11</a>	<a href="#">Alg11.cSep08</a>	<a href="#">361174</a>	4382	734	3	148	asparagine-linked glycosylation 11 homolog (yeast, alpha-1,2-mannosyltransferase) (Alg11) alternative variant cSep08, mRNA.
<a href="#">Alg11</a>	<a href="#">Alg11.eSep08</a>	<a href="#">361174</a>	990	117	2	22	asparagine-linked glycosylation 11 homolog (yeast, alpha-1,2-mannosyltransferase) (Alg11) alternative variant eSep08, mRNA.
<a href="#">Alg12</a>	<a href="#">Alg12.aSep08</a>	<a href="#">315212</a>	14185	2038	10	488	asparagine-linked glycosylation 12 homolog (yeast, alpha-1,6-mannosyltransferase) (54.7 kD) (Alg12) alternative variant aSep08, mRNA.
<a href="#">Alg14</a>	<a href="#">Alg14.bSep08</a>	<a href="#">362031</a>	89052	824	2	83	asparagine-linked glycosylation 14 homolog ( <i>S. cerevisiae</i> ) (9.7 kD) (Alg14) alternative variant bSep08, mRNA.
<a href="#">Alkbh</a>	<a href="#">Alkbh.bSep08</a>	<a href="#">362766</a>	4098	458	2	89	alkB, alkylation repair homolog ( <i>E. coli</i> ) (Alkbh) alternative variant bSep08, mRNA.
<a href="#">Alkbh2</a>	<a href="#">Alkbh2.aSep08</a>	<a href="#">304578</a>	4623	1796	2	591	alkB, alkylation repair homolog 2 ( <i>E. coli</i> ) (Alkbh2) alternative variant aSep08, mRNA.
<a href="#">Alkbh3</a>	<a href="#">Alkbh3.bSep08</a>	<a href="#">362169</a>	14905	802	8	205	alkB, alkylation repair homolog 3 ( <i>E. coli</i> ) (Alkbh3) alternative variant bSep08, mRNA.
<a href="#">Alkbh3</a>	<a href="#">Alkbh3.cSep08</a>	<a href="#">362169</a>	17775	807	3	187	alkB, alkylation repair homolog 3 ( <i>E. coli</i> ) (Alkbh3) alternative variant cSep08, mRNA.
<a href="#">Alkbh3</a>	<a href="#">Alkbh3.dSep08</a>	<a href="#">362169</a>	24528	766	4	154	alkB, alkylation repair homolog 3 ( <i>E. coli</i> ) (17.7 kD) (Alkbh3) alternative variant dSep08, mRNA.
<a href="#">Alkbh3</a>	<a href="#">Alkbh3.eSep08</a>	<a href="#">362169</a>	32192	879	8	143	alkB, alkylation repair homolog 3 ( <i>E. coli</i> ) (Alkbh3) alternative variant eSep08, mRNA.
<a href="#">Alkbh5</a>	<a href="#">Alkbh5.aSep08</a>	<a href="#">303193</a>	2254	1893		104	alkB, alkylation repair homolog 5 ( <i>E. coli</i> ) (Alkbh5) mRNA.
<a href="#">Alkbh6</a>	<a href="#">Alkbh6.bSep08</a>	<a href="#">292780</a>	5370	897	7	177	alkB, alkylation repair homolog 6 ( <i>E. coli</i> ) (19.7 kD) (Alkbh6) alternative variant bSep08, complete mRNA.

<a href="#">Alkbh6</a>	<a href="#">Alkbh6.cSep08</a>	<a href="#">292780</a>	4582	690	5	163	alkB, alkylation repair homolog 6 (E. coli) (17.8 kD) (Alkbh6) alternative variant cSep08, mRNA.
<a href="#">Alkbh6</a>	<a href="#">Alkbh6.dSep08</a>	<a href="#">292780</a>	5334	1442	5	85	alkB, alkylation repair homolog 6 (E. coli) (Alkbh6) alternative variant dSep08, mRNA.
<a href="#">Alkbh6</a>	<a href="#">Alkbh6.eSep08</a>	<a href="#">292780</a>	1945	639	4	60	alkB, alkylation repair homolog 6 (E. coli) (Alkbh6) alternative variant eSep08, mRNA.
<a href="#">Alkbh7</a>	<a href="#">Alkbh7.bSep08</a>	<a href="#">679944</a>	954	680	3	79	alkB, alkylation repair homolog 7 (E. coli) (9.0 kD) (Alkbh7) alternative variant bSep08, mRNA.
<a href="#">Alkbh7</a>	<a href="#">Alkbh7.cSep08</a>	<a href="#">679944</a>	2050	609	2	77	alkB, alkylation repair homolog 7 (E. coli) (Alkbh7) alternative variant cSep08, mRNA.
<a href="#">Alkbh8</a>	<a href="#">Alkbh8.aSep08</a>	<a href="#">366783</a>	41175	730	3	243	alkB, alkylation repair homolog 8 (E. coli) (Alkbh8) alternative variant aSep08, mRNA.
<a href="#">Alkbh8</a>	<a href="#">Alkbh8.bSep08</a>	<a href="#">366783</a>	82299	740	3	176	alkB, alkylation repair homolog 8 (E. coli) (Alkbh8) alternative variant bSep08, mRNA.
<a href="#">Allc</a>	<a href="#">Allc.bSep08</a>	<a href="#">246758</a>	1359	400		104	allantoicase (Allc) alternative variant bSep08, mRNA.
<a href="#">Alms1</a>	<a href="#">Alms1.bSep08</a>	<a href="#">297408</a>	22208	1717	5	529	alstrom syndrome 1 homolog (human) (Alms1) alternative variant bSep08, mRNA.
<a href="#">Alms1</a>	<a href="#">Alms1.cSep08</a>	<a href="#">297408</a>	15109	636	4	212	alstrom syndrome 1 homolog (human) (Alms1) alternative variant cSep08, mRNA.
<a href="#">Alms1</a>	<a href="#">Alms1.dSep08</a>	<a href="#">297408</a>	5941	601	5	162	alstrom syndrome 1 homolog (human) (Alms1) alternative variant dSep08, mRNA.
<a href="#">Alms1</a>	<a href="#">Alms1.eSep08</a>	<a href="#">297408</a>	1690	604	2	124	alstrom syndrome 1 homolog (human) (Alms1) alternative variant eSep08, mRNA.
<a href="#">Alox12</a>	<a href="#">Alox12.bSep08</a>	<a href="#">287454</a>	9640	1825	12	514	arachidonate 12-lipoxygenase (58.4 kD) (Alox12) alternative variant bSep08, complete mRNA.
<a href="#">Alox15</a>	<a href="#">Alox15.bSep08</a>	<a href="#">81639</a>	1663	346	4	76	arachidonate 15-lipoxygenase (Alox15) alternative variant bSep08, mRNA.
<a href="#">Aloxe3</a>	<a href="#">Aloxe3.bSep08</a>	<a href="#">287424</a>	1795	585	3	133	arachidonate lipoxygenase 3 (Aloxe3) alternative variant bSep08, mRNA.
<a href="#">Aloxe3</a>	<a href="#">Aloxe3.cSep08</a>	<a href="#">287424</a>	506	424	2	97	arachidonate lipoxygenase 3 (Aloxe3) alternative variant cSep08, mRNA.
<a href="#">Aloxe3</a>	<a href="#">Aloxe3.dSep08</a>	<a href="#">287424</a>	1655	193	2	64	arachidonate lipoxygenase 3 (Aloxe3) alternative variant dSep08, mRNA.
<a href="#">Alpha-amylase_C.0</a>	<a href="#">Alpha-amylase_C.0.aSep08</a>		27561	1179		164	glucan branching enzyme 1 (Alpha-amylase_C.0) mRNA.
<a href="#">Alpha-amylase_C.1</a>	<a href="#">Alpha-amylase_C.1.aSep08</a>		1986	348		104	alpha-amylase (Alpha-amylase_C.1) mRNA.
<a href="#">Alpha_adaptinC2.0</a>	<a href="#">Alpha_adaptinC2.0.aSep08</a>		7574	2843	19	329	CRA b (36.3 kD) (Alpha_adaptinC2.0) alternative variant aSep08, mRNA.
<a href="#">Alpha_adaptinC2.0</a>	<a href="#">Alpha_adaptinC2.0.bSep08</a>		995	695	3	156	adaptor-related protein complex 1 gamma (Alpha_adaptinC2.0) alternative variant bSep08, mRNA.
<a href="#">Alpha_adaptinC2.0</a>	<a href="#">Alpha_adaptinC2.0.cSep08</a>		1188	828	3	119	CRA a (13.3 kD) (Alpha_adaptinC2.0) alternative variant cSep08, mRNA.
<a href="#">Alpha_adaptinC2.0</a>	<a href="#">Alpha_adaptinC2.0.dSep08</a>		1554	775	3	101	protein complex 1 gamma precursor (11.1 kD) (Alpha_adaptinC2.0) alternative variant dSep08, mRNA.
<a href="#">Alpk1</a>	<a href="#">Alpk1.aSep08</a>	<a href="#">310879</a>	558	450		112	alpha-kinase 1 (Alpk1) mRNA.
<a href="#">Alpk2</a>	<a href="#">Alpk2.aSep08</a>	<a href="#">498875</a>	16050	785		261	alpha-kinase 2 (Alpk2) mRNA.

<a href="#">Alpk3</a>	<a href="#">Alpk3.aSep08</a>	<a href="#">365298</a>	7388	1035		345	alpha-kinase 3 (Alpk3) mRNA.
<a href="#">Alpl</a>	<a href="#">Alpl.bSep08</a>	<a href="#">25586</a>	11408	609	3	106	alkaline phosphatase, liver/bone/kidney (Alpl) alternative variant bSep08, mRNA.
<a href="#">Alpl</a>	<a href="#">Alpl.cSep08</a>	<a href="#">25586</a>	3227	396	3	79	alkaline phosphatase, liver/bone/kidney (Alpl) alternative variant cSep08, mRNA.
<a href="#">Alpl</a>	<a href="#">Alpl.dSep08</a>	<a href="#">25586</a>	13489	438	4	63	alkaline phosphatase, liver/bone/kidney (Alpl) alternative variant dSep08, mRNA.
<a href="#">Alpl</a>	<a href="#">Alpl.eSep08</a>	<a href="#">25586</a>	15864	391	3	53	alkaline phosphatase, liver/bone/kidney (Alpl) alternative variant eSep08, mRNA.
<a href="#">Als2</a>	<a href="#">Als2.bSep08</a>	<a href="#">363235</a>	14508	1783	7	403	amyotrophic lateral sclerosis 2 (juvenile) homolog (human) (Als2) alternative variant bSep08, mRNA.
<a href="#">Als2cl</a>	<a href="#">Als2cl.aSep08</a>	<a href="#">316017</a>	5615	1196		375	ALS2 C-terminal like (Als2cl) mRNA.
<a href="#">Als2cr2</a>	<a href="#">Als2cr2.aSep08</a>	<a href="#">501146</a>	18911	2359	11	418	protein kinase and tyrosine protein kinase (46.9 kD) (Als2cr2) mRNA.
<a href="#">Als2cr4</a>	<a href="#">Als2cr4.aSep08</a>	<a href="#">316412</a>	17769	908	4	288	amyotrophic lateral sclerosis 2 region candidate 4 (Als2cr4) alternative variant aSep08, mRNA.
<a href="#">Als2cr4</a>	<a href="#">Als2cr4.bSep08</a>	<a href="#">316412</a>	13475	752	1	130	amyotrophic lateral sclerosis 2 region candidate 4 (14.4 kD) (Als2cr4) alternative variant bSep08, complete mRNA.
<a href="#">Als2cr12</a>	<a href="#">Als2cr12.bSep08</a>	<a href="#">316413</a>	6821	1776	1	19	putative protein (2.0 kD) (Als2cr12) alternative variant bSep08, complete mRNA.
<a href="#">Amacr</a>	<a href="#">Amacr.bSep08</a>	<a href="#">25284</a>	11464	741	4	196	alpha-methylacyl-CoA racemase (20.9 kD) (Amacr) alternative variant bSep08, mRNA.
<a href="#">Amacr</a>	<a href="#">Amacr.cSep08</a>	<a href="#">25284</a>	5158	960	2	158	alpha-methylacyl-CoA racemase (Amacr) alternative variant cSep08, mRNA.
<a href="#">Ambn</a>	<a href="#">Ambn.bSep08</a>	<a href="#">25376</a>	6418	377	1	125	ameloblastin (Ambn) alternative variant bSep08, mRNA.
<a href="#">Ambp</a>	<a href="#">Ambp.bSep08</a>	<a href="#">25377</a>	8940	895	8	263	alpha 1 microglobulin/bikunin (29.8 kD) (Ambp) alternative variant bSep08, mRNA.
<a href="#">Ambp</a>	<a href="#">Ambp.cSep08</a>	<a href="#">25377</a>	3489	709	3	100	alpha 1 microglobulin/bikunin (Ambp) alternative variant cSep08, mRNA.
<a href="#">Ambp</a>	<a href="#">Ambp.dSep08</a>	<a href="#">25377</a>	454	276	2	37	alpha 1 microglobulin/bikunin (Ambp) alternative variant dSep08, mRNA.
<a href="#">Amd1</a>	<a href="#">Amd1.bSep08</a>	<a href="#">81640</a>	12339	701	6	233	s-adenosylmethionine decarboxylase (Amd1) alternative variant bSep08, mRNA.
<a href="#">Amd1</a>	<a href="#">Amd1.cSep08</a>	<a href="#">81640</a>	11919	697	6	167	s-adenosylmethionine decarboxylase (Amd1) alternative variant cSep08, mRNA.
<a href="#">Amd1</a>	<a href="#">Amd1.dSep08</a>	<a href="#">81640</a>	1710	1610	2	110	S-adenosylmethionine decarboxylase (12.4 kD) (Amd1) alternative variant dSep08, mRNA.
<a href="#">Amd1</a>	<a href="#">Amd1.eSep08</a>	<a href="#">81640</a>	1871	902	2	87	S-adenosylmethionine decarboxylase (10.2 kD) (Amd1) alternative variant eSep08, mRNA.
<a href="#">Amdhd1</a>	<a href="#">Amdhd1.aSep08</a>	<a href="#">299735</a>	13882	1167	6	388	amidohydrolase 1 and amidohydrolase 3 (Amdhd1) alternative variant aSep08, mRNA.
<a href="#">Amdhd1</a>	<a href="#">Amdhd1.bSep08</a>	<a href="#">299735</a>	3278	479	1	154	putative protein of ancient origin (Amdhd1) alternative variant bSep08, mRNA.
<a href="#">Amdhd2</a>	<a href="#">Amdhd2.bSep08</a>	<a href="#">302972</a>	8496	1551	10	289	putative protein of ancient origin (31.1 kD) (Amdhd2) alternative variant bSep08, complete mRNA.
<a href="#">Amdhd2</a>	<a href="#">Amdhd2.cSep08</a>	<a href="#">302972</a>	6125	680	6	226	putative protein of ancient origin (Amdhd2) alternative variant cSep08, mRNA.



<a href="#">Amdhd2</a>	<a href="#">Amdhd2.dSep08</a>	<a href="#">302972</a>	6054	649	5	169	putative protein of ancient origin (Amdhd2) alternative variant dSep08, mRNA.
<a href="#">Amdhd2</a>	<a href="#">Amdhd2.fSep08</a>	<a href="#">302972</a>	1191	657	2	32	putative protein (3.7 kD) (Amdhd2) alternative variant fSep08, mRNA.
<a href="#">Amelx</a>	<a href="#">Amelx.bSep08</a>	<a href="#">29160</a>	5426	504	1	168	amelogenin (Amelx) alternative variant bSep08, mRNA.
<a href="#">Amelx</a>	<a href="#">Amelx.cSep08</a>	<a href="#">29160</a>	5331	481	1	160	amelogenin (Amelx) alternative variant cSep08, mRNA.
<a href="#">Amfr</a>	<a href="#">Amfr.aSep08</a>	<a href="#">361367</a>	33725	2933	8	763	autocrine motility factor receptor (Amfr) alternative variant aSep08, mRNA.
<a href="#">Amfr</a>	<a href="#">Amfr.bSep08</a>	<a href="#">361367</a>	15254	3204	7	539	autocrine motility factor receptor (Amfr) alternative variant bSep08, mRNA.
<a href="#">Amfr</a>	<a href="#">Amfr.cSep08</a>	<a href="#">361367</a>	8918	1585	5	527	autocrine motility factor receptor (Amfr) alternative variant cSep08, mRNA.
<a href="#">Amfr</a>	<a href="#">Amfr.dSep08</a>	<a href="#">361367</a>	10133	1561	6	520	autocrine motility factor receptor (Amfr) alternative variant dSep08, mRNA.
<a href="#">Amfr</a>	<a href="#">Amfr.eSep08</a>	<a href="#">361367</a>	1628	904	2	198	autocrine motility factor receptor (22.6 kD) (Amfr) alternative variant eSep08, mRNA.
<a href="#">Amhr2</a>	<a href="#">Amhr2.bSep08</a>	<a href="#">29530</a>	403	287	2	95	anti-Mullerian hormone type 2 receptor (Amhr2) alternative variant bSep08, mRNA.
<a href="#">Amica1</a>	<a href="#">Amica1.aSep08</a>	<a href="#">315610</a>	7064	715		123	adhesion molecule, interacts with CXADR antigen 1 (Amica1) mRNA.
<a href="#">Amigo2</a>	<a href="#">Amigo2.bSep08</a>	<a href="#">300186</a>	896	514	2	89	adhesion molecule with Ig like domain 2 (Amigo2) alternative variant bSep08, mRNA.
<a href="#">Amino_oxidase.0</a>	<a href="#">Amino_oxidase.0.bSep08</a>		1666	739	3	245	polyamine oxidase (Amino_oxidase.0) alternative variant bSep08, mRNA.
<a href="#">Amino_oxidase.0</a>	<a href="#">Amino_oxidase.0.cSep08</a>		2027	543	2	180	polyamine oxidase CRA b (Amino_oxidase.0) alternative variant cSep08, mRNA.
<a href="#">Ammecr1l</a>	<a href="#">Ammecr1l.bSep08</a>	<a href="#">307526</a>	13468	779	4	259	AMME chromosomal region gene 1-like (Ammecr1l) alternative variant bSep08, mRNA.
<a href="#">Ammecr1l</a>	<a href="#">Ammecr1l.cSep08</a>	<a href="#">307526</a>	11833	1581	3	209	AMME chromosomal region gene 1-like (22.6 kD) (Ammecr1l) alternative variant cSep08, mRNA.
<a href="#">Ammecr1l</a>	<a href="#">Ammecr1l.dSep08</a>	<a href="#">307526</a>	11735	507	3	103	AMME chromosomal region gene 1-like (Ammecr1l) alternative variant dSep08, mRNA.
<a href="#">Ammecr1l</a>	<a href="#">Ammecr1l.eSep08</a>	<a href="#">307526</a>	2071	445	3	34	AMME chromosomal region gene 1-like (Ammecr1l) alternative variant eSep08, mRNA.
<a href="#">Amn</a>	<a href="#">Amn.bSep08</a>	<a href="#">314459</a>	647	476	1	115	amnionless (Amn) alternative variant bSep08, mRNA.
<a href="#">Amot</a>	<a href="#">Amot.aSep08</a>	<a href="#">300289</a>	24995	773	1	166	angiominin (Amot) alternative variant aSep08, mRNA.
<a href="#">Amot</a>	<a href="#">Amot.bSep08</a>	<a href="#">300289</a>	17191	371	2	72	angiominin (Amot) alternative variant bSep08, mRNA.
<a href="#">Amotl1</a>	<a href="#">Amotl1.bSep08</a>	<a href="#">315430</a>	53283	429	4	142	angiominin-like 1 (Amotl1) alternative variant bSep08, mRNA.
<a href="#">Amotl1</a>	<a href="#">Amotl1.cSep08</a>	<a href="#">315430</a>	22849	509	3	97	angiominin-like 1 (Amotl1) alternative variant cSep08, mRNA.
<a href="#">Amotl2</a>	<a href="#">Amotl2.aSep08</a>	<a href="#">65157</a>	9350	2998	8	440	angiominin like 2 (Amotl2) alternative variant aSep08, mRNA.
<a href="#">Amotl2</a>	<a href="#">Amotl2.bSep08</a>	<a href="#">65157</a>	1380	748	2	104	angiominin like 2 (Amotl2) alternative variant bSep08, mRNA.

<a href="#">Ampd1</a>	<a href="#">Ampd1.bSep08</a>	<a href="#">25028</a>	4862	698	1	98	adenosine monophosphate deaminase 1 (isoform M) (Ampd1) alternative variant bSep08, mRNA.
<a href="#">Ampd2</a>	<a href="#">Ampd2.bSep08</a>	<a href="#">362015</a>	993	634	3	118	adenosine monophosphate deaminase 2 (isoform L) (Ampd2) alternative variant bSep08, mRNA.
<a href="#">Ampd2</a>	<a href="#">Ampd2.cSep08</a>	<a href="#">362015</a>	5021	775	3	117	adenosine monophosphate deaminase 2 (isoform L) (12.9 kD) (Ampd2) alternative variant cSep08, mRNA.
<a href="#">Ampd2</a>	<a href="#">Ampd2.eSep08</a>	<a href="#">362015</a>	2351	414	3	60	adenosine monophosphate deaminase 2 (isoform L) (Ampd2) alternative variant eSep08, mRNA.
<a href="#">Ampd3</a>	<a href="#">Ampd3.bSep08</a>	<a href="#">25095</a>	6040	601	5	117	adenosine monophosphate deaminase 3 (Ampd3) alternative variant bSep08, mRNA.
<a href="#">Ampd3</a>	<a href="#">Ampd3.dSep08</a>	<a href="#">25095</a>	2100	425	2	45	adenosine monophosphate deaminase 3 (Ampd3) alternative variant dSep08, mRNA.
<a href="#">Amt</a>	<a href="#">Amt.bSep08</a>	<a href="#">306586</a>	1510	753	1	126	aminomethyltransferase (glycine cleavage system protein T) (Amt) alternative variant bSep08, mRNA.
<a href="#">Amy2</a>	<a href="#">Amy2.aSep08</a>	<a href="#">497039</a>	8566	974		324	amylase 2, pancreatic (Amy2) mRNA.
<a href="#">Amy2-3</a>	<a href="#">Amy2-3.aSep08</a>	<a href="#">365914</a>	4192	571		190	amylase 2-3, pancreatic (Amy2-3) mRNA.
<a href="#">Amy2-4</a>	<a href="#">Amy2-4.aSep08</a>	<a href="#">499693</a>	2133	519		94	amylase 2-4, pancreatic (9.8 kD) (Amy2-4) mRNA.
<a href="#">Amz2</a>	<a href="#">Amz2.bSep08</a>	<a href="#">360650</a>	1516	449	3	135	archaelysin family metallopeptidase 2 (Amz2) alternative variant bSep08, mRNA.
<a href="#">Amz2</a>	<a href="#">Amz2.cSep08</a>	<a href="#">360650</a>	2691	615	6	117	archaelysin family metallopeptidase 2 (Amz2) alternative variant cSep08, mRNA.
<a href="#">Amz2</a>	<a href="#">Amz2.dSep08</a>	<a href="#">360650</a>	2603	615	5	71	archaelysin family metallopeptidase 2 (Amz2) alternative variant dSep08, mRNA.
<a href="#">Anapc1</a>	<a href="#">Anapc1.bSep08</a>	<a href="#">311412</a>	5083	580	5	193	anaphase promoting complex subunit 1 (Anapc1) alternative variant bSep08, mRNA.
<a href="#">Anapc2</a>	<a href="#">Anapc2.bSep08</a>	<a href="#">296558</a>	6412	979	1	288	anaphase promoting complex subunit 2 (Anapc2) alternative variant bSep08, mRNA.
<a href="#">Anapc4</a>	<a href="#">Anapc4.bSep08</a>	<a href="#">305420</a>	9933	890	8	296	anaphase promoting complex (Anapc4) alternative variant bSep08, mRNA.
<a href="#">Anapc4</a>	<a href="#">Anapc4.cSep08</a>	<a href="#">305420</a>	1125	726	2	103	anaphase promoting complex (11.7 kD) (Anapc4) alternative variant cSep08, mRNA.
<a href="#">Anapc4</a>	<a href="#">Anapc4.dSep08</a>	<a href="#">305420</a>	1578	738	2	102	anaphase promoting complex (Anapc4) alternative variant dSep08, mRNA.
<a href="#">Anapc4</a>	<a href="#">Anapc4.eSep08</a>	<a href="#">305420</a>	8362	408	6	88	anaphase promoting complex CRA b (Anapc4) alternative variant eSep08, mRNA.
<a href="#">Anapc4</a>	<a href="#">Anapc4.fSep08</a>	<a href="#">305420</a>	546	464	2	31	anaphase promoting complex (Anapc4) alternative variant fSep08, mRNA.
<a href="#">Anapc5</a>	<a href="#">Anapc5.bSep08</a>	<a href="#">288671</a>	27968	1522	13	507	anaphase-promoting complex subunit 5 (Anapc5) alternative variant bSep08, mRNA.
<a href="#">Anapc5</a>	<a href="#">Anapc5.cSep08</a>	<a href="#">288671</a>	12538	933	6	196	anaphase-promoting complex subunit 5 (Anapc5) alternative variant cSep08, mRNA.
<a href="#">Anapc7</a>	<a href="#">Anapc7.bSep08</a>	<a href="#">304490</a>	13894	427	5	78	anaphase promoting complex subunit 7 (Anapc7) alternative variant bSep08, mRNA.
<a href="#">Anapc10</a>	<a href="#">Anapc10.bSep08</a>	<a href="#">361389</a>	52107	976	1	83	anaphase promoting complex subunit 10 (9.3 kD) (Anapc10) alternative variant bSep08, mRNA.
<a href="#">Anapc10</a>	<a href="#">Anapc10.cSep08</a>	<a href="#">361389</a>	51818	770	2	70	anaphase promoting complex subunit 10 (7.8 kD) (Anapc10) alternative variant cSep08, complete mRNA.

<a href="#">Anapc11</a>	<a href="#">Anapc11.bSep08</a>	<a href="#">498030</a>	8382	978	2	84	anaphase promoting complex subunit 11 homolog (yeast) (9.8 kD) (Anapc11) alternative variant bSep08, mRNA.
<a href="#">Anapc11</a>	<a href="#">Anapc11.cSep08</a>	<a href="#">498030</a>	8717	852	3	84	anaphase promoting complex subunit 11 homolog (yeast) (9.8 kD) (Anapc11) alternative variant cSep08, mRNA.
<a href="#">Andpro</a>	<a href="#">Andpro.bSep08</a>	<a href="#">25030</a>	6346	682	4	168	androgen regulated 20 kDa protein (Andpro) alternative variant bSep08, mRNA.
<a href="#">Andpro</a>	<a href="#">Andpro.cSep08</a>	<a href="#">25030</a>	6346	662	4	106	androgen regulated 20 kDa protein (12.7 kD) (Andpro) alternative variant cSep08, mRNA.
<a href="#">Andpro</a>	<a href="#">Andpro.dSep08</a>	<a href="#">25030</a>	6337	530	4	87	androgen regulated 20 kDa protein (10.2 kD) (Andpro) alternative variant dSep08, mRNA.
<a href="#">Andpro</a>	<a href="#">Andpro.eSep08</a>	<a href="#">25030</a>	2580	482	2	79	androgen regulated 20 kDa protein (Andpro) alternative variant eSep08, mRNA.
<a href="#">Angel1</a>	<a href="#">Angel1.bSep08</a>	<a href="#">362765</a>	14705	890	2	135	angel homolog 1 (Drosophila) (Angel1) alternative variant bSep08, mRNA.
<a href="#">Angel2</a>	<a href="#">Angel2.aSep08</a>	<a href="#">305035</a>	7843	3108		195	angel homolog 2 (Drosophila) (Angel2) mRNA.
<a href="#">Angpt4</a>	<a href="#">Angpt4.bSep08</a>	<a href="#">296269</a>	10046	686	2	116	angiopoietin 4 (Angpt4) alternative variant bSep08, mRNA.
<a href="#">Angptl1</a>	<a href="#">Angptl1.aSep08</a>	<a href="#">679942</a>	12913	1501		343	angiopoietin-like 1 (Angptl1) mRNA.
<a href="#">Angptl2</a>	<a href="#">Angptl2.bSep08</a>	<a href="#">171100</a>	5650	2325	2	191	angiopoietin-like 2 (22.7 kD) (Angptl2) alternative variant bSep08, mRNA.
<a href="#">Angptl3</a>	<a href="#">Angptl3.bSep08</a>	<a href="#">502970</a>	4669	904	5	220	angiopoietin-like 3 (Angptl3) alternative variant bSep08, mRNA.
<a href="#">Angptl3</a>	<a href="#">Angptl3.cSep08</a>	<a href="#">502970</a>	1266	472	3	89	angiopoietin-like 3 (Angptl3) alternative variant cSep08, mRNA.
<a href="#">Angptl4</a>	<a href="#">Angptl4.bSep08</a>	<a href="#">362850</a>	2022	760	4	121	angiopoietin-like 4 (Angptl4) alternative variant bSep08, mRNA.
<a href="#">Angptl6</a>	<a href="#">Angptl6.cSep08</a>	<a href="#">298698</a>	1723	712	2		
<a href="#">Ank</a>	<a href="#">Ank.bSep08</a>	<a href="#">114506</a>	91349	742	3	219	progressive ankylosis (Ank) alternative variant bSep08, mRNA.
<a href="#">Ank</a>	<a href="#">Ank.cSep08</a>	<a href="#">114506</a>	77291	553	2	171	progressive ankylosis (Ank) alternative variant cSep08, mRNA.
<a href="#">Ank.1</a>	<a href="#">Ank.1.aSep08</a>		2006	1499		313	ankyrin (34.1 kD) (Ank.1) mRNA.
<a href="#">Ank.2</a>	<a href="#">Ank.2.aSep08</a>		8081	345		115	CRA a (Ank.2) mRNA.
<a href="#">Ank.3</a>	<a href="#">Ank.3.aSep08</a>		22192	1142	2	380	CRA a (Ank.3) alternative variant aSep08, mRNA.
<a href="#">Ank.3</a>	<a href="#">Ank.3.bSep08</a>		17065	621	1	207	CRA a (Ank.3) alternative variant bSep08, mRNA.
<a href="#">Ank.4</a>	<a href="#">Ank.4.aSep08</a>		9906	924		308	ankyrin (Ank.4) mRNA.
<a href="#">Ank.5</a>	<a href="#">Ank.5.aSep08</a>	<a href="#">502141</a>	24570	250		54	ankyrin (Ank.5) mRNA.
<a href="#">Ank.7</a>	<a href="#">Ank.7.aSep08</a>		11547	633		106	ankyrin (Ank.7) mRNA.
<a href="#">Ank.8</a>	<a href="#">Ank.8.aSep08</a>		2884	896		112	ankyrin (11.8 kD) (Ank.8) mRNA.
<a href="#">Ank.9</a>	<a href="#">Ank.9.aSep08</a>		30493	373		124	ankyrin (Ank.9) mRNA.
<a href="#">Ank.10</a>	<a href="#">Ank.10.aSep08</a>		25648	2181	8	478	putative protein of metazoan origin (Ank.10) alternative variant aSep08, mRNA.
<a href="#">Ank.10</a>	<a href="#">Ank.10.bSep08</a>		3261	641	3	160	putative protein of vertebrate origin (Ank.10) alternative variant bSep08, mRNA.
<a href="#">Ank.10</a>	<a href="#">Ank.10.cSep08</a>		27980	774	4	87	putative protein of vertebrate origin (9.5 kD) (Ank.10) alternative variant cSep08, mRNA.

<a href="#">Ank.10</a>	<a href="#">Ank.10.dSep08</a>		3490	2379	3	84	ankyrin (Ank.10) alternative variant dSep08, mRNA.
<a href="#">Ank.11</a>	<a href="#">Ank.11.aSep08</a>		59815	607		202	ankyrin (Ank.11) mRNA.
<a href="#">Ank.12</a>	<a href="#">Ank.12.aSep08</a>		11989	809		144	CRA b (Ank.12) mRNA.
<a href="#">Ank.13</a>	<a href="#">Ank.13.aSep08</a>		4581	1272		401	ankyrin (Ank.13) mRNA.
<a href="#">Ank.14</a>	<a href="#">Ank.14.aSep08</a>		3873	528		175	ankyrin (Ank.14) mRNA.
<a href="#">Ank.15</a>	<a href="#">Ank.15.aSep08</a>		1638	750		245	ankyrin (Ank.15) mRNA.
<a href="#">Ank.16</a>	<a href="#">Ank.16.aSep08</a>		10112	639		148	ankyrin (Ank.16) mRNA.
<a href="#">Ank.17</a>	<a href="#">Ank.17.aSep08</a>		2297	1410	1	444	mindbomb homolog 2 CRA a (Ank.17) alternative variant aSep08, mRNA.
<a href="#">Ank.17</a>	<a href="#">Ank.17.bSep08</a>		1456	696		133	mindbomb homolog 2 CRA a (Ank.17) alternative variant bSep08, mRNA.
<a href="#">Ank.19</a>	<a href="#">Ank.19.aSep08</a>		3972	449		113	ankyrin B (Ank.19) mRNA.
<a href="#">Ank.20</a>	<a href="#">Ank.20.aSep08</a>		21449	720		150	TNNI3 interacting kinase (16.8 kD) (Ank.20) mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.bSep08</a>	<a href="#">306570</a>	8460	535	2	177	ankyrin (Ank1) alternative variant bSep08, mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.cSep08</a>	<a href="#">306570</a>	8638	1110	5	155	ankyrin 1 precursor (17.5 kD) (Ank1) alternative variant cSep08, mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.dSep08</a>	<a href="#">306570</a>	27661	505	5	137	ankyrin (Ank1) alternative variant dSep08, mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.fSep08</a>	<a href="#">306570</a>	7525	567	4	91	ankyrin (Ank1) alternative variant fSep08, mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.gSep08</a>	<a href="#">306570</a>	1860	421	3	81	ankyrin (Ank1) alternative variant gSep08, mRNA.
<a href="#">Ank1</a>	<a href="#">Ank1.hSep08</a>	<a href="#">306570</a>	7182	1228	3	56	putative protein (6.2 kD) (Ank1) alternative variant hSep08, mRNA.
<a href="#">Ank2</a>	<a href="#">Ank2.aSep08</a>	<a href="#">362036</a>	12001	1631	6	278	ankyrin 2, neuronal (Ank2) alternative variant aSep08, mRNA.
<a href="#">Ank2</a>	<a href="#">Ank2.bSep08</a>	<a href="#">362036</a>	37120	1781	6	167	ankyrin 2, neuronal (Ank2) alternative variant bSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.cSep08</a>	<a href="#">361833</a>	13214	1796	6	598	ankyrin 3 epithelial CRA c (Ank3) alternative variant cSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.dSep08</a>	<a href="#">361833</a>	54589	3041	11	484	ankyrin 3 epithelial CRA j (Ank3) alternative variant dSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.eSep08</a>	<a href="#">361833</a>	46093	1893	8	453	ankyrin G (Ank3) alternative variant eSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.fSep08</a>	<a href="#">361833</a>	20408	1008	9	335	ankyrin 3 (Ank3) alternative variant fSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.gSep08</a>	<a href="#">361833</a>	10380	3240	5	296	ankyrin G (Ank3) alternative variant gSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.hSep08</a>	<a href="#">361833</a>	32975	804	6	268	ankyrin G1 (Ank3) alternative variant hSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.iSep08</a>	<a href="#">361833</a>	10310	549	3	183	ankyrin G (Ank3) alternative variant iSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.jSep08</a>	<a href="#">361833</a>	5457	450	4	149	ankyrin 3 (Ank3) alternative variant jSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.kSep08</a>	<a href="#">361833</a>	17074	391	4	82	ankyrin 3 epithelial CRA g (Ank3) alternative variant kSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.lSep08</a>	<a href="#">361833</a>	4795	395	3	71	ankyrin G1 (Ank3) alternative variant lSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.nSep08</a>	<a href="#">361833</a>	1679	414	3	58	ankyrin 3 (Ank3) alternative variant nSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.oSep08</a>	<a href="#">361833</a>	10205	501	4	54	ankyrin G1 (Ank3) alternative variant oSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.pSep08</a>	<a href="#">361833</a>	831	661	2	39	ankyrin 3 epithelial CRA i (Ank3) alternative variant pSep08, mRNA.
<a href="#">Ank3</a>	<a href="#">Ank3.qSep08</a>	<a href="#">361833</a>	64434	1111	2	49	putative protein (5.3 kD) (Ank3) alternative variant qSep08, mRNA.

<a href="#">Ankar</a>	<a href="#">Ankar.aSep08</a>	<a href="#">501138</a>	5571	601		200	ankyrin and armadillo repeat containing (Ankar) mRNA.
<a href="#">Ankfy1</a>	<a href="#">Ankfy1.bSep08</a>	<a href="#">303292</a>	11475	625	5	208	ankyrin (Ankfy1) alternative variant bSep08, mRNA.
<a href="#">Ankib1</a>	<a href="#">Ankib1.bSep08</a>	<a href="#">368062</a>	65796	1247	6	276	ankyrin (31.1 kD) (Ankib1) alternative variant bSep08, complete mRNA.
<a href="#">Ankib1</a>	<a href="#">Ankib1.cSep08</a>	<a href="#">368062</a>	11708	555	6	185	putative protein of eukaryotic origin (Ankib1) alternative variant cSep08, mRNA.
<a href="#">Ankle2</a>	<a href="#">Ankle2.bSep08</a>	<a href="#">360829</a>	25603	1787	6	528	putative protein of metazoan origin (Ankle2) alternative variant bSep08, mRNA.
<a href="#">Ankle2</a>	<a href="#">Ankle2.cSep08</a>	<a href="#">360829</a>	7388	1154	3	384	lamino-associated polypeptide 2/emerin (Ankle2) alternative variant cSep08, mRNA.
<a href="#">Ankle2</a>	<a href="#">Ankle2.dSep08</a>	<a href="#">360829</a>	17185	954	4	318	putative protein of metazoan origin (Ankle2) alternative variant dSep08, mRNA.
<a href="#">Ankle2</a>	<a href="#">Ankle2.eSep08</a>	<a href="#">360829</a>	2906	2004	3	274	putative protein of bilateral origin (Ankle2) alternative variant eSep08, mRNA.
<a href="#">Ankmy2</a>	<a href="#">Ankmy2.aSep08</a>	<a href="#">314046</a>	42803	2543	11	468	ankyrin and zinc finger, MYND-type (Ankmy2) alternative variant aSep08, mRNA.
<a href="#">Ankmy2</a>	<a href="#">Ankmy2.dSep08</a>	<a href="#">314046</a>	9238	398	2	132	putative protein (Ankmy2) alternative variant dSep08, mRNA.
<a href="#">Ankmy2</a>	<a href="#">Ankmy2.fSep08</a>	<a href="#">314046</a>	3125	934	3	65	zinc finger, MYND-type (Ankmy2) alternative variant fSep08, mRNA.
<a href="#">Ankra2</a>	<a href="#">Ankra2.bSep08</a>	<a href="#">294679</a>	8639	737	5	192	ankyrin (Ankra2) alternative variant bSep08, mRNA.
<a href="#">Ankra2</a>	<a href="#">Ankra2.cSep08</a>	<a href="#">294679</a>	6627	1401	5	165	ankyrin (Ankra2) alternative variant cSep08, mRNA.
<a href="#">Ankra2</a>	<a href="#">Ankra2.dSep08</a>	<a href="#">294679</a>	1607	644	3	78	ankyrin (Ankra2) alternative variant dSep08, mRNA.
<a href="#">Ankrd1</a>	<a href="#">Ankrd1.bSep08</a>	<a href="#">27064</a>	15509	931	9	310	ankyrin (Ankrd1) alternative variant bSep08, mRNA.
<a href="#">Ankrd1</a>	<a href="#">Ankrd1.cSep08</a>	<a href="#">27064</a>	4243	256	3	65	putative protein (Ankrd1) alternative variant cSep08, mRNA.
<a href="#">Ankrd1</a>	<a href="#">Ankrd1.dSep08</a>	<a href="#">27064</a>	6243	567	5	49	putative protein (5.4 kD) (Ankrd1) alternative variant dSep08, mRNA.
<a href="#">Ankrd2</a>	<a href="#">Ankrd2.bSep08</a>	<a href="#">309374</a>	3239	758	5	166	ankyrin (18.5 kD) (Ankrd2) alternative variant bSep08, mRNA.
<a href="#">Ankrd5</a>	<a href="#">Ankrd5.bSep08</a>	<a href="#">296184</a>	34561	2250	12	256	ankyrin (29.1 kD) (Ankrd5) alternative variant bSep08, mRNA.
<a href="#">Ankrd5</a>	<a href="#">Ankrd5.cSep08</a>	<a href="#">296184</a>	2327	449	3	101	putative protein (Ankrd5) alternative variant cSep08, mRNA.
<a href="#">Ankrd6</a>	<a href="#">Ankrd6.aSep08</a>	<a href="#">500430</a>	139100	4586	17	713	ankyrin (78.1 kD) (Ankrd6) alternative variant aSep08, mRNA.
<a href="#">Ankrd6</a>	<a href="#">Ankrd6.bSep08</a>	<a href="#">500430</a>	2044	775	2	98	putative protein, with a coiled coil domain, of vertebrate origin (Ankrd6) alternative variant bSep08, mRNA.
<a href="#">Ankrd6</a>	<a href="#">Ankrd6.cSep08</a>	<a href="#">500430</a>	1475	245	2	81	putative protein of vertebrate origin (Ankrd6) alternative variant cSep08, mRNA.
<a href="#">Ankrd10</a>	<a href="#">Ankrd10.aSep08</a>	<a href="#">361183</a>	8189	3005	3	232	putative nuclear protein of vertebrate origin (24.7 kD) (Ankrd10) alternative variant aSep08, mRNA.
<a href="#">Ankrd10</a>	<a href="#">Ankrd10.bSep08</a>	<a href="#">361183</a>	16426	667	4	222	ankyrin (Ankrd10) alternative variant bSep08, mRNA.
<a href="#">Ankrd10</a>	<a href="#">Ankrd10.cSep08</a>	<a href="#">361183</a>	10328	909	4	219	putative protein of vertebrate origin (Ankrd10) alternative variant cSep08, mRNA.
<a href="#">Ankrd10</a>	<a href="#">Ankrd10.dSep08</a>	<a href="#">361183</a>	22588	1781	6	195	ankyrin (Ankrd10) alternative variant dSep08, mRNA.

<a href="#">Ankrd11</a>	<a href="#">Ankrd11.bSep08</a>	<a href="#">365023</a>	8243	3176	4	626	putative protein of bilateral origin (Ankrd11) alternative variant bSep08, mRNA.
<a href="#">Ankrd11</a>	<a href="#">Ankrd11.dSep08</a>	<a href="#">365023</a>	157390	1206	7	270	putative protein (Ankrd11) alternative variant dSep08, mRNA.
<a href="#">Ankrd12</a>	<a href="#">Ankrd12.bSep08</a>	<a href="#">316775</a>	25772	524	4	157	putative protein (Ankrd12) alternative variant bSep08, mRNA.
<a href="#">Ankrd13a</a>	<a href="#">Ankrd13a.bSep08</a>	<a href="#">360823</a>	8515	738	8	246	putative protein of eukaryotic origin (Ankrd13a) alternative variant bSep08, mRNA.
<a href="#">Ankrd13c</a>	<a href="#">Ankrd13c.aSep08</a>	<a href="#">685374</a>	8043	2383		124	putative protein of eukaryotic origin (Ankrd13c) mRNA.
<a href="#">Ankrd13d</a>	<a href="#">Ankrd13d.dSep08</a>	<a href="#">361699</a>	479	385	2	83	putative protein of metazoan origin (Ankrd13d) alternative variant dSep08, mRNA.
<a href="#">Ankrd15</a>	<a href="#">Ankrd15.bSep08</a>	<a href="#">309429</a>	25540	3431	7	1000	putative protein, with 2 coiled coil domains, of bilateral origin (Ankrd15) alternative variant bSep08, mRNA.
<a href="#">Ankrd15</a>	<a href="#">Ankrd15.cSep08</a>	<a href="#">309429</a>	5714	360	4	119	putative protein of vertebrate origin (Ankrd15) alternative variant cSep08, mRNA.
<a href="#">Ankrd15</a>	<a href="#">Ankrd15.dSep08</a>	<a href="#">309429</a>	2950	981	2	105	ankyrin (11.5 kD) (Ankrd15) alternative variant dSep08, mRNA.
<a href="#">Ankrd15</a>	<a href="#">Ankrd15.gSep08</a>	<a href="#">309429</a>	5134	580	3	72	putative protein (Ankrd15) alternative variant gSep08, mRNA.
<a href="#">Ankrd15</a>	<a href="#">Ankrd15.hSep08</a>	<a href="#">309429</a>	2320	486	2	99	putative protein of vertebrate origin (Ankrd15) alternative variant hSep08, mRNA.
<a href="#">Ankrd17</a>	<a href="#">Ankrd17.bSep08</a>	<a href="#">289521</a>	53054	5239	21	1680	ankyrin and KH, type 1 (Ankrd17) alternative variant bSep08, mRNA.
<a href="#">Ankrd17</a>	<a href="#">Ankrd17.cSep08</a>	<a href="#">289521</a>	16763	1321	5	440	ankyrin (Ankrd17) alternative variant cSep08, mRNA.
<a href="#">Ankrd17</a>	<a href="#">Ankrd17.dSep08</a>	<a href="#">289521</a>	19222	908	8	302	ankyrin (Ankrd17) alternative variant dSep08, mRNA.
<a href="#">Ankrd17</a>	<a href="#">Ankrd17.eSep08</a>	<a href="#">289521</a>	9573	359	3	119	ankyrin (Ankrd17) alternative variant eSep08, mRNA.
<a href="#">Ankrd17</a>	<a href="#">Ankrd17.fSep08</a>	<a href="#">289521</a>	1345	423	3	83	putative protein, with a coiled coil domain (Ankrd17) alternative variant fSep08, mRNA.
<a href="#">Ankrd23</a>	<a href="#">Ankrd23.aSep08</a>	<a href="#">316330</a>	5834	2765		309	ankyrin (Ankrd23) mRNA.
<a href="#">Ankrd24</a>	<a href="#">Ankrd24.bSep08</a>	<a href="#">299639</a>	4412	1930	5	508	putative protein, with 3 coiled coil domains, of fungal and metazoan origin (Ankrd24) alternative variant bSep08, mRNA.
<a href="#">Ankrd24</a>	<a href="#">Ankrd24.cSep08</a>	<a href="#">299639</a>	644	532	2		
<a href="#">Ankrd26</a>	<a href="#">Ankrd26.aSep08</a>	<a href="#">312667</a>	7618	393		130	putative protein, with 2 coiled coil domains, of mammalian origin (Ankrd26) mRNA.
<a href="#">Ankrd27</a>	<a href="#">Ankrd27.aSep08</a>	<a href="#">361555</a>	21943	1238		372	repeat-containing protein (Ankrd27) mRNA.
<a href="#">Ankrd28</a>	<a href="#">Ankrd28.aSep08</a>	<a href="#">306264</a>	47101	1786		451	ankyrin (Ankrd28) alternative variant aSep08, mRNA.
<a href="#">Ankrd28</a>	<a href="#">Ankrd28.bSep08</a>	<a href="#">306264</a>	15773	671		223	ankyrin (Ankrd28) alternative variant bSep08, mRNA.
<a href="#">Ankrd28</a>	<a href="#">Ankrd28.cSep08</a>	<a href="#">306264</a>	16062	513		171	ankyrin (Ankrd28) alternative variant cSep08, mRNA.
<a href="#">Ankrd28</a>	<a href="#">Ankrd28.dSep08</a>	<a href="#">306264</a>	10758	3685		139	ankyrin (Ankrd28) alternative variant dSep08, mRNA.
<a href="#">Ankrd34b</a>	<a href="#">Ankrd34b.bSep08</a>	<a href="#">499506</a>	1863	345	1	91	putative protein (Ankrd34b) alternative variant bSep08, mRNA.
<a href="#">Ankrd35</a>	<a href="#">Ankrd35.aSep08</a>	<a href="#">365881</a>	5925	463	5	89	putative protein, with a coiled coil domain, of vertebrate origin (Ankrd35) alternative variant aSep08, mRNA.
<a href="#">Ankrd36</a>	<a href="#">Ankrd36.aSep08</a>	<a href="#">305491</a>	25707	1194	9	397	ankyrin (Ankrd36) alternative variant aSep08, mRNA.

<a href="#">Ankrd36</a>	<a href="#">Ankrd36.bSep08</a>	<a href="#">305491</a>	9711	1307	4	372	ankyrin (Ankrd36) alternative variant bSep08, mRNA.
<a href="#">Ankrd36</a>	<a href="#">Ankrd36.cSep08</a>	<a href="#">305491</a>	25708	1945	7	227	putative protein of mammalian origin (25.7 kD) (Ankrd36) alternative variant cSep08, mRNA.
<a href="#">Ankrd37</a>	<a href="#">Ankrd37.bSep08</a>	<a href="#">361149</a>	1118	642	1	42	putative protein of mammalian origin (4.9 kD) (Ankrd37) alternative variant bSep08, mRNA.
<a href="#">Ankrd39</a>	<a href="#">Ankrd39.aSep08</a>	<a href="#">367251</a>	9620	1795		522	ankyrin (Ankrd39) alternative variant aSep08, mRNA.
<a href="#">Ankrd41</a>	<a href="#">Ankrd41.aSep08</a>	<a href="#">361122</a>	3866	2895		467	CRA c (51.5 kD) (Ankrd41) mRNA.
<a href="#">Ankrd42</a>	<a href="#">Ankrd42.aSep08</a>	<a href="#">293117</a>	30455	1503	8	403	ankyrin (Ankrd42) alternative variant aSep08, mRNA.
<a href="#">Ankrd42</a>	<a href="#">Ankrd42.bSep08</a>	<a href="#">293117</a>	3507	786	1	30	putative protein (3.5 kD) (Ankrd42) alternative variant bSep08, mRNA.
<a href="#">Ankrd44</a>	<a href="#">Ankrd44.aSep08</a>	<a href="#">301415</a>	69738	743		247	ankyrin (Ankrd44) mRNA.
<a href="#">Ankrd45</a>	<a href="#">Ankrd45.aSep08</a>	<a href="#">289152</a>	12677	611		193	ankyrin (Ankrd45) mRNA.
<a href="#">Ankrd46</a>	<a href="#">Ankrd46.aSep08</a>	<a href="#">299982</a>	19492	807	1	250	ankyrin (Ankrd46) alternative variant aSep08, mRNA.
<a href="#">Ankrd49</a>	<a href="#">Ankrd49.bSep08</a>	<a href="#">315434</a>	1597	612	2	172	ankyrin (Ankrd49) alternative variant bSep08, mRNA.
<a href="#">Ankrd49</a>	<a href="#">Ankrd49.cSep08</a>	<a href="#">315434</a>	1525	400	2	48	putative protein of mammalian origin (5.5 kD) (Ankrd49) alternative variant cSep08, mRNA.
<a href="#">Ankrd54</a>	<a href="#">Ankrd54.bSep08</a>	<a href="#">362957</a>	11933	1459	6	179	ankyrin (19.8 kD) (Ankrd54) alternative variant bSep08, mRNA.
<a href="#">Ankrd54</a>	<a href="#">Ankrd54.cSep08</a>	<a href="#">362957</a>	6495	2658	2	142	ankyrin (16.0 kD) (Ankrd54) alternative variant cSep08, mRNA.
<a href="#">Ankrd55</a>	<a href="#">Ankrd55.aSep08</a>	<a href="#">361898</a>	40608	826	6	249	ankyrin (Ankrd55) alternative variant aSep08, mRNA.
<a href="#">Ankrd55</a>	<a href="#">Ankrd55.bSep08</a>	<a href="#">361898</a>	20034	498	4	101	ankyrin (Ankrd55) alternative variant bSep08, mRNA.
<a href="#">Ankrd55</a>	<a href="#">Ankrd55.cSep08</a>	<a href="#">361898</a>	8973	889	4	84	ankyrin (Ankrd55) alternative variant cSep08, mRNA.
<a href="#">Ankrd60</a>	<a href="#">Ankrd60.aSep08</a>	<a href="#">296417</a>	4181	703		198	ankyrin (Ankrd60) mRNA.
<a href="#">Anks1</a>	<a href="#">Anks1.bSep08</a>	<a href="#">309639</a>	1126	387	2	116	putative protein of vertebrate origin (Anks1) alternative variant bSep08, mRNA.
<a href="#">Anks1</a>	<a href="#">Anks1.dSep08</a>	<a href="#">309639</a>	1436	442	2	41	putative protein (Anks1) alternative variant dSep08, mRNA.
<a href="#">Anks3</a>	<a href="#">Anks3.bSep08</a>	<a href="#">302937</a>	9061	751	6	249	ankyrin (Anks3) alternative variant bSep08, mRNA.
<a href="#">Anks3</a>	<a href="#">Anks3.cSep08</a>	<a href="#">302937</a>	4266	954	7	153	putative protein, with 2 coiled coil domains (Anks3) alternative variant cSep08, mRNA.
<a href="#">Anks3</a>	<a href="#">Anks3.dSep08</a>	<a href="#">302937</a>	1741	733	5	114	putative protein, with a coiled coil domain, of mammalian origin (Anks3) alternative variant dSep08, mRNA.
<a href="#">Ankzf1</a>	<a href="#">Ankzf1.bSep08</a>	<a href="#">363255</a>	3730	1768	5	241	putative cytoplasmic protein of eukaryotic origin (26.3 kD) (Ankzf1) alternative variant bSep08, mRNA.
<a href="#">Ankzf1</a>	<a href="#">Ankzf1.cSep08</a>	<a href="#">363255</a>	6690	578	2	141	putative protein of mammalian origin (Ankzf1) alternative variant cSep08, mRNA.
<a href="#">Ankzf1</a>	<a href="#">Ankzf1.dSep08</a>	<a href="#">363255</a>	1093	753	3	140	putative protein of eukaryotic origin (16.1 kD) (Ankzf1) alternative variant dSep08, mRNA.
<a href="#">Ankzf1</a>	<a href="#">Ankzf1.eSep08</a>	<a href="#">363255</a>	616	418	2	116	putative protein of eukaryotic origin (Ankzf1) alternative variant eSep08, mRNA.
<a href="#">Annexin.2</a>	<a href="#">Annexin.2.aSep08</a>		37970	422		140	annexin A13 (Annexin.2) mRNA.
<a href="#">Anp32a</a>	<a href="#">Anp32a.bSep08</a>	<a href="#">25379</a>	7908	766	4	155	acidic (leucine-rich) nuclear phosphoprotein 32 family, member A (Anp32a) alternative variant bSep08, mRNA.
<a href="#">Anp32a</a>	<a href="#">Anp32a.cSep08</a>	<a href="#">25379</a>	31101	333	3	110	acidic (leucine-rich) nuclear phosphoprotein 32 family, member A (Anp32a) alternative variant cSep08, mRNA.

<a href="#">Anp32a</a>	<a href="#">Anp32a.dSep08</a>	<a href="#">25379</a>	24717	342	3	25	acidic (leucine-rich) nuclear phosphoprotein 32 family, member A (Anp32a) alternative variant dSep08, mRNA.
<a href="#">Anp32a</a>	<a href="#">Anp32a.eSep08</a>	<a href="#">25379</a>	500	314	2	50	acidic (leucine-rich) nuclear phosphoprotein 32 family, member A (Anp32a) alternative variant eSep08, mRNA.
<a href="#">Anp32e</a>	<a href="#">Anp32e.bSep08</a>	<a href="#">361999</a>	7169	858	3	211	putative protein (Anp32e) alternative variant bSep08, mRNA.
<a href="#">Anp32e</a>	<a href="#">Anp32e.cSep08</a>	<a href="#">361999</a>	4372	715	2	90	acidic nuclear phosphoprotein 32 family member E CRA e (10.5 kD) (Anp32e) alternative variant cSep08, mRNA.
<a href="#">Anp32e</a>	<a href="#">Anp32e.dSep08</a>	<a href="#">361999</a>	3103	2298	2	74	putative nuclear protein (8.8 kD) (Anp32e) alternative variant dSep08, mRNA.
<a href="#">Anpep</a>	<a href="#">Anpep.bSep08</a>	<a href="#">81641</a>	732	365	2	57	alanyl (membrane) aminopeptidase (Anpep) alternative variant bSep08, mRNA.
<a href="#">Antxrl</a>	<a href="#">Antxrl.aSep08</a>	<a href="#">364513</a>	6552	741		239	anthrax toxin receptor-like (Antxrl) mRNA.
<a href="#">Anubl1</a>	<a href="#">Anubl1.bSep08</a>	<a href="#">286998</a>	51360	999	6	308	AN1, ubiquitin-like, homolog (Xenopus laevis) (Anubl1) alternative variant bSep08, mRNA.
<a href="#">Anubl1</a>	<a href="#">Anubl1.dSep08</a>	<a href="#">286998</a>	27850	758	4	180	AN1, ubiquitin-like, homolog (Xenopus laevis) (Anubl1) alternative variant dSep08, mRNA.
<a href="#">Anubl1</a>	<a href="#">Anubl1.eSep08</a>	<a href="#">286998</a>	27131	374	4	105	AN1, ubiquitin-like, homolog (Xenopus laevis) (Anubl1) alternative variant eSep08, mRNA.
<a href="#">Anxa1</a>	<a href="#">Anxa1.bSep08</a>	<a href="#">25380</a>	12744	963	11	269	annexin A1 (30.2 kD) (Anxa1) alternative variant bSep08, mRNA.
<a href="#">Anxa1</a>	<a href="#">Anxa1.cSep08</a>	<a href="#">25380</a>	9872	661	8	168	annexin A1 (Anxa1) alternative variant cSep08, mRNA.
<a href="#">Anxa1</a>	<a href="#">Anxa1.dSep08</a>	<a href="#">25380</a>	2332	1471	2	78	putative protein (9.2 kD) (Anxa1) alternative variant dSep08, mRNA.
<a href="#">Anxa1</a>	<a href="#">Anxa1.hSep08</a>	<a href="#">25380</a>	7580	508	6	65	annexin A1 CRA b (Anxa1) alternative variant hSep08, mRNA.
<a href="#">Anxa2</a>	<a href="#">Anxa2.bSep08</a>	<a href="#">56611</a>	30556	776	9	258	annexin A2 (Anxa2) alternative variant bSep08, mRNA.
<a href="#">Anxa2</a>	<a href="#">Anxa2.cSep08</a>	<a href="#">56611</a>	6021	1491	6	167	annexin A2 (Anxa2) alternative variant cSep08, mRNA.
<a href="#">Anxa2</a>	<a href="#">Anxa2.dSep08</a>	<a href="#">56611</a>	879	298	3	27	annexin A2 (Anxa2) alternative variant dSep08, mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.aSep08</a>	<a href="#">25291</a>	43383	1191	13	212	annexin A3 (24.1 kD) (Anxa3) alternative variant aSep08, complete mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.bSep08</a>	<a href="#">25291</a>	52764	2281	14	199	annexin A3 (22.6 kD) (Anxa3) alternative variant bSep08, complete mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.cSep08</a>	<a href="#">25291</a>	52768	418	4	131	annexin A3 (Anxa3) alternative variant cSep08, mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.dSep08</a>	<a href="#">25291</a>	31391	698	9	126	annexin A3 (13.9 kD) (Anxa3) alternative variant dSep08, mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.eSep08</a>	<a href="#">25291</a>	30542	740	8	126	annexin A3 (13.9 kD) (Anxa3) alternative variant eSep08, mRNA.
<a href="#">Anxa3</a>	<a href="#">Anxa3.fSep08</a>	<a href="#">25291</a>	18003	709	7	116	annexin A3 (Anxa3) alternative variant fSep08, mRNA.
<a href="#">Anxa4</a>	<a href="#">Anxa4.bSep08</a>	<a href="#">79124</a>	25213	703	8	170	annexin A4 (Anxa4) alternative variant bSep08, mRNA.
<a href="#">Anxa4</a>	<a href="#">Anxa4.cSep08</a>	<a href="#">79124</a>	27930	548	7	157	annexin A4 (Anxa4) alternative variant cSep08, mRNA.
<a href="#">Anxa4</a>	<a href="#">Anxa4.dSep08</a>	<a href="#">79124</a>	4511	877	2	48	annexin A4 (5.4 kD) (Anxa4) alternative variant dSep08, mRNA.
<a href="#">Anxa5</a>	<a href="#">Anxa5.aSep08</a>	<a href="#">25673</a>	30672	1543	8	377	annexin A5 (Anxa5) alternative variant aSep08, mRNA.
<a href="#">Anxa5</a>	<a href="#">Anxa5.bSep08</a>	<a href="#">25673</a>	31292	1201	8	353	annexin A5 (Anxa5) alternative variant bSep08, mRNA.



<a href="#">Anxa5</a>	<a href="#">Anxa5.cSep08</a>	<a href="#">25673</a>	27485	765	4	254	annexin A5 (Anxa5) alternative variant cSep08, mRNA.
<a href="#">Anxa5</a>	<a href="#">Anxa5.dSep08</a>	<a href="#">25673</a>	28093	742	4	205	annexin A5 (Anxa5) alternative variant dSep08, mRNA.
<a href="#">Anxa5</a>	<a href="#">Anxa5.eSep08</a>	<a href="#">25673</a>	23026	688	2	137	annexin A5 (Anxa5) alternative variant eSep08, mRNA.
<a href="#">Anxa6</a>	<a href="#">Anxa6.aSep08</a>	<a href="#">79125</a>	55856	2772	10	714	annexin A6 (Anxa6) alternative variant aSep08, mRNA.
<a href="#">Anxa6</a>	<a href="#">Anxa6.bSep08</a>	<a href="#">79125</a>	20198	1384	9	330	annexin A6 (Anxa6) alternative variant bSep08, mRNA.
<a href="#">Anxa6</a>	<a href="#">Anxa6.cSep08</a>	<a href="#">79125</a>	12110	749	2	249	annexin A6 (Anxa6) alternative variant cSep08, mRNA.
<a href="#">Anxa6</a>	<a href="#">Anxa6.dSep08</a>	<a href="#">79125</a>	8346	1010	4	103	annexin A6 (Anxa6) alternative variant dSep08, mRNA.
<a href="#">Anxa7</a>	<a href="#">Anxa7.bSep08</a>	<a href="#">155423</a>	17016	741	2	230	annexin A7 (Anxa7) alternative variant bSep08, mRNA.
<a href="#">Anxa7</a>	<a href="#">Anxa7.cSep08</a>	<a href="#">155423</a>	20462	707	2	229	annexin A7 (Anxa7) alternative variant cSep08, mRNA.
<a href="#">Anxa7</a>	<a href="#">Anxa7.dSep08</a>	<a href="#">155423</a>	20328	715	2	212	annexin A7 (Anxa7) alternative variant dSep08, mRNA.
<a href="#">Anxa9</a>	<a href="#">Anxa9.bSep08</a>	<a href="#">689830</a>	9087	1776	7	68	annexin A9 (7.2 kD) (Anxa9) alternative variant bSep08, mRNA.
<a href="#">Anxa9</a>	<a href="#">Anxa9.cSep08</a>	<a href="#">689830</a>	938	646	2	55	annexin A9 (Anxa9) alternative variant cSep08, mRNA.
<a href="#">Anxa11</a>	<a href="#">Anxa11.bSep08</a>	<a href="#">290527</a>	11196	1816	9	236	annexin A11 (26.8 kD) (Anxa11) alternative variant bSep08, mRNA.
<a href="#">Anxa11</a>	<a href="#">Anxa11.dSep08</a>	<a href="#">290527</a>	9004	1315	8	105	annexin A11 (11.9 kD) (Anxa11) alternative variant dSep08, mRNA.
<a href="#">Anxa13</a>	<a href="#">Anxa13.aSep08</a>	<a href="#">362915</a>	7144	923		98	annexin A13 (Anxa13) mRNA.
<a href="#">Aoah</a>	<a href="#">Aoah.aSep08</a>	<a href="#">498757</a>	67033	414		137	acyloxyacyl hydrolase (Aoah) mRNA.
<a href="#">Aof2</a>	<a href="#">Aof2.aSep08</a>	<a href="#">500569</a>	22597	1643	15	547	amine oxidase (flavin containing) domain 2 (Aof2) alternative variant aSep08, mRNA.
<a href="#">Aof2</a>	<a href="#">Aof2.cSep08</a>	<a href="#">500569</a>	37164	799	7	208	amine oxidase (flavin containing) domain 2 (Aof2) alternative variant cSep08, mRNA.
<a href="#">Aof2</a>	<a href="#">Aof2.dSep08</a>	<a href="#">500569</a>	3539	1778	5	138	amine oxidase (flavin containing) domain 2 (14.7 kD) (Aof2) alternative variant dSep08, mRNA.
<a href="#">Aox1</a>	<a href="#">Aox1.bSep08</a>	<a href="#">54349</a>	15635	1009	6	335	aldehyde oxidase 1 (Aox1) alternative variant bSep08, mRNA.
<a href="#">Aox1</a>	<a href="#">Aox1.cSep08</a>	<a href="#">54349</a>	10675	1050	8	275	aldehyde oxidase 1 (Aox1) alternative variant cSep08, mRNA.
<a href="#">Aox1</a>	<a href="#">Aox1.dSep08</a>	<a href="#">54349</a>	8586	568	4	188	aldehyde oxidase 1 (Aox1) alternative variant dSep08, mRNA.
<a href="#">Aox1</a>	<a href="#">Aox1.eSep08</a>	<a href="#">54349</a>	3093	520	4	172	aldehyde oxidase 1 (Aox1) alternative variant eSep08, mRNA.
<a href="#">Aox3</a>	<a href="#">Aox3.bSep08</a>	<a href="#">493909</a>	7343	785	6	261	aldehyde oxidase (Aox3) alternative variant bSep08, mRNA.
<a href="#">Aox3</a>	<a href="#">Aox3.cSep08</a>	<a href="#">493909</a>	1545	365	2	111	putative protein (Aox3) alternative variant cSep08, mRNA.
<a href="#">Aox3</a>	<a href="#">Aox3.dSep08</a>	<a href="#">493909</a>	14406	327	3	80	aldehyde oxidase (Aox3) alternative variant dSep08, mRNA.
<a href="#">Aox3</a>	<a href="#">Aox3.eSep08</a>	<a href="#">493909</a>	9808	279	4	32	putative protein (3.6 kD) (Aox3) alternative variant eSep08, mRNA.
<a href="#">Aox4</a>	<a href="#">Aox4.aSep08</a>	<a href="#">316424</a>	3589	1051		96	aldehyde oxidase 4 (Aox4) mRNA.
<a href="#">Ap1b1</a>	<a href="#">Ap1b1.bSep08</a>	<a href="#">29663</a>	7935	827	6	275	adaptor protein complex AP-1, beta 1 subunit (Ap1b1) alternative variant bSep08, mRNA.
<a href="#">Ap1b1</a>	<a href="#">Ap1b1.cSep08</a>	<a href="#">29663</a>	28260	749	7	194	adaptor protein complex AP-1, beta 1 subunit (Ap1b1) alternative variant cSep08, mRNA.

<a href="#">Ap1b1</a>	<a href="#">Ap1b1.dSep08</a>	<a href="#">29663</a>	25544	357	4	71	adaptor protein complex AP-1, beta 1 subunit (Ap1b1) alternative variant dSep08, mRNA.
<a href="#">Ap1gbp1</a>	<a href="#">Ap1gbp1.aSep08</a>	<a href="#">84479</a>	35611	2005	9	600	AP1 gamma subunit binding protein 1 (Ap1gbp1) alternative variant aSep08, mRNA.
<a href="#">Ap1gbp1</a>	<a href="#">Ap1gbp1.cSep08</a>	<a href="#">84479</a>	25815	1915	7	266	AP1 gamma subunit binding protein 1 (Ap1gbp1) alternative variant cSep08, mRNA.
<a href="#">Ap1gbp1</a>	<a href="#">Ap1gbp1.dSep08</a>	<a href="#">84479</a>	21779	556	5	69	AP1 gamma subunit binding protein 1 (Ap1gbp1) alternative variant dSep08, mRNA.
<a href="#">Ap1s1</a>	<a href="#">Ap1s1.bSep08</a>	<a href="#">360785</a>	10447	1302	5	158	adaptor protein complex AP-1, sigma 1 (18.7 kD) (Ap1s1) alternative variant bSep08, complete mRNA.
<a href="#">Ap1s1</a>	<a href="#">Ap1s1.dSep08</a>	<a href="#">360785</a>	6050	711	3	34	adaptor protein complex AP-1, sigma 1 (Ap1s1) alternative variant dSep08, mRNA.
<a href="#">Ap1s2</a>	<a href="#">Ap1s2.aSep08</a>	<a href="#">302671</a>	26182	1169	2	204	adaptor-related protein complex 1, sigma 2 subunit (Ap1s2) alternative variant aSep08, mRNA.
<a href="#">Ap1s2</a>	<a href="#">Ap1s2.cSep08</a>	<a href="#">302671</a>	22904	2025	1	189	adaptor-related protein complex 1, sigma 2 subunit (Ap1s2) alternative variant cSep08, mRNA.
<a href="#">Ap1s2</a>	<a href="#">Ap1s2.dSep08</a>	<a href="#">302671</a>	64012	751	1	185	adaptor-related protein complex 1, sigma 2 subunit (Ap1s2) alternative variant dSep08, mRNA.
<a href="#">Ap2a1</a>	<a href="#">Ap2a1.aSep08</a>	<a href="#">308578</a>	30184	2927	24	972	adaptor protein complex AP-2, alpha 1 subunit (Ap2a1) alternative variant aSep08, mRNA.
<a href="#">Ap2a1</a>	<a href="#">Ap2a1.bSep08</a>	<a href="#">308578</a>	3404	1137	9	299	adaptor protein complex AP-2, alpha 1 subunit (Ap2a1) alternative variant bSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.bSep08</a>	<a href="#">81637</a>	73215	2591	12	631	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant bSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.cSep08</a>	<a href="#">81637</a>	14937	1663	10	554	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant cSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.dSep08</a>	<a href="#">81637</a>	6496	740	5	246	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant dSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.eSep08</a>	<a href="#">81637</a>	2842	1238	5	173	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant eSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.fSep08</a>	<a href="#">81637</a>	3024	512	3	155	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant fSep08, mRNA.
<a href="#">Ap2a2</a>	<a href="#">Ap2a2.hSep08</a>	<a href="#">81637</a>	41549	393	3	93	adaptor protein complex AP-2, alpha 2 subunit (Ap2a2) alternative variant hSep08, mRNA.
<a href="#">Ap2b1</a>	<a href="#">Ap2b1.bSep08</a>	<a href="#">140670</a>	44685	879	6	292	adaptor-related protein complex 2 beta (Ap2b1) alternative variant bSep08, mRNA.
<a href="#">Ap2b1</a>	<a href="#">Ap2b1.cSep08</a>	<a href="#">140670</a>	28757	712	7	200	adaptor-related protein complex 2 beta CRA d (Ap2b1) alternative variant cSep08, mRNA.
<a href="#">Ap2b1</a>	<a href="#">Ap2b1.eSep08</a>	<a href="#">140670</a>	16720	248	4	52	putative protein (Ap2b1) alternative variant eSep08, mRNA.
<a href="#">Ap2b1</a>	<a href="#">Ap2b1.gSep08</a>	<a href="#">140670</a>	13295	602	4	65	putative protein (7.1 kD) (Ap2b1) alternative variant gSep08, mRNA.
<a href="#">Ap2m1</a>	<a href="#">Ap2m1.bSep08</a>	<a href="#">116563</a>	4573	726	3	132	adaptor-related protein complex 2 mu (15.2 kD) (Ap2m1) alternative variant bSep08, mRNA.
<a href="#">Ap2m1</a>	<a href="#">Ap2m1.cSep08</a>	<a href="#">116563</a>	1465	658	5	110	adaptor-related protein complex 2 mu (Ap2m1) alternative variant cSep08, mRNA.

<a href="#">Ap2m1</a>	<a href="#">Ap2m1.dSep08</a>	<a href="#">116563</a>	916	523	2	108	A Mu2 Adaptin Of Ap2 adaptor Complexed With Internalization Peptide (Ap2m1) alternative variant dSep08, mRNA.
<a href="#">Ap2m1</a>	<a href="#">Ap2m1.fSep08</a>	<a href="#">116563</a>	1363	912	2	78	adaptor-related protein complex 2 mu (9.1 kD) (Ap2m1) alternative variant fSep08, mRNA.
<a href="#">Ap2m1</a>	<a href="#">Ap2m1.hSep08</a>	<a href="#">116563</a>	4152	450	4	42	adaptor-related protein complex 2 mu (4.5 kD) (Ap2m1) alternative variant hSep08, mRNA.
<a href="#">Ap2s1</a>	<a href="#">Ap2s1.aSep08</a>	<a href="#">65046</a>	11479	938	5	167	adaptor-related protein complex 2, sigma 1 subunit (20.0 kD) (Ap2s1) alternative variant aSep08, mRNA.
<a href="#">Ap2s1</a>	<a href="#">Ap2s1.cSep08</a>	<a href="#">65046</a>	1952	918	3	128	adaptor-related protein complex 2, sigma 1 subunit (14.9 kD) (Ap2s1) alternative variant cSep08, mRNA.
<a href="#">Ap2s1</a>	<a href="#">Ap2s1.dSep08</a>	<a href="#">65046</a>	7055	645	4	121	adaptor-related protein complex 2, sigma 1 subunit (Ap2s1) alternative variant dSep08, mRNA.
<a href="#">Ap3b1</a>	<a href="#">Ap3b1.bSep08</a>	<a href="#">309969</a>	80324	1965	2	338	adaptor-related protein complex 3, beta 1 subunit (36.4 kD) (Ap3b1) alternative variant bSep08, mRNA.
<a href="#">Ap3b2</a>	<a href="#">Ap3b2.aSep08</a>	<a href="#">308777</a>	16293	2773	20	845	adaptor-related protein complex 3 beta (Ap3b2) alternative variant aSep08, mRNA.
<a href="#">Ap3b2</a>	<a href="#">Ap3b2.cSep08</a>	<a href="#">308777</a>	3210	425	4	137	adaptor-related protein complex 3 beta (Ap3b2) alternative variant cSep08, mRNA.
<a href="#">Ap3b2</a>	<a href="#">Ap3b2.eSep08</a>	<a href="#">308777</a>	748	307	2	101	adaptor-related protein complex 3 beta CRA b (Ap3b2) alternative variant eSep08, mRNA.
<a href="#">Ap3d1</a>	<a href="#">Ap3d1.bSep08</a>	<a href="#">314633</a>	3265	903	6	300	adaptor-related protein complex 3, delta 1 subunit (Ap3d1) alternative variant bSep08, mRNA.
<a href="#">Ap3d1</a>	<a href="#">Ap3d1.cSep08</a>	<a href="#">314633</a>	2825	647	4	210	adaptor-related protein complex 3, delta 1 subunit (Ap3d1) alternative variant cSep08, mRNA.
<a href="#">Ap3d1</a>	<a href="#">Ap3d1.dSep08</a>	<a href="#">314633</a>	2068	829	5	191	adaptor-related protein complex 3, delta 1 subunit (Ap3d1) alternative variant dSep08, mRNA.
<a href="#">Ap3d1</a>	<a href="#">Ap3d1.eSep08</a>	<a href="#">314633</a>	8755	498	6	73	adaptor-related protein complex 3, delta 1 subunit (Ap3d1) alternative variant eSep08, mRNA.
<a href="#">Ap3m1</a>	<a href="#">Ap3m1.bSep08</a>	<a href="#">171126</a>	11396	717	4	238	adaptor-related protein complex 3, mu 1 subunit (Ap3m1) alternative variant bSep08, mRNA.
<a href="#">Ap3m1</a>	<a href="#">Ap3m1.cSep08</a>	<a href="#">171126</a>	739	639	2	28	adaptor-related protein complex 3, mu 1 subunit (3.0 kD) (Ap3m1) alternative variant cSep08, mRNA.
<a href="#">Ap3m2</a>	<a href="#">Ap3m2.bSep08</a>	<a href="#">140667</a>	6560	980	3	296	adaptor-related protein complex 3 mu (Ap3m2) alternative variant bSep08, mRNA.
<a href="#">Ap3m2</a>	<a href="#">Ap3m2.cSep08</a>	<a href="#">140667</a>	4652	597	2	181	putative protein of eukaryotic origin (Ap3m2) alternative variant cSep08, mRNA.
<a href="#">Ap3m2</a>	<a href="#">Ap3m2.dSep08</a>	<a href="#">140667</a>	13891	775	5	152	adaptor-related protein complex 3 mu (17.6 kD) (Ap3m2) alternative variant dSep08, mRNA.
<a href="#">Ap3m2</a>	<a href="#">Ap3m2.eSep08</a>	<a href="#">140667</a>	5700	663	3	146	adaptor-related protein complex 3 mu (Ap3m2) alternative variant eSep08, mRNA.
<a href="#">Ap3m2</a>	<a href="#">Ap3m2.fSep08</a>	<a href="#">140667</a>	1888	485	2	102	putative protein (Ap3m2) alternative variant fSep08, mRNA.
<a href="#">Ap3s1</a>	<a href="#">Ap3s1.aSep08</a>	<a href="#">302290</a>	72070	1255	6	225	adaptor-related protein complex 3, sigma 1 subunit (Ap3s1) alternative variant aSep08, mRNA.
<a href="#">Ap3s1</a>	<a href="#">Ap3s1.cSep08</a>	<a href="#">302290</a>	71742	968	7	162	adaptor-related protein complex 3, sigma 1 subunit (18.8 kD) (Ap3s1) alternative variant cSep08, complete mRNA.

<a href="#">Ap3s1</a>	<a href="#">Ap3s1.dSep08</a>	<a href="#">302290</a>	52775	557	4	134	adaptor-related protein complex 3, sigma 1 subunit (Ap3s1) alternative variant dSep08, mRNA.
<a href="#">Ap4b1</a>	<a href="#">Ap4b1.bSep08</a>	<a href="#">310746</a>	11984	1782	5	321	adaptor-related protein complex AP-4, beta 1 (Ap4b1) alternative variant bSep08, mRNA.
<a href="#">Ap4b1</a>	<a href="#">Ap4b1.cSep08</a>	<a href="#">310746</a>	3587	731	5	174	adaptor-related protein complex AP-4, beta 1 (Ap4b1) alternative variant cSep08, mRNA.
<a href="#">Ap4e1</a>	<a href="#">Ap4e1.aSep08</a>	<a href="#">311404</a>	5025	1028		277	adaptor-related protein complex AP-4, epsilon 1 (Ap4e1) alternative variant aSep08, mRNA.
<a href="#">Ap4e1</a>	<a href="#">Ap4e1.bSep08</a>	<a href="#">311404</a>	4732	734		145	adaptor-related protein complex AP-4, epsilon 1 (Ap4e1) alternative variant bSep08, mRNA.
<a href="#">Ap4m1</a>	<a href="#">Ap4m1.bSep08</a>	<a href="#">304344</a>	8239	3845	16	325	adaptor-related protein complex AP-4, mu 1 (35.6 kD) (Ap4m1) alternative variant bSep08, complete mRNA.
<a href="#">Ap4m1</a>	<a href="#">Ap4m1.cSep08</a>	<a href="#">304344</a>	1426	380	5	126	adaptor-related protein complex AP-4, mu 1 (Ap4m1) alternative variant cSep08, mRNA.
<a href="#">Ap4m1</a>	<a href="#">Ap4m1.dSep08</a>	<a href="#">304344</a>	1324	380	3	84	adaptor-related protein complex AP-4, mu 1 (9.5 kD) (Ap4m1) alternative variant dSep08, mRNA.
<a href="#">Ap4m1</a>	<a href="#">Ap4m1.eSep08</a>	<a href="#">304344</a>	314	214	2	70	adaptor-related protein complex AP-4, mu 1 (Ap4m1) alternative variant eSep08, mRNA.
<a href="#">Ap4s1</a>	<a href="#">Ap4s1.bSep08</a>	<a href="#">366618</a>	42170	1321	3	101	adaptor-related protein complex AP-4, sigma 1 (12.1 kD) (Ap4s1) alternative variant bSep08, complete mRNA.
<a href="#">Ap4s1</a>	<a href="#">Ap4s1.cSep08</a>	<a href="#">366618</a>	41048	584	2	67	adaptor-related protein complex AP-4, sigma 1 (7.8 kD) (Ap4s1) alternative variant cSep08, complete mRNA.
<a href="#">Apaf1</a>	<a href="#">Apaf1.bSep08</a>	<a href="#">78963</a>	36162	1801	1	261	apoptotic peptidase activating factor 1 (Apaf1) alternative variant bSep08, mRNA.
<a href="#">Apaf1</a>	<a href="#">Apaf1.cSep08</a>	<a href="#">78963</a>	5252	2026	1	97	apoptotic peptidase activating factor 1 (Apaf1) alternative variant cSep08, mRNA.
<a href="#">Apba1</a>	<a href="#">Apba1.bSep08</a>	<a href="#">83589</a>	25066	814	7	271	amyloid beta (A4) precursor protein-binding, family A, member 1 (Apba1) alternative variant bSep08, mRNA.
<a href="#">Apba1</a>	<a href="#">Apba1.cSep08</a>	<a href="#">83589</a>	21615	398	5	132	amyloid beta (A4) precursor protein-binding, family A, member 1 (Apba1) alternative variant cSep08, mRNA.
<a href="#">Apba2</a>	<a href="#">Apba2.bSep08</a>	<a href="#">83610</a>	13955	775	6	258	amyloid beta protein-binding family A member 2 CRA b like (Apba2) alternative variant bSep08, mRNA.
<a href="#">Apba2</a>	<a href="#">Apba2.cSep08</a>	<a href="#">83610</a>	67388	759	2	125	putative protein (Apba2) alternative variant cSep08, mRNA.
<a href="#">Apba3</a>	<a href="#">Apba3.bSep08</a>	<a href="#">83611</a>	2097	1289	6	379	amyloid beta (A4) precursor protein-binding, family A, member 3 (41.2 kD) (Apba3) alternative variant bSep08, mRNA.
<a href="#">Apba3</a>	<a href="#">Apba3.cSep08</a>	<a href="#">83611</a>	2999	878	5	251	amyloid beta (A4) precursor protein-binding, family A, member 3 (Apba3) alternative variant cSep08, mRNA.
<a href="#">Apba3</a>	<a href="#">Apba3.dSep08</a>	<a href="#">83611</a>	1792	1087	5	217	amyloid beta (A4) precursor protein-binding, family A, member 3 (Apba3) alternative variant dSep08, mRNA.
<a href="#">Apbb1</a>	<a href="#">Apbb1.bSep08</a>	<a href="#">29722</a>	7779	760	1	252	amyloid beta (A4) precursor protein-binding, family B, member 1 (Apbb1) alternative variant bSep08, mRNA.
<a href="#">Apbb1</a>	<a href="#">Apbb1.cSep08</a>	<a href="#">29722</a>	2060	311	1	63	amyloid beta (A4) precursor protein-binding, family B, member 1 (Apbb1) alternative variant cSep08, mRNA.
<a href="#">Apbb1ip</a>	<a href="#">Apbb1ip.aSep08</a>	<a href="#">307171</a>	47795	2429		645	amyloid beta (A4) precursor protein-binding, family B, member 1 interacting protein (Apbb1ip) mRNA.

<a href="#">Apbb2</a>	<a href="#">Apbb2.aSep08</a>	<a href="#">305338</a>	33799	4130	7	265	amyloid beta (A4) precursor protein-binding, family B, member 2 (Apbb2) alternative variant aSep08, mRNA.
<a href="#">Apbb2</a>	<a href="#">Apbb2.bSep08</a>	<a href="#">305338</a>	5870	439	5	146	amyloid beta (A4) precursor protein-binding, family B, member 2 (Apbb2) alternative variant bSep08, mRNA.
<a href="#">Apbb2</a>	<a href="#">Apbb2.cSep08</a>	<a href="#">305338</a>	10451	675	2	50	amyloid beta (A4) precursor protein-binding, family B, member 2 (Apbb2) alternative variant cSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.cSep08</a>	<a href="#">117026</a>	3253	1019	2	182	amyloid beta protein-binding family B member 3 CRA b like precursor (19.1 kD) (Apbb3andSra1) alternative variant cSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.cSep08</a>	<a href="#">252891</a>	3253	1019	2	182	amyloid beta protein-binding family B member 3 CRA b like precursor (19.1 kD) (Apbb3andSra1) alternative variant cSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.dSep08</a>	<a href="#">117026</a>	2319	742	3	147	steroid receptor RNA activator 1 (15.3 kD) (Apbb3andSra1) alternative variant dSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.dSep08</a>	<a href="#">252891</a>	2319	742	3	147	steroid receptor RNA activator 1 (15.3 kD) (Apbb3andSra1) alternative variant dSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.eSep08</a>	<a href="#">117026</a>	1445	723	5	94	amyloid beta protein-binding family B member 3 like (Apbb3andSra1) alternative variant eSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.eSep08</a>	<a href="#">252891</a>	1445	723	5	94	amyloid beta protein-binding family B member 3 like (Apbb3andSra1) alternative variant eSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.gSep08</a>	<a href="#">117026</a>	1642	1514	2	74	steroid receptor RNA activator 1 CRA b (8.4 kD) (Apbb3andSra1) alternative variant gSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.gSep08</a>	<a href="#">252891</a>	1642	1514	2	74	steroid receptor RNA activator 1 CRA b (8.4 kD) (Apbb3andSra1) alternative variant gSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.hSep08</a>	<a href="#">117026</a>	1284	576	3	64	steroid receptor RNA activator 1 (Apbb3andSra1) alternative variant hSep08, mRNA.
<a href="#">Apbb3andSra1</a>	<a href="#">Apbb3andSra1.hSep08</a>	<a href="#">252891</a>	1284	576	3	64	steroid receptor RNA activator 1 (Apbb3andSra1) alternative variant hSep08, mRNA.
<a href="#">Apc</a>	<a href="#">Apc.bSep08</a>	<a href="#">24205</a>	51250	823	6	235	adenomatosis polyposis coli (26.6 kD) (Apc) alternative variant bSep08, mRNA.
<a href="#">Apccd1</a>	<a href="#">Apccd1.aSep08</a>	<a href="#">689616</a>	2435	873		215	adenomatosis polyposis coli down-regulated 1 (Apccd1) mRNA.
<a href="#">Apeh</a>	<a href="#">Apeh.bSep08</a>	<a href="#">24206</a>	5138	1464	10	304	acylpeptide hydrolase (33.9 kD) (Apeh) alternative variant bSep08, mRNA.
<a href="#">Apeh</a>	<a href="#">Apeh.dSep08</a>	<a href="#">24206</a>	555	480	2	50	acylpeptide hydrolase (5.4 kD) (Apeh) alternative variant dSep08, mRNA.
<a href="#">Apex1</a>	<a href="#">Apex1.aSep08</a>	<a href="#">79116</a>	2567	1950	2	347	apurinic/apurimidinic endonuclease 1 (38.9 kD) (Apex1) alternative variant aSep08, mRNA.
<a href="#">Apex1</a>	<a href="#">Apex1.cSep08</a>	<a href="#">79116</a>	1668	785	4	228	apurinic/apurimidinic endonuclease 1 (Apex1) alternative variant cSep08, mRNA.
<a href="#">Aph1a</a>	<a href="#">Aph1a.bSep08</a>	<a href="#">365872</a>	435	286	1	95	anterior pharynx defective 1 homolog a (Aph1a) alternative variant bSep08, mRNA.
<a href="#">Aph1b.1</a>	<a href="#">Aph1b.1.aSep08</a>	<a href="#">300802</a>	21014	1249	6	292	anterior pharynx defective 1b homolog (Aph1b.1) alternative variant aSep08, mRNA.
<a href="#">Aph1b.1</a>	<a href="#">Aph1b.1.bSep08</a>	<a href="#">300802</a>	12059	628	5	152	anterior pharynx defective 1b homolog (Aph1b.1) alternative variant bSep08, mRNA.

<a href="#">Aph1b.1</a>	<a href="#">Aph1b.1.cSep08</a>	<a href="#">300802</a>	7904	960	2	96	anterior pharynx defective 1b homolog (11.1 kD) (Aph1b.1) alternative variant cSep08, mRNA.
<a href="#">Api5</a>	<a href="#">Api5.bSep08</a>	<a href="#">362170</a>	7601	2407	5	145	apoptosis inhibitor 5 (Api5) alternative variant bSep08, mRNA.
<a href="#">Api5</a>	<a href="#">Api5.cSep08</a>	<a href="#">362170</a>	9882	813	2	80	apoptosis inhibitor 5 (9.0 kD) (Api5) alternative variant cSep08, mRNA.
<a href="#">Api5</a>	<a href="#">Api5.dSep08</a>	<a href="#">362170</a>	2423	713	2	64	apoptosis inhibitor 5 (Api5) alternative variant dSep08, mRNA.
<a href="#">Apip</a>	<a href="#">Apip.bSep08</a>	<a href="#">295961</a>	8727	1032	5	165	APAF1 interacting protein (Apip) alternative variant bSep08, mRNA.
<a href="#">Aplp1</a>	<a href="#">Aplp1.aSep08</a>	<a href="#">502317</a>	8613	1671	3	556	amyloid beta (A4) precursor-like protein 1 (Aplp1) alternative variant aSep08, mRNA.
<a href="#">Aplp1</a>	<a href="#">Aplp1.bSep08</a>	<a href="#">502317</a>	6001	1268	2	323	amyloid beta (A4) precursor-like protein 1 (Aplp1) alternative variant bSep08, mRNA.
<a href="#">Aplp1</a>	<a href="#">Aplp1.cSep08</a>	<a href="#">502317</a>	3324	782	1	260	amyloid beta (A4) precursor-like protein 1 (Aplp1) alternative variant cSep08, mRNA.
<a href="#">Aplp1</a>	<a href="#">Aplp1.dSep08</a>	<a href="#">502317</a>	1404	512	1	106	amyloid beta (A4) precursor-like protein 1 (Aplp1) alternative variant dSep08, mRNA.
<a href="#">Aplp2</a>	<a href="#">Aplp2.aSep08</a>	<a href="#">64312</a>	22454	3141	11	631	amyloid beta (A4) precursor-like protein 2 (Aplp2) alternative variant aSep08, mRNA.
<a href="#">Aplp2</a>	<a href="#">Aplp2.bSep08</a>	<a href="#">64312</a>	21424	1907	9	552	amyloid beta (A4) precursor-like protein 2 (Aplp2) alternative variant bSep08, mRNA.
<a href="#">Aplp2</a>	<a href="#">Aplp2.cSep08</a>	<a href="#">64312</a>	19361	397	2	132	amyloid beta (A4) precursor-like protein 2 (Aplp2) alternative variant cSep08, mRNA.
<a href="#">Aplp2</a>	<a href="#">Aplp2.dSep08</a>	<a href="#">64312</a>	3398	858	2	68	amyloid beta (A4) precursor-like protein 2 (Aplp2) alternative variant dSep08, mRNA.
<a href="#">Apoa1bp</a>	<a href="#">Apoa1bp.aSep08</a>	<a href="#">295229</a>	2052	921	6	282	apolipoprotein A-I binding protein (30.9 kD) (Apoa1bp) alternative variant aSep08, complete mRNA.
<a href="#">Apoa1bp</a>	<a href="#">Apoa1bp.cSep08</a>	<a href="#">295229</a>	1304	1030	4	169	apolipoprotein A-I binding protein (Apoa1bp) alternative variant cSep08, mRNA.
<a href="#">Apoa1bp</a>	<a href="#">Apoa1bp.dSep08</a>	<a href="#">295229</a>	953	865	2	139	apolipoprotein A-I binding protein (Apoa1bp) alternative variant dSep08, mRNA.
<a href="#">Apoa2</a>	<a href="#">Apoa2.aSep08</a>	<a href="#">25649</a>	1723	497	1	129	apolipoprotein A-II (Apoa2) alternative variant aSep08, mRNA.
<a href="#">Apoa2</a>	<a href="#">Apoa2.bSep08</a>	<a href="#">25649</a>	1844	465	2	117	apolipoprotein A-II (Apoa2) alternative variant bSep08, mRNA.
<a href="#">Apob48r</a>	<a href="#">Apob48r.bSep08</a>	<a href="#">499264</a>	4178	1181		137	apolipoprotein B48 receptor (Apob48r) alternative variant bSep08, mRNA.
<a href="#">Apobec1</a>	<a href="#">Apobec1.aSep08</a>	<a href="#">25383</a>	3245	1216	1	89	apolipoprotein B editing complex 1 (10.6 kD) (Apobec1) alternative variant aSep08, mRNA.
<a href="#">Apobec1</a>	<a href="#">Apobec1.bSep08</a>	<a href="#">25383</a>	26648	1389	2	86	apolipoprotein B editing complex 1 (10.3 kD) (Apobec1) alternative variant bSep08, mRNA.
<a href="#">Apobec3</a>	<a href="#">Apobec3.bSep08</a>	<a href="#">315137</a>	10233	774	4	172	apolipoprotein B editing complex 3 (Apobec3) alternative variant bSep08, mRNA.
<a href="#">Apoe</a>	<a href="#">Apoe.bSep08</a>	<a href="#">25728</a>	2446	820	4	273	apolipoprotein E (Apoe) alternative variant bSep08, mRNA.
<a href="#">Apoe</a>	<a href="#">Apoe.cSep08</a>	<a href="#">25728</a>	2104	599	4	173	apolipoprotein E (Apoe) alternative variant cSep08, mRNA.
<a href="#">Apol8</a>	<a href="#">Apol8.aSep08</a>	<a href="#">315111</a>	5695	557		185	apolipoprotein L 8 (Apol8) mRNA.

<a href="#">Apol9a</a>	<a href="#">Apol9a.aSep08</a>	<a href="#">503164</a>	16599	1849	5	366	apolipoprotein L 7 (39.9 kD) (Apol9a) alternative variant aSep08, mRNA.
<a href="#">Apol9a</a>	<a href="#">Apol9a.cSep08</a>	<a href="#">503164</a>	15382	729	5	191	apolipoprotein L 7 (Apol9a) alternative variant cSep08, mRNA.
<a href="#">Apol9a</a>	<a href="#">Apol9a.dSep08</a>	<a href="#">503164</a>	17452	750	6	154	apolipoprotein L 7 (Apol9a) alternative variant dSep08, mRNA.
<a href="#">Apol9a</a>	<a href="#">Apol9a.eSep08</a>	<a href="#">503164</a>	15284	756	6	121	apolipoprotein L 7 (Apol9a) alternative variant eSep08, mRNA.
<a href="#">Apol9a</a>	<a href="#">Apol9a.fSep08</a>	<a href="#">503164</a>	1198	735	2	66	apolipoprotein L 9 (Apol9a) alternative variant fSep08, mRNA.
<a href="#">Apol9a</a>	<a href="#">Apol9a.gSep08</a>	<a href="#">503164</a>	12435	788	3	50	apolipoprotein L 7e like (5.8 kD) (Apol9a) alternative variant gSep08, mRNA.
<a href="#">Apom</a>	<a href="#">Apom.bSep08</a>	<a href="#">55939</a>	863	597	1	90	apolipoprotein M (10.2 kD) (Apom) alternative variant bSep08, mRNA.
<a href="#">Apool</a>	<a href="#">Apool.aSep08</a>	<a href="#">317191</a>	68127	1965	1	474	apolipoprotein O-like (Apool) alternative variant aSep08, mRNA.
<a href="#">App</a>	<a href="#">App.bSep08</a>	<a href="#">54226</a>	136655	1868	11	603	amyloid beta protein (App) alternative variant bSep08, mRNA.
<a href="#">App</a>	<a href="#">App.cSep08</a>	<a href="#">54226</a>	145602	2562	12	554	amyloid protein (App) alternative variant cSep08, mRNA.
<a href="#">App</a>	<a href="#">App.dSep08</a>	<a href="#">54226</a>	50952	550	5	178	amyloid protein (App) alternative variant dSep08, mRNA.
<a href="#">App</a>	<a href="#">App.eSep08</a>	<a href="#">54226</a>	51464	404	2	134	amyloid protein (App) alternative variant eSep08, mRNA.
<a href="#">App</a>	<a href="#">App.fSep08</a>	<a href="#">54226</a>	6277	648	2	111	amyloid beta protein (12.4 kD) (App) alternative variant fSep08, mRNA.
<a href="#">App</a>	<a href="#">App.gSep08</a>	<a href="#">54226</a>	12107	419	3	86	amyloid protein (App) alternative variant gSep08, mRNA.
<a href="#">Appl2</a>	<a href="#">Appl2.aSep08</a>	<a href="#">362860</a>	49153	2962	21	662	adaptor protein, phosphotyrosine interaction, PH domain and leucine zipper containing 2 (74.1 kD) (Appl2) alternative variant aSep08, complete mRNA.
<a href="#">Appl2</a>	<a href="#">Appl2.bSep08</a>	<a href="#">362860</a>	9555	1123	9	374	adaptor protein, phosphotyrosine interaction, PH domain and leucine zipper containing 2 (Appl2) alternative variant bSep08, mRNA.
<a href="#">Appl2</a>	<a href="#">Appl2.cSep08</a>	<a href="#">362860</a>	5165	536	6	178	adaptor protein, phosphotyrosine interaction, PH domain and leucine zipper containing 2 (Appl2) alternative variant cSep08, mRNA.
<a href="#">Appl2</a>	<a href="#">Appl2.dSep08</a>	<a href="#">362860</a>	3486	1799	3	55	adaptor protein, phosphotyrosine interaction, PH domain and leucine zipper containing 2 (6.1 kD) (Appl2) alternative variant dSep08, mRNA.
<a href="#">Appl2</a>	<a href="#">Appl2.eSep08</a>	<a href="#">362860</a>	2462	561	3	48	adaptor protein, phosphotyrosine interaction, PH domain and leucine zipper containing 2 (Appl2) alternative variant eSep08, mRNA.
<a href="#">Aprt</a>	<a href="#">Aprt.bSep08</a>	<a href="#">292072</a>	2138	843	4	117	adenine phosphoribosyl transferase (12.6 kD) (Aprt) alternative variant bSep08, mRNA.
<a href="#">Aqp2</a>	<a href="#">Aqp2.bSep08</a>	<a href="#">25386</a>	3419	487	1	108	aquaporin 2 (Aqp2) alternative variant bSep08, mRNA.
<a href="#">Aqp4</a>	<a href="#">Aqp4.bSep08</a>	<a href="#">25293</a>	4406	617	3	205	aquaporin 4 (Aqp4) alternative variant bSep08, mRNA.
<a href="#">Aqp5</a>	<a href="#">Aqp5.bSep08</a>	<a href="#">25241</a>	2071	993	1	134	aquaporin 5 (14.9 kD) (Aqp5) alternative variant bSep08, mRNA.
<a href="#">Aqp7</a>	<a href="#">Aqp7.bSep08</a>	<a href="#">29171</a>	9160	388	1	129	aquaporin 7 (Aqp7) alternative variant bSep08, mRNA.

<a href="#">Aqp8</a>	<a href="#">Aqp8.bSep08</a>	<a href="#">29172</a>	3059	751	4	202	aquaporin 8 (21.3 kD) (Aqp8) alternative variant bSep08, mRNA.
<a href="#">Aqp9</a>	<a href="#">Aqp9.bSep08</a>	<a href="#">65054</a>	31519	972	5	237	aquaporin 9 (Aqp9) alternative variant bSep08, mRNA.
<a href="#">Aqr</a>	<a href="#">Aqr.bSep08</a>	<a href="#">366163</a>	7275	558	3	147	aquarius (Aqr) alternative variant bSep08, mRNA.
<a href="#">Aqr</a>	<a href="#">Aqr.cSep08</a>	<a href="#">366163</a>	1838	810	2	102	aquarius (Aqr) alternative variant cSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.bSep08</a>	<a href="#">64363</a>	4960	1091	7	363	v-raf murine sarcoma 3611 viral oncogene homolog (Araf) alternative variant bSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.cSep08</a>	<a href="#">64363</a>	2395	912	6	295	v-raf murine sarcoma 3611 viral oncogene homolog (Araf) alternative variant cSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.dSep08</a>	<a href="#">64363</a>	4551	874	5	291	v-raf murine sarcoma 3611 viral oncogene homolog (Araf) alternative variant dSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.eSep08</a>	<a href="#">64363</a>	7107	1036	7	219	v-raf murine sarcoma 3611 viral oncogene homolog (24.7 kD) (Araf) alternative variant eSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.gSep08</a>	<a href="#">64363</a>	2136	929	4	172	v-raf murine sarcoma 3611 viral oncogene homolog (Araf) alternative variant gSep08, mRNA.
<a href="#">Araf</a>	<a href="#">Araf.hSep08</a>	<a href="#">64363</a>	3742	1052	6	170	v-raf murine sarcoma 3611 viral oncogene homolog (19.0 kD) (Araf) alternative variant hSep08, complete mRNA.
<a href="#">Araf</a>	<a href="#">Araf.iSep08</a>	<a href="#">64363</a>	3482	925	5	153	v-raf murine sarcoma 3611 viral oncogene homolog (17.2 kD) (Araf) alternative variant iSep08, complete mRNA.
<a href="#">Araf</a>	<a href="#">Araf.kSep08</a>	<a href="#">64363</a>	1220	1056	2	62	v-raf murine sarcoma 3611 viral oncogene homolog (Araf) alternative variant kSep08, mRNA.
<a href="#">Arbp</a>	<a href="#">Arbp.bSep08</a>	<a href="#">64205</a>	2232	861	5	271	acidic ribosomal phosphoprotein P0 (Arbp) alternative variant bSep08, mRNA.
<a href="#">Arcn1</a>	<a href="#">Arcn1.bSep08</a>	<a href="#">300674</a>	11372	755	4	204	archain 1 (Arcn1) alternative variant bSep08, mRNA.
<a href="#">Ard1</a>	<a href="#">Ard1.aSep08</a>	<a href="#">363518</a>	5132	873	8	265	N-acetyltransferase ARD1 homolog (S. cerevisiae) (Ard1) alternative variant aSep08, mRNA.
<a href="#">Ard1</a>	<a href="#">Ard1.bSep08</a>	<a href="#">363518</a>	4163	758	6	185	N-acetyltransferase ARD1 homolog (S. cerevisiae) (Ard1) alternative variant bSep08, mRNA.
<a href="#">Ard1</a>	<a href="#">Ard1.cSep08</a>	<a href="#">363518</a>	2815	543	3	154	N-acetyltransferase ARD1 homolog (S. cerevisiae) (Ard1) alternative variant cSep08, mRNA.
<a href="#">Ard1</a>	<a href="#">Ard1.dSep08</a>	<a href="#">363518</a>	2749	568	3	121	N-acetyltransferase ARD1 homolog (S. cerevisiae) (Ard1) alternative variant dSep08, mRNA.
<a href="#">Ard1</a>	<a href="#">Ard1.fSep08</a>	<a href="#">363518</a>	2312	759	2	68	N-acetyltransferase ARD1 homolog (S. cerevisiae) (Ard1) alternative variant fSep08, mRNA.
<a href="#">Arf.0</a>	<a href="#">Arf.0.aSep08</a>		3522	2404		97	tripartite motif protein 23 CRA a (Arf.0) mRNA.
<a href="#">Arf1</a>	<a href="#">Arf1.aSep08</a>	<a href="#">64310</a>	16476	1774	5	214	ADP-ribosylation factor 1 (Arf1) alternative variant aSep08, mRNA.
<a href="#">Arf1</a>	<a href="#">Arf1.bSep08</a>	<a href="#">64310</a>	15425	786	6	195	ADP-ribosylation factor 1 (22.4 kD) (Arf1) alternative variant bSep08, mRNA.
<a href="#">Arf1</a>	<a href="#">Arf1.dSep08</a>	<a href="#">64310</a>	15827	1014	4	144	ADP-ribosylation factor 1 (16.2 kD) (Arf1) alternative variant dSep08, mRNA.
<a href="#">Arf1</a>	<a href="#">Arf1.eSep08</a>	<a href="#">64310</a>	14716	405	2	75	ADP-ribosylation factor 1 (Arf1) alternative variant eSep08, mRNA.
<a href="#">Arf2</a>	<a href="#">Arf2.aSep08</a>	<a href="#">79119</a>	20513	1316	2	189	ADP-ribosylation factor 2 (21.6 kD) (Arf2) alternative variant aSep08, mRNA.



<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.aSep08</a>	<a href="#">140940</a>	3437	720	6	201	FK506 binding protein 11 like (22.2 kD) (Arf3andFkbp11) alternative variant aSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.aSep08</a>	<a href="#">300211</a>	3437	720	6	201	FK506 binding protein 11 like (22.2 kD) (Arf3andFkbp11) alternative variant aSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.cSep08</a>	<a href="#">140940</a>	3404	516	3	144	FK506 binding protein 11 like (16.0 kD) (Arf3andFkbp11) alternative variant cSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.cSep08</a>	<a href="#">300211</a>	3404	516	3	144	FK506 binding protein 11 like (16.0 kD) (Arf3andFkbp11) alternative variant cSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.dSep08</a>	<a href="#">140940</a>	2719	450	5	142	FK506 binding protein 11 like (Arf3andFkbp11) alternative variant dSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.dSep08</a>	<a href="#">300211</a>	2719	450	5	142	FK506 binding protein 11 like (Arf3andFkbp11) alternative variant dSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.eSep08</a>	<a href="#">140940</a>	17968	588	4	123	ADP-ribosylation factor (Arf3andFkbp11) alternative variant eSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.eSep08</a>	<a href="#">300211</a>	17968	588	4	123	ADP-ribosylation factor (Arf3andFkbp11) alternative variant eSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.gSep08</a>	<a href="#">140940</a>	1275	535	2	88	putative protein human specific (Arf3andFkbp11) alternative variant gSep08, mRNA.
<a href="#">Arf3andFkbp11</a>	<a href="#">Arf3andFkbp11.gSep08</a>	<a href="#">300211</a>	1275	535	2	88	putative protein human specific (Arf3andFkbp11) alternative variant gSep08, mRNA.
<a href="#">Arf4</a>	<a href="#">Arf4.aSep08</a>	<a href="#">79120</a>	13682	897	5	186	ADP-ribosylation factor 4 (20.9 kD) (Arf4) alternative variant aSep08, mRNA.
<a href="#">Arf4l</a>	<a href="#">Arf4l.bSep08</a>	<a href="#">303559</a>	7470	500	2	48	ADP-ribosylation factor 4-like (Arf4l) alternative variant bSep08, mRNA.
<a href="#">Arf4l</a>	<a href="#">Arf4l.cSep08</a>	<a href="#">303559</a>	13066	296	2	49	ADP-ribosylation factor 4-like (Arf4l) alternative variant cSep08, mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.bSep08</a>	<a href="#">246310</a>	26715	2964	12	330	ADP-ribosylation factor GTPase activating protein 1 (36.0 kD) (Arfgap1) alternative variant bSep08, complete mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.cSep08</a>	<a href="#">246310</a>	5402	685	6	204	ADP-ribosylation factor GTPase activating protein 1 (Arfgap1) alternative variant cSep08, mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.dSep08</a>	<a href="#">246310</a>	18660	535	7	178	ADP-ribosylation factor GTPase activating protein 1 (Arfgap1) alternative variant dSep08, mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.eSep08</a>	<a href="#">246310</a>	2211	1386	2	103	ADP-ribosylation factor GTPase activating protein 1 (Arfgap1) alternative variant eSep08, mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.fSep08</a>	<a href="#">246310</a>	2594	1633	3	59	ADP-ribosylation factor GTPase activating protein 1 (6.5 kD) (Arfgap1) alternative variant fSep08, mRNA.
<a href="#">Arfgap1</a>	<a href="#">Arfgap1.hSep08</a>	<a href="#">246310</a>	567	358	2	88	ADP-ribosylation factor GTPase activating protein 1 (Arfgap1) alternative variant hSep08, mRNA.
<a href="#">Arfgap2</a>	<a href="#">Arfgap2.bSep08</a>	<a href="#">362162</a>	2751	503		139	ADP-ribosylation factor GTPase activating protein 2 (Arfgap2) alternative variant bSep08, mRNA.
<a href="#">Arfgef1</a>	<a href="#">Arfgef1.aSep08</a>	<a href="#">312915</a>	43731	4131	22	958	ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-inhibited) (Arfgef1) alternative variant aSep08, mRNA.
<a href="#">Arfgef1</a>	<a href="#">Arfgef1.bSep08</a>	<a href="#">312915</a>	4874	885	4	230	ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-inhibited) (Arfgef1) alternative variant bSep08, mRNA.

<a href="#">Arfp1</a>	<a href="#">Arfp1.bSep08</a>	<a href="#">60382</a>	53150	472	5	157	ADP-ribosylation factor interacting protein 1 (Arfp1) alternative variant bSep08, mRNA.
<a href="#">Arfp1</a>	<a href="#">Arfp1.cSep08</a>	<a href="#">60382</a>	53120	584	6	119	ADP-ribosylation factor interacting protein 1 (Arfp1) alternative variant cSep08, mRNA.
<a href="#">Arfp1</a>	<a href="#">Arfp1.dSep08</a>	<a href="#">60382</a>	16175	1846	2	62	ADP-ribosylation factor interacting protein 1 (Arfp1) alternative variant dSep08, mRNA.
<a href="#">Arfp2</a>	<a href="#">Arfp2.bSep08</a>	<a href="#">293344</a>	3198	1008	6	233	ADP-ribosylation factor interacting protein 2 (25.0 kD) (Arfp2) alternative variant bSep08, mRNA.
<a href="#">Arfp2</a>	<a href="#">Arfp2.cSep08</a>	<a href="#">293344</a>	3068	671	6	191	ADP-ribosylation factor interacting protein 2 (Arfp2) alternative variant cSep08, mRNA.
<a href="#">Arfp2</a>	<a href="#">Arfp2.dSep08</a>	<a href="#">293344</a>	2527	663	5	188	ADP-ribosylation factor interacting protein 2 (Arfp2) alternative variant dSep08, mRNA.
<a href="#">Arfp2</a>	<a href="#">Arfp2.eSep08</a>	<a href="#">293344</a>	1847	1000	3	116	ADP-ribosylation factor interacting protein 2 (Arfp2) alternative variant eSep08, mRNA.
<a href="#">Arfp2</a>	<a href="#">Arfp2.hSep08</a>	<a href="#">293344</a>	1413	396	3	87	ADP-ribosylation factor interacting protein 2 (9.0 kD) (Arfp2) alternative variant hSep08, complete mRNA.
<a href="#">Arfrp1</a>	<a href="#">Arfrp1.bSep08</a>	<a href="#">117051</a>	5555	628	6	157	ADP-ribosylation factor related protein 1 (Arfrp1) alternative variant bSep08, mRNA.
<a href="#">Arfrp1</a>	<a href="#">Arfrp1.cSep08</a>	<a href="#">117051</a>	7534	2368	5	126	ADP-ribosylation factor related protein 1 (14.5 kD) (Arfrp1) alternative variant cSep08, complete mRNA.
<a href="#">Arfrp1</a>	<a href="#">Arfrp1.dSep08</a>	<a href="#">117051</a>	5895	830	6	121	ADP-ribosylation factor related protein 1 (13.9 kD) (Arfrp1) alternative variant dSep08, mRNA.
<a href="#">Arfrp1</a>	<a href="#">Arfrp1.eSep08</a>	<a href="#">117051</a>	5816	1783	4	83	ADP-ribosylation factor related protein 1 (Arfrp1) alternative variant eSep08, mRNA.
<a href="#">Arfrp1</a>	<a href="#">Arfrp1.fSep08</a>	<a href="#">117051</a>	786	713	2	57	ADP-ribosylation factor related protein 1 (Arfrp1) alternative variant fSep08, mRNA.
<a href="#">Argbp2</a>	<a href="#">Argbp2.aSep08</a>	<a href="#">114901</a>	53630	3696	11	514	arg/Abl-interacting protein ArgBP2 (Argbp2) alternative variant aSep08, mRNA.
<a href="#">Argbp2</a>	<a href="#">Argbp2.bSep08</a>	<a href="#">114901</a>	33694	718	7	238	arg/Abl-interacting protein ArgBP2 (Argbp2) alternative variant bSep08, mRNA.
<a href="#">Argbp2</a>	<a href="#">Argbp2.cSep08</a>	<a href="#">114901</a>	22949	917	2	199	arg/Abl-interacting protein ArgBP2 (Argbp2) alternative variant cSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.aSep08</a>	<a href="#">311193</a>	21934	2880	13	484	rho GTPase activating protein 1 (Arhgap1) alternative variant aSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.bSep08</a>	<a href="#">311193</a>	1976	930	6	253	rho GTPase activating protein 1 (Arhgap1) alternative variant bSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.cSep08</a>	<a href="#">311193</a>	18113	751	5	250	rho GTPase activating protein 1 (Arhgap1) alternative variant cSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.dSep08</a>	<a href="#">311193</a>	18751	802	9	225	rho GTPase activating protein 1 (Arhgap1) alternative variant dSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.eSep08</a>	<a href="#">311193</a>	18673	761	9	215	rho GTPase activating protein 1 (Arhgap1) alternative variant eSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.fSep08</a>	<a href="#">311193</a>	18371	759	9	204	rho GTPase activating protein 1 (Arhgap1) alternative variant fSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.gSep08</a>	<a href="#">311193</a>	18783	771	8	187	rho GTPase activating protein 1 (21.3 kD) (Arhgap1) alternative variant gSep08, mRNA.

<a href="#">Arhgap1</a>	<a href="#">Arhgap1.hSep08</a>	<a href="#">311193</a>	17823	460	6	130	rho GTPase activating protein 1 (Arhgap1) alternative variant hSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.iSep08</a>	<a href="#">311193</a>	15575	647	6	115	rho GTPase activating protein 1 (12.9 kD) (Arhgap1) alternative variant iSep08, complete mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.jSep08</a>	<a href="#">311193</a>	12081	524	4	106	rho GTPase activating protein 1 (12.0 kD) (Arhgap1) alternative variant jSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.kSep08</a>	<a href="#">311193</a>	4242	293	4	42	putative protein (4.5 kD) (Arhgap1) alternative variant kSep08, mRNA.
<a href="#">Arhgap1</a>	<a href="#">Arhgap1.lSep08</a>	<a href="#">311193</a>	1319	982	3	53	rho GTPase activating protein 1 (5.8 kD) (Arhgap1) alternative variant lSep08, mRNA.
<a href="#">Arhgap4</a>	<a href="#">Arhgap4.bSep08</a>	<a href="#">246249</a>	2562	1096	6	170	rho GTPase activating protein 4 (19.3 kD) (Arhgap4) alternative variant bSep08, mRNA.
<a href="#">Arhgap4</a>	<a href="#">Arhgap4.cSep08</a>	<a href="#">246249</a>	1934	653	3	151	rho GTPase activating protein 4 (Arhgap4) alternative variant cSep08, mRNA.
<a href="#">Arhgap4</a>	<a href="#">Arhgap4.dSep08</a>	<a href="#">246249</a>	363	288	2	95	rho GTPase activating protein 4 (Arhgap4) alternative variant dSep08, mRNA.
<a href="#">Arhgap5</a>	<a href="#">Arhgap5.aSep08</a>	<a href="#">299012</a>	48097	5049	5	1290	rho GTPase activating protein 5 (Arhgap5) alternative variant aSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.cSep08</a>	<a href="#">362893</a>	2225	729	8	242	rho GTPase activating protein 9 (Arhgap9) alternative variant cSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.dSep08</a>	<a href="#">362893</a>	1574	784	6	182	rho GTPase activating protein 9 (Arhgap9) alternative variant dSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.eSep08</a>	<a href="#">362893</a>	2205	742	5	157	rho GTPase activating protein 9 (Arhgap9) alternative variant eSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.fSep08</a>	<a href="#">362893</a>	3008	785	8	154	rho GTPase activating protein 9 (Arhgap9) alternative variant fSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.hSep08</a>	<a href="#">362893</a>	2559	694	6	90	rho GTPase activating protein 9 (Arhgap9) alternative variant hSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.iSep08</a>	<a href="#">362893</a>	2433	693	5	86	rho GTPase activating protein 9 (9.2 kD) (Arhgap9) alternative variant iSep08, mRNA.
<a href="#">Arhgap9</a>	<a href="#">Arhgap9.jSep08</a>	<a href="#">362893</a>	1220	644	2	104	rho GTPase activating protein 9 (11.1 kD) (Arhgap9) alternative variant jSep08, mRNA.
<a href="#">Arhgap10</a>	<a href="#">Arhgap10.bSep08</a>	<a href="#">688429</a>	62059	707	9	235	rho GTPase activating protein 10 (Arhgap10) alternative variant bSep08, mRNA.
<a href="#">Arhgap10</a>	<a href="#">Arhgap10.cSep08</a>	<a href="#">688429</a>	70037	400	4	86	rho GTPase activating protein 10 (Arhgap10) alternative variant cSep08, mRNA.
<a href="#">Arhgap11a</a>	<a href="#">Arhgap11a.aSep08</a>	<a href="#">296060</a>	8481	1019		325	rho GTPase activating protein 11A (Arhgap11a) mRNA.
<a href="#">Arhgap12</a>	<a href="#">Arhgap12.bSep08</a>	<a href="#">307016</a>	4422	729	4	158	rho GTPase activating protein 12 (Arhgap12) alternative variant bSep08, mRNA.
<a href="#">Arhgap17</a>	<a href="#">Arhgap17.bSep08</a>	<a href="#">63994</a>	21904	4305	8	440	rho GTPase activating protein 17 (Arhgap17) alternative variant bSep08, mRNA.
<a href="#">Arhgap18</a>	<a href="#">Arhgap18.aSep08</a>	<a href="#">293947</a>	148407	2300	14	692	rho GTPase activating protein 18 (Arhgap18) alternative variant aSep08, mRNA.
<a href="#">Arhgap18</a>	<a href="#">Arhgap18.bSep08</a>	<a href="#">293947</a>	49603	788	6	235	rho GTPase activating protein 18 (Arhgap18) alternative variant bSep08, mRNA.
<a href="#">Arhgap21</a>	<a href="#">Arhgap21.aSep08</a>	<a href="#">307178</a>	15054	1851	11	616	rho GTPase activating protein 21 (Arhgap21) alternative variant aSep08, mRNA.

<a href="#">Arhgap21</a>	<a href="#">Arhgap21.bSep08</a>	<a href="#">307178</a>	12665	3040	6	320	rho GTPase activating protein 21 (Arhgap21) alternative variant bSep08, mRNA.
<a href="#">Arhgap21</a>	<a href="#">Arhgap21.cSep08</a>	<a href="#">307178</a>	4781	434	1	118	rho GTPase activating protein 21 (Arhgap21) alternative variant cSep08, mRNA.
<a href="#">Arhgap22</a>	<a href="#">Arhgap22.bSep08</a>	<a href="#">306279</a>	15506	1471	3	490	rho GTPase activating protein 22 (Arhgap22) alternative variant bSep08, mRNA.
<a href="#">Arhgap23</a>	<a href="#">Arhgap23.aSep08</a>	<a href="#">303501</a>	39983	2258	10	364	rho GTPase activating protein 23 (Arhgap23) alternative variant aSep08, mRNA.
<a href="#">Arhgap23</a>	<a href="#">Arhgap23.bSep08</a>	<a href="#">303501</a>	11282	379		126	rho GTPase activating protein 23 (Arhgap23) alternative variant bSep08, mRNA.
<a href="#">Arhgap24</a>	<a href="#">Arhgap24.bSep08</a>	<a href="#">305156</a>	42079	699	6	214	rho GTPase activating protein 24 (Arhgap24) alternative variant bSep08, mRNA.
<a href="#">Arhgap24</a>	<a href="#">Arhgap24.cSep08</a>	<a href="#">305156</a>	152149	801	6	185	rho GTPase activating protein 24 (Arhgap24) alternative variant cSep08, mRNA.
<a href="#">Arhgap24</a>	<a href="#">Arhgap24.dSep08</a>	<a href="#">305156</a>	355122	597	5	158	rho GTPase activating protein 24 (Arhgap24) alternative variant dSep08, mRNA.
<a href="#">Arhgap24</a>	<a href="#">Arhgap24.eSep08</a>	<a href="#">305156</a>	103165	580	4	115	rho GTPase activating protein 24 (Arhgap24) alternative variant eSep08, mRNA.
<a href="#">Arhgap24</a>	<a href="#">Arhgap24.gSep08</a>	<a href="#">305156</a>	133564	389	4	66	rho GTPase activating protein 24 (Arhgap24) alternative variant gSep08, mRNA.
<a href="#">Arhgap27</a>	<a href="#">Arhgap27.bSep08</a>	<a href="#">303583</a>	4009	1361	4	183	rho GTPase activating protein 27 (Arhgap27) alternative variant bSep08, mRNA.
<a href="#">Arhgap27</a>	<a href="#">Arhgap27.cSep08</a>	<a href="#">303583</a>	2506	400	3	91	rho GTPase activating protein 27 (Arhgap27) alternative variant cSep08, mRNA.
<a href="#">Arhgap28</a>	<a href="#">Arhgap28.aSep08</a>	<a href="#">301709</a>	28869	2846		336	rho GTPase activating protein 28 (Arhgap28) mRNA.
<a href="#">Arhgap29</a>	<a href="#">Arhgap29.bSep08</a>	<a href="#">310833</a>	36164	754	6	186	rho GTPase activating protein 29 (Arhgap29) alternative variant bSep08, mRNA.
<a href="#">Arhgap29</a>	<a href="#">Arhgap29.cSep08</a>	<a href="#">310833</a>	6758	354	2	68	rho GTPase activating protein 29 (Arhgap29) alternative variant cSep08, mRNA.
<a href="#">Arhgdia</a>	<a href="#">Arhgdia.aSep08</a>	<a href="#">360678</a>	1915	721	6	240	rho GDP dissociation inhibitor (GDI) alpha (Arhgdia) alternative variant aSep08, mRNA.
<a href="#">Arhgdia</a>	<a href="#">Arhgdia.cSep08</a>	<a href="#">360678</a>	1609	566	5	127	rho GDP dissociation inhibitor (GDI) alpha (Arhgdia) alternative variant cSep08, mRNA.
<a href="#">Arhgdia</a>	<a href="#">Arhgdia.dSep08</a>	<a href="#">360678</a>	514	428	2	92	rho GDP dissociation inhibitor (GDI) alpha (Arhgdia) alternative variant dSep08, mRNA.
<a href="#">Arhgdib</a>	<a href="#">Arhgdib.aSep08</a>	<a href="#">362456</a>	18019	747	4	200	rho, GDP dissociation inhibitor (GDI) beta (22.9 kD) (Arhgdib) alternative variant aSep08, mRNA.
<a href="#">Arhgdib</a>	<a href="#">Arhgdib.bSep08</a>	<a href="#">362456</a>	18449	743	4	200	rho, GDP dissociation inhibitor (GDI) beta (22.9 kD) (Arhgdib) alternative variant bSep08, mRNA.
<a href="#">Arhgdib</a>	<a href="#">Arhgdib.cSep08</a>	<a href="#">362456</a>	17590	753	4	200	rho, GDP dissociation inhibitor (GDI) beta (22.9 kD) (Arhgdib) alternative variant cSep08, mRNA.
<a href="#">Arhgdib</a>	<a href="#">Arhgdib.dSep08</a>	<a href="#">362456</a>	18092	780	4	200	rho, GDP dissociation inhibitor (GDI) beta (22.9 kD) (Arhgdib) alternative variant dSep08, mRNA.
<a href="#">Arhgdib</a>	<a href="#">Arhgdib.fSep08</a>	<a href="#">362456</a>	12772	907	2	161	rho, GDP dissociation inhibitor (GDI) beta (18.0 kD) (Arhgdib) alternative variant fSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.aSep08</a>	<a href="#">60323</a>	16341	2993	26	884	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant aSep08, mRNA.

<a href="#">Arhgef1</a>	<a href="#">Arhgef1.bSep08</a>	<a href="#">60323</a>	9701	796	9	265	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant bSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.cSep08</a>	<a href="#">60323</a>	4590	744	5	219	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant cSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.dSep08</a>	<a href="#">60323</a>	2368	655	7	218	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant dSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.eSep08</a>	<a href="#">60323</a>	987	915	2	155	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant eSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.fSep08</a>	<a href="#">60323</a>	3009	1664	6	142	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant fSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.gSep08</a>	<a href="#">60323</a>	4412	1041	3	126	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant gSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.hSep08</a>	<a href="#">60323</a>	1289	898	3	98	rho guanine nucleotide exchange factor 1 (10.5 kD) (Arhgef1) alternative variant hSep08, mRNA.
<a href="#">Arhgef1</a>	<a href="#">Arhgef1.iSep08</a>	<a href="#">60323</a>	902	416	4	83	rho guanine nucleotide exchange factor 1 (Arhgef1) alternative variant iSep08, mRNA.
<a href="#">Arhgef2</a>	<a href="#">Arhgef2.aSep08</a>	<a href="#">310635</a>	27394	4238	20	1026	rho/rac guanine nucleotide exchange factor (GEF) 2 (Arhgef2) alternative variant aSep08, mRNA.
<a href="#">Arhgef2</a>	<a href="#">Arhgef2.bSep08</a>	<a href="#">310635</a>	17380	775	5	258	rho/rac guanine nucleotide exchange factor (GEF) 2 (Arhgef2) alternative variant bSep08, mRNA.
<a href="#">Arhgef2</a>	<a href="#">Arhgef2.cSep08</a>	<a href="#">310635</a>	17198	433	3	143	rho/rac guanine nucleotide exchange factor (GEF) 2 (Arhgef2) alternative variant cSep08, mRNA.
<a href="#">Arhgef3</a>	<a href="#">Arhgef3.bSep08</a>	<a href="#">290541</a>	275996	831	7	168	rho guanine nucleotide exchange factor (GEF) 3 (Arhgef3) alternative variant bSep08, mRNA.
<a href="#">Arhgef4</a>	<a href="#">Arhgef4.aSep08</a>	<a href="#">301334</a>	6502	2124	8	406	rho guanine nucleotide exchange factor (GEF) 4 (Arhgef4) alternative variant aSep08, mRNA.
<a href="#">Arhgef5</a>	<a href="#">Arhgef5.aSep08</a>	<a href="#">140898</a>	6334	867		288	rho guanine nucleotide exchange factor (GEF) 5 (Arhgef5) mRNA.
<a href="#">Arhgef6</a>	<a href="#">Arhgef6.bSep08</a>	<a href="#">363509</a>	45161	1299	13	406	rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 (Arhgef6) alternative variant bSep08, mRNA.
<a href="#">Arhgef6</a>	<a href="#">Arhgef6.cSep08</a>	<a href="#">363509</a>	48893	583	5	194	rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 (Arhgef6) alternative variant cSep08, mRNA.
<a href="#">Arhgef6</a>	<a href="#">Arhgef6.dSep08</a>	<a href="#">363509</a>	7658	386	4	128	rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 (Arhgef6) alternative variant dSep08, mRNA.
<a href="#">Arhgef6</a>	<a href="#">Arhgef6.eSep08</a>	<a href="#">363509</a>	27163	384	4	87	rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 (Arhgef6) alternative variant eSep08, mRNA.
<a href="#">Arhgef7</a>	<a href="#">Arhgef7.fSep08</a>	<a href="#">114559</a>	2637	396	2	39	rho guanine nucleotide exchange factor (GEF7) (Arhgef7) alternative variant fSep08, mRNA.
<a href="#">Arhgef9</a>	<a href="#">Arhgef9.bSep08</a>	<a href="#">66013</a>	13416	606	4	110	cdc42 guanine nucleotide exchange factor (GEF) 9 (Arhgef9) alternative variant bSep08, mRNA.
<a href="#">Arhgef9</a>	<a href="#">Arhgef9.cSep08</a>	<a href="#">66013</a>	3188	369	2	79	cdc42 guanine nucleotide exchange factor (GEF) 9 (Arhgef9) alternative variant cSep08, mRNA.
<a href="#">Arhgef9</a>	<a href="#">Arhgef9.dSep08</a>	<a href="#">66013</a>	33871	774	2	47	cdc42 guanine nucleotide exchange factor (GEF) 9 (Arhgef9) alternative variant dSep08, mRNA.
<a href="#">Arhgef9</a>	<a href="#">Arhgef9.fSep08</a>	<a href="#">66013</a>	33491	383	2	40	cdc42 guanine nucleotide exchange factor (GEF) 9 (Arhgef9) alternative variant fSep08, mRNA.

<a href="#">Arhgef11</a>	<a href="#">Arhgef11.bSep08</a>	<a href="#">78966</a>	6093	1213	7	389	rho guanine nucleotide exchange factor 11 (Arhgef11) alternative variant bSep08, mRNA.
<a href="#">Arhgef11</a>	<a href="#">Arhgef11.cSep08</a>	<a href="#">78966</a>	2969	696	4	215	rho guanine nucleotide exchange factor 11 (Arhgef11) alternative variant cSep08, mRNA.
<a href="#">Arhgef11</a>	<a href="#">Arhgef11.dSep08</a>	<a href="#">78966</a>	2876	1507	3	151	rho guanine nucleotide exchange factor 11 (16.0 kD) (Arhgef11) alternative variant dSep08, mRNA.
<a href="#">Arhgef11</a>	<a href="#">Arhgef11.eSep08</a>	<a href="#">78966</a>	8319	372	4	123	rho guanine nucleotide exchange factor 11 (Arhgef11) alternative variant eSep08, mRNA.
<a href="#">Arhgef11</a>	<a href="#">Arhgef11.fSep08</a>	<a href="#">78966</a>	619	369	2	116	rho guanine nucleotide exchange factor 11 (Arhgef11) alternative variant fSep08, mRNA.
<a href="#">Arhgef12</a>	<a href="#">Arhgef12.aSep08</a>	<a href="#">367072</a>	14694	3186	11	465	rho guanine nucleotide exchange factor 12 (Arhgef12) alternative variant aSep08, mRNA.
<a href="#">Arhgef12</a>	<a href="#">Arhgef12.bSep08</a>	<a href="#">367072</a>	3772	1963	3	129	putative mitochondrial protein (14.4 kD) (Arhgef12) alternative variant bSep08, mRNA.
<a href="#">Arhgef12</a>	<a href="#">Arhgef12.cSep08</a>	<a href="#">367072</a>	1018	529	2	79	rho guanine nucleotide exchange factor 12 (9.0 kD) (Arhgef12) alternative variant cSep08, mRNA.
<a href="#">Arhgef12</a>	<a href="#">Arhgef12.dSep08</a>	<a href="#">367072</a>	3067	1192	2	53	rho guanine nucleotide exchange factor 12 (Arhgef12) alternative variant dSep08, mRNA.
<a href="#">Arhgef17</a>	<a href="#">Arhgef17.bSep08</a>	<a href="#">308862</a>	9792	1782	6	387	rho guanine nucleotide exchange factor (GEF) 17 (Arhgef17) alternative variant bSep08, mRNA.
<a href="#">Arhgef17</a>	<a href="#">Arhgef17.cSep08</a>	<a href="#">308862</a>	2436	774	6	258	rho guanine nucleotide exchange factor (GEF) 17 (Arhgef17) alternative variant cSep08, mRNA.
<a href="#">Arhgef18</a>	<a href="#">Arhgef18.bSep08</a>	<a href="#">304193</a>	22962	4344	16	846	rho/rac guanine nucleotide exchange factor (GEF) 18 (Arhgef18) alternative variant bSep08, mRNA.
<a href="#">Arhgef18</a>	<a href="#">Arhgef18.dSep08</a>	<a href="#">304193</a>	2001	305	3	101	rho/rac guanine nucleotide exchange factor (GEF) 18 (Arhgef18) alternative variant dSep08, mRNA.
<a href="#">Arhgef18</a>	<a href="#">Arhgef18.eSep08</a>	<a href="#">304193</a>	4413	339	3	16	rho/rac guanine nucleotide exchange factor (GEF) 18 (1.8 kD) (Arhgef18) alternative variant eSep08, mRNA.
<a href="#">Arhgef19</a>	<a href="#">Arhgef19.bSep08</a>	<a href="#">362648</a>	6739	384	3	103	putative protein (Arhgef19) alternative variant bSep08, mRNA.
<a href="#">Arhgef19</a>	<a href="#">Arhgef19.cSep08</a>	<a href="#">362648</a>	1455	686	2	96	putative nuclear protein (11.1 kD) (Arhgef19) alternative variant cSep08, mRNA.
<a href="#">Arhgef19</a>	<a href="#">Arhgef19.dSep08</a>	<a href="#">362648</a>	3547	421	2	90	putative protein (9.7 kD) (Arhgef19) alternative variant dSep08, mRNA.
<a href="#">Arhgef19</a>	<a href="#">Arhgef19.eSep08</a>	<a href="#">362648</a>	468	388	2	36	putative protein (Arhgef19) alternative variant eSep08, mRNA.
<a href="#">Arid1a</a>	<a href="#">Arid1a.bSep08</a>	<a href="#">297867</a>	2035	743	4	226	AT rich interactive domain 1A (Swi1 like) (Arid1a) alternative variant bSep08, mRNA.
<a href="#">Arid1a</a>	<a href="#">Arid1a.cSep08</a>	<a href="#">297867</a>	1598	627	5	208	AT rich interactive domain 1A (Swi1 like) (Arid1a) alternative variant cSep08, mRNA.
<a href="#">Arid1b</a>	<a href="#">Arid1b.aSep08</a>	<a href="#">282546</a>	91554	518		164	AT rich interactive domain 1B (Swi1 like) (Arid1b) mRNA.
<a href="#">Arid2</a>	<a href="#">Arid2.aSep08</a>	<a href="#">366980</a>	32105	1738	7	508	AT rich interactive domain 2 (Arid-rfx like) (Arid2) alternative variant aSep08, mRNA.

<a href="#">Arid3a</a>	<a href="#">Arid3a.aSep08</a>	<a href="#">314616</a>	21494	2347	8	655	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant aSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.aSep08</a>	<a href="#">690999</a>	21494	2347	8	655	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant aSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.bSep08</a>	<a href="#">314616</a>	25427	1793	3	346	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant bSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.bSep08</a>	<a href="#">690999</a>	25427	1793	3	346	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant bSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.dSep08</a>	<a href="#">314616</a>	4093	353	2	71	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant dSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.dSep08</a>	<a href="#">690999</a>	4093	353	2	71	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant dSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.eSep08</a>	<a href="#">314616</a>	802	176	2	35	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant eSep08, mRNA.
<a href="#">Arid3a</a>	<a href="#">Arid3a.eSep08</a>	<a href="#">690999</a>	802	176	2	35	AT rich interactive domain 3A (Bright like) and similar to AT-rich interactive domain-containing protein 3A (ARID domain-containing protein 3A) (Dead ringer-like protein 1) (B-cell regulator of IgH transcription) (Bright) (Arid3a) alternative variant eSep08, mRNA.
<a href="#">Arid4a</a>	<a href="#">Arid4a.bSep08</a>	<a href="#">314205</a>	40877	4709	14	1021	AT rich interactive domain 4A (Rbp1 like) (Arid4a) alternative variant bSep08, mRNA.
<a href="#">Arid4a</a>	<a href="#">Arid4a.cSep08</a>	<a href="#">314205</a>	21823	539	7	179	AT rich interactive domain 4A (Rbp1 like) (Arid4a) alternative variant cSep08, mRNA.
<a href="#">Arid4a</a>	<a href="#">Arid4a.dSep08</a>	<a href="#">314205</a>	14018	545	5	96	AT rich interactive domain 4A (Rbp1 like) (Arid4a) alternative variant dSep08, mRNA.

<a href="#">Arid4b</a>	<a href="#">Arid4b.bSep08</a>	<a href="#">84481</a>	31273	1348	5	449	AT rich interactive domain 4B (Rbp1 like) (Arid4b) alternative variant bSep08, mRNA.
<a href="#">Arid4b</a>	<a href="#">Arid4b.cSep08</a>	<a href="#">84481</a>	2170	667	2	125	AT rich interactive domain 4B (Rbp1 like) (14.7 kD) (Arid4b) alternative variant cSep08, mRNA.
<a href="#">Arid5a</a>	<a href="#">Arid5a.bSep08</a>	<a href="#">316327</a>	10458	588	6	196	AT rich interactive domain 5A (Mrf1 like) (Arid5a) alternative variant bSep08, mRNA.
<a href="#">Arid5a</a>	<a href="#">Arid5a.cSep08</a>	<a href="#">316327</a>	1234	396	2	121	AT rich interactive domain 5A (Mrf1 like) (Arid5a) alternative variant cSep08, mRNA.
<a href="#">Arid5b</a>	<a href="#">Arid5b.bSep08</a>	<a href="#">309728</a>	32172	1142	5	380	AT rich interactive domain 5B (Mrf1 like) (Arid5b) alternative variant bSep08, mRNA.
<a href="#">Arid5b</a>	<a href="#">Arid5b.cSep08</a>	<a href="#">309728</a>	1835	390	2	79	AT rich interactive domain 5B (Mrf1 like) (Arid5b) alternative variant cSep08, mRNA.
<a href="#">Arih2</a>	<a href="#">Arih2.aSep08</a>	<a href="#">316005</a>	58922	3754	16	492	ariadne homolog 2 (Drosophila) (57.7 kD) (Arih2) alternative variant aSep08, complete mRNA.
<a href="#">Arih2</a>	<a href="#">Arih2.bSep08</a>	<a href="#">316005</a>	7137	708	7	234	ariadne homolog 2 (Drosophila) (Arih2) alternative variant bSep08, mRNA.
<a href="#">Arih2</a>	<a href="#">Arih2.cSep08</a>	<a href="#">316005</a>	8784	822	7	228	ariadne homolog 2 (Drosophila) (Arih2) alternative variant cSep08, mRNA.
<a href="#">Arih2</a>	<a href="#">Arih2.dSep08</a>	<a href="#">316005</a>	4027	729	5	120	ariadne homolog 2 (Drosophila) (Arih2) alternative variant dSep08, mRNA.
<a href="#">Arih2</a>	<a href="#">Arih2.fSep08</a>	<a href="#">316005</a>	7257	663	3	98	ariadne homolog 2 (Drosophila) (10.8 kD) (Arih2) alternative variant fSep08, complete mRNA.
<a href="#">Arl2</a>	<a href="#">Arl2.bSep08</a>	<a href="#">65142</a>	3008	675	3	127	ADP-ribosylation factor-like 2 (Arl2) alternative variant bSep08, mRNA.
<a href="#">Arl2</a>	<a href="#">Arl2.cSep08</a>	<a href="#">65142</a>	1051	962	2	125	ADP-ribosylation factor-like 2 (13.9 kD) (Arl2) alternative variant cSep08, mRNA.
<a href="#">Arl2</a>	<a href="#">Arl2.dSep08</a>	<a href="#">65142</a>	1755	714	2	95	ADP-ribosylation factor-like 2 (10.5 kD) (Arl2) alternative variant dSep08, mRNA.
<a href="#">Arl2bp</a>	<a href="#">Arl2bp.aSep08</a>	<a href="#">498910</a>	8619	1029	6	163	ADP-ribosylation factor-like 2 binding protein (18.7 kD) (Arl2bp) alternative variant aSep08, mRNA.
<a href="#">Arl2bp</a>	<a href="#">Arl2bp.bSep08</a>	<a href="#">498910</a>	7489	744	5	138	ADP-ribosylation factor-like 2 binding protein (16.0 kD) (Arl2bp) alternative variant bSep08, mRNA.
<a href="#">Arl2bp</a>	<a href="#">Arl2bp.dSep08</a>	<a href="#">498910</a>	8064	621	7	119	ADP-ribosylation factor-like 2 binding protein (13.8 kD) (Arl2bp) alternative variant dSep08, mRNA.
<a href="#">Arl3</a>	<a href="#">Arl3.aSep08</a>	<a href="#">64664</a>	115072	1209	2	186	ADP-ribosylation factor-like 3 (20.9 kD) (Arl3) alternative variant aSep08, mRNA.
<a href="#">Arl4a</a>	<a href="#">Arl4a.aSep08</a>	<a href="#">29308</a>	1546	1117	1	200	ADP-ribosylation factor-like 4A (22.6 kD) (Arl4a) alternative variant aSep08, mRNA.
<a href="#">Arl4c</a>	<a href="#">Arl4c.aSep08</a>	<a href="#">367311</a>	3588	770	2	105	ADP-ribosylation factor-like 4C (11.6 kD) (Arl4c) alternative variant aSep08, mRNA.
<a href="#">Arl5b</a>	<a href="#">Arl5b.aSep08</a>	<a href="#">364788</a>	24578	3366	2	245	ADP-ribosylation factor-like 5B (Arl5b) alternative variant aSep08, mRNA.
<a href="#">Arl5c</a>	<a href="#">Arl5c.bSep08</a>	<a href="#">497990</a>	10065	1674	6	116	ADP-ribosylation factor-like 5C (13.0 kD) (Arl5c) alternative variant bSep08, mRNA.
<a href="#">Arl5c</a>	<a href="#">Arl5c.cSep08</a>	<a href="#">497990</a>	2043	670	2	91	ADP-ribosylation factor-like 5C (10.2 kD) (Arl5c) alternative variant cSep08, mRNA.



<a href="#">Arl6</a>	<a href="#">Arl6.bSep08</a>	<a href="#">363760</a>	26333	1856	8	186	ADP-ribosylation factor-like 6 (21.0 kD) (Arl6) alternative variant bSep08, mRNA.
<a href="#">Arl6</a>	<a href="#">Arl6.cSep08</a>	<a href="#">363760</a>	12314	680	5	155	ADP-ribosylation factor-like 6 (Arl6) alternative variant cSep08, mRNA.
<a href="#">Arl6ip1</a>	<a href="#">Arl6ip1.bSep08</a>	<a href="#">293551</a>	3438	494	3	97	ADP-ribosylation factor-like 6 interacting protein 1 (10.9 kD) (Arl6ip1) alternative variant bSep08, mRNA.
<a href="#">Arl6ip2</a>	<a href="#">Arl6ip2.aSep08</a>	<a href="#">298757</a>	16752	2996	12	461	ADP-ribosylation factor-like 6 interacting protein 2 (Arl6ip2) alternative variant aSep08, mRNA.
<a href="#">Arl6ip2</a>	<a href="#">Arl6ip2.bSep08</a>	<a href="#">298757</a>	5243	1216	4	225	ADP-ribosylation factor-like 6 interacting protein 2 (Arl6ip2) alternative variant bSep08, mRNA.
<a href="#">Arl6ip2</a>	<a href="#">Arl6ip2.cSep08</a>	<a href="#">298757</a>	26089	611	4	203	ADP-ribosylation factor-like 6 interacting protein 2 (Arl6ip2) alternative variant cSep08, mRNA.
<a href="#">Arl6ip2</a>	<a href="#">Arl6ip2.dSep08</a>	<a href="#">298757</a>	1808	1711	2	44	ADP-ribosylation factor-like 6 interacting protein 2 (4.8 kD) (Arl6ip2) alternative variant dSep08, mRNA.
<a href="#">Arl6ip5</a>	<a href="#">Arl6ip5.bSep08</a>	<a href="#">66028</a>	19630	846	1	132	ADP-ribosylation factor-like 6 interacting protein 5 (Arl6ip5) alternative variant bSep08, mRNA.
<a href="#">Arl6ip6</a>	<a href="#">Arl6ip6.bSep08</a>	<a href="#">499798</a>	27508	1038	4	205	ADP-ribosylation factor-like 6 interacting protein 6 (22.6 kD) (Arl6ip6) alternative variant bSep08, complete mRNA.
<a href="#">Arl6ip6</a>	<a href="#">Arl6ip6.cSep08</a>	<a href="#">499798</a>	25118	841	4	147	ADP-ribosylation factor-like 6 interacting protein 6 (Arl6ip6) alternative variant cSep08, mRNA.
<a href="#">Arl8b</a>	<a href="#">Arl8b.aSep08</a>	<a href="#">500282</a>	43065	745	6	200	ADP-ribosylation factor-like 8B (Arl8b) alternative variant aSep08, mRNA.
<a href="#">Arl8b</a>	<a href="#">Arl8b.cSep08</a>	<a href="#">500282</a>	43436	189	1	63	ADP-ribosylation factor-like 8B (Arl8b) alternative variant cSep08, mRNA.
<a href="#">Arl10</a>	<a href="#">Arl10.aSep08</a>	<a href="#">306767</a>	6341	595	1	198	ADP-ribosylation factor-like 10 (Arl10) alternative variant aSep08, mRNA.
<a href="#">Arl13b</a>	<a href="#">Arl13b.bSep08</a>	<a href="#">304037</a>	8661	1643	3	153	ADP-ribosylation factor-like 1 (Arl13b) alternative variant bSep08, mRNA.
<a href="#">Arl13b</a>	<a href="#">Arl13b.cSep08</a>	<a href="#">304037</a>	6719	270	2	54	ADP-ribosylation factor-like 13 (Arl13b) alternative variant cSep08, mRNA.
<a href="#">Arl15</a>	<a href="#">Arl15.aSep08</a>	<a href="#">689079</a>	253623	3178		191	ADP-ribosylation factor-like 15 (Arl15) mRNA.
<a href="#">Arm.0</a>	<a href="#">Arm.0.aSep08</a>		32265	599		199	armadillo repeat containing 3 CRA b (Arm.0) mRNA.
<a href="#">Arm.1</a>	<a href="#">Arm.1.aSep08</a>		2141	1096		365	karyopherin alpha 2 (Arm.1) mRNA.
<a href="#">Armc2</a>	<a href="#">Armc2.aSep08</a>	<a href="#">499470</a>	73972	1705	8	568	armadillo repeat containing 2 (Armc2) alternative variant aSep08, mRNA.
<a href="#">Armc2</a>	<a href="#">Armc2.bSep08</a>	<a href="#">499470</a>	19150	1787	3	511	armadillo repeat containing 2 (Armc2) alternative variant bSep08, mRNA.
<a href="#">Armc2</a>	<a href="#">Armc2.cSep08</a>	<a href="#">499470</a>	11288	767	1	255	armadillo repeat containing 2 (Armc2) alternative variant cSep08, mRNA.
<a href="#">Armc2</a>	<a href="#">Armc2.dSep08</a>	<a href="#">499470</a>	14072	783	1	191	armadillo repeat containing 2 (Armc2) alternative variant dSep08, mRNA.
<a href="#">Armc4</a>	<a href="#">Armc4.aSep08</a>	<a href="#">307036</a>	42624	533		125	armadillo repeat containing 4 (Armc4) mRNA.
<a href="#">Armc5</a>	<a href="#">Armc5.bSep08</a>	<a href="#">361653</a>	1523	1431	2	306	armadillo repeat containing 5 (32.0 kD) (Armc5) alternative variant bSep08, mRNA.
<a href="#">Armc5</a>	<a href="#">Armc5.cSep08</a>	<a href="#">361653</a>	4989	2061	6	160	armadillo repeat containing 5 CRA c (16.6 kD) (Armc5) alternative variant cSep08, mRNA.

<a href="#">Armc6</a>	<a href="#">Armc6.aSep08</a>	<a href="#">306352</a>	15269	1885	7	468	armadillo repeat containing 6 (50.6 kD) (Armc6) alternative variant aSep08, complete mRNA.
<a href="#">Armc6</a>	<a href="#">Armc6.cSep08</a>	<a href="#">306352</a>	5462	335	2	111	armadillo repeat containing 6 (Armc6) alternative variant cSep08, mRNA.
<a href="#">Armc8</a>	<a href="#">Armc8.aSep08</a>	<a href="#">315949</a>	94311	4390	22	684	armadillo repeat containing 8 (Armc8) alternative variant aSep08, mRNA.
<a href="#">Armc8</a>	<a href="#">Armc8.bSep08</a>	<a href="#">315949</a>	44909	675	7	178	armadillo repeat containing 8 (Armc8) alternative variant bSep08, mRNA.
<a href="#">Armc8</a>	<a href="#">Armc8.cSep08</a>	<a href="#">315949</a>	3293	667	2	67	armadillo repeat containing 8 (Armc8) alternative variant cSep08, mRNA.
<a href="#">Armc9</a>	<a href="#">Armc9.bSep08</a>	<a href="#">301579</a>	70132	1594	13	436	armadillo repeat containing 9 (Armc9) alternative variant bSep08, mRNA.
<a href="#">Armc9</a>	<a href="#">Armc9.cSep08</a>	<a href="#">301579</a>	40296	407	4	104	armadillo repeat containing 9 (Armc9) alternative variant cSep08, mRNA.
<a href="#">Armc9</a>	<a href="#">Armc9.dSep08</a>	<a href="#">301579</a>	6639	310	3	86	armadillo repeat containing 9 (Armc9) alternative variant dSep08, mRNA.
<a href="#">Armc9</a>	<a href="#">Armc9.fSep08</a>	<a href="#">301579</a>	1155	600	2	63	armadillo repeat containing 9 (Armc9) alternative variant fSep08, mRNA.
<a href="#">Armcx1</a>	<a href="#">Armcx1.bSep08</a>	<a href="#">501619</a>	1459	405	2	98	armadillo repeat containing, X-linked 1 (Armcx1) alternative variant bSep08, mRNA.
<a href="#">Armcx1</a>	<a href="#">Armcx1.cSep08</a>	<a href="#">501619</a>	2342	489	1	73	armadillo repeat containing, X-linked 1 (Armcx1) alternative variant cSep08, mRNA.
<a href="#">Armcx3</a>	<a href="#">Armcx3.aSep08</a>	<a href="#">367902</a>	4915	3353	2	105	armadillo repeat containing, X-linked 3 (11.9 kD) (Armcx3) alternative variant aSep08, mRNA.
<a href="#">Armcx6</a>	<a href="#">Armcx6.bSep08</a>	<a href="#">363496</a>	1769	800	4	156	armadillo repeat containing, X-linked 6 (Armcx6) alternative variant bSep08, mRNA.
<a href="#">Armcx6</a>	<a href="#">Armcx6.cSep08</a>	<a href="#">363496</a>	1433	765	3	140	armadillo repeat containing, X-linked 6 (Armcx6) alternative variant cSep08, mRNA.
<a href="#">Armcx6</a>	<a href="#">Armcx6.dSep08</a>	<a href="#">363496</a>	1668	723	4	116	armadillo repeat containing, X-linked 6 (Armcx6) alternative variant dSep08, mRNA.
<a href="#">Armet</a>	<a href="#">Armet.aSep08</a>	<a href="#">315989</a>	3235	876	2	200	arginine-rich, mutated in early stage tumors (Armet) alternative variant aSep08, mRNA.
<a href="#">Arnt</a>	<a href="#">Arnt.bSep08</a>	<a href="#">25242</a>	5084	1386	4	187	aryl hydrocarbon receptor nuclear translocator (Arnt) alternative variant bSep08, mRNA.
<a href="#">Arnt</a>	<a href="#">Arnt.cSep08</a>	<a href="#">25242</a>	33429	613	7	165	aryl hydrocarbon receptor nuclear translocator (Arnt) alternative variant cSep08, mRNA.
<a href="#">Arnt</a>	<a href="#">Arnt.dSep08</a>	<a href="#">25242</a>	723	289	3	96	aryl hydrocarbon receptor nuclear translocator (Arnt) alternative variant dSep08, mRNA.
<a href="#">Arntl</a>	<a href="#">Arntl.bSep08</a>	<a href="#">29657</a>	11258	893	7	297	aryl hydrocarbon receptor nuclear translocator-like (Arntl) alternative variant bSep08, mRNA.
<a href="#">Arntl</a>	<a href="#">Arntl.cSep08</a>	<a href="#">29657</a>	10984	738	6	204	aryl hydrocarbon receptor nuclear translocator-like (Arntl) alternative variant cSep08, mRNA.
<a href="#">Arntl</a>	<a href="#">Arntl.dSep08</a>	<a href="#">29657</a>	71941	617	6	122	aryl hydrocarbon receptor nuclear translocator-like (Arntl) alternative variant dSep08, mRNA.
<a href="#">Arpc1a</a>	<a href="#">Arpc1a.aSep08</a>	<a href="#">81824</a>	22483	1796	5	537	actin related protein 2/3 complex, subunit 1A (Arpc1a) alternative variant aSep08, complete mRNA.

<a href="#">Arpc1a</a>	<a href="#">Arpc1a.bSep08</a>	<a href="#">81824</a>	17696	937	7	257	actin related protein 2/3 complex, subunit 1A (29.4 kD) (Arpc1a) alternative variant bSep08, mRNA.
<a href="#">Arpc1b</a>	<a href="#">Arpc1b.bSep08</a>	<a href="#">54227</a>	2180	2048	2	93	actin related protein 2/3 complex, subunit 1B (9.9 kD) (Arpc1b) alternative variant bSep08, mRNA.
<a href="#">Arpc1b</a>	<a href="#">Arpc1b.dSep08</a>	<a href="#">54227</a>	682	475	3	60	actin related protein 2/3 complex, subunit 1B (Arpc1b) alternative variant dSep08, mRNA.
<a href="#">Arpc2</a>	<a href="#">Arpc2.bSep08</a>	<a href="#">301511</a>	14736	524	4	174	actin related protein 2/3 complex, subunit 2 (Arpc2) alternative variant bSep08, mRNA.
<a href="#">Arpc2</a>	<a href="#">Arpc2.cSep08</a>	<a href="#">301511</a>	13240	463	3	71	actin related protein 2/3 complex, subunit 2 (Arpc2) alternative variant cSep08, mRNA.
<a href="#">Arpc2</a>	<a href="#">Arpc2.dSep08</a>	<a href="#">301511</a>	7320	784	2	62	actin related protein 2/3 complex, subunit 2 (Arpc2) alternative variant dSep08, mRNA.
<a href="#">Arpc2</a>	<a href="#">Arpc2.fSep08</a>	<a href="#">301511</a>	3870	771	3	35	actin related protein 2/3 complex, subunit 2 (4.0 kD) (Arpc2) alternative variant fSep08, mRNA.
<a href="#">Arpc5</a>	<a href="#">Arpc5.bSep08</a>	<a href="#">360854</a>	6079	1045	2	116	actin related protein 2/3 complex, subunit 5 (Arpc5) alternative variant bSep08, mRNA.
<a href="#">Arpc5</a>	<a href="#">Arpc5.cSep08</a>	<a href="#">360854</a>	3366	689	1	90	actin related protein 2/3 complex, subunit 5 (9.9 kD) (Arpc5) alternative variant cSep08, mRNA.
<a href="#">Arpc5l</a>	<a href="#">Arpc5l.aSep08</a>	<a href="#">296710</a>	8094	1740	3	176	actin related protein 2/3 complex, subunit 5-like (Arpc5l) alternative variant aSep08, mRNA.
<a href="#">Arpp19</a>	<a href="#">Arpp19.aSep08</a>	<a href="#">60336</a>	25779	3968	4	137	cAMP-regulated phosphoprotein 19 (Arpp19) alternative variant aSep08, mRNA.
<a href="#">Arpp19</a>	<a href="#">Arpp19.bSep08</a>	<a href="#">60336</a>	22601	794	4	112	cAMP-regulated phosphoprotein 19 (12.3 kD) (Arpp19) alternative variant bSep08, complete mRNA.
<a href="#">Arpp19</a>	<a href="#">Arpp19.dSep08</a>	<a href="#">60336</a>	7054	462	2	96	cAMP-regulated phosphoprotein 19 (10.6 kD) (Arpp19) alternative variant dSep08, mRNA.
<a href="#">Arpp21</a>	<a href="#">Arpp21.aSep08</a>	<a href="#">363153</a>	60952	1551	6	344	cyclic AMP-regulated phosphoprotein, 21 (Arpp21) alternative variant aSep08, mRNA.
<a href="#">Arpp21</a>	<a href="#">Arpp21.bSep08</a>	<a href="#">363153</a>	1835	706	3	98	cyclic AMP-regulated phosphoprotein, 21 (Arpp21) alternative variant bSep08, mRNA.
<a href="#">Arr3</a>	<a href="#">Arr3.aSep08</a>	<a href="#">171107</a>	3881	389	6	95	arrestin 3, retinal (Arr3) alternative variant aSep08, mRNA.
<a href="#">Arrb1</a>	<a href="#">Arrb1.aSep08</a>	<a href="#">25387</a>	89195	1793	13	553	arrestin, beta 1 (Arrb1) alternative variant aSep08, mRNA.
<a href="#">Arrb1</a>	<a href="#">Arrb1.cSep08</a>	<a href="#">25387</a>	1899	415	2	50	arrestin, beta 1 (Arrb1) alternative variant cSep08, mRNA.
<a href="#">Arrb2</a>	<a href="#">Arrb2.bSep08</a>	<a href="#">25388</a>	947	761	2	77	arrestin, beta 2 (Arrb2) alternative variant bSep08, mRNA.
<a href="#">Arrb2</a>	<a href="#">Arrb2.cSep08</a>	<a href="#">25388</a>	1060	697	3	44	arrestin, beta 2 (Arrb2) alternative variant cSep08, mRNA.
<a href="#">Arrdc1</a>	<a href="#">Arrdc1.aSep08</a>	<a href="#">366001</a>	7182	1702	6	434	arrestin, N-terminal and arrestin, C-terminal (46.1 kD) (Arrdc1) alternative variant aSep08, mRNA.
<a href="#">Arrdc1</a>	<a href="#">Arrdc1.bSep08</a>	<a href="#">366001</a>	6989	1479	4	359	arrestin, N-terminal and arrestin, C-terminal (Arrdc1) alternative variant bSep08, mRNA.
<a href="#">Arrdc1</a>	<a href="#">Arrdc1.cSep08</a>	<a href="#">366001</a>	1007	760	1	253	arrestin, C-terminal (Arrdc1) alternative variant cSep08, mRNA.
<a href="#">Arrdc2</a>	<a href="#">Arrdc2.aSep08</a>	<a href="#">306344</a>	1939	927	4	268	arrestin, N-terminal and arrestin, C-terminal (Arrdc2) alternative variant aSep08, mRNA.
<a href="#">Arrdc2</a>	<a href="#">Arrdc2.bSep08</a>	<a href="#">306344</a>	1642	712	3	237	arrestin, N-terminal and arrestin, C-terminal (Arrdc2) alternative variant bSep08, mRNA.

<a href="#">Arrdc3</a>	<a href="#">Arrdc3.bSep08</a>	<a href="#">309945</a>	4701	3026	2	102	putative nuclear protein of vertebrate origin (11.5 kD) (Arrdc3) alternative variant bSep08, mRNA.
<a href="#">Arrdc3</a>	<a href="#">Arrdc3.cSep08</a>	<a href="#">309945</a>	3117	887	4	81	putative protein of vertebrate origin (Arrdc3) alternative variant cSep08, mRNA.
<a href="#">Arrdc4</a>	<a href="#">Arrdc4.bSep08</a>	<a href="#">293019</a>	4587	790	2	214	arrestin, C-terminal (Arrdc4) alternative variant bSep08, mRNA.
<a href="#">Arsa</a>	<a href="#">Arsa.bSep08</a>	<a href="#">315222</a>	3244	2168	7	360	arylsulfatase A (Arsa) alternative variant bSep08, mRNA.
<a href="#">Arsa</a>	<a href="#">Arsa.cSep08</a>	<a href="#">315222</a>	1194	696	4	104	arylsulfatase A (Arsa) alternative variant cSep08, mRNA.
<a href="#">Arzb</a>	<a href="#">Arzb.bSep08</a>	<a href="#">25227</a>	136335	486		151	arylsulfatase B (Arzb) alternative variant bSep08, mRNA.
<a href="#">Arse</a>	<a href="#">Arse.bSep08</a>	<a href="#">310326</a>	7690	3342	8	287	arylsulfatase E (chondrodysplasia punctata 1) (30.6 kD) (Arse) alternative variant bSep08, mRNA.
<a href="#">Arse</a>	<a href="#">Arse.cSep08</a>	<a href="#">310326</a>	1296	342	2	47	arylsulfatase E (chondrodysplasia punctata 1) (4.8 kD) (Arse) alternative variant cSep08, mRNA.
<a href="#">Arsg</a>	<a href="#">Arsg.bSep08</a>	<a href="#">303631</a>	8746	385	1	96	arylsulfatase G (Arsg) alternative variant bSep08, mRNA.
<a href="#">Arsk</a>	<a href="#">Arsk.bSep08</a>	<a href="#">365619</a>	2680	410	2	40	arylsulfatase K (Arsk) alternative variant bSep08, mRNA.
<a href="#">Art2b</a>	<a href="#">Art2b.bSep08</a>	<a href="#">293152</a>	9675	1129	3	266	ADP-ribosyltransferase 2b (Art2b) alternative variant bSep08, mRNA.
<a href="#">Art2b</a>	<a href="#">Art2b.cSep08</a>	<a href="#">293152</a>	10578	722	5	97	ADP-ribosyltransferase 2b (11.1 kD) (Art2b) alternative variant cSep08, mRNA.
<a href="#">Art3</a>	<a href="#">Art3.aSep08</a>	<a href="#">305235</a>	28070	1466	9	403	ADP-ribosyltransferase 3 (Art3) alternative variant aSep08, mRNA.
<a href="#">Art3</a>	<a href="#">Art3.cSep08</a>	<a href="#">305235</a>	22666	936	3	259	ADP-ribosyltransferase 3 (Art3) alternative variant cSep08, mRNA.
<a href="#">Art4</a>	<a href="#">Art4.aSep08</a>	<a href="#">312806</a>	9222	1323		311	ADP-ribosyltransferase 4 (Art4) mRNA.
<a href="#">Art5</a>	<a href="#">Art5.bSep08</a>	<a href="#">259167</a>	1233	440	1	76	ADP-ribosyltransferase 5 (Art5) alternative variant bSep08, mRNA.
<a href="#">Artn</a>	<a href="#">Artn.bSep08</a>	<a href="#">362572</a>	1249	369	1	91	artemin (Artn) alternative variant bSep08, mRNA.
<a href="#">Arv1</a>	<a href="#">Arv1.bSep08</a>	<a href="#">292097</a>	11759	1059	6	279	ARV1 homolog (yeast) (30.9 kD) (Arv1) alternative variant bSep08, complete mRNA.
<a href="#">Arv1</a>	<a href="#">Arv1.cSep08</a>	<a href="#">292097</a>	11737	971	6	223	ARV1 homolog (yeast) (Arv1) alternative variant cSep08, mRNA.
<a href="#">Arv1</a>	<a href="#">Arv1.dSep08</a>	<a href="#">292097</a>	9033	743	5	175	ARV1 homolog (yeast) (Arv1) alternative variant dSep08, mRNA.
<a href="#">Arvcf</a>	<a href="#">Arvcf.bSep08</a>	<a href="#">303798</a>	5131	794	3	264	armadillo repeat gene deleted in velo-cardio-facial syndrome (Arvcf) alternative variant bSep08, mRNA.
<a href="#">Arvcf</a>	<a href="#">Arvcf.cSep08</a>	<a href="#">303798</a>	2499	786	8	217	armadillo repeat gene deleted in velo-cardio-facial syndrome (Arvcf) alternative variant cSep08, mRNA.
<a href="#">Arvcf</a>	<a href="#">Arvcf.dSep08</a>	<a href="#">303798</a>	1187	618	4	110	armadillo repeat gene deleted in velo-cardio-facial syndrome (Arvcf) alternative variant dSep08, mRNA.
<a href="#">Arx</a>	<a href="#">Arx.bSep08</a>	<a href="#">317268</a>	7055	1751	3	441	aristaless related homeobox gene (Drosophila) (Arx) alternative variant bSep08, mRNA.
<a href="#">Arx</a>	<a href="#">Arx.cSep08</a>	<a href="#">317268</a>	6446	1558	3	201	aristaless related homeobox gene (Drosophila) (Arx) alternative variant cSep08, mRNA.
<a href="#">Arx</a>	<a href="#">Arx.dSep08</a>	<a href="#">317268</a>	2097	522	1	108	aristaless related homeobox gene (Drosophila) (Arx) alternative variant dSep08, mRNA.

<a href="#">As3mt</a>	<a href="#">As3mt.bSep08</a>	<a href="#">140925</a>	27141	715	6	231	methyltransferase (As3mt) alternative variant bSep08, mRNA.
<a href="#">As3mt</a>	<a href="#">As3mt.cSep08</a>	<a href="#">140925</a>	7282	755	7	231	methyltransferase (As3mt) alternative variant cSep08, mRNA.
<a href="#">As3mt</a>	<a href="#">As3mt.dSep08</a>	<a href="#">140925</a>	17677	733	8	205	methyltransferase (23.0 kD) (As3mt) alternative variant dSep08, complete mRNA.
<a href="#">As3mt</a>	<a href="#">As3mt.eSep08</a>	<a href="#">140925</a>	2019	547	4	117	methyltransferase (13.1 kD) (As3mt) alternative variant eSep08, mRNA.
<a href="#">As3mt</a>	<a href="#">As3mt.hSep08</a>	<a href="#">140925</a>	974	234	2	40	putative protein (As3mt) alternative variant hSep08, mRNA.
<a href="#">Asah1</a>	<a href="#">Asah1.bSep08</a>	<a href="#">84431</a>	25719	591	8	196	N-acylsphingosine amidohydrolase 1 (Asah1) alternative variant bSep08, mRNA.
<a href="#">Asah1</a>	<a href="#">Asah1.cSep08</a>	<a href="#">84431</a>	1733	711	2	43	N-acylsphingosine amidohydrolase 1 (Asah1) alternative variant cSep08, mRNA.
<a href="#">Asah2</a>	<a href="#">Asah2.bSep08</a>	<a href="#">114104</a>	37604	809	4	68	N-acylsphingosine amidohydrolase 2 (6.9 kD) (Asah2) alternative variant bSep08, mRNA.
<a href="#">Asah3</a>	<a href="#">Asah3.bSep08</a>	<a href="#">301118</a>	25496	519	3	115	N-acylsphingosine amidohydrolase (alkaline ceramidase) 3 (Asah3) alternative variant bSep08, mRNA.
<a href="#">Asah3l</a>	<a href="#">Asah3l.bSep08</a>	<a href="#">313339</a>	22542	380	3	126	N-acylsphingosine amidohydrolase 3-like (Asah3l) alternative variant bSep08, mRNA.
<a href="#">Asah3l</a>	<a href="#">Asah3l.cSep08</a>	<a href="#">313339</a>	2509	266	2	65	N-acylsphingosine amidohydrolase 3-like (Asah3l) alternative variant cSep08, mRNA.
<a href="#">Asb1</a>	<a href="#">Asb1.bSep08</a>	<a href="#">316628</a>	15015	1473	2	332	ankyrin and SOCS protein, C-terminal (37.1 kD) (Asb1) alternative variant bSep08, mRNA.
<a href="#">Asb2</a>	<a href="#">Asb2.bSep08</a>	<a href="#">299266</a>	5185	734	4	244	ankyrin (Asb2) alternative variant bSep08, mRNA.
<a href="#">Asb2</a>	<a href="#">Asb2.cSep08</a>	<a href="#">299266</a>	10454	709	2	206	ankyrin (Asb2) alternative variant cSep08, mRNA.
<a href="#">Asb2</a>	<a href="#">Asb2.fSep08</a>	<a href="#">299266</a>	10011	486	3	79	putative protein of mammalian origin (8.8 kD) (Asb2) alternative variant fSep08, mRNA.
<a href="#">Asb2</a>	<a href="#">Asb2.gSep08</a>	<a href="#">299266</a>	8318	549	2	73	putative protein of mammalian origin (Asb2) alternative variant gSep08, mRNA.
<a href="#">Asb3</a>	<a href="#">Asb3.bSep08</a>	<a href="#">364227</a>	74532	754	6	251	ankyrin (Asb3) alternative variant bSep08, mRNA.
<a href="#">Asb6</a>	<a href="#">Asb6.bSep08</a>	<a href="#">296627</a>	3308	919	1	288	ankyrin (Asb6) alternative variant bSep08, mRNA.
<a href="#">Asb6</a>	<a href="#">Asb6.cSep08</a>	<a href="#">296627</a>	3079	690	1	197	ankyrin (Asb6) alternative variant cSep08, mRNA.
<a href="#">Asb8</a>	<a href="#">Asb8.aSep08</a>	<a href="#">315287</a>	10229	2111	4	288	ankyrin and SOCS protein, C-terminal (31.7 kD) (Asb8) alternative variant aSep08, mRNA.
<a href="#">Asb8</a>	<a href="#">Asb8.cSep08</a>	<a href="#">315287</a>	9268	986	3	262	ankyrin and SOCS protein, C-terminal (Asb8) alternative variant cSep08, mRNA.
<a href="#">Asb8</a>	<a href="#">Asb8.dSep08</a>	<a href="#">315287</a>	34357	637	5	88	putative protein (Asb8) alternative variant dSep08, mRNA.
<a href="#">Asb9</a>	<a href="#">Asb9.aSep08</a>	<a href="#">367785</a>	12191	696		201	ankyrin and SOCS protein, C-terminal (Asb9) mRNA.
<a href="#">Asb11</a>	<a href="#">Asb11.aSep08</a>	<a href="#">302666</a>	23294	1667	2	323	ankyrin and SOCS protein, C-terminal (35.3 kD) (Asb11) alternative variant aSep08, complete mRNA.
<a href="#">Asb12</a>	<a href="#">Asb12.aSep08</a>	<a href="#">503446</a>	12196	1651	2	443	ankyrin and SOCS protein, C-terminal (Asb12) alternative variant aSep08, mRNA.
<a href="#">Asb12</a>	<a href="#">Asb12.cSep08</a>	<a href="#">503446</a>	1472	454	1	106	CRA b (Asb12) alternative variant cSep08, mRNA.
<a href="#">Asb13</a>	<a href="#">Asb13.aSep08</a>	<a href="#">361268</a>	18923	2127	6	278	ankyrin and SOCS protein, C-terminal (30.0 kD) (Asb13) alternative variant aSep08, mRNA.
<a href="#">Asb13</a>	<a href="#">Asb13.bSep08</a>	<a href="#">361268</a>	29425	723	5	240	ankyrin (Asb13) alternative variant bSep08, mRNA.

<a href="#">Asb13</a>	<a href="#">Asb13.cSep08</a>	<a href="#">361268</a>	8072	406	2	93	ankyrin (Asb13) alternative variant cSep08, mRNA.
<a href="#">Asb14</a>	<a href="#">Asb14.aSep08</a>	<a href="#">680076</a>	2983	602	3	186	SOCS protein, C-terminal (Asb14) alternative variant aSep08, mRNA.
<a href="#">Asb14</a>	<a href="#">Asb14.cSep08</a>	<a href="#">680076</a>	2816	468	3	118	SOCS protein, C-terminal (Asb14) alternative variant cSep08, mRNA.
<a href="#">Asb15</a>	<a href="#">Asb15.aSep08</a>	<a href="#">500050</a>	35851	934		289	ankyrin (Asb15) mRNA.
<a href="#">Ascc1</a>	<a href="#">Ascc1.bSep08</a>	<a href="#">294512</a>	45925	764	7	254	activating signal cointegrator 1 complex subunit 1 (Ascc1) alternative variant bSep08, mRNA.
<a href="#">Ascc1</a>	<a href="#">Ascc1.cSep08</a>	<a href="#">294512</a>	36141	735	6	244	activating signal cointegrator 1 complex subunit 1 (Ascc1) alternative variant cSep08, mRNA.
<a href="#">Ascc1</a>	<a href="#">Ascc1.dSep08</a>	<a href="#">294512</a>	45763	767	7	232	activating signal cointegrator 1 complex subunit 1 (Ascc1) alternative variant dSep08, mRNA.
<a href="#">Ascc1</a>	<a href="#">Ascc1.eSep08</a>	<a href="#">294512</a>	9157	672	5	151	activating signal cointegrator 1 complex subunit 1 (Ascc1) alternative variant eSep08, mRNA.
<a href="#">Ascc1</a>	<a href="#">Ascc1.fSep08</a>	<a href="#">294512</a>	31135	746	4	94	activating signal cointegrator 1 complex subunit 1 (10.9 kD) (Ascc1) alternative variant fSep08, mRNA.
<a href="#">Ascc2</a>	<a href="#">Ascc2.bSep08</a>	<a href="#">498402</a>	10074	645	7	109	activating signal cointegrator 1 complex subunit 2 (Ascc2) alternative variant bSep08, mRNA.
<a href="#">Ascc3</a>	<a href="#">Ascc3.bSep08</a>	<a href="#">309887</a>	125171	2916	16	769	activating signal cointegrator 1 complex subunit 3 (Ascc3) alternative variant bSep08, mRNA.
<a href="#">Ascc3</a>	<a href="#">Ascc3.cSep08</a>	<a href="#">309887</a>	22387	578	3	92	activating signal cointegrator 1 complex subunit 3 (10.6 kD) (Ascc3) alternative variant cSep08, mRNA.
<a href="#">Ascc3</a>	<a href="#">Ascc3.dSep08</a>	<a href="#">309887</a>	11631	282	3	46	activating signal cointegrator 1 complex subunit 3 (Ascc3) alternative variant dSep08, mRNA.
<a href="#">ASD2.0</a>	<a href="#">ASD2.0.aSep08</a>		4116	405		135	shroom family member 4 (ASD2.0) mRNA.
<a href="#">ASD2.1</a>	<a href="#">ASD2.1.aSep08</a>		1145	1035		149	apical protein 2 (ASD2.1) mRNA.
<a href="#">Asf1a</a>	<a href="#">Asf1a.aSep08</a>	<a href="#">294408</a>	14847	2693	2	204	ASF1 anti-silencing function 1 homolog A ( <i>S. cerevisiae</i> ) (22.9 kD) (Asf1a) alternative variant aSep08, complete mRNA.
<a href="#">Asf1a</a>	<a href="#">Asf1a.bSep08</a>	<a href="#">294408</a>	12867	595	2	134	ASF1 anti-silencing function 1 homolog A ( <i>S. cerevisiae</i> ) (15.0 kD) (Asf1a) alternative variant bSep08, mRNA.
<a href="#">Asgr1</a>	<a href="#">Asgr1.bSep08</a>	<a href="#">24210</a>	3924	1185	3	236	asialoglycoprotein receptor 1 (27.3 kD) (Asgr1) alternative variant bSep08, complete mRNA.
<a href="#">Asgr1</a>	<a href="#">Asgr1.cSep08</a>	<a href="#">24210</a>	2856	678	1	197	asialoglycoprotein receptor 1 (Asgr1) alternative variant cSep08, mRNA.
<a href="#">Asgr2</a>	<a href="#">Asgr2.bSep08</a>	<a href="#">29403</a>	11761	728	3	236	asialoglycoprotein receptor 2 (Asgr2) alternative variant bSep08, mRNA.
<a href="#">Asgr2</a>	<a href="#">Asgr2.cSep08</a>	<a href="#">29403</a>	6588	835	4	204	asialoglycoprotein receptor 2 (Asgr2) alternative variant cSep08, mRNA.
<a href="#">Ash1l</a>	<a href="#">Ash1l.bSep08</a>	<a href="#">310638</a>	32873	4614	3	1537	ash1 (absent, small, or homeotic)-like ( <i>Drosophila</i> ) (Ash1l) alternative variant bSep08, mRNA.
<a href="#">Ash1l</a>	<a href="#">Ash1l.cSep08</a>	<a href="#">310638</a>	2236	617	3	152	ash1 (absent, small, or homeotic)-like ( <i>Drosophila</i> ) (Ash1l) alternative variant cSep08, mRNA.
<a href="#">Ash1l</a>	<a href="#">Ash1l.dSep08</a>	<a href="#">310638</a>	12900	427	5	141	ash1 (absent, small, or homeotic)-like ( <i>Drosophila</i> ) (Ash1l) alternative variant dSep08, mRNA.
<a href="#">Ash1l</a>	<a href="#">Ash1l.eSep08</a>	<a href="#">310638</a>	2708	372	3	124	ash1 (absent, small, or homeotic)-like ( <i>Drosophila</i> ) (Ash1l) alternative variant eSep08, mRNA.

<a href="#">Ash2l</a>	<a href="#">Ash2l.bSep08</a>	<a href="#">290829</a>	15448	1351	13	449	ash2 (absent, small, or homeotic)-like (Drosophila) (Ash2l) alternative variant bSep08, mRNA.
<a href="#">Ash2l</a>	<a href="#">Ash2l.cSep08</a>	<a href="#">290829</a>	22976	3828	15	308	ash2 (absent, small, or homeotic)-like (Drosophila) (33.0 kD) (Ash2l) alternative variant cSep08, complete mRNA.
<a href="#">Ash2l</a>	<a href="#">Ash2l.dSep08</a>	<a href="#">290829</a>	21373	814	9	271	ash2 (absent, small, or homeotic)-like (Drosophila) (Ash2l) alternative variant dSep08, mRNA.
<a href="#">Asl</a>	<a href="#">Asl.bSep08</a>	<a href="#">59085</a>	13101	1078	9	196	argininosuccinate lyase (Asl) alternative variant bSep08, mRNA.
<a href="#">Asl</a>	<a href="#">Asl.cSep08</a>	<a href="#">59085</a>	5115	415	3	137	putative protein (Asl) alternative variant cSep08, mRNA.
<a href="#">Asl</a>	<a href="#">Asl.dSep08</a>	<a href="#">59085</a>	5149	484	3	123	putative protein (Asl) alternative variant dSep08, mRNA.
<a href="#">Asl</a>	<a href="#">Asl.eSep08</a>	<a href="#">59085</a>	1043	266	4	88	argininosuccinate lyase (Asl) alternative variant eSep08, mRNA.
<a href="#">Asl</a>	<a href="#">Asl.fSep08</a>	<a href="#">59085</a>	5055	605	3	21	putative protein (Asl) alternative variant fSep08, mRNA.
<a href="#">Asmtl</a>	<a href="#">Asmtl.bSep08</a>	<a href="#">288527</a>	553	439	2	103	acetylserotonin O-methyltransferase-like (Asmtl) alternative variant bSep08, mRNA.
<a href="#">Asmtl</a>	<a href="#">Asmtl.cSep08</a>	<a href="#">288527</a>	2659	1239	6	98	acetylserotonin O-methyltransferase-like (10.6 kD) (Asmtl) alternative variant cSep08, mRNA.
<a href="#">Asmtl</a>	<a href="#">Asmtl.dSep08</a>	<a href="#">288527</a>	822	464	3	23	acetylserotonin O-methyltransferase-like (Asmtl) alternative variant dSep08, mRNA.
<a href="#">Asna1</a>	<a href="#">Asna1.aSep08</a>	<a href="#">288919</a>	8088	1351	7	348	arsA arsenite transporter, ATP-binding, homolog 1 (bacterial) (38.8 kD) (Asna1) alternative variant aSep08, complete mRNA.
<a href="#">Asns</a>	<a href="#">Asns.bSep08</a>	<a href="#">25612</a>	10606	873	5	245	asparagine synthetase (Asns) alternative variant bSep08, mRNA.
<a href="#">Asnsd1</a>	<a href="#">Asnsd1.bSep08</a>	<a href="#">299507</a>	16273	2094	2	111	putative protein, with a coiled coil domain (Asnsd1) alternative variant bSep08, mRNA.
<a href="#">Aspa</a>	<a href="#">Aspa.bSep08</a>	<a href="#">79251</a>	53844	1646	7	262	aspartoacylase (29.9 kD) (Aspa) alternative variant bSep08, mRNA.
<a href="#">Asph</a>	<a href="#">Asph.bSep08</a>	<a href="#">312981</a>	76386	795	6	202	aspartate-beta-hydroxylase (Asph) alternative variant bSep08, mRNA.
<a href="#">Asph</a>	<a href="#">Asph.dSep08</a>	<a href="#">312981</a>	43627	478	3	101	aspartate-beta-hydroxylase (Asph) alternative variant dSep08, mRNA.
<a href="#">Asphd2</a>	<a href="#">Asphd2.bSep08</a>	<a href="#">364948</a>	1787	533	2	32	putative protein (3.7 kD) (Asphd2) alternative variant bSep08, mRNA.
<a href="#">Aspn</a>	<a href="#">Aspn.aSep08</a>	<a href="#">306805</a>	23988	1784	1	451	asporin (Aspn) alternative variant aSep08, mRNA.
<a href="#">Aspscr1</a>	<a href="#">Aspscr1.aSep08</a>	<a href="#">691026</a>	33261	1924		448	putative protein, with a coiled coil domain, of eukaryotic origin (Aspscr1) mRNA.
<a href="#">Asp_Arg_Hydrox.0</a>	<a href="#">Asp_Arg_Hydrox.0.aSep08</a>		27229	2621		155	aspartate-beta-hydroxylase CRA a (Asp_Arg_Hydrox.0) mRNA.
<a href="#">Ass1</a>	<a href="#">Ass1.bSep08</a>	<a href="#">25698</a>	3340	729	2	59	argininosuccinate synthetase (Ass1) alternative variant bSep08, mRNA.
<a href="#">Ass1</a>	<a href="#">Ass1.cSep08</a>	<a href="#">25698</a>	1808	515	2	36	argininosuccinate synthetase (4.1 kD) (Ass1) alternative variant cSep08, mRNA.
<a href="#">Aste1</a>	<a href="#">Aste1.aSep08</a>	<a href="#">363130</a>	2375	837	2	139	asteroid homolog 1 (15.7 kD) (Aste1) alternative variant aSep08, mRNA.
<a href="#">Astl</a>	<a href="#">Astl.bSep08</a>	<a href="#">296129</a>	2023	747	2	183	astacin-like metalloendopeptidase (M12 family) (Astl) alternative variant bSep08, mRNA.

<a href="#">Astn1</a>	<a href="#">Astn1.aSep08</a>	<a href="#">304900</a>	32001	1160	6	368	astrotactin 1 (Astn1) alternative variant aSep08, mRNA.
<a href="#">Astn1</a>	<a href="#">Astn1.bSep08</a>	<a href="#">304900</a>	7344	471	1	95	astrotactin 1 (Astn1) alternative variant bSep08, mRNA.
<a href="#">Asxl2</a>	<a href="#">Asxl2.dSep08</a>	<a href="#">313922</a>	7425	1117	2	64	additional sex combs like 2 (Drosophila) (7.2 kD) (Asxl2) alternative variant dSep08, mRNA.
<a href="#">Atad1</a>	<a href="#">Atad1.bSep08</a>	<a href="#">309532</a>	68770	3933	11	361	neuroprotective protein 6 (40.7 kD) (Atad1) alternative variant bSep08, mRNA.
<a href="#">Atad2</a>	<a href="#">Atad2.bSep08</a>	<a href="#">314993</a>	15514	3395	13	544	bromodomain containing protein (62.2 kD) (Atad2) alternative variant bSep08, mRNA.
<a href="#">Atad3a</a>	<a href="#">Atad3a.bSep08</a>	<a href="#">298682</a>	12308	1307	11	302	aaa-atpase tob3 (33.5 kD) (Atad3a) alternative variant bSep08, mRNA.
<a href="#">Atad5</a>	<a href="#">Atad5.aSep08</a>	<a href="#">303348</a>	13054	716		238	putative protein of eukaryotic origin (Atad5) mRNA.
<a href="#">Atcay</a>	<a href="#">Atcay.bSep08</a>	<a href="#">362826</a>	1713	436	3	66	ataxia, cerebellar, Cayman type (Atcay) alternative variant bSep08, mRNA.
<a href="#">Ate1</a>	<a href="#">Ate1.bSep08</a>	<a href="#">293526</a>	122960	1276	7	189	arginine-tRNA-protein transferase 1 (21.6 kD) (Ate1) alternative variant bSep08, mRNA.
<a href="#">Ate1</a>	<a href="#">Ate1.cSep08</a>	<a href="#">293526</a>	10818	446	4	87	arginine-tRNA-protein transferase 1 (Ate1) alternative variant cSep08, mRNA.
<a href="#">Atf1</a>	<a href="#">Atf1.aSep08</a>	<a href="#">315305</a>	43635	2328	3	359	activating transcription factor 1 (Atf1) alternative variant aSep08, mRNA.
<a href="#">Atf1</a>	<a href="#">Atf1.bSep08</a>	<a href="#">315305</a>	40647	1177	4	224	activating transcription factor 1 (23.9 kD) (Atf1) alternative variant bSep08, mRNA.
<a href="#">Atf1</a>	<a href="#">Atf1.cSep08</a>	<a href="#">315305</a>	35808	723	2	211	activating transcription factor 1 (Atf1) alternative variant cSep08, mRNA.
<a href="#">Atf2</a>	<a href="#">Atf2.bSep08</a>	<a href="#">81647</a>	69111	1154	11	314	activating transcription factor 2 (Atf2) alternative variant bSep08, mRNA.
<a href="#">Atf3</a>	<a href="#">Atf3.bSep08</a>	<a href="#">25389</a>	87075	591	3	196	activating transcription factor 3 (Atf3) alternative variant bSep08, mRNA.
<a href="#">Atf3</a>	<a href="#">Atf3.cSep08</a>	<a href="#">25389</a>	29494	347	2	115	activating transcription factor 3 (Atf3) alternative variant cSep08, mRNA.
<a href="#">Atf4</a>	<a href="#">Atf4.bSep08</a>	<a href="#">79255</a>	1514	975	2	92	putative mitochondrial protein (9.9 kD) (Atf4) alternative variant bSep08, mRNA.
<a href="#">Atf5</a>	<a href="#">Atf5.bSep08</a>	<a href="#">282840</a>	3277	643	3	114	activating transcription factor 5 (Atf5) alternative variant bSep08, mRNA.
<a href="#">Atf7</a>	<a href="#">Atf7.bSep08</a>	<a href="#">315333</a>	108569	1781	12	483	activating transcription factor 7 (51.8 kD) (Atf7) alternative variant bSep08, mRNA.
<a href="#">Atf7</a>	<a href="#">Atf7.cSep08</a>	<a href="#">315333</a>	74548	417	5	107	activating transcription factor 7 (Atf7) alternative variant cSep08, mRNA.
<a href="#">Atf7ip</a>	<a href="#">Atf7ip.aSep08</a>	<a href="#">312800</a>	82894	4504	13	1283	activating transcription factor 7 interacting protein (136.2 kD) (Atf7ip) alternative variant aSep08, mRNA.
<a href="#">Atf7ip</a>	<a href="#">Atf7ip.bSep08</a>	<a href="#">312800</a>	39261	1484	6	436	activating transcription factor 7 interacting protein (Atf7ip) alternative variant bSep08, mRNA.
<a href="#">Atf7ip</a>	<a href="#">Atf7ip.dSep08</a>	<a href="#">312800</a>	14647	634	5	211	activating transcription factor 7 interacting protein (Atf7ip) alternative variant dSep08, mRNA.
<a href="#">Atf7ip</a>	<a href="#">Atf7ip.eSep08</a>	<a href="#">312800</a>	6408	557	2	156	activating transcription factor 7 interacting protein (Atf7ip) alternative variant eSep08, mRNA.
<a href="#">Atf7ip</a>	<a href="#">Atf7ip.fSep08</a>	<a href="#">312800</a>	8099	366	3	121	activating transcription factor 7 interacting protein (Atf7ip) alternative variant fSep08, mRNA.



<a href="#">Atf7ip</a>	<a href="#">Atf7ip.gSep08</a>	<a href="#">312800</a>	6166	375	3	75	activating transcription factor 7 interacting protein (Atf7ip) alternative variant gSep08, mRNA.
<a href="#">Atf7ip2</a>	<a href="#">Atf7ip2.bSep08</a>	<a href="#">497859</a>	15398	373	1	124	activating transcription factor 7 interacting protein 2 (Atf7ip2) alternative variant bSep08, mRNA.
<a href="#">Atg2a</a>	<a href="#">Atg2a.bSep08</a>	<a href="#">689688</a>	1693	755	1	131	ATG2 autophagy related 2 homolog A ( <i>S. cerevisiae</i> ) (Atg2a) alternative variant bSep08, mRNA.
<a href="#">Atg3</a>	<a href="#">Atg3.aSep08</a>	<a href="#">171415</a>	12663	1776	6	437	autophagy-related 3 (yeast) (Atg3) alternative variant aSep08, mRNA.
<a href="#">Atg3</a>	<a href="#">Atg3.bSep08</a>	<a href="#">171415</a>	28312	1374	11	325	autophagy-related 3 (yeast) (37.1 kD) (Atg3) alternative variant bSep08, mRNA.
<a href="#">Atg4b</a>	<a href="#">Atg4b.aSep08</a>	<a href="#">316640</a>	30758	1690	9	360	autophagy-related 4B (yeast) (Atg4b) alternative variant aSep08, mRNA.
<a href="#">Atg4b</a>	<a href="#">Atg4b.cSep08</a>	<a href="#">316640</a>	27433	894	6	143	autophagy-related 4B (yeast) (16.8 kD) (Atg4b) alternative variant cSep08, mRNA.
<a href="#">Atg4c</a>	<a href="#">Atg4c.bSep08</a>	<a href="#">313391</a>	7102	702	4	176	autophagy-related 4C (yeast) (Atg4c) alternative variant bSep08, mRNA.
<a href="#">Atg7</a>	<a href="#">Atg7.bSep08</a>	<a href="#">312647</a>	157478	1777	5	510	ATG7 autophagy related 7 homolog ( <i>S. cerevisiae</i> ) (Atg7) alternative variant bSep08, mRNA.
<a href="#">Atg7</a>	<a href="#">Atg7.cSep08</a>	<a href="#">312647</a>	19208	724	5	171	ATG7 autophagy related 7 homolog ( <i>S. cerevisiae</i> ) (Atg7) alternative variant cSep08, mRNA.
<a href="#">Atg7</a>	<a href="#">Atg7.dSep08</a>	<a href="#">312647</a>	1097	573	2	68	ATG7 autophagy related 7 homolog ( <i>S. cerevisiae</i> ) (7.7 kD) (Atg7) alternative variant dSep08, mRNA.
<a href="#">Atg9a</a>	<a href="#">Atg9a.bSep08</a>	<a href="#">363254</a>	5533	1445	8	397	autophagy-related 9A CRA a (Atg9a) alternative variant bSep08, mRNA.
<a href="#">Atg9a</a>	<a href="#">Atg9a.cSep08</a>	<a href="#">363254</a>	1903	1289	3	141	putative protein (15.3 kD) (Atg9a) alternative variant cSep08, mRNA.
<a href="#">Atg9a</a>	<a href="#">Atg9a.dSep08</a>	<a href="#">363254</a>	842	735	2	83	autophagy-related 9A CRA a (Atg9a) alternative variant dSep08, mRNA.
<a href="#">Atg9a</a>	<a href="#">Atg9a.eSep08</a>	<a href="#">363254</a>	1600	242	2	80	autophagy-related 9A CRA a (Atg9a) alternative variant eSep08, mRNA.
<a href="#">Atg9a</a>	<a href="#">Atg9a.gSep08</a>	<a href="#">363254</a>	2346	405	6	36	putative protein (3.9 kD) (Atg9a) alternative variant gSep08, mRNA.
<a href="#">Atg10</a>	<a href="#">Atg10.bSep08</a>	<a href="#">688555</a>	68222	328	3	89	autophagy-related 10 ( <i>S. cerevisiae</i> ) (Atg10) alternative variant bSep08, mRNA.
<a href="#">Atg10</a>	<a href="#">Atg10.cSep08</a>	<a href="#">688555</a>	42408	461	2	45	autophagy-related 10 ( <i>S. cerevisiae</i> ) (Atg10) alternative variant cSep08, mRNA.
<a href="#">Atg16l1</a>	<a href="#">Atg16l1.bSep08</a>	<a href="#">363278</a>	17369	843	9	280	1 16 (Atg16l1) alternative variant bSep08, mRNA.
<a href="#">Atg16l1</a>	<a href="#">Atg16l1.cSep08</a>	<a href="#">363278</a>	9958	522	5	115	1 16 (Atg16l1) alternative variant cSep08, mRNA.
<a href="#">Atg16l1</a>	<a href="#">Atg16l1.dSep08</a>	<a href="#">363278</a>	1335	727	2	83	putative protein (9.1 kD) (Atg16l1) alternative variant dSep08, mRNA.
<a href="#">Atg16l2</a>	<a href="#">Atg16l2.aSep08</a>	<a href="#">308865</a>	3991	962	7	231	ATG16 autophagy related 16-like 2 (Atg16l2) alternative variant aSep08, mRNA.
<a href="#">Atg16l2</a>	<a href="#">Atg16l2.bSep08</a>	<a href="#">308865</a>	2535	823	5	151	ATG16 autophagy related 16-like 2 (16.9 kD) (Atg16l2) alternative variant bSep08, mRNA.
<a href="#">Atg16l2</a>	<a href="#">Atg16l2.cSep08</a>	<a href="#">308865</a>	2322	582	5	116	CRA b (Atg16l2) alternative variant cSep08, mRNA.
<a href="#">Atg16l2</a>	<a href="#">Atg16l2.dSep08</a>	<a href="#">308865</a>	724	491	2	88	ATG16 autophagy related 16-like 2 (Atg16l2) alternative variant dSep08, mRNA.

<a href="#">Ath1</a>	<a href="#">Ath1.aSep08</a>	<a href="#">309103</a>	2446	888	7	295	ATH1, acid trehalase-like 1 (yeast) (Ath1) alternative variant aSep08, mRNA.
<a href="#">Ath1</a>	<a href="#">Ath1.bSep08</a>	<a href="#">309103</a>	1184	771	4	169	ATH1, acid trehalase-like 1 (yeast) (Ath1) alternative variant bSep08, mRNA.
<a href="#">Ath1</a>	<a href="#">Ath1.cSep08</a>	<a href="#">309103</a>	1893	1396	3	133	ATH1, acid trehalase-like 1 (yeast) (15.1 kD) (Ath1) alternative variant cSep08, mRNA.
<a href="#">Atic</a>	<a href="#">Atic.bSep08</a>	<a href="#">81643</a>	6014	858	5	216	5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase (Atic) alternative variant bSep08, mRNA.
<a href="#">Atic</a>	<a href="#">Atic.cSep08</a>	<a href="#">81643</a>	6984	731	8	153	5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase (Atic) alternative variant cSep08, mRNA.
<a href="#">Atm</a>	<a href="#">Atm.bSep08</a>	<a href="#">300711</a>	11846	921	5	181	ataxia telangiectasia mutated homolog (human) (21.2 kD) (Atm) alternative variant bSep08, mRNA.
<a href="#">Atm</a>	<a href="#">Atm.cSep08</a>	<a href="#">300711</a>	6190	480	4	146	ataxia telangiectasia mutated homolog (human) (17.4 kD) (Atm) alternative variant cSep08, mRNA.
<a href="#">Atm</a>	<a href="#">Atm.dSep08</a>	<a href="#">300711</a>	2677	527	2	75	ataxia telangiectasia mutated homolog (human) (Atm) alternative variant dSep08, mRNA.
<a href="#">Atn1</a>	<a href="#">Atn1.bSep08</a>	<a href="#">29515</a>	3437	1729	2	325	atrophin 1 (Atn1) alternative variant bSep08, mRNA.
<a href="#">Atp1a1</a>	<a href="#">Atp1a1.bSep08</a>	<a href="#">24211</a>	7177	1569	10	208	na+ K+ -ATPase alpha (24.4 kD) (Atp1a1) alternative variant bSep08, mRNA.
<a href="#">Atp1a1</a>	<a href="#">Atp1a1.cSep08</a>	<a href="#">24211</a>	1108	709	2	125	putative mitochondrial protein (13.6 kD) (Atp1a1) alternative variant cSep08, mRNA.
<a href="#">Atp1a1</a>	<a href="#">Atp1a1.dSep08</a>	<a href="#">24211</a>	4167	360	3	119	na+ K+ -ATPase alpha (Atp1a1) alternative variant dSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.cSep08</a>	<a href="#">24212</a>	7001	767	5	255	ATPase Na+ K+ transporting alpha 4 polypeptide (Atp1a2andAtp1a4) alternative variant cSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.cSep08</a>	<a href="#">29132</a>	7001	767	5	255	ATPase Na+ K+ transporting alpha 4 polypeptide (Atp1a2andAtp1a4) alternative variant cSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.dSep08</a>	<a href="#">24212</a>	40242	712	5	237	na+ K+ (Atp1a2andAtp1a4) alternative variant dSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.dSep08</a>	<a href="#">29132</a>	40242	712	5	237	na+ K+ (Atp1a2andAtp1a4) alternative variant dSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.eSep08</a>	<a href="#">24212</a>	7845	1797	5	134	ATPase Na+ K+ alpha (Atp1a2andAtp1a4) alternative variant eSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.eSep08</a>	<a href="#">29132</a>	7845	1797	5	134	ATPase Na+ K+ alpha (Atp1a2andAtp1a4) alternative variant eSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.fSep08</a>	<a href="#">24212</a>	1253	519	4	134	ATPase Na+ K+ alpha (Atp1a2andAtp1a4) alternative variant fSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.fSep08</a>	<a href="#">29132</a>	1253	519	4	134	ATPase Na+ K+ alpha (Atp1a2andAtp1a4) alternative variant fSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.gSep08</a>	<a href="#">24212</a>	511	397	2	87	na+ K+ -ATPase (Atp1a2andAtp1a4) alternative variant gSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.gSep08</a>	<a href="#">29132</a>	511	397	2	87	na+ K+ -ATPase (Atp1a2andAtp1a4) alternative variant gSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.hSep08</a>	<a href="#">24212</a>	1090	576	3	60	ATPase Na+ K+ transporting alpha 2 polypeptide (Atp1a2andAtp1a4) alternative variant hSep08, mRNA.

<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.hSep08</a>	<a href="#">29132</a>	1090	576	3	60	ATPase Na <sup>+</sup> K <sup>+</sup> transporting alpha 2 polypeptide (Atp1a2andAtp1a4) alternative variant hSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.iSep08</a>	<a href="#">24212</a>	604	351	2	50	putative protein (Atp1a2andAtp1a4) alternative variant iSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.iSep08</a>	<a href="#">29132</a>	604	351	2	50	putative protein (Atp1a2andAtp1a4) alternative variant iSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.jSep08</a>	<a href="#">24212</a>	443	257	2	33	ATPase Na <sup>+</sup> K <sup>+</sup> transporting alpha polypeptide (Atp1a2andAtp1a4) alternative variant jSep08, mRNA.
<a href="#">Atp1a2andAtp1a4</a>	<a href="#">Atp1a2andAtp1a4.jSep08</a>	<a href="#">29132</a>	443	257	2	33	ATPase Na <sup>+</sup> K <sup>+</sup> transporting alpha polypeptide (Atp1a2andAtp1a4) alternative variant jSep08, mRNA.
<a href="#">Atp1b1</a>	<a href="#">Atp1b1.bSep08</a>	<a href="#">25650</a>	14381	921	1	143	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide (Atp1b1) alternative variant bSep08, mRNA.
<a href="#">Atp1b2</a>	<a href="#">Atp1b2.bSep08</a>	<a href="#">24214</a>	3430	2166		210	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 2 polypeptide (Atp1b2) alternative variant bSep08, mRNA.
<a href="#">Atp1b3</a>	<a href="#">Atp1b3.bSep08</a>	<a href="#">25390</a>	11888	962	4	83	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 3 polypeptide (Atp1b3) alternative variant bSep08, mRNA.
<a href="#">Atp1b3</a>	<a href="#">Atp1b3.dSep08</a>	<a href="#">25390</a>	11864	293	3	54	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 3 polypeptide (Atp1b3) alternative variant dSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.aSep08</a>	<a href="#">29693</a>	13152	4000	12	648	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant aSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.bSep08</a>	<a href="#">29693</a>	30621	1767	10	482	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (52.7 kD) (Atp2a2) alternative variant bSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.cSep08</a>	<a href="#">29693</a>	10174	1947	6	441	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant cSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.dSep08</a>	<a href="#">29693</a>	6605	1559	7	295	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant dSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.eSep08</a>	<a href="#">29693</a>	35306	1278	8	250	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant eSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.fSep08</a>	<a href="#">29693</a>	5613	883	6	227	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant fSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.hSep08</a>	<a href="#">29693</a>	707	623	2	130	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant hSep08, mRNA.
<a href="#">Atp2a2</a>	<a href="#">Atp2a2.iSep08</a>	<a href="#">29693</a>	912	297	2	97	ATPase Ca <sup>++</sup> transporting cardiac muscle slow twitch 2 (Atp2a2) alternative variant iSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.bSep08</a>	<a href="#">29598</a>	75988	1783	5	479	ATPase Plasma membrane (Atp2b1) alternative variant bSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.cSep08</a>	<a href="#">29598</a>	23494	5325	10	314	ATPase plasma membrane 1 (Atp2b1) alternative variant cSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.dSep08</a>	<a href="#">29598</a>	11248	688	4	228	ATPase Plasma membrane 1 (Atp2b1) alternative variant dSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.eSep08</a>	<a href="#">29598</a>	3676	704	3	184	membrane calcium-transporting ATPase 1 (Atp2b1) alternative variant eSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.fSep08</a>	<a href="#">29598</a>	5497	796	3	172	plasma membrane ATPase (19.3 kD) (Atp2b1) alternative variant fSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.gSep08</a>	<a href="#">29598</a>	1151	749	2	112	plasma membrane ATPase (Atp2b1) alternative variant gSep08, mRNA.

<a href="#">Atp2b1</a>	<a href="#">Atp2b1.hSep08</a>	<a href="#">29598</a>	53375	621	2	103	putative protein (Atp2b1) alternative variant hSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.jSep08</a>	<a href="#">29598</a>	3822	763	2	76	ATPase plasma membrane (8.2 kD) (Atp2b1) alternative variant jSep08, mRNA.
<a href="#">Atp2b1</a>	<a href="#">Atp2b1.nSep08</a>	<a href="#">29598</a>	3691	478	4	80	putative protein (Atp2b1) alternative variant nSep08, mRNA.
<a href="#">Atp2b2</a>	<a href="#">Atp2b2.bSep08</a>	<a href="#">24215</a>	15347	4395	5	291	ATPase, Ca <sup>++</sup> transporting, plasma membrane 2 (Atp2b2) alternative variant bSep08, mRNA.
<a href="#">Atp2b2</a>	<a href="#">Atp2b2.cSep08</a>	<a href="#">24215</a>	44242	1652	3	54	ATPase, Ca <sup>++</sup> transporting, plasma membrane 2 (5.6 kD) (Atp2b2) alternative variant cSep08, mRNA.
<a href="#">Atp2b3</a>	<a href="#">Atp2b3.bSep08</a>	<a href="#">29599</a>	10014	1784	6	554	ATPase, Ca <sup>++</sup> transporting, plasma membrane 3 (Atp2b3) alternative variant bSep08, mRNA.
<a href="#">Atp2b3</a>	<a href="#">Atp2b3.cSep08</a>	<a href="#">29599</a>	5052	386	2	115	ATPase, Ca <sup>++</sup> transporting, plasma membrane 3 (Atp2b3) alternative variant cSep08, mRNA.
<a href="#">Atp2b3</a>	<a href="#">Atp2b3.dSep08</a>	<a href="#">29599</a>	1029	408	2	74	ATPase, Ca <sup>++</sup> transporting, plasma membrane 3 (Atp2b3) alternative variant dSep08, mRNA.
<a href="#">Atp2b4</a>	<a href="#">Atp2b4.bSep08</a>	<a href="#">29600</a>	6121	2175	2	74	ATPase, Ca <sup>++</sup> transporting, plasma membrane 4 (Atp2b4) alternative variant bSep08, mRNA.
<a href="#">Atp2c1</a>	<a href="#">Atp2c1.bSep08</a>	<a href="#">170699</a>	56569	4152	24	820	ATPase, Ca <sup>++</sup> transporting, type 2C, member 1 (Atp2c1) alternative variant bSep08, mRNA.
<a href="#">Atp2c1</a>	<a href="#">Atp2c1.cSep08</a>	<a href="#">170699</a>	4655	375	3	92	ATPase, Ca <sup>++</sup> transporting, type 2C, member 1 (Atp2c1) alternative variant cSep08, mRNA.
<a href="#">Atp2c1</a>	<a href="#">Atp2c1.dSep08</a>	<a href="#">170699</a>	29784	566	4	98	ATPase, Ca <sup>++</sup> transporting, type 2C, member 1 (Atp2c1) alternative variant dSep08, mRNA.
<a href="#">Atp4a</a>	<a href="#">Atp4a.aSep08</a>	<a href="#">24216</a>	1763	652		216	ATPase, H <sup>+</sup> /K <sup>+</sup> exchanging, alpha polypeptide (Atp4a) mRNA.
<a href="#">Atp5a1</a>	<a href="#">Atp5a1.bSep08</a>	<a href="#">65262</a>	3522	435	2	137	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, alpha subunit 1, cardiac muscle (Atp5a1) alternative variant bSep08, mRNA.
<a href="#">Atp5b</a>	<a href="#">Atp5b.bSep08</a>	<a href="#">171374</a>	2473	1320	5	283	ATP synthase, H <sup>+</sup> transporting mitochondrial F1 complex, beta subunit (29.7 kD) (Atp5b) alternative variant bSep08, mRNA.
<a href="#">Atp5b</a>	<a href="#">Atp5b.cSep08</a>	<a href="#">171374</a>	2122	520	3	122	ATP synthase, H <sup>+</sup> transporting mitochondrial F1 complex, beta subunit (Atp5b) alternative variant cSep08, mRNA.
<a href="#">Atp5b</a>	<a href="#">Atp5b.fSep08</a>	<a href="#">171374</a>	1158	493	3	81	ATP synthase, H <sup>+</sup> transporting mitochondrial F1 complex, beta subunit (8.8 kD) (Atp5b) alternative variant fSep08, mRNA.
<a href="#">Atp5c1</a>	<a href="#">Atp5c1.bSep08</a>	<a href="#">116550</a>	21908	1436	8	273	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, gamma polypeptide 1 (30.2 kD) (Atp5c1) alternative variant bSep08, mRNA.
<a href="#">Atp5c1</a>	<a href="#">Atp5c1.cSep08</a>	<a href="#">116550</a>	3985	374	2	44	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, gamma polypeptide 1 (Atp5c1) alternative variant cSep08, mRNA.
<a href="#">Atp5d</a>	<a href="#">Atp5d.bSep08</a>	<a href="#">245965</a>	5327	2473	4	184	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, delta subunit and hypothetical protein LOC690935 (19.3 kD) (Atp5d) alternative variant bSep08, mRNA.

<a href="#">Atp5d</a>	<a href="#">Atp5d.bSep08</a>	<a href="#">690935</a>	5327	2473	4	184	ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit and hypothetical protein LOC690935 (19.3 kD) (Atp5d) alternative variant bSep08, mRNA.
<a href="#">Atp5f1</a>	<a href="#">Atp5f1.aSep08</a>	<a href="#">171375</a>	10634	1090	6	314	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit B1 (35.5 kD) (Atp5f1) alternative variant aSep08, mRNA.
<a href="#">Atp5f1</a>	<a href="#">Atp5f1.cSep08</a>	<a href="#">171375</a>	4270	627	3	119	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit B1 (13.1 kD) (Atp5f1) alternative variant cSep08, mRNA.
<a href="#">Atp5f1</a>	<a href="#">Atp5f1.fSep08</a>	<a href="#">171375</a>	595	345	2	58	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit B1 (Atp5f1) alternative variant fSep08, mRNA.
<a href="#">Atp5g1</a>	<a href="#">Atp5g1.aSep08</a>	<a href="#">29754</a>	2584	1103	4	145	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C1 (subunit 9) (15.2 kD) (Atp5g1) alternative variant aSep08, mRNA.
<a href="#">Atp5g1</a>	<a href="#">Atp5g1.bSep08</a>	<a href="#">29754</a>	2648	558	5	136	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C1 (subunit 9) (14.2 kD) (Atp5g1) alternative variant bSep08, mRNA.
<a href="#">Atp5g1</a>	<a href="#">Atp5g1.dSep08</a>	<a href="#">29754</a>	2275	384	5	111	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C1 (subunit 9) (Atp5g1) alternative variant dSep08, mRNA.
<a href="#">Atp5g1</a>	<a href="#">Atp5g1.eSep08</a>	<a href="#">29754</a>	1812	886	2	57	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C1 (subunit 9) (6.0 kD) (Atp5g1) alternative variant eSep08, mRNA.
<a href="#">Atp5g2</a>	<a href="#">Atp5g2.aSep08</a>	<a href="#">171082</a>	7581	545	5	141	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C2 (subunit 9) (14.9 kD) (Atp5g2) alternative variant aSep08, mRNA.
<a href="#">Atp5g3</a>	<a href="#">Atp5g3.aSep08</a>	<a href="#">114630</a>	2854	930	4	185	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9) (Atp5g3) alternative variant aSep08, mRNA.
<a href="#">Atp5g3</a>	<a href="#">Atp5g3.bSep08</a>	<a href="#">114630</a>	2625	691	4	179	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9) (Atp5g3) alternative variant bSep08, mRNA.
<a href="#">Atp5g3</a>	<a href="#">Atp5g3.cSep08</a>	<a href="#">114630</a>	2102	603	3	139	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9) (Atp5g3) alternative variant cSep08, mRNA.
<a href="#">Atp5g3</a>	<a href="#">Atp5g3.dSep08</a>	<a href="#">114630</a>	2045	862	2	101	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9) (Atp5g3) alternative variant dSep08, mRNA.
<a href="#">Atp5h</a>	<a href="#">Atp5h.bSep08</a>	<a href="#">641434</a>	4395	940	4	108	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d (12.0 kD) (Atp5h) alternative variant bSep08, complete mRNA.
<a href="#">Atp5h</a>	<a href="#">Atp5h.cSep08</a>	<a href="#">641434</a>	4984	348	5	107	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d (Atp5h) alternative variant cSep08, mRNA.
<a href="#">Atp5h</a>	<a href="#">Atp5h.dSep08</a>	<a href="#">641434</a>	589	459	2	82	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d (8.9 kD) (Atp5h) alternative variant dSep08, mRNA.

<a href="#">Atp5i</a>	<a href="#">Atp5i.bSep08</a>	<a href="#">140608</a>	1119	524	3	49	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit E (5.9 kD) (Atp5i) alternative variant bSep08, complete mRNA.
<a href="#">Atp5j</a>	<a href="#">Atp5j.aSep08</a>	<a href="#">94271</a>	7099	617	4	108	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6 (12.5 kD) (Atp5j) alternative variant aSep08, mRNA.
<a href="#">Atp5j</a>	<a href="#">Atp5j.bSep08</a>	<a href="#">94271</a>	7084	594	4	108	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6 (12.5 kD) (Atp5j) alternative variant bSep08, mRNA.
<a href="#">Atp5j2</a>	<a href="#">Atp5j2.aSep08</a>	<a href="#">690441</a>	6373	441	4	88	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F2 (10.5 kD) (Atp5j2) alternative variant aSep08, complete mRNA.
<a href="#">Atp5l</a>	<a href="#">Atp5l.bSep08</a>	<a href="#">300677</a>	5825	732	2	81	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit G (9.0 kD) (Atp5l) alternative variant bSep08, mRNA.
<a href="#">Atp5o</a>	<a href="#">Atp5o.bSep08</a>	<a href="#">192241</a>	1963	792	4	177	synthase mitochondrial (19.5 kD) (Atp5o) alternative variant bSep08, mRNA.
<a href="#">Atp5o</a>	<a href="#">Atp5o.cSep08</a>	<a href="#">192241</a>	2918	463	3	92	synthase mitochondrial (Atp5o) alternative variant cSep08, complete mRNA.
<a href="#">Atp5sl</a>	<a href="#">Atp5sl.bSep08</a>	<a href="#">361520</a>	4908	792	2	143	ATP5S-like (Atp5sl) alternative variant bSep08, mRNA.
<a href="#">Atp6ap1</a>	<a href="#">Atp6ap1.bSep08</a>	<a href="#">83615</a>	3788	1331	7	300	ATPase, H+ transporting, lysosomal accessory protein 1 (Atp6ap1) alternative variant bSep08, mRNA.
<a href="#">Atp6ap1</a>	<a href="#">Atp6ap1.cSep08</a>	<a href="#">83615</a>	4808	1336	6	190	ATPase, H+ transporting, lysosomal accessory protein 1 (20.6 kD) (Atp6ap1) alternative variant cSep08, mRNA.
<a href="#">Atp6ap2</a>	<a href="#">Atp6ap2.aSep08</a>	<a href="#">302526</a>	27917	2126	1	390	ATPase, H+ transporting, lysosomal accessory protein 2 (Atp6ap2) alternative variant aSep08, mRNA.
<a href="#">Atp6ap2</a>	<a href="#">Atp6ap2.bSep08</a>	<a href="#">302526</a>	48449	669	2	101	ATPase, H+ transporting, lysosomal accessory protein 2 (Atp6ap2) alternative variant bSep08, mRNA.
<a href="#">Atp6v0a1</a>	<a href="#">Atp6v0a1.aSep08</a>	<a href="#">29757</a>	27141	2771	12	497	ATPase, H+ transporting, lysosomal V0 subunit A1 (Atp6v0a1) alternative variant aSep08, mRNA.
<a href="#">Atp6v0a1</a>	<a href="#">Atp6v0a1.cSep08</a>	<a href="#">29757</a>	20808	1855	8	306	ATPase, H+ transporting, lysosomal V0 subunit A1 (Atp6v0a1) alternative variant cSep08, mRNA.
<a href="#">Atp6v0a1</a>	<a href="#">Atp6v0a1.dSep08</a>	<a href="#">29757</a>	4562	537	3	64	ATPase, H+ transporting, lysosomal V0 subunit A1 (Atp6v0a1) alternative variant dSep08, mRNA.
<a href="#">Atp6v0a1</a>	<a href="#">Atp6v0a1.eSep08</a>	<a href="#">29757</a>	8651	430	4	37	ATPase, H+ transporting, lysosomal V0 subunit A1 (Atp6v0a1) alternative variant eSep08, mRNA.
<a href="#">Atp6v0a2</a>	<a href="#">Atp6v0a2.aSep08</a>	<a href="#">116455</a>	31742	3502	19	886	ATPase, H+ transporting, lysosomal V0 subunit A2 (Atp6v0a2) alternative variant aSep08, mRNA.
<a href="#">Atp6v0a2</a>	<a href="#">Atp6v0a2.cSep08</a>	<a href="#">116455</a>	10184	2213	10	463	ATPase, H+ transporting, lysosomal V0 subunit A2 (Atp6v0a2) alternative variant cSep08, mRNA.
<a href="#">Atp6v0a2</a>	<a href="#">Atp6v0a2.dSep08</a>	<a href="#">116455</a>	3955	748	5	248	ATPase, H+ transporting, lysosomal V0 subunit A2 (Atp6v0a2) alternative variant dSep08, mRNA.
<a href="#">Atp6v0a4</a>	<a href="#">Atp6v0a4.bSep08</a>	<a href="#">296981</a>	37951	2606	16	657	ATPase, H+ transporting, lysosomal V0 subunit A4 (76.0 kD) (Atp6v0a4) alternative variant bSep08, mRNA.
<a href="#">Atp6v0a4</a>	<a href="#">Atp6v0a4.cSep08</a>	<a href="#">296981</a>	28397	697	7	213	ATPase, H+ transporting, lysosomal V0 subunit A4 (Atp6v0a4) alternative variant cSep08, mRNA.

<a href="#">Atp6v0b</a>	<a href="#">Atp6v0b.aSep08</a>	<a href="#">298451</a>	3028	1005	8	239	ATPase, H+ transporting, lysosomal V0 subunit B (Atp6v0b) alternative variant aSep08, mRNA.
<a href="#">Atp6v0b</a>	<a href="#">Atp6v0b.cSep08</a>	<a href="#">298451</a>	1615	1167	4	83	ATPase, H+ transporting, lysosomal V0 subunit B (Atp6v0b) alternative variant cSep08, mRNA.
<a href="#">Atp6v0b</a>	<a href="#">Atp6v0b.dSep08</a>	<a href="#">298451</a>	1304	723	2	69	ATPase, H+ transporting, lysosomal V0 subunit B (6.9 kD) (Atp6v0b) alternative variant dSep08, mRNA.
<a href="#">Atp6v0c</a>	<a href="#">Atp6v0c.aSep08</a>	<a href="#">170667</a>	5089	878	4	182	ATPase, H+ transporting, lysosomal V0 subunit C (Atp6v0c) alternative variant aSep08, mRNA.
<a href="#">Atp6v0c</a>	<a href="#">Atp6v0c.bSep08</a>	<a href="#">170667</a>	3983	620	3	164	ATPase, H+ transporting, lysosomal V0 subunit C (Atp6v0c) alternative variant bSep08, mRNA.
<a href="#">Atp6v0c</a>	<a href="#">Atp6v0c.dSep08</a>	<a href="#">170667</a>	2369	2276	2	128	ATPase, H+ transporting, lysosomal V0 subunit C (13.0 kD) (Atp6v0c) alternative variant dSep08, mRNA.
<a href="#">Atp6v0d1</a>	<a href="#">Atp6v0d1.bSep08</a>	<a href="#">291969</a>	43972	1488	7	323	ATPase, H+ transporting, lysosomal V0 subunit D1 (37.2 kD) (Atp6v0d1) alternative variant bSep08, complete mRNA.
<a href="#">Atp6v0d1</a>	<a href="#">Atp6v0d1.cSep08</a>	<a href="#">291969</a>	9661	1537	2	136	ATPase, H+ transporting, lysosomal V0 subunit D1 (Atp6v0d1) alternative variant cSep08, mRNA.
<a href="#">Atp6v0d1</a>	<a href="#">Atp6v0d1.dSep08</a>	<a href="#">291969</a>	821	685	2	77	ATPase, H+ transporting, lysosomal V0 subunit D1 (Atp6v0d1) alternative variant dSep08, mRNA.
<a href="#">Atp6v0d1</a>	<a href="#">Atp6v0d1.fSep08</a>	<a href="#">291969</a>	42695	897	5	64	ATPase, H+ transporting, lysosomal V0 subunit D1 (7.3 kD) (Atp6v0d1) alternative variant fSep08, mRNA.
<a href="#">Atp6v0d2</a>	<a href="#">Atp6v0d2.bSep08</a>	<a href="#">297932</a>	1508	265	1	26	ATPase, H+ transporting, lysosomal V0 subunit D2 (Atp6v0d2) alternative variant bSep08, mRNA.
<a href="#">Atp6v0e1</a>	<a href="#">Atp6v0e1.bSep08</a>	<a href="#">94170</a>	22693	327	2	57	ATPase, H+ transporting, lysosomal V0 subunit E1 (6.3 kD) (Atp6v0e1) alternative variant bSep08, mRNA.
<a href="#">Atp6v0e1</a>	<a href="#">Atp6v0e1.cSep08</a>	<a href="#">94170</a>	5048	835	1	38	ATPase, H+ transporting, lysosomal V0 subunit E1 (4.2 kD) (Atp6v0e1) alternative variant cSep08, mRNA.
<a href="#">Atp6v0e2</a>	<a href="#">Atp6v0e2.bSep08</a>	<a href="#">436582</a>	1147	757	2	42	ATPase, H+ transporting, lysosomal V0 subunit E2 (4.7 kD) (Atp6v0e2) alternative variant bSep08, mRNA.
<a href="#">Atp6v1b2</a>	<a href="#">Atp6v1b2.aSep08</a>	<a href="#">117596</a>	24093	2767	14	536	vacuolar H+ATPase B2 (Atp6v1b2) alternative variant aSep08, mRNA.
<a href="#">Atp6v1b2</a>	<a href="#">Atp6v1b2.bSep08</a>	<a href="#">117596</a>	1566	683	5	165	vacuolar H+ATPase B2 (Atp6v1b2) alternative variant bSep08, mRNA.
<a href="#">Atp6v1b2</a>	<a href="#">Atp6v1b2.cSep08</a>	<a href="#">117596</a>	2535	698	2	70	vacuolar H+ATPase B2 (Atp6v1b2) alternative variant cSep08, mRNA.
<a href="#">Atp6v1c1</a>	<a href="#">Atp6v1c1.bSep08</a>	<a href="#">299971</a>	3961	1149	3	64	ATPase, H+ transporting, lysosomal V1 subunit C1 (7.5 kD) (Atp6v1c1) alternative variant bSep08, mRNA.
<a href="#">Atp6v1c1</a>	<a href="#">Atp6v1c1.cSep08</a>	<a href="#">299971</a>	1546	261	2	51	ATPase, H+ transporting, lysosomal V1 subunit C1 (Atp6v1c1) alternative variant cSep08, mRNA.
<a href="#">Atp6v1c2</a>	<a href="#">Atp6v1c2.bSep08</a>	<a href="#">362802</a>	14893	2818	7	229	vacuolar H+ ATPase C2 (25.8 kD) (Atp6v1c2) alternative variant bSep08, mRNA.
<a href="#">Atp6v1c2</a>	<a href="#">Atp6v1c2.cSep08</a>	<a href="#">362802</a>	16548	1048	9	176	ATPase H+ (20.7 kD) (Atp6v1c2) alternative variant cSep08, mRNA.
<a href="#">Atp6v1c2</a>	<a href="#">Atp6v1c2.dSep08</a>	<a href="#">362802</a>	3178	657	3	98	H+ ATPase (11.2 kD) (Atp6v1c2) alternative variant dSep08, mRNA.
<a href="#">Atp6v1d</a>	<a href="#">Atp6v1d.bSep08</a>	<a href="#">299159</a>	1970	513	2	78	ATPase, H+ transporting, lysosomal V1 subunit D (9.4 kD) (Atp6v1d) alternative variant bSep08, mRNA.

<a href="#">Atp6v1d</a>	<a href="#">Atp6v1d.cSep08</a>	<a href="#">299159</a>	11526	1494	6	81	ATPase, H+ transporting, lysosomal V1 subunit D (Atp6v1d) alternative variant cSep08, mRNA.
<a href="#">Atp6v1e1</a>	<a href="#">Atp6v1e1.aSep08</a>	<a href="#">297566</a>	19206	725	1	241	ATPase, H+ transporting, lysosomal V1 subunit E1 (Atp6v1e1) alternative variant aSep08, mRNA.
<a href="#">Atp6v1g1</a>	<a href="#">Atp6v1g1.cSep08</a>	<a href="#">298103</a>	5631	505	3	47	ATPase, H transporting, lysosomal V1 subunit G1 (Atp6v1g1) alternative variant cSep08, mRNA.
<a href="#">Atp6v1g2</a>	<a href="#">Atp6v1g2.bSep08</a>	<a href="#">368044</a>	2229	1946	2	91	ATPase, H+ transporting, V1 subunit G isoform 2 (10.3 kD) (Atp6v1g2) alternative variant bSep08, complete mRNA.
<a href="#">Atp6v1g2</a>	<a href="#">Atp6v1g2.cSep08</a>	<a href="#">368044</a>	1384	722	3	77	ATPase, H+ transporting, V1 subunit G isoform 2 (9.0 kD) (Atp6v1g2) alternative variant cSep08, mRNA.
<a href="#">Atp6v1g3</a>	<a href="#">Atp6v1g3.bSep08</a>	<a href="#">289407</a>	3368	342	2	69	ATPase, H+ transporting, lysosomal V1 subunit G3 (Atp6v1g3) alternative variant bSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.aSep08</a>	<a href="#">297797</a>	94183	1952	14	483	ATPase H+ transporting lysosomal V1 (55.9 kD) (Atp6v1h) alternative variant aSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.aSep08</a>	<a href="#">681270</a>	94183	1952	14	483	ATPase H+ transporting lysosomal V1 (55.9 kD) (Atp6v1h) alternative variant aSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.cSep08</a>	<a href="#">297797</a>	50545	984	8	272	ATPase H+ transporting lysosomal V1 (Atp6v1h) alternative variant cSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.cSep08</a>	<a href="#">681270</a>	50545	984	8	272	ATPase H+ transporting lysosomal V1 (Atp6v1h) alternative variant cSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.dSep08</a>	<a href="#">297797</a>	32075	614	6	159	ATPase H+ transporting lysosomal V1 (Atp6v1h) alternative variant dSep08, mRNA.
<a href="#">Atp6v1h</a>	<a href="#">Atp6v1h.dSep08</a>	<a href="#">681270</a>	32075	614	6	159	ATPase H+ transporting lysosomal V1 (Atp6v1h) alternative variant dSep08, mRNA.
<a href="#">Atp7b</a>	<a href="#">Atp7b.bSep08</a>	<a href="#">24218</a>	2727	2127	2	122	ATPase, Cu++ transporting, beta polypeptide (13.5 kD) (Atp7b) alternative variant bSep08, mRNA.
<a href="#">Atp8a1</a>	<a href="#">Atp8a1.aSep08</a>	<a href="#">289615</a>	50113	5694		417	ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1 (Atp8a1) mRNA.
<a href="#">Atp8b1</a>	<a href="#">Atp8b1.bSep08</a>	<a href="#">291555</a>	21948	3450	2	400	ATPase, Class I, type 8B, member 1 (Atp8b1) alternative variant bSep08, mRNA.
<a href="#">Atp8b2</a>	<a href="#">Atp8b2.bSep08</a>	<a href="#">361984</a>	34747	1552	1	467	atpase, class I, type 8B, member 2 (Atp8b2) alternative variant bSep08, mRNA.
<a href="#">Atp8b3</a>	<a href="#">Atp8b3.aSep08</a>	<a href="#">299616</a>	12331	2342	13	725	ATPase, Class I, type 8B, member 3 (Atp8b3) alternative variant aSep08, mRNA.
<a href="#">Atp8b3</a>	<a href="#">Atp8b3.bSep08</a>	<a href="#">299616</a>	2472	702	1	233	ATPase, Class I, type 8B, member 3 (Atp8b3) alternative variant bSep08, mRNA.
<a href="#">Atp8b4</a>	<a href="#">Atp8b4.aSep08</a>	<a href="#">311396</a>	21081	1606		258	ATPase, class I, type 8B, member 4 (29.3 kD) (Atp8b4) complete mRNA.
<a href="#">Atp9b</a>	<a href="#">Atp9b.bSep08</a>	<a href="#">291411</a>	11740	915	4	149	ATPase, class II, type 9B (Atp9b) alternative variant bSep08, mRNA.
<a href="#">Atp9b</a>	<a href="#">Atp9b.cSep08</a>	<a href="#">291411</a>	73156	403	6	113	ATPase, class II, type 9B (Atp9b) alternative variant cSep08, mRNA.
<a href="#">Atp9b</a>	<a href="#">Atp9b.dSep08</a>	<a href="#">291411</a>	54096	533	6	84	ATPase, class II, type 9B (Atp9b) alternative variant dSep08, mRNA.
<a href="#">Atp10a</a>	<a href="#">Atp10a.aSep08</a>	<a href="#">365266</a>	19405	1033		344	ATPase, class V, type 10A (Atp10a) mRNA.
<a href="#">Atp10d</a>	<a href="#">Atp10d.aSep08</a>	<a href="#">360932</a>	54281	1430		476	ATPase, class V, type 10D (Atp10d) mRNA.



<a href="#">Atp11a</a>	<a href="#">Atp11a.bSep08</a>	<a href="#">306600</a>	31961	1893	12	564	ATPase, class VI, type 11A (Atp11a) alternative variant bSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.cSep08</a>	<a href="#">306600</a>	15542	1160	8	235	ATPase, class VI, type 11A (Atp11a) alternative variant cSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.dSep08</a>	<a href="#">306600</a>	61906	795	6	234	ATPase, class VI, type 11A (Atp11a) alternative variant dSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.eSep08</a>	<a href="#">306600</a>	61755	521	5	173	ATPase, class VI, type 11A (Atp11a) alternative variant eSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.fSep08</a>	<a href="#">306600</a>	13057	768	6	149	ATPase, class VI, type 11A (Atp11a) alternative variant fSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.gSep08</a>	<a href="#">306600</a>	7548	476	5	123	ATPase, class VI, type 11A (Atp11a) alternative variant gSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.hSep08</a>	<a href="#">306600</a>	3827	389	3	122	ATPase, class VI, type 11A (Atp11a) alternative variant hSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.iSep08</a>	<a href="#">306600</a>	54355	631	4	117	ATPase, class VI, type 11A (Atp11a) alternative variant iSep08, mRNA.
<a href="#">Atp11a</a>	<a href="#">Atp11a.jSep08</a>	<a href="#">306600</a>	1927	336	3	112	ATPase, class VI, type 11A (Atp11a) alternative variant jSep08, mRNA.
<a href="#">Atp11b</a>	<a href="#">Atp11b.aSep08</a>	<a href="#">361929</a>	25872	1764		587	ATPase, class VI, type 11B (Atp11b) mRNA.
<a href="#">Atp11c</a>	<a href="#">Atp11c.aSep08</a>	<a href="#">317599</a>	21705	642	4	191	atpase, class VI, type 11C (Atp11c) alternative variant aSep08, mRNA.
<a href="#">Atp11c</a>	<a href="#">Atp11c.bSep08</a>	<a href="#">317599</a>	14149	466	1	99	atpase, class VI, type 11C (Atp11c) alternative variant bSep08, mRNA.
<a href="#">Atp13a1</a>	<a href="#">Atp13a1.bSep08</a>	<a href="#">290673</a>	16048	3873	24	507	ATPase type 13a1 (Atp13a1) alternative variant bSep08, mRNA.
<a href="#">Atp13a1</a>	<a href="#">Atp13a1.cSep08</a>	<a href="#">290673</a>	3583	709	5	134	ATPase type 13a1 (Atp13a1) alternative variant cSep08, mRNA.
<a href="#">Atp13a1</a>	<a href="#">Atp13a1.dSep08</a>	<a href="#">290673</a>	1376	566	2	86	ATPase type 13a1 (Atp13a1) alternative variant dSep08, mRNA.
<a href="#">Atp13a2</a>	<a href="#">Atp13a2.aSep08</a>	<a href="#">362645</a>	6636	2491	15	749	ATPase type 13A2 (Atp13a2) alternative variant aSep08, mRNA.
<a href="#">Atp13a2</a>	<a href="#">Atp13a2.bSep08</a>	<a href="#">362645</a>	3692	2076	6	328	ATPase type 13A2 (34.8 kD) (Atp13a2) alternative variant bSep08, mRNA.
<a href="#">Atp13a2</a>	<a href="#">Atp13a2.cSep08</a>	<a href="#">362645</a>	2019	687	4	228	ATPase type 13A2 (Atp13a2) alternative variant cSep08, mRNA.
<a href="#">Atp13a2</a>	<a href="#">Atp13a2.dSep08</a>	<a href="#">362645</a>	1541	730	5	200	ATPase type 13A2 (Atp13a2) alternative variant dSep08, mRNA.
<a href="#">Atp13a2</a>	<a href="#">Atp13a2.eSep08</a>	<a href="#">362645</a>	1067	964	1	86	ATPase type 13A2 (Atp13a2) alternative variant eSep08, mRNA.
<a href="#">Atp13a4</a>	<a href="#">Atp13a4.aSep08</a>	<a href="#">288026</a>	11544	836		171	ATPase type 13A4 (Atp13a4) mRNA.
<a href="#">Atp13a5</a>	<a href="#">Atp13a5.aSep08</a>	<a href="#">303856</a>	25564	294		58	ATPase type 13A5 (Atp13a5) mRNA.
<a href="#">Atpaf1</a>	<a href="#">Atpaf1.bSep08</a>	<a href="#">313510</a>	5089	398	3	132	ATP synthase mitochondrial F1 complex assembly factor 1 (Atpaf1) alternative variant bSep08, mRNA.
<a href="#">Atpaf2</a>	<a href="#">Atpaf2.bSep08</a>	<a href="#">303190</a>	781	393	2	86	ATP synthase mitochondrial F1 complex assembly factor 2 (9.6 kD) (Atpaf2) alternative variant bSep08, mRNA.
<a href="#">Atpaf2</a>	<a href="#">Atpaf2.eSep08</a>	<a href="#">303190</a>	4970	692	6	40	ATP synthase mitochondrial F1 complex assembly factor 2 (4.7 kD) (Atpaf2) alternative variant eSep08, mRNA.

<a href="#">Atpbd4</a>	<a href="#">Atpbd4.aSep08</a>	<a href="#">362191</a>	131456	877		129	ATP binding domain 4 (14.6 kD) (Atpbd4) mRNA.
<a href="#">Atrip</a>	<a href="#">Atrip.bSep08</a>	<a href="#">301014</a>	2059	1556	1	98	ATR interacting protein (11.1 kD) (Atrip) alternative variant bSep08, mRNA.
<a href="#">Atrn</a>	<a href="#">Atrn.bSep08</a>	<a href="#">83526</a>	11203	1319	5	163	atractin (Atrn) alternative variant bSep08, mRNA.
<a href="#">Atrn</a>	<a href="#">Atrn.cSep08</a>	<a href="#">83526</a>	65671	630	3	95	atractin (Atrn) alternative variant cSep08, mRNA.
<a href="#">Atrn</a>	<a href="#">Atrn.dSep08</a>	<a href="#">83526</a>	4031	431	2	90	atractin (Atrn) alternative variant dSep08, mRNA.
<a href="#">Atrnl1</a>	<a href="#">Atrnl1.aSep08</a>	<a href="#">307992</a>	413456	2785		206	atractin like 1 (Atrnl1) mRNA.
<a href="#">Atrx</a>	<a href="#">Atrx.aSep08</a>	<a href="#">246284</a>	29585	4389		495	alpha thalassemia/mental retardation syndrome X-linked homolog (human) (Atrx) mRNA.
<a href="#">Atxn2</a>	<a href="#">Atxn2.aSep08</a>	<a href="#">288663</a>	41579	3564	18	918	ataxin 2 (Atxn2) alternative variant aSep08, mRNA.
<a href="#">Atxn2</a>	<a href="#">Atxn2.bSep08</a>	<a href="#">288663</a>	27304	1901	11	511	ataxin 2 (Atxn2) alternative variant bSep08, mRNA.
<a href="#">Atxn2</a>	<a href="#">Atxn2.cSep08</a>	<a href="#">288663</a>	21111	1306	10	434	ataxin 2 (Atxn2) alternative variant cSep08, mRNA.
<a href="#">Atxn2</a>	<a href="#">Atxn2.eSep08</a>	<a href="#">288663</a>	15801	706	7	235	ataxin 2 (Atxn2) alternative variant eSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.aSep08</a>	<a href="#">361649</a>	12283	5171	21	1178	ataxin 2-like CRA f (Atxn2l) alternative variant aSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.cSep08</a>	<a href="#">361649</a>	8285	1867	9	455	ataxin 2-like CRA f (Atxn2l) alternative variant cSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.dSep08</a>	<a href="#">361649</a>	2978	1974	6	425	ataxin 2-like CRA g (Atxn2l) alternative variant dSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.eSep08</a>	<a href="#">361649</a>	1464	762	5	253	ataxin 2-like CRA g (Atxn2l) alternative variant eSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.fSep08</a>	<a href="#">361649</a>	1729	722	5	142	ataxin 2-like CRA e (Atxn2l) alternative variant fSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.gSep08</a>	<a href="#">361649</a>	793	559	2	128	putative protein (14.1 kD) (Atxn2l) alternative variant gSep08, mRNA.
<a href="#">Atxn2l</a>	<a href="#">Atxn2l.hSep08</a>	<a href="#">361649</a>	1194	597	2	61	putative protein (Atxn2l) alternative variant hSep08, mRNA.
<a href="#">Atxn3</a>	<a href="#">Atxn3.bSep08</a>	<a href="#">60331</a>	22170	1038	10	332	ataxin 3 (Atxn3) alternative variant bSep08, mRNA.
<a href="#">Atxn7</a>	<a href="#">Atxn7.aSep08</a>	<a href="#">361015</a>	7906	1368		389	ataxin 7 (Atxn7) mRNA.
<a href="#">Atxn7l1</a>	<a href="#">Atxn7l1.aSep08</a>	<a href="#">314033</a>	4433	774	1	257	ataxin 7-like 1 (Atxn7l1) alternative variant aSep08, mRNA.
<a href="#">Atxn7l1</a>	<a href="#">Atxn7l1.bSep08</a>	<a href="#">314033</a>	4692	727		241	ataxin 7-like 1 (Atxn7l1) alternative variant bSep08, mRNA.
<a href="#">Atxn7l2</a>	<a href="#">Atxn7l2.aSep08</a>	<a href="#">310781</a>	8258	2358	11	719	ataxin 7-like 2 (Atxn7l2) alternative variant aSep08, mRNA.
<a href="#">Atxn7l2</a>	<a href="#">Atxn7l2.cSep08</a>	<a href="#">310781</a>	1664	424	2	77	ataxin 7-like 2 (Atxn7l2) alternative variant cSep08, mRNA.
<a href="#">Atxn10</a>	<a href="#">Atxn10.bSep08</a>	<a href="#">170821</a>	28506	877	8	292	ataxin 10 (Atxn10) alternative variant bSep08, mRNA.
<a href="#">Atxn10</a>	<a href="#">Atxn10.cSep08</a>	<a href="#">170821</a>	39799	866	6	288	ataxin 10 (Atxn10) alternative variant cSep08, mRNA.
<a href="#">Atxn10</a>	<a href="#">Atxn10.dSep08</a>	<a href="#">170821</a>	39306	861	5	215	ataxin 10 (Atxn10) alternative variant dSep08, mRNA.
<a href="#">Atxn10</a>	<a href="#">Atxn10.eSep08</a>	<a href="#">170821</a>	535	338	2	44	ataxin 10 (Atxn10) alternative variant eSep08, mRNA.
<a href="#">AT_hook.0</a>	<a href="#">AT_hook.0.aSep08</a>		2094	1494		497	proline rich 12 (AT_hook.0) mRNA.
<a href="#">Auh</a>	<a href="#">Auh.aSep08</a>	<a href="#">361215</a>	109838	1196	9	315	AU RNA binding protein/enoyl-coenzyme A hydratase (33.3 kD) (Auh) alternative variant aSep08, complete mRNA.
<a href="#">Auh</a>	<a href="#">Auh.cSep08</a>	<a href="#">361215</a>	97985	1259	9	238	AU RNA binding protein/enoyl-coenzyme A hydratase (Auh) alternative variant cSep08, mRNA.
<a href="#">Auh</a>	<a href="#">Auh.dSep08</a>	<a href="#">361215</a>	109594	559	4	181	AU RNA binding protein/enoyl-coenzyme A hydratase (Auh) alternative variant dSep08, mRNA.

<a href="#">Auh</a>	<a href="#">Auh.eSep08</a>	<a href="#">361215</a>	62214	457	2	39	AU RNA binding protein/enoyl-coenzyme A hydratase (Auh) alternative variant eSep08, mRNA.
<a href="#">Aup1</a>	<a href="#">Aup1.bSep08</a>	<a href="#">680423</a>	1695	662	7	210	ancient ubiquitous protein 1 (Aup1) alternative variant bSep08, mRNA.
<a href="#">Aup1</a>	<a href="#">Aup1.cSep08</a>	<a href="#">680423</a>	1257	840	4	127	ancient ubiquitous protein 1 (Aup1) alternative variant cSep08, mRNA.
<a href="#">Aup1</a>	<a href="#">Aup1.dSep08</a>	<a href="#">680423</a>	3079	2644	4	114	ancient ubiquitous protein 1 CRA c (13.0 kD) (Aup1) alternative variant dSep08, mRNA.
<a href="#">Aup1</a>	<a href="#">Aup1.eSep08</a>	<a href="#">680423</a>	1066	630	5	102	ancient ubiquitous protein 1 CRA c (Aup1) alternative variant eSep08, mRNA.
<a href="#">Aup1</a>	<a href="#">Aup1.fSep08</a>	<a href="#">680423</a>	682	586	2	92	ancient ubiquitous protein 1 CRA c precursor (10.2 kD) (Aup1) alternative variant fSep08, mRNA.
<a href="#">Aurka</a>	<a href="#">Aurka.bSep08</a>	<a href="#">261730</a>	12510	1217	7	276	aurora kinase A (Aurka) alternative variant bSep08, mRNA.
<a href="#">Aurka</a>	<a href="#">Aurka.cSep08</a>	<a href="#">261730</a>	816	728	2	55	aurora kinase A (Aurka) alternative variant cSep08, mRNA.
<a href="#">Aurkaip1</a>	<a href="#">Aurkaip1.bSep08</a>	<a href="#">298687</a>	1079	496	4	133	aurora kinase A interacting protein 1 (Aurkaip1) alternative variant bSep08, mRNA.
<a href="#">Aurkaip1</a>	<a href="#">Aurkaip1.cSep08</a>	<a href="#">298687</a>	1073	806	2	49	aurora kinase A interacting protein 1 (Aurkaip1) alternative variant cSep08, mRNA.
<a href="#">Aurkb</a>	<a href="#">Aurkb.bSep08</a>	<a href="#">114592</a>	3742	628	1	209	aurora kinase B (Aurkb) alternative variant bSep08, mRNA.
<a href="#">Auts2l</a>	<a href="#">Auts2l.aSep08</a>	<a href="#">304419</a>	162455	856		86	autism susceptibility candidate 2-like (Auts2l) mRNA.
<a href="#">Aven</a>	<a href="#">Aven.aSep08</a>	<a href="#">311299</a>	134530	1185	3	282	apoptosis, caspase activation inhibitor (Aven) alternative variant aSep08, mRNA.
<a href="#">Aven</a>	<a href="#">Aven.cSep08</a>	<a href="#">311299</a>	105160	463	2	38	apoptosis, caspase activation inhibitor (4.5 kD) (Aven) alternative variant cSep08, mRNA.
<a href="#">Avpi1</a>	<a href="#">Avpi1.cSep08</a>	<a href="#">171386</a>	754	365	2	11	arginine vasopressin-induced 1 (1.2 kD) (Avpi1) alternative variant cSep08, mRNA.
<a href="#">Axin2</a>	<a href="#">Axin2.bSep08</a>	<a href="#">29134</a>	8353	1784	1	426	axin2 (Axin2) alternative variant bSep08, mRNA.
<a href="#">Axin2</a>	<a href="#">Axin2.cSep08</a>	<a href="#">29134</a>	2835	403	2	39	axin2 (Axin2) alternative variant cSep08, mRNA.
<a href="#">Axl</a>	<a href="#">Axl.aSep08</a>	<a href="#">308444</a>	28350	3574	17	702	axl receptor tyrosine kinase (Axl) alternative variant aSep08, mRNA.
<a href="#">Axl</a>	<a href="#">Axl.bSep08</a>	<a href="#">308444</a>	7846	712	5	237	axl receptor tyrosine kinase (Axl) alternative variant bSep08, mRNA.
<a href="#">Axl</a>	<a href="#">Axl.dSep08</a>	<a href="#">308444</a>	1139	581	2	70	axl receptor tyrosine kinase (Axl) alternative variant dSep08, mRNA.
<a href="#">Azgp1</a>	<a href="#">Azgp1.bSep08</a>	<a href="#">25294</a>	3529	639	2	210	alpha-2-glycoprotein 1, zinc (Azgp1) alternative variant bSep08, mRNA.
<a href="#">Azi1</a>	<a href="#">Azi1.bSep08</a>	<a href="#">360672</a>	954	724	3	127	5-azacytidine induced gene 1 (Azi1) alternative variant bSep08, mRNA.
<a href="#">Azi1</a>	<a href="#">Azi1.cSep08</a>	<a href="#">360672</a>	533	461	2	94	5-azacytidine induced gene 1 (9.5 kD) (Azi1) alternative variant cSep08, mRNA.
<a href="#">Azi2</a>	<a href="#">Azi2.bSep08</a>	<a href="#">316051</a>	15900	1379	7	237	5-azacytidine induced gene 2 (27.4 kD) (Azi2) alternative variant bSep08, mRNA.
<a href="#">Azi2</a>	<a href="#">Azi2.cSep08</a>	<a href="#">316051</a>	1747	411	3	130	5-azacytidine induced gene 2 (Azi2) alternative variant cSep08, mRNA.
<a href="#">Azin1</a>	<a href="#">Azin1.bSep08</a>	<a href="#">58961</a>	28105	1889	12	437	antizyme inhibitor 1 (Azin1) alternative variant bSep08, mRNA.

<a href="#">Azin1</a>	<a href="#">Azin1.cSep08</a>	<a href="#">58961</a>	26327	1693	11	324	antizyme inhibitor 1 (Azin1) alternative variant cSep08, mRNA.
<a href="#">B3galnt1</a>	<a href="#">B3galnt1.aSep08</a>	<a href="#">310508</a>	29989	832	2	128	UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide 1 (13.5 kD) (B3galnt1) alternative variant aSep08, mRNA.
<a href="#">B3galnt1</a>	<a href="#">B3galnt1.cSep08</a>	<a href="#">310508</a>	29446	713	2	94	UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide 1 (B3galnt1) alternative variant cSep08, mRNA.
<a href="#">B3galtl</a>	<a href="#">B3galtl.aSep08</a>	<a href="#">689765</a>	82689	2299		670	beta 1,3-galactosyltransferase-like (B3galtl) alternative variant aSep08, mRNA.
<a href="#">B3galtl</a>	<a href="#">B3galtl.bSep08</a>	<a href="#">689765</a>	7342	1816		83	beta 1,3-galactosyltransferase-like (B3galtl) alternative variant bSep08, mRNA.
<a href="#">B3gat1</a>	<a href="#">B3gat1.cSep08</a>	<a href="#">117108</a>	1848	406	2	58	beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P) (6.5 kD) (B3gat1) alternative variant cSep08, mRNA.
<a href="#">B3gnt2</a>	<a href="#">B3gnt2.bSep08</a>	<a href="#">305571</a>	22311	393	2	41	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 2 (B3gnt2) alternative variant bSep08, mRNA.
<a href="#">B3gnt1</a>	<a href="#">B3gnt1.aSep08</a>	<a href="#">367384</a>	60232	2905	2	653	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase-like 1 (B3gnt1) alternative variant aSep08, mRNA.
<a href="#">B4galnt1</a>	<a href="#">B4galnt1.bSep08</a>	<a href="#">64828</a>	1448	676	4	207	beta-1,4-N-acetyl-galactosaminyl transferase 1 (B4galnt1) alternative variant bSep08, mRNA.
<a href="#">B4galnt1</a>	<a href="#">B4galnt1.dSep08</a>	<a href="#">64828</a>	584	397	2	96	beta-1,4-N-acetyl-galactosaminyl transferase 1 (B4galnt1) alternative variant dSep08, mRNA.
<a href="#">B4galnt4</a>	<a href="#">B4galnt4.bSep08</a>	<a href="#">309105</a>	3576	1715	4	479	beta-1 4-N-acetyl-galactosaminyl transferase 4 (B4galnt4) alternative variant bSep08, complete mRNA.
<a href="#">B4galnt4</a>	<a href="#">B4galnt4.cSep08</a>	<a href="#">309105</a>	1455	943	6	198	beta-1 4-N-acetyl-galactosaminyl transferase 4 CRA b (B4galnt4) alternative variant cSep08, mRNA.
<a href="#">B4galt1</a>	<a href="#">B4galt1.bSep08</a>	<a href="#">24390</a>	42841	476	3	98	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1 (B4galt1) alternative variant bSep08, mRNA.
<a href="#">B4galt1</a>	<a href="#">B4galt1.cSep08</a>	<a href="#">24390</a>	2845	829	2	44	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1 (B4galt1) alternative variant cSep08, mRNA.
<a href="#">B4galt2</a>	<a href="#">B4galt2.bSep08</a>	<a href="#">313536</a>	6060	1056	6	346	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2 (B4galt2) alternative variant bSep08, mRNA.
<a href="#">B4galt3</a>	<a href="#">B4galt3.bSep08</a>	<a href="#">494342</a>	5497	1262	7	198	beta 1 4- galactosyltransferase 3 (B4galt3) alternative variant bSep08, mRNA.
<a href="#">B4galt3</a>	<a href="#">B4galt3.cSep08</a>	<a href="#">494342</a>	3108	2593	2	70	putative protein (7.4 kD) (B4galt3) alternative variant cSep08, mRNA.
<a href="#">B4galt5</a>	<a href="#">B4galt5.bSep08</a>	<a href="#">362275</a>	1917	361	3	120	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5 (B4galt5) alternative variant bSep08, mRNA.
<a href="#">B4galt7</a>	<a href="#">B4galt7.bSep08</a>	<a href="#">364675</a>	1060	405	1	123	xylosylprotein beta1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I) (B4galt7) alternative variant bSep08, mRNA.
<a href="#">B9d1</a>	<a href="#">B9d1.bSep08</a>	<a href="#">287383</a>	7557	902	7	136	protein-like precursor (15.2 kD) (B9d1) alternative variant bSep08, complete mRNA.
<a href="#">B9d1</a>	<a href="#">B9d1.cSep08</a>	<a href="#">287383</a>	3461	492	3	118	b9 protein domain 1 (B9d1) alternative variant cSep08, mRNA.

<a href="#">B9d1</a>	<a href="#">B9d1.dSep08</a>	<a href="#">287383</a>	3060	412	2	67	putative protein (B9d1) alternative variant dSep08, mRNA.
<a href="#">B9d2andTgfb1</a>	<a href="#">B9d2andTgfb1.bSep08</a>	<a href="#">59086</a>	5511	568	4	158	b9 protein domain 2 (B9d2andTgfb1) alternative variant bSep08, mRNA.
<a href="#">B9d2andTgfb1</a>	<a href="#">B9d2andTgfb1.bSep08</a>	<a href="#">308443</a>	5511	568	4	158	b9 protein domain 2 (B9d2andTgfb1) alternative variant bSep08, mRNA.
<a href="#">B12-binding.0</a>	<a href="#">B12-binding.0.aSep08</a>	<a href="#">301276</a>	14007	2187		288	methylmalonyl-CoA mutase (B12-binding.0) mRNA.
<a href="#">Baalc</a>	<a href="#">Baalc.bSep08</a>	<a href="#">140720</a>	9142	622	2	56	brain and acute leukemia, cytoplasmic (Baalc) alternative variant bSep08, mRNA.
<a href="#">baby</a>	<a href="#">baby.aSep08</a>		4325	625		208	odd Oz ten-m homolog (baby) mRNA.
<a href="#">Bace1</a>	<a href="#">Bace1.bSep08</a>	<a href="#">29392</a>	2622	821	3	108	beta-site APP cleaving enzyme 1 (Bace1) alternative variant bSep08, mRNA.
<a href="#">Bace2</a>	<a href="#">Bace2.bSep08</a>	<a href="#">288227</a>	1249	742	1	132	beta-site APP-cleaving enzyme 2 (Bace2) alternative variant bSep08, mRNA.
<a href="#">Bach1</a>	<a href="#">Bach1.bSep08</a>	<a href="#">304127</a>	17815	599	1	108	BTB and CNC homology 1 (12.1 kD) (Bach1) alternative variant bSep08, complete mRNA.
<a href="#">bachu</a>	<a href="#">bachu.aSep08</a>		1228	420		89	regulator of telomere elongation helicase 1 (bachu) mRNA.
<a href="#">bachy</a>	<a href="#">bachy.aSep08</a>		13362	442	3	42	putative protein (bachy) alternative variant aSep08, mRNA.
<a href="#">BACK.0</a>	<a href="#">BACK.0.aSep08</a>		34362	2612		509	kelch-like 13 CRA b (BACK.0) mRNA.
<a href="#">BACK.1</a>	<a href="#">BACK.1.aSep08</a>		22252	490	3	144	kelch-like 3 (BACK.1) alternative variant aSep08, mRNA.
<a href="#">BACK.2</a>	<a href="#">BACK.2.aSep08</a>		1778	741		247	kelch-like 30 (BACK.2) mRNA.
<a href="#">BACK.3</a>	<a href="#">BACK.3.aSep08</a>		17548	1800	4	463	kelch-like 18 (BACK.3) alternative variant aSep08, mRNA.
<a href="#">BACK.3</a>	<a href="#">BACK.3.bSep08</a>		1914	430	1	70	kelch-like 18 CRA c (BACK.3) alternative variant bSep08, mRNA.
<a href="#">Bad</a>	<a href="#">Bad.bSep08</a>	<a href="#">64639</a>	2327	1891	2	114	bcl2-antagonist of cell death (12.0 kD) (Bad) alternative variant bSep08, mRNA.
<a href="#">Bad</a>	<a href="#">Bad.cSep08</a>	<a href="#">64639</a>	1135	359	2	82	bcl2-antagonist of cell death (Bad) alternative variant cSep08, mRNA.
<a href="#">Bad</a>	<a href="#">Bad.dSep08</a>	<a href="#">64639</a>	8634	1047	5	71	bcl2-antagonist of cell death (Bad) alternative variant dSep08, mRNA.
<a href="#">bafer</a>	<a href="#">bafer.aSep08</a>		498	389	2	116	putative protein (bafer) alternative variant aSep08, mRNA.
<a href="#">bafer</a>	<a href="#">bafer.bSep08</a>		12580	593	3	64	CRA b like (7.5 kD) (bafer) alternative variant bSep08, mRNA.
<a href="#">bafer</a>	<a href="#">bafer.cSep08</a>		17168	781	4	64	CRA b like (7.5 kD) (bafer) alternative variant cSep08, mRNA.
<a href="#">bafer</a>	<a href="#">bafer.dSep08</a>		2160	615	2	39	putative protein (3.9 kD) (bafer) alternative variant dSep08, mRNA.
<a href="#">baflo</a>	<a href="#">baflo.aSep08</a>		848	409		80	putative protein (9.5 kD) (baflo) mRNA.
<a href="#">baflu</a>	<a href="#">baflu.aSep08</a>		2551	373		123	ryanodine receptor (baflu) mRNA.
<a href="#">Bag1</a>	<a href="#">Bag1.bSep08</a>	<a href="#">297994</a>	22335	671	6	159	bcl2-associated athanogene 1 (Bag1) alternative variant bSep08, mRNA.
<a href="#">Bag1</a>	<a href="#">Bag1.cSep08</a>	<a href="#">297994</a>	6324	469	4	99	bcl2-associated athanogene 1 (Bag1) alternative variant cSep08, mRNA.
<a href="#">Bag1</a>	<a href="#">Bag1.dSep08</a>	<a href="#">297994</a>	2391	356	2	70	bcl2-associated athanogene 1 (Bag1) alternative variant dSep08, mRNA.

<a href="#">Bag5</a>	<a href="#">Bag5.bSep08</a>	<a href="#">366734</a>	1863	679	2	125	BCL2-associated athanogene 5 (Bag5) alternative variant bSep08, mRNA.
<a href="#">BAH.0</a>	<a href="#">BAH.0.aSep08</a>		3863	1818		152	bah domain coiled-coil containing 1 (BAH.0) alternative variant aSep08, mRNA.
<a href="#">Bahd1</a>	<a href="#">Bahd1.aSep08</a>	<a href="#">362194</a>	7118	2872		308	bromo adjacent region (Bahd1) mRNA.
<a href="#">Bai1</a>	<a href="#">Bai1.aSep08</a>	<a href="#">362931</a>	6936	2413	2	381	brain-specific angiogenesis inhibitor 1 (41.6 kD) (Bai1) alternative variant aSep08, mRNA.
<a href="#">Bai1</a>	<a href="#">Bai1.bSep08</a>	<a href="#">362931</a>	1495	686	1	228	brain-specific angiogenesis inhibitor 1 (Bai1) alternative variant bSep08, mRNA.
<a href="#">Bai2</a>	<a href="#">Bai2.bSep08</a>	<a href="#">313058</a>	1452	607	1	103	brain-specific angiogenesis inhibitor 2 (Bai2) alternative variant bSep08, mRNA.
<a href="#">Baiap2</a>	<a href="#">Baiap2.aSep08</a>	<a href="#">117542</a>	58907	1358	8	420	brain-specific angiogenesis inhibitor 1-associated protein 2 (Baiap2) alternative variant aSep08, mRNA.
<a href="#">Baiap2</a>	<a href="#">Baiap2.bSep08</a>	<a href="#">117542</a>	57798	771	6	214	brain-specific angiogenesis inhibitor 1-associated protein 2 (24.5 kD) (Baiap2) alternative variant bSep08, mRNA.
<a href="#">Baiap2</a>	<a href="#">Baiap2.cSep08</a>	<a href="#">117542</a>	18885	604	1	128	brain-specific angiogenesis inhibitor 1-associated protein 2 (14.5 kD) (Baiap2) alternative variant cSep08, mRNA.
<a href="#">Baiap2</a>	<a href="#">Baiap2.eSep08</a>	<a href="#">117542</a>	41373	626	5	81	brain-specific angiogenesis inhibitor 1-associated protein 2 (9.1 kD) (Baiap2) alternative variant eSep08, mRNA.
<a href="#">Baiap2l1</a>	<a href="#">Baiap2l1.bSep08</a>	<a href="#">304282</a>	71379	1231	3	409	BAI1-associated protein 2-like 1 (Baiap2l1) alternative variant bSep08, mRNA.
<a href="#">Baiap2l1</a>	<a href="#">Baiap2l1.cSep08</a>	<a href="#">304282</a>	12905	779	1	199	BAI1-associated protein 2-like 1 (Baiap2l1) alternative variant cSep08, mRNA.
<a href="#">Baiap2l2</a>	<a href="#">Baiap2l2.aSep08</a>	<a href="#">685357</a>	14569	617		205	BAI1-associated protein 2-like 2 (Baiap2l2) mRNA.
<a href="#">Bak1</a>	<a href="#">Bak1.aSep08</a>	<a href="#">116502</a>	8646	1941		209	BCL2-antagonist/killer 1 (23.2 kD) (Bak1) mRNA.
<a href="#">baloy</a>	<a href="#">baloy.aSep08</a>		2014	345		28	putative protein (3.2 kD) (baloy) mRNA.
<a href="#">Bambi</a>	<a href="#">Bambi.bSep08</a>	<a href="#">83837</a>	1385	953	2	107	BMP and activin membrane-bound inhibitor, homolog (Xenopus laevis) (Bambi) alternative variant bSep08, mRNA.
<a href="#">bamer</a>	<a href="#">bamer.aSep08</a>		5401	349		56	putative protein (bamer) mRNA.
<a href="#">Banf2</a>	<a href="#">Banf2.aSep08</a>	<a href="#">296198</a>	13601	451		130	barrier to autointegration factor 2 (Banf2) mRNA.
<a href="#">banoy</a>	<a href="#">banoy.aSep08</a>		7881	223		55	putative protein (banoy) mRNA.
<a href="#">Banp</a>	<a href="#">Banp.aSep08</a>	<a href="#">292064</a>	75112	2361	14	548	btg3 associated nuclear protein (59.7 kD) (Banp) alternative variant aSep08, complete mRNA.
<a href="#">Banp</a>	<a href="#">Banp.bSep08</a>	<a href="#">292064</a>	26487	1138	8	253	btg3 associated nuclear protein (27.3 kD) (Banp) alternative variant bSep08, mRNA.
<a href="#">Banp</a>	<a href="#">Banp.cSep08</a>	<a href="#">292064</a>	43899	774	6	212	btg3 associated nuclear protein (Banp) alternative variant cSep08, mRNA.
<a href="#">Banp</a>	<a href="#">Banp.dSep08</a>	<a href="#">292064</a>	2527	988	2	62	btg3 associated nuclear protein (7.0 kD) (Banp) alternative variant dSep08, mRNA.
<a href="#">Bap1</a>	<a href="#">Bap1.bSep08</a>	<a href="#">306257</a>	1132	774	3	257	brca1 associated (Bap1) alternative variant bSep08, mRNA.
<a href="#">Bap1</a>	<a href="#">Bap1.cSep08</a>	<a href="#">306257</a>	899	785	2	210	brca1 associated (Bap1) alternative variant cSep08, mRNA.
<a href="#">Bap1</a>	<a href="#">Bap1.dSep08</a>	<a href="#">306257</a>	3550	1299	7	149	brca1 associated (Bap1) alternative variant dSep08, mRNA.

<a href="#">Bap1</a>	<a href="#">Bap1.eSep08</a>	<a href="#">306257</a>	1359	733	3	105	brca1 associated (Bap1) alternative variant eSep08, mRNA.
<a href="#">Bap1</a>	<a href="#">Bap1.fSep08</a>	<a href="#">306257</a>	593	404	2	67	putative protein (Bap1) alternative variant fSep08, mRNA.
<a href="#">bapor</a>	<a href="#">bapor.aSep08</a>		3612	306		13	putative protein (1.5 kD) (bapor) mRNA.
<a href="#">barby</a>	<a href="#">barby.aSep08</a>		10017	552		184	THO complex 2 (barby) mRNA.
<a href="#">barchy</a>	<a href="#">barchy.aSep08</a>		16332	573	3	91	putative protein (barchy) mRNA.
<a href="#">Bard1</a>	<a href="#">Bard1.bSep08</a>	<a href="#">64557</a>	8933	416	2	138	BRCA1 associated RING domain 1 (Bard1) alternative variant bSep08, mRNA.
<a href="#">barfer</a>	<a href="#">barfer.aSep08</a>		22321	407		51	putative protein (5.4 kD) (barfer) mRNA.
<a href="#">barflo</a>	<a href="#">barflo.aSep08</a>		3749	455		151	CRA b (barflo) mRNA.
<a href="#">barflu</a>	<a href="#">barflu.aSep08</a>		9069	414		137	CRA b like (barflu) mRNA.
<a href="#">barloy</a>	<a href="#">barloy.aSep08</a>		24805	735	2	37	putative protein (4.3 kD) (barloy) alternative variant aSep08, mRNA.
<a href="#">barloy</a>	<a href="#">barloy.bSep08</a>		21054	564	1	76	putative cytoplasmic protein (8.6 kD) (barloy) alternative variant bSep08, mRNA.
<a href="#">barmer</a>	<a href="#">barmer.aSep08</a>		753	506		35	putative protein (barmer) mRNA.
<a href="#">barnoy</a>	<a href="#">barnoy.aSep08</a>		1430	1337		91	putative cytoplasmic protein (10.1 kD) (barnoy) mRNA.
<a href="#">barpor</a>	<a href="#">barpor.aSep08</a>		2172	382		19	putative protein (barpor) mRNA.
<a href="#">barsa</a>	<a href="#">barsa.aSep08</a>		1367	400	2	133	putative protein (barsa) alternative variant aSep08, mRNA.
<a href="#">barshee</a>	<a href="#">barshee.aSep08</a>		2780	646		86	putative protein (9.6 kD) (barshee) mRNA.
<a href="#">barto</a>	<a href="#">barto.aSep08</a>		5845	797	4	45	putative protein (5.3 kD) (barto) alternative variant aSep08, mRNA.
<a href="#">barto</a>	<a href="#">barto.bSep08</a>		1203	399	2	43	putative protein (barto) alternative variant bSep08, mRNA.
<a href="#">barvar</a>	<a href="#">barvar.aSep08</a>		1499	767		99	putative protein (11.2 kD) (barvar) mRNA.
<a href="#">barwey</a>	<a href="#">barwey.aSep08</a>		4444	346		67	CRA c like (barwey) mRNA.
<a href="#">basa</a>	<a href="#">basa.aSep08</a>		928	653		82	putative protein (9.2 kD) (basa) mRNA.
<a href="#">bashee</a>	<a href="#">bashee.aSep08</a>		3085	453		150	putative protein of vertebrate origin (bashee) mRNA.
<a href="#">Bat1a</a>	<a href="#">Bat1a.bSep08</a>	<a href="#">114612</a>	3702	2336	4	254	HLA-B associated transcript 1 (28.9 kD) (Bat1a) alternative variant bSep08, mRNA.
<a href="#">Bat1a</a>	<a href="#">Bat1a.cSep08</a>	<a href="#">114612</a>	2802	613	4	167	CRA c like (Bat1a) alternative variant cSep08, mRNA.
<a href="#">Bat2</a>	<a href="#">Bat2.aSep08</a>	<a href="#">294250</a>	7221	3577	13	1128	HLA-B associated transcript 2 (Bat2) alternative variant aSep08, mRNA.
<a href="#">Bat2</a>	<a href="#">Bat2.bSep08</a>	<a href="#">294250</a>	4914	2233	9	487	HLA-B associated transcript 2 (50.6 kD) (Bat2) alternative variant bSep08, mRNA.
<a href="#">Bat2</a>	<a href="#">Bat2.cSep08</a>	<a href="#">294250</a>	1161	777	2	182	HLA-B associated transcript 2 (Bat2) alternative variant cSep08, mRNA.
<a href="#">Bat2</a>	<a href="#">Bat2.dSep08</a>	<a href="#">294250</a>	1286	509	2	169	HLA-B associated transcript 2 (Bat2) alternative variant dSep08, mRNA.
<a href="#">Bat2d</a>	<a href="#">Bat2d.aSep08</a>	<a href="#">360865</a>	28701	1335	9	366	BAT2, N-terminal (Bat2d) alternative variant aSep08, mRNA.
<a href="#">Bat2d</a>	<a href="#">Bat2d.bSep08</a>	<a href="#">360865</a>	3212	395	2	121	BAT2, N-terminal (Bat2d) alternative variant bSep08, mRNA.
<a href="#">Bat2d</a>	<a href="#">Bat2d.cSep08</a>	<a href="#">360865</a>	19951	540	2	83	putative protein (Bat2d) alternative variant cSep08, mRNA.
<a href="#">BAT2_N.0</a>	<a href="#">BAT2_N.0.aSep08</a>		3139	593		143	hla-b associated transcript 2 (BAT2_N.0) mRNA.

<a href="#">Bat3</a>	<a href="#">Bat3.cSep08</a>	<a href="#">94342</a>	8569	2087	13	587	HLA-B-associated transcript 3 (Bat3) alternative variant cSep08, mRNA.
<a href="#">Bat3</a>	<a href="#">Bat3.dSep08</a>	<a href="#">94342</a>	2034	915	7	304	HLA-B-associated transcript 3 (Bat3) alternative variant dSep08, mRNA.
<a href="#">Bat3</a>	<a href="#">Bat3.eSep08</a>	<a href="#">94342</a>	4125	697	6	221	HLA-B-associated transcript 3 (Bat3) alternative variant eSep08, mRNA.
<a href="#">Bat3</a>	<a href="#">Bat3.fSep08</a>	<a href="#">94342</a>	1686	698	6	208	HLA-B-associated transcript 3 (Bat3) alternative variant fSep08, mRNA.
<a href="#">Bat3</a>	<a href="#">Bat3.gSep08</a>	<a href="#">94342</a>	981	772	3	144	HLA-B-associated transcript 3 (Bat3) alternative variant gSep08, mRNA.
<a href="#">Bat4</a>	<a href="#">Bat4.bSep08</a>	<a href="#">415064</a>	1879	797	1	202	bat4 gene (Bat4) alternative variant bSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.bSep08</a>	<a href="#">361796</a>	4155	1192	12	193	hla-b associated transcript 5 (22.4 kD) (Bat5) alternative variant bSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.cSep08</a>	<a href="#">361796</a>	7242	403	5	133	hla-b associated transcript 5 CRA d (Bat5) alternative variant cSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.dSep08</a>	<a href="#">361796</a>	6184	758	9	129	hla-b associated transcript 5 CRA c (Bat5) alternative variant dSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.eSep08</a>	<a href="#">361796</a>	2439	1360	9	117	hla-b associated transcript 5 CRA d (13.8 kD) (Bat5) alternative variant eSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.fSep08</a>	<a href="#">361796</a>	2926	771	6	116	hla-b associated transcript 5 CRA c (Bat5) alternative variant fSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.gSep08</a>	<a href="#">361796</a>	1610	738	2	106	hla-b associated transcript 5 CRA f (11.9 kD) (Bat5) alternative variant gSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.hSep08</a>	<a href="#">361796</a>	5435	327	5	99	hla-b associated transcript 5 CRA a (Bat5) alternative variant hSep08, mRNA.
<a href="#">Bat5</a>	<a href="#">Bat5.jSep08</a>	<a href="#">361796</a>	5612	406	4	65	hla-b associated transcript 5 CRA f (Bat5) alternative variant jSep08, mRNA.
<a href="#">bato</a>	<a href="#">bato.aSep08</a>		39466	1802		63	putative protein (bato) mRNA.
<a href="#">bavar</a>	<a href="#">bavar.aSep08</a>		63607	790	3	92	putative nuclear protein (10.1 kD) (bavar) alternative variant aSep08, mRNA.
<a href="#">bavar</a>	<a href="#">bavar.bSep08</a>		837	299	1	46	putative protein (5.2 kD) (bavar) alternative variant bSep08, mRNA.
<a href="#">bawby</a>	<a href="#">bawby.aSep08</a>		16694	557		185	THO complex 2 (bawby) mRNA.
<a href="#">bawchy</a>	<a href="#">bawchy.aSep08</a>		2014	718		64	putative protein (bawchy) mRNA.
<a href="#">bawey</a>	<a href="#">bawey.aSep08</a>		8587	643		86	putative protein (bawey) mRNA.
<a href="#">bawfer</a>	<a href="#">bawfer.aSep08</a>		784	462		28	putative protein (3.0 kD) (bawfer) mRNA.
<a href="#">bawflo</a>	<a href="#">bawflo.aSep08</a>		89552	345		36	putative protein (bawflo) mRNA.
<a href="#">bawflu</a>	<a href="#">bawflu.aSep08</a>		837	526	2	125	HAI-2 related small like (13.0 kD) (bawflu) alternative variant aSep08, mRNA.
<a href="#">bawflu</a>	<a href="#">bawflu.bSep08</a>		678	518	1	115	CRA b like (12.2 kD) (bawflu) alternative variant bSep08, mRNA.
<a href="#">bawflu</a>	<a href="#">bawflu.cSep08</a>		965	535	3	104	HAI-2 related small (10.7 kD) (bawflu) alternative variant cSep08, mRNA.
<a href="#">bawkee</a>	<a href="#">bawkee.aSep08</a>		6457	325		108	contactin associated protein-like 5A (bawkee) mRNA.
<a href="#">bawloy</a>	<a href="#">bawloy.bSep08</a>		1350	646	2	77	putative protein (8.8 kD) (bawloy) alternative variant bSep08, mRNA.



<a href="#">bawmer</a>	<a href="#">bawmer.aSep08</a>		2031	657		218	CRA b (bawmer) mRNA.
<a href="#">bawnoy</a>	<a href="#">bawnoy.aSep08</a>		566	411		102	putative protein (bawnoy) mRNA.
<a href="#">bawpor</a>	<a href="#">bawpor.aSep08</a>		859	683		47	putative protein (bawpor) mRNA.
<a href="#">bawsa</a>	<a href="#">bawsa.aSep08</a>		721	510		169	methyl-CpG binding domain protein 6 like (bawsa) mRNA.
<a href="#">bawshee</a>	<a href="#">bawshee.aSep08</a>		732	463		13	putative protein (bawshee) mRNA.
<a href="#">bawto</a>	<a href="#">bawto.aSep08</a>		1211	406		22	putative protein (bawto) mRNA.
<a href="#">bawvar</a>	<a href="#">bawvar.aSep08</a>		15187	347		110	putative protein (bawvar) mRNA.
<a href="#">bawwey</a>	<a href="#">bawwey.aSep08</a>		1349	570		36	putative protein (4.1 kD) (bawwey) mRNA.
<a href="#">Bax</a>	<a href="#">Bax.aSep08</a>	<a href="#">24887</a>	5407	832	6	192	bcl2-associated X protein (21.4 kD) (Bax) alternative variant aSep08, mRNA.
<a href="#">Bax</a>	<a href="#">Bax.bSep08</a>	<a href="#">24887</a>	4086	628	5	167	bcl2-associated X protein (18.8 kD) (Bax) alternative variant bSep08, complete mRNA.
<a href="#">Bax</a>	<a href="#">Bax.cSep08</a>	<a href="#">24887</a>	1636	352	4	117	bcl2-associated X protein (Bax) alternative variant cSep08, mRNA.
<a href="#">Bax</a>	<a href="#">Bax.dSep08</a>	<a href="#">24887</a>	1705	925	2	70	bcl2-associated X protein (Bax) alternative variant dSep08, mRNA.
<a href="#">Bax</a>	<a href="#">Bax.eSep08</a>	<a href="#">24887</a>	851	758	2	52	bcl2-associated X protein (Bax) alternative variant eSep08, mRNA.
<a href="#">Baz1a</a>	<a href="#">Baz1a.aSep08</a>	<a href="#">314126</a>	5099	1592	4	284	bromodomain adjacent to zinc finger domain, 1A (Baz1a) alternative variant aSep08, mRNA.
<a href="#">Baz1a</a>	<a href="#">Baz1a.bSep08</a>	<a href="#">314126</a>	2023	741	3	56	bromodomain adjacent to zinc finger domain, 1A (6.2 kD) (Baz1a) alternative variant bSep08, mRNA.
<a href="#">Baz1b</a>	<a href="#">Baz1b.aSep08</a>	<a href="#">368002</a>	5907	2612	4	229	bromodomain adjacent to zinc finger domain, 1B (Baz1b) alternative variant aSep08, mRNA.
<a href="#">Baz1b</a>	<a href="#">Baz1b.bSep08</a>	<a href="#">368002</a>	18697	801	5	199	bromodomain adjacent to zinc finger domain, 1B (Baz1b) alternative variant bSep08, mRNA.
<a href="#">Baz2a</a>	<a href="#">Baz2a.bSep08</a>	<a href="#">304601</a>	7248	4823	10	559	bromodomain adjacent to zinc finger domain, 2A (Baz2a) alternative variant bSep08, mRNA.
<a href="#">Baz2a</a>	<a href="#">Baz2a.cSep08</a>	<a href="#">304601</a>	875	667	3	177	bromodomain adjacent to zinc finger domain, 2A (Baz2a) alternative variant cSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.bSep08</a>	<a href="#">317627</a>	13940	2825	9	522	bromodomain adjacent to zinc finger domain, 2B (Baz2b) alternative variant bSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.cSep08</a>	<a href="#">317627</a>	232969	1590	9	372	bromodomain adjacent to zinc finger domain, 2B (Baz2b) alternative variant cSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.dSep08</a>	<a href="#">317627</a>	22983	697	7	232	bromodomain adjacent to zinc finger domain, 2B (Baz2b) alternative variant dSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.eSep08</a>	<a href="#">317627</a>	5058	846	3	161	bromodomain adjacent to zinc finger domain, 2B (Baz2b) alternative variant eSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.fSep08</a>	<a href="#">317627</a>	848	752	2	81	bromodomain adjacent to zinc finger domain, 2B (Baz2b) alternative variant fSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.gSep08</a>	<a href="#">317627</a>	1946	390	2	45	bromodomain adjacent to zinc finger domain, 2B (4.8 kD) (Baz2b) alternative variant gSep08, mRNA.
<a href="#">Baz2b</a>	<a href="#">Baz2b.hSep08</a>	<a href="#">317627</a>	134425	377	4	48	bromodomain adjacent to zinc finger domain, 2B (5.7 kD) (Baz2b) alternative variant hSep08, mRNA.
<a href="#">Bbs1</a>	<a href="#">Bbs1.bSep08</a>	<a href="#">309156</a>	4854	697	4	136	bardet-Biedl syndrome 1 homolog (human) (Bbs1) alternative variant bSep08, mRNA.

<a href="#">Bbs1</a>	<a href="#">Bbs1.cSep08</a>	<a href="#">309156</a>	3347	355	5	44	bardet-Biedl syndrome 1 homolog (human) (Bbs1) alternative variant cSep08, mRNA.
<a href="#">Bbs1</a>	<a href="#">Bbs1.dSep08</a>	<a href="#">309156</a>	612	369	2	35	bardet-Biedl syndrome 1 homolog (human) (Bbs1) alternative variant dSep08, mRNA.
<a href="#">Bbs2</a>	<a href="#">Bbs2.bSep08</a>	<a href="#">113948</a>	6605	692	6	198	bardet-Biedl syndrome 2 like (Bbs2) alternative variant bSep08, mRNA.
<a href="#">Bbs2</a>	<a href="#">Bbs2.cSep08</a>	<a href="#">113948</a>	13189	746	7	125	bardet-Biedl syndrome 2 like (Bbs2) alternative variant cSep08, mRNA.
<a href="#">Bbs2</a>	<a href="#">Bbs2.dSep08</a>	<a href="#">113948</a>	683	514	2	85	bardet-Biedl syndrome 2 like (Bbs2) alternative variant dSep08, mRNA.
<a href="#">Bbs4</a>	<a href="#">Bbs4.bSep08</a>	<a href="#">300754</a>	21345	1536	10	417	bardet-Biedl syndrome 4 homolog (human) (47.4 kD) (Bbs4) alternative variant bSep08, mRNA.
<a href="#">Bbs4</a>	<a href="#">Bbs4.cSep08</a>	<a href="#">300754</a>	5766	507	1	54	bardet-Biedl syndrome 4 homolog (human) (Bbs4) alternative variant cSep08, mRNA.
<a href="#">Bbs5</a>	<a href="#">Bbs5.bSep08</a>	<a href="#">362142</a>	8450	639	5	148	bardet-Biedl syndrome 5 (human) (Bbs5) alternative variant bSep08, mRNA.
<a href="#">Bbs5</a>	<a href="#">Bbs5.cSep08</a>	<a href="#">362142</a>	3179	683	4	121	bardet-Biedl syndrome 5 (human) (Bbs5) alternative variant cSep08, mRNA.
<a href="#">Bbs5</a>	<a href="#">Bbs5.dSep08</a>	<a href="#">362142</a>	14864	1020	4	68	bardet-Biedl syndrome 5 (human) (8.0 kD) (Bbs5) alternative variant dSep08, mRNA.
<a href="#">Bbs7</a>	<a href="#">Bbs7.bSep08</a>	<a href="#">361930</a>	4337	595	2	110	bardet-Biedl syndrome 7 (12.4 kD) (Bbs7) alternative variant bSep08, mRNA.
<a href="#">Bbs9</a>	<a href="#">Bbs9.aSep08</a>	<a href="#">315484</a>	241031	1914	11	511	bardet-Biedl syndrome 9 (Bbs9) alternative variant aSep08, mRNA.
<a href="#">Bbs9</a>	<a href="#">Bbs9.bSep08</a>	<a href="#">315484</a>	48051	378	1	60	bardet-Biedl syndrome 9 (Bbs9) alternative variant bSep08, mRNA.
<a href="#">Bbx</a>	<a href="#">Bbx.bSep08</a>	<a href="#">303970</a>	86029	572	4	60	bobby sox homolog (Drosophila) (Bbx) alternative variant bSep08, mRNA.
<a href="#">Bbx</a>	<a href="#">Bbx.cSep08</a>	<a href="#">303970</a>	152454	348	3	34	bobby sox homolog (Drosophila) (4.0 kD) (Bbx) alternative variant cSep08, mRNA.
<a href="#">Bcam</a>	<a href="#">Bcam.bSep08</a>	<a href="#">78958</a>	4835	792	1	190	basal cell adhesion molecule (Bcam) alternative variant bSep08, mRNA.
<a href="#">Bcan</a>	<a href="#">Bcan.cSep08</a>	<a href="#">25393</a>	1682	418	3	139	brevican (Bcan) alternative variant cSep08, mRNA.
<a href="#">Bcan</a>	<a href="#">Bcan.fSep08</a>	<a href="#">25393</a>	1347	312	3	90	brevican (Bcan) alternative variant fSep08, mRNA.
<a href="#">Bcan</a>	<a href="#">Bcan.hSep08</a>	<a href="#">25393</a>	1445	651	2	25	brevican (Bcan) alternative variant hSep08, mRNA.
<a href="#">Bcap29</a>	<a href="#">Bcap29.bSep08</a>	<a href="#">298943</a>	34451	833	1	227	B-cell receptor-associated protein 29 (Bcap29) alternative variant bSep08, mRNA.
<a href="#">Bcap31</a>	<a href="#">Bcap31.bSep08</a>	<a href="#">293852</a>	28929	866	7	233	B-cell receptor-associated protein 31 (Bcap31) alternative variant bSep08, mRNA.
<a href="#">Bcap31</a>	<a href="#">Bcap31.cSep08</a>	<a href="#">293852</a>	28698	1548	6	227	B-cell receptor-associated protein 31 (25.8 kD) (Bcap31) alternative variant cSep08, mRNA.
<a href="#">Bcap31</a>	<a href="#">Bcap31.dSep08</a>	<a href="#">293852</a>	27741	716	6	178	B-cell receptor-associated protein 31 (Bcap31) alternative variant dSep08, mRNA.
<a href="#">Bcap31</a>	<a href="#">Bcap31.eSep08</a>	<a href="#">293852</a>	28457	677	5	159	B-cell receptor-associated protein 31 (Bcap31) alternative variant eSep08, mRNA.
<a href="#">Bcap31</a>	<a href="#">Bcap31.gSep08</a>	<a href="#">293852</a>	2035	658	2	35	B-cell receptor-associated protein 31 (4.1 kD) (Bcap31) alternative variant gSep08, mRNA.

<a href="#">Bcar1</a>	<a href="#">Bcar1.aSep08</a>	<a href="#">25414</a>	23107	3115	5	874	breast cancer anti-estrogen resistance 1 (94.5 kD) (Bcar1) alternative variant aSep08, mRNA.
<a href="#">Bcar1</a>	<a href="#">Bcar1.bSep08</a>	<a href="#">25414</a>	12441	472	2	131	breast cancer anti-estrogen resistance 1 (Bcar1) alternative variant bSep08, mRNA.
<a href="#">Bcas1</a>	<a href="#">Bcas1.bSep08</a>	<a href="#">246755</a>	63013	2470	7	470	breast carcinoma amplified sequence 1 (Bcas1) alternative variant bSep08, mRNA.
<a href="#">Bcas1</a>	<a href="#">Bcas1.cSep08</a>	<a href="#">246755</a>	62580	1803	5	392	breast carcinoma amplified sequence 1 (Bcas1) alternative variant cSep08, mRNA.
<a href="#">Bcas1</a>	<a href="#">Bcas1.dSep08</a>	<a href="#">246755</a>	62580	1761	4	378	breast carcinoma amplified sequence 1 (Bcas1) alternative variant dSep08, mRNA.
<a href="#">Bcas1</a>	<a href="#">Bcas1.eSep08</a>	<a href="#">246755</a>	62580	1734	5	369	breast carcinoma amplified sequence 1 (Bcas1) alternative variant eSep08, mRNA.
<a href="#">Bcas1</a>	<a href="#">Bcas1.fSep08</a>	<a href="#">246755</a>	22642	423	1	141	breast carcinoma amplified sequence 1 (Bcas1) alternative variant fSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.aSep08</a>	<a href="#">363662</a>	301809	1981	7	543	breast carcinoma amplified sequence 3 like (Bcas3) alternative variant aSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.bSep08</a>	<a href="#">363662</a>	251058	920	8	305	breast carcinoma amplified sequence 3 like (Bcas3) alternative variant bSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.cSep08</a>	<a href="#">363662</a>	4557	1532	2	142	breast carcinoma amplified sequence 3 like (15.8 kD) (Bcas3) alternative variant cSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.dSep08</a>	<a href="#">363662</a>	86521	1084	3	130	breast carcinoma amplified sequence 3 CRA c like (Bcas3) alternative variant dSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.eSep08</a>	<a href="#">363662</a>	18588	417	1	70	breast carcinoma amplified sequence 3 like (Bcas3) alternative variant eSep08, mRNA.
<a href="#">Bcas3</a>	<a href="#">Bcas3.fSep08</a>	<a href="#">363662</a>	87525	747	4	52	breast carcinoma amplified sequence 3 like (Bcas3) alternative variant fSep08, mRNA.
<a href="#">Bcat1</a>	<a href="#">Bcat1.bSep08</a>	<a href="#">29592</a>	84681	1551	8	306	branched chain aminotransferase 1, cytosolic (34.2 kD) (Bcat1) alternative variant bSep08, mRNA.
<a href="#">Bcat1</a>	<a href="#">Bcat1.cSep08</a>	<a href="#">29592</a>	28611	869	6	217	branched chain aminotransferase 1, cytosolic (Bcat1) alternative variant cSep08, mRNA.
<a href="#">Bcat1</a>	<a href="#">Bcat1.dSep08</a>	<a href="#">29592</a>	16528	390	4	129	branched chain aminotransferase 1, cytosolic (Bcat1) alternative variant dSep08, mRNA.
<a href="#">Bcat1</a>	<a href="#">Bcat1.eSep08</a>	<a href="#">29592</a>	50104	595	5	122	branched chain aminotransferase 1, cytosolic (Bcat1) alternative variant eSep08, mRNA.
<a href="#">Bcat2</a>	<a href="#">Bcat2.bSep08</a>	<a href="#">64203</a>	13142	691	6	227	branched chain aminotransferase 2, mitochondrial (Bcat2) alternative variant bSep08, mRNA.
<a href="#">Bccip</a>	<a href="#">Bccip.aSep08</a>	<a href="#">361666</a>	12998	1815	7	370	BRCA2 and CDKN1A interacting protein (41.7 kD) (Bccip) alternative variant aSep08, mRNA.
<a href="#">Bckdha</a>	<a href="#">Bckdha.bSep08</a>	<a href="#">25244</a>	1619	1306	3	310	branched chain ketoacid dehydrogenase E1, alpha polypeptide (Bckdha) alternative variant bSep08, mRNA.
<a href="#">Bckdha</a>	<a href="#">Bckdha.cSep08</a>	<a href="#">25244</a>	1589	718	4	233	branched chain ketoacid dehydrogenase E1, alpha polypeptide (Bckdha) alternative variant cSep08, mRNA.
<a href="#">Bckdha</a>	<a href="#">Bckdha.dSep08</a>	<a href="#">25244</a>	9893	693	4	167	branched chain ketoacid dehydrogenase E1, alpha polypeptide (Bckdha) alternative variant dSep08, mRNA.
<a href="#">Bckdk</a>	<a href="#">Bckdk.bSep08</a>	<a href="#">29603</a>	4277	1577	10	365	branched chain ketoacid dehydrogenase kinase (41.2 kD) (Bckdk) alternative variant bSep08, mRNA.

<a href="#">Bckdk</a>	<a href="#">Bckdk.cSep08</a>	<a href="#">29603</a>	2911	957	7	248	branched chain ketoacid dehydrogenase kinase (Bckdk) alternative variant cSep08, mRNA.
<a href="#">Bckdk</a>	<a href="#">Bckdk.dSep08</a>	<a href="#">29603</a>	892	502	2	98	branched chain ketoacid dehydrogenase kinase (Bckdk) alternative variant dSep08, mRNA.
<a href="#">Bcl2l1</a>	<a href="#">Bcl2l1.dSep08</a>	<a href="#">24888</a>	1241	976	2	62	bcl2-like 1 (7.1 kD) (Bcl2l1) alternative variant dSep08, mRNA.
<a href="#">Bcl2l2</a>	<a href="#">Bcl2l2.cSep08</a>	<a href="#">60434</a>	1066	510	3	64	bcl2-like 2 (Bcl2l2) alternative variant cSep08, mRNA.
<a href="#">Bcl2l2</a>	<a href="#">Bcl2l2.dSep08</a>	<a href="#">60434</a>	702	417	2	45	bcl2-like 2 (Bcl2l2) alternative variant dSep08, mRNA.
<a href="#">Bcl2l12</a>	<a href="#">Bcl2l12.bSep08</a>	<a href="#">361567</a>	3286	662	3	173	BCL2-like 12 (proline rich) (Bcl2l12) alternative variant bSep08, mRNA.
<a href="#">Bcl2l12</a>	<a href="#">Bcl2l12.cSep08</a>	<a href="#">361567</a>	3076	637	3	165	BCL2-like 12 (proline rich) (Bcl2l12) alternative variant cSep08, mRNA.
<a href="#">Bcl2l12</a>	<a href="#">Bcl2l12.eSep08</a>	<a href="#">361567</a>	6075	449	4	110	BCL2-like 12 (proline rich) (Bcl2l12) alternative variant eSep08, mRNA.
<a href="#">Bcl2l13</a>	<a href="#">Bcl2l13.aSep08</a>	<a href="#">312682</a>	42812	2877	2	444	BCL2-like 13 (apoptosis facilitator) (Bcl2l13) alternative variant aSep08, mRNA.
<a href="#">Bcl2l14</a>	<a href="#">Bcl2l14.bSep08</a>	<a href="#">500348</a>	3621	774	3	55	bcl2-like 14 (apoptosis facilitator) (6.1 kD) (Bcl2l14) alternative variant bSep08, mRNA.
<a href="#">Bcl2l14</a>	<a href="#">Bcl2l14.cSep08</a>	<a href="#">500348</a>	39980	558	3	18	bcl2-like 14 (apoptosis facilitator) (Bcl2l14) alternative variant cSep08, mRNA.
<a href="#">Bcl2l14</a>	<a href="#">Bcl2l14.dSep08</a>	<a href="#">500348</a>	8665	394	2	58	bcl2-like 14 (apoptosis facilitator) (Bcl2l14) alternative variant dSep08, mRNA.
<a href="#">Bcl3</a>	<a href="#">Bcl3.bSep08</a>	<a href="#">680611</a>	1991	590	2	96	B-cell leukemia/lymphoma 3 (Bcl3) alternative variant bSep08, mRNA.
<a href="#">Bcl3</a>	<a href="#">Bcl3.cSep08</a>	<a href="#">680611</a>	2627	617	2	83	B-cell leukemia/lymphoma 3 (Bcl3) alternative variant cSep08, mRNA.
<a href="#">Bcl6b</a>	<a href="#">Bcl6b.bSep08</a>	<a href="#">360551</a>	4487	976	3	209	B-cell CLL/lymphoma 6, member B (Bcl6b) alternative variant bSep08, mRNA.
<a href="#">Bcl7b</a>	<a href="#">Bcl7b.bSep08</a>	<a href="#">368001</a>	4768	1525	1	126	B-cell CLL/lymphoma 7B (Bcl7b) alternative variant bSep08, mRNA.
<a href="#">Bcl7c</a>	<a href="#">Bcl7c.bSep08</a>	<a href="#">293514</a>	2560	545	4	157	B-cell CLL/lymphoma 7C (Bcl7c) alternative variant bSep08, mRNA.
<a href="#">Bcl7c</a>	<a href="#">Bcl7c.cSep08</a>	<a href="#">293514</a>	1325	372	4	123	B-cell CLL/lymphoma 7C (Bcl7c) alternative variant cSep08, mRNA.
<a href="#">Bcl9</a>	<a href="#">Bcl9.bSep08</a>	<a href="#">310704</a>	7040	4774	1	628	B-cell CLL/lymphoma 9 (64.3 kD) (Bcl9) alternative variant bSep08, mRNA.
<a href="#">Bcl10</a>	<a href="#">Bcl10.cSep08</a>	<a href="#">83477</a>	8616	350	2	33	B-cell leukemia/lymphoma 10 (3.7 kD) (Bcl10) alternative variant cSep08, mRNA.
<a href="#">Bcl11a</a>	<a href="#">Bcl11a.aSep08</a>	<a href="#">305589</a>	93720	1486	5	240	B-cell CLL/lymphoma 11A (zinc finger protein) (26.4 kD) (Bcl11a) alternative variant aSep08, mRNA.
<a href="#">Bcl11a</a>	<a href="#">Bcl11a.cSep08</a>	<a href="#">305589</a>	86203	617	2	64	B-cell CLL/lymphoma 11A (zinc finger protein) (Bcl11a) alternative variant cSep08, mRNA.
<a href="#">Bcl11a</a>	<a href="#">Bcl11a.dSep08</a>	<a href="#">305589</a>	8486	597	2	60	B-cell CLL/lymphoma 11A (zinc finger protein) (Bcl11a) alternative variant dSep08, mRNA.
<a href="#">Bcl11b</a>	<a href="#">Bcl11b.bSep08</a>	<a href="#">314423</a>	76178	2567	3	796	B-cell leukemia/lymphoma 11B (Bcl11b) alternative variant bSep08, mRNA.

<a href="#">Bclaf1</a>	<a href="#">Bclaf1.bSep08</a>	<a href="#">293017</a>	10884	819	4	216	transcription factor 1 (Bclaf1) alternative variant bSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.cSep08</a>	<a href="#">293017</a>	7584	632	5	210	transcription factor 1 (Bclaf1) alternative variant cSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.dSep08</a>	<a href="#">293017</a>	14418	4603	7	129	transcription factor 1 (15.4 kD) (Bclaf1) alternative variant dSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.eSep08</a>	<a href="#">293017</a>	6114	742	3	89	transcription factor 1 (10.5 kD) (Bclaf1) alternative variant eSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.fSep08</a>	<a href="#">293017</a>	10655	492	3	83	aa2-041 like (Bclaf1) alternative variant fSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.gSep08</a>	<a href="#">293017</a>	6497	565	4	54	putative protein (Bclaf1) alternative variant gSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.hSep08</a>	<a href="#">293017</a>	6793	543	4	34	transcription factor 1 like (Bclaf1) alternative variant hSep08, mRNA.
<a href="#">Bclaf1</a>	<a href="#">Bclaf1.iSep08</a>	<a href="#">293017</a>	9825	485	2	46	putative protein (Bclaf1) alternative variant iSep08, mRNA.
<a href="#">Bco2</a>	<a href="#">Bco2.bSep08</a>	<a href="#">315644</a>	1644	682	1	87	beta-carotene oxygenase 2 (Bco2) alternative variant bSep08, mRNA.
<a href="#">Bcor</a>	<a href="#">Bcor.aSep08</a>	<a href="#">317346</a>	20179	4396	1	1175	bcl6 interacting corepressor (Bcor) alternative variant aSep08, mRNA.
<a href="#">Bcor</a>	<a href="#">Bcor.bSep08</a>	<a href="#">317346</a>	1736	723	1	241	bcl6 interacting corepressor (Bcor) alternative variant bSep08, mRNA.
<a href="#">Bcorl1</a>	<a href="#">Bcorl1.aSep08</a>	<a href="#">302810</a>	30093	3532		552	BCL6 co-repressor-like 1 (Bcorl1) mRNA.
<a href="#">Bcr</a>	<a href="#">Bcr.aSep08</a>	<a href="#">309696</a>	48003	1783		472	breakpoint cluster region homolog (Bcr) alternative variant aSep08, mRNA.
<a href="#">Bcr</a>	<a href="#">Bcr.bSep08</a>	<a href="#">309696</a>	25951	1177		389	breakpoint cluster region homolog (Bcr) alternative variant bSep08, mRNA.
<a href="#">Bcr</a>	<a href="#">Bcr.cSep08</a>	<a href="#">309696</a>	9314	1665	1	307	breakpoint cluster region homolog (Bcr) alternative variant cSep08, mRNA.
<a href="#">Bcr</a>	<a href="#">Bcr.dSep08</a>	<a href="#">309696</a>	8270	1077	1	282	breakpoint cluster region homolog (Bcr) alternative variant dSep08, mRNA.
<a href="#">Bcr</a>	<a href="#">Bcr.eSep08</a>	<a href="#">309696</a>	14421	405	2	135	breakpoint cluster region homolog (Bcr) alternative variant eSep08, mRNA.
<a href="#">Bdh1</a>	<a href="#">Bdh1.bSep08</a>	<a href="#">117099</a>	41950	3061	9	344	3-hydroxybutyrate dehydrogenase, type 1 (38.3 kD) (Bdh1) alternative variant bSep08, mRNA.
<a href="#">Bdh1</a>	<a href="#">Bdh1.cSep08</a>	<a href="#">117099</a>	38057	769	6	185	3-hydroxybutyrate dehydrogenase, type 1 (Bdh1) alternative variant cSep08, mRNA.
<a href="#">Bdh1</a>	<a href="#">Bdh1.dSep08</a>	<a href="#">117099</a>	2406	727	2	176	3-hydroxybutyrate dehydrogenase, type 1 (Bdh1) alternative variant dSep08, mRNA.
<a href="#">Bdh1</a>	<a href="#">Bdh1.eSep08</a>	<a href="#">117099</a>	32581	719	6	152	3-hydroxybutyrate dehydrogenase, type 1 (Bdh1) alternative variant eSep08, mRNA.
<a href="#">Bdh2</a>	<a href="#">Bdh2.bSep08</a>	<a href="#">295458</a>	20541	1069	10	228	3-hydroxybutyrate dehydrogenase, type 2 (24.8 kD) (Bdh2) alternative variant bSep08, mRNA.
<a href="#">Bdh2</a>	<a href="#">Bdh2.cSep08</a>	<a href="#">295458</a>	2392	733	2	71	3-hydroxybutyrate dehydrogenase, type 2 (Bdh2) alternative variant cSep08, mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.aSep08</a>	<a href="#">24225</a>	31261	4006	2	362	brain derived neurotrophic factor (40.6 kD) (Bdnf) alternative variant aSep08, mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.dSep08</a>	<a href="#">24225</a>	16132	3890	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant dSep08, mRNA.

<a href="#">Bdnf</a>	<a href="#">Bdnf.eSep08</a>	<a href="#">24225</a>	31723	3694	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant eSep08, mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.fSep08</a>	<a href="#">24225</a>	48197	3839	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant fSep08, mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.gSep08</a>	<a href="#">24225</a>	32310	3953	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant gSep08, mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.hSep08</a>	<a href="#">24225</a>	49333	4112	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant hSep08, complete mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.iSep08</a>	<a href="#">24225</a>	49333	4029	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant iSep08, complete mRNA.
<a href="#">Bdnf</a>	<a href="#">Bdnf.jSep08</a>	<a href="#">24225</a>	49333	3817	2	249	brain derived neurotrophic factor (28.1 kD) (Bdnf) alternative variant jSep08, complete mRNA.
<a href="#">Bdp1</a>	<a href="#">Bdp1.bSep08</a>	<a href="#">294687</a>	12615	1253	6	310	B double prime 1, subunit of RNA polymerase III transcription initiation factor IIIB (34.2 kD) (Bdp1) alternative variant bSep08, mRNA.
<a href="#">Bdp1</a>	<a href="#">Bdp1.cSep08</a>	<a href="#">294687</a>	20354	537	6	179	B double prime 1, subunit of RNA polymerase III transcription initiation factor IIIB (Bdp1) alternative variant cSep08, mRNA.
<a href="#">Bdp1</a>	<a href="#">Bdp1.dSep08</a>	<a href="#">294687</a>	6110	269	3	29	B double prime 1, subunit of RNA polymerase III transcription initiation factor IIIB (Bdp1) alternative variant dSep08, mRNA.
<a href="#">Beach.0</a>	<a href="#">Beach.0.aSep08</a>		8114	893		259	beige/BEACH (Beach.0) mRNA.
<a href="#">Beach.1</a>	<a href="#">Beach.1.aSep08</a>		5094	395		131	neurobeachin like 1 (Beach.1) mRNA.
<a href="#">Beach.2</a>	<a href="#">Beach.2.aSep08</a>		958	472	4	156	CRA a (Beach.2) alternative variant aSep08, mRNA.
<a href="#">Beach.3</a>	<a href="#">Beach.3.aSep08</a>		16416	421		119	neurobeachin (Beach.3) mRNA.
<a href="#">Beach.4</a>	<a href="#">Beach.4.aSep08</a>		37279	413		137	neurobeachin (Beach.4) mRNA.
<a href="#">Bean</a>	<a href="#">Bean.aSep08</a>	<a href="#">361358</a>	35411	2617	3	222	brain expressed, associated with Nedd4 (24.5 kD) (Bean) alternative variant aSep08, mRNA.
<a href="#">Bean</a>	<a href="#">Bean.bSep08</a>	<a href="#">361358</a>	38703	395	1	131	brain expressed, associated with Nedd4 (Bean) alternative variant bSep08, mRNA.
<a href="#">Becn1</a>	<a href="#">Becn1.cSep08</a>	<a href="#">114558</a>	1649	729	3	156	beclin 1, autophagy related (Becn1) alternative variant cSep08, mRNA.
<a href="#">Becn1</a>	<a href="#">Becn1.dSep08</a>	<a href="#">114558</a>	7959	695	5	116	beclin 1, autophagy related (12.8 kD) (Becn1) alternative variant dSep08, mRNA.
<a href="#">Becn1</a>	<a href="#">Becn1.eSep08</a>	<a href="#">114558</a>	4069	821	3	107	beclin 1, autophagy related (Becn1) alternative variant eSep08, mRNA.
<a href="#">Becn1</a>	<a href="#">Becn1.fSep08</a>	<a href="#">114558</a>	7293	428	4	95	beclin 1, autophagy related (Becn1) alternative variant fSep08, mRNA.
<a href="#">Becn1</a>	<a href="#">Becn1.hSep08</a>	<a href="#">114558</a>	7304	371	3	46	beclin 1, autophagy related (5.1 kD) (Becn1) alternative variant hSep08, complete mRNA.
<a href="#">beeby</a>	<a href="#">beeby.aSep08</a>		14331	1190		396	THO complex 2 (beeby) mRNA.
<a href="#">beechy</a>	<a href="#">beechy.aSep08</a>		5715	766		41	putative protein (5.0 kD) (beechy) mRNA.
<a href="#">beefef</a>	<a href="#">beefef.aSep08</a>		10266	268		52	putative protein (6.0 kD) (beefef) mRNA.
<a href="#">beeflo</a>	<a href="#">beeflo.aSep08</a>		19759	369		63	putative protein of mammalian origin (7.0 kD) (beeflo) mRNA.
<a href="#">beeflu</a>	<a href="#">beeflu.aSep08</a>		5125	653		75	putative protein (beeflu) mRNA.

<a href="#">beekee</a>	<a href="#">beekee.aSep08</a>		6179	758		63	putative protein (beekee) mRNA.
<a href="#">beeloy</a>	<a href="#">beeloy.aSep08</a>		1195	418		64	putative protein (beeloy) mRNA.
<a href="#">beemer</a>	<a href="#">beemer.aSep08</a>		5263	963		321	rho GTPase activating protein 23 (beemer) alternative variant aSep08, mRNA.
<a href="#">beemer</a>	<a href="#">beemer.bSep08</a>		4735	434		144	rho GTPase activating protein 23 (beemer) alternative variant bSep08, mRNA.
<a href="#">beenoy</a>	<a href="#">beenoy.aSep08</a>		435	258		79	putative protein (beenoy) mRNA.
<a href="#">beepor</a>	<a href="#">beepor.aSep08</a>		2190	536		91	putative protein (beepor) mRNA.
<a href="#">beesa</a>	<a href="#">beesa.aSep08</a>		1794	381		126	low density lipoprotein-related protein 1B (beesa) mRNA.
<a href="#">beeshee</a>	<a href="#">beeshee.aSep08</a>		14645	373		89	putative mitochondrial protein (10.0 kD) (beeshee) mRNA.
<a href="#">beeto</a>	<a href="#">beeto.aSep08</a>		2441	1336	2	72	putative protein (8.2 kD) (beeto) alternative variant aSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.bSep08</a>		2137	531	3	48	putative protein (beeto) alternative variant bSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.cSep08</a>		15844	893	3	39	putative protein (beeto) alternative variant cSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.dSep08</a>		15644	3017	8	91	putative protein (10.7 kD) (beeto) alternative variant dSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.eSep08</a>		3230	1617	3	89	putative cytoplasmic protein (10.5 kD) (beeto) alternative variant eSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.fSep08</a>		42420	1522	12		
<a href="#">beeto</a>	<a href="#">beeto.gSep08</a>		8862	1004	5	47	putative protein (beeto) alternative variant gSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.iSep08</a>		26150	553	4	38	putative protein (4.2 kD) (beeto) alternative variant iSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.kSep08</a>		21841	459	5	34	putative protein (beeto) alternative variant kSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.lSep08</a>		14296	239	3	23	putative protein (beeto) alternative variant lSep08, mRNA.
<a href="#">beeto</a>	<a href="#">beeto.mSep08</a>		914	113	2	37	putative protein (beeto) alternative variant mSep08, mRNA.
<a href="#">beevar</a>	<a href="#">beevar.aSep08</a>		19947	432		107	putative protein (beevar) mRNA.
<a href="#">beewey</a>	<a href="#">beewey.aSep08</a>		2021	438		96	putative protein of metazoan origin (beewey) mRNA.
<a href="#">Begain</a>	<a href="#">Begain.bSep08</a>	<a href="#">79146</a>	5064	2537	3	501	brain-enriched guanylate kinase-associated (53.7 kD) (Begain) alternative variant bSep08, mRNA.
<a href="#">Begain</a>	<a href="#">Begain.eSep08</a>	<a href="#">79146</a>	1944	402	2	103	brain-enriched guanylate kinase-associated (Begain) alternative variant eSep08, mRNA.
<a href="#">berby</a>	<a href="#">berby.aSep08</a>		1169	514		115	putative protein, with a coiled coil domain, of mammalian origin (berby) mRNA.
<a href="#">berchy</a>	<a href="#">berchy.aSep08</a>		5535	261		28	putative protein (berchy) mRNA.
<a href="#">berfer</a>	<a href="#">berfer.aSep08</a>		11799	865		55	putative protein (6.5 kD) (berfer) mRNA.
<a href="#">berflo</a>	<a href="#">berflo.aSep08</a>		20285	427	3	82	putative protein of mammalian origin (berflo) alternative variant aSep08, mRNA.
<a href="#">berflo</a>	<a href="#">berflo.bSep08</a>		29918	279	1	74	putative protein (berflo) alternative variant bSep08, mRNA.
<a href="#">berflu</a>	<a href="#">berflu.aSep08</a>		8492	307		47	zinc finger protein 585b (berflu) mRNA.
<a href="#">berkee</a>	<a href="#">berkee.aSep08</a>		2347	1100		42	putative protein (berkee) mRNA.
<a href="#">berloy</a>	<a href="#">berloy.aSep08</a>		43372	560		186	putative protein of mammalian origin (berloy) mRNA.
<a href="#">bermer</a>	<a href="#">bermer.aSep08</a>		1392	559		115	putative protein (bermer) mRNA.
<a href="#">bernoy</a>	<a href="#">bernoy.aSep08</a>		9002	858		245	type IX alpha 1 (bernoy) mRNA.

<a href="#">berpor</a>	<a href="#">berpor.aSep08</a>		484	410		109	putative protein of metazoan origin (berpor) mRNA.
<a href="#">bersa</a>	<a href="#">bersa.aSep08</a>		2433	800		266	myosin-ia (bersa) mRNA.
<a href="#">bershee</a>	<a href="#">bershee.aSep08</a>		8059	584		49	putative protein (bershee) mRNA.
<a href="#">berto</a>	<a href="#">berto.aSep08</a>		4030	548		89	CRA a like (9.9 kD) (berto) mRNA.
<a href="#">bervar</a>	<a href="#">bervar.aSep08</a>		3839	477		158	ubiquitin 24 (bervar) mRNA.
<a href="#">berwey</a>	<a href="#">berwey.aSep08</a>		68738	406	2	83	putative protein (9.6 kD) (berwey) alternative variant aSep08, mRNA.
<a href="#">berwey</a>	<a href="#">berwey.bSep08</a>		239423	1443	5	65	putative protein (berwey) alternative variant bSep08, mRNA.
<a href="#">Bet1</a>	<a href="#">Bet1.bSep08</a>	<a href="#">29631</a>	9368	626	5	96	blocked early in transport 1 homolog (S. cerevisiae) (10.7 kD) (Bet1) alternative variant bSep08, complete mRNA.
<a href="#">Bet1</a>	<a href="#">Bet1.cSep08</a>	<a href="#">29631</a>	9586	787	5	77	blocked early in transport 1 homolog (S. cerevisiae) (8.6 kD) (Bet1) alternative variant cSep08, complete mRNA.
<a href="#">Bet1</a>	<a href="#">Bet1.dSep08</a>	<a href="#">29631</a>	9596	689	4	73	blocked early in transport 1 homolog (S. cerevisiae) (8.0 kD) (Bet1) alternative variant dSep08, mRNA.
<a href="#">Bet1</a>	<a href="#">Bet1.eSep08</a>	<a href="#">29631</a>	10351	1396	4	52	blocked early in transport 1 homolog (S. cerevisiae) (6.1 kD) (Bet1) alternative variant eSep08, complete mRNA.
<a href="#">Bet1l</a>	<a href="#">Bet1l.aSep08</a>	<a href="#">54400</a>	2172	774	2	125	blocked early in transport 1 homolog (S. cerevisiae) like (Bet1l) alternative variant aSep08, mRNA.
<a href="#">beyby</a>	<a href="#">beyby.aSep08</a>		1377	403	3	116	upf3 regulator of nonsense transcripts homolog B (beyby) alternative variant aSep08, mRNA.
<a href="#">beychy</a>	<a href="#">beychy.aSep08</a>		1049	549		90	putative protein (beychy) mRNA.
<a href="#">beyfer</a>	<a href="#">beyfer.aSep08</a>		33959	626		93	putative cytoplasmic protein (10.7 kD) (beyfer) mRNA.
<a href="#">beyflo</a>	<a href="#">beyflo.aSep08</a>		1220	706		53	putative protein (beyflo) mRNA.
<a href="#">beyflu</a>	<a href="#">beyflu.aSep08</a>		9733	461	4	33	putative protein (beyflu) alternative variant aSep08, mRNA.
<a href="#">beyflu</a>	<a href="#">beyflu.cSep08</a>		2463	242	2	29	putative protein (beyflu) alternative variant cSep08, mRNA.
<a href="#">beykee</a>	<a href="#">beykee.aSep08</a>		2187	656		66	contactin associated protein-like 5C (7.5 kD) (beykee) mRNA.
<a href="#">beyloy</a>	<a href="#">beyloy.aSep08</a>		4679	1349		61	putative protein (7.2 kD) (beyloy) mRNA.
<a href="#">beymer</a>	<a href="#">beymer.aSep08</a>		6819	401		88	putative protein (beymer) mRNA.
<a href="#">beynoy</a>	<a href="#">beynoy.aSep08</a>		555	377		55	putative protein (6.4 kD) (beynoy) mRNA.
<a href="#">beypor</a>	<a href="#">beypor.aSep08</a>		1861	620		64	putative protein (7.2 kD) (beypor) mRNA.
<a href="#">beysa</a>	<a href="#">beysa.aSep08</a>		2879	400		54	putative protein (5.8 kD) (beysa) mRNA.
<a href="#">beyshee</a>	<a href="#">beyshee.aSep08</a>		1505	284		60	putative protein (beyshee) mRNA.
<a href="#">beyto</a>	<a href="#">beyto.aSep08</a>		1546	732		56	CRA b like (6.2 kD) (beyto) mRNA.
<a href="#">beyvar</a>	<a href="#">beyvar.aSep08</a>		7835	748		44	putative protein (beyvar) mRNA.
<a href="#">beywey</a>	<a href="#">beywey.aSep08</a>		10799	372		37	putative protein (beywey) mRNA.
<a href="#">Bfar</a>	<a href="#">Bfar.bSep08</a>	<a href="#">304709</a>	6033	942	2	147	bifunctional apoptosis regulator (17.8 kD) (Bfar) alternative variant bSep08, mRNA.
<a href="#">Bgn</a>	<a href="#">Bgn.bSep08</a>	<a href="#">25181</a>	2836	1099	5	239	biglycan (Bgn) alternative variant bSep08, mRNA.
<a href="#">Bgn</a>	<a href="#">Bgn.cSep08</a>	<a href="#">25181</a>	5529	763	3	141	biglycan (Bgn) alternative variant cSep08, mRNA.
<a href="#">Bhlhb2</a>	<a href="#">Bhlhb2.bSep08</a>	<a href="#">79431</a>	3487	1512		291	orange (Bhlhb2) alternative variant bSep08, mRNA.
<a href="#">Bhlhb9</a>	<a href="#">Bhlhb9.bSep08</a>	<a href="#">317407</a>	1777	614	3	52	putative protein (Bhlhb9) alternative variant bSep08, mRNA.



<a href="#">Bhmt2</a>	<a href="#">Bhmt2.bSep08</a>	<a href="#">365972</a>	6175	1615		245	betaine-homocysteine methyltransferase 2 (Bhmt2) alternative variant bSep08, mRNA.
<a href="#">Bicd1</a>	<a href="#">Bicd1.bSep08</a>	<a href="#">362466</a>	2946	690	2	149	bicaudal D homolog 1 (Drosophila) (Bicd1) alternative variant bSep08, mRNA.
<a href="#">Bicd2</a>	<a href="#">Bicd2.bSep08</a>	<a href="#">306809</a>	6989	710	4	182	bicaudal D homolog 2 (Drosophila) (Bicd2) alternative variant bSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.aSep08</a>	<a href="#">117028</a>	59536	2391	18	557	bridging integrator 1 (61.1 kD) (Bin1) alternative variant aSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.bSep08</a>	<a href="#">117028</a>	35669	1139	12	340	bridging integrator 1 (Bin1) alternative variant bSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.cSep08</a>	<a href="#">117028</a>	35885	1249	13	325	bridging integrator 1 (Bin1) alternative variant cSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.dSep08</a>	<a href="#">117028</a>	24849	1164	9	311	bridging integrator 1 (Bin1) alternative variant dSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.eSep08</a>	<a href="#">117028</a>	22880	852	8	263	bridging integrator 1 (Bin1) alternative variant eSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.fSep08</a>	<a href="#">117028</a>	20834	986	11	260	bridging integrator 1 (Bin1) alternative variant fSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.gSep08</a>	<a href="#">117028</a>	5864	413	5	137	bridging integrator 1 (Bin1) alternative variant gSep08, mRNA.
<a href="#">Bin1</a>	<a href="#">Bin1.iSep08</a>	<a href="#">117028</a>	5270	2105	3	72	bridging integrator 1 (8.4 kD) (Bin1) alternative variant iSep08, mRNA.
<a href="#">Bin2</a>	<a href="#">Bin2.bSep08</a>	<a href="#">366988</a>	20978	737	9	238	bridging integrator 2 (Bin2) alternative variant bSep08, mRNA.
<a href="#">Bin2</a>	<a href="#">Bin2.dSep08</a>	<a href="#">366988</a>	12966	465	5	150	bridging integrator 2 (Bin2) alternative variant dSep08, mRNA.
<a href="#">Bin2a</a>	<a href="#">Bin2a.bSep08</a>	<a href="#">494244</a>	24939	657	5	170	beta-galactosidase-like protein (Bin2a) alternative variant bSep08, mRNA.
<a href="#">Bin2a</a>	<a href="#">Bin2a.cSep08</a>	<a href="#">494244</a>	2659	402	2	51	beta-galactosidase-like protein (Bin2a) alternative variant cSep08, mRNA.
<a href="#">Bin3</a>	<a href="#">Bin3.bSep08</a>	<a href="#">361065</a>	38037	747	8	213	bridging integrator 3 (Bin3) alternative variant bSep08, mRNA.
<a href="#">Bin3</a>	<a href="#">Bin3.cSep08</a>	<a href="#">361065</a>	11127	832	5	198	bridging integrator 3 (Bin3) alternative variant cSep08, mRNA.
<a href="#">Bin3</a>	<a href="#">Bin3.dSep08</a>	<a href="#">361065</a>	4860	345	1	59	bridging integrator 3 (Bin3) alternative variant dSep08, mRNA.
<a href="#">BIR.0</a>	<a href="#">BIR.0.aSep08</a>		2230	364		121	neuronal apoptosis inhibitory protein 2 (BIR.0) mRNA.
<a href="#">BIR.1</a>	<a href="#">BIR.1.aSep08</a>		684	232		77	neuronal apoptosis inhibitory protein 2 (BIR.1) mRNA.
<a href="#">Birc2</a>	<a href="#">Birc2.bSep08</a>	<a href="#">60371</a>	21010	2443	7	227	baculoviral IAP repeat-containing 2 (25.6 kD) (Birc2) alternative variant bSep08, mRNA.
<a href="#">Birc2</a>	<a href="#">Birc2.cSep08</a>	<a href="#">60371</a>	16571	386	4	128	baculoviral IAP repeat-containing 2 (Birc2) alternative variant cSep08, mRNA.
<a href="#">Birc3</a>	<a href="#">Birc3.bSep08</a>	<a href="#">78971</a>	5401	1779	2	498	baculoviral IAP repeat-containing 3 (Birc3) alternative variant bSep08, mRNA.
<a href="#">Birc6</a>	<a href="#">Birc6.aSep08</a>	<a href="#">313876</a>	42696	3274		603	baculoviral IAP repeat-containing 6 (Birc6) mRNA.
<a href="#">Birc7</a>	<a href="#">Birc7.aSep08</a>	<a href="#">296468</a>	2974	694		164	baculoviral IAP repeat-containing 7 (livin) (Birc7) mRNA.

<a href="#">blabor</a>	<a href="#">blabor.aSep08</a>		18812	865		65	putative protein (7.3 kD) (blabor) mRNA.
<a href="#">blachy</a>	<a href="#">blachy.aSep08</a>		14505	822		57	ac1262 like (6.3 kD) (blachy) mRNA.
<a href="#">bladoy</a>	<a href="#">bladoy.aSep08</a>		757	638		97	putative protein (bladoy) mRNA.
<a href="#">blafly</a>	<a href="#">blafly.aSep08</a>		444	389		34	putative protein (3.6 kD) (blafly) mRNA.
<a href="#">blafly</a>	<a href="#">blafly.aSep08</a>		24564	438		56	putative protein (5.9 kD) (blafly) mRNA.
<a href="#">blagar</a>	<a href="#">blagar.aSep08</a>		8732	652	3	61	putative protein (blagar) alternative variant aSep08, mRNA.
<a href="#">blaja</a>	<a href="#">blaja.aSep08</a>		2647	229		43	putative protein (4.9 kD) (blaja) mRNA.
<a href="#">blajey</a>	<a href="#">blajey.aSep08</a>		1888	481	2	118	putative protein (blajey) alternative variant aSep08, mRNA.
<a href="#">blakee</a>	<a href="#">blakee.aSep08</a>		1981	358		38	mitochondrial -trna synthetase (blakee) mRNA.
<a href="#">blalo</a>	<a href="#">blalo.aSep08</a>		39157	714		72	putative nuclear protein (8.2 kD) (blalo) mRNA.
<a href="#">blamee</a>	<a href="#">blamee.aSep08</a>		965	692		52	putative protein (5.7 kD) (blamee) mRNA.
<a href="#">blanoy</a>	<a href="#">blanoy.aSep08</a>		5972	783		47	putative protein (blanoy) mRNA.
<a href="#">blapor</a>	<a href="#">blapor.aSep08</a>		5553	580		86	putative protein of mammalian origin (blapor) mRNA.
<a href="#">blarbor</a>	<a href="#">blarbor.aSep08</a>		5397	2235		75	putative protein (8.3 kD) (blarbor) mRNA.
<a href="#">blarchy</a>	<a href="#">blarchy.aSep08</a>		2048	757		44	putative protein (5.1 kD) (blarchy) mRNA.
<a href="#">blardoy</a>	<a href="#">blardoy.aSep08</a>		3170	764		54	putative protein (5.7 kD) (blardoy) mRNA.
<a href="#">blarflu</a>	<a href="#">blarflu.aSep08</a>		754	261		41	putative protein (blarflu) mRNA.
<a href="#">blarfly</a>	<a href="#">blarfly.aSep08</a>		3428	434		70	putative protein (7.6 kD) (blarfly) mRNA.
<a href="#">blargar</a>	<a href="#">blargar.aSep08</a>		4833	672		125	polyprotein (13.6 kD) (blargar) mRNA.
<a href="#">blarja</a>	<a href="#">blarja.aSep08</a>		8035	715	3	81	putative secreted or extracellular protein precursor (9.1 kD) (blarja) alternative variant aSep08, mRNA.
<a href="#">blarja</a>	<a href="#">blarja.bSep08</a>		14772	658	1	70	putative protein (7.6 kD) (blarja) alternative variant bSep08, mRNA.
<a href="#">blarjey</a>	<a href="#">blarjey.aSep08</a>		17701	344		74	putative mitochondrial protein (8.8 kD) (blarjey) mRNA.
<a href="#">blarkee</a>	<a href="#">blarkee.aSep08</a>		3552	749	3	125	putative protein (13.9 kD) (blarkee) alternative variant aSep08, mRNA.
<a href="#">blarlo</a>	<a href="#">blarlo.aSep08</a>		1644	362		34	putative protein (3.8 kD) (blarlo) mRNA.
<a href="#">blarmee</a>	<a href="#">blarmee.aSep08</a>		1383	493	2	108	putative protein (blarmee) alternative variant aSep08, mRNA.
<a href="#">blarmee</a>	<a href="#">blarmee.bSep08</a>		1903	455	3	37	putative protein (blarmee) alternative variant bSep08, mRNA.
<a href="#">blarnoy</a>	<a href="#">blarnoy.aSep08</a>		1901	673		28	putative protein (9.4 kD) (blarnoy) mRNA.
<a href="#">blaroy</a>	<a href="#">blaroy.aSep08</a>		1373	585	3	60	putative protein (blaroy) alternative variant aSep08, mRNA.
<a href="#">blaroy</a>	<a href="#">blaroy.bSep08</a>		1242	523	1	46	putative protein (5.1 kD) (blaroy) alternative variant bSep08, mRNA.
<a href="#">blarpor</a>	<a href="#">blarpor.aSep08</a>		882	695		110	voltage-dependent calcium channel (blarpor) mRNA.
<a href="#">blarroy</a>	<a href="#">blarroy.aSep08</a>		2971	398		66	putative protein (blarroy) mRNA.
<a href="#">blarsa</a>	<a href="#">blarsa.aSep08</a>		913	767		30	putative protein (blarsa) mRNA.
<a href="#">blarshee</a>	<a href="#">blarshee.aSep08</a>		1699	276		24	putative protein (blarshee) mRNA.
<a href="#">blartu</a>	<a href="#">blartu.aSep08</a>		14070	573	5	129	putative protein (blartu) alternative variant aSep08, mRNA.
<a href="#">blartu</a>	<a href="#">blartu.cSep08</a>		37290	1347	5	89	putative protein (9.9 kD) (blartu) alternative variant cSep08, mRNA.

<a href="#">blartu</a>	<a href="#">blartu.dSep08</a>		23044	994	7	89	putative protein (9.9 kD) (blartu) alternative variant dSep08, mRNA.
<a href="#">blarvo</a>	<a href="#">blarvo.aSep08</a>		5133	841		46	putative protein (5.3 kD) (blarvo) mRNA.
<a href="#">blarwer</a>	<a href="#">blarwer.aSep08</a>		661	188			
<a href="#">blarwey</a>	<a href="#">blarwey.aSep08</a>		35655	412		137	atp-binding cassette sub-family c member 9 like (blarwey) mRNA.
<a href="#">blasa</a>	<a href="#">blasa.aSep08</a>		3810	331		75	putative nuclear protein (8.7 kD) (blasa) mRNA.
<a href="#">blashee</a>	<a href="#">blashee.aSep08</a>		20407	1036		202	homeodomain transcription factor 1 CRA a (blashee) mRNA.
<a href="#">blatu</a>	<a href="#">blatu.aSep08</a>		12500	1036	5	209	putative cytoplasmic protein, with a coiled coil domain, of metazoan origin (23.8 kD) (blatu) alternative variant aSep08, mRNA.
<a href="#">blatu</a>	<a href="#">blatu.bSep08</a>		12166	537	3	150	putative protein, with a coiled coil domain, of vertebrate origin (blatu) alternative variant bSep08, mRNA.
<a href="#">blavo</a>	<a href="#">blavo.aSep08</a>		103671	514		41	CRA b like (4.9 kD) (blavo) mRNA.
<a href="#">blawbor</a>	<a href="#">blawbor.aSep08</a>		8584	422		140	collagen type VI alpha 2 CRA c (blawbor) mRNA.
<a href="#">blawchy</a>	<a href="#">blawchy.aSep08</a>		22502	402		64	CRA a like (blawchy) mRNA.
<a href="#">blawdoy</a>	<a href="#">blawdoy.aSep08</a>		1577	350		71	putative protein of vertebrate origin (blawdoy) mRNA.
<a href="#">blawer</a>	<a href="#">blawer.aSep08</a>		8587	578		138	F-box leucine-rich repeat protein 13 (blawer) mRNA.
<a href="#">blawey</a>	<a href="#">blawey.aSep08</a>		1308	621		159	putative protein (16.0 kD) (blawey) mRNA.
<a href="#">blawflu</a>	<a href="#">blawflu.aSep08</a>		5255	548		35	putative protein (blawflu) mRNA.
<a href="#">blawfly</a>	<a href="#">blawfly.aSep08</a>		2982	590		71	putative cytoplasmic protein (8.5 kD) (blawfly) mRNA.
<a href="#">blawgar</a>	<a href="#">blawgar.aSep08</a>		779	379		50	putative protein (blawgar) mRNA.
<a href="#">blawja</a>	<a href="#">blawja.aSep08</a>		4717	363		63	a disintegrin metalloproteinase domain 28 CRA c (blawja) mRNA.
<a href="#">blawjey</a>	<a href="#">blawjey.aSep08</a>		1945	734	4	145	sfi1 homolog spindle assembly associated (17.4 kD) (blawjey) alternative variant aSep08, mRNA.
<a href="#">blawjey</a>	<a href="#">blawjey.bSep08</a>		6236	352	5	117	sfi1 homolog spindle assembly associated (blawjey) alternative variant bSep08, mRNA.
<a href="#">blawkee</a>	<a href="#">blawkee.aSep08</a>		18375	264		43	putative protein (blawkee) mRNA.
<a href="#">blawlo</a>	<a href="#">blawlo.aSep08</a>		1369	786		49	putative protein (blawlo) mRNA.
<a href="#">blawmee</a>	<a href="#">blawmee.aSep08</a>		4493	605		70	putative protein (7.9 kD) (blawmee) mRNA.
<a href="#">blawnoy</a>	<a href="#">blawnoy.aSep08</a>		5158	616		92	putative mitochondrial protein (10.2 kD) (blawnoy) mRNA.
<a href="#">blawpor</a>	<a href="#">blawpor.aSep08</a>		5661	885		41	putative protein (blawpor) alternative variant aSep08, mRNA.
<a href="#">blawpor</a>	<a href="#">blawpor.cSep08</a>		4995	218		17	putative protein (2.1 kD) (blawpor) alternative variant cSep08, mRNA.
<a href="#">blawroy</a>	<a href="#">blawroy.aSep08</a>		602	421		90	putative secreted or extracellular protein precursor (9.5 kD) (blawroy) mRNA.
<a href="#">blawsa</a>	<a href="#">blawsa.aSep08</a>		2488	876		104	dip2 disco-interacting protein 2 homolog b (blawsa) mRNA.
<a href="#">blawshee</a>	<a href="#">blawshee.aSep08</a>		22562	409		117	putative protein (blawshee) mRNA.
<a href="#">blawtu</a>	<a href="#">blawtu.aSep08</a>		413	313		98	putative protein (blawtu) mRNA.
<a href="#">blawvo</a>	<a href="#">blawvo.aSep08</a>		989	314			
<a href="#">blawwer</a>	<a href="#">blawwer.aSep08</a>		1828	388		32	putative protein (3.6 kD) (blawwer) mRNA.

<a href="#">blawwey</a>	<a href="#">blawwey.aSep08</a>		3545	358		119	glycogen synthase (blawwey) mRNA.
<a href="#">bleebor</a>	<a href="#">bleebor.aSep08</a>		6406	407		135	pericentrin (bleebor) mRNA.
<a href="#">bleechy</a>	<a href="#">bleechy.aSep08</a>		2404	738		57	putative protein (6.5 kD) (bleechy) mRNA.
<a href="#">bleedoy</a>	<a href="#">bleedoy.aSep08</a>		2116	652		82	putative protein (9.1 kD) (bleedoy) mRNA.
<a href="#">bleeflu</a>	<a href="#">bleeflu.aSep08</a>		5932	523		55	putative protein (5.9 kD) (bleeflu) mRNA.
<a href="#">bleefly</a>	<a href="#">bleefly.aSep08</a>		24310	859		160	putative protein (bleefly) mRNA.
<a href="#">bleegar</a>	<a href="#">bleegar.aSep08</a>		2995	763		127	CRA a like (bleegar) mRNA.
<a href="#">bleeja</a>	<a href="#">bleeja.aSep08</a>		10801	367		44	putative protein (bleeja) mRNA.
<a href="#">bleejey</a>	<a href="#">bleejey.aSep08</a>		9185	706		80	sfi1 homolog spindle assembly associated (bleejey) mRNA.
<a href="#">bleekee</a>	<a href="#">bleekee.aSep08</a>		19262	639		58	putative protein (bleekee) mRNA.
<a href="#">bleelo</a>	<a href="#">bleelo.aSep08</a>		5648	789		92	putative protein (9.9 kD) (bleelo) mRNA.
<a href="#">bleemee</a>	<a href="#">bleemee.aSep08</a>		33798	673		7	putative protein (0.7 kD) (bleemee) mRNA.
<a href="#">bleenoy</a>	<a href="#">bleenoy.bSep08</a>		1564	308	2	37	putative protein (bleenoy) alternative variant bSep08, mRNA.
<a href="#">bleepor</a>	<a href="#">bleepor.bSep08</a>		1136	501	2	63	putative protein of mammalian origin (bleepor) alternative variant bSep08, mRNA.
<a href="#">bleeroy</a>	<a href="#">bleeroy.bSep08</a>		1605	446	2	64	putative protein (bleeroy) alternative variant bSep08, mRNA.
<a href="#">bleesa</a>	<a href="#">bleesa.aSep08</a>		18806	772		257	solute carrier family 11 member 2 CRA c (bleesa) alternative variant aSep08, mRNA.
<a href="#">bleshee</a>	<a href="#">bleshee.bSep08</a>		1566	411	2	30	putative protein (3.6 kD) (bleshee) alternative variant bSep08, mRNA.
<a href="#">bleetu</a>	<a href="#">bleetu.aSep08</a>		5218	613		75	putative cytoplasmic protein of ancient origin (8.9 kD) (bleetu) mRNA.
<a href="#">bleevo</a>	<a href="#">bleevo.aSep08</a>		4433	257		36	putative protein (bleevo) mRNA.
<a href="#">bleewer</a>	<a href="#">bleewer.aSep08</a>		28499	745		98	putative protein (bleewer) mRNA.
<a href="#">blerbor</a>	<a href="#">blerbor.aSep08</a>		2181	572		190	pericentrin (blerbor) mRNA.
<a href="#">blerchy</a>	<a href="#">blerchy.aSep08</a>		4823	973		100	csrp2 binding protein like (11.2 kD) (blerchy) mRNA.
<a href="#">blerdoy</a>	<a href="#">blerdoy.aSep08</a>		1192	246		72	putative protein (blerdoy) mRNA.
<a href="#">blerflu</a>	<a href="#">blerflu.aSep08</a>		7035	678		50	putative protein (blerflu) mRNA.
<a href="#">blerfly</a>	<a href="#">blerfly.aSep08</a>		2400	424		74	putative protein (8.5 kD) (blerfly) mRNA.
<a href="#">blergar</a>	<a href="#">blergar.aSep08</a>		5712	1173		235	putative protein (blergar) alternative variant aSep08, mRNA.
<a href="#">blerja</a>	<a href="#">blerja.aSep08</a>		26572	767	2	67	putative protein (blerja) alternative variant aSep08, mRNA.
<a href="#">blerjey</a>	<a href="#">blerjey.aSep08</a>		43627	781		78	putative protein (blerjey) mRNA.
<a href="#">blerkee</a>	<a href="#">blerkee.aSep08</a>		1111	415		46	putative protein (4.8 kD) (blerkee) mRNA.
<a href="#">blerlo</a>	<a href="#">blerlo.aSep08</a>		1005	728	2	90	dead box polypeptide 55 (blerlo) alternative variant aSep08, mRNA.
<a href="#">blermee</a>	<a href="#">blermee.aSep08</a>		1813	1107		187	putative protein (20.3 kD) (blermee) mRNA.
<a href="#">blernoy</a>	<a href="#">blernoy.aSep08</a>		56799	565		69	putative protein of mammalian origin (blernoy) mRNA.
<a href="#">blerpor</a>	<a href="#">blerpor.aSep08</a>		4121	453		30	putative protein (blerpor) mRNA.
<a href="#">blerroy</a>	<a href="#">blerroy.aSep08</a>		789	684		56	putative protein (blerroy) mRNA.

<a href="#">blersa</a>	<a href="#">blersa.aSep08</a>		12789	516	4	122	POU domain-containing transcription factor (blersa) alternative variant aSep08, mRNA.
<a href="#">blersa</a>	<a href="#">blersa.cSep08</a>		5826	328	2	67	CRA a like (blersa) alternative variant cSep08, mRNA.
<a href="#">blershee</a>	<a href="#">blershee.aSep08</a>		8403	284		25	putative protein (blershee) mRNA.
<a href="#">blertu</a>	<a href="#">blertu.aSep08</a>		1437	698		53	putative protein (blertu) mRNA.
<a href="#">blervo</a>	<a href="#">blervo.aSep08</a>		5888	400		133	CRA a (blervo) mRNA.
<a href="#">blerwer</a>	<a href="#">blerwer.aSep08</a>		1289	652		42	putative protein (5.0 kD) (blerwer) mRNA.
<a href="#">Bles03</a>	<a href="#">Bles03.aSep08</a>	<a href="#">266609</a>	2251	1595	2	292	basophilic leukemia expressed protein BLES03 (31.4 kD) (Bles03) alternative variant aSep08, mRNA.
<a href="#">bleybor</a>	<a href="#">bleybor.aSep08</a>		7242	944		314	pericentrin (bleybor) mRNA.
<a href="#">bleychy</a>	<a href="#">bleychy.aSep08</a>		13422	421		140	csrp2 binding protein CRA b like (bleychy) mRNA.
<a href="#">bleydoy</a>	<a href="#">bleydoy.aSep08</a>		42634	736		45	putative protein (5.2 kD) (bleydoy) mRNA.
<a href="#">bleyflu</a>	<a href="#">bleyflu.aSep08</a>		4515	564		98	putative protein (10.9 kD) (bleyflu) mRNA.
<a href="#">bleyfly</a>	<a href="#">bleyfly.aSep08</a>		24741	602		160	T-cell lymphoma invasion metastasis 2 like (bleyfly) mRNA.
<a href="#">bleygar</a>	<a href="#">bleygar.aSep08</a>		314	204		67	g-protein coupled receptor 124 (bleygar) mRNA.
<a href="#">bleyja</a>	<a href="#">bleyja.aSep08</a>		1083	394		64	putative protein (bleyja) mRNA.
<a href="#">bleyjey</a>	<a href="#">bleyjey.aSep08</a>		3685	486		31	putative protein (3.5 kD) (bleyjey) mRNA.
<a href="#">bleykee</a>	<a href="#">bleykee.aSep08</a>		1199	729		62	putative protein (bleykee) mRNA.
<a href="#">bleylo</a>	<a href="#">bleylo.aSep08</a>		4981	657		199	M-phase phosphoprotein 9 (bleylo) mRNA.
<a href="#">bleymee</a>	<a href="#">bleymee.aSep08</a>		4553	1954		246	phospholipase D2 (bleymee) mRNA.
<a href="#">bleynoy</a>	<a href="#">bleynoy.aSep08</a>		44372	684		69	CRA a (7.8 kD) (bleynoy) mRNA.
<a href="#">bleypor</a>	<a href="#">bleypor.aSep08</a>		3967	370		55	putative protein (6.4 kD) (bleypor) mRNA.
<a href="#">bleyroy</a>	<a href="#">bleyroy.aSep08</a>		6041	1384		461	ATPase class I type 8B member 3 (bleyroy) mRNA.
<a href="#">bleysa</a>	<a href="#">bleysa.aSep08</a>		1533	311		98	putative protein (bleysa) mRNA.
<a href="#">bleyshee</a>	<a href="#">bleyshee.aSep08</a>		1985	1010	4	145	putative protein of mammalian origin (bleyshee) alternative variant aSep08, mRNA.
<a href="#">bleyshee</a>	<a href="#">bleyshee.bSep08</a>		2160	384	4	116	putative protein of mammalian origin (bleyshee) alternative variant bSep08, mRNA.
<a href="#">bleyshee</a>	<a href="#">bleyshee.cSep08</a>		4957	640	3	73	CRA b like (7.5 kD) (bleyshee) alternative variant cSep08, mRNA.
<a href="#">bleytu</a>	<a href="#">bleytu.aSep08</a>		7706	317	2	105	ctage family member (bleytu) alternative variant aSep08, mRNA.
<a href="#">bleytu</a>	<a href="#">bleytu.bSep08</a>		9287	293	1	60	meningioma expressed antigen 6 CRA b like (bleytu) alternative variant bSep08, mRNA.
<a href="#">bleyvo</a>	<a href="#">bleyvo.aSep08</a>		1836	687		88	putative protein (9.5 kD) (bleyvo) mRNA.
<a href="#">bleywer</a>	<a href="#">bleywer.aSep08</a>		76713	403		40	calcium channel voltage-dependent CRA c (bleywer) mRNA.
<a href="#">Blk</a>	<a href="#">Blk.bSep08</a>	<a href="#">364403</a>	1829	688	2	65	B lymphoid kinase (Blk) alternative variant bSep08, mRNA.
<a href="#">Blm</a>	<a href="#">Blm.bSep08</a>	<a href="#">308755</a>	29653	1844	10	551	bloom syndrome homolog (human) (Blm) alternative variant bSep08, mRNA.
<a href="#">Blmh</a>	<a href="#">Blmh.bSep08</a>	<a href="#">287552</a>	687	386	2	128	bleomycin hydrolase (Blmh) alternative variant bSep08, mRNA.
<a href="#">Blmh</a>	<a href="#">Blmh.cSep08</a>	<a href="#">287552</a>	7013	574	5	93	bleomycin hydrolase (Blmh) alternative variant cSep08, mRNA.

<a href="#">Blnk</a>	<a href="#">Blnk.aSep08</a>	<a href="#">499356</a>	56653	1090	12	362	B-cell linker (Blnk) alternative variant aSep08, mRNA.
<a href="#">Blnk</a>	<a href="#">Blnk.bSep08</a>	<a href="#">499356</a>	18879	882	7	135	putative protein (Blnk) alternative variant bSep08, mRNA.
<a href="#">Blnk</a>	<a href="#">Blnk.cSep08</a>	<a href="#">499356</a>	6877	736	2	42	B-cell linker protein like (5.1 kD) (Blnk) alternative variant cSep08, mRNA.
<a href="#">blobor</a>	<a href="#">blobor.aSep08</a>		2432	507		141	putative nuclear protein (15.6 kD) (blobor) mRNA.
<a href="#">Bloc1s1</a>	<a href="#">Bloc1s1.bSep08</a>	<a href="#">288785</a>	1124	649	1	75	biogenesis of lysosome-related organelles complex-1, subunit 1 (8.6 kD) (Bloc1s1) alternative variant bSep08, mRNA.
<a href="#">Bloc1s2</a>	<a href="#">Bloc1s2.aSep08</a>	<a href="#">293938</a>	7025	929	6	168	biogenesis of lysosome-related organelles complex-1, subunit 2 (18.9 kD) (Bloc1s2) alternative variant aSep08, complete mRNA.
<a href="#">Bloc1s2</a>	<a href="#">Bloc1s2.bSep08</a>	<a href="#">293938</a>	3693	737	3	144	biogenesis of lysosome-related organelles complex-1, subunit 2 (Bloc1s2) alternative variant bSep08, mRNA.
<a href="#">Bloc1s2</a>	<a href="#">Bloc1s2.dSep08</a>	<a href="#">293938</a>	7018	878	5	140	biogenesis of lysosome-related organelles complex-1, subunit 2 (15.7 kD) (Bloc1s2) alternative variant dSep08, mRNA.
<a href="#">Bloc1s2</a>	<a href="#">Bloc1s2.eSep08</a>	<a href="#">293938</a>	6923	633	5	140	biogenesis of lysosome-related organelles complex-1, subunit 2 (15.6 kD) (Bloc1s2) alternative variant eSep08, mRNA.
<a href="#">Bloc1s2</a>	<a href="#">Bloc1s2.fSep08</a>	<a href="#">293938</a>	6750	856	4	99	biogenesis of lysosome-related organelles complex-1, subunit 2 (11.5 kD) (Bloc1s2) alternative variant fSep08, mRNA.
<a href="#">Bloc1s3</a>	<a href="#">Bloc1s3.aSep08</a>	<a href="#">680476</a>	1655	832		212	biogenesis of lysosome-related organelles complex-1, subunit 3 (Bloc1s3) alternative variant aSep08, mRNA.
<a href="#">blochy</a>	<a href="#">blochy.aSep08</a>		45873	279		92	putative protein of mammalian origin (blochy) mRNA.
<a href="#">blodoy</a>	<a href="#">blodoy.aSep08</a>		11786	385	1	84	putative protein (9.6 kD) (blodoy) alternative variant aSep08, mRNA.
<a href="#">blodoy</a>	<a href="#">blodoy.bSep08</a>		12841	353	1	37	putative protein (4.3 kD) (blodoy) alternative variant bSep08, mRNA.
<a href="#">bloflu</a>	<a href="#">bloflu.aSep08</a>		16289	361		119	TEA domain family member 1 (bloflu) mRNA.
<a href="#">blofly</a>	<a href="#">blofly.aSep08</a>		1584	807		70	GTPase activating protein testicular GAP1 like (8.0 kD) (blofly) mRNA.
<a href="#">blogar</a>	<a href="#">blogar.aSep08</a>		1512	1007		46	putative protein (5.1 kD) (blogar) mRNA.
<a href="#">bloja</a>	<a href="#">bloja.aSep08</a>		512860	293		46	putative protein (5.0 kD) (bloja) mRNA.
<a href="#">blojey</a>	<a href="#">blojey.aSep08</a>		3067	199		63	gag protein like (blojey) mRNA.
<a href="#">blokee</a>	<a href="#">blokee.aSep08</a>		10585	601		64	putative protein (blokee) mRNA.
<a href="#">blolo</a>	<a href="#">blolo.aSep08</a>		4056	328	2	109	putative protein (blolo) alternative variant aSep08, mRNA.
<a href="#">blolo</a>	<a href="#">blolo.bSep08</a>		4074	478	3	77	putative protein (8.7 kD) (blolo) alternative variant bSep08, mRNA.
<a href="#">blomee</a>	<a href="#">blomee.aSep08</a>		798	697		77	putative mitochondrial protein (8.9 kD) (blomee) mRNA.
<a href="#">blonoy</a>	<a href="#">blonoy.aSep08</a>		13844	756	5	103	putative protein (11.5 kD) (blonoy) alternative variant aSep08, mRNA.
<a href="#">blonoy</a>	<a href="#">blonoy.bSep08</a>		12189	769	7	55	putative protein (blonoy) alternative variant bSep08, mRNA.
<a href="#">blopor</a>	<a href="#">blopor.aSep08</a>		3977	402		134	vpr binding protein like (blopor) mRNA.

<a href="#">blorbor</a>	<a href="#">blorbor.aSep08</a>		6392	377		125	disco-interacting protein 2 homolog a (blorbor) mRNA.
<a href="#">blorchy</a>	<a href="#">blorchy.aSep08</a>		3663	800	2	123	putative protein (14.0 kD) (blorchy) alternative variant aSep08, mRNA.
<a href="#">blordoy</a>	<a href="#">blordoy.aSep08</a>		9010	346		114	putative protein of mammalian origin (blordoy) mRNA.
<a href="#">blorflu</a>	<a href="#">blorflu.aSep08</a>		3074	573		190	PI-3-kinase-related kinase smg-1 (blorflu) mRNA.
<a href="#">blorfly</a>	<a href="#">blorfly.aSep08</a>		5058	971	6	323	T-cell lymphoma invasion metastasis 2 like (blorfly) alternative variant aSep08, mRNA.
<a href="#">blorgar</a>	<a href="#">blorgar.aSep08</a>		3308	1085		171	G-protein coupled receptor 124 (blorgar) mRNA.
<a href="#">blorja</a>	<a href="#">blorja.aSep08</a>		3779	239		40	putative protein (blorja) mRNA.
<a href="#">blorjey</a>	<a href="#">blorjey.aSep08</a>		472	389		82	putative protein (blorjey) mRNA.
<a href="#">blorkee</a>	<a href="#">blorkee.aSep08</a>		4865	551		77	putative protein (blorkee) mRNA.
<a href="#">blorlo</a>	<a href="#">blorlo.aSep08</a>		1113	481		57	putative protein (blorlo) mRNA.
<a href="#">blormee</a>	<a href="#">blormee.aSep08</a>		5920	1212	2	403	misshapen-like kinase 1 (blormee) alternative variant aSep08, mRNA.
<a href="#">blormee</a>	<a href="#">blormee.bSep08</a>		1805	709	1	235	misshapen-like kinase 1 (blormee) alternative variant bSep08, mRNA.
<a href="#">blornoy</a>	<a href="#">blornoy.aSep08</a>		6651	575		44	putative protein (blornoy) mRNA.
<a href="#">bloroy</a>	<a href="#">bloroy.aSep08</a>		995	655		29	putative protein (3.5 kD) (bloroy) mRNA.
<a href="#">blorpor</a>	<a href="#">blorpor.aSep08</a>		4312	432		100	putative protein (blorpor) mRNA.
<a href="#">blorroy</a>	<a href="#">blorroy.aSep08</a>		682	459		84	putative protein (blorroy) mRNA.
<a href="#">blorsa</a>	<a href="#">blorsa.aSep08</a>		12129	745		105	N-acetylgalactosaminyltransferase (blorsa) mRNA.
<a href="#">blorshee</a>	<a href="#">blorshee.aSep08</a>		78234	1191	7	232	adenosine A3 receptor (25.7 kD) (blorshee) alternative variant aSep08, complete mRNA.
<a href="#">blorshee</a>	<a href="#">blorshee.bSep08</a>		8279	842	4	67	adenosine A3 receptor like (blorshee) alternative variant bSep08, mRNA.
<a href="#">blorshee</a>	<a href="#">blorshee.cSep08</a>		50744	751	4	96	adenosine A3 receptor (blorshee) alternative variant cSep08, mRNA.
<a href="#">blorshee</a>	<a href="#">blorshee.dSep08</a>		836	244	2	41	putative protein (4.8 kD) (blorshee) alternative variant dSep08, mRNA.
<a href="#">blorvo</a>	<a href="#">blorvo.aSep08</a>		1205	352		44	putative protein (4.8 kD) (blorvo) mRNA.
<a href="#">blorwer</a>	<a href="#">blorwer.aSep08</a>		1203	771		77	CRA d like (8.4 kD) (blorwer) mRNA.
<a href="#">blosa</a>	<a href="#">blosa.aSep08</a>		26145	380		126	dip2 disco-interacting protein 2 homolog (blosa) mRNA.
<a href="#">bloshee</a>	<a href="#">bloshee.aSep08</a>		6142	604		56	putative protein (6.2 kD) (bloshee) mRNA.
<a href="#">blotu</a>	<a href="#">blotu.aSep08</a>		810	441		72	putative protein (blotu) mRNA.
<a href="#">blovo</a>	<a href="#">blovo.aSep08</a>		7994	869		46	putative protein (blovo) mRNA.
<a href="#">blower</a>	<a href="#">blower.aSep08</a>		10514	260		62	putative protein (blower) mRNA.
<a href="#">blowey</a>	<a href="#">blowey.aSep08</a>		2285	475	2	111	putative protein (blowey) alternative variant aSep08, mRNA.
<a href="#">bloybor</a>	<a href="#">bloybor.aSep08</a>		5262	2174		211	CRA b (bloybor) mRNA.
<a href="#">bloychy</a>	<a href="#">bloychy.aSep08</a>		14541	515	5	171	putative protein of eukaryotic origin (bloychy) alternative variant aSep08, mRNA.
<a href="#">bloydoy</a>	<a href="#">bloydoy.aSep08</a>		38857	525	1	71	putative protein (bloydoy) alternative variant aSep08, mRNA.

<a href="#">bloydoy</a>	<a href="#">bloydoy.bSep08</a>		13078	249	1	49	putative protein (bloydoy) alternative variant bSep08, mRNA.
<a href="#">bloyflu</a>	<a href="#">bloyflu.aSep08</a>		2633	371		123	kinase (bloyflu) mRNA.
<a href="#">bloyfly</a>	<a href="#">bloyfly.aSep08</a>		14432	403		81	putative protein (bloyfly) mRNA.
<a href="#">bloygar</a>	<a href="#">bloygar.aSep08</a>		18541	568	3	48	CRA a like (5.3 kD) (bloygar) alternative variant aSep08, mRNA.
<a href="#">bloygar</a>	<a href="#">bloygar.bSep08</a>		6894	397	1	71	CRA b like (bloygar) alternative variant bSep08, mRNA.
<a href="#">bloyja</a>	<a href="#">bloyja.aSep08</a>		3479	1755	8	342	putative protein, with a coiled coil domain, of bilateral origin (bloyja) alternative variant aSep08, mRNA.
<a href="#">bloyja</a>	<a href="#">bloyja.bSep08</a>		2807	1609	7	296	putative protein, with a coiled coil domain, of bilateral origin (bloyja) alternative variant bSep08, mRNA.
<a href="#">bloyja</a>	<a href="#">bloyja.cSep08</a>		1680	1272	4	176	putative protein, with a coiled coil domain, of vertebrate origin (20.0 kD) (bloyja) alternative variant cSep08, mRNA.
<a href="#">bloyjey</a>	<a href="#">bloyjey.aSep08</a>		9160	425		59	putative protein (6.8 kD) (bloyjey) mRNA.
<a href="#">bloykee</a>	<a href="#">bloykee.aSep08</a>		984	509		130	putative protein of mammalian origin (bloykee) mRNA.
<a href="#">bloylo</a>	<a href="#">bloylo.aSep08</a>		1882	451		149	DDHD (bloylo) mRNA.
<a href="#">bloymee</a>	<a href="#">bloymee.aSep08</a>		961	693	2	221	CRA c (bloymee) alternative variant aSep08, mRNA.
<a href="#">bloymee</a>	<a href="#">bloymee.bSep08</a>		674	388	1	128	CRA b (bloymee) alternative variant bSep08, mRNA.
<a href="#">bloynoy</a>	<a href="#">bloynoy.aSep08</a>		828	729		78	putative protein (bloynoy) mRNA.
<a href="#">bloypor</a>	<a href="#">bloypor.aSep08</a>		2230	618	5	194	CRA d (bloypor) alternative variant aSep08, mRNA.
<a href="#">bloyroy</a>	<a href="#">bloyroy.aSep08</a>		2108	685	4	167	midnolin (bloyroy) alternative variant aSep08, mRNA.
<a href="#">bloysa</a>	<a href="#">bloysa.aSep08</a>		4816	340		28	putative protein (3.4 kD) (bloysa) mRNA.
<a href="#">bloyshee</a>	<a href="#">bloyshee.aSep08</a>		27275	716		36	putative protein (3.8 kD) (bloyshee) mRNA.
<a href="#">bloytu</a>	<a href="#">bloytu.aSep08</a>		14973	293		97	fam179b (bloytu) mRNA.
<a href="#">bloyvo</a>	<a href="#">bloyvo.aSep08</a>		3821	313		33	putative protein (bloyvo) mRNA.
<a href="#">bloywer</a>	<a href="#">bloywer.aSep08</a>		2755	573		40	putative protein (bloywer) mRNA.
<a href="#">blubor</a>	<a href="#">blubor.aSep08</a>		15166	636		70	putative protein (blubor) mRNA.
<a href="#">bluchy</a>	<a href="#">bluchy.aSep08</a>		38524	808	2	127	CRA a (14.5 kD) (bluchy) alternative variant aSep08, mRNA.
<a href="#">bluchy</a>	<a href="#">bluchy.bSep08</a>		70757	274	2	91	putative protein of vertebrate origin (bluchy) alternative variant bSep08, mRNA.
<a href="#">bludoy</a>	<a href="#">bludoy.bSep08</a>		1606	629	2	20	putative protein (2.2 kD) (bludoy) alternative variant bSep08, mRNA.
<a href="#">blufly</a>	<a href="#">blufly.aSep08</a>		5487	1345		79	putative protein (blufly) mRNA.
<a href="#">blufly</a>	<a href="#">blufly.aSep08</a>		2374	523	1	173	putative protein of mammalian origin (blufly) alternative variant aSep08, mRNA.
<a href="#">blufly</a>	<a href="#">blufly.bSep08</a>		32638	901	6	100	antigen-like 2 (blufly) alternative variant bSep08, mRNA.
<a href="#">blugar</a>	<a href="#">blugar.aSep08</a>		16594	640		117	werner syndrome like (blugar) mRNA.
<a href="#">bluja</a>	<a href="#">bluja.aSep08</a>		2130	526	1	39	CRA a like (4.4 kD) (bluja) alternative variant aSep08, mRNA.
<a href="#">bluja</a>	<a href="#">bluja.bSep08</a>		2556	528	1	31	CRA b like (3.3 kD) (bluja) alternative variant bSep08, mRNA.
<a href="#">blujey</a>	<a href="#">blujey.aSep08</a>		10992	722		90	putative protein (10.5 kD) (blujey) mRNA.
<a href="#">blukee</a>	<a href="#">blukee.aSep08</a>		5633	615		41	putative protein (blukee) mRNA.



<a href="#">blulo</a>	<a href="#">blulo.aSep08</a>		5624	313		40	putative protein (4.2 kD) (blulo) mRNA.
<a href="#">blumee</a>	<a href="#">blumee.aSep08</a>		1631	1454	2	83	Y box protein 2 (blumee) alternative variant aSep08, mRNA.
<a href="#">blumee</a>	<a href="#">blumee.bSep08</a>		1351	774	2	71	putative protein of metazoan origin (blumee) alternative variant bSep08, mRNA.
<a href="#">blunoy</a>	<a href="#">blunoy.aSep08</a>		12989	415		83	uncharacterized protein like (blunoy) mRNA.
<a href="#">blupor</a>	<a href="#">blupor.aSep08</a>		3450	302		100	vpr binding protein like (blupor) mRNA.
<a href="#">bluroy</a>	<a href="#">bluroy.aSep08</a>		2043	413	2	66	putative protein (bluroy) alternative variant aSep08, mRNA.
<a href="#">blusa</a>	<a href="#">blusa.aSep08</a>		10476	293	2	62	uncharacterized protein like (blusa) alternative variant aSep08, mRNA.
<a href="#">blushee</a>	<a href="#">blushee.aSep08</a>		11193	397		78	putative secreted or extracellular protein precursor (8.6 kD) (blushee) mRNA.
<a href="#">blutu</a>	<a href="#">blutu.aSep08</a>		60238	816		271	GTPase activating RanGAP domain-like 1 (blutu) mRNA.
<a href="#">bluvo</a>	<a href="#">bluvo.aSep08</a>		1763	766		42	putative protein (bluvo) mRNA.
<a href="#">bluwer</a>	<a href="#">bluwer.aSep08</a>		717	332		36	putative protein (4.3 kD) (bluwer) mRNA.
<a href="#">bluwey</a>	<a href="#">bluwey.aSep08</a>		37962	906		68	repeat containing 1 like (bluwey) mRNA.
<a href="#">BlvrB</a>	<a href="#">BlvrB.bSep08</a>	<a href="#">292737</a>	17586	716	3	200	biliverdin reductase B (flavin reductase (NADPH)) (BlvrB) alternative variant bSep08, mRNA.
<a href="#">BlvrB</a>	<a href="#">BlvrB.cSep08</a>	<a href="#">292737</a>	4919	553	2	120	biliverdin reductase B (flavin reductase (NADPH)) (13.4 kD) (BlvrB) alternative variant cSep08, mRNA.
<a href="#">blybor</a>	<a href="#">blybor.aSep08</a>		597	416		45	putative protein (5.3 kD) (blybor) mRNA.
<a href="#">blychy</a>	<a href="#">blychy.aSep08</a>		11693	1204		43	putative protein (blychy) mRNA.
<a href="#">blydoy</a>	<a href="#">blydoy.aSep08</a>		10151	746		180	putative protein (blydoy) mRNA.
<a href="#">blyflu</a>	<a href="#">blyflu.aSep08</a>		1785	333		52	putative protein (6.1 kD) (blyflu) mRNA.
<a href="#">blyfly</a>	<a href="#">blyfly.aSep08</a>		1125	201		35	putative protein (blyfly) mRNA.
<a href="#">blygar</a>	<a href="#">blygar.aSep08</a>		4213	355	4	64	putative protein (blygar) alternative variant aSep08, mRNA.
<a href="#">blyja</a>	<a href="#">blyja.aSep08</a>		1041	570		25	putative protein (blyja) mRNA.
<a href="#">blyjey</a>	<a href="#">blyjey.aSep08</a>		25792	473		63	putative protein (7.4 kD) (blyjey) mRNA.
<a href="#">blykee</a>	<a href="#">blykee.aSep08</a>		7899	580		119	putative protein (13.1 kD) (blykee) mRNA.
<a href="#">blylo</a>	<a href="#">blylo.aSep08</a>		554	442		78	putative nuclear protein (9.1 kD) (blylo) mRNA.
<a href="#">blymee</a>	<a href="#">blymee.aSep08</a>		833	327		108	polymerase II (blymee) mRNA.
<a href="#">blynoy</a>	<a href="#">blynoy.aSep08</a>		6216	580		127	uncharacterized protein like (blynoy) mRNA.
<a href="#">blypor</a>	<a href="#">blypor.aSep08</a>		778	270		29	putative protein (3.0 kD) (blypor) mRNA.
<a href="#">blyroy</a>	<a href="#">blyroy.aSep08</a>		1637	701		89	putative protein (blyroy) mRNA.
<a href="#">blysa</a>	<a href="#">blysa.aSep08</a>		27388	387		67	putative protein (blysa) mRNA.
<a href="#">blyshee</a>	<a href="#">blyshee.aSep08</a>		28354	570		131	putative protein (blyshee) mRNA.
<a href="#">blytu</a>	<a href="#">blytu.aSep08</a>		38604	1100	7	366	GTPase activating RanGAP domain-like 1 (blytu) alternative variant aSep08, mRNA.
<a href="#">blyvo</a>	<a href="#">blyvo.aSep08</a>		3347	343		69	putative protein (blyvo) mRNA.
<a href="#">blywer</a>	<a href="#">blywer.aSep08</a>		1948	1342		48	homeodomain transcription factor 2 CRA c (blywer) mRNA.
<a href="#">blywey</a>	<a href="#">blywey.aSep08</a>		1216	236		72	putative protein (blywey) mRNA.
<a href="#">Bmi1</a>	<a href="#">Bmi1.bSep08</a>	<a href="#">307151</a>	2959	2728	1	66	bmi1 polycomb ring finger oncogene (Bmi1) alternative variant bSep08, mRNA.

<a href="#">Bmp1</a>	<a href="#">Bmp1.aSep08</a>	<a href="#">83470</a>	32520	3047	17	829	bone morphogenetic protein 1 (Bmp1) alternative variant aSep08, mRNA.
<a href="#">Bmp1</a>	<a href="#">Bmp1.bSep08</a>	<a href="#">83470</a>	11490	1450	6	212	bone morphogenetic protein 1 (Bmp1) alternative variant bSep08, mRNA.
<a href="#">Bmp1</a>	<a href="#">Bmp1.cSep08</a>	<a href="#">83470</a>	11671	721	5	199	bone morphogenetic protein 1 (Bmp1) alternative variant cSep08, mRNA.
<a href="#">Bmp2k</a>	<a href="#">Bmp2k.aSep08</a>	<a href="#">498333</a>	62926	2972	11	322	BMP-2 inducible kinase (Bmp2k) alternative variant aSep08, mRNA.
<a href="#">Bmp2k</a>	<a href="#">Bmp2k.bSep08</a>	<a href="#">498333</a>	30713	2341	6	183	BMP-2 inducible kinase (Bmp2k) alternative variant bSep08, mRNA.
<a href="#">Bmp2k</a>	<a href="#">Bmp2k.cSep08</a>	<a href="#">498333</a>	15585	708	1	116	BMP-2 inducible kinase (Bmp2k) alternative variant cSep08, mRNA.
<a href="#">Bmp4</a>	<a href="#">Bmp4.bSep08</a>	<a href="#">25296</a>	4650	601	3	69	bone morphogenetic protein 4 (Bmp4) alternative variant bSep08, mRNA.
<a href="#">Bmp6</a>	<a href="#">Bmp6.bSep08</a>	<a href="#">25644</a>	15872	2624	2	228	bone morphogenetic protein 6 (Bmp6) alternative variant bSep08, mRNA.
<a href="#">Bmp7</a>	<a href="#">Bmp7.aSep08</a>	<a href="#">85272</a>	10202	1274		180	bone morphogenetic protein 7 (Bmp7) mRNA.
<a href="#">Bmper</a>	<a href="#">Bmper.aSep08</a>	<a href="#">300455</a>	70851	744		178	BMP-binding endothelial regulator (Bmper) mRNA.
<a href="#">Bmpr1a</a>	<a href="#">Bmpr1a.bSep08</a>	<a href="#">81507</a>	483	351	2	77	bone morphogenetic protein receptor, type 1A (8.7 kD) (Bmpr1a) alternative variant bSep08, mRNA.
<a href="#">Bmpr2</a>	<a href="#">Bmpr2.aSep08</a>	<a href="#">140590</a>	109512	2772		718	bone morphogenetic protein receptor, type II (serine/threonine kinase) (Bmpr2) alternative variant aSep08, mRNA.
<a href="#">Bms1</a>	<a href="#">Bms1.aSep08</a>	<a href="#">362426</a>	26129	2968		897	BMS1 homolog, ribosome assembly protein (yeast) (Bms1) alternative variant aSep08, mRNA.
<a href="#">Bms1</a>	<a href="#">Bms1.bSep08</a>	<a href="#">362426</a>	1109	662		128	BMS1 homolog, ribosome assembly protein (yeast) (Bms1) alternative variant bSep08, mRNA.
<a href="#">Bmx</a>	<a href="#">Bmx.bSep08</a>	<a href="#">367786</a>	12696	905		301	BMX non-receptor tyrosine kinase (Bmx) alternative variant bSep08, mRNA.
<a href="#">Bnc2</a>	<a href="#">Bnc2.bSep08</a>	<a href="#">298189</a>	9445	610	1	135	basonuclin 2 and hypothetical protein LOC678750 (Bnc2) alternative variant bSep08, mRNA.
<a href="#">Bnc2</a>	<a href="#">Bnc2.bSep08</a>	<a href="#">678750</a>	9445	610	1	135	basonuclin 2 and hypothetical protein LOC678750 (Bnc2) alternative variant bSep08, mRNA.
<a href="#">Bnip1</a>	<a href="#">Bnip1.bSep08</a>	<a href="#">140932</a>	11991	916	4	209	BCL2/adenovirus E1B 19kDa-interacting protein 1 (23.9 kD) (Bnip1) alternative variant bSep08, complete mRNA.
<a href="#">Bnip1</a>	<a href="#">Bnip1.cSep08</a>	<a href="#">140932</a>	10866	682	3	192	BCL2/adenovirus E1B 19kDa-interacting protein 1 (Bnip1) alternative variant cSep08, mRNA.
<a href="#">Bnip1</a>	<a href="#">Bnip1.dSep08</a>	<a href="#">140932</a>	6361	591	1	129	BCL2/adenovirus E1B 19kDa-interacting protein 1 (Bnip1) alternative variant dSep08, mRNA.
<a href="#">Bnip2</a>	<a href="#">Bnip2.aSep08</a>	<a href="#">300811</a>	10982	1779	6	507	BCL2/adenovirus E1B interacting protein 2 (Bnip2) alternative variant aSep08, mRNA.
<a href="#">Bnip2</a>	<a href="#">Bnip2.cSep08</a>	<a href="#">300811</a>	9455	1872	8	242	BCL2/adenovirus E1B interacting protein 2 (Bnip2) alternative variant cSep08, mRNA.
<a href="#">Bnip2</a>	<a href="#">Bnip2.dSep08</a>	<a href="#">300811</a>	7827	963	6	215	BCL2/adenovirus E1B interacting protein 2 (Bnip2) alternative variant dSep08, mRNA.
<a href="#">Bnip2</a>	<a href="#">Bnip2.eSep08</a>	<a href="#">300811</a>	8465	732	5	183	BCL2/adenovirus E1B interacting protein 2 (Bnip2) alternative variant eSep08, mRNA.

<a href="#">Bnip2</a>	<a href="#">Bnip2.fSep08</a>	<a href="#">300811</a>	3976	318	3	83	BCL2/adenovirus E1B interacting protein 2 (Bnip2) alternative variant fSep08, mRNA.
<a href="#">Bnip3</a>	<a href="#">Bnip3.aSep08</a>	<a href="#">84480</a>	17104	1619	6	227	BCL2/adenovirus E1B 19 kDa-interacting protein 3 (Bnip3) alternative variant aSep08, mRNA.
<a href="#">Bnip3</a>	<a href="#">Bnip3.bSep08</a>	<a href="#">84480</a>	6855	431	4	129	BCL2/adenovirus E1B 19 kDa-interacting protein 3 (Bnip3) alternative variant bSep08, mRNA.
<a href="#">Bnip3</a>	<a href="#">Bnip3.cSep08</a>	<a href="#">84480</a>	9142	669	2	60	BCL2/adenovirus E1B 19 kDa-interacting protein 3 (6.5 kD) (Bnip3) alternative variant cSep08, mRNA.
<a href="#">Bnip3l</a>	<a href="#">Bnip3l.aSep08</a>	<a href="#">140923</a>	21108	1301	6	271	BCL2/adenovirus E1B interacting protein 3-like (Bnip3l) alternative variant aSep08, mRNA.
<a href="#">Bnip3l</a>	<a href="#">Bnip3l.cSep08</a>	<a href="#">140923</a>	13373	946	6	179	BCL2/adenovirus E1B interacting protein 3-like (19.4 kD) (Bnip3l) alternative variant cSep08, mRNA.
<a href="#">Bnip3l</a>	<a href="#">Bnip3l.dSep08</a>	<a href="#">140923</a>	14774	1901	4	6	BCL2/adenovirus E1B interacting protein 3-like (Bnip3l) alternative variant dSep08, mRNA.
<a href="#">boby</a>	<a href="#">boby.aSep08</a>		59760	726		110	putative nuclear protein (12.1 kD) (boby) mRNA.
<a href="#">Boc</a>	<a href="#">Boc.aSep08</a>	<a href="#">360715</a>	75385	4491	19	685	biregional cell adhesion molecule-related down-regulated by oncogenes binding protein CRA a like (75.1 kD) (Boc) alternative variant aSep08, mRNA.
<a href="#">bochy</a>	<a href="#">bochy.aSep08</a>		4385	706		104	complement component 5 (bochy) mRNA.
<a href="#">bofer</a>	<a href="#">bofer.aSep08</a>		23853	373	2	32	putative protein (bofer) alternative variant aSep08, mRNA.
<a href="#">bofer</a>	<a href="#">bofer.bSep08</a>		10610	338	1	29	putative protein (bofer) alternative variant bSep08, mRNA.
<a href="#">boflo</a>	<a href="#">boflo.aSep08</a>		9913	219		72	CRA b like (boflo) mRNA.
<a href="#">boflu</a>	<a href="#">boflu.aSep08</a>		3266	1615	3	181	putative protein (20.6 kD) (boflu) alternative variant aSep08, mRNA.
<a href="#">boflu</a>	<a href="#">boflu.bSep08</a>		2430	731	3	83	uncharacterized Protein homolog like (boflu) alternative variant bSep08, mRNA.
<a href="#">Bok</a>	<a href="#">Bok.aSep08</a>	<a href="#">29884</a>	10312	1326	4	213	ovarian killer (23.5 kD) (Bok) alternative variant aSep08, mRNA.
<a href="#">Bok</a>	<a href="#">Bok.cSep08</a>	<a href="#">29884</a>	9538	750	5	120	ovarian killer (13.2 kD) (Bok) alternative variant cSep08, mRNA.
<a href="#">Bok</a>	<a href="#">Bok.dSep08</a>	<a href="#">29884</a>	802	394	2	62	putative protein (6.5 kD) (Bok) alternative variant dSep08, mRNA.
<a href="#">BoIA.0</a>	<a href="#">BoIA.0.aSep08</a>		1415	1077	3	86	bola-Like Protein 2 (10.2 kD) (BoIA.0) alternative variant aSep08, mRNA.
<a href="#">BoIA.0</a>	<a href="#">BoIA.0.bSep08</a>		554	299	2	41	putative protein (BoIA.0) alternative variant bSep08, mRNA.
<a href="#">boloy</a>	<a href="#">boloy.aSep08</a>		1448	476		158	down syndrome cell adhesion molecule CHD2-42 like (boloy) mRNA.
<a href="#">bomer</a>	<a href="#">bomer.aSep08</a>		7493	431	3	61	putative protein (bomer) alternative variant aSep08, mRNA.
<a href="#">bomer</a>	<a href="#">bomer.bSep08</a>		7120	1106	3	45	putative protein (5.1 kD) (bomer) alternative variant bSep08, mRNA.
<a href="#">bomer</a>	<a href="#">bomer.cSep08</a>		6073	1087	3	48	putative protein (5.3 kD) (bomer) alternative variant cSep08, mRNA.
<a href="#">bomer</a>	<a href="#">bomer.dSep08</a>		19660	665	3	45	putative protein (5.1 kD) (bomer) alternative variant dSep08, mRNA.

<a href="#">bomer</a>	<a href="#">bomer.eSep08</a>		6500	485	2	45	putative protein (5.1 kD) (bomer) alternative variant eSep08, mRNA.
<a href="#">bonoy</a>	<a href="#">bonoy.aSep08</a>		3180	477		58	putative protein (bonoy) mRNA.
<a href="#">Bop1</a>	<a href="#">Bop1.bSep08</a>	<a href="#">300050</a>	1338	938	1	312	block of proliferation 1 (Bop1) alternative variant bSep08, mRNA.
<a href="#">bopor</a>	<a href="#">bopor.aSep08</a>		5552	249		53	CRA a like (bopor) mRNA.
<a href="#">borby</a>	<a href="#">borby.aSep08</a>		1555	570		58	putative protein (6.8 kD) (borby) mRNA.
<a href="#">borchy</a>	<a href="#">borchy.aSep08</a>		4014	597		123	putative protein (13.5 kD) (borchy) mRNA.
<a href="#">borfer</a>	<a href="#">borfer.aSep08</a>		13423	1091		48	myosin light kinase (borfer) mRNA.
<a href="#">borflo</a>	<a href="#">borflo.aSep08</a>		31766	343		37	putative protein (borflo) mRNA.
<a href="#">borflu</a>	<a href="#">borflu.aSep08</a>		14500	726	3	132	putative protein (borflu) alternative variant aSep08, mRNA.
<a href="#">borflu</a>	<a href="#">borflu.bSep08</a>		1884	345	2	62	putative protein (6.8 kD) (borflu) alternative variant bSep08, mRNA.
<a href="#">borkee</a>	<a href="#">borkee.aSep08</a>		8196	339	1	88	putative protein (borkee) alternative variant aSep08, mRNA.
<a href="#">borkee</a>	<a href="#">borkee.bSep08</a>		13316	649	2	46	putative protein (5.1 kD) (borkee) alternative variant bSep08, mRNA.
<a href="#">borloy</a>	<a href="#">borloy.aSep08</a>		4805	713		104	putative protein (12.0 kD) (borloy) mRNA.
<a href="#">bormer</a>	<a href="#">bormer.aSep08</a>		2321	459		152	myeloid lymphoid mixed-lineage leukemia like (bormer) mRNA.
<a href="#">bornoy</a>	<a href="#">bornoy.aSep08</a>		969	286		37	putative protein (bornoy) mRNA.
<a href="#">borpor</a>	<a href="#">borpor.aSep08</a>		422	312		86	putative protein (borpor) mRNA.
<a href="#">borsa</a>	<a href="#">borsa.bSep08</a>		2240	822	2	72	putative protein (borsa) alternative variant bSep08, mRNA.
<a href="#">borshee</a>	<a href="#">borshee.aSep08</a>		6501	703		50	putative protein (5.8 kD) (borshee) mRNA.
<a href="#">borto</a>	<a href="#">borto.aSep08</a>		3320	1350		44	putative protein (borto) mRNA.
<a href="#">borvar</a>	<a href="#">borvar.aSep08</a>		28493	4019	17	511	ubiquitin 24 (borvar) alternative variant aSep08, mRNA.
<a href="#">borvar</a>	<a href="#">borvar.bSep08</a>		849	715	2	54	putative protein (borvar) alternative variant bSep08, mRNA.
<a href="#">borwey</a>	<a href="#">borwey.aSep08</a>		120423	1761		51	putative protein (borwey) mRNA.
<a href="#">bosa</a>	<a href="#">bosa.aSep08</a>		2218	644	3	109	deltex 3 (bosa) alternative variant aSep08, mRNA.
<a href="#">bosa</a>	<a href="#">bosa.bSep08</a>		488	317	1	29	putative protein (bosa) alternative variant bSep08, mRNA.
<a href="#">boshee</a>	<a href="#">boshee.aSep08</a>		3149	777		54	putative protein (6.2 kD) (boshee) mRNA.
<a href="#">boto</a>	<a href="#">boto.aSep08</a>		7419	261		27	putative protein (boto) mRNA.
<a href="#">bovar</a>	<a href="#">bovar.aSep08</a>		4281	1049		237	interacting protein 1 (bovar) mRNA.
<a href="#">bowey</a>	<a href="#">bowey.bSep08</a>		16326	623	2	67	putative protein (bowey) alternative variant bSep08, mRNA.
<a href="#">bowey</a>	<a href="#">bowey.cSep08</a>		1290	469	1	43	putative protein (4.9 kD) (bowey) alternative variant cSep08, mRNA.
<a href="#">boyby</a>	<a href="#">boyby.aSep08</a>		8719	455		32	putative protein (3.7 kD) (boyby) mRNA.
<a href="#">boychy</a>	<a href="#">boychy.aSep08</a>		3677	393		130	putative protein (boychy) mRNA.
<a href="#">boyfer</a>	<a href="#">boyfer.aSep08</a>		1285	595		58	putative protein (6.5 kD) (boyfer) mRNA.
<a href="#">boyflo</a>	<a href="#">boyflo.aSep08</a>		6042	715		100	putative protein (boyflo) mRNA.
<a href="#">boyflu</a>	<a href="#">boyflu.aSep08</a>		1820	411		136	sorting nexin 26 (boyflu) mRNA.
<a href="#">boykee</a>	<a href="#">boykee.aSep08</a>		29164	355		50	putative protein (5.7 kD) (boykee) mRNA.

<a href="#">boyloy</a>	<a href="#">boyloy.aSep08</a>		2366	555		98	putative protein (boyloy) mRNA.
<a href="#">boymer</a>	<a href="#">boymer.aSep08</a>		551	395		86	putative protein (boymer) mRNA.
<a href="#">boynoy</a>	<a href="#">boynoy.aSep08</a>		1942	490		29	putative protein (3.4 kD) (boynoy) mRNA.
<a href="#">boypor</a>	<a href="#">boypor.aSep08</a>		13491	1418	11	472	pleckstrin homology-like domain family b member 1 (boypor) alternative variant aSep08, mRNA.
<a href="#">boypor</a>	<a href="#">boypor.bSep08</a>		6660	1236	5	318	pleckstrin homology-like domain family B member 1 (34.3 kD) (boypor) alternative variant bSep08, mRNA.
<a href="#">boypor</a>	<a href="#">boypor.cSep08</a>		1128	435	3	144	pleckstrin homology-like domain family b member 1 (boypor) alternative variant cSep08, mRNA.
<a href="#">boypor</a>	<a href="#">boypor.dSep08</a>		4020	381	3	126	pleckstrin homology-like domain family B member 1 CRA a (boypor) alternative variant dSep08, mRNA.
<a href="#">boysa</a>	<a href="#">boysa.aSep08</a>		6928	842	1	166	putative mitochondrial protein (11.9 kD) (boysa) alternative variant aSep08, mRNA.
<a href="#">boysa</a>	<a href="#">boysa.bSep08</a>		2657	514	1	113	regulator of G-protein 22 (boysa) alternative variant bSep08, mRNA.
<a href="#">boyshee</a>	<a href="#">boyshee.aSep08</a>		824	278		30	putative protein (3.3 kD) (boyshee) mRNA.
<a href="#">boyto</a>	<a href="#">boyto.aSep08</a>		826	260		49	putative protein (boyto) mRNA.
<a href="#">boyvar</a>	<a href="#">boyvar.aSep08</a>		8467	1239	2	40	putative protein (4.2 kD) (boyvar) alternative variant aSep08, mRNA.
<a href="#">boyvar</a>	<a href="#">boyvar.bSep08</a>		5100	710	1	81	putative protein (8.5 kD) (boyvar) alternative variant bSep08, mRNA.
<a href="#">boywey</a>	<a href="#">boywey.aSep08</a>		18569	509		44	putative protein (4.8 kD) (boywey) mRNA.
<a href="#">Bpgm</a>	<a href="#">Bpgm.bSep08</a>	<a href="#">296973</a>	28641	848	1	214	2,3-bisphosphoglycerate mutase (Bpgm) alternative variant bSep08, mRNA.
<a href="#">Bphl</a>	<a href="#">Bphl.bSep08</a>	<a href="#">361239</a>	13789	455	3	151	biphenyl hydrolase-like CRA a (Bphl) alternative variant bSep08, mRNA.
<a href="#">Bphl</a>	<a href="#">Bphl.cSep08</a>	<a href="#">361239</a>	28274	813	4	143	biphenyl hydrolase-like CRA d (Bphl) alternative variant cSep08, mRNA.
<a href="#">Bphl</a>	<a href="#">Bphl.dSep08</a>	<a href="#">361239</a>	9850	388	4	128	biphenyl hydrolase-like CRA a (Bphl) alternative variant dSep08, mRNA.
<a href="#">Bphl</a>	<a href="#">Bphl.eSep08</a>	<a href="#">361239</a>	16025	418	4	89	biphenyl hydrolase-like CRA a (Bphl) alternative variant eSep08, mRNA.
<a href="#">Bphl</a>	<a href="#">Bphl.fSep08</a>	<a href="#">361239</a>	1023	934	2	82	biphenyl hydrolase-like CRA d (Bphl) alternative variant fSep08, mRNA.
<a href="#">Bpnt1</a>	<a href="#">Bpnt1.bSep08</a>	<a href="#">64473</a>	24708	713	5	237	bisphosphate 3'-nucleotidase 1 (Bpnt1) alternative variant bSep08, mRNA.
<a href="#">Bpnt1</a>	<a href="#">Bpnt1.cSep08</a>	<a href="#">64473</a>	12782	639	5	212	bisphosphate 3'-nucleotidase 1 (Bpnt1) alternative variant cSep08, mRNA.
<a href="#">Bpnt1</a>	<a href="#">Bpnt1.dSep08</a>	<a href="#">64473</a>	8626	1090	5	196	bisphosphate 3'-nucleotidase 1 (Bpnt1) alternative variant dSep08, mRNA.
<a href="#">Bpnt1</a>	<a href="#">Bpnt1.eSep08</a>	<a href="#">64473</a>	9012	1431	4	164	bisphosphate 3'-nucleotidase 1 (Bpnt1) alternative variant eSep08, mRNA.
<a href="#">Bpnt1</a>	<a href="#">Bpnt1.fSep08</a>	<a href="#">64473</a>	1791	242	2	80	bisphosphate 3'-nucleotidase 1 (Bpnt1) alternative variant fSep08, mRNA.
<a href="#">Braf</a>	<a href="#">Braf.aSep08</a>	<a href="#">114486</a>	32361	2058		234	v-raf murine sarcoma viral oncogene homolog B1 (Braf) mRNA.

<a href="#">Brca1</a>	<a href="#">Brca1.cSep08</a>	<a href="#">497672</a>	1718	457	2	64	breast cancer 1 (Brca1) alternative variant cSep08, mRNA.
<a href="#">Brca2</a>	<a href="#">Brca2.bSep08</a>	<a href="#">360254</a>	1904	965	2	238	breast cancer 2 (25.4 kD) (Brca2) alternative variant bSep08, mRNA.
<a href="#">BRCT.0</a>	<a href="#">BRCT.0.aSep08</a>		15850	613		204	ADP-ribosyltransferase -like 1 (BRCT.0) mRNA.
<a href="#">BRCT.1</a>	<a href="#">BRCT.1.aSep08</a>		2194	401		132	topoisomerase II binding protein 1 like (BRCT.1) mRNA.
<a href="#">Brd1</a>	<a href="#">Brd1.aSep08</a>	<a href="#">315210</a>	48051	4875	2	1189	zinc finger, PHD-type and bromodomain containing protein (133.5 kD) (Brd1) alternative variant aSep08, complete mRNA.
<a href="#">Brd1</a>	<a href="#">Brd1.bSep08</a>	<a href="#">315210</a>	16621	397	2	131	putative protein of metazoan origin (Brd1) alternative variant bSep08, mRNA.
<a href="#">Brd1</a>	<a href="#">Brd1.cSep08</a>	<a href="#">315210</a>	852	410	1	62	putative protein of mammalian origin (Brd1) alternative variant cSep08, mRNA.
<a href="#">Brd2</a>	<a href="#">Brd2.bSep08</a>	<a href="#">294276</a>	9158	3737	13	636	bromodomain containing protein (Brd2) alternative variant bSep08, mRNA.
<a href="#">Brd2</a>	<a href="#">Brd2.cSep08</a>	<a href="#">294276</a>	3609	1375	5	286	bromodomain containing protein (Brd2) alternative variant cSep08, mRNA.
<a href="#">Brd2</a>	<a href="#">Brd2.dSep08</a>	<a href="#">294276</a>	2383	1958	4	237	bromodomain containing protein (25.4 kD) (Brd2) alternative variant dSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.bSep08</a>	<a href="#">362092</a>	27369	2184	8	497	bromodomain containing protein (Brd3) alternative variant bSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.cSep08</a>	<a href="#">362092</a>	6798	735	5	244	putative protein of eukaryotic origin (Brd3) alternative variant cSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.dSep08</a>	<a href="#">362092</a>	40838	765	5	206	bromodomain containing protein (Brd3) alternative variant dSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.eSep08</a>	<a href="#">362092</a>	7389	3459	4	195	putative protein, with a coiled coil domain, of eukaryotic origin (Brd3) alternative variant eSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.fSep08</a>	<a href="#">362092</a>	4231	520	2	121	putative protein, with a coiled coil domain, of eukaryotic origin (Brd3) alternative variant fSep08, mRNA.
<a href="#">Brd3</a>	<a href="#">Brd3.gSep08</a>	<a href="#">362092</a>	1819	602	2	16	putative protein (1.9 kD) (Brd3) alternative variant gSep08, mRNA.
<a href="#">Brd4</a>	<a href="#">Brd4.aSep08</a>	<a href="#">362844</a>	74006	3027	14	978	bromodomain containing protein (Brd4) alternative variant aSep08, mRNA.
<a href="#">Brd4</a>	<a href="#">Brd4.cSep08</a>	<a href="#">362844</a>	54138	1235	6	146	bromodomain containing protein (16.9 kD) (Brd4) alternative variant cSep08, mRNA.
<a href="#">Brd4</a>	<a href="#">Brd4.dSep08</a>	<a href="#">362844</a>	46755	373	2	124	putative protein of eukaryotic origin (Brd4) alternative variant dSep08, mRNA.
<a href="#">Brd7</a>	<a href="#">Brd7.aSep08</a>	<a href="#">361374</a>	27162	1967	15	585	bromodomain containing protein (Brd7) alternative variant aSep08, mRNA.
<a href="#">Brd7</a>	<a href="#">Brd7.cSep08</a>	<a href="#">361374</a>	3768	681	5	168	putative protein, with a coiled coil domain, of metazoan origin (Brd7) alternative variant cSep08, mRNA.
<a href="#">Brd7</a>	<a href="#">Brd7.eSep08</a>	<a href="#">361374</a>	6167	809	4	121	putative protein of metazoan origin (Brd7) alternative variant eSep08, mRNA.
<a href="#">Brd8</a>	<a href="#">Brd8.bSep08</a>	<a href="#">291691</a>	15363	841	9	280	putative protein, with 2 coiled coil domains, of metazoan origin (Brd8) alternative variant bSep08, mRNA.
<a href="#">Brd8</a>	<a href="#">Brd8.cSep08</a>	<a href="#">291691</a>	16068	762	9	253	putative protein, with 2 coiled coil domains, of metazoan origin (Brd8) alternative variant cSep08, mRNA.

<a href="#">Brd8</a>	<a href="#">Brd8.dSep08</a>	<a href="#">291691</a>	1957	662	3	182	putative protein of vertebrate origin (Brd8) alternative variant dSep08, mRNA.
<a href="#">Brd8</a>	<a href="#">Brd8.eSep08</a>	<a href="#">291691</a>	2576	621	7	134	bromodomain containing protein (Brd8) alternative variant eSep08, mRNA.
<a href="#">Brd8</a>	<a href="#">Brd8.gSep08</a>	<a href="#">291691</a>	716	360	2	101	putative protein of vertebrate origin (Brd8) alternative variant gSep08, mRNA.
<a href="#">Brd8</a>	<a href="#">Brd8.hSep08</a>	<a href="#">291691</a>	4077	773	7	127	putative protein of vertebrate origin (Brd8) alternative variant hSep08, mRNA.
<a href="#">Brd9</a>	<a href="#">Brd9.bSep08</a>	<a href="#">308067</a>	11552	737	7	245	putative protein of metazoan origin (Brd9) alternative variant bSep08, mRNA.
<a href="#">Brd9</a>	<a href="#">Brd9.cSep08</a>	<a href="#">308067</a>	11381	771	7	233	putative protein of metazoan origin (Brd9) alternative variant cSep08, mRNA.
<a href="#">Brd9</a>	<a href="#">Brd9.dSep08</a>	<a href="#">308067</a>	2964	598	5	199	putative protein, with a coiled coil domain, of bilateral origin (Brd9) alternative variant dSep08, mRNA.
<a href="#">Brdt</a>	<a href="#">Brdt.bSep08</a>	<a href="#">305123</a>	19745	775	1	146	bromodomain, testis-specific (Brdt) alternative variant bSep08, mRNA.
<a href="#">Bre</a>	<a href="#">Bre.bSep08</a>	<a href="#">362704</a>	219385	481	5	130	brain reproductive organ-expressed (Bre) alternative variant bSep08, mRNA.
<a href="#">Bre</a>	<a href="#">Bre.cSep08</a>	<a href="#">362704</a>	92752	464	5	100	brain reproductive organ-expressed (Bre) alternative variant cSep08, mRNA.
<a href="#">Bre</a>	<a href="#">Bre.dSep08</a>	<a href="#">362704</a>	8193	463	4	65	mitochondrial ribosomal protein l33 CRA b (7.5 kD) (Bre) alternative variant dSep08, mRNA.
<a href="#">Bre</a>	<a href="#">Bre.eSep08</a>	<a href="#">362704</a>	6029	761	2	50	putative protein (5.9 kD) (Bre) alternative variant eSep08, mRNA.
<a href="#">Bre</a>	<a href="#">Bre.fSep08</a>	<a href="#">362704</a>	135436	646	7	72	brain reproductive organ-expressed (7.8 kD) (Bre) alternative variant fSep08, mRNA.
<a href="#">Brf1</a>	<a href="#">Brf1.bSep08</a>	<a href="#">299347</a>	30615	870	7	289	BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S. cerevisiae) (Brf1) alternative variant bSep08, mRNA.
<a href="#">Brf1</a>	<a href="#">Brf1.cSep08</a>	<a href="#">299347</a>	7662	779	6	188	BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S. cerevisiae) (Brf1) alternative variant cSep08, mRNA.
<a href="#">Brf1</a>	<a href="#">Brf1.dSep08</a>	<a href="#">299347</a>	18516	729	5	170	BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S. cerevisiae) (Brf1) alternative variant dSep08, mRNA.
<a href="#">Brf2</a>	<a href="#">Brf2.bSep08</a>	<a href="#">306542</a>	2103	1125	1	127	BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like (Brf2) alternative variant bSep08, mRNA.
<a href="#">Bri3</a>	<a href="#">Bri3.bSep08</a>	<a href="#">304284</a>	8221	566	1	98	brain protein l3 (Bri3) alternative variant bSep08, mRNA.
<a href="#">Brip1</a>	<a href="#">Brip1.aSep08</a>	<a href="#">360588</a>	46245	1470		489	BRCA1 interacting protein C-terminal helicase 1 (Brip1) mRNA.
<a href="#">BRK.0</a>	<a href="#">BRK.0.aSep08</a>		14902	647	4	215	chromodomain helicase DNA binding protein 9 CRA b like (BRK.0) alternative variant aSep08, mRNA.
<a href="#">BRK.0</a>	<a href="#">BRK.0.bSep08</a>		836	356	1	42	putative protein (5.1 kD) (BRK.0) alternative variant bSep08, mRNA.
<a href="#">Brms1</a>	<a href="#">Brms1.bSep08</a>	<a href="#">293668</a>	1602	782	6	213	breast cancer metastasis-suppressor 1 (Brms1) alternative variant bSep08, mRNA.

<a href="#">Brms1</a>	<a href="#">Brms1.cSep08</a>	<a href="#">293668</a>	5966	769	5	180	breast cancer metastasis-suppressor 1 (Brms1) alternative variant cSep08, mRNA.
<a href="#">Brms1</a>	<a href="#">Brms1.dSep08</a>	<a href="#">293668</a>	1035	679	4	150	breast cancer metastasis-suppressor 1 (Brms1) alternative variant dSep08, mRNA.
<a href="#">Brms1</a>	<a href="#">Brms1.fSep08</a>	<a href="#">293668</a>	4325	733	6	91	breast cancer metastasis-suppressor 1 (Brms1) alternative variant fSep08, mRNA.
<a href="#">Brms1</a>	<a href="#">Brms1.gSep08</a>	<a href="#">293668</a>	795	690	2	87	breast cancer metastasis-suppressor 1 (9.9 kD) (Brms1) alternative variant gSep08, mRNA.
<a href="#">Brms1l</a>	<a href="#">Brms1l.aSep08</a>	<a href="#">299053</a>	33957	2582	10	323	breast cancer metastasis-suppressor 1-like (37.6 kD) (Brms1l) alternative variant aSep08, mRNA.
<a href="#">Brms1l</a>	<a href="#">Brms1l.bSep08</a>	<a href="#">299053</a>	29011	773	9	225	breast cancer metastasis-suppressor 1-like (Brms1l) alternative variant bSep08, mRNA.
<a href="#">Brms1l</a>	<a href="#">Brms1l.cSep08</a>	<a href="#">299053</a>	1720	660	2	37	breast cancer metastasis-suppressor 1-like (Brms1l) alternative variant cSep08, mRNA.
<a href="#">Bromodomain.0</a>	<a href="#">Bromodomain.0.aSep08</a>		14802	750	5	249	polybromo 1 CRA a (Bromodomain.0) alternative variant aSep08, mRNA.
<a href="#">Bromodomain.0</a>	<a href="#">Bromodomain.0.bSep08</a>		19713	652	5	217	polybromo 1 CRA c (Bromodomain.0) alternative variant bSep08, mRNA.
<a href="#">Bromodomain.1</a>	<a href="#">Bromodomain.1.aSep08</a>		27808	1379		459	polybromo 1 CRA c (Bromodomain.1) mRNA.
<a href="#">Bromodomain.3</a>	<a href="#">Bromodomain.3.aSep08</a>		2335	570		189	tripartite motif-containing 66 (Bromodomain.3) mRNA.
<a href="#">Brp16</a>	<a href="#">Brp16.bSep08</a>	<a href="#">315094</a>	1040	832	3	252	brain protein 16 (26.2 kD) (Brp16) alternative variant bSep08, mRNA.
<a href="#">Brpf1</a>	<a href="#">Brpf1.aSep08</a>	<a href="#">679713</a>	3716	1567	5	476	bromodomain and PHD finger containing, 1 (Brpf1) alternative variant aSep08, mRNA.
<a href="#">Brpf1</a>	<a href="#">Brpf1.bSep08</a>	<a href="#">679713</a>	2131	734	3	244	bromodomain and PHD finger containing, 1 (Brpf1) alternative variant bSep08, mRNA.
<a href="#">Brs3</a>	<a href="#">Brs3.bSep08</a>	<a href="#">260319</a>	22022	760	2	222	bombesin-like receptor 3 (Brs3) alternative variant bSep08, mRNA.
<a href="#">Brsk1</a>	<a href="#">Brsk1.aSep08</a>	<a href="#">499073</a>	25739	2798	17	764	BR serine/threonine kinase 1 (Brsk1) alternative variant aSep08, mRNA.
<a href="#">Brsk1</a>	<a href="#">Brsk1.bSep08</a>	<a href="#">499073</a>	2189	726	4	241	BR serine/threonine kinase 1 (Brsk1) alternative variant bSep08, mRNA.
<a href="#">Brsk1</a>	<a href="#">Brsk1.cSep08</a>	<a href="#">499073</a>	2101	884	2	239	BR serine/threonine kinase 1 (27.3 kD) (Brsk1) alternative variant cSep08, mRNA.
<a href="#">Brsk2</a>	<a href="#">Brsk2.aSep08</a>	<a href="#">293631</a>	3954	428		142	brain serine/threonine kinase 2 (Brsk2) mRNA.
<a href="#">Brunol4</a>	<a href="#">Brunol4.aSep08</a>	<a href="#">307540</a>	28329	3134	10	337	bruno-like 4, RNA binding protein (Drosophila) (Brunol4) alternative variant aSep08, mRNA.
<a href="#">Brunol4</a>	<a href="#">Brunol4.bSep08</a>	<a href="#">307540</a>	164420	597	4	168	bruno-like 4, RNA binding protein (Drosophila) (Brunol4) alternative variant bSep08, mRNA.
<a href="#">Brunol4</a>	<a href="#">Brunol4.cSep08</a>	<a href="#">307540</a>	76286	774	1	125	bruno-like 4, RNA binding protein (Drosophila) (Brunol4) alternative variant cSep08, mRNA.
<a href="#">Brunol5</a>	<a href="#">Brunol5.aSep08</a>	<a href="#">314647</a>	12649	457	4	152	bruno-like 5, RNA binding protein (Drosophila) (Brunol5) alternative variant aSep08, mRNA.
<a href="#">Brunol5</a>	<a href="#">Brunol5.bSep08</a>	<a href="#">314647</a>	2328	418	4	138	bruno-like 5, RNA binding protein (Drosophila) (Brunol5) alternative variant bSep08, mRNA.



<a href="#">Brunol5</a>	<a href="#">Brunol5.cSep08</a>	<a href="#">314647</a>	6231	591	6	108	bruno-like 5, RNA binding protein (Drosophila) (11.8 kD) (Brunol5) alternative variant cSep08, mRNA.
<a href="#">Brunol6</a>	<a href="#">Brunol6.bSep08</a>	<a href="#">300758</a>	698	383	4	127	bruno-like 6, RNA binding protein (Drosophila) (Brunol6) alternative variant bSep08, mRNA.
<a href="#">Brunol6</a>	<a href="#">Brunol6.cSep08</a>	<a href="#">300758</a>	16844	363	4	61	bruno-like 6, RNA binding protein (Drosophila) (Brunol6) alternative variant cSep08, mRNA.
<a href="#">Brwd1</a>	<a href="#">Brwd1.bSep08</a>	<a href="#">304061</a>	5905	433	5	144	putative protein of eukaryotic origin (Brwd1) alternative variant bSep08, mRNA.
<a href="#">Brwd2</a>	<a href="#">Brwd2.aSep08</a>	<a href="#">309016</a>	28262	3023	20	749	putative protein of ancient origin (Brwd2) alternative variant aSep08, mRNA.
<a href="#">Brwd2</a>	<a href="#">Brwd2.bSep08</a>	<a href="#">309016</a>	5317	631	5	176	putative protein of metazoan origin (Brwd2) alternative variant bSep08, mRNA.
<a href="#">Brwd2</a>	<a href="#">Brwd2.cSep08</a>	<a href="#">309016</a>	3243	684	5	164	putative endoplasmic reticulum protein of eukaryotic origin (18.1 kD) (Brwd2) alternative variant cSep08, mRNA.
<a href="#">Brwd3</a>	<a href="#">Brwd3.aSep08</a>	<a href="#">317213</a>	5624	521		173	bromodomain containing protein (Brwd3) mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.bSep08</a>	<a href="#">361722</a>	10564	1699	6	384	bernardinelli-Seip congenital lipodystrophy 2 (Bsc12) alternative variant bSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.cSep08</a>	<a href="#">361722</a>	2724	665	6	186	seipin (Bsc12) alternative variant cSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.dSep08</a>	<a href="#">361722</a>	2726	525	5	139	seipin (Bsc12) alternative variant dSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.eSep08</a>	<a href="#">361722</a>	1598	1372	3	121	congenital lipodystrophy 2 (13.1 kD) (Bsc12) alternative variant eSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.fSep08</a>	<a href="#">361722</a>	2501	610	3	101	putative protein (Bsc12) alternative variant fSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.gSep08</a>	<a href="#">361722</a>	632	398	2	81	putative mitochondrial protein (9.3 kD) (Bsc12) alternative variant gSep08, mRNA.
<a href="#">Bsc12</a>	<a href="#">Bsc12.hSep08</a>	<a href="#">361722</a>	895	405	2	61	putative protein (6.9 kD) (Bsc12) alternative variant hSep08, mRNA.
<a href="#">Bsd1</a>	<a href="#">Bsd1.bSep08</a>	<a href="#">297890</a>	18085	2314	6	310	BSD (Bsd1) alternative variant bSep08, mRNA.
<a href="#">Bsd1</a>	<a href="#">Bsd1.cSep08</a>	<a href="#">297890</a>	1187	366	1	9	putative protein (Bsd1) alternative variant cSep08, mRNA.
<a href="#">Bsn</a>	<a href="#">Bsn.bSep08</a>	<a href="#">29138</a>	481	365	1	95	bassoon (10.8 kD) (Bsn) alternative variant bSep08, mRNA.
<a href="#">Bst1</a>	<a href="#">Bst1.bSep08</a>	<a href="#">81506</a>	8314	696	2	82	bone marrow stromal cell antigen 1 (Bst1) alternative variant bSep08, mRNA.
<a href="#">Bst2</a>	<a href="#">Bst2.bSep08</a>	<a href="#">378947</a>	562	374		37	bone marrow stromal cell antigen 2 (Bst2) alternative variant bSep08, mRNA.
<a href="#">Btaf1</a>	<a href="#">Btaf1.aSep08</a>	<a href="#">368042</a>	13488	2593	7	408	BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, (Mot1 homolog, S. cerevisiae) (Btaf1) alternative variant aSep08, mRNA.
<a href="#">Btaf1</a>	<a href="#">Btaf1.bSep08</a>	<a href="#">368042</a>	1094	1006	1	117	BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, (Mot1 homolog, S. cerevisiae) (Btaf1) alternative variant bSep08, mRNA.
<a href="#">BTB.0</a>	<a href="#">BTB.0.aSep08</a>		1102	858		285	BTB/POZ (BTB.0) mRNA.
<a href="#">BTB.1</a>	<a href="#">BTB.1.aSep08</a>		44769	922	1	244	BTB/POZ (BTB.1) alternative variant aSep08, mRNA.
<a href="#">BTB.1</a>	<a href="#">BTB.1.bSep08</a>		40407	645	1	182	BTB/POZ (BTB.1) alternative variant bSep08, mRNA.
<a href="#">BTB.2</a>	<a href="#">BTB.2.aSep08</a>		860	493		164	BTB/POZ (BTB.2) mRNA.
<a href="#">Btbd1</a>	<a href="#">Btbd1.bSep08</a>	<a href="#">293060</a>	27829	1417	9	206	putative cytoplasmic protein of eukaryotic origin (23.3 kD) (Btbd1) alternative variant bSep08, mRNA.

<a href="#">Btbd1</a>	<a href="#">Btbd1.cSep08</a>	<a href="#">293060</a>	5479	2063	3	73	putative protein of metazoan origin (8.0 kD) (Btbd1) alternative variant cSep08, mRNA.
<a href="#">Btbd2</a>	<a href="#">Btbd2.aSep08</a>	<a href="#">500793</a>	5667	2245	2	397	CRA a (Btbd2) alternative variant aSep08, mRNA.
<a href="#">Btbd2</a>	<a href="#">Btbd2.bSep08</a>	<a href="#">500793</a>	2434	1360	3	259	CRA a (Btbd2) alternative variant bSep08, mRNA.
<a href="#">Btbd2</a>	<a href="#">Btbd2.cSep08</a>	<a href="#">500793</a>	1174	473	1	157	CRA a (Btbd2) alternative variant cSep08, mRNA.
<a href="#">Btbd4</a>	<a href="#">Btbd4.bSep08</a>	<a href="#">311718</a>	12932	1560	2	462	BTB/POZ (50.3 kD) (Btbd4) alternative variant bSep08, mRNA.
<a href="#">Btbd4</a>	<a href="#">Btbd4.cSep08</a>	<a href="#">311718</a>	6029	2159	2	173	zinc finger BTB domain-containing protein 46 (19.0 kD) (Btbd4) alternative variant cSep08, mRNA.
<a href="#">Btbd6</a>	<a href="#">Btbd6.bSep08</a>	<a href="#">690367</a>	2031	1704	4	424	BTB domain protein (Btbd6) alternative variant bSep08, mRNA.
<a href="#">Btbd6</a>	<a href="#">Btbd6.cSep08</a>	<a href="#">690367</a>	1260	1081	3	225	BTB domain 6 (Btbd6) alternative variant cSep08, mRNA.
<a href="#">Btbd7</a>	<a href="#">Btbd7.bSep08</a>	<a href="#">362772</a>	11057	758	5	252	putative protein of metazoan origin (Btbd7) alternative variant bSep08, mRNA.
<a href="#">Btbd9</a>	<a href="#">Btbd9.aSep08</a>	<a href="#">294318</a>	119195	1632	4	231	putative protein of ancient origin (Btbd9) alternative variant aSep08, mRNA.
<a href="#">Btbd9</a>	<a href="#">Btbd9.bSep08</a>	<a href="#">294318</a>	436	239	1	49	putative protein (Btbd9) alternative variant bSep08, mRNA.
<a href="#">Btbd10</a>	<a href="#">Btbd10.bSep08</a>	<a href="#">308890</a>	44101	719	5	217	putative protein of eukaryotic origin (Btbd10) alternative variant bSep08, mRNA.
<a href="#">Btbd10</a>	<a href="#">Btbd10.cSep08</a>	<a href="#">308890</a>	6702	504	2	33	putative protein (3.7 kD) (Btbd10) alternative variant cSep08, mRNA.
<a href="#">Btbd11</a>	<a href="#">Btbd11.bSep08</a>	<a href="#">314675</a>	26227	334	1	111	putative protein of metazoan origin (Btbd11) alternative variant bSep08, mRNA.
<a href="#">Btbd12</a>	<a href="#">Btbd12.aSep08</a>	<a href="#">302953</a>	6679	2358	4	572	putative protein of metazoan origin (Btbd12) alternative variant aSep08, mRNA.
<a href="#">Btbd14a</a>	<a href="#">Btbd14a.aSep08</a>	<a href="#">296583</a>	62944	2314	4	585	btbde BTB POZ domain-containing protein 14A (63.1 kD) (Btbd14a) alternative variant aSep08, mRNA.
<a href="#">Btbd14a</a>	<a href="#">Btbd14a.bSep08</a>	<a href="#">296583</a>	24696	588	1	180	btbde BTB POZ domain-containing protein 14A (Btbd14a) alternative variant bSep08, mRNA.
<a href="#">Btbd14b</a>	<a href="#">Btbd14b.bSep08</a>	<a href="#">171454</a>	5770	2054	4	508	transcriptional repressor NAC1 (Btbd14b) alternative variant bSep08, mRNA.
<a href="#">Btbd14b</a>	<a href="#">Btbd14b.cSep08</a>	<a href="#">171454</a>	1626	700	1	168	containing BTB domain 14 (Btbd14b) alternative variant cSep08, mRNA.
<a href="#">Btbd16</a>	<a href="#">Btbd16.bSep08</a>	<a href="#">361658</a>	5047	723	3	130	putative protein of metazoan origin (Btbd16) alternative variant bSep08, mRNA.
<a href="#">Btbd16</a>	<a href="#">Btbd16.cSep08</a>	<a href="#">361658</a>	5924	991	3	56	putative protein of mammalian origin (6.6 kD) (Btbd16) alternative variant cSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.bSep08</a>	<a href="#">306262</a>	31803	880	4	271	biotinidase (Btd) alternative variant bSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.cSep08</a>	<a href="#">306262</a>	31587	742	5	247	biotinidase (Btd) alternative variant cSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.dSep08</a>	<a href="#">306262</a>	31453	726	7	241	biotinidase (Btd) alternative variant dSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.eSep08</a>	<a href="#">306262</a>	31577	395	3	131	biotinidase (Btd) alternative variant eSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.fSep08</a>	<a href="#">306262</a>	30518	739	4	126	biotinidase (Btd) alternative variant fSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.gSep08</a>	<a href="#">306262</a>	30479	783	4	113	biotinidase (Btd) alternative variant gSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.iSep08</a>	<a href="#">306262</a>	26790	626	4	75	putative protein (Btd) alternative variant iSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.jSep08</a>	<a href="#">306262</a>	26654	907	4	69	putative protein (Btd) alternative variant jSep08, mRNA.

<a href="#">Btd</a>	<a href="#">Btd.kSep08</a>	<a href="#">306262</a>	26911	789	5	56	putative protein (Btd) alternative variant kSep08, mRNA.
<a href="#">Btd</a>	<a href="#">Btd.mSep08</a>	<a href="#">306262</a>	26473	546	2	70	putative protein (Btd) alternative variant mSep08, mRNA.
<a href="#">Btf3</a>	<a href="#">Btf3.aSep08</a>	<a href="#">294680</a>	6849	841	6	206	basic transcription factor 3 (22.1 kD) (Btf3) alternative variant aSep08, mRNA.
<a href="#">Btf3</a>	<a href="#">Btf3.cSep08</a>	<a href="#">294680</a>	3982	743	3	99	basic transcription factor 3 (10.7 kD) (Btf3) alternative variant cSep08, mRNA.
<a href="#">Btf3l4</a>	<a href="#">Btf3l4.aSep08</a>	<a href="#">366434</a>	17831	1022		182	basic transcription factor 3-like 4 (Btf3l4) mRNA.
<a href="#">Btg2</a>	<a href="#">Btg2.bSep08</a>	<a href="#">29619</a>	2014	773		139	B-cell translocation gene 2, anti-proliferative (Btg2) alternative variant bSep08, mRNA.
<a href="#">Btg3</a>	<a href="#">Btg3.cSep08</a>	<a href="#">54230</a>	12761	978	4	80	B-cell translocation gene 3 (Btg3) alternative variant cSep08, mRNA.
<a href="#">Btk</a>	<a href="#">Btk.bSep08</a>	<a href="#">367901</a>	4827	695	1	107	bruton agammaglobulinemia tyrosine kinase (Btk) alternative variant bSep08, mRNA.
<a href="#">Btn2a2</a>	<a href="#">Btn2a2.aSep08</a>	<a href="#">306957</a>	3752	743		160	butyrophilin, subfamily 2, member A2 (Btn2a2) mRNA.
<a href="#">Btrc</a>	<a href="#">Btrc.bSep08</a>	<a href="#">361765</a>	138277	926	6	212	beta-transducin repeat containing (Btrc) alternative variant bSep08, mRNA.
<a href="#">Btrc</a>	<a href="#">Btrc.cSep08</a>	<a href="#">361765</a>	146123	790	7	179	beta-transducin repeat containing (Btrc) alternative variant cSep08, mRNA.
<a href="#">Btrc</a>	<a href="#">Btrc.dSep08</a>	<a href="#">361765</a>	136649	388	4	128	beta-transducin repeat containing (Btrc) alternative variant dSep08, mRNA.
<a href="#">Btrc</a>	<a href="#">Btrc.eSep08</a>	<a href="#">361765</a>	12747	1353	3	114	beta-transducin repeat containing (Btrc) alternative variant eSep08, mRNA.
<a href="#">Bub1</a>	<a href="#">Bub1.bSep08</a>	<a href="#">296137</a>	7096	759	6	252	budding uninhibited by benzimidazoles 1 homolog (S. cerevisiae) (Bub1) alternative variant bSep08, mRNA.
<a href="#">Bub1</a>	<a href="#">Bub1.cSep08</a>	<a href="#">296137</a>	5905	744	5	209	budding uninhibited by benzimidazoles 1 homolog (S. cerevisiae) (Bub1) alternative variant cSep08, mRNA.
<a href="#">Bub1</a>	<a href="#">Bub1.dSep08</a>	<a href="#">296137</a>	5714	658	5	172	budding uninhibited by benzimidazoles 1 homolog (S. cerevisiae) (Bub1) alternative variant dSep08, mRNA.
<a href="#">Bub1b</a>	<a href="#">Bub1b.aSep08</a>	<a href="#">171576</a>	31871	2174	14	645	budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae) (Bub1b) alternative variant aSep08, mRNA.
<a href="#">Bub1b</a>	<a href="#">Bub1b.bSep08</a>	<a href="#">171576</a>	12952	1793	3	509	budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae) (Bub1b) alternative variant bSep08, mRNA.
<a href="#">Bub1b</a>	<a href="#">Bub1b.cSep08</a>	<a href="#">171576</a>	7502	1835	2	235	budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae) (26.9 kD) (Bub1b) alternative variant cSep08, mRNA.
<a href="#">Bub3</a>	<a href="#">Bub3.bSep08</a>	<a href="#">361662</a>	10511	1335	8	254	budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) (29.0 kD) (Bub3) alternative variant bSep08, complete mRNA.
<a href="#">Bub3</a>	<a href="#">Bub3.eSep08</a>	<a href="#">361662</a>	3518	743	2	78	budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) (Bub3) alternative variant eSep08, mRNA.
<a href="#">buby</a>	<a href="#">buby.aSep08</a>		4096	419		139	stromal antigen 2 like (buby) mRNA.
<a href="#">buchy</a>	<a href="#">buchy.aSep08</a>		4360	1785		489	complement component (buchy) mRNA.
<a href="#">Bud31</a>	<a href="#">Bud31.aSep08</a>	<a href="#">89819</a>	7427	803	2	144	BUD31 homolog (yeast) (17.0 kD) (Bud31) alternative variant aSep08, complete mRNA.
<a href="#">Bud31</a>	<a href="#">Bud31.bSep08</a>	<a href="#">89819</a>	8221	1756	2	144	BUD31 homolog (yeast) (17.0 kD) (Bud31) alternative variant bSep08, complete mRNA.

<a href="#">Bud31</a>	<a href="#">Bud31.dSep08</a>	<a href="#">89819</a>	5641	778	2	113	BUD31 homolog (yeast) (Bud31) alternative variant dSep08, mRNA.
<a href="#">bufer</a>	<a href="#">bufer.aSep08</a>		7378	715		47	putative protein (bufer) mRNA.
<a href="#">buflo</a>	<a href="#">buflo.aSep08</a>		1451	608		202	putative protein of vertebrate origin (buflo) mRNA.
<a href="#">buflu</a>	<a href="#">buflu.aSep08</a>		4110	323		107	ryanodine receptor (buflu) mRNA.
<a href="#">buloy</a>	<a href="#">buloy.aSep08</a>		1264	716		41	putative protein (buloy) mRNA.
<a href="#">bumer</a>	<a href="#">bumer.aSep08</a>		2417	639		88	putative nuclear protein (9.8 kD) (bumer) mRNA.
<a href="#">bunoy</a>	<a href="#">bunoy.aSep08</a>		2072	686	2	80	putative protein (8.4 kD) (bunoy) alternative variant aSep08, mRNA.
<a href="#">bunoy</a>	<a href="#">bunoy.bSep08</a>		2365	350	2	9	putative protein (bunoy) alternative variant bSep08, mRNA.
<a href="#">bupor</a>	<a href="#">bupor.aSep08</a>		590	512		61	putative protein (bupor) mRNA.
<a href="#">busa</a>	<a href="#">busa.aSep08</a>		4916	447	3	54	centaurin gamma 1 like (busa) alternative variant aSep08, mRNA.
<a href="#">bushee</a>	<a href="#">bushee.aSep08</a>		12948	483		161	putative protein of eukaryotic origin (bushee) mRNA.
<a href="#">buto</a>	<a href="#">buto.aSep08</a>		1465	653		32	putative protein (buto) mRNA.
<a href="#">buvar</a>	<a href="#">buvar.aSep08</a>		22212	482		105	interacting protein 1 (buvar) mRNA.
<a href="#">buwey</a>	<a href="#">buwey.aSep08</a>		2571	1053		112	putative protein of metazoan origin (buwey) mRNA.
<a href="#">Bxdc2</a>	<a href="#">Bxdc2.bSep08</a>	<a href="#">294799</a>	8964	734	9	152	brix (Bxdc2) alternative variant bSep08, mRNA.
<a href="#">Bxdc5</a>	<a href="#">Bxdc5.aSep08</a>	<a href="#">499725</a>	15050	1818	9	367	brix (42.2 kD) (Bxdc5) alternative variant aSep08, mRNA.
<a href="#">Bxdc5</a>	<a href="#">Bxdc5.bSep08</a>	<a href="#">499725</a>	2328	732	2	101	putative protein, with a coiled coil domain, of vertebrate origin (Bxdc5) alternative variant bSep08, mRNA.
<a href="#">Bxdc5</a>	<a href="#">Bxdc5.cSep08</a>	<a href="#">499725</a>	1148	565	3	87	putative protein of eukaryotic origin (Bxdc5) alternative variant cSep08, mRNA.
<a href="#">Bxdc5</a>	<a href="#">Bxdc5.dSep08</a>	<a href="#">499725</a>	1070	928	2	84	putative protein of eukaryotic origin (10.1 kD) (Bxdc5) alternative variant dSep08, mRNA.
<a href="#">byby</a>	<a href="#">byby.aSep08</a>		12287	612		204	stromal antigen 2 like (byby) mRNA.
<a href="#">bychu</a>	<a href="#">bychu.aSep08</a>		22136	777	5	118	epididymal protein 52 like (13.1 kD) (bychu) alternative variant aSep08, mRNA.
<a href="#">bychu</a>	<a href="#">bychu.bSep08</a>		5586	657	3	76	putative protein (bychu) alternative variant bSep08, mRNA.
<a href="#">bychy</a>	<a href="#">bychy.aSep08</a>		4249	1313		79	putative protein (bychy) mRNA.
<a href="#">byfer</a>	<a href="#">byfer.aSep08</a>		663	365		36	putative protein (byfer) mRNA.
<a href="#">byflo</a>	<a href="#">byflo.aSep08</a>		1909	536		178	putative protein of ancient origin (byflo) mRNA.
<a href="#">byflu</a>	<a href="#">byflu.aSep08</a>		461	369		122	ryanodine receptor (byflu) mRNA.
<a href="#">byloy</a>	<a href="#">byloy.aSep08</a>		2981	290		33	putative protein (byloy) mRNA.
<a href="#">bymer</a>	<a href="#">bymer.aSep08</a>		534	350		62	putative protein (7.1 kD) (bymer) mRNA.
<a href="#">bynoy</a>	<a href="#">bynoy.aSep08</a>		5701	674		224	putative protein, with a coiled coil domain, of ancient origin (bynoy) mRNA.
<a href="#">bypor</a>	<a href="#">bypor.aSep08</a>		4122	426		56	putative protein (6.5 kD) (bypor) mRNA.
<a href="#">bysa</a>	<a href="#">bysa.aSep08</a>		1212	172		57	centaurin gamma 1 PIKE-L (bysa) mRNA.
<a href="#">byshee</a>	<a href="#">byshee.aSep08</a>		5782	501		166	putative protein of mammalian origin (byshee) alternative variant aSep08, mRNA.
<a href="#">byshee</a>	<a href="#">byshee.bSep08</a>		13768	403		134	putative protein of mammalian origin (byshee) alternative variant bSep08, mRNA.

<a href="#">byto</a>	<a href="#">byto.bSep08</a>		2618	770	2	62	putative protein of mammalian origin (7.4 kD) (byto) alternative variant bSep08, mRNA.
<a href="#">byvar</a>	<a href="#">byvar.aSep08</a>		6309	300		100	interacting protein 1 (byvar) mRNA.
<a href="#">bywey</a>	<a href="#">bywey.aSep08</a>		7029	418		116	putative protein (bywey) mRNA.
<a href="#">Bzrap1</a>	<a href="#">Bzrap1.aSep08</a>	<a href="#">287609</a>	7789	1800		537	benzodiazapine receptor associated protein 1 (Bzrap1) alternative variant aSep08, mRNA.
<a href="#">Bzrap1</a>	<a href="#">Bzrap1.bSep08</a>	<a href="#">287609</a>	1728	423		140	benzodiazapine receptor associated protein 1 (Bzrap1) alternative variant bSep08, mRNA.
<a href="#">Bzw1</a>	<a href="#">Bzw1.bSep08</a>	<a href="#">363232</a>	12716	2100	7	311	basic leucine zipper and W2 domains 1 (Bzw1) alternative variant bSep08, mRNA.
<a href="#">Bzw1</a>	<a href="#">Bzw1.cSep08</a>	<a href="#">363232</a>	5601	430	4	143	basic leucine zipper and W2 domains 1 (Bzw1) alternative variant cSep08, mRNA.
<a href="#">Bzw1</a>	<a href="#">Bzw1.eSep08</a>	<a href="#">363232</a>	1971	680	2	27	basic leucine zipper and W2 domains 1 (Bzw1) alternative variant eSep08, mRNA.
<a href="#">Bzw2</a>	<a href="#">Bzw2.bSep08</a>	<a href="#">171439</a>	2122	1486	2	62	basic leucine zipper and W2 domains 2 (Bzw2) alternative variant bSep08, mRNA.
<a href="#">C1-set.0</a>	<a href="#">C1-set.0.bSep08</a>		2721	802	3	128	T-cell receptor like (C1-set.0) alternative variant bSep08, mRNA.
<a href="#">C1-set.1</a>	<a href="#">C1-set.1.aSep08</a>		1855	492	2	117	lambda chain (C1-set.1) alternative variant aSep08, mRNA.
<a href="#">C1-set.1</a>	<a href="#">C1-set.1.bSep08</a>		3340	531	2	110	lambda chain (11.9 kD) (C1-set.1) alternative variant bSep08, mRNA.
<a href="#">C1-set.3</a>	<a href="#">C1-set.3.aSep08</a>		671	587		195	ig epsilon (C1-set.3) mRNA.
<a href="#">C1-set.4</a>	<a href="#">C1-set.4.aSep08</a>		1627	1116		336	immunoglobulin (C1-set.4) mRNA.
<a href="#">C1-set.5</a>	<a href="#">C1-set.5.aSep08</a>		1646	1078		322	immunoglobulin (C1-set.5) mRNA.
<a href="#">C1-set.6</a>	<a href="#">C1-set.6.aSep08</a>		3922	1390	3	400	ig region (C1-set.6) alternative variant aSep08, mRNA.
<a href="#">C1-set.6</a>	<a href="#">C1-set.6.bSep08</a>		799	700	1	200	immunoglobulin heavy chain (C1-set.6) alternative variant bSep08, mRNA.
<a href="#">C1-set.7</a>	<a href="#">C1-set.7.aSep08</a>		7974	1481	3	162	ig delta region (C1-set.7) alternative variant aSep08, mRNA.
<a href="#">C1-set.7</a>	<a href="#">C1-set.7.bSep08</a>		5069	386	2	75	delta secreted form (C1-set.7) alternative variant bSep08, mRNA.
<a href="#">C1-set.8</a>	<a href="#">C1-set.8.aSep08</a>		330240	778	4	259	T-cell receptor like (C1-set.8) alternative variant aSep08, mRNA.
<a href="#">C1-set.8</a>	<a href="#">C1-set.8.bSep08</a>		327935	727	3	213	T-cell receptor like (C1-set.8) alternative variant bSep08, mRNA.
<a href="#">C1-set.8</a>	<a href="#">C1-set.8.cSep08</a>		4488	766	5	179	T-cell receptor like (C1-set.8) alternative variant cSep08, mRNA.
<a href="#">C1galt1c1</a>	<a href="#">C1galt1c1.aSep08</a>	<a href="#">302499</a>	4928	1862	2	316	C1GALT1-specific chaperone 1 (36.0 kD) (C1galt1c1) alternative variant aSep08, complete mRNA.
<a href="#">C1galt1c1</a>	<a href="#">C1galt1c1.cSep08</a>	<a href="#">302499</a>	705	401	2	43	C1GALT1-specific chaperone 1 (C1galt1c1) alternative variant cSep08, mRNA.
<a href="#">C1qbp</a>	<a href="#">C1qbp.bSep08</a>	<a href="#">29681</a>	4251	795	6	233	complement component 1, q subcomponent binding protein (C1qbp) alternative variant bSep08, mRNA.
<a href="#">C1qbp</a>	<a href="#">C1qbp.cSep08</a>	<a href="#">29681</a>	3941	682	4	145	complement component 1, q subcomponent binding protein (C1qbp) alternative variant cSep08, mRNA.

<a href="#">C1ql3</a>	<a href="#">C1ql3.aSep08</a>	<a href="#">680404</a>	8693	2965	2	98	complement component 1, q subcomponent-like 3 (C1ql3) alternative variant aSep08, mRNA.
<a href="#">C1qtnf1</a>	<a href="#">C1qtnf1.aSep08</a>	<a href="#">303701</a>	6834	2491	2	315	c1q and tumor necrosis factor related protein 1 (C1qtnf1) alternative variant aSep08, mRNA.
<a href="#">C1qtnf1</a>	<a href="#">C1qtnf1.cSep08</a>	<a href="#">303701</a>	13856	827	2	259	c1q and tumor necrosis factor related protein 1 (C1qtnf1) alternative variant cSep08, mRNA.
<a href="#">C1qtnf2</a>	<a href="#">C1qtnf2.aSep08</a>	<a href="#">497886</a>	17664	1216		294	c1q and tumor necrosis factor related protein 2 (31.0 kD) (C1qtnf2) mRNA.
<a href="#">C1qtnf3</a>	<a href="#">C1qtnf3.bSep08</a>	<a href="#">294806</a>	13359	577	1	192	c1q and tumor necrosis factor related protein 3 (C1qtnf3) alternative variant bSep08, mRNA.
<a href="#">C1qtnf4</a>	<a href="#">C1qtnf4.bSep08</a>	<a href="#">311184</a>	2320	667	1	61	c1q and tumor necrosis factor related protein 4 (C1qtnf4) alternative variant bSep08, mRNA.
<a href="#">C1qtnf5</a>	<a href="#">C1qtnf5.aSep08</a>	<a href="#">315598</a>	1262	933	1	311	c1q and tumor necrosis factor related protein 5 (C1qtnf5) alternative variant aSep08, mRNA.
<a href="#">C1qtnf7</a>	<a href="#">C1qtnf7.bSep08</a>	<a href="#">305423</a>	112799	1785	1	296	c1q and tumor necrosis factor related protein 7 (31.4 kD) (C1qtnf7) alternative variant bSep08, mRNA.
<a href="#">C1r</a>	<a href="#">C1r.bSep08</a>	<a href="#">312705</a>	1186	728	2	52	complement component 1, r subcomponent (C1r) alternative variant bSep08, mRNA.
<a href="#">C1sandRGD1561715</a>	<a href="#">C1sandRGD1561715.bSep08</a>	<a href="#">192262</a>	2614	1012	3	252	complement component 1 s subcomponent (C1sandRGD1561715) alternative variant bSep08, mRNA.
<a href="#">C1sandRGD1561715</a>	<a href="#">C1sandRGD1561715.bSep08</a>	<a href="#">500313</a>	2614	1012	3	252	complement component 1 s subcomponent (C1sandRGD1561715) alternative variant bSep08, mRNA.
<a href="#">C1sandRGD1561715</a>	<a href="#">C1sandRGD1561715.cSep08</a>	<a href="#">192262</a>	379569	728	2	209	complement component C1SA (C1sandRGD1561715) alternative variant cSep08, mRNA.
<a href="#">C1sandRGD1561715</a>	<a href="#">C1sandRGD1561715.cSep08</a>	<a href="#">500313</a>	379569	728	2	209	complement component C1SA (C1sandRGD1561715) alternative variant cSep08, mRNA.
<a href="#">C1_1.0</a>	<a href="#">C1_1.0.aSep08</a>		82772	795		203	protein kinase C theta (C1_1.0) mRNA.
<a href="#">C1_1.1</a>	<a href="#">C1_1.1.aSep08</a>		2798	428		142	diacylglycerol kinase theta (C1_1.1) mRNA.
<a href="#">C1_1.2</a>	<a href="#">C1_1.2.aSep08</a>		1339	292		97	diacylglycerol kinase eta (C1_1.2) mRNA.
<a href="#">C1_1.3</a>	<a href="#">C1_1.3.aSep08</a>		6954	995		295	kinase d1 (C1_1.3) mRNA.
<a href="#">C2</a>	<a href="#">C2.aSep08</a>	<a href="#">24231</a>	4893	1155	9	304	complement component 2 (C2) alternative variant aSep08, mRNA.
<a href="#">C2</a>	<a href="#">C2.bSep08</a>	<a href="#">24231</a>	9862	824	6	274	complement component 2 (C2) alternative variant bSep08, mRNA.
<a href="#">C2</a>	<a href="#">C2.eSep08</a>	<a href="#">24231</a>	8770	691	3		
<a href="#">C2.1</a>	<a href="#">C2.1.aSep08</a>		8389	337	3	112	unc-13 homolog B CRA b (C2.1) alternative variant aSep08, mRNA.
<a href="#">C2.2</a>	<a href="#">C2.2.aSep08</a>		16253	963		315	RAS p21 protein activator 4 (C2.2) mRNA.
<a href="#">C2.3</a>	<a href="#">C2.3.aSep08</a>		2788	702		121	intersectin 1 (C2.3) mRNA.
<a href="#">C2.4</a>	<a href="#">C2.4.aSep08</a>		5185	3104		90	DNA segment Chr 12 ERATO Doi 551 expressed (C2.4) mRNA.
<a href="#">C2.5</a>	<a href="#">C2.5.aSep08</a>		12208	329		97	regulator of G-protein 3 (C2.5) mRNA.
<a href="#">C2.6</a>	<a href="#">C2.6.aSep08</a>		12761	653		217	dysferlin (C2.6) mRNA.
<a href="#">C2.7</a>	<a href="#">C2.7.aSep08</a>		8312	738		245	synaptotagmin XVII (C2.7) mRNA.
<a href="#">C2.8</a>	<a href="#">C2.8.aSep08</a>		43202	1717		572	myoferlin (C2.8) mRNA.

<a href="#">C2cd3</a>	<a href="#">C2cd3.aSep08</a>	<a href="#">293148</a>	15922	1527		299	CRA b (C2cd3) mRNA.
<a href="#">C4.0</a>	<a href="#">C4.0.aSep08</a>		14501	3300		773	procollagen type IV alpha 2 (C4.0) mRNA.
<a href="#">C4.1</a>	<a href="#">C4.1.aSep08</a>		20795	3184	6	371	IV alpha (C4.1) alternative variant aSep08, mRNA.
<a href="#">C4.1</a>	<a href="#">C4.1.bSep08</a>		5744	708	1	58	putative protein (C4.1) alternative variant bSep08, mRNA.
<a href="#">C4.2</a>	<a href="#">C4.2.aSep08</a>		14018	771		257	alpha IV (C4.2) mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.cSep08</a>	<a href="#">24233</a>	1907	898	6	195	complement component (C4aandC4-2) alternative variant cSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.cSep08</a>	<a href="#">406161</a>	1907	898	6	195	complement component (C4aandC4-2) alternative variant cSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.dSep08</a>	<a href="#">24233</a>	847	512	4	163	complement component (C4aandC4-2) alternative variant dSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.dSep08</a>	<a href="#">406161</a>	847	512	4	163	complement component (C4aandC4-2) alternative variant dSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.eSep08</a>	<a href="#">24233</a>	843	466	4	155	complement component (C4aandC4-2) alternative variant eSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.eSep08</a>	<a href="#">406161</a>	843	466	4	155	complement component (C4aandC4-2) alternative variant eSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.fSep08</a>	<a href="#">24233</a>	1712	398	6	104	complement component (C4aandC4-2) alternative variant fSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.fSep08</a>	<a href="#">406161</a>	1712	398	6	104	complement component (C4aandC4-2) alternative variant fSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.gSep08</a>	<a href="#">24233</a>	567	433	2	59	complement component 4a (6.7 kD) (C4aandC4-2) alternative variant gSep08, mRNA.
<a href="#">C4aandC4-2</a>	<a href="#">C4aandC4-2.gSep08</a>	<a href="#">406161</a>	567	433	2	59	complement component 4a (6.7 kD) (C4aandC4-2) alternative variant gSep08, mRNA.
<a href="#">C4bpb</a>	<a href="#">C4bpb.bSep08</a>	<a href="#">24236</a>	1629	284	1	81	complement component 4 binding protein, beta (C4bpb) alternative variant bSep08, mRNA.
<a href="#">C5</a>	<a href="#">C5.aSep08</a>	<a href="#">362119</a>	40720	2828	20	824	complement component (C5) alternative variant aSep08, mRNA.
<a href="#">C5</a>	<a href="#">C5.bSep08</a>	<a href="#">362119</a>	20821	1802	8	435	complement component (C5) alternative variant bSep08, mRNA.
<a href="#">C5</a>	<a href="#">C5.cSep08</a>	<a href="#">362119</a>	8557	886	3	116	complement component (C5) alternative variant cSep08, mRNA.
<a href="#">C5</a>	<a href="#">C5.dSep08</a>	<a href="#">362119</a>	7477	928	5	109	complement component C5 (C5) alternative variant dSep08, mRNA.
<a href="#">C6</a>	<a href="#">C6.bSep08</a>	<a href="#">24237</a>	10573	830	2	169	complement component 6 (18.9 kD) (C6) alternative variant bSep08, mRNA.
<a href="#">C7</a>	<a href="#">C7.aSep08</a>	<a href="#">117517</a>	72587	1616	8	367	complement component 7 (C7) alternative variant aSep08, mRNA.
<a href="#">C8.0</a>	<a href="#">C8.0.aSep08</a>		5118	587		195	IgG binding protein like (C8.0) mRNA.
<a href="#">C8b</a>	<a href="#">C8b.aSep08</a>	<a href="#">313421</a>	37470	2377		583	complement component 8, beta polypeptide (C8b) mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.aSep08</a>	<a href="#">296545</a>	10088	1300	7	271	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant aSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.aSep08</a>	<a href="#">680602</a>	10088	1300	7	271	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant aSep08, mRNA.

<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.bSep08</a>	<a href="#">296545</a>	2980	704	6	208	lipocalin 12 (C8gandLcn12) alternative variant bSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.bSep08</a>	<a href="#">680602</a>	2980	704	6	208	lipocalin 12 (C8gandLcn12) alternative variant bSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.cSep08</a>	<a href="#">296545</a>	1545	753	7	199	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant cSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.cSep08</a>	<a href="#">680602</a>	1545	753	7	199	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant cSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.eSep08</a>	<a href="#">296545</a>	1572	523	5	164	complement component (C8gandLcn12) alternative variant eSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.eSep08</a>	<a href="#">680602</a>	1572	523	5	164	complement component (C8gandLcn12) alternative variant eSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.gSep08</a>	<a href="#">296545</a>	917	576	3	121	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant gSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.gSep08</a>	<a href="#">680602</a>	917	576	3	121	complement component 8 gamma polypeptide CRA b (C8gandLcn12) alternative variant gSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.hSep08</a>	<a href="#">296545</a>	2357	370	4	95	lipocalin 12 (C8gandLcn12) alternative variant hSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.hSep08</a>	<a href="#">680602</a>	2357	370	4	95	lipocalin 12 (C8gandLcn12) alternative variant hSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.iSep08</a>	<a href="#">296545</a>	1086	374	2	79	lipocalin 12 (C8gandLcn12) alternative variant iSep08, mRNA.
<a href="#">C8gandLcn12</a>	<a href="#">C8gandLcn12.iSep08</a>	<a href="#">680602</a>	1086	374	2	79	lipocalin 12 (C8gandLcn12) alternative variant iSep08, mRNA.
<a href="#">C9</a>	<a href="#">C9.bSep08</a>	<a href="#">117512</a>	6223	1339	3	446	complement component 9 (C9) alternative variant bSep08, mRNA.
<a href="#">Cab39</a>	<a href="#">Cab39.aSep08</a>	<a href="#">301574</a>	59832	1937	9	341	calcium binding protein 39 (39.9 kD) (Cab39) alternative variant aSep08, mRNA.
<a href="#">Cab39</a>	<a href="#">Cab39.bSep08</a>	<a href="#">301574</a>	55861	771	7	230	calcium binding protein 39 (Cab39) alternative variant bSep08, mRNA.
<a href="#">Cab39</a>	<a href="#">Cab39.cSep08</a>	<a href="#">301574</a>	53581	739	6	209	calcium binding protein 39 (Cab39) alternative variant cSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.bSep08</a>	<a href="#">290291</a>	89838	2739	9	289	calcium binding protein 39-like (33.7 kD) (Cab39I) alternative variant bSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.cSep08</a>	<a href="#">290291</a>	74988	931	8	186	calcium binding protein 39-like (Cab39I) alternative variant cSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.dSep08</a>	<a href="#">290291</a>	64392	684	6	121	calcium binding protein 39-like (Cab39I) alternative variant dSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.eSep08</a>	<a href="#">290291</a>	8833	699	2	93	calcium binding protein 39-like (11.0 kD) (Cab39I) alternative variant eSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.fSep08</a>	<a href="#">290291</a>	64361	811	7	116	calcium binding protein 39-like (Cab39I) alternative variant fSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.gSep08</a>	<a href="#">290291</a>	49694	618	6	23	calcium binding protein 39-like (Cab39I) alternative variant gSep08, mRNA.
<a href="#">Cab39I</a>	<a href="#">Cab39I.hSep08</a>	<a href="#">290291</a>	15210	412	2	49	calcium binding protein 39-like (Cab39I) alternative variant hSep08, mRNA.



<a href="#">Cabc1</a>	<a href="#">Cabc1.aSep08</a>	<a href="#">360887</a>	29581	2676	1	649	chaperone, ABC1 activity of bc1 complex like (S. pombe) (72.2 kD) (Cabc1) alternative variant aSep08, mRNA.
<a href="#">Cabc1</a>	<a href="#">Cabc1.bSep08</a>	<a href="#">360887</a>	18711	766	1	240	chaperone, ABC1 activity of bc1 complex like (S. pombe) (Cabc1) alternative variant bSep08, mRNA.
<a href="#">Cabin1</a>	<a href="#">Cabin1.bSep08</a>	<a href="#">94165</a>	11344	1055	4	258	calcineurin binding protein 1 like (Cabin1) alternative variant bSep08, mRNA.
<a href="#">Cabin1</a>	<a href="#">Cabin1.cSep08</a>	<a href="#">94165</a>	4682	596	3	136	calcineurin binding protein 1 like (Cabin1) alternative variant cSep08, mRNA.
<a href="#">Cables2</a>	<a href="#">Cables2.aSep08</a>	<a href="#">311703</a>	991	734	2	244	cdk5 and Abl enzyme substrate 2 (Cables2) alternative variant aSep08, mRNA.
<a href="#">Cables2</a>	<a href="#">Cables2.bSep08</a>	<a href="#">311703</a>	928	816	1	123	cdk5 and Abl enzyme substrate 2 (Cables2) alternative variant bSep08, mRNA.
<a href="#">Cabp1</a>	<a href="#">Cabp1.cSep08</a>	<a href="#">171051</a>	3887	1060	1	108	calcium binding protein 1 (Cabp1) alternative variant cSep08, mRNA.
<a href="#">Cabp2</a>	<a href="#">Cabp2.aSep08</a>	<a href="#">499298</a>	4867	562		34	calcium binding protein 2 (3.4 kD) (Cabp2) mRNA.
<a href="#">Cachd1</a>	<a href="#">Cachd1.aSep08</a>	<a href="#">298267</a>	140368	469		156	VWA N-terminal (Cachd1) mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.bSep08</a>	<a href="#">25398</a>	36845	1796	16	371	calcium channel (Cacna1a) alternative variant bSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.cSep08</a>	<a href="#">25398</a>	15534	1348	6	236	calcium channel (27.0 kD) (Cacna1a) alternative variant cSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.dSep08</a>	<a href="#">25398</a>	17135	791	7	192	calcium channel alpha (Cacna1a) alternative variant dSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.eSep08</a>	<a href="#">25398</a>	9830	485	3	139	calcium channel BI-1 (Cacna1a) alternative variant eSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.gSep08</a>	<a href="#">25398</a>	710	341	2	113	calcium channel (Cacna1a) alternative variant gSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.hSep08</a>	<a href="#">25398</a>	2393	500	2	110	calcium channel (Cacna1a) alternative variant hSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.iSep08</a>	<a href="#">25398</a>	1408	746	2	91	calcium channel alpha (Cacna1a) alternative variant iSep08, mRNA.
<a href="#">Cacna1a</a>	<a href="#">Cacna1a.jSep08</a>	<a href="#">25398</a>	970	590	3	73	calcium channel BI-2 (Cacna1a) alternative variant jSep08, mRNA.
<a href="#">Cacna1c</a>	<a href="#">Cacna1c.aSep08</a>	<a href="#">24239</a>	526188	6908	48	2271	calcium channel, voltage-dependent, L type, alpha 1C subunit (Cacna1c) alternative variant aSep08, mRNA.
<a href="#">Cacna1c</a>	<a href="#">Cacna1c.cSep08</a>	<a href="#">24239</a>	496271	7206	48	2258	calcium channel, voltage-dependent, L type, alpha 1C subunit (Cacna1c) alternative variant cSep08, mRNA.
<a href="#">Cacna1c</a>	<a href="#">Cacna1c.dSep08</a>	<a href="#">24239</a>	7917	852	5	283	calcium channel, voltage-dependent, L type, alpha 1C subunit (Cacna1c) alternative variant dSep08, mRNA.
<a href="#">Cacna1c</a>	<a href="#">Cacna1c.fSep08</a>	<a href="#">24239</a>	3076	306	2	52	calcium channel, voltage-dependent, L type, alpha 1C subunit (Cacna1c) alternative variant fSep08, mRNA.
<a href="#">Cacna1d</a>	<a href="#">Cacna1d.bSep08</a>	<a href="#">29716</a>	32282	502	3	142	calcium channel, voltage-dependent, L type, alpha 1D subunit (Cacna1d) alternative variant bSep08, mRNA.
<a href="#">Cacna1d</a>	<a href="#">Cacna1d.cSep08</a>	<a href="#">29716</a>	8711	910	3	51	calcium channel, voltage-dependent, L type, alpha 1D subunit (Cacna1d) alternative variant cSep08, mRNA.
<a href="#">Cacna1e</a>	<a href="#">Cacna1e.bSep08</a>	<a href="#">54234</a>	4572	361	2	59	calcium channel, voltage-dependent, R type, alpha 1E subunit (Cacna1e) alternative variant bSep08, mRNA.

<a href="#">Cacna1f</a>	<a href="#">Cacna1f.bSep08</a>	<a href="#">114493</a>	1656	355	4	116	calcium channel, voltage-dependent, alpha 1F subunit (Cacna1f) alternative variant bSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.bSep08</a>	<a href="#">29717</a>	11346	1057	7	351	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant bSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.cSep08</a>	<a href="#">29717</a>	10195	882	9	293	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant cSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.dSep08</a>	<a href="#">29717</a>	4958	867	4	288	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant dSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.eSep08</a>	<a href="#">29717</a>	2059	381	5	127	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant eSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.fSep08</a>	<a href="#">29717</a>	2753	406	2	106	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant fSep08, mRNA.
<a href="#">Cacna1g</a>	<a href="#">Cacna1g.hSep08</a>	<a href="#">29717</a>	885	419	2	86	calcium channel, voltage-dependent, T type, alpha 1G subunit (Cacna1g) alternative variant hSep08, mRNA.
<a href="#">Cacna1h</a>	<a href="#">Cacna1h.cSep08</a>	<a href="#">114862</a>	787	505	2	79	calcium channel, voltage-dependent, T type, alpha 1H subunit (Cacna1h) alternative variant cSep08, mRNA.
<a href="#">Cacna1i</a>	<a href="#">Cacna1i.aSep08</a>	<a href="#">56827</a>	4025	532		176	calcium channel, voltage-dependent, alpha 1I subunit (Cacna1i) mRNA.
<a href="#">Cacna2d1</a>	<a href="#">Cacna2d1.cSep08</a>	<a href="#">25399</a>	18203	1371	10	319	calcium channel, voltage-dependent, alpha2/delta subunit 1 (Cacna2d1) alternative variant cSep08, mRNA.
<a href="#">Cacna2d1</a>	<a href="#">Cacna2d1.dSep08</a>	<a href="#">25399</a>	2173	394	3	103	calcium channel, voltage-dependent, alpha2/delta subunit 1 (Cacna2d1) alternative variant dSep08, mRNA.
<a href="#">Cacna2d1</a>	<a href="#">Cacna2d1.eSep08</a>	<a href="#">25399</a>	15304	1045	2	49	calcium channel, voltage-dependent, alpha2/delta subunit 1 (Cacna2d1) alternative variant eSep08, mRNA.
<a href="#">Cacna2d2</a>	<a href="#">Cacna2d2.aSep08</a>	<a href="#">300992</a>	2673	501		166	calcium channel, voltage-dependent, alpha 2/delta subunit 2 (Cacna2d2) mRNA.
<a href="#">Cacna2d3</a>	<a href="#">Cacna2d3.bSep08</a>	<a href="#">306243</a>	401844	634	7	120	calcium channel, voltage-dependent, alpha2/delta subunit 3 (Cacna2d3) alternative variant bSep08, mRNA.
<a href="#">Cacna2d4</a>	<a href="#">Cacna2d4.aSep08</a>	<a href="#">312668</a>	3961	392		130	calcium channel, voltage-dependent, alpha 2/delta subunit 4 (Cacna2d4) mRNA.
<a href="#">Cacnb1</a>	<a href="#">Cacnb1.bSep08</a>	<a href="#">50688</a>	5548	403	4	134	calcium channel, voltage-dependent, beta 1 subunit (Cacnb1) alternative variant bSep08, mRNA.
<a href="#">Cacnb1</a>	<a href="#">Cacnb1.fSep08</a>	<a href="#">50688</a>	1139	510	3	84	calcium channel, voltage-dependent, beta 1 subunit (Cacnb1) alternative variant fSep08, mRNA.
<a href="#">Cacnb2</a>	<a href="#">Cacnb2.bSep08</a>	<a href="#">116600</a>	18100	403	1	134	calcium channel, voltage-dependent, beta 2 subunit (Cacnb2) alternative variant bSep08, mRNA.
<a href="#">Cacnb3</a>	<a href="#">Cacnb3.bSep08</a>	<a href="#">25297</a>	1707	779	4	239	calcium channel, voltage-dependent, beta 3 subunit (Cacnb3) alternative variant bSep08, mRNA.
<a href="#">Cacnb3</a>	<a href="#">Cacnb3.cSep08</a>	<a href="#">25297</a>	8410	687	6	147	calcium channel, voltage-dependent, beta 3 subunit (Cacnb3) alternative variant cSep08, mRNA.
<a href="#">Cacng6</a>	<a href="#">Cacng6.bSep08</a>	<a href="#">140727</a>	12728	734	1	158	calcium channel, voltage-dependent, gamma subunit 6 (Cacng6) alternative variant bSep08, mRNA.
<a href="#">Cacybp</a>	<a href="#">Cacybp.aSep08</a>	<a href="#">289144</a>	8081	696	5	231	calcyclin binding protein (Cacybp) alternative variant aSep08, mRNA.
<a href="#">Cacybp</a>	<a href="#">Cacybp.cSep08</a>	<a href="#">289144</a>	4985	939	2	89	calcyclin binding protein (10.1 kD) (Cacybp) alternative variant cSep08, mRNA.

<a href="#">Cad</a>	<a href="#">Cad.bSep08</a>	<a href="#">24240</a>	2374	906	6	302	carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase (Cad) alternative variant bSep08, mRNA.
<a href="#">Cad</a>	<a href="#">Cad.cSep08</a>	<a href="#">24240</a>	1130	873	1	161	carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase (Cad) alternative variant cSep08, mRNA.
<a href="#">Cadherin.0</a>	<a href="#">Cadherin.0.aSep08</a>		116023	1038	6	208	cadherin 11 (Cadherin.0) alternative variant aSep08, mRNA.
<a href="#">Cadherin.1</a>	<a href="#">Cadherin.1.aSep08</a>		9735	466		155	cadherin 11 (Cadherin.1) mRNA.
<a href="#">Cadherin.2</a>	<a href="#">Cadherin.2.aSep08</a>		9391	1129		370	beta protocadherin 1 (Cadherin.2) mRNA.
<a href="#">Cadherin.3</a>	<a href="#">Cadherin.3.aSep08</a>		8741	1256	2	391	protocadherin 24 (Cadherin.3) alternative variant aSep08, mRNA.
<a href="#">Cadherin.3</a>	<a href="#">Cadherin.3.bSep08</a>		4534	742	1	90	protocadherin LKC CRA a (Cadherin.3) alternative variant bSep08, mRNA.
<a href="#">Cadherin.6</a>	<a href="#">Cadherin.6.aSep08</a>		36762	550		183	cadherin 18 type 2 (Cadherin.6) mRNA.
<a href="#">Cadherin_C.0</a>	<a href="#">Cadherin_C.0.aSep08</a>		29963	3499	6	397	cadherin 11 CRA b (Cadherin_C.0) alternative variant aSep08, mRNA.
<a href="#">Cadherin_pro.0</a>	<a href="#">Cadherin_pro.0.aSep08</a>		11846	827		182	desmocollin 2 (Cadherin_pro.0) mRNA.
<a href="#">Cadm1</a>	<a href="#">Cadm1.bSep08</a>	<a href="#">363058</a>	38926	606	6	201	cell adhesion molecule 1 (Cadm1) alternative variant bSep08, mRNA.
<a href="#">Cadm1</a>	<a href="#">Cadm1.cSep08</a>	<a href="#">363058</a>	53056	3529	6	189	cell adhesion molecule 1 (Cadm1) alternative variant cSep08, mRNA.
<a href="#">Cadm1</a>	<a href="#">Cadm1.dSep08</a>	<a href="#">363058</a>	35957	701	5	181	cell adhesion molecule 1 (Cadm1) alternative variant dSep08, mRNA.
<a href="#">Cadm1</a>	<a href="#">Cadm1.eSep08</a>	<a href="#">363058</a>	31315	687	3	99	cell adhesion molecule 1 (Cadm1) alternative variant eSep08, mRNA.
<a href="#">Cadm3</a>	<a href="#">Cadm3.bSep08</a>	<a href="#">360882</a>	22754	726	1	241	cell adhesion molecule 3 (Cadm3) alternative variant bSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.bSep08</a>	<a href="#">26989</a>	147441	2030	19	676	secretion activator (Cadps) alternative variant bSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.cSep08</a>	<a href="#">26989</a>	83601	941	6	276	secretion activator CRA a (Cadps) alternative variant cSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.dSep08</a>	<a href="#">26989</a>	86977	800	2	221	secretion activator CRA b (Cadps) alternative variant dSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.eSep08</a>	<a href="#">26989</a>	16329	683	4	194	secretion activator CRA e (Cadps) alternative variant eSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.fSep08</a>	<a href="#">26989</a>	23947	483	6	160	activator secretion (Cadps) alternative variant fSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.gSep08</a>	<a href="#">26989</a>	3523	645	2	98	secretion activator CRA b (Cadps) alternative variant gSep08, mRNA.
<a href="#">Cadps</a>	<a href="#">Cadps.iSep08</a>	<a href="#">26989</a>	591	276	2	59	putative protein (Cadps) alternative variant iSep08, mRNA.
<a href="#">Cadps2</a>	<a href="#">Cadps2.aSep08</a>	<a href="#">312166</a>	69570	1523	6	256	ca2+-dependent activator protein for secretion 2 (Cadps2) alternative variant aSep08, mRNA.
<a href="#">Cadps2</a>	<a href="#">Cadps2.bSep08</a>	<a href="#">312166</a>	29407	410	1	108	ca2+-dependent activator protein for secretion 2 (Cadps2) alternative variant bSep08, mRNA.
<a href="#">CAF1.0</a>	<a href="#">CAF1.0.aSep08</a>		23544	1119		256	poly-specific ribonuclease CRA c (CAF1.0) mRNA.

<a href="#">Calb1</a>	<a href="#">Calb1.bSep08</a>	<a href="#">83839</a>	23229	745	9	246	calbindin 1 (Calb1) alternative variant bSep08, mRNA.
<a href="#">Calb2</a>	<a href="#">Calb2.bSep08</a>	<a href="#">117059</a>	15510	597	1	113	calbindin 2 (12.8 kD) (Calb2) alternative variant bSep08, mRNA.
<a href="#">Calca</a>	<a href="#">Calca.aSep08</a>	<a href="#">24241</a>	4002	587	4	172	calcitonin/calcitonin-related polypeptide, alpha (Calca) alternative variant aSep08, mRNA.
<a href="#">Calcoco1</a>	<a href="#">Calcoco1.bSep08</a>	<a href="#">246047</a>	2872	412	3	78	calcium binding and coiled coil domain 1 (Calcoco1) alternative variant bSep08, mRNA.
<a href="#">Calcl</a>	<a href="#">Calcl.bSep08</a>	<a href="#">25029</a>	50974	1772	8	432	calcitonin receptor-like (Calcl) alternative variant bSep08, mRNA.
<a href="#">Calcl</a>	<a href="#">Calcl.cSep08</a>	<a href="#">25029</a>	54330	767	6	101	calcitonin receptor-like (Calcl) alternative variant cSep08, mRNA.
<a href="#">Calcl</a>	<a href="#">Calcl.dSep08</a>	<a href="#">25029</a>	2662	366	3	92	calcitonin receptor-like (Calcl) alternative variant dSep08, mRNA.
<a href="#">Cald1</a>	<a href="#">Cald1.bSep08</a>	<a href="#">25687</a>	149130	760	4	204	caldesmon 1 (Cald1) alternative variant bSep08, mRNA.
<a href="#">Cald1</a>	<a href="#">Cald1.cSep08</a>	<a href="#">25687</a>	2140	1098	2	141	caldesmon 1 (14.8 kD) (Cald1) alternative variant cSep08, mRNA.
<a href="#">Cald1</a>	<a href="#">Cald1.dSep08</a>	<a href="#">25687</a>	7289	418	3	139	caldesmon 1 (Cald1) alternative variant dSep08, mRNA.
<a href="#">Cald1</a>	<a href="#">Cald1.eSep08</a>	<a href="#">25687</a>	62374	577	4	86	caldesmon 1 (Cald1) alternative variant eSep08, mRNA.
<a href="#">Calm3</a>	<a href="#">Calm3.cSep08</a>	<a href="#">24244</a>	6759	474	1	119	calmodulin 3 (Calm3) alternative variant cSep08, mRNA.
<a href="#">Calr</a>	<a href="#">Calr.bSep08</a>	<a href="#">64202</a>	4338	1654	8	366	calreticulin (Calr) alternative variant bSep08, mRNA.
<a href="#">Calr</a>	<a href="#">Calr.cSep08</a>	<a href="#">64202</a>	2108	709	4	166	calreticulin (Calr) alternative variant cSep08, mRNA.
<a href="#">Calr</a>	<a href="#">Calr.dSep08</a>	<a href="#">64202</a>	580	414	3	137	calreticulin (Calr) alternative variant dSep08, mRNA.
<a href="#">Calr4</a>	<a href="#">Calr4.aSep08</a>	<a href="#">689537</a>	9199	1052		178	calreticulin 4 (21.2 kD) (Calr4) mRNA.
<a href="#">Calu</a>	<a href="#">Calu.cSep08</a>	<a href="#">64366</a>	17306	368		96	calumenin (Calu) alternative variant cSep08, mRNA.
<a href="#">Calu</a>	<a href="#">Calu.cSep08</a>	<a href="#">360380</a>	17306	368		96	calumenin (Calu) alternative variant cSep08, mRNA.
<a href="#">Calx-beta.0</a>	<a href="#">Calx-beta.0.aSep08</a>		3530	433		144	fras1 related extracellular matrix protein 2 (Calx-beta.0) alternative variant aSep08, mRNA.
<a href="#">Caly</a>	<a href="#">Caly.bSep08</a>	<a href="#">192349</a>	10584	636	5	212	calcyon neuron-specific vesicular protein (Caly) alternative variant bSep08, mRNA.
<a href="#">Camk1</a>	<a href="#">Camk1.aSep08</a>	<a href="#">171503</a>	6405	864		255	calcium/calmodulin-dependent protein kinase I (Camk1) alternative variant aSep08, mRNA.
<a href="#">Camk1</a>	<a href="#">Camk1.bSep08</a>	<a href="#">171503</a>	2062	498		165	calcium/calmodulin-dependent protein kinase I (Camk1) alternative variant bSep08, mRNA.
<a href="#">Camk1g</a>	<a href="#">Camk1g.bSep08</a>	<a href="#">171358</a>	2358	1003	3	88	calcium/calmodulin-dependent protein kinase I gamma (Camk1g) alternative variant bSep08, mRNA.
<a href="#">Camk1g</a>	<a href="#">Camk1g.cSep08</a>	<a href="#">171358</a>	1833	413	2	80	calcium/calmodulin-dependent protein kinase I gamma (Camk1g) alternative variant cSep08, mRNA.
<a href="#">Camk2a</a>	<a href="#">Camk2a.aSep08</a>	<a href="#">25400</a>	4000	517	5	152	calcium/calmodulin-dependent protein kinase II, alpha (Camk2a) alternative variant aSep08, mRNA.
<a href="#">Camk2a</a>	<a href="#">Camk2a.cSep08</a>	<a href="#">25400</a>	13947	573	4		
<a href="#">Camk2b</a>	<a href="#">Camk2b.dSep08</a>	<a href="#">24245</a>	4424	712	1	144	calcium/calmodulin-dependent protein kinase II, beta (Camk2b) alternative variant dSep08, mRNA.
<a href="#">Camk2d</a>	<a href="#">Camk2d.bSep08</a>	<a href="#">24246</a>	71597	1264	11	317	calcium/calmodulin-dependent protein kinase II, delta (Camk2d) alternative variant bSep08, mRNA.

<a href="#">Camk2d</a>	<a href="#">Camk2d.cSep08</a>	<a href="#">24246</a>	178286	1500	8	301	calcium/calmodulin-dependent protein kinase II, delta (Camk2d) alternative variant cSep08, mRNA.
<a href="#">Camk2d</a>	<a href="#">Camk2d.dSep08</a>	<a href="#">24246</a>	52823	340	6	112	calcium/calmodulin-dependent protein kinase II, delta (Camk2d) alternative variant dSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.bSep08</a>	<a href="#">171140</a>	36683	3341	17	451	calcium calmodulin-dependent protein kinase II gamma CRA o (Camk2g) alternative variant bSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.cSep08</a>	<a href="#">171140</a>	52690	1140	15	322	calcium calmodulin-dependent protein kinase II gamma (Camk2g) alternative variant cSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.dSep08</a>	<a href="#">171140</a>	28375	918	9	277	calcium calmodulin-dependent protein kinase II gamma (Camk2g) alternative variant dSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.eSep08</a>	<a href="#">171140</a>	23573	507	10	169	calcium calmodulin-dependent protein kinase II gamma (Camk2g) alternative variant eSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.fSep08</a>	<a href="#">171140</a>	1653	1153	2	137	putative protein (Camk2g) alternative variant fSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.gSep08</a>	<a href="#">171140</a>	8544	748	5	115	calcium calmodulin-dependent protein kinase II gamma (12.9 kD) (Camk2g) alternative variant gSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.hSep08</a>	<a href="#">171140</a>	816	325	2	107	putative protein (Camk2g) alternative variant hSep08, mRNA.
<a href="#">Camk2g</a>	<a href="#">Camk2g.jSep08</a>	<a href="#">171140</a>	10336	589	8	57	putative protein (6.2 kD) (Camk2g) alternative variant jSep08, mRNA.
<a href="#">Camk4</a>	<a href="#">Camk4.bSep08</a>	<a href="#">25050</a>	31582	695	5	166	calcium calmodulin-dependent protein kinase IV (Camk4) alternative variant bSep08, mRNA.
<a href="#">Camk4</a>	<a href="#">Camk4.cSep08</a>	<a href="#">25050</a>	113257	414	3	138	calcium calmodulin-dependent protein kinase IV (Camk4) alternative variant cSep08, mRNA.
<a href="#">Camk4</a>	<a href="#">Camk4.dSep08</a>	<a href="#">25050</a>	1088	733	2	89	calmodulin-dependent protein kinase IV (Camk4) alternative variant dSep08, mRNA.
<a href="#">CaMKII_AD.0</a>	<a href="#">CaMKII_AD.0.aSep08</a>		7975	3384		87	calcium calmodulin-dependent protein kinase II alpha (CaMKII_AD.0) mRNA.
<a href="#">Camkk1</a>	<a href="#">Camkk1.aSep08</a>	<a href="#">60341</a>	19505	1415	13	277	calcium/calmodulin-dependent protein kinase kinase 1, alpha (Camkk1) alternative variant aSep08, mRNA.
<a href="#">Camkk1</a>	<a href="#">Camkk1.bSep08</a>	<a href="#">60341</a>	4760	2235	3	144	calcium/calmodulin-dependent protein kinase kinase 1, alpha (Camkk1) alternative variant bSep08, mRNA.
<a href="#">Camkk1</a>	<a href="#">Camkk1.cSep08</a>	<a href="#">60341</a>	2765	738	1	108	calcium/calmodulin-dependent protein kinase kinase 1, alpha (Camkk1) alternative variant cSep08, mRNA.
<a href="#">Camkk2</a>	<a href="#">Camkk2.cSep08</a>	<a href="#">83506</a>	1230	448	2	41	calcium/calmodulin-dependent protein kinase kinase 2, beta (Camkk2) alternative variant cSep08, mRNA.
<a href="#">Camkk2</a>	<a href="#">Camkk2.dSep08</a>	<a href="#">83506</a>	6645	573	3		
<a href="#">Camkv</a>	<a href="#">Camkv.bSep08</a>	<a href="#">79011</a>	10078	620	1	149	CaM kinase-like vesicle-associated (Camkv) alternative variant bSep08, mRNA.
<a href="#">Camlg</a>	<a href="#">Camlg.aSep08</a>	<a href="#">81715</a>	10848	1384	4	295	calcium modulating ligand (32.8 kD) (Camlg) alternative variant aSep08, complete mRNA.
<a href="#">Camlg</a>	<a href="#">Camlg.cSep08</a>	<a href="#">81715</a>	10704	1652	4	229	calcium modulating ligand (25.5 kD) (Camlg) alternative variant cSep08, mRNA.
<a href="#">Camsap1</a>	<a href="#">Camsap1.aSep08</a>	<a href="#">296580</a>	12632	4136	6	886	calmodulin regulated spectrin-associated protein 1 (Camsap1) alternative variant aSep08, mRNA.
<a href="#">Camsap1</a>	<a href="#">Camsap1.bSep08</a>	<a href="#">296580</a>	6196	619	2	117	calmodulin regulated spectrin-associated protein 1 (Camsap1) alternative variant bSep08, mRNA.

<a href="#">Camsap1</a>	<a href="#">Camsap1.cSep08</a>	<a href="#">296580</a>	5500	455	1	58	calmodulin regulated spectrin-associated protein 1 (6.4 kD) (Camsap1) alternative variant cSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.bSep08</a>	<a href="#">287462</a>	4895	1146	7	381	calmodulin binding transcription activator 2 like (Camta2andSpag7) alternative variant bSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.bSep08</a>	<a href="#">303260</a>	4895	1146	7	381	calmodulin binding transcription activator 2 like (Camta2andSpag7) alternative variant bSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.dSep08</a>	<a href="#">287462</a>	4926	656	7	180	calmodulin binding transcription activator 2 like (Camta2andSpag7) alternative variant dSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.dSep08</a>	<a href="#">303260</a>	4926	656	7	180	calmodulin binding transcription activator 2 like (Camta2andSpag7) alternative variant dSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.eSep08</a>	<a href="#">287462</a>	5697	810	7	176	sperm associated antigen 7 like (20.1 kD) (Camta2andSpag7) alternative variant eSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.eSep08</a>	<a href="#">303260</a>	5697	810	7	176	sperm associated antigen 7 like (20.1 kD) (Camta2andSpag7) alternative variant eSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.fSep08</a>	<a href="#">287462</a>	857	371	3	95	calmodulin binding transcription activator 2 CRA d like (Camta2andSpag7) alternative variant fSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.fSep08</a>	<a href="#">303260</a>	857	371	3	95	calmodulin binding transcription activator 2 CRA d like (Camta2andSpag7) alternative variant fSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.gSep08</a>	<a href="#">287462</a>	899	802	2	46	sperm associated antigen 7 like (5.2 kD) (Camta2andSpag7) alternative variant gSep08, mRNA.
<a href="#">Camta2andSpag7</a>	<a href="#">Camta2andSpag7.gSep08</a>	<a href="#">303260</a>	899	802	2	46	sperm associated antigen 7 like (5.2 kD) (Camta2andSpag7) alternative variant gSep08, mRNA.
<a href="#">Cand2</a>	<a href="#">Cand2.aSep08</a>	<a href="#">192226</a>	13292	3408	6	677	cullin-associated and neddylation-dissociated 2 (putative) (Cand2) alternative variant aSep08, mRNA.
<a href="#">Cand2</a>	<a href="#">Cand2.bSep08</a>	<a href="#">192226</a>	13159	963	3	34	cullin-associated and neddylation-dissociated 2 (putative) (Cand2) alternative variant bSep08, mRNA.
<a href="#">Cant1</a>	<a href="#">Cant1.bSep08</a>	<a href="#">246272</a>	9425	647	2	215	calcium activated nucleotidase 1 (Cant1) alternative variant bSep08, mRNA.
<a href="#">Cant1</a>	<a href="#">Cant1.cSep08</a>	<a href="#">246272</a>	5531	404	2	134	calcium activated nucleotidase 1 (Cant1) alternative variant cSep08, mRNA.
<a href="#">Canx</a>	<a href="#">Canx.bSep08</a>	<a href="#">29144</a>	24430	1618	9	343	calnexin precursor (39.0 kD) (Canx) alternative variant bSep08, mRNA.
<a href="#">Canx</a>	<a href="#">Canx.cSep08</a>	<a href="#">29144</a>	18375	853	7	235	calnexin (Canx) alternative variant cSep08, mRNA.
<a href="#">Canx</a>	<a href="#">Canx.dSep08</a>	<a href="#">29144</a>	3835	896	5	193	calnexin (Canx) alternative variant dSep08, mRNA.
<a href="#">Canx</a>	<a href="#">Canx.eSep08</a>	<a href="#">29144</a>	18201	711	7	187	calnexin (Canx) alternative variant eSep08, mRNA.
<a href="#">Canx</a>	<a href="#">Canx.fSep08</a>	<a href="#">29144</a>	6639	3703	5	175	calnexin (19.8 kD) (Canx) alternative variant fSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.aSep08</a>	<a href="#">64185</a>	26691	2574	6	474	CAP, adenylate cyclase-associated protein 1 (yeast) (51.6 kD) (Cap1) alternative variant aSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.bSep08</a>	<a href="#">64185</a>	21620	848	3	282	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant bSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.cSep08</a>	<a href="#">64185</a>	21521	771	3	239	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant cSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.dSep08</a>	<a href="#">64185</a>	21449	696	3	214	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant dSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.eSep08</a>	<a href="#">64185</a>	18561	495	3	165	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant eSep08, mRNA.

<a href="#">Cap1</a>	<a href="#">Cap1.fSep08</a>	<a href="#">64185</a>	2590	745	4	139	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant fSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.gSep08</a>	<a href="#">64185</a>	1097	814	1	118	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant gSep08, mRNA.
<a href="#">Cap1</a>	<a href="#">Cap1.hSep08</a>	<a href="#">64185</a>	752	552	2	43	CAP, adenylate cyclase-associated protein 1 (yeast) (Cap1) alternative variant hSep08, mRNA.
<a href="#">Capg</a>	<a href="#">Capg.bSep08</a>	<a href="#">297339</a>	13386	692	7	199	capping protein (actin filament), gelsolin-like (Capg) alternative variant bSep08, mRNA.
<a href="#">Capg</a>	<a href="#">Capg.cSep08</a>	<a href="#">297339</a>	6199	442	5	146	capping protein (actin filament), gelsolin-like (Capg) alternative variant cSep08, mRNA.
<a href="#">Capg</a>	<a href="#">Capg.dSep08</a>	<a href="#">297339</a>	11880	509	6	130	capping protein (actin filament), gelsolin-like (Capg) alternative variant dSep08, mRNA.
<a href="#">Capg</a>	<a href="#">Capg.eSep08</a>	<a href="#">297339</a>	1547	411	2	98	capping protein (actin filament), gelsolin-like (Capg) alternative variant eSep08, mRNA.
<a href="#">Capn1</a>	<a href="#">Capn1.aSep08</a>	<a href="#">29153</a>	21088	1855	18	618	calpain 1 (Capn1) alternative variant aSep08, mRNA.
<a href="#">Capn1</a>	<a href="#">Capn1.bSep08</a>	<a href="#">29153</a>	4382	772	6	230	calpain 1 (Capn1) alternative variant bSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.bSep08</a>	<a href="#">29154</a>	15016	703	6	180	calpain 2 (Capn2) alternative variant bSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.bSep08</a>	<a href="#">690477</a>	15016	703	6	180	calpain 2 (Capn2) alternative variant bSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.cSep08</a>	<a href="#">29154</a>	9079	690	7	121	calpain 2 (14.2 kD) (Capn2) alternative variant cSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.cSep08</a>	<a href="#">690477</a>	9079	690	7	121	calpain 2 (14.2 kD) (Capn2) alternative variant cSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.eSep08</a>	<a href="#">29154</a>	21626	478	3	92	putative protein (Capn2) alternative variant eSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.eSep08</a>	<a href="#">690477</a>	21626	478	3	92	putative protein (Capn2) alternative variant eSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.fSep08</a>	<a href="#">29154</a>	5135	521	3	88	calpain 2 (Capn2) alternative variant fSep08, mRNA.
<a href="#">Capn2</a>	<a href="#">Capn2.fSep08</a>	<a href="#">690477</a>	5135	521	3	88	calpain 2 (Capn2) alternative variant fSep08, mRNA.
<a href="#">Capn3</a>	<a href="#">Capn3.bSep08</a>	<a href="#">29155</a>	8521	718	1	239	calpain 3 (Capn3) alternative variant bSep08, mRNA.
<a href="#">Capn5</a>	<a href="#">Capn5.bSep08</a>	<a href="#">171495</a>	11345	354	3	84	calpain 5 (Capn5) alternative variant bSep08, mRNA.
<a href="#">Capn7</a>	<a href="#">Capn7.bSep08</a>	<a href="#">306260</a>	2329	743	4	160	calpain 7 (Capn7) alternative variant bSep08, mRNA.
<a href="#">Capn7</a>	<a href="#">Capn7.cSep08</a>	<a href="#">306260</a>	2316	401	2	107	putative protein (Capn7) alternative variant cSep08, mRNA.
<a href="#">Capn7</a>	<a href="#">Capn7.dSep08</a>	<a href="#">306260</a>	10549	713	4	103	calpain 7 (Capn7) alternative variant dSep08, mRNA.
<a href="#">Capn7</a>	<a href="#">Capn7.eSep08</a>	<a href="#">306260</a>	2980	350	3	79	calpain 7 (Capn7) alternative variant eSep08, mRNA.
<a href="#">Capn12</a>	<a href="#">Capn12.bSep08</a>	<a href="#">308476</a>	1249	633	4	210	calpain 12 (Capn12) alternative variant bSep08, mRNA.
<a href="#">Capn13</a>	<a href="#">Capn13.bSep08</a>	<a href="#">362701</a>	3577	294	1	89	calpain 13 (Capn13) alternative variant bSep08, mRNA.
<a href="#">Capns1</a>	<a href="#">Capns1.bSep08</a>	<a href="#">29156</a>	6052	675	8	224	calpain, small subunit 1 (Capns1) alternative variant bSep08, mRNA.
<a href="#">Capns1</a>	<a href="#">Capns1.cSep08</a>	<a href="#">29156</a>	1015	400	1	104	calpain, small subunit 1 (Capns1) alternative variant cSep08, mRNA.
<a href="#">Caprin1</a>	<a href="#">Caprin1.bSep08</a>	<a href="#">362173</a>	25476	1381	11	460	cell cycle associated protein 1 (Caprin1) alternative variant bSep08, mRNA.
<a href="#">Caprin1</a>	<a href="#">Caprin1.cSep08</a>	<a href="#">362173</a>	7651	1084	6	349	cell cycle associated protein 1 (Caprin1) alternative variant cSep08, mRNA.

<a href="#">Caprin1</a>	<a href="#">Caprin1.dSep08</a>	<a href="#">362173</a>	5043	1837	4	178	cell cycle associated protein 1 (Caprin1) alternative variant dSep08, mRNA.
<a href="#">Caprin1</a>	<a href="#">Caprin1.eSep08</a>	<a href="#">362173</a>	1002	489	2	163	cell cycle associated protein 1 (Caprin1) alternative variant eSep08, mRNA.
<a href="#">Caprin1</a>	<a href="#">Caprin1.fSep08</a>	<a href="#">362173</a>	3703	772	3	133	cell cycle associated protein 1 (Caprin1) alternative variant fSep08, mRNA.
<a href="#">Caprin1</a>	<a href="#">Caprin1.gSep08</a>	<a href="#">362173</a>	1842	1266	3	50	cell cycle associated protein 1 (5.2 kD) (Caprin1) alternative variant gSep08, mRNA.
<a href="#">Caps2</a>	<a href="#">Caps2.bSep08</a>	<a href="#">366891</a>	28244	636	3	211	calcyphosphine 2 (Caps2) alternative variant bSep08, mRNA.
<a href="#">Caps2</a>	<a href="#">Caps2.cSep08</a>	<a href="#">366891</a>	16395	755	2	175	calcyphosphine 2 (Caps2) alternative variant cSep08, mRNA.
<a href="#">Capza1</a>	<a href="#">Capza1.bSep08</a>	<a href="#">310756</a>	2781	693	5	230	capping protein (actin filament) muscle Z-line, alpha 1 (Capza1) alternative variant bSep08, mRNA.
<a href="#">Capza1</a>	<a href="#">Capza1.cSep08</a>	<a href="#">310756</a>	2301	978	6	216	capping protein (actin filament) muscle Z-line, alpha 1 (Capza1) alternative variant cSep08, mRNA.
<a href="#">Capza1</a>	<a href="#">Capza1.dSep08</a>	<a href="#">310756</a>	813	683	2	140	capping protein (actin filament) muscle Z-line, alpha 1 (15.8 kD) (Capza1) alternative variant dSep08, mRNA.
<a href="#">Capza1</a>	<a href="#">Capza1.eSep08</a>	<a href="#">310756</a>	645	391	2	129	capping protein (actin filament) muscle Z-line, alpha 1 (Capza1) alternative variant eSep08, mRNA.
<a href="#">Capza1</a>	<a href="#">Capza1.fSep08</a>	<a href="#">310756</a>	1183	1083	2	49	capping protein (actin filament) muscle Z-line, alpha 1 (5.1 kD) (Capza1) alternative variant fSep08, mRNA.
<a href="#">Capza2</a>	<a href="#">Capza2.aSep08</a>	<a href="#">493810</a>	36049	2464	6	309	capping protein (actin filament) muscle Z-line, alpha 2 (Capza2) alternative variant aSep08, mRNA.
<a href="#">Capza2</a>	<a href="#">Capza2.bSep08</a>	<a href="#">493810</a>	19300	735	1	161	capping protein (actin filament) muscle Z-line, alpha 2 (18.5 kD) (Capza2) alternative variant bSep08, mRNA.
<a href="#">Capzb</a>	<a href="#">Capzb.bSep08</a>	<a href="#">298584</a>	99228	1161	10	277	capping protein (actin filament) muscle Z-line, beta (31.3 kD) (Capzb) alternative variant bSep08, complete mRNA.
<a href="#">Capzb</a>	<a href="#">Capzb.cSep08</a>	<a href="#">298584</a>	96891	768	8	239	capping protein (actin filament) muscle Z-line, beta (27.0 kD) (Capzb) alternative variant cSep08, mRNA.
<a href="#">Capzb</a>	<a href="#">Capzb.dSep08</a>	<a href="#">298584</a>	64760	733	3	85	capping protein (actin filament) muscle Z-line, beta (Capzb) alternative variant dSep08, mRNA.
<a href="#">CAP_GLY.0</a>	<a href="#">CAP_GLY.0.aSep08</a>		10584	963		254	ubiquitin carboxyl-terminal hydrolase cyld (CAP_GLY.0) mRNA.
<a href="#">Car2</a>	<a href="#">Car2.bSep08</a>	<a href="#">54231</a>	14326	742	7	247	carbonic anhydrase II and hypothetical protein LOC686037 (Car2) alternative variant bSep08, mRNA.
<a href="#">Car2</a>	<a href="#">Car2.bSep08</a>	<a href="#">686037</a>	14326	742	7	247	carbonic anhydrase II and hypothetical protein LOC686037 (Car2) alternative variant bSep08, mRNA.
<a href="#">Car2</a>	<a href="#">Car2.dSep08</a>	<a href="#">54231</a>	7772	378	3	99	carbonic anhydrase II and hypothetical protein LOC686037 (10.8 kD) (Car2) alternative variant dSep08, complete mRNA.
<a href="#">Car2</a>	<a href="#">Car2.dSep08</a>	<a href="#">686037</a>	7772	378	3	99	carbonic anhydrase II and hypothetical protein LOC686037 (10.8 kD) (Car2) alternative variant dSep08, complete mRNA.
<a href="#">Car4</a>	<a href="#">Car4.bSep08</a>	<a href="#">29242</a>	6441	1581	6	184	carbonic anhydrase 4 (20.7 kD) (Car4) alternative variant bSep08, mRNA.



<a href="#">Car5b</a>	<a href="#">Car5b.bSep08</a>	<a href="#">302669</a>	26436	680	1	167	carbonic anhydrase 5b, mitochondrial (Car5b) alternative variant bSep08, mRNA.
<a href="#">Car5b</a>	<a href="#">Car5b.cSep08</a>	<a href="#">302669</a>	33239	804	1	167	carbonic anhydrase 5b, mitochondrial (19.5 kD) (Car5b) alternative variant cSep08, mRNA.
<a href="#">Car6</a>	<a href="#">Car6.bSep08</a>	<a href="#">298657</a>	7531	738	4	91	carbonic anhydrase 6 (Car6) alternative variant bSep08, mRNA.
<a href="#">Car6</a>	<a href="#">Car6.cSep08</a>	<a href="#">298657</a>	3980	522	2	56	carbonic anhydrase 6 (Car6) alternative variant cSep08, mRNA.
<a href="#">Car7</a>	<a href="#">Car7.bSep08</a>	<a href="#">291819</a>	2482	421	1	140	carbonic anhydrase 7 (Car7) alternative variant bSep08, mRNA.
<a href="#">Car11</a>	<a href="#">Car11.bSep08</a>	<a href="#">308588</a>	4142	828	7	252	carbonic anhydrase 11 (27.9 kD) (Car11) alternative variant bSep08, mRNA.
<a href="#">Car11</a>	<a href="#">Car11.cSep08</a>	<a href="#">308588</a>	5296	1637	8	193	carbonic anhydrase 11 (20.8 kD) (Car11) alternative variant cSep08, mRNA.
<a href="#">Car11</a>	<a href="#">Car11.dSep08</a>	<a href="#">308588</a>	4813	1410	5	177	carbonic anhydrase 11 (Car11) alternative variant dSep08, mRNA.
<a href="#">Car12</a>	<a href="#">Car12.bSep08</a>	<a href="#">363085</a>	38059	1271	10	336	carbonic anhydrase 12 (37.4 kD) (Car12) alternative variant bSep08, mRNA.
<a href="#">Car12</a>	<a href="#">Car12.cSep08</a>	<a href="#">363085</a>	12448	686	4	127	carbonic anhydrase 12 (Car12) alternative variant cSep08, mRNA.
<a href="#">Car13</a>	<a href="#">Car13.aSep08</a>	<a href="#">499566</a>	37411	1964		262	carbonic anhydrase 13 (29.6 kD) (Car13) mRNA.
<a href="#">Car14</a>	<a href="#">Car14.aSep08</a>	<a href="#">791259</a>	5386	934	9	311	carbonic anhydrase 14 (Car14) alternative variant aSep08, mRNA.
<a href="#">Car14</a>	<a href="#">Car14.bSep08</a>	<a href="#">791259</a>	2824	658	6	219	carbonic anhydrase 14 (Car14) alternative variant bSep08, mRNA.
<a href="#">Car14</a>	<a href="#">Car14.cSep08</a>	<a href="#">791259</a>	4915	842	6	176	carbonic anhydrase 14 (Car14) alternative variant cSep08, mRNA.
<a href="#">Car14</a>	<a href="#">Car14.dSep08</a>	<a href="#">791259</a>	1037	372	2	97	carbonic anhydrase 14 (Car14) alternative variant dSep08, mRNA.
<a href="#">Car14</a>	<a href="#">Car14.eSep08</a>	<a href="#">791259</a>	2708	531	2	57	carbonic anhydrase 14 (6.2 kD) (Car14) alternative variant eSep08, complete mRNA.
<a href="#">Car14</a>	<a href="#">Car14.fSep08</a>	<a href="#">791259</a>	6943	1784	3	55	carbonic anhydrase 14 (Car14) alternative variant fSep08, mRNA.
<a href="#">Car15</a>	<a href="#">Car15.bSep08</a>	<a href="#">288360</a>	1177	753	2	82	carbonic anhydrase 15 (9.2 kD) (Car15) alternative variant bSep08, mRNA.
<a href="#">CARD.0</a>	<a href="#">CARD.0.aSep08</a>		1135	732		110	NACHT- LRR- PYD-containing protein 1 paralog B (CARD.0) mRNA.
<a href="#">Card9</a>	<a href="#">Card9.bSep08</a>	<a href="#">64171</a>	3317	663	6	177	caspase recruitment domain family, member 9 (Card9) alternative variant bSep08, mRNA.
<a href="#">Card10</a>	<a href="#">Card10.bSep08</a>	<a href="#">315120</a>	2809	739	5	234	caspase recruitment domain family, member 10 (Card10) alternative variant bSep08, mRNA.
<a href="#">Carf</a>	<a href="#">Carf.bSep08</a>	<a href="#">301446</a>	27314	958	7	150	putative protein of vertebrate origin (Carf) alternative variant bSep08, mRNA.
<a href="#">Carf</a>	<a href="#">Carf.cSep08</a>	<a href="#">301446</a>	7312	429	4	107	putative protein of metazoan origin (Carf) alternative variant cSep08, mRNA.
<a href="#">Carhsp1</a>	<a href="#">Carhsp1.bSep08</a>	<a href="#">260416</a>	8393	322	3	107	calcium regulated heat stable protein 1 (Carhsp1) alternative variant bSep08, mRNA.

<a href="#">Carhsp1</a>	<a href="#">Carhsp1.cSep08</a>	<a href="#">260416</a>	11300	446	1	88	calcium regulated heat stable protein 1 (9.2 kD) (Carhsp1) alternative variant cSep08, complete mRNA.
<a href="#">Carm1</a>	<a href="#">Carm1.cSep08</a>	<a href="#">363026</a>	8579	733	1	209	coactivator-associated arginine methyltransferase 1 (Carm1) alternative variant cSep08, mRNA.
<a href="#">Cars</a>	<a href="#">Cars.bSep08</a>	<a href="#">293638</a>	37952	2024	19	650	cysteinyI-tRNA synthetase (Cars) alternative variant bSep08, mRNA.
<a href="#">Cars</a>	<a href="#">Cars.cSep08</a>	<a href="#">293638</a>	42217	1949	8	539	cysteinyI-tRNA synthetase (Cars) alternative variant cSep08, mRNA.
<a href="#">Cars</a>	<a href="#">Cars.eSep08</a>	<a href="#">293638</a>	1349	697	4	128	cysteinyI-tRNA synthetase (Cars) alternative variant eSep08, mRNA.
<a href="#">Cartpt</a>	<a href="#">Cartpt.bSep08</a>	<a href="#">29131</a>	1695	475	2	122	CART prepropeptide (Cartpt) alternative variant bSep08, mRNA.
<a href="#">Casc1</a>	<a href="#">Casc1.bSep08</a>	<a href="#">297720</a>	6830	1533	4	348	cancer susceptibility candidate 1 (Casc1) alternative variant bSep08, mRNA.
<a href="#">Casc1</a>	<a href="#">Casc1.cSep08</a>	<a href="#">297720</a>	7810	556	3	66	cancer susceptibility candidate 1 (Casc1) alternative variant cSep08, mRNA.
<a href="#">Casc3</a>	<a href="#">Casc3.bSep08</a>	<a href="#">259170</a>	10066	1111	6	369	cancer susceptibility candidate 3 (Casc3) alternative variant bSep08, mRNA.
<a href="#">Casc3</a>	<a href="#">Casc3.dSep08</a>	<a href="#">259170</a>	416	263	2	45	cancer susceptibility candidate 3 (Casc3) alternative variant dSep08, mRNA.
<a href="#">Casc4</a>	<a href="#">Casc4.aSep08</a>	<a href="#">362204</a>	33097	2862	2	171	cancer susceptibility candidate 4 (Casc4) alternative variant aSep08, mRNA.
<a href="#">Casc4</a>	<a href="#">Casc4.bSep08</a>	<a href="#">362204</a>	25182	929	1	153	cancer susceptibility candidate 4 (Casc4) alternative variant bSep08, mRNA.
<a href="#">Casc5</a>	<a href="#">Casc5.aSep08</a>	<a href="#">311327</a>	16879	1791		543	cancer susceptibility candidate 5 (Casc5) alternative variant aSep08, mRNA.
<a href="#">Casc5</a>	<a href="#">Casc5.bSep08</a>	<a href="#">311327</a>	25932	1200		299	cancer susceptibility candidate 5 (35.0 kD) (Casc5) alternative variant bSep08, mRNA.
<a href="#">Cask</a>	<a href="#">Cask.bSep08</a>	<a href="#">29647</a>	44661	2569	12	464	calcium/calmodulin-dependent serine protein kinase (MAGUK family) (Cask) alternative variant bSep08, mRNA.
<a href="#">Cask</a>	<a href="#">Cask.cSep08</a>	<a href="#">29647</a>	287403	1197	12	334	calcium/calmodulin-dependent serine protein kinase (MAGUK family) (Cask) alternative variant cSep08, mRNA.
<a href="#">Cask</a>	<a href="#">Cask.dSep08</a>	<a href="#">29647</a>	747	653	2	44	calcium/calmodulin-dependent serine protein kinase (MAGUK family) (4.6 kD) (Cask) alternative variant dSep08, mRNA.
<a href="#">Caskin1</a>	<a href="#">Caskin1.bSep08</a>	<a href="#">140722</a>	2232	2151	2	613	cask interacting protein 1 CRA b (Caskin1) alternative variant bSep08, mRNA.
<a href="#">Caskin1</a>	<a href="#">Caskin1.cSep08</a>	<a href="#">140722</a>	791	605	3	170	cask interacting protein 1 CRA c (Caskin1) alternative variant cSep08, mRNA.
<a href="#">Caskin1</a>	<a href="#">Caskin1.dSep08</a>	<a href="#">140722</a>	678	588	2	139	cask interacting protein 1 CRA b (Caskin1) alternative variant dSep08, mRNA.
<a href="#">Caskin2</a>	<a href="#">Caskin2.bSep08</a>	<a href="#">303678</a>	1711	544	6	181	cask-interacting protein 2 (Caskin2) alternative variant bSep08, mRNA.
<a href="#">Casp1</a>	<a href="#">Casp1.bSep08</a>	<a href="#">25166</a>	3475	670	1	183	caspase 1 (Casp1) alternative variant bSep08, mRNA.
<a href="#">Casp2</a>	<a href="#">Casp2.bSep08</a>	<a href="#">64314</a>	2749	2271	2	63	caspase 2 (7.2 kD) (Casp2) alternative variant bSep08, mRNA.

<a href="#">Casp3</a>	<a href="#">Casp3.aSep08</a>	<a href="#">25402</a>	16869	1141	6	288	caspace 3, apoptosis related cysteine protease (Casp3) alternative variant aSep08, mRNA.
<a href="#">Casp3</a>	<a href="#">Casp3.cSep08</a>	<a href="#">25402</a>	1401	413	1	100	caspace 3, apoptosis related cysteine protease (Casp3) alternative variant cSep08, mRNA.
<a href="#">Casp4</a>	<a href="#">Casp4.bSep08</a>	<a href="#">114555</a>	25250	1421	3	141	caspace 4, apoptosis-related cysteine peptidase (16.4 kD) (Casp4) alternative variant bSep08, mRNA.
<a href="#">Casp4</a>	<a href="#">Casp4.cSep08</a>	<a href="#">114555</a>	21617	413	1	126	caspace 4, apoptosis-related cysteine peptidase (Casp4) alternative variant cSep08, mRNA.
<a href="#">Casp4</a>	<a href="#">Casp4.dSep08</a>	<a href="#">114555</a>	700	299	2	32	caspace 4, apoptosis-related cysteine peptidase (3.8 kD) (Casp4) alternative variant dSep08, mRNA.
<a href="#">Casp6</a>	<a href="#">Casp6.bSep08</a>	<a href="#">83584</a>	11619	704	7	217	caspace 6 (Casp6) alternative variant bSep08, mRNA.
<a href="#">Casp6</a>	<a href="#">Casp6.cSep08</a>	<a href="#">83584</a>	5266	789	4	153	caspace 6 (17.7 kD) (Casp6) alternative variant cSep08, mRNA.
<a href="#">Casp7</a>	<a href="#">Casp7.bSep08</a>	<a href="#">64026</a>	14596	601	1	67	caspace 7 (Casp7) alternative variant bSep08, mRNA.
<a href="#">Casp12</a>	<a href="#">Casp12.bSep08</a>	<a href="#">156117</a>	11408	1890	2	325	caspace 12 (Casp12) alternative variant bSep08, mRNA.
<a href="#">Casq1</a>	<a href="#">Casq1.aSep08</a>	<a href="#">686019</a>	6964	1789	2	468	calsequestrin 1 (Casq1) alternative variant aSep08, mRNA.
<a href="#">Casq1</a>	<a href="#">Casq1.bSep08</a>	<a href="#">686019</a>	5102	1089	4	335	calsequestrin 1 (Casq1) alternative variant bSep08, mRNA.
<a href="#">Casq1</a>	<a href="#">Casq1.cSep08</a>	<a href="#">686019</a>	3780	887	5	140	calsequestrin 1 (Casq1) alternative variant cSep08, mRNA.
<a href="#">Casq1</a>	<a href="#">Casq1.dSep08</a>	<a href="#">686019</a>	1557	581	1	104	calsequestrin 1 (Casq1) alternative variant dSep08, mRNA.
<a href="#">Cast</a>	<a href="#">Cast.aSep08</a>	<a href="#">25403</a>	111473	2667	29	800	calpastatin (86.4 kD) (Cast) alternative variant aSep08, complete mRNA.
<a href="#">Cast</a>	<a href="#">Cast.bSep08</a>	<a href="#">25403</a>	111348	2473	28	777	calpastatin (83.8 kD) (Cast) alternative variant bSep08, complete mRNA.
<a href="#">Cast</a>	<a href="#">Cast.cSep08</a>	<a href="#">25403</a>	111348	2389	27	749	calpastatin (80.8 kD) (Cast) alternative variant cSep08, complete mRNA.
<a href="#">Cast</a>	<a href="#">Cast.fSep08</a>	<a href="#">25403</a>	37138	935	10	299	calpastatin (Cast) alternative variant fSep08, mRNA.
<a href="#">Cast</a>	<a href="#">Cast.gSep08</a>	<a href="#">25403</a>	59099	711	9	237	calpastatin (Cast) alternative variant gSep08, mRNA.
<a href="#">Cast</a>	<a href="#">Cast.hSep08</a>	<a href="#">25403</a>	55319	730	7	187	calpastatin (Cast) alternative variant hSep08, mRNA.
<a href="#">Cast</a>	<a href="#">Cast.iSep08</a>	<a href="#">25403</a>	7483	404	6	117	calpastatin (Cast) alternative variant iSep08, mRNA.
<a href="#">Cat</a>	<a href="#">Cat.bSep08</a>	<a href="#">24248</a>	12464	834	5	200	catalase (Cat) alternative variant bSep08, mRNA.
<a href="#">Cat</a>	<a href="#">Cat.cSep08</a>	<a href="#">24248</a>	602	258	2	53	catalase (Cat) alternative variant cSep08, complete mRNA.
<a href="#">Catsper2</a>	<a href="#">Catsper2.bSep08</a>	<a href="#">366174</a>	3300	471	2	50	cation channel, sperm associated 2 (6.3 kD) (Catsper2) alternative variant bSep08, mRNA.
<a href="#">Catsper4</a>	<a href="#">Catsper4.aSep08</a>	<a href="#">362623</a>	13406	728		242	channel, sperm associated 4 (Catsper4) mRNA.
<a href="#">Cav1</a>	<a href="#">Cav1.aSep08</a>	<a href="#">25404</a>	31048	655	3	203	caveolin, caveolae protein 1 (Cav1) alternative variant aSep08, mRNA.
<a href="#">Cav1</a>	<a href="#">Cav1.dSep08</a>	<a href="#">25404</a>	36833	736	3	146	caveolin, caveolae protein 1 (Cav1) alternative variant dSep08, mRNA.
<a href="#">Cav2</a>	<a href="#">Cav2.aSep08</a>	<a href="#">363425</a>	7479	2591	2	162	caveolin 2 (18.2 kD) (Cav2) alternative variant aSep08, mRNA.
<a href="#">Cav2</a>	<a href="#">Cav2.bSep08</a>	<a href="#">363425</a>	4899	377	1	125	caveolin 2 (Cav2) alternative variant bSep08, mRNA.
<a href="#">Cav2.1</a>	<a href="#">Cav2.1.aSep08</a>	<a href="#">363425</a>	7479	2591	2	162	caveolin 2 (18.2 kD) (Cav2.1) alternative variant aSep08, mRNA.
<a href="#">Cav2.1</a>	<a href="#">Cav2.1.bSep08</a>	<a href="#">363425</a>	4899	377	1	125	caveolin 2 (Cav2.1) alternative variant bSep08, mRNA.

<a href="#">CB741658</a>	<a href="#">CB741658.aSep08</a>	<a href="#">415051</a>	1652	1285	2	355	CB741658 gene (CB741658) alternative variant aSep08, mRNA.
<a href="#">CB741658</a>	<a href="#">CB741658.cSep08</a>	<a href="#">415051</a>	3252	454	2	150	CB741658 gene (CB741658) alternative variant cSep08, mRNA.
<a href="#">CB741658</a>	<a href="#">CB741658.dSep08</a>	<a href="#">415051</a>	5416	411	2	59	CB741658 gene (CB741658) alternative variant dSep08, mRNA.
<a href="#">Cbfa2t2</a>	<a href="#">Cbfa2t2.aSep08</a>	<a href="#">296293</a>	12160	752		173	core-binding factor, runt domain, alpha subunit 2, translocated to, 2 (human) (18.4 kD) (Cbfa2t2) mRNA.
<a href="#">Cbfa2t3</a>	<a href="#">Cbfa2t3.bSep08</a>	<a href="#">361431</a>	39359	600	5	200	core-binding factor, runt domain, alpha subunit 2, translocated to, 3 (human) (Cbfa2t3) alternative variant bSep08, mRNA.
<a href="#">Cbfb</a>	<a href="#">Cbfb.aSep08</a>	<a href="#">361391</a>	41642	949	2	186	core binding factor beta (22.0 kD) (Cbfb) alternative variant aSep08, mRNA.
<a href="#">Cbl</a>	<a href="#">Cbl.aSep08</a>	<a href="#">500985</a>	6907	626		208	casitas B-lineage lymphoma (Cbl) mRNA.
<a href="#">Cbhc</a>	<a href="#">Cbhc.bSep08</a>	<a href="#">292699</a>	5460	532	5	156	casitas B-lineage lymphoma c (Cbhc) alternative variant bSep08, mRNA.
<a href="#">Cbhc</a>	<a href="#">Cbhc.cSep08</a>	<a href="#">292699</a>	4289	422	4	134	casitas B-lineage lymphoma c (Cbhc) alternative variant cSep08, mRNA.
<a href="#">Cbl1</a>	<a href="#">Cbl1.bSep08</a>	<a href="#">314028</a>	11049	1113	4	370	casitas B-lineage lymphoma-like 1 (Cbl1) alternative variant bSep08, mRNA.
<a href="#">Cbl1</a>	<a href="#">Cbl1.cSep08</a>	<a href="#">314028</a>	6390	741	3	246	casitas B-lineage lymphoma-like 1 (Cbl1) alternative variant cSep08, mRNA.
<a href="#">Cbr1</a>	<a href="#">Cbr1.bSep08</a>	<a href="#">29224</a>	1526	1056	2	207	carbonyl reductase 1 (22.9 kD) (Cbr1) alternative variant bSep08, mRNA.
<a href="#">Cbr4</a>	<a href="#">Cbr4.aSep08</a>	<a href="#">359725</a>	6201	448	3	132	carbonyl reductase 4 (Cbr4) alternative variant aSep08, mRNA.
<a href="#">Cbr4</a>	<a href="#">Cbr4.bSep08</a>	<a href="#">359725</a>	12747	605	2	82	carbonyl reductase 4 (Cbr4) alternative variant bSep08, mRNA.
<a href="#">Cbs</a>	<a href="#">Cbs.bSep08</a>	<a href="#">24250</a>	8686	782	6	220	cystathionine beta synthase (Cbs) alternative variant bSep08, mRNA.
<a href="#">Cbs</a>	<a href="#">Cbs.cSep08</a>	<a href="#">24250</a>	2710	632	1	122	cystathionine beta synthase (Cbs) alternative variant cSep08, mRNA.
<a href="#">Cbwd1</a>	<a href="#">Cbwd1.bSep08</a>	<a href="#">171057</a>	15745	401	6	133	cobalamin (vitamin B12) biosynthesis P47K (Cbwd1) alternative variant bSep08, mRNA.
<a href="#">Cbwd1</a>	<a href="#">Cbwd1.cSep08</a>	<a href="#">171057</a>	5749	555	2	41	putative protein of vertebrate origin (Cbwd1) alternative variant cSep08, mRNA.
<a href="#">Cbx1</a>	<a href="#">Cbx1.aSep08</a>	<a href="#">360609</a>	15128	746	4	206	chromobox homolog 1 (HP1 beta homolog Drosophila ) (Cbx1) alternative variant aSep08, mRNA.
<a href="#">Cbx1</a>	<a href="#">Cbx1.bSep08</a>	<a href="#">360609</a>	6870	591	3	95	chromobox homolog 1 (HP1 beta homolog Drosophila ) (Cbx1) alternative variant bSep08, mRNA.
<a href="#">Cbx1</a>	<a href="#">Cbx1.cSep08</a>	<a href="#">360609</a>	4689	846	2	78	chromobox homolog 1 (HP1 beta homolog Drosophila ) (Cbx1) alternative variant cSep08, mRNA.
<a href="#">Cbx1</a>	<a href="#">Cbx1.dSep08</a>	<a href="#">360609</a>	1780	381	2	66	chromobox homolog 1 (HP1 beta homolog Drosophila ) (Cbx1) alternative variant dSep08, mRNA.
<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.aSep08</a>	<a href="#">297093</a>	14250	1792	7	183	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (20.8 kD) (Cbx3.1) alternative variant aSep08, mRNA.

<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.bSep08</a>	<a href="#">297093</a>	11599	1222	5	183	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (20.8 kD) (Cbx3.1) alternative variant bSep08, mRNA.
<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.cSep08</a>	<a href="#">297093</a>	12955	1444	6	183	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (20.8 kD) (Cbx3.1) alternative variant cSep08, mRNA.
<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.dSep08</a>	<a href="#">297093</a>	12566	1050	6	183	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (20.8 kD) (Cbx3.1) alternative variant dSep08, mRNA.
<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.eSep08</a>	<a href="#">297093</a>	12016	1310	6	158	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (Cbx3.1) alternative variant eSep08, mRNA.
<a href="#">Cbx3.1</a>	<a href="#">Cbx3.1.hSep08</a>	<a href="#">297093</a>	7729	395	4	23	chromobox homolog 3 (HP1 gamma homolog, Drosophila) (Cbx3.1) alternative variant hSep08, mRNA.
<a href="#">Cbx4</a>	<a href="#">Cbx4.aSep08</a>	<a href="#">501403</a>	510	363		38	chromobox homolog 4 (Drosophila Pc class) (4.5 kD) (Cbx4) mRNA.
<a href="#">Cbx5</a>	<a href="#">Cbx5.bSep08</a>	<a href="#">300266</a>	35214	462	3	104	chromobox homolog 5 (Drosophila HP1a) (Cbx5) alternative variant bSep08, mRNA.
<a href="#">Cbx5</a>	<a href="#">Cbx5.cSep08</a>	<a href="#">300266</a>	11002	478	3	100	chromobox homolog 5 (Drosophila HP1a) (Cbx5) alternative variant cSep08, mRNA.
<a href="#">Cbx6</a>	<a href="#">Cbx6.bSep08</a>	<a href="#">315136</a>	5540	931	1	278	chromobox homolog 6 (Cbx6) alternative variant bSep08, mRNA.
<a href="#">Cbx6</a>	<a href="#">Cbx6.cSep08</a>	<a href="#">315136</a>	10020	4981	2	225	chromobox homolog 6 (Cbx6) alternative variant cSep08, mRNA.
<a href="#">Cbx7</a>	<a href="#">Cbx7.bSep08</a>	<a href="#">362962</a>	17308	2967	7	158	chromobox homolog 7 (18.0 kD) (Cbx7) alternative variant bSep08, complete mRNA.
<a href="#">Cbx7</a>	<a href="#">Cbx7.cSep08</a>	<a href="#">362962</a>	15416	803	5	151	chromobox homolog 7 (17.4 kD) (Cbx7) alternative variant cSep08, mRNA.
<a href="#">Cbx7</a>	<a href="#">Cbx7.dSep08</a>	<a href="#">362962</a>	12206	339	4	112	chromobox homolog 7 (Cbx7) alternative variant dSep08, mRNA.
<a href="#">Cby1</a>	<a href="#">Cby1.aSep08</a>	<a href="#">246768</a>	3691	437	2	145	chibby homolog 1 (Drosophila) (Cby1) alternative variant aSep08, mRNA.
<a href="#">Cc2d1a</a>	<a href="#">Cc2d1a.bSep08</a>	<a href="#">288908</a>	14991	1931	11	466	putative protein, with a coiled coil domain, of metazoan origin (Cc2d1a) alternative variant bSep08, complete mRNA.
<a href="#">Cc2d1a</a>	<a href="#">Cc2d1a.eSep08</a>	<a href="#">288908</a>	1258	661	4	67	putative protein of vertebrate origin (7.8 kD) (Cc2d1a) alternative variant eSep08, mRNA.
<a href="#">Cc2d1b</a>	<a href="#">Cc2d1b.aSep08</a>	<a href="#">313478</a>	4548	843	10	255	c2 calcium-dependent membrane targeting (Cc2d1b) alternative variant aSep08, mRNA.
<a href="#">Cc2d1b</a>	<a href="#">Cc2d1b.bSep08</a>	<a href="#">313478</a>	3776	1255	7	158	c2 calcium-dependent membrane targeting (18.2 kD) (Cc2d1b) alternative variant bSep08, mRNA.
<a href="#">Ccar1</a>	<a href="#">Ccar1.bSep08</a>	<a href="#">361849</a>	17950	2053	11	587	cell division cycle and apoptosis regulator 1 (Ccar1) alternative variant bSep08, mRNA.
<a href="#">Ccar1</a>	<a href="#">Ccar1.cSep08</a>	<a href="#">361849</a>	9506	1587	4	237	cell division cycle and apoptosis regulator 1 (28.4 kD) (Ccar1) alternative variant cSep08, mRNA.
<a href="#">Ccar1</a>	<a href="#">Ccar1.dSep08</a>	<a href="#">361849</a>	6221	656	5	155	cell division cycle and apoptosis regulator 1 (Ccar1) alternative variant dSep08, mRNA.
<a href="#">Ccar1</a>	<a href="#">Ccar1.hSep08</a>	<a href="#">361849</a>	2227	494	2	54	cell division cycle and apoptosis regulator 1 (Ccar1) alternative variant hSep08, mRNA.
<a href="#">Ccar1</a>	<a href="#">Ccar1.jSep08</a>	<a href="#">361849</a>	6267	258	3	50	cell division cycle and apoptosis regulator 1 (Ccar1) alternative variant jSep08, mRNA.

<a href="#">Ccbe1</a>	<a href="#">Ccbe1.aSep08</a>	<a href="#">361341</a>	18631	410		136	collagen and calcium binding EGF domains 1 (Ccbe1) mRNA.
<a href="#">Ccbl1</a>	<a href="#">Ccbl1.bSep08</a>	<a href="#">311844</a>	64281	1802	7	426	cysteine conjugate-beta lyase 1 (Ccbl1) alternative variant bSep08, mRNA.
<a href="#">Ccbl1</a>	<a href="#">Ccbl1.cSep08</a>	<a href="#">311844</a>	67491	2034	12	241	cysteine conjugate-beta lyase 1 (26.7 kD) (Ccbl1) alternative variant cSep08, mRNA.
<a href="#">Ccbl1</a>	<a href="#">Ccbl1.dSep08</a>	<a href="#">311844</a>	62547	848	8	240	cysteine conjugate-beta lyase 1 (Ccbl1) alternative variant dSep08, mRNA.
<a href="#">Ccbl1</a>	<a href="#">Ccbl1.eSep08</a>	<a href="#">311844</a>	1139	891	2	96	cysteine conjugate-beta lyase 1 (10.3 kD) (Ccbl1) alternative variant eSep08, mRNA.
<a href="#">Ccdc5</a>	<a href="#">Ccdc5.bSep08</a>	<a href="#">192228</a>	4485	290	5	89	putative protein (Ccdc5) alternative variant bSep08, mRNA.
<a href="#">Ccdc6</a>	<a href="#">Ccdc6.aSep08</a>	<a href="#">691155</a>	21689	418		139	putative protein, with 2 coiled coil domains, of metazoan origin (Ccdc6) mRNA.
<a href="#">Ccdc11</a>	<a href="#">Ccdc11.aSep08</a>	<a href="#">364899</a>	52705	658		187	putative protein, with 2 coiled coil domains, of vertebrate origin (Ccdc11) mRNA.
<a href="#">Ccdc12</a>	<a href="#">Ccdc12.bSep08</a>	<a href="#">363151</a>	50639	675		115	mRNA splicing factor, Cwf18 (13.1 kD) (Ccdc12) alternative variant bSep08, mRNA.
<a href="#">Ccdc14</a>	<a href="#">Ccdc14.aSep08</a>	<a href="#">288054</a>	1325	397		131	putative protein, with a coiled coil domain, of metazoan origin (Ccdc14) mRNA.
<a href="#">Ccdc17</a>	<a href="#">Ccdc17.bSep08</a>	<a href="#">500528</a>	2673	650	3	144	putative protein of metazoan origin (Ccdc17) alternative variant bSep08, mRNA.
<a href="#">Ccdc17</a>	<a href="#">Ccdc17.cSep08</a>	<a href="#">500528</a>	1634	1289	4	122	putative protein of vertebrate origin (Ccdc17) alternative variant cSep08, mRNA.
<a href="#">Ccdc17</a>	<a href="#">Ccdc17.dSep08</a>	<a href="#">500528</a>	1219	485	6	87	putative protein, with a coiled coil domain, of mammalian origin (Ccdc17) alternative variant dSep08, mRNA.
<a href="#">Ccdc17</a>	<a href="#">Ccdc17.eSep08</a>	<a href="#">500528</a>	1563	764	3	86	putative protein of mammalian origin (Ccdc17) alternative variant eSep08, mRNA.
<a href="#">Ccdc18</a>	<a href="#">Ccdc18.aSep08</a>	<a href="#">305628</a>	15359	719		124	putative protein, with 2 coiled coil domains (Ccdc18) mRNA.
<a href="#">Ccdc19</a>	<a href="#">Ccdc19.bSep08</a>	<a href="#">304984</a>	7887	506	1	109	putative protein of mammalian origin (Ccdc19) alternative variant bSep08, mRNA.
<a href="#">Ccdc21</a>	<a href="#">Ccdc21.bSep08</a>	<a href="#">362622</a>	16524	1786	8	418	putative protein, with 3 coiled coil domains, of metazoan origin (Ccdc21) alternative variant bSep08, mRNA.
<a href="#">Ccdc21</a>	<a href="#">Ccdc21.cSep08</a>	<a href="#">362622</a>	14555	1082	7	289	putative protein, with 3 coiled coil domains, of metazoan origin (Ccdc21) alternative variant cSep08, mRNA.
<a href="#">Ccdc21</a>	<a href="#">Ccdc21.dSep08</a>	<a href="#">362622</a>	2379	837	2	102	putative nuclear protein, with a coiled coil domain, of metazoan origin (11.2 kD) (Ccdc21) alternative variant dSep08, mRNA.
<a href="#">Ccdc22</a>	<a href="#">Ccdc22.aSep08</a>	<a href="#">317381</a>	7047	1911	3	553	putative protein, with 3 coiled coil domains, of eukaryotic origin (Ccdc22) alternative variant aSep08, mRNA.
<a href="#">Ccdc22</a>	<a href="#">Ccdc22.bSep08</a>	<a href="#">317381</a>	1495	1136	1	146	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc22) alternative variant bSep08, mRNA.
<a href="#">Ccdc25</a>	<a href="#">Ccdc25.aSep08</a>	<a href="#">361059</a>	32963	2402		234	putative protein, with 2 coiled coil domains, of ancient origin (Ccdc25) mRNA.
<a href="#">Ccdc27</a>	<a href="#">Ccdc27.aSep08</a>	<a href="#">679584</a>	1300	264		56	putative protein of mammalian origin (Ccdc27) mRNA.
<a href="#">Ccdc28a</a>	<a href="#">Ccdc28a.aSep08</a>	<a href="#">361454</a>	15208	1793	2	553	putative protein of metazoan origin (Ccdc28a) alternative variant aSep08, mRNA.

<a href="#">Ccgc28a</a>	<a href="#">Ccgc28a.cSep08</a>	<a href="#">361454</a>	8735	544	1	140	putative protein of metazoan origin (Ccgc28a) alternative variant cSep08, mRNA.
<a href="#">Ccgc33</a>	<a href="#">Ccgc33.bSep08</a>	<a href="#">315712</a>	2946	444	1	119	putative protein, with a coiled coil domain, of mammalian origin (Ccgc33) alternative variant bSep08, mRNA.
<a href="#">Ccgc34</a>	<a href="#">Ccgc34.bSep08</a>	<a href="#">362187</a>	5128	753	3	165	putative protein, with a coiled coil domain, of vertebrate origin (Ccgc34) alternative variant bSep08, mRNA.
<a href="#">Ccgc34</a>	<a href="#">Ccgc34.cSep08</a>	<a href="#">362187</a>	30514	632	3	95	putative protein, with a coiled coil domain, of mammalian origin (Ccgc34) alternative variant cSep08, mRNA.
<a href="#">Ccgc37</a>	<a href="#">Ccgc37.aSep08</a>	<a href="#">297444</a>	2581	678		177	putative protein, with a coiled coil domain, of metazoan origin (Ccgc37) mRNA.
<a href="#">Ccgc39</a>	<a href="#">Ccgc39.bSep08</a>	<a href="#">310315</a>	5624	632	3	140	putative protein, with 2 coiled coil domains, of eukaryotic origin (16.3 kD) (Ccgc39) alternative variant bSep08, mRNA.
<a href="#">Ccgc39</a>	<a href="#">Ccgc39.cSep08</a>	<a href="#">310315</a>	4014	362	2	99	putative protein, with 2 coiled coil domains (Ccgc39) alternative variant cSep08, mRNA.
<a href="#">Ccgc40</a>	<a href="#">Ccgc40.cSep08</a>	<a href="#">287867</a>	560	477	2	101	putative protein, with 2 coiled coil domains (Ccgc40) alternative variant cSep08, mRNA.
<a href="#">Ccgc40</a>	<a href="#">Ccgc40.dSep08</a>	<a href="#">287867</a>	3659	627	2	64	putative protein (Ccgc40) alternative variant dSep08, mRNA.
<a href="#">Ccgc40</a>	<a href="#">Ccgc40.eSep08</a>	<a href="#">287867</a>	535	333	2	41	putative protein, with a coiled coil domain (Ccgc40) alternative variant eSep08, mRNA.
<a href="#">Ccgc41</a>	<a href="#">Ccgc41.bSep08</a>	<a href="#">366872</a>	41756	768	4	176	putative nuclear protein, with 2 coiled coil domains, of vertebrate origin (21.0 kD) (Ccgc41) alternative variant bSep08, mRNA.
<a href="#">Ccgc41</a>	<a href="#">Ccgc41.cSep08</a>	<a href="#">366872</a>	25009	674	2	174	putative nuclear protein, with 2 coiled coil domains, of vertebrate origin (20.7 kD) (Ccgc41) alternative variant cSep08, mRNA.
<a href="#">Ccgc41</a>	<a href="#">Ccgc41.dSep08</a>	<a href="#">366872</a>	29272	442	2	147	putative protein, with a coiled coil domain, of vertebrate origin (Ccgc41) alternative variant dSep08, mRNA.
<a href="#">Ccgc41</a>	<a href="#">Ccgc41.eSep08</a>	<a href="#">366872</a>	4234	885	1	132	putative protein, with a coiled coil domain, of vertebrate origin (Ccgc41) alternative variant eSep08, mRNA.
<a href="#">Ccgc41</a>	<a href="#">Ccgc41.fSep08</a>	<a href="#">366872</a>	24974	635	2	114	putative protein, with a coiled coil domain, of vertebrate origin (Ccgc41) alternative variant fSep08, mRNA.
<a href="#">Ccgc43</a>	<a href="#">Ccgc43.aSep08</a>	<a href="#">360637</a>	10690	1085	5	333	putative protein, with 2 coiled coil domains, of bilateral origin (Ccgc43) alternative variant aSep08, mRNA.
<a href="#">Ccgc44</a>	<a href="#">Ccgc44.bSep08</a>	<a href="#">360645</a>	5573	672	3	169	putative mitochondrial protein of ancient origin (18.6 kD) (Ccgc44) alternative variant bSep08, mRNA.
<a href="#">Ccgc44</a>	<a href="#">Ccgc44.cSep08</a>	<a href="#">360645</a>	8078	1505	4	169	putative protein of ancient origin (18.6 kD) (Ccgc44) alternative variant cSep08, mRNA.
<a href="#">Ccgc45</a>	<a href="#">Ccgc45.bSep08</a>	<a href="#">287766</a>	3503	1174	4	222	putative nuclear protein, with a coiled coil domain, of metazoan origin (Ccgc45) alternative variant bSep08, mRNA.
<a href="#">Ccgc45</a>	<a href="#">Ccgc45.cSep08</a>	<a href="#">287766</a>	2099	368	2	102	putative protein of mammalian origin (Ccgc45) alternative variant cSep08, mRNA.
<a href="#">Ccgc45</a>	<a href="#">Ccgc45.eSep08</a>	<a href="#">287766</a>	2964	697	3	52	putative protein (Ccgc45) alternative variant eSep08, mRNA.

<a href="#">Ccdc45</a>	<a href="#">Ccdc45.gSep08</a>	<a href="#">287766</a>	532	307	2	32	putative protein (Ccdc45) alternative variant gSep08, mRNA.
<a href="#">Ccdc46</a>	<a href="#">Ccdc46.bSep08</a>	<a href="#">287776</a>	204050	2258	19	334	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc46) alternative variant bSep08, mRNA.
<a href="#">Ccdc49</a>	<a href="#">Ccdc49.bSep08</a>	<a href="#">360613</a>	8140	676	1	143	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc49) alternative variant bSep08, mRNA.
<a href="#">Ccdc50</a>	<a href="#">Ccdc50.bSep08</a>	<a href="#">288022</a>	18337	426	3	80	putative nuclear protein of bilateral origin (9.2 kD) (Ccdc50) alternative variant bSep08, complete mRNA.
<a href="#">Ccdc50</a>	<a href="#">Ccdc50.cSep08</a>	<a href="#">288022</a>	1524	765	3	70	putative protein of vertebrate origin (Ccdc50) alternative variant cSep08, mRNA.
<a href="#">Ccdc51</a>	<a href="#">Ccdc51.bSep08</a>	<a href="#">316008</a>	15754	572	1	121	putative protein of vertebrate origin (Ccdc51) alternative variant bSep08, mRNA.
<a href="#">Ccdc52</a>	<a href="#">Ccdc52.bSep08</a>	<a href="#">288111</a>	8101	1170	5	257	putative protein, with a coiled coil domain, of vertebrate origin (Ccdc52) alternative variant bSep08, mRNA.
<a href="#">Ccdc53</a>	<a href="#">Ccdc53.bSep08</a>	<a href="#">299707</a>	47281	615	7	186	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc53) alternative variant bSep08, mRNA.
<a href="#">Ccdc53</a>	<a href="#">Ccdc53.cSep08</a>	<a href="#">299707</a>	34461	662	6	105	putative protein of metazoan origin (Ccdc53) alternative variant cSep08, mRNA.
<a href="#">Ccdc53</a>	<a href="#">Ccdc53.dSep08</a>	<a href="#">299707</a>	8162	383	2	60	putative protein (Ccdc53) alternative variant dSep08, mRNA.
<a href="#">Ccdc55</a>	<a href="#">Ccdc55.bSep08</a>	<a href="#">303346</a>	30897	622	6	207	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc55) alternative variant bSep08, mRNA.
<a href="#">Ccdc58</a>	<a href="#">Ccdc58.aSep08</a>	<a href="#">288065</a>	10458	681	3	147	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc58) alternative variant aSep08, mRNA.
<a href="#">Ccdc58</a>	<a href="#">Ccdc58.cSep08</a>	<a href="#">288065</a>	20402	385	1	50	putative protein of metazoan origin (Ccdc58) alternative variant cSep08, mRNA.
<a href="#">Ccdc60</a>	<a href="#">Ccdc60.bSep08</a>	<a href="#">498190</a>	41634	688		228	putative protein of metazoan origin (Ccdc60) alternative variant bSep08, mRNA.
<a href="#">Ccdc62</a>	<a href="#">Ccdc62.bSep08</a>	<a href="#">689909</a>	25556	1400	9	147	putative protein, with 2 coiled coil domains, of metazoan origin (17.5 kD) (Ccdc62) alternative variant bSep08, mRNA.
<a href="#">Ccdc62</a>	<a href="#">Ccdc62.cSep08</a>	<a href="#">689909</a>	9167	483	3	114	putative protein, with a coiled coil domain, of metazoan origin (Ccdc62) alternative variant cSep08, mRNA.
<a href="#">Ccdc63</a>	<a href="#">Ccdc63.aSep08</a>	<a href="#">304484</a>	4653	676	3	205	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc63) alternative variant aSep08, mRNA.
<a href="#">Ccdc63</a>	<a href="#">Ccdc63.bSep08</a>	<a href="#">304484</a>	1480	406	1	79	putative protein of mammalian origin (Ccdc63) alternative variant bSep08, mRNA.
<a href="#">Ccdc64</a>	<a href="#">Ccdc64.aSep08</a>	<a href="#">304537</a>	82036	2512	8	408	putative protein, with 5 coiled coil domains, of metazoan origin (Ccdc64) alternative variant aSep08, mRNA.
<a href="#">Ccdc64</a>	<a href="#">Ccdc64.bSep08</a>	<a href="#">304537</a>	19206	983	6	279	putative protein, with 4 coiled coil domains, of vertebrate origin (Ccdc64) alternative variant bSep08, mRNA.
<a href="#">Ccdc64</a>	<a href="#">Ccdc64.cSep08</a>	<a href="#">304537</a>	7328	1050	4	93	putative protein, with a coiled coil domain, of bilateral origin (10.9 kD) (Ccdc64) alternative variant cSep08, mRNA.
<a href="#">Ccdc65</a>	<a href="#">Ccdc65.bSep08</a>	<a href="#">362994</a>	9168	841	4	204	putative protein, with 3 coiled coil domains, of eukaryotic origin (24.4 kD) (Ccdc65) alternative variant bSep08, complete mRNA.



<a href="#">Ccfdc65</a>	<a href="#">Ccfdc65.cSep08</a>	<a href="#">362994</a>	4892	580	3	186	putative protein, with a coiled coil domain, of eukaryotic origin (Ccfdc65) alternative variant cSep08, mRNA.
<a href="#">Ccfdc65</a>	<a href="#">Ccfdc65.eSep08</a>	<a href="#">362994</a>	2481	927	4	111	putative protein, with a coiled coil domain, of eukaryotic origin (Ccfdc65) alternative variant eSep08, mRNA.
<a href="#">Ccfdc65</a>	<a href="#">Ccfdc65.fSep08</a>	<a href="#">362994</a>	4239	583	2	83	putative protein (Ccfdc65) alternative variant fSep08, mRNA.
<a href="#">Ccfdc65</a>	<a href="#">Ccfdc65.gSep08</a>	<a href="#">362994</a>	499	333	2	80	putative protein (Ccfdc65) alternative variant gSep08, mRNA.
<a href="#">Ccfdc67</a>	<a href="#">Ccfdc67.bSep08</a>	<a href="#">315438</a>	21698	709	5	199	putative protein, with a coiled coil domain, of vertebrate origin (Ccfdc67) alternative variant bSep08, mRNA.
<a href="#">Ccfdc67</a>	<a href="#">Ccfdc67.cSep08</a>	<a href="#">315438</a>	25789	672	6	170	putative protein, with 2 coiled coil domains, of metazoan origin (Ccfdc67) alternative variant cSep08, mRNA.
<a href="#">Ccfdc67</a>	<a href="#">Ccfdc67.dSep08</a>	<a href="#">315438</a>	2399	261	3	69	putative protein of mammalian origin (Ccfdc67) alternative variant dSep08, mRNA.
<a href="#">Ccfdc68</a>	<a href="#">Ccfdc68.bSep08</a>	<a href="#">291530</a>	21431	1140	2	227	putative protein, with 3 coiled coil domains, of vertebrate origin (Ccfdc68) alternative variant bSep08, mRNA.
<a href="#">Ccfdc69</a>	<a href="#">Ccfdc69.bSep08</a>	<a href="#">497906</a>	26013	756	6	184	putative protein, with a coiled coil domain, of vertebrate origin (Ccfdc69) alternative variant bSep08, mRNA.
<a href="#">Ccfdc71</a>	<a href="#">Ccfdc71.cSep08</a>	<a href="#">498678</a>	610	300	2	51	putative protein (Ccfdc71) alternative variant cSep08, mRNA.
<a href="#">Ccfdc77</a>	<a href="#">Ccfdc77.aSep08</a>	<a href="#">312677</a>	15902	605	4	201	putative protein, with a coiled coil domain, of eukaryotic origin (Ccfdc77) alternative variant aSep08, mRNA.
<a href="#">Ccfdc77</a>	<a href="#">Ccfdc77.bSep08</a>	<a href="#">312677</a>	7666	388	3	129	putative protein, with a coiled coil domain, of metazoan origin (Ccfdc77) alternative variant bSep08, mRNA.
<a href="#">Ccfdc79</a>	<a href="#">Ccfdc79.aSep08</a>	<a href="#">307615</a>	5171	389		129	putative protein of vertebrate origin (Ccfdc79) mRNA.
<a href="#">Ccfdc81</a>	<a href="#">Ccfdc81.bSep08</a>	<a href="#">308810</a>	2176	630	1	85	putative protein of metazoan origin (Ccfdc81) alternative variant bSep08, mRNA.
<a href="#">Ccfdc82</a>	<a href="#">Ccfdc82.bSep08</a>	<a href="#">300359</a>	4230	467	1	86	putative protein of mammalian origin (Ccfdc82) alternative variant bSep08, mRNA.
<a href="#">Ccfdc84</a>	<a href="#">Ccfdc84.aSep08</a>	<a href="#">689046</a>	6572	1048	8	179	putative secreted or extracellular protein precursor of eukaryotic origin (19.6 kD) (Ccfdc84) alternative variant aSep08, mRNA.
<a href="#">Ccfdc84</a>	<a href="#">Ccfdc84.bSep08</a>	<a href="#">689046</a>	5942	1276	6	162	putative secreted or extracellular protein precursor of eukaryotic origin (17.7 kD) (Ccfdc84) alternative variant bSep08, mRNA.
<a href="#">Ccfdc84</a>	<a href="#">Ccfdc84.cSep08</a>	<a href="#">689046</a>	1628	584	4	110	putative nuclear protein of metazoan origin (12.6 kD) (Ccfdc84) alternative variant cSep08, mRNA.
<a href="#">Ccfdc84</a>	<a href="#">Ccfdc84.dSep08</a>	<a href="#">689046</a>	2108	1932	3	88	putative nuclear protein of metazoan origin (10.1 kD) (Ccfdc84) alternative variant dSep08, mRNA.
<a href="#">Ccfdc84</a>	<a href="#">Ccfdc84.fSep08</a>	<a href="#">689046</a>	775	632	3	48	putative protein (5.0 kD) (Ccfdc84) alternative variant fSep08, mRNA.
<a href="#">Ccfdc85a</a>	<a href="#">Ccfdc85a.aSep08</a>	<a href="#">289855</a>	229619	757		233	putative protein of vertebrate origin (Ccfdc85a) mRNA.
<a href="#">Ccfdc88a</a>	<a href="#">Ccfdc88a.aSep08</a>	<a href="#">305605</a>	17298	1299	9	433	girdin (Ccfdc88a) alternative variant aSep08, mRNA.
<a href="#">Ccfdc88a</a>	<a href="#">Ccfdc88a.bSep08</a>	<a href="#">305605</a>	3283	1212	3	404	putative protein of vertebrate origin (Ccfdc88a) alternative variant bSep08, mRNA.
<a href="#">Ccfdc88a</a>	<a href="#">Ccfdc88a.cSep08</a>	<a href="#">305605</a>	4734	714	3	138	girdin (Ccfdc88a) alternative variant cSep08, mRNA.
<a href="#">Ccfdc88a</a>	<a href="#">Ccfdc88a.dSep08</a>	<a href="#">305605</a>	4081	659	2	86	girdin (Ccfdc88a) alternative variant dSep08, mRNA.

<a href="#">Ccdc88b</a>	<a href="#">Ccdc88b.aSep08</a>	<a href="#">361715</a>	10730	3087	3	784	protein kinase and protein kinase, C-terminal and tyrosine protein kinase (Ccdc88b) alternative variant aSep08, mRNA.
<a href="#">Ccdc88b</a>	<a href="#">Ccdc88b.bSep08</a>	<a href="#">361715</a>	2305	539	2	179	putative protein of eukaryotic origin (Ccdc88b) alternative variant bSep08, mRNA.
<a href="#">Ccdc88b</a>	<a href="#">Ccdc88b.cSep08</a>	<a href="#">361715</a>	3172	763	3	171	protein kinase and tyrosine protein kinase (Ccdc88b) alternative variant cSep08, mRNA.
<a href="#">Ccdc88c</a>	<a href="#">Ccdc88c.aSep08</a>	<a href="#">362770</a>	62911	561		187	CRA a (Ccdc88c) mRNA.
<a href="#">Ccdc90a</a>	<a href="#">Ccdc90a.aSep08</a>	<a href="#">291034</a>	16866	773	8	249	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc90a) alternative variant aSep08, mRNA.
<a href="#">Ccdc90a</a>	<a href="#">Ccdc90a.bSep08</a>	<a href="#">291034</a>	1754	627	1	32	putative protein of vertebrate origin (3.7 kD) (Ccdc90a) alternative variant bSep08, mRNA.
<a href="#">Ccdc90b</a>	<a href="#">Ccdc90b.bSep08</a>	<a href="#">308820</a>	7491	1535	4	149	putative mitochondrial protein, with a coiled coil domain, of metazoan origin (17.3 kD) (Ccdc90b) alternative variant bSep08, mRNA.
<a href="#">Ccdc91</a>	<a href="#">Ccdc91.bSep08</a>	<a href="#">312863</a>	70999	1698		250	protein CRA c (Ccdc91) alternative variant bSep08, mRNA.
<a href="#">Ccdc93</a>	<a href="#">Ccdc93.bSep08</a>	<a href="#">304743</a>	1458	282	1	54	putative protein, with a coiled coil domain, of metazoan origin (Ccdc93) alternative variant bSep08, mRNA.
<a href="#">Ccdc95</a>	<a href="#">Ccdc95.bSep08</a>	<a href="#">293494</a>	9419	869	7	242	putative nuclear protein, with a coiled coil domain, of metazoan origin (26.2 kD) (Ccdc95) alternative variant bSep08, mRNA.
<a href="#">Ccdc95</a>	<a href="#">Ccdc95.cSep08</a>	<a href="#">293494</a>	9106	603	5	155	putative protein of bilateral origin (Ccdc95) alternative variant cSep08, mRNA.
<a href="#">Ccdc95</a>	<a href="#">Ccdc95.dSep08</a>	<a href="#">293494</a>	10565	571	5	124	putative protein (Ccdc95) alternative variant dSep08, mRNA.
<a href="#">Ccdc95</a>	<a href="#">Ccdc95.eSep08</a>	<a href="#">293494</a>	1099	600	2	83	putative protein of mammalian origin (Ccdc95) alternative variant eSep08, mRNA.
<a href="#">Ccdc97</a>	<a href="#">Ccdc97.bSep08</a>	<a href="#">292724</a>	3323	677	1	193	putative protein of metazoan origin (Ccdc97) alternative variant bSep08, mRNA.
<a href="#">Ccdc99</a>	<a href="#">Ccdc99.bSep08</a>	<a href="#">303037</a>	3708	1219	2	145	putative nuclear protein of mammalian origin (16.0 kD) (Ccdc99) alternative variant bSep08, mRNA.
<a href="#">Ccdc100</a>	<a href="#">Ccdc100.aSep08</a>	<a href="#">307302</a>	41787	3194	12	563	protein CRA a (Ccdc100) alternative variant aSep08, mRNA.
<a href="#">Ccdc101</a>	<a href="#">Ccdc101.bSep08</a>	<a href="#">293488</a>	7833	824	7	242	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc101) alternative variant bSep08, mRNA.
<a href="#">Ccdc101</a>	<a href="#">Ccdc101.cSep08</a>	<a href="#">293488</a>	660	376	4	93	putative protein of metazoan origin (Ccdc101) alternative variant cSep08, mRNA.
<a href="#">Ccdc101</a>	<a href="#">Ccdc101.dSep08</a>	<a href="#">293488</a>	30627	2404	5	82	putative protein, with a coiled coil domain, of metazoan origin (9.3 kD) (Ccdc101) alternative variant dSep08, mRNA.
<a href="#">Ccdc102a</a>	<a href="#">Ccdc102a.bSep08</a>	<a href="#">361363</a>	6400	1379		392	putative protein, with at least 3 transmembrane domains, 2 coiled coil domains, of metazoan origin (Ccdc102a) alternative variant bSep08, mRNA.
<a href="#">Ccdc102a</a>	<a href="#">Ccdc102a.cSep08</a>	<a href="#">361363</a>	8112	858		154	putative protein, with a coiled coil domain, of metazoan origin (Ccdc102a) alternative variant cSep08, mRNA.
<a href="#">Ccdc102a</a>	<a href="#">Ccdc102a.dSep08</a>	<a href="#">361363</a>	4247	680		94	putative protein of vertebrate origin (Ccdc102a) alternative variant dSep08, mRNA.

<a href="#">Ccdc104</a>	<a href="#">Ccdc104.bSep08</a>	<a href="#">289859</a>	3349	716	3	102	putative cytoplasmic protein, with a coiled coil domain, of eukaryotic origin (11.8 kD) (Ccdc104) alternative variant bSep08, complete mRNA.
<a href="#">Ccdc104</a>	<a href="#">Ccdc104.cSep08</a>	<a href="#">289859</a>	25476	1792	6	79	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc104) alternative variant cSep08, mRNA.
<a href="#">Ccdc104</a>	<a href="#">Ccdc104.dSep08</a>	<a href="#">289859</a>	1953	502	2	54	putative protein (Ccdc104) alternative variant dSep08, mRNA.
<a href="#">Ccdc105</a>	<a href="#">Ccdc105.bSep08</a>	<a href="#">500800</a>	2271	881	1	255	putative protein, with a coiled coil domain, of metazoan origin (Ccdc105) alternative variant bSep08, mRNA.
<a href="#">Ccdc107</a>	<a href="#">Ccdc107.aSep08</a>	<a href="#">313496</a>	3384	1626	6	266	putative protein, with a coiled coil domain, of mammalian origin (Ccdc107) alternative variant aSep08, mRNA.
<a href="#">Ccdc107</a>	<a href="#">Ccdc107.bSep08</a>	<a href="#">313496</a>	3418	1153	6	261	putative protein, with a coiled coil domain, of mammalian origin (Ccdc107) alternative variant bSep08, mRNA.
<a href="#">Ccdc107</a>	<a href="#">Ccdc107.cSep08</a>	<a href="#">313496</a>	2653	898	6	253	putative protein, with a coiled coil domain, of mammalian origin (Ccdc107) alternative variant cSep08, mRNA.
<a href="#">Ccdc107</a>	<a href="#">Ccdc107.eSep08</a>	<a href="#">313496</a>	2294	1590	3	107	protein CRA e (Ccdc107) alternative variant eSep08, mRNA.
<a href="#">Ccdc107</a>	<a href="#">Ccdc107.fSep08</a>	<a href="#">313496</a>	585	396	3	46	putative protein (5.3 kD) (Ccdc107) alternative variant fSep08, mRNA.
<a href="#">Ccdc109b</a>	<a href="#">Ccdc109b.aSep08</a>	<a href="#">295462</a>	3862	678		181	putative protein of eukaryotic origin (Ccdc109b) mRNA.
<a href="#">Ccdc110</a>	<a href="#">Ccdc110.aSep08</a>	<a href="#">290755</a>	5814	2427	5	800	putative protein, with 4 coiled coil domains, of vertebrate origin (Ccdc110) alternative variant aSep08, mRNA.
<a href="#">Ccdc110</a>	<a href="#">Ccdc110.bSep08</a>	<a href="#">290755</a>	12152	2636	4	776	putative protein, with 4 coiled coil domains, of vertebrate origin (Ccdc110) alternative variant bSep08, mRNA.
<a href="#">Ccdc111</a>	<a href="#">Ccdc111.aSep08</a>	<a href="#">361147</a>	16256	1018		264	DNA primase, small subunit (Ccdc111) mRNA.
<a href="#">Ccdc112</a>	<a href="#">Ccdc112.bSep08</a>	<a href="#">498858</a>	26711	642		213	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc112) alternative variant bSep08, mRNA.
<a href="#">Ccdc113</a>	<a href="#">Ccdc113.aSep08</a>	<a href="#">291847</a>	19953	1783	6	576	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc113) alternative variant aSep08, mRNA.
<a href="#">Ccdc113</a>	<a href="#">Ccdc113.bSep08</a>	<a href="#">291847</a>	8805	682	5	209	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc113) alternative variant bSep08, mRNA.
<a href="#">Ccdc113</a>	<a href="#">Ccdc113.cSep08</a>	<a href="#">291847</a>	7029	790	4	131	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc113) alternative variant cSep08, mRNA.
<a href="#">Ccdc113</a>	<a href="#">Ccdc113.dSep08</a>	<a href="#">291847</a>	1655	299	2	80	putative protein, with a coiled coil domain, of metazoan origin (Ccdc113) alternative variant dSep08, mRNA.
<a href="#">Ccdc113</a>	<a href="#">Ccdc113.eSep08</a>	<a href="#">291847</a>	1605	608	2	46	putative protein of mammalian origin (Ccdc113) alternative variant eSep08, mRNA.
<a href="#">Ccdc115</a>	<a href="#">Ccdc115.bSep08</a>	<a href="#">363213</a>	3770	986	4	165	putative protein of metazoan origin (17.9 kD) (Ccdc115) alternative variant bSep08, mRNA.
<a href="#">Ccdc115</a>	<a href="#">Ccdc115.cSep08</a>	<a href="#">363213</a>	2011	848	4	147	putative protein, with a coiled coil domain, of eukaryotic origin (Ccdc115) alternative variant cSep08, mRNA.
<a href="#">Ccdc115</a>	<a href="#">Ccdc115.dSep08</a>	<a href="#">363213</a>	2314	613	2	119	putative protein of metazoan origin (13.4 kD) (Ccdc115) alternative variant dSep08, mRNA.
<a href="#">Ccdc117</a>	<a href="#">Ccdc117.aSep08</a>	<a href="#">498404</a>	10430	1797	3	489	putative protein of mammalian origin (Ccdc117) alternative variant aSep08, mRNA.
<a href="#">Ccdc117</a>	<a href="#">Ccdc117.cSep08</a>	<a href="#">498404</a>	4268	445	1	86	putative cytoplasmic protein (9.8 kD) (Ccdc117) alternative variant cSep08, mRNA.

<a href="#">Ccdc120</a>	<a href="#">Ccdc120.aSep08</a>	<a href="#">317377</a>	5443	3033	3	505	putative protein of vertebrate origin (Ccdc120) alternative variant aSep08, mRNA.
<a href="#">Ccdc120</a>	<a href="#">Ccdc120.bSep08</a>	<a href="#">317377</a>	1978	621	1	140	putative protein of vertebrate origin (Ccdc120) alternative variant bSep08, mRNA.
<a href="#">Ccdc124</a>	<a href="#">Ccdc124.bSep08</a>	<a href="#">290642</a>	4766	672	2	194	putative protein, with 2 coiled coil domains, of eukaryotic origin (Ccdc124) alternative variant bSep08, mRNA.
<a href="#">Ccdc125</a>	<a href="#">Ccdc125.bSep08</a>	<a href="#">499518</a>	12865	1978	3	326	putative nuclear protein, with 3 coiled coil domains, of metazoan origin (36.8 kD) (Ccdc125) alternative variant bSep08, mRNA.
<a href="#">Ccdc125</a>	<a href="#">Ccdc125.cSep08</a>	<a href="#">499518</a>	6042	409	1	83	putative protein, with a coiled coil domain (Ccdc125) alternative variant cSep08, mRNA.
<a href="#">Ccdc129</a>	<a href="#">Ccdc129.aSep08</a>	<a href="#">500139</a>	10318	966		191	putative protein, with a coiled coil domain, of mammalian origin (21.7 kD) (Ccdc129) mRNA.
<a href="#">Ccdc130</a>	<a href="#">Ccdc130.bSep08</a>	<a href="#">304656</a>	9657	726	8	149	putative protein of eukaryotic origin (Ccdc130) alternative variant bSep08, mRNA.
<a href="#">Ccdc130</a>	<a href="#">Ccdc130.cSep08</a>	<a href="#">304656</a>	4675	763	5	140	putative protein of eukaryotic origin (Ccdc130) alternative variant cSep08, mRNA.
<a href="#">Ccdc130</a>	<a href="#">Ccdc130.dSep08</a>	<a href="#">304656</a>	860	771	2	83	putative nuclear protein of metazoan origin (9.3 kD) (Ccdc130) alternative variant dSep08, mRNA.
<a href="#">Ccdc132</a>	<a href="#">Ccdc132.aSep08</a>	<a href="#">312083</a>	48677	2364	13	543	putative protein of eukaryotic origin (Ccdc132) alternative variant aSep08, mRNA.
<a href="#">Ccdc132</a>	<a href="#">Ccdc132.bSep08</a>	<a href="#">312083</a>	1144	518	2	50	putative protein of metazoan origin (Ccdc132) alternative variant bSep08, mRNA.
<a href="#">Ccdc132</a>	<a href="#">Ccdc132.cSep08</a>	<a href="#">312083</a>	1882	704	2	57	putative protein (Ccdc132) alternative variant cSep08, mRNA.
<a href="#">Ccdc136</a>	<a href="#">Ccdc136.aSep08</a>	<a href="#">362331</a>	29739	1637	9	470	protein CRA b (Ccdc136) alternative variant aSep08, mRNA.
<a href="#">Ccdc136</a>	<a href="#">Ccdc136.bSep08</a>	<a href="#">362331</a>	12317	1482	5	418	CRA b (Ccdc136) alternative variant bSep08, mRNA.
<a href="#">Ccdc136</a>	<a href="#">Ccdc136.cSep08</a>	<a href="#">362331</a>	17825	852	5	172	protein CRA b (Ccdc136) alternative variant cSep08, mRNA.
<a href="#">Ccdc136</a>	<a href="#">Ccdc136.dSep08</a>	<a href="#">362331</a>	7517	948	2	123	CRA a like (14.2 kD) (Ccdc136) alternative variant dSep08, mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.bSep08</a>	<a href="#">297380</a>	2329	1837	5	119	mitochondrial ribosomal protein L53 (12.7 kD) (Ccdc142andMrpl53) alternative variant bSep08, mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.bSep08</a>	<a href="#">362388</a>	2329	1837	5	119	mitochondrial ribosomal protein L53 (12.7 kD) (Ccdc142andMrpl53) alternative variant bSep08, mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.cSep08</a>	<a href="#">297380</a>	612	347	4	115	putative protein of mammalian origin (Ccdc142andMrpl53) alternative variant cSep08, mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.cSep08</a>	<a href="#">362388</a>	612	347	4	115	putative protein of mammalian origin (Ccdc142andMrpl53) alternative variant cSep08, mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.eSep08</a>	<a href="#">297380</a>	886	656	3	51	mitochondrial ribosomal protein L53 (5.7 kD) (Ccdc142andMrpl53) alternative variant eSep08, complete mRNA.
<a href="#">Ccdc142andMrpl53</a>	<a href="#">Ccdc142andMrpl53.eSep08</a>	<a href="#">362388</a>	886	656	3	51	mitochondrial ribosomal protein L53 (5.7 kD) (Ccdc142andMrpl53) alternative variant eSep08, complete mRNA.

<a href="#">Ccdc147</a>	<a href="#">Ccdc147.aSep08</a>	<a href="#">309467</a>	63497	760		252	putative protein, with 2 coiled coil domains, of eukaryotic origin (Ccdc147) mRNA.
<a href="#">Ccdc148</a>	<a href="#">Ccdc148.aSep08</a>	<a href="#">311051</a>	101170	1776	7	485	putative protein, with a coiled coil domain, of vertebrate origin (Ccdc148) alternative variant aSep08, mRNA.
<a href="#">Ccdc148</a>	<a href="#">Ccdc148.cSep08</a>	<a href="#">311051</a>	101281	1492	10	263	putative nuclear protein, with 2 coiled coil domains, of metazoan origin (30.7 kD) (Ccdc148) alternative variant cSep08, complete mRNA.
<a href="#">Ccdc150</a>	<a href="#">Ccdc150.aSep08</a>	<a href="#">316399</a>	1898	244		81	putative protein, with a coiled coil domain, of mammalian origin (Ccdc150) mRNA.
<a href="#">Ccdc151</a>	<a href="#">Ccdc151.aSep08</a>	<a href="#">315465</a>	3911	798		265	putative protein, with 2 coiled coil domains, of eukaryotic origin (Ccdc151) mRNA.
<a href="#">Ccl1</a>	<a href="#">Ccl1.aSep08</a>	<a href="#">688605</a>	1586	293		39	chemokine (C-C motif) ligand 1 (Ccl1) mRNA.
<a href="#">Ccl9</a>	<a href="#">Ccl9.bSep08</a>	<a href="#">360579</a>	4569	917		98	chemokine (C-C motif) ligand 9 (11.0 kD) (Ccl9) alternative variant bSep08, mRNA.
<a href="#">Ccl25</a>	<a href="#">Ccl25.bSep08</a>	<a href="#">360750</a>	4292	413	1	137	chemokine (C-C motif) ligand 25 (Ccl25) alternative variant bSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.bSep08</a>	<a href="#">362505</a>	1710	1210	3	147	putative protein (Ccl27) alternative variant bSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.cSep08</a>	<a href="#">362505</a>	4474	454	6	131	chemokine ligand 27 CRA b (Ccl27) alternative variant cSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.dSep08</a>	<a href="#">362505</a>	1855	575	3	127	chemokine ligand 27 (14.6 kD) (Ccl27) alternative variant dSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.eSep08</a>	<a href="#">362505</a>	725	412	3	126	chemokine ligand 27 (Ccl27) alternative variant eSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.fSep08</a>	<a href="#">362505</a>	4225	425	5	112	chemokine ligand 27 (Ccl27) alternative variant fSep08, mRNA.
<a href="#">Ccl27</a>	<a href="#">Ccl27.gSep08</a>	<a href="#">362505</a>	930	316	4	87	chemokine ligand 27 CRA g (10.1 kD) (Ccl27) alternative variant gSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.bSep08</a>	<a href="#">295052</a>	9482	1317	8	377	cyclin A1 (42.6 kD) (Ccna1) alternative variant bSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.cSep08</a>	<a href="#">295052</a>	5334	750	5	194	cyclin A1 (Ccna1) alternative variant cSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.dSep08</a>	<a href="#">295052</a>	5161	777	4	175	cyclin A1 (Ccna1) alternative variant dSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.eSep08</a>	<a href="#">295052</a>	4119	613	4	164	cyclin A1 (Ccna1) alternative variant eSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.fSep08</a>	<a href="#">295052</a>	6236	761	4	155	cyclin A1 (Ccna1) alternative variant fSep08, mRNA.
<a href="#">Ccna1</a>	<a href="#">Ccna1.gSep08</a>	<a href="#">295052</a>	4174	725	2	135	cyclin A1 (Ccna1) alternative variant gSep08, mRNA.
<a href="#">Ccna2</a>	<a href="#">Ccna2.bSep08</a>	<a href="#">114494</a>	4439	1090	6	260	cyclin A2 (Ccna2) alternative variant bSep08, mRNA.
<a href="#">Ccna2</a>	<a href="#">Ccna2.cSep08</a>	<a href="#">114494</a>	1101	452	2	127	cyclin A2 (Ccna2) alternative variant cSep08, mRNA.
<a href="#">Ccnb1</a>	<a href="#">Ccnb1.bSep08</a>	<a href="#">25203</a>	2062	1204	2	302	cyclin B1 (Ccnb1) alternative variant bSep08, mRNA.
<a href="#">Ccnb2</a>	<a href="#">Ccnb2.aSep08</a>	<a href="#">363088</a>	14100	1514	1	430	cyclin B2 (Ccnb2) alternative variant aSep08, mRNA.
<a href="#">Ccnb2</a>	<a href="#">Ccnb2.bSep08</a>	<a href="#">363088</a>	9946	734	1	225	cyclin B2 (Ccnb2) alternative variant bSep08, mRNA.
<a href="#">Ccnc</a>	<a href="#">Ccnc.aSep08</a>	<a href="#">114839</a>	15038	940	12	313	cyclin C (Ccnc) alternative variant aSep08, mRNA.
<a href="#">Ccnc</a>	<a href="#">Ccnc.bSep08</a>	<a href="#">114839</a>	8221	601	9	167	cyclin C (Ccnc) alternative variant bSep08, mRNA.
<a href="#">Ccnc</a>	<a href="#">Ccnc.cSep08</a>	<a href="#">114839</a>	15549	1504	13	151	cyclin C (17.7 kD) (Ccnc) alternative variant cSep08, mRNA.
<a href="#">Ccnd2</a>	<a href="#">Ccnd2.aSep08</a>	<a href="#">64033</a>	27494	6223	5	384	cyclin D2 (Ccnd2) alternative variant aSep08, mRNA.
<a href="#">Ccnd2</a>	<a href="#">Ccnd2.bSep08</a>	<a href="#">64033</a>	2377	711	2	153	cyclin D2 (Ccnd2) alternative variant bSep08, mRNA.

<a href="#">Ccnd2</a>	<a href="#">Ccnd2.cSep08</a>	<a href="#">64033</a>	3119	498	2	79	cyclin D2 (Ccnd2) alternative variant cSep08, mRNA.
<a href="#">Ccndbp1</a>	<a href="#">Ccndbp1.bSep08</a>	<a href="#">362201</a>	9618	1318	7	235	cyclin D-type binding-protein 1 (25.8 kD) (Ccndbp1) alternative variant bSep08, mRNA.
<a href="#">Ccndbp1</a>	<a href="#">Ccndbp1.cSep08</a>	<a href="#">362201</a>	848	697	2	104	cyclin D-type binding-protein 1 (Ccndbp1) alternative variant cSep08, mRNA.
<a href="#">Ccne1</a>	<a href="#">Ccne1.bSep08</a>	<a href="#">25729</a>	9153	1958	11	411	cyclin E1 (47.4 kD) (Ccne1) alternative variant bSep08, mRNA.
<a href="#">Ccne1</a>	<a href="#">Ccne1.cSep08</a>	<a href="#">25729</a>	6182	639	5	210	cyclin E1 (Ccne1) alternative variant cSep08, mRNA.
<a href="#">Ccne1</a>	<a href="#">Ccne1.dSep08</a>	<a href="#">25729</a>	8573	462	2	37	cyclin E1 (Ccne1) alternative variant dSep08, mRNA.
<a href="#">Ccne2</a>	<a href="#">Ccne2.bSep08</a>	<a href="#">362485</a>	867	770	2	38	cyclin E2 (Ccne2) alternative variant bSep08, mRNA.
<a href="#">Ccng1</a>	<a href="#">Ccng1.aSep08</a>	<a href="#">25405</a>	6400	3223	5	294	cyclin G1 (33.9 kD) (Ccng1) alternative variant aSep08, mRNA.
<a href="#">Ccng1</a>	<a href="#">Ccng1.bSep08</a>	<a href="#">25405</a>	2817	734	3	235	cyclin G1 (Ccng1) alternative variant bSep08, mRNA.
<a href="#">Ccng1</a>	<a href="#">Ccng1.cSep08</a>	<a href="#">25405</a>	2740	847	3	153	cyclin G1 (Ccng1) alternative variant cSep08, mRNA.
<a href="#">Ccnh</a>	<a href="#">Ccnh.bSep08</a>	<a href="#">84389</a>	12539	748	5	218	cyclin H (Ccnh) alternative variant bSep08, mRNA.
<a href="#">Ccni</a>	<a href="#">Ccni.bSep08</a>	<a href="#">289500</a>	15500	905	4	106	cyclin I (Ccni) alternative variant bSep08, mRNA.
<a href="#">Ccni</a>	<a href="#">Ccni.cSep08</a>	<a href="#">289500</a>	1457	1374	2	62	cyclin I (Ccni) alternative variant cSep08, mRNA.
<a href="#">Ccnk</a>	<a href="#">Ccnk.bSep08</a>	<a href="#">500715</a>	7023	1065	4	254	cyclin K (Ccnk) alternative variant bSep08, mRNA.
<a href="#">Ccnl1</a>	<a href="#">Ccnl1.bSep08</a>	<a href="#">114121</a>	12548	2442	12	371	cyclin L1 (43.3 kD) (Ccnl1) alternative variant bSep08, complete mRNA.
<a href="#">Ccnl1</a>	<a href="#">Ccnl1.cSep08</a>	<a href="#">114121</a>	2968	1620	6	305	cyclin L1 (35.6 kD) (Ccnl1) alternative variant cSep08, mRNA.
<a href="#">Ccnl1</a>	<a href="#">Ccnl1.dSep08</a>	<a href="#">114121</a>	10414	1318	10	242	cyclin L1 (Ccnl1) alternative variant dSep08, mRNA.
<a href="#">Ccnl1</a>	<a href="#">Ccnl1.gSep08</a>	<a href="#">114121</a>	725	469	2	63	cyclin L1 (7.1 kD) (Ccnl1) alternative variant gSep08, mRNA.
<a href="#">Ccnl1</a>	<a href="#">Ccnl1.hSep08</a>	<a href="#">114121</a>	3074	359	5	22	cyclin L1 (Ccnl1) alternative variant hSep08, mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.aSep08</a>	<a href="#">298686</a>	11251	2213	11	520	cyclin L2 CRA c (58.2 kD) (Ccnl2) alternative variant aSep08, complete mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.bSep08</a>	<a href="#">298686</a>	3744	1885	6	349	cyclin L2 (39.8 kD) (Ccnl2) alternative variant bSep08, mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.cSep08</a>	<a href="#">298686</a>	3823	2046	5	327	cyclin L2 CRA d (36.7 kD) (Ccnl2) alternative variant cSep08, mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.dSep08</a>	<a href="#">298686</a>	7955	793	7	233	cyclin L2 (Ccnl2) alternative variant dSep08, mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.fSep08</a>	<a href="#">298686</a>	3686	1283	7	209	cyclin L2 CRA c (23.3 kD) (Ccnl2) alternative variant fSep08, mRNA.
<a href="#">Ccnl2</a>	<a href="#">Ccnl2.gSep08</a>	<a href="#">298686</a>	2472	853	5	198	cyclin L2 CRA c (Ccnl2) alternative variant gSep08, mRNA.
<a href="#">Ccno</a>	<a href="#">Ccno.aSep08</a>	<a href="#">499528</a>	3284	2069	2	352	cyclin O (38.9 kD) (Ccno) alternative variant aSep08, mRNA.
<a href="#">Ccnt2</a>	<a href="#">Ccnt2.bSep08</a>	<a href="#">304758</a>	1253	1155	2	344	cyclin T2 (Ccnt2) alternative variant bSep08, mRNA.
<a href="#">Ccny</a>	<a href="#">Ccny.aSep08</a>	<a href="#">361261</a>	102578	984	3	328	cyclin Y (Ccny) alternative variant aSep08, mRNA.
<a href="#">Ccny</a>	<a href="#">Ccny.bSep08</a>	<a href="#">361261</a>	94356	424	2	93	cyclin Y (Ccny) alternative variant bSep08, mRNA.
<a href="#">Ccnyl1</a>	<a href="#">Ccnyl1.aSep08</a>	<a href="#">316452</a>	24020	2819	7	217	cyclin Y-like 1 (25.5 kD) (Ccnyl1) alternative variant aSep08, mRNA.
<a href="#">Ccnyl1</a>	<a href="#">Ccnyl1.bSep08</a>	<a href="#">316452</a>	2321	377	2	36	cyclin Y-like 1 (Ccnyl1) alternative variant bSep08, mRNA.

<a href="#">Ccp1</a>	<a href="#">Ccp1.aSep08</a>	<a href="#">363098</a>	17562	2782	8	760	cell cycle progression 1 (Ccp1) alternative variant aSep08, mRNA.
<a href="#">Ccp1</a>	<a href="#">Ccp1.cSep08</a>	<a href="#">363098</a>	3081	490	3	137	cell cycle progression 1 (Ccp1) alternative variant cSep08, mRNA.
<a href="#">Ccrk</a>	<a href="#">Ccrk.bSep08</a>	<a href="#">364666</a>	4769	962	1	187	cell cycle related kinase (20.8 kD) (Ccrk) alternative variant bSep08, mRNA.
<a href="#">Cct2</a>	<a href="#">Cct2.bSep08</a>	<a href="#">299809</a>	2589	693	6	174	chaperonin containing TCP1, subunit 2 (beta) (Cct2) alternative variant bSep08, mRNA.
<a href="#">Cct2</a>	<a href="#">Cct2.cSep08</a>	<a href="#">299809</a>	1493	751	2	103	chaperonin containing TCP1, subunit 2 (beta) (Cct2) alternative variant cSep08, mRNA.
<a href="#">Cct2</a>	<a href="#">Cct2.dSep08</a>	<a href="#">299809</a>	2126	649	5	76	chaperonin containing TCP1, subunit 2 (beta) (Cct2) alternative variant dSep08, mRNA.
<a href="#">Cct3</a>	<a href="#">Cct3.bSep08</a>	<a href="#">295230</a>	43430	913	7	304	chaperonin subunit 3 (gamma) (Cct3) alternative variant bSep08, mRNA.
<a href="#">Cct3</a>	<a href="#">Cct3.eSep08</a>	<a href="#">295230</a>	4851	651	4	49	chaperonin subunit 3 (gamma) (Cct3) alternative variant eSep08, mRNA.
<a href="#">Cct4</a>	<a href="#">Cct4.cSep08</a>	<a href="#">29374</a>	1231	751	3	60	chaperonin subunit 4 (delta) (Cct4) alternative variant cSep08, mRNA.
<a href="#">Cct5</a>	<a href="#">Cct5.bSep08</a>	<a href="#">294864</a>	11791	2969	10	248	chaperonin containing Tcp1 (27.0 kD) (Cct5) alternative variant bSep08, complete mRNA.
<a href="#">Cct5</a>	<a href="#">Cct5.cSep08</a>	<a href="#">294864</a>	4286	529	3	165	chaperonin CRA e (17.9 kD) (Cct5) alternative variant cSep08, mRNA.
<a href="#">Cct5</a>	<a href="#">Cct5.dSep08</a>	<a href="#">294864</a>	2223	955	3	163	chaperonin containing Tcp1 (Cct5) alternative variant dSep08, mRNA.
<a href="#">Cct5</a>	<a href="#">Cct5.eSep08</a>	<a href="#">294864</a>	6540	1246	5	153	chaperonin CRA c (16.4 kD) (Cct5) alternative variant eSep08, complete mRNA.
<a href="#">Cct6a</a>	<a href="#">Cct6a.bSep08</a>	<a href="#">288620</a>	4692	691	7	105	chaperonin subunit 6a (zeta) (Cct6a) alternative variant bSep08, mRNA.
<a href="#">Cct7</a>	<a href="#">Cct7.bSep08</a>	<a href="#">297406</a>	15045	1330	10	380	chaperonin subunit 7 (eta) (41.7 kD) (Cct7) alternative variant bSep08, complete mRNA.
<a href="#">Cct7</a>	<a href="#">Cct7.cSep08</a>	<a href="#">297406</a>	10106	704	4	143	chaperonin subunit 7 (eta) (15.3 kD) (Cct7) alternative variant cSep08, mRNA.
<a href="#">Cct8</a>	<a href="#">Cct8.bSep08</a>	<a href="#">288305</a>	10250	1524	14	507	chaperonin subunit 8 (theta) (Cct8) alternative variant bSep08, mRNA.
<a href="#">Cct8</a>	<a href="#">Cct8.cSep08</a>	<a href="#">288305</a>	1708	563	5	178	chaperonin subunit 8 (theta) (Cct8) alternative variant cSep08, mRNA.
<a href="#">Cct8</a>	<a href="#">Cct8.dSep08</a>	<a href="#">288305</a>	1565	723	3	145	chaperonin subunit 8 (theta) (Cct8) alternative variant dSep08, mRNA.
<a href="#">Cd2ap</a>	<a href="#">Cd2ap.bSep08</a>	<a href="#">316258</a>	34645	761	4	61	CD2-associated protein (6.9 kD) (Cd2ap) alternative variant bSep08, mRNA.
<a href="#">Cd2bp2</a>	<a href="#">Cd2bp2.bSep08</a>	<a href="#">293505</a>	1140	457	3	106	CD2 antigen (cytoplasmic tail) binding protein 2 (Cd2bp2) alternative variant bSep08, mRNA.
<a href="#">Cd2bp2</a>	<a href="#">Cd2bp2.dSep08</a>	<a href="#">293505</a>	365	221	2	40	CD2 antigen (cytoplasmic tail) binding protein 2 (4.3 kD) (Cd2bp2) alternative variant dSep08, complete mRNA.
<a href="#">Cd3d</a>	<a href="#">Cd3d.bSep08</a>	<a href="#">25710</a>	622	324	1	79	CD3 molecule delta polypeptide (8.8 kD) (Cd3d) alternative variant bSep08, mRNA.

<a href="#">Cd3e</a>	<a href="#">Cd3e.aSep08</a>	<a href="#">315609</a>	9037	504	6	167	CD3 molecule, epsilon polypeptide (Cd3e) alternative variant aSep08, mRNA.
<a href="#">Cd3e</a>	<a href="#">Cd3e.bSep08</a>	<a href="#">315609</a>	8736	1033	5	140	CD3 molecule, epsilon polypeptide (15.7 kD) (Cd3e) alternative variant bSep08, mRNA.
<a href="#">Cd3g</a>	<a href="#">Cd3g.aSep08</a>	<a href="#">300678</a>	6598	756	7	203	CD3 molecule, gamma polypeptide (Cd3g) alternative variant aSep08, mRNA.
<a href="#">Cd3g</a>	<a href="#">Cd3g.cSep08</a>	<a href="#">300678</a>	1347	342	2	44	CD3 molecule, gamma polypeptide (Cd3g) alternative variant cSep08, mRNA.
<a href="#">Cd4</a>	<a href="#">Cd4.bSep08</a>	<a href="#">24932</a>	776	648	2	98	CD4 antigen (Cd4) alternative variant bSep08, mRNA.
<a href="#">Cd5</a>	<a href="#">Cd5.bSep08</a>	<a href="#">54236</a>	15771	1031	7	307	CD5 antigen (Cd5) alternative variant bSep08, mRNA.
<a href="#">Cd6</a>	<a href="#">Cd6.bSep08</a>	<a href="#">25752</a>	4345	767	3	222	CD6 antigen (Cd6) alternative variant bSep08, mRNA.
<a href="#">Cd6</a>	<a href="#">Cd6.cSep08</a>	<a href="#">25752</a>	4102	483	2	160	CD6 antigen (Cd6) alternative variant cSep08, mRNA.
<a href="#">Cd8a</a>	<a href="#">Cd8a.aSep08</a>	<a href="#">24930</a>	4828	2065	5	239	CD8 antigen, alpha chain (26.4 kD) (Cd8a) alternative variant aSep08, mRNA.
<a href="#">Cd8a</a>	<a href="#">Cd8a.cSep08</a>	<a href="#">24930</a>	798	712	2	141	CD8 antigen, alpha chain (Cd8a) alternative variant cSep08, mRNA.
<a href="#">Cd8a</a>	<a href="#">Cd8a.dSep08</a>	<a href="#">24930</a>	1734	750	2	100	CD8 antigen, alpha chain (Cd8a) alternative variant dSep08, mRNA.
<a href="#">Cd14</a>	<a href="#">Cd14.bSep08</a>	<a href="#">60350</a>	1595	1544	2	369	CD14 molecule (39.7 kD) (Cd14) alternative variant bSep08, mRNA.
<a href="#">Cd14</a>	<a href="#">Cd14.cSep08</a>	<a href="#">60350</a>	31733	315	2	56	CD14 molecule (Cd14) alternative variant cSep08, mRNA.
<a href="#">Cd19</a>	<a href="#">Cd19.aSep08</a>	<a href="#">365367</a>	5679	2167	13	470	CD19 molecule (51.5 kD) (Cd19) alternative variant aSep08, mRNA.
<a href="#">Cd19</a>	<a href="#">Cd19.bSep08</a>	<a href="#">365367</a>	3245	814	9	271	CD19 molecule (Cd19) alternative variant bSep08, mRNA.
<a href="#">Cd19</a>	<a href="#">Cd19.cSep08</a>	<a href="#">365367</a>	1468	988	3	58	CD19 molecule (Cd19) alternative variant cSep08, mRNA.
<a href="#">Cd19</a>	<a href="#">Cd19.dSep08</a>	<a href="#">365367</a>	1216	1086	2	52	CD19 molecule (5.7 kD) (Cd19) alternative variant dSep08, mRNA.
<a href="#">Cd22</a>	<a href="#">Cd22.bSep08</a>	<a href="#">308501</a>	2670	1882	2	44	CD22 molecule (5.0 kD) (Cd22) alternative variant bSep08, mRNA.
<a href="#">Cd24</a>	<a href="#">Cd24.aSep08</a>	<a href="#">25145</a>	3027	418	2	118	CD24 molecule (Cd24) alternative variant aSep08, mRNA.
<a href="#">Cd24</a>	<a href="#">Cd24.dSep08</a>	<a href="#">25145</a>	5212	492	3	70	CD24 molecule (Cd24) alternative variant dSep08, mRNA.
<a href="#">Cd27</a>	<a href="#">Cd27.aSep08</a>	<a href="#">500318</a>	3912	818	5	252	CD27 molecule (Cd27) alternative variant aSep08, mRNA.
<a href="#">Cd27</a>	<a href="#">Cd27.cSep08</a>	<a href="#">500318</a>	1728	796	4	202	CD27 molecule (Cd27) alternative variant cSep08, mRNA.
<a href="#">Cd27</a>	<a href="#">Cd27.dSep08</a>	<a href="#">500318</a>	2869	554	3	180	CD27 molecule (Cd27) alternative variant dSep08, mRNA.
<a href="#">Cd28</a>	<a href="#">Cd28.bSep08</a>	<a href="#">25660</a>	87433	732	5	65	cd28 molecule (Cd28) alternative variant bSep08, mRNA.
<a href="#">Cd28</a>	<a href="#">Cd28.cSep08</a>	<a href="#">25660</a>	1368	535	2	44	cd28 molecule (4.8 kD) (Cd28) alternative variant cSep08, mRNA.
<a href="#">Cd34</a>	<a href="#">Cd34.bSep08</a>	<a href="#">305081</a>	19295	1478	1	384	CD34 molecule (Cd34) alternative variant bSep08, mRNA.
<a href="#">Cd36</a>	<a href="#">Cd36.bSep08</a>	<a href="#">29184</a>	17550	791	7	263	cd36 antigen (Cd36) alternative variant bSep08, mRNA.
<a href="#">Cd36</a>	<a href="#">Cd36.cSep08</a>	<a href="#">29184</a>	36740	776	5	230	cd36 antigen (Cd36) alternative variant cSep08, mRNA.
<a href="#">Cd36</a>	<a href="#">Cd36.dSep08</a>	<a href="#">29184</a>	35624	737	5	185	cd36 antigen (Cd36) alternative variant dSep08, mRNA.
<a href="#">Cd36</a>	<a href="#">Cd36.eSep08</a>	<a href="#">29184</a>	9577	506	3	105	cd36 antigen (Cd36) alternative variant eSep08, mRNA.
<a href="#">Cd37</a>	<a href="#">Cd37.bSep08</a>	<a href="#">29185</a>	4633	1961	2	192	CD37 antigen like precursor (21.2 kD) (Cd37) alternative variant bSep08, mRNA.



<a href="#">Cd37</a>	<a href="#">Cd37.cSep08</a>	<a href="#">29185</a>	1471	798	2	69	CD37 antigen like (7.9 kD) (Cd37) alternative variant cSep08, mRNA.
<a href="#">Cd40</a>	<a href="#">Cd40.bSep08</a>	<a href="#">171369</a>	13442	768	7	234	CD40 molecule, TNF receptor superfamily member 5 (25.6 kD) (Cd40) alternative variant bSep08, mRNA.
<a href="#">Cd40</a>	<a href="#">Cd40.cSep08</a>	<a href="#">171369</a>	10716	603	6	194	CD40 molecule, TNF receptor superfamily member 5 (Cd40) alternative variant cSep08, mRNA.
<a href="#">Cd40</a>	<a href="#">Cd40.dSep08</a>	<a href="#">171369</a>	1527	677	4	107	CD40 molecule, TNF receptor superfamily member 5 (Cd40) alternative variant dSep08, mRNA.
<a href="#">Cd44</a>	<a href="#">Cd44.aSep08</a>	<a href="#">25406</a>	88627	3984	5	410	CD44 antigen (Cd44) alternative variant aSep08, mRNA.
<a href="#">Cd44</a>	<a href="#">Cd44.bSep08</a>	<a href="#">25406</a>	18162	537	2	50	CD44 antigen (Cd44) alternative variant bSep08, mRNA.
<a href="#">Cd46</a>	<a href="#">Cd46.bSep08</a>	<a href="#">29333</a>	9378	487	1	42	CD46 antigen, complement regulatory protein (Cd46) alternative variant bSep08, mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.aSep08</a>	<a href="#">29364</a>	59069	1368	12	359	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (38.9 kD) (Cd47) alternative variant aSep08, complete mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.cSep08</a>	<a href="#">29364</a>	13832	735	3	225	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (24.6 kD) (Cd47) alternative variant cSep08, complete mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.dSep08</a>	<a href="#">29364</a>	17165	691	7	132	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (Cd47) alternative variant dSep08, mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.eSep08</a>	<a href="#">29364</a>	4551	819	2	49	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (5.8 kD) (Cd47) alternative variant eSep08, mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.fSep08</a>	<a href="#">29364</a>	14951	482	6	61	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (Cd47) alternative variant fSep08, mRNA.
<a href="#">Cd47</a>	<a href="#">Cd47.gSep08</a>	<a href="#">29364</a>	14895	401	5	42	CD47 antigen (Rh-related antigen, integrin-associated signal transducer) (Cd47) alternative variant gSep08, mRNA.
<a href="#">Cd48</a>	<a href="#">Cd48.bSep08</a>	<a href="#">245962</a>	18198	654	3	192	CD48 antigen (Cd48) alternative variant bSep08, mRNA.
<a href="#">Cd48</a>	<a href="#">Cd48.cSep08</a>	<a href="#">245962</a>	17550	537	3	177	CD48 antigen (Cd48) alternative variant cSep08, mRNA.
<a href="#">Cd48</a>	<a href="#">Cd48.dSep08</a>	<a href="#">245962</a>	23544	1275	4	117	CD48 antigen (13.5 kD) (Cd48) alternative variant dSep08, complete mRNA.
<a href="#">Cd55</a>	<a href="#">Cd55.aSep08</a>	<a href="#">64036</a>	16764	1765	9	430	CD55 antigen (48.5 kD) (Cd55) alternative variant aSep08, mRNA.
<a href="#">Cd55</a>	<a href="#">Cd55.cSep08</a>	<a href="#">64036</a>	11277	1421	3	381	CD55 antigen (Cd55) alternative variant cSep08, mRNA.
<a href="#">Cd55</a>	<a href="#">Cd55.dSep08</a>	<a href="#">64036</a>	1810	704	2	218	CD55 antigen (Cd55) alternative variant dSep08, mRNA.
<a href="#">Cd55</a>	<a href="#">Cd55.eSep08</a>	<a href="#">64036</a>	20521	1370	6	189	CD55 antigen (Cd55) alternative variant eSep08, mRNA.
<a href="#">Cd63</a>	<a href="#">Cd63.bSep08</a>	<a href="#">29186</a>	1425	752		218	CD63 antigen (Cd63) alternative variant bSep08, mRNA.
<a href="#">Cd68</a>	<a href="#">Cd68.bSep08</a>	<a href="#">287435</a>	835	704	1	234	CD68 antigen (Cd68) alternative variant bSep08, mRNA.
<a href="#">Cd69</a>	<a href="#">Cd69.bSep08</a>	<a href="#">29187</a>	7179	741		158	CD69 antigen (17.9 kD) (Cd69) alternative variant bSep08, mRNA.
<a href="#">Cd72</a>	<a href="#">Cd72.bSep08</a>	<a href="#">313498</a>	1575	932	1	144	CD72 antigen (15.7 kD) (Cd72) alternative variant bSep08, mRNA.

<a href="#">Cd74</a>	<a href="#">Cd74.aSep08</a>	<a href="#">25599</a>	9202	1394	9	312	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (Cd74) alternative variant aSep08, mRNA.
<a href="#">Cd74</a>	<a href="#">Cd74.bSep08</a>	<a href="#">25599</a>	8651	2341	6	297	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (Cd74) alternative variant bSep08, mRNA.
<a href="#">Cd74</a>	<a href="#">Cd74.dSep08</a>	<a href="#">25599</a>	828	652	2	165	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (17.1 kD) (Cd74) alternative variant dSep08, mRNA.
<a href="#">Cd74</a>	<a href="#">Cd74.fSep08</a>	<a href="#">25599</a>	3418	768	5	103	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (Cd74) alternative variant fSep08, mRNA.
<a href="#">Cd79a</a>	<a href="#">Cd79a.aSep08</a>	<a href="#">681236</a>	2702	936	2	311	CD79A antigen (immunoglobulin-associated alpha) (Cd79a) alternative variant aSep08, mRNA.
<a href="#">Cd79a</a>	<a href="#">Cd79a.bSep08</a>	<a href="#">681236</a>	4337	1196	2	94	CD79A antigen (immunoglobulin-associated alpha) (10.2 kD) (Cd79a) alternative variant bSep08, mRNA.
<a href="#">Cd81</a>	<a href="#">Cd81.bSep08</a>	<a href="#">25621</a>	14034	681	4	168	CD 81 antigen (Cd81) alternative variant bSep08, mRNA.
<a href="#">Cd81</a>	<a href="#">Cd81.cSep08</a>	<a href="#">25621</a>	2210	1452	4	123	CD 81 antigen (13.6 kD) (Cd81) alternative variant cSep08, mRNA.
<a href="#">Cd81</a>	<a href="#">Cd81.dSep08</a>	<a href="#">25621</a>	5924	2327	6	101	CD 81 antigen (11.0 kD) (Cd81) alternative variant dSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.bSep08</a>	<a href="#">83628</a>	43078	1034	8	238	cd82 antigen CRA c like (Cd82) alternative variant bSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.cSep08</a>	<a href="#">83628</a>	43038	753	8	202	cd82 antigen CRA c like (Cd82) alternative variant cSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.dSep08</a>	<a href="#">83628</a>	6625	892	5	150	cd82 antigen like (16.8 kD) (Cd82) alternative variant dSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.eSep08</a>	<a href="#">83628</a>	33225	380	5	126	putative protein (Cd82) alternative variant eSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.fSep08</a>	<a href="#">83628</a>	33538	391	5	95	cd82 antigen CRA c like (Cd82) alternative variant fSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.gSep08</a>	<a href="#">83628</a>	1909	822	2	95	cd82 antigen like (Cd82) alternative variant gSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.hSep08</a>	<a href="#">83628</a>	29750	401	4	83	putative protein (Cd82) alternative variant hSep08, mRNA.
<a href="#">Cd82</a>	<a href="#">Cd82.iSep08</a>	<a href="#">83628</a>	827	533	2	51	cd82 antigen like (5.7 kD) (Cd82) alternative variant iSep08, mRNA.
<a href="#">Cd96</a>	<a href="#">Cd96.bSep08</a>	<a href="#">498079</a>	31459	846	1	158	CD96 antigen (17.8 kD) (Cd96) alternative variant bSep08, mRNA.
<a href="#">Cd97</a>	<a href="#">Cd97.bSep08</a>	<a href="#">361383</a>	2391	844	5	274	CD97 antigen (Cd97) alternative variant bSep08, mRNA.
<a href="#">Cd97</a>	<a href="#">Cd97.cSep08</a>	<a href="#">361383</a>	5788	710	6	236	CD97 antigen (Cd97) alternative variant cSep08, mRNA.
<a href="#">Cd97</a>	<a href="#">Cd97.eSep08</a>	<a href="#">361383</a>	843	763	2	99	CD97 antigen (11.2 kD) (Cd97) alternative variant eSep08, mRNA.
<a href="#">Cd99I2</a>	<a href="#">Cd99I2.bSep08</a>	<a href="#">171485</a>	3496	284	1	54	cd99 antigen-like 2 (6.2 kD) (Cd99I2) alternative variant bSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.aSep08</a>	<a href="#">64315</a>	4013	1782	6	493	CD151 molecule (Raph blood group) (Cd151) alternative variant aSep08, complete mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.bSep08</a>	<a href="#">64315</a>	2645	1574	6	253	CD151 molecule (Raph blood group) (28.4 kD) (Cd151) alternative variant bSep08, mRNA.

<a href="#">Cd151</a>	<a href="#">Cd151.cSep08</a>	<a href="#">64315</a>	2938	1540	6	253	CD151 molecule (Raph blood group) (28.4 kD) (Cd151) alternative variant cSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.eSep08</a>	<a href="#">64315</a>	2884	717	5	172	CD151 molecule (Raph blood group) (18.9 kD) (Cd151) alternative variant eSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.fSep08</a>	<a href="#">64315</a>	2680	744	6	167	CD151 molecule (Raph blood group) (Cd151) alternative variant fSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.gSep08</a>	<a href="#">64315</a>	3395	1430	5	123	CD151 molecule (Raph blood group) (13.2 kD) (Cd151) alternative variant gSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.hSep08</a>	<a href="#">64315</a>	1703	526	3	68	CD151 molecule (Raph blood group) (Cd151) alternative variant hSep08, mRNA.
<a href="#">Cd151</a>	<a href="#">Cd151.iSep08</a>	<a href="#">64315</a>	1299	389	2	41	CD151 molecule (Raph blood group) (4.4 kD) (Cd151) alternative variant iSep08, mRNA.
<a href="#">Cd163</a>	<a href="#">Cd163.bSep08</a>	<a href="#">312701</a>	2499	1419	2	51	CD163 molecule (Cd163) alternative variant bSep08, mRNA.
<a href="#">Cd163</a>	<a href="#">Cd163.cSep08</a>	<a href="#">312701</a>	3940	325	3	32	CD163 molecule (Cd163) alternative variant cSep08, mRNA.
<a href="#">Cd177</a>	<a href="#">Cd177.aSep08</a>	<a href="#">499099</a>	1272	640		128	CD177 antigen (Cd177) mRNA.
<a href="#">Cd200</a>	<a href="#">Cd200.aSep08</a>	<a href="#">24560</a>	32663	1105	5	313	cd200 molecule (Cd200) alternative variant aSep08, mRNA.
<a href="#">Cd200</a>	<a href="#">Cd200.cSep08</a>	<a href="#">24560</a>	17680	936	5	184	cd200 molecule (Cd200) alternative variant cSep08, mRNA.
<a href="#">Cd200</a>	<a href="#">Cd200.dSep08</a>	<a href="#">24560</a>	15330	666	3	157	cd200 molecule (Cd200) alternative variant dSep08, mRNA.
<a href="#">Cd200</a>	<a href="#">Cd200.eSep08</a>	<a href="#">24560</a>	13775	787	4	140	cd200 molecule (Cd200) alternative variant eSep08, mRNA.
<a href="#">Cd209b</a>	<a href="#">Cd209b.aSep08</a>	<a href="#">288378</a>	8540	1148		333	CD209b antigen (Cd209b) mRNA.
<a href="#">Cd209g</a>	<a href="#">Cd209g.aSep08</a>	<a href="#">688750</a>	1510	393		130	CD209g molecule (Cd209g) mRNA.
<a href="#">CD225.0</a>	<a href="#">CD225.0.aSep08</a>		780	612		91	interferon-induced transmembrane protein (CD225.0) mRNA.
<a href="#">CD225.1</a>	<a href="#">CD225.1.aSep08</a>		1339	791		134	interferon induced transmembrane protein 5 like (14.6 kD) (CD225.1) mRNA.
<a href="#">Cd226</a>	<a href="#">Cd226.bSep08</a>	<a href="#">307199</a>	72384	388	1	68	CD226 molecule (Cd226) alternative variant bSep08, mRNA.
<a href="#">Cd247</a>	<a href="#">Cd247.bSep08</a>	<a href="#">25300</a>	78452	743		137	cd247 molecule (15.4 kD) (Cd247) alternative variant bSep08, mRNA.
<a href="#">Cd276</a>	<a href="#">Cd276.bSep08</a>	<a href="#">315716</a>	3005	331	3	36	cd276 molecule (Cd276) alternative variant bSep08, mRNA.
<a href="#">Cd276</a>	<a href="#">Cd276.cSep08</a>	<a href="#">315716</a>	2128	714	2	44	cd276 molecule (5.0 kD) (Cd276) alternative variant cSep08, mRNA.
<a href="#">Cd300le</a>	<a href="#">Cd300le.aSep08</a>	<a href="#">360655</a>	5632	899		206	CD300 molecule-like family member E (Cd300le) mRNA.
<a href="#">Cd300lf</a>	<a href="#">Cd300lf.aSep08</a>	<a href="#">287818</a>	15908	1174	1	370	CD300 molecule-like family member f (Cd300lf) alternative variant aSep08, mRNA.
<a href="#">Cd320</a>	<a href="#">Cd320.bSep08</a>	<a href="#">362851</a>	4889	716	3	203	CD320 molecule (Cd320) alternative variant bSep08, mRNA.
<a href="#">Cd320</a>	<a href="#">Cd320.cSep08</a>	<a href="#">362851</a>	2538	1017	4	130	CD320 molecule (Cd320) alternative variant cSep08, complete mRNA.

<a href="#">Cd320</a>	<a href="#">Cd320.dSep08</a>	<a href="#">362851</a>	1108	603	3	91	CD320 molecule (Cd320) alternative variant dSep08, mRNA.
<a href="#">Cdadc1</a>	<a href="#">Cdadc1.aSep08</a>	<a href="#">361052</a>	28110	2209	9	478	CMP/dCMP deaminase, zinc-binding (54.2 kD) (Cdadc1) alternative variant aSep08, mRNA.
<a href="#">Cdadc1</a>	<a href="#">Cdadc1.cSep08</a>	<a href="#">361052</a>	11352	772	4	151	putative protein of ancient origin (Cdadc1) alternative variant cSep08, mRNA.
<a href="#">Cdadc1</a>	<a href="#">Cdadc1.dSep08</a>	<a href="#">361052</a>	6613	741	3	147	putative protein of ancient origin (Cdadc1) alternative variant dSep08, mRNA.
<a href="#">Cdadc1</a>	<a href="#">Cdadc1.eSep08</a>	<a href="#">361052</a>	615	427	2	83	putative protein of mammalian origin (9.2 kD) (Cdadc1) alternative variant eSep08, mRNA.
<a href="#">Cdadc1</a>	<a href="#">Cdadc1.gSep08</a>	<a href="#">361052</a>	4686	441	3	51	putative protein of vertebrate origin (Cdadc1) alternative variant gSep08, mRNA.
<a href="#">Cdc2a</a>	<a href="#">Cdc2a.bSep08</a>	<a href="#">54237</a>	7048	800	6	220	cell division cycle 2 homolog A (S. pombe) (Cdc2a) alternative variant bSep08, mRNA.
<a href="#">Cdc2a</a>	<a href="#">Cdc2a.cSep08</a>	<a href="#">54237</a>	12416	797	7	204	cell division cycle 2 homolog A (S. pombe) (Cdc2a) alternative variant cSep08, mRNA.
<a href="#">Cdc2l1</a>	<a href="#">Cdc2l1.aSep08</a>	<a href="#">252879</a>	25175	2027	15	563	cell division cycle 2 homolog (S.pombe)-like 1 (Cdc2l1) alternative variant aSep08, mRNA.
<a href="#">Cdc2l1</a>	<a href="#">Cdc2l1.bSep08</a>	<a href="#">252879</a>	2396	1143	8	330	cell division cycle 2 homolog (S.pombe)-like 1 (Cdc2l1) alternative variant bSep08, mRNA.
<a href="#">Cdc2l1</a>	<a href="#">Cdc2l1.cSep08</a>	<a href="#">252879</a>	5374	368	4	122	cell division cycle 2 homolog (S.pombe)-like 1 (Cdc2l1) alternative variant cSep08, mRNA.
<a href="#">Cdc2l1</a>	<a href="#">Cdc2l1.dSep08</a>	<a href="#">252879</a>	1036	726	2	64	cell division cycle 2 homolog (S.pombe)-like 1 (Cdc2l1) alternative variant dSep08, mRNA.
<a href="#">Cdc2l5</a>	<a href="#">Cdc2l5.aSep08</a>	<a href="#">306998</a>	53229	3281	5	803	cell division cycle 2-like 5 (cholinesterase-related cell division controller) (Cdc2l5) alternative variant aSep08, mRNA.
<a href="#">Cdc2l5</a>	<a href="#">Cdc2l5.bSep08</a>	<a href="#">306998</a>	9611	919	5	306	cell division cycle 2-like 5 (cholinesterase-related cell division controller) (Cdc2l5) alternative variant bSep08, mRNA.
<a href="#">Cdc2l5</a>	<a href="#">Cdc2l5.cSep08</a>	<a href="#">306998</a>	12209	773	1	51	cell division cycle 2-like 5 (cholinesterase-related cell division controller) (Cdc2l5) alternative variant cSep08, mRNA.
<a href="#">Cdc2l6</a>	<a href="#">Cdc2l6.bSep08</a>	<a href="#">309804</a>	97216	735	4	111	cell division cycle 2-like 6 (CDK8-like) (12.4 kD) (Cdc2l6) alternative variant bSep08, mRNA.
<a href="#">Cdc2l6</a>	<a href="#">Cdc2l6.eSep08</a>	<a href="#">309804</a>	110859	1007	6	87	cell division cycle 2-like 6 (CDK8-like) (9.5 kD) (Cdc2l6) alternative variant eSep08, mRNA.
<a href="#">Cdc2l6</a>	<a href="#">Cdc2l6.fSep08</a>	<a href="#">309804</a>	3045	722	3	30	cell division cycle 2-like 6 (CDK8-like) (Cdc2l6) alternative variant fSep08, mRNA.
<a href="#">Cdc5l</a>	<a href="#">Cdc5l.bSep08</a>	<a href="#">85434</a>	22787	1021	6	340	cell division cycle 5-like (S. pombe) (Cdc5l) alternative variant bSep08, mRNA.
<a href="#">Cdc5l</a>	<a href="#">Cdc5l.cSep08</a>	<a href="#">85434</a>	11289	754	2	100	cell division cycle 5-like (S. pombe) (12.0 kD) (Cdc5l) alternative variant cSep08, mRNA.
<a href="#">Cdc7</a>	<a href="#">Cdc7.bSep08</a>	<a href="#">360908</a>	7881	862	2	205	cell division cycle 7 (S. cerevisiae) (Cdc7) alternative variant bSep08, mRNA.
<a href="#">Cdc7</a>	<a href="#">Cdc7.cSep08</a>	<a href="#">360908</a>	9305	1493	3	88	cell division cycle 7 (S. cerevisiae) (9.4 kD) (Cdc7) alternative variant cSep08, mRNA.

<a href="#">Cdc14a</a>	<a href="#">Cdc14a.cSep08</a>	<a href="#">310806</a>	41638	384	4	43	CDC14 cell division cycle 14 homolog A ( <i>S. cerevisiae</i> ) (5.0 kD) (Cdc14a) alternative variant cSep08, mRNA.
<a href="#">Cdc14b</a>	<a href="#">Cdc14b.aSep08</a>	<a href="#">361195</a>	25569	748		212	CDC14 cell division cycle 14 homolog B ( <i>S. cerevisiae</i> ) (Cdc14b) mRNA.
<a href="#">Cdc16</a>	<a href="#">Cdc16.bSep08</a>	<a href="#">290875</a>	9904	731	6	212	CDC16 cell division cycle 16 homolog ( <i>S. cerevisiae</i> ) (Cdc16) alternative variant bSep08, mRNA.
<a href="#">Cdc16</a>	<a href="#">Cdc16.cSep08</a>	<a href="#">290875</a>	5615	865	4	129	CDC16 cell division cycle 16 homolog ( <i>S. cerevisiae</i> ) (14.8 kD) (Cdc16) alternative variant cSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.bSep08</a>	<a href="#">291689</a>	6013	955	8	205	CDC23 (cell division cycle 23, yeast, homolog) (Cdc23) alternative variant bSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.cSep08</a>	<a href="#">291689</a>	8595	819	5	176	CDC23 (cell division cycle 23, yeast, homolog) (Cdc23) alternative variant cSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.dSep08</a>	<a href="#">291689</a>	1303	748	3	86	CDC23 (cell division cycle 23, yeast, homolog) (Cdc23) alternative variant dSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.fSep08</a>	<a href="#">291689</a>	2123	851	2	38	CDC23 (cell division cycle 23, yeast, homolog) (Cdc23) alternative variant fSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.hSep08</a>	<a href="#">291689</a>	1841	564	2	16	CDC23 (cell division cycle 23, yeast, homolog) (2.0 kD) (Cdc23) alternative variant hSep08, mRNA.
<a href="#">Cdc23</a>	<a href="#">Cdc23.iSep08</a>	<a href="#">291689</a>	3621	539	3	54	CDC23 (cell division cycle 23, yeast, homolog) (Cdc23) alternative variant iSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.bSep08</a>	<a href="#">171103</a>	4741	772	7	232	cell division cycle 25 homolog B CRA b (Cdc25b) alternative variant bSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.cSep08</a>	<a href="#">171103</a>	3235	728	4	152	M-phase inducer phosphatase (Cdc25b) alternative variant cSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.dSep08</a>	<a href="#">171103</a>	2008	441	5	146	cell division cycle 25 homolog B (Cdc25b) alternative variant dSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.eSep08</a>	<a href="#">171103</a>	1731	753	3	125	cell division cycle 25 homolog B CRA b (Cdc25b) alternative variant eSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.fSep08</a>	<a href="#">171103</a>	1414	671	2	86	cell division cycle 25 homolog B CRA a (Cdc25b) alternative variant fSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.gSep08</a>	<a href="#">171103</a>	501	415	2	82	cell division cycle 25 homolog B CRA b (Cdc25b) alternative variant gSep08, mRNA.
<a href="#">Cdc25b</a>	<a href="#">Cdc25b.hSep08</a>	<a href="#">171103</a>	1309	517	2	64	cell division cycle 25 homolog B (Cdc25b) alternative variant hSep08, mRNA.
<a href="#">Cdc25c</a>	<a href="#">Cdc25c.bSep08</a>	<a href="#">307511</a>	5472	677	5	138	cell division cycle 25C (Cdc25c) alternative variant bSep08, mRNA.
<a href="#">Cdc25c</a>	<a href="#">Cdc25c.cSep08</a>	<a href="#">307511</a>	14272	1799	1	95	cdc25 like (10.4 kD) (Cdc25c) alternative variant cSep08, mRNA.
<a href="#">Cdc26</a>	<a href="#">Cdc26.aSep08</a>	<a href="#">366381</a>	14911	2712	2	85	cell division cycle 26 (9.8 kD) (Cdc26) alternative variant aSep08, complete mRNA.
<a href="#">Cdc26</a>	<a href="#">Cdc26.cSep08</a>	<a href="#">366381</a>	12886	723	3	85	cell division cycle 26 (9.8 kD) (Cdc26) alternative variant cSep08, complete mRNA.
<a href="#">Cdc26</a>	<a href="#">Cdc26.eSep08</a>	<a href="#">366381</a>	8834	398	2	19	cell division cycle 26 (Cdc26) alternative variant eSep08, mRNA.
<a href="#">Cdc27</a>	<a href="#">Cdc27.bSep08</a>	<a href="#">360643</a>	9598	858	6	259	cell division cycle 27 homolog ( <i>S. cerevisiae</i> ) (Cdc27) alternative variant bSep08, mRNA.

<a href="#">Cdc27</a>	<a href="#">Cdc27.cSep08</a>	<a href="#">360643</a>	11151	735	6	244	cell division cycle 27 homolog ( <i>S. cerevisiae</i> ) (Cdc27) alternative variant cSep08, mRNA.
<a href="#">Cdc34</a>	<a href="#">Cdc34.bSep08</a>	<a href="#">299602</a>	3169	1262	2	153	cell division cycle 34 homolog ( <i>S. cerevisiae</i> ) (17.1 kD) (Cdc34) alternative variant bSep08, mRNA.
<a href="#">Cdc34</a>	<a href="#">Cdc34.cSep08</a>	<a href="#">299602</a>	2106	393	4	130	cell division cycle 34 homolog ( <i>S. cerevisiae</i> ) (Cdc34) alternative variant cSep08, mRNA.
<a href="#">Cdc37</a>	<a href="#">Cdc37.bSep08</a>	<a href="#">114562</a>	8330	560	3	174	cell division cycle 37 homolog ( <i>S. cerevisiae</i> ) (Cdc37) alternative variant bSep08, mRNA.
<a href="#">Cdc37</a>	<a href="#">Cdc37.cSep08</a>	<a href="#">114562</a>	1018	846	3	168	cell division cycle 37 homolog ( <i>S. cerevisiae</i> ) (19.6 kD) (Cdc37) alternative variant cSep08, mRNA.
<a href="#">Cdc37</a>	<a href="#">Cdc37.dSep08</a>	<a href="#">114562</a>	9994	1374	5	159	cell division cycle 37 homolog ( <i>S. cerevisiae</i> ) (Cdc37) alternative variant dSep08, mRNA.
<a href="#">Cdc3711</a>	<a href="#">Cdc3711.bSep08</a>	<a href="#">293886</a>	9034	458	4	115	cell division cycle 37 homolog ( <i>S. cerevisiae</i> )-like 1 (Cdc3711) alternative variant bSep08, mRNA.
<a href="#">Cdc3711</a>	<a href="#">Cdc3711.cSep08</a>	<a href="#">293886</a>	10605	723	3	72	cell division cycle 37 homolog ( <i>S. cerevisiae</i> )-like 1 (Cdc3711) alternative variant cSep08, mRNA.
<a href="#">Cdc3711</a>	<a href="#">Cdc3711.fSep08</a>	<a href="#">293886</a>	8244	514	3	44	cell division cycle 37 homolog ( <i>S. cerevisiae</i> )-like 1 (Cdc3711) alternative variant fSep08, mRNA.
<a href="#">Cdc40</a>	<a href="#">Cdc40.aSep08</a>	<a href="#">361859</a>	22111	2669	12	444	cell division cycle 40 homolog (yeast) (Cdc40) alternative variant aSep08, mRNA.
<a href="#">Cdc42</a>	<a href="#">Cdc42.aSep08</a>	<a href="#">64465</a>	35130	2573	2	191	cell division cycle 42 homolog ( <i>S. cerevisiae</i> ) (21.3 kD) (Cdc42) alternative variant aSep08, mRNA.
<a href="#">Cdc42</a>	<a href="#">Cdc42.cSep08</a>	<a href="#">64465</a>	37145	2603	2	191	cell division cycle 42 homolog ( <i>S. cerevisiae</i> ) (21.3 kD) (Cdc42) alternative variant cSep08, mRNA.
<a href="#">Cdc42bpa</a>	<a href="#">Cdc42bpa.bSep08</a>	<a href="#">114116</a>	33255	868	9	289	CDC42 binding protein kinase alpha (Cdc42bpa) alternative variant bSep08, mRNA.
<a href="#">Cdc42bpb</a>	<a href="#">Cdc42bpb.bSep08</a>	<a href="#">113960</a>	1370	616	3	152	cdc42 binding protein kinase beta (Cdc42bpb) alternative variant bSep08, mRNA.
<a href="#">Cdc42bpb</a>	<a href="#">Cdc42bpb.cSep08</a>	<a href="#">113960</a>	495	446	2	57	cdc42 binding protein kinase beta (Cdc42bpb) alternative variant cSep08, mRNA.
<a href="#">Cdc42bpg</a>	<a href="#">Cdc42bpg.bSep08</a>	<a href="#">293693</a>	4915	495	1	165	CDC42 binding protein kinase gamma (DMPK-like) (Cdc42bpg) alternative variant bSep08, mRNA.
<a href="#">Cdc42ep1</a>	<a href="#">Cdc42ep1.bSep08</a>	<a href="#">315121</a>	4208	494	3	48	putative protein (Cdc42ep1) alternative variant bSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.bSep08</a>	<a href="#">309175</a>	8865	1507	2	214	putative protein (Cdc42ep2) alternative variant bSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.cSep08</a>	<a href="#">309175</a>	7150	603	2	117	putative nuclear protein (12.8 kD) (Cdc42ep2) alternative variant cSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.dSep08</a>	<a href="#">309175</a>	7822	642	3	114	putative protein (12.6 kD) (Cdc42ep2) alternative variant dSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.eSep08</a>	<a href="#">309175</a>	7656	1005	3	88	putative nuclear protein (9.6 kD) (Cdc42ep2) alternative variant eSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.fSep08</a>	<a href="#">309175</a>	4638	703	3	85	putative protein (9.3 kD) (Cdc42ep2) alternative variant fSep08, mRNA.
<a href="#">Cdc42ep2</a>	<a href="#">Cdc42ep2.gSep08</a>	<a href="#">309175</a>	7496	361	3	71	putative protein (Cdc42ep2) alternative variant gSep08, mRNA.

<a href="#">Cdc42ep4</a>	<a href="#">Cdc42ep4.bSep08</a>	<a href="#">303653</a>	22141	1187	1	92	CDC42 effector protein (Rho GTPase binding) 4 (10.4 kD) (Cdc42ep4) alternative variant bSep08, mRNA.
<a href="#">Cdc42ep4</a>	<a href="#">Cdc42ep4.cSep08</a>	<a href="#">303653</a>	10192	754	1	203	CDC42 effector protein (Rho GTPase binding) 4 (Cdc42ep4) alternative variant cSep08, mRNA.
<a href="#">Cdc42ep4</a>	<a href="#">Cdc42ep4.dSep08</a>	<a href="#">303653</a>	658	347	1	62	CDC42 effector protein (Rho GTPase binding) 4 (Cdc42ep4) alternative variant dSep08, mRNA.
<a href="#">Cdc42ep5</a>	<a href="#">Cdc42ep5.aSep08</a>	<a href="#">361505</a>	1729	620	2	161	CDC42 effector protein (Rho GTPase binding) 5 (Cdc42ep5) alternative variant aSep08, mRNA.
<a href="#">Cdc42ep5</a>	<a href="#">Cdc42ep5.bSep08</a>	<a href="#">361505</a>	2124	765	2	153	CDC42 effector protein (Rho GTPase binding) 5 (15.7 kD) (Cdc42ep5) alternative variant bSep08, mRNA.
<a href="#">Cdc42se1</a>	<a href="#">Cdc42se1.aSep08</a>	<a href="#">499672</a>	3957	740	5	79	cdc42 small effector 1 (9.0 kD) (Cdc42se1) alternative variant aSep08, mRNA.
<a href="#">Cdc42se1</a>	<a href="#">Cdc42se1.cSep08</a>	<a href="#">499672</a>	3527	736	2	76	putative protein (Cdc42se1) alternative variant cSep08, mRNA.
<a href="#">Cdc42se1</a>	<a href="#">Cdc42se1.dSep08</a>	<a href="#">499672</a>	3181	1257	3	33	cdc42 small effector 1 (3.5 kD) (Cdc42se1) alternative variant dSep08, mRNA.
<a href="#">Cdc42se2</a>	<a href="#">Cdc42se2.aSep08</a>	<a href="#">502327</a>	1293	1047		84	CDC42 small effector 2 (9.2 kD) (Cdc42se2) mRNA.
<a href="#">Cdc45l</a>	<a href="#">Cdc45l.cSep08</a>	<a href="#">287961</a>	2976	783	2	94	cell division cycle 45 homolog (S. cerevisiae)-like (Cdc45l) alternative variant cSep08, mRNA.
<a href="#">Cdc45l</a>	<a href="#">Cdc45l.dSep08</a>	<a href="#">287961</a>	807	527	2	56	cell division cycle 45 homolog (S. cerevisiae)-like (6.4 kD) (Cdc45l) alternative variant dSep08, mRNA.
<a href="#">Cdc9111</a>	<a href="#">Cdc9111.aSep08</a>	<a href="#">353304</a>	17636	777	4	160	CDC91 cell division cycle 91-like 1 (S. cerevisiae) (Cdc9111) alternative variant aSep08, mRNA.
<a href="#">Cdc123</a>	<a href="#">Cdc123.bSep08</a>	<a href="#">116656</a>	42715	1464	11	219	cell division cycle 123 (25.7 kD) (Cdc123) alternative variant bSep08, complete mRNA.
<a href="#">Cdc123</a>	<a href="#">Cdc123.cSep08</a>	<a href="#">116656</a>	10617	879	5	122	cell division cycle 123 (13.9 kD) (Cdc123) alternative variant cSep08, mRNA.
<a href="#">Cdc123</a>	<a href="#">Cdc123.dSep08</a>	<a href="#">116656</a>	812	274	2	44	cell division cycle 123 (Cdc123) alternative variant dSep08, mRNA.
<a href="#">Cdca2</a>	<a href="#">Cdca2.bSep08</a>	<a href="#">305984</a>	25579	1599	8	532	cell division cycle associated 2 (Cdca2) alternative variant bSep08, mRNA.
<a href="#">Cdca2</a>	<a href="#">Cdca2.cSep08</a>	<a href="#">305984</a>	8864	742	5	180	cell division cycle associated 2 (Cdca2) alternative variant cSep08, mRNA.
<a href="#">Cdca2</a>	<a href="#">Cdca2.dSep08</a>	<a href="#">305984</a>	9221	691	5	149	cell division cycle associated 2 (Cdca2) alternative variant dSep08, mRNA.
<a href="#">Cdca3</a>	<a href="#">Cdca3.aSep08</a>	<a href="#">297594</a>	3344	1342	4	300	cell division cycle associated 3 (Cdca3) alternative variant aSep08, mRNA.
<a href="#">Cdca3</a>	<a href="#">Cdca3.cSep08</a>	<a href="#">297594</a>	2557	744	4	128	cell division cycle associated 3 (Cdca3) alternative variant cSep08, mRNA.
<a href="#">Cdca3</a>	<a href="#">Cdca3.dSep08</a>	<a href="#">297594</a>	1381	731	1	111	cell division cycle associated 3 (Cdca3) alternative variant dSep08, mRNA.
<a href="#">Cdca4</a>	<a href="#">Cdca4.aSep08</a>	<a href="#">500727</a>	7143	1738	2	243	cell division cycle associated 4 (26.8 kD) (Cdca4) alternative variant aSep08, mRNA.
<a href="#">Cdca7</a>	<a href="#">Cdca7.bSep08</a>	<a href="#">311742</a>	7178	750	5	250	cell division cycle associated 7 (Cdca7) alternative variant bSep08, mRNA.
<a href="#">Cdca7</a>	<a href="#">Cdca7.cSep08</a>	<a href="#">311742</a>	6981	985	4	208	cell division cycle associated 7 (Cdca7) alternative variant cSep08, mRNA.

<a href="#">Cdca7l</a>	<a href="#">Cdca7l.bSep08</a>	<a href="#">619566</a>	41021	799		251	cell division cycle associated 7 like (Cdca7l) alternative variant bSep08, mRNA.
<a href="#">Cdca8</a>	<a href="#">Cdca8.aSep08</a>	<a href="#">500545</a>	25275	2208	8	180	cell division cycle associated 8 (19.7 kD) (Cdca8) alternative variant aSep08, mRNA.
<a href="#">Cdca8</a>	<a href="#">Cdca8.bSep08</a>	<a href="#">500545</a>	19721	497	6	165	cell division cycle associated 8 (Cdca8) alternative variant bSep08, mRNA.
<a href="#">Cdca8</a>	<a href="#">Cdca8.eSep08</a>	<a href="#">500545</a>	2614	929	3	125	cell division cycle associated 8 (13.4 kD) (Cdca8) alternative variant eSep08, mRNA.
<a href="#">Cdh1</a>	<a href="#">Cdh1.bSep08</a>	<a href="#">83502</a>	10957	809	1	182	cadherin 1 (Cdh1) alternative variant bSep08, mRNA.
<a href="#">Cdh2</a>	<a href="#">Cdh2.bSep08</a>	<a href="#">83501</a>	2364	545	1	89	cadherin 2 (Cdh2) alternative variant bSep08, mRNA.
<a href="#">Cdh3</a>	<a href="#">Cdh3.aSep08</a>	<a href="#">116777</a>	14332	1955		436	cadherin 3, type 1, P-cadherin (placental) (Cdh3) mRNA.
<a href="#">Cdh5</a>	<a href="#">Cdh5.bSep08</a>	<a href="#">307618</a>	12494	3139	4	514	cadherin 5 (Cdh5) alternative variant bSep08, mRNA.
<a href="#">Cdh9</a>	<a href="#">Cdh9.aSep08</a>	<a href="#">29163</a>	28235	1917		638	cadherin 9 (Cdh9) mRNA.
<a href="#">Cdh10</a>	<a href="#">Cdh10.aSep08</a>	<a href="#">29181</a>	6752	559		186	cadherin 10 (Cdh10) mRNA.
<a href="#">Cdh17</a>	<a href="#">Cdh17.bSep08</a>	<a href="#">117048</a>	3635	582	4	102	cadherin 17 (Cdh17) alternative variant bSep08, mRNA.
<a href="#">Cdh18</a>	<a href="#">Cdh18.aSep08</a>	<a href="#">310174</a>	31667	1129		77	cadherin 18, type 2 (Cdh18) mRNA.
<a href="#">Cdh19</a>	<a href="#">Cdh19.bSep08</a>	<a href="#">360835</a>	48251	1777	1	454	cadherin 19, type 2 (Cdh19) alternative variant bSep08, mRNA.
<a href="#">Cdh22</a>	<a href="#">Cdh22.bSep08</a>	<a href="#">29182</a>	4968	1424	2	262	cadherin 22 (Cdh22) alternative variant bSep08, mRNA.
<a href="#">Cdh23</a>	<a href="#">Cdh23.cSep08</a>	<a href="#">114102</a>	854	386	2	128	cadherin 23 (otocadherin) (Cdh23) alternative variant cSep08, mRNA.
<a href="#">Cdh24</a>	<a href="#">Cdh24.aSep08</a>	<a href="#">498515</a>	1951	430		139	cadherin-like 24 (Cdh24) mRNA.
<a href="#">Cdig2</a>	<a href="#">Cdig2.aSep08</a>	<a href="#">266732</a>	10622	2931	12	566	cdig2 protein (Cdig2) alternative variant aSep08, mRNA.
<a href="#">Cdipt</a>	<a href="#">Cdipt.bSep08</a>	<a href="#">192260</a>	3590	1410	5	185	CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase) (20.3 kD) (Cdipt) alternative variant bSep08, mRNA.
<a href="#">Cdipt</a>	<a href="#">Cdipt.cSep08</a>	<a href="#">192260</a>	4074	828	6	103	CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase) (11.7 kD) (Cdipt) alternative variant cSep08, complete mRNA.
<a href="#">Cdk2</a>	<a href="#">Cdk2.bSep08</a>	<a href="#">362817</a>	5086	742	5	139	cyclin dependent kinase 2 (Cdk2) alternative variant bSep08, mRNA.
<a href="#">Cdk2</a>	<a href="#">Cdk2.cSep08</a>	<a href="#">362817</a>	1168	915	2	132	cyclin dependent kinase 2 (Cdk2) alternative variant cSep08, mRNA.
<a href="#">Cdk2ap2</a>	<a href="#">Cdk2ap2.aSep08</a>	<a href="#">688405</a>	1334	429	3	128	CDK2-associated protein 2 (Cdk2ap2) alternative variant aSep08, mRNA.
<a href="#">Cdk2ap2</a>	<a href="#">Cdk2ap2.bSep08</a>	<a href="#">688405</a>	1197	830	2	128	CDK2-associated protein 2 (Cdk2ap2) alternative variant bSep08, mRNA.
<a href="#">Cdk2ap2</a>	<a href="#">Cdk2ap2.dSep08</a>	<a href="#">688405</a>	1112	400	3	60	CDK2-associated protein 2 (Cdk2ap2) alternative variant dSep08, mRNA.
<a href="#">Cdk4</a>	<a href="#">Cdk4.bSep08</a>	<a href="#">94201</a>	1142	348	3	47	cyclin-dependent kinase 4 (Cdk4) alternative variant bSep08, mRNA.
<a href="#">Cdk5</a>	<a href="#">Cdk5.bSep08</a>	<a href="#">140908</a>	2534	1524	6	191	cyclin-dependent kinase 5 (Cdk5) alternative variant bSep08, mRNA.
<a href="#">Cdk5</a>	<a href="#">Cdk5.cSep08</a>	<a href="#">140908</a>	3021	548	8	92	cyclin-dependent kinase 5 (10.5 kD) (Cdk5) alternative variant cSep08, mRNA.



<a href="#">Cdk5</a>	<a href="#">Cdk5.dSep08</a>	<a href="#">140908</a>	823	696	2	74	putative protein (7.7 kD) (Cdk5) alternative variant dSep08, mRNA.
<a href="#">Cdk5rap1</a>	<a href="#">Cdk5rap1.bSep08</a>	<a href="#">252827</a>	18678	758	6	201	CDK5 regulatory subunit associated protein 1 (Cdk5rap1) alternative variant bSep08, mRNA.
<a href="#">Cdk5rap1</a>	<a href="#">Cdk5rap1.cSep08</a>	<a href="#">252827</a>	2306	481	2	131	CDK5 regulatory subunit associated protein 1 (14.7 kD) (Cdk5rap1) alternative variant cSep08, complete mRNA.
<a href="#">Cdk5rap1</a>	<a href="#">Cdk5rap1.dSep08</a>	<a href="#">252827</a>	8360	460	3	53	CDK5 regulatory subunit associated protein 1 (Cdk5rap1) alternative variant dSep08, mRNA.
<a href="#">Cdk5rap2</a>	<a href="#">Cdk5rap2.aSep08</a>	<a href="#">286919</a>	42822	2215	9	725	CDK5 regulatory subunit associated protein 2 (Cdk5rap2) alternative variant aSep08, mRNA.
<a href="#">Cdk5rap2</a>	<a href="#">Cdk5rap2.bSep08</a>	<a href="#">286919</a>	20717	1217	4	377	CDK5 regulatory subunit associated protein 2 (Cdk5rap2) alternative variant bSep08, mRNA.
<a href="#">Cdk5rap2</a>	<a href="#">Cdk5rap2.cSep08</a>	<a href="#">286919</a>	13168	862	2	111	CDK5 regulatory subunit associated protein 2 (12.6 kD) (Cdk5rap2) alternative variant cSep08, mRNA.
<a href="#">Cdk6</a>	<a href="#">Cdk6.aSep08</a>	<a href="#">114483</a>	254613	765	2	237	cyclin-dependent kinase 6 (Cdk6) alternative variant aSep08, mRNA.
<a href="#">Cdk6</a>	<a href="#">Cdk6.bSep08</a>	<a href="#">114483</a>	253274	682	2	152	cyclin-dependent kinase 6 (Cdk6) alternative variant bSep08, mRNA.
<a href="#">Cdk7</a>	<a href="#">Cdk7.aSep08</a>	<a href="#">171150</a>	24679	1340	12	346	cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) (39.0 kD) (Cdk7) alternative variant aSep08, mRNA.
<a href="#">Cdk7</a>	<a href="#">Cdk7.bSep08</a>	<a href="#">171150</a>	24548	1775	8	78	cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) (Cdk7) alternative variant bSep08, mRNA.
<a href="#">Cdk7</a>	<a href="#">Cdk7.eSep08</a>	<a href="#">171150</a>	1212	316	2	53	cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) (5.6 kD) (Cdk7) alternative variant eSep08, mRNA.
<a href="#">Cdk9</a>	<a href="#">Cdk9.bSep08</a>	<a href="#">362110</a>	1577	1088	3	204	cyclin-dependent kinase 9 (CDC2-related kinase) (Cdk9) alternative variant bSep08, mRNA.
<a href="#">Cdk9</a>	<a href="#">Cdk9.cSep08</a>	<a href="#">362110</a>	3128	1098	5	118	cyclin-dependent kinase 9 (CDC2-related kinase) (Cdk9) alternative variant cSep08, mRNA.
<a href="#">Cdk10</a>	<a href="#">Cdk10.aSep08</a>	<a href="#">361434</a>	6587	1784	4	541	cyclin-dependent kinase (CDC2-like) 10 (Cdk10) alternative variant aSep08, mRNA.
<a href="#">Cdk10</a>	<a href="#">Cdk10.eSep08</a>	<a href="#">361434</a>	1829	699	2	90	cyclin-dependent kinase (CDC2-like) 10 (Cdk10) alternative variant eSep08, mRNA.
<a href="#">Cdk10</a>	<a href="#">Cdk10.fSep08</a>	<a href="#">361434</a>	3840	553	4	84	cyclin-dependent kinase (CDC2-like) 10 (Cdk10) alternative variant fSep08, mRNA.
<a href="#">Cdk10</a>	<a href="#">Cdk10.gSep08</a>	<a href="#">361434</a>	1942	908	2	70	cyclin-dependent kinase (CDC2-like) 10 (Cdk10) alternative variant gSep08, mRNA.
<a href="#">Cdkal1</a>	<a href="#">Cdkal1.aSep08</a>	<a href="#">361243</a>	356977	1179	8	387	CDK5 regulatory subunit associated protein 1-like 1 (Cdkal1) alternative variant aSep08, mRNA.
<a href="#">Cdkl2</a>	<a href="#">Cdkl2.bSep08</a>	<a href="#">305242</a>	14049	973	6	213	cyclin-dependent kinase-like 2 (CDC2-related kinase) (24.6 kD) (Cdkl2) alternative variant bSep08, mRNA.
<a href="#">Cdkl2</a>	<a href="#">Cdkl2.cSep08</a>	<a href="#">305242</a>	12740	2397	6	148	cyclin-dependent kinase-like 2 (CDC2-related kinase) (Cdkl2) alternative variant cSep08, mRNA.
<a href="#">Cdkl3</a>	<a href="#">Cdkl3.bSep08</a>	<a href="#">60396</a>	63454	1214	9	344	cyclin-dependent kinase-like 3 (Cdkl3) alternative variant bSep08, mRNA.

<a href="#">Cdkl3</a>	<a href="#">Cdkl3.cSep08</a>	<a href="#">60396</a>	23817	1075	6	266	cyclin-dependent kinase-like 3 (30.6 kD) (Cdkl3) alternative variant cSep08, mRNA.
<a href="#">Cdkl3</a>	<a href="#">Cdkl3.dSep08</a>	<a href="#">60396</a>	20599	718	5	191	cyclin-dependent kinase-like 3 (Cdkl3) alternative variant dSep08, mRNA.
<a href="#">Cdkl4</a>	<a href="#">Cdkl4.aSep08</a>	<a href="#">503009</a>	12358	367		114	cyclin-dependent kinase-like 4 (Cdkl4) mRNA.
<a href="#">Cdkn1a</a>	<a href="#">Cdkn1a.aSep08</a>	<a href="#">114851</a>	6269	795	3	196	cyclin-dependent kinase inhibitor 1A (p21, Cip1) (Cdkn1a) alternative variant aSep08, mRNA.
<a href="#">Cdkn1a</a>	<a href="#">Cdkn1a.bSep08</a>	<a href="#">114851</a>	1296	714	2	173	cyclin-dependent kinase inhibitor 1A (p21, Cip1) (Cdkn1a) alternative variant bSep08, mRNA.
<a href="#">Cdkn1c</a>	<a href="#">Cdkn1c.cSep08</a>	<a href="#">246060</a>	1442	1268	2	255	cyclin-dependent kinase inhibitor 1C (P57) (Cdkn1c) alternative variant cSep08, mRNA.
<a href="#">Cdkn1c</a>	<a href="#">Cdkn1c.dSep08</a>	<a href="#">246060</a>	1166	989	2	255	cyclin-dependent kinase inhibitor 1C (P57) (Cdkn1c) alternative variant dSep08, mRNA.
<a href="#">Cdkn2aip</a>	<a href="#">Cdkn2aip.cSep08</a>	<a href="#">306455</a>	3629	2574	3	164	CDKN2A interacting protein (Cdkn2aip) alternative variant cSep08, mRNA.
<a href="#">Cdkn2b</a>	<a href="#">Cdkn2b.bSep08</a>	<a href="#">25164</a>	4821	1382	3	189	cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4) (19.6 kD) (Cdkn2b) alternative variant bSep08, mRNA.
<a href="#">Cdkn2b</a>	<a href="#">Cdkn2b.cSep08</a>	<a href="#">25164</a>	5224	762	2	107	cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4) (Cdkn2b) alternative variant cSep08, mRNA.
<a href="#">Cdkn2c</a>	<a href="#">Cdkn2c.aSep08</a>	<a href="#">54238</a>	4322	1051	2	168	cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) (18.2 kD) (Cdkn2c) alternative variant aSep08, mRNA.
<a href="#">Cdkn3</a>	<a href="#">Cdkn3.aSep08</a>	<a href="#">289993</a>	11329	798	8	212	cyclin-dependent kinase inhibitor 3 (23.8 kD) (Cdkn3) alternative variant aSep08, complete mRNA.
<a href="#">Cdo1</a>	<a href="#">Cdo1.bSep08</a>	<a href="#">81718</a>	12495	753	4	188	cysteine dioxygenase 1, cytosolic (Cdo1) alternative variant bSep08, mRNA.
<a href="#">Cdr2l</a>	<a href="#">Cdr2l.aSep08</a>	<a href="#">360656</a>	21793	1802	2	495	cerebellar degeneration-related protein 2-like (Cdr2l) alternative variant aSep08, mRNA.
<a href="#">Cdr2l</a>	<a href="#">Cdr2l.bSep08</a>	<a href="#">360656</a>	4105	745	3	100	cerebellar degeneration-related protein 2-like (Cdr2l) alternative variant bSep08, mRNA.
<a href="#">Cds1</a>	<a href="#">Cds1.bSep08</a>	<a href="#">81925</a>	36633	431	5	143	CDP-diacylglycerol synthase 1 (Cds1) alternative variant bSep08, mRNA.
<a href="#">Cds1</a>	<a href="#">Cds1.cSep08</a>	<a href="#">81925</a>	6879	1210	4	112	CDP-diacylglycerol synthase 1 (12.9 kD) (Cds1) alternative variant cSep08, mRNA.
<a href="#">Cdx2</a>	<a href="#">Cdx2.bSep08</a>	<a href="#">66019</a>	6789	1575		174	caudal type homeo box 2 (Cdx2) alternative variant bSep08, mRNA.
<a href="#">Ceacam1</a>	<a href="#">Ceacam1.eSep08</a>	<a href="#">81613</a>	1925	272	3	72	CEA-related cell adhesion molecule 1 (Ceacam1) alternative variant eSep08, mRNA.
<a href="#">Ceacam1</a>	<a href="#">Ceacam1.fSep08</a>	<a href="#">81613</a>	2313	1244	2	91	CEA-related cell adhesion molecule 1 (9.6 kD) (Ceacam1) alternative variant fSep08, mRNA.
<a href="#">Ceacam6</a>	<a href="#">Ceacam6.aSep08</a>	<a href="#">100125369</a>	16341	1618	1	452	carcinoembryonic antigen-related cell adhesion molecule 6 (50.9 kD) (Ceacam6) alternative variant aSep08, mRNA.
<a href="#">Ceacam16</a>	<a href="#">Ceacam16.aSep08</a>	<a href="#">292700</a>	9896	1615		538	CEA-related cell adhesion molecule 16 (Ceacam16) mRNA.
<a href="#">Ceacam20</a>	<a href="#">Ceacam20.aSep08</a>	<a href="#">292701</a>	3744	288		77	CEA-related cell adhesion molecule 20 (Ceacam20) mRNA.
<a href="#">Cebpz</a>	<a href="#">Cebpz.bSep08</a>	<a href="#">362686</a>	3910	1017	6	197	CCAAT/enhancer binding protein zeta (22.2 kD) (Cebpz) alternative variant bSep08, mRNA.

<a href="#">Cebpz</a>	<a href="#">Cebpz.cSep08</a>	<a href="#">362686</a>	2296	435	2	58	CCAAT/enhancer binding protein zeta (7.1 kD) (Cebpz) alternative variant cSep08, mRNA.
<a href="#">Cecr2</a>	<a href="#">Cecr2.aSep08</a>	<a href="#">500308</a>	3608	741		221	putative protein of vertebrate origin (Cecr2) mRNA.
<a href="#">Cecr5</a>	<a href="#">Cecr5.bSep08</a>	<a href="#">312680</a>	39923	873	8	254	putative protein of eukaryotic origin (Cecr5) alternative variant bSep08, mRNA.
<a href="#">Cecr5</a>	<a href="#">Cecr5.cSep08</a>	<a href="#">312680</a>	22706	687	4	96	putative protein of eukaryotic origin (Cecr5) alternative variant cSep08, mRNA.
<a href="#">Cel</a>	<a href="#">Cel.bSep08</a>	<a href="#">24254</a>	3447	494	2	135	carboxyl ester lipase (Cel) alternative variant bSep08, mRNA.
<a href="#">Celsr1</a>	<a href="#">Celsr1.aSep08</a>	<a href="#">300128</a>	2479	2260		69	cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila) (Celsr1) mRNA.
<a href="#">Celsr2</a>	<a href="#">Celsr2.aSep08</a>	<a href="#">83465</a>	3687	1916	7	597	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (Celsr2) alternative variant aSep08, mRNA.
<a href="#">Celsr2</a>	<a href="#">Celsr2.bSep08</a>	<a href="#">83465</a>	5156	1785	8	554	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (Celsr2) alternative variant bSep08, mRNA.
<a href="#">Celsr2</a>	<a href="#">Celsr2.cSep08</a>	<a href="#">83465</a>	1762	1103	3	213	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (22.5 kD) (Celsr2) alternative variant cSep08, mRNA.
<a href="#">Celsr2</a>	<a href="#">Celsr2.dSep08</a>	<a href="#">83465</a>	1402	1191	1	180	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (Celsr2) alternative variant dSep08, mRNA.
<a href="#">Celsr2</a>	<a href="#">Celsr2.eSep08</a>	<a href="#">83465</a>	815	324	2	67	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (Celsr2) alternative variant eSep08, mRNA.
<a href="#">Celsr3</a>	<a href="#">Celsr3.bSep08</a>	<a href="#">83466</a>	3955	2264	2	204	cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) (Celsr3) alternative variant bSep08, mRNA.
<a href="#">Celsr3</a>	<a href="#">Celsr3.cSep08</a>	<a href="#">83466</a>	934	638	2	152	cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) (15.7 kD) (Celsr3) alternative variant cSep08, mRNA.
<a href="#">Cenpa</a>	<a href="#">Cenpa.bSep08</a>	<a href="#">298850</a>	7797	1865	3	159	centromere protein A (17.8 kD) (Cenpa) alternative variant bSep08, mRNA.
<a href="#">Cenpa</a>	<a href="#">Cenpa.cSep08</a>	<a href="#">298850</a>	6793	1761	2	125	centromere protein A (Cenpa) alternative variant cSep08, mRNA.
<a href="#">Cenpc1</a>	<a href="#">Cenpc1.bSep08</a>	<a href="#">305270</a>	1680	595	2	29	centromere protein C 1 (3.6 kD) (Cenpc1) alternative variant bSep08, mRNA.
<a href="#">Cenpe</a>	<a href="#">Cenpe.aSep08</a>	<a href="#">362044</a>	19132	1833		400	centromere protein E (45.9 kD) (Cenpe) mRNA.
<a href="#">Cenpf</a>	<a href="#">Cenpf.aSep08</a>	<a href="#">257649</a>	10930	3422		559	centromere protein F (62.4 kD) (Cenpf) mRNA.
<a href="#">Cenpi</a>	<a href="#">Cenpi.bSep08</a>	<a href="#">25448</a>	17633	890	5	209	centromere protein I (Cenpi) alternative variant bSep08, mRNA.
<a href="#">Cenpj</a>	<a href="#">Cenpj.bSep08</a>	<a href="#">305909</a>	51327	2056	9	631	centromere protein J (Cenpj) alternative variant bSep08, mRNA.
<a href="#">Cenpj</a>	<a href="#">Cenpj.cSep08</a>	<a href="#">305909</a>	6269	1086	5	138	centromere protein J CRA b (15.9 kD) (Cenpj) alternative variant cSep08, mRNA.

<a href="#">Cenpj</a>	<a href="#">Cenpj.dSep08</a>	<a href="#">305909</a>	3019	401	2	79	centromere protein J (Cenpj) alternative variant dSep08, mRNA.
<a href="#">Cenpj</a>	<a href="#">Cenpj.eSep08</a>	<a href="#">305909</a>	1008	437	2	36	putative protein (Cenpj) alternative variant eSep08, mRNA.
<a href="#">Cenpk</a>	<a href="#">Cenpk.bSep08</a>	<a href="#">294712</a>	25215	956	10	286	centromere protein K (Cenpk) alternative variant bSep08, mRNA.
<a href="#">Cenpk</a>	<a href="#">Cenpk.cSep08</a>	<a href="#">294712</a>	16852	571	7	103	centromere protein K (12.2 kD) (Cenpk) alternative variant cSep08, mRNA.
<a href="#">Cenpl</a>	<a href="#">Cenpl.bSep08</a>	<a href="#">289150</a>	14865	1625	5	236	centromere protein L (26.9 kD) (Cenpl) alternative variant bSep08, complete mRNA.
<a href="#">Cenpl</a>	<a href="#">Cenpl.cSep08</a>	<a href="#">289150</a>	5584	721	2	179	centromere protein L (Cenpl) alternative variant cSep08, mRNA.
<a href="#">Cenpl</a>	<a href="#">Cenpl.dSep08</a>	<a href="#">289150</a>	6521	789	3	124	centromere protein L (Cenpl) alternative variant dSep08, mRNA.
<a href="#">Cenpm</a>	<a href="#">Cenpm.bSep08</a>	<a href="#">315164</a>	6534	923	2	83	centromere protein M (9.1 kD) (Cenpm) alternative variant bSep08, mRNA.
<a href="#">Cenpn</a>	<a href="#">Cenpn.bSep08</a>	<a href="#">361416</a>	9219	978	7	251	centromere protein N (Cenpn) alternative variant bSep08, mRNA.
<a href="#">Cenpn</a>	<a href="#">Cenpn.cSep08</a>	<a href="#">361416</a>	12651	773	7	197	centromere protein N (Cenpn) alternative variant cSep08, mRNA.
<a href="#">Cenpn</a>	<a href="#">Cenpn.dSep08</a>	<a href="#">361416</a>	15968	733	7	191	centromere protein N (Cenpn) alternative variant dSep08, mRNA.
<a href="#">Cenpn</a>	<a href="#">Cenpn.eSep08</a>	<a href="#">361416</a>	12737	639	6	116	centromere protein N (Cenpn) alternative variant eSep08, mRNA.
<a href="#">Cenpn</a>	<a href="#">Cenpn.fSep08</a>	<a href="#">361416</a>	7831	716	4	90	centromere protein N (10.6 kD) (Cenpn) alternative variant fSep08, mRNA.
<a href="#">Cenpq</a>	<a href="#">Cenpq.aSep08</a>	<a href="#">363198</a>	14804	1145	3	224	centromere protein Q (Cenpq) alternative variant aSep08, mRNA.
<a href="#">Cenpq</a>	<a href="#">Cenpq.bSep08</a>	<a href="#">363198</a>	10814	787	2	202	centromere protein Q (23.2 kD) (Cenpq) alternative variant bSep08, mRNA.
<a href="#">Cenpt</a>	<a href="#">Cenpt.bSep08</a>	<a href="#">307805</a>	591	432	2	143	centromere protein T (Cenpt) alternative variant bSep08, mRNA.
<a href="#">Cenpt</a>	<a href="#">Cenpt.cSep08</a>	<a href="#">307805</a>	888	596	3	65	centromere protein T (Cenpt) alternative variant cSep08, mRNA.
<a href="#">Centa1</a>	<a href="#">Centa1.aSep08</a>	<a href="#">171097</a>	52627	2360	11	374	centaurin, alpha 1 (43.4 kD) (Centa1) alternative variant aSep08, mRNA.
<a href="#">Centa1</a>	<a href="#">Centa1.bSep08</a>	<a href="#">171097</a>	6654	739	6	157	centaurin, alpha 1 (Centa1) alternative variant bSep08, mRNA.
<a href="#">Centa1</a>	<a href="#">Centa1.cSep08</a>	<a href="#">171097</a>	1102	579	2	80	centaurin, alpha 1 (Centa1) alternative variant cSep08, mRNA.
<a href="#">Centb1</a>	<a href="#">Centb1.aSep08</a>	<a href="#">287443</a>	10385	1825	13	337	centaurin beta 1 CRA a (Centb1) alternative variant aSep08, mRNA.
<a href="#">Centb1</a>	<a href="#">Centb1.bSep08</a>	<a href="#">287443</a>	5063	1006	8	335	centaurin beta 1 CRA a (Centb1) alternative variant bSep08, mRNA.
<a href="#">Centb1</a>	<a href="#">Centb1.cSep08</a>	<a href="#">287443</a>	5490	839	7	218	centaurin beta 1 CRA a (Centb1) alternative variant cSep08, mRNA.
<a href="#">Centb1</a>	<a href="#">Centb1.eSep08</a>	<a href="#">287443</a>	4239	734	5	117	centaurin beta (13.3 kD) (Centb1) alternative variant eSep08, mRNA.

<a href="#">Centb2</a>	<a href="#">Centb2.bSep08</a>	<a href="#">619382</a>	16916	951	6	190	centaurin, beta 2 (22.3 kD) (Centb2) alternative variant bSep08, mRNA.
<a href="#">Centb5</a>	<a href="#">Centb5.bSep08</a>	<a href="#">313772</a>	1089	905	3	129	centaurin, beta 5 (Centb5) alternative variant bSep08, mRNA.
<a href="#">Centb5</a>	<a href="#">Centb5.cSep08</a>	<a href="#">313772</a>	5876	397	5	105	centaurin, beta 5 (Centb5) alternative variant cSep08, mRNA.
<a href="#">Centb5</a>	<a href="#">Centb5.dSep08</a>	<a href="#">313772</a>	2666	327	2	97	centaurin, beta 5 (Centb5) alternative variant dSep08, mRNA.
<a href="#">Centd1</a>	<a href="#">Centd1.bSep08</a>	<a href="#">305367</a>	45011	879	10	293	centaurin, delta 1 (Centd1) alternative variant bSep08, mRNA.
<a href="#">Centd1</a>	<a href="#">Centd1.cSep08</a>	<a href="#">305367</a>	5626	672	4	175	centaurin, delta 1 (Centd1) alternative variant cSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.aSep08</a>	<a href="#">361617</a>	65464	5486	32	943	centaurin, delta 2 (103.7 kD) (Centd2) alternative variant aSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.bSep08</a>	<a href="#">361617</a>	12516	2183	16	538	centaurin, delta 2 (Centd2) alternative variant bSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.cSep08</a>	<a href="#">361617</a>	7835	1368	7	391	centaurin, delta 2 (Centd2) alternative variant cSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.dSep08</a>	<a href="#">361617</a>	10306	1005	10	335	centaurin, delta 2 (Centd2) alternative variant dSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.eSep08</a>	<a href="#">361617</a>	4113	396	4	132	centaurin, delta 2 (Centd2) alternative variant eSep08, mRNA.
<a href="#">Centd2</a>	<a href="#">Centd2.iSep08</a>	<a href="#">361617</a>	6661	409	2	70	centaurin, delta 2 (Centd2) alternative variant iSep08, mRNA.
<a href="#">Centd3</a>	<a href="#">Centd3.aSep08</a>	<a href="#">361314</a>	16443	1942	1	516	centaurin, delta 3 (Centd3) alternative variant aSep08, mRNA.
<a href="#">Centd3</a>	<a href="#">Centd3.bSep08</a>	<a href="#">361314</a>	1689	921	4	239	centaurin, delta 3 (Centd3) alternative variant bSep08, mRNA.
<a href="#">Centd3</a>	<a href="#">Centd3.cSep08</a>	<a href="#">361314</a>	1960	1526	2	160	centaurin, delta 3 (17.6 kD) (Centd3) alternative variant cSep08, mRNA.
<a href="#">Centg1</a>	<a href="#">Centg1.aSep08</a>	<a href="#">65218</a>	2058	401		133	centaurin, gamma 1 (Centg1) mRNA.
<a href="#">Centg2</a>	<a href="#">Centg2.bSep08</a>	<a href="#">316611</a>	180519	1442	11	401	centaurin, gamma 2 (Centg2) alternative variant bSep08, mRNA.
<a href="#">Centg2</a>	<a href="#">Centg2.cSep08</a>	<a href="#">316611</a>	98909	821	5	273	centaurin, gamma 2 (Centg2) alternative variant cSep08, mRNA.
<a href="#">Centg3</a>	<a href="#">Centg3.aSep08</a>	<a href="#">362300</a>	25655	1836	14	612	centaurin gamma 3 CRA b (Centg3) alternative variant aSep08, mRNA.
<a href="#">Centg3</a>	<a href="#">Centg3.cSep08</a>	<a href="#">362300</a>	6289	1631	6	179	centaurin gamma 3 (Centg3) alternative variant cSep08, mRNA.
<a href="#">Cep57</a>	<a href="#">Cep57.bSep08</a>	<a href="#">315423</a>	9005	706	6	160	centrosomal protein 57 (Cep57) alternative variant bSep08, mRNA.
<a href="#">Cep57</a>	<a href="#">Cep57.cSep08</a>	<a href="#">315423</a>	8396	741	7	115	centrosomal protein 57 (Cep57) alternative variant cSep08, mRNA.
<a href="#">Cep57</a>	<a href="#">Cep57.dSep08</a>	<a href="#">315423</a>	2438	1416	2	93	centrosomal protein 57 (10.7 kD) (Cep57) alternative variant dSep08, mRNA.
<a href="#">Cep57</a>	<a href="#">Cep57.eSep08</a>	<a href="#">315423</a>	1503	816	2	71	centrosomal protein 57 (Cep57) alternative variant eSep08, mRNA.

<a href="#">Cep57</a>	<a href="#">Cep57.fSep08</a>	<a href="#">315423</a>	2660	193	2	36	centrosomal protein 57 (Cep57) alternative variant fSep08, mRNA.
<a href="#">Cep68</a>	<a href="#">Cep68.aSep08</a>	<a href="#">289822</a>	10103	871	5	228	centrosomal protein 68 (Cep68) alternative variant aSep08, mRNA.
<a href="#">Cep68</a>	<a href="#">Cep68.bSep08</a>	<a href="#">289822</a>	1298	676	2	211	centrosomal protein 68 (Cep68) alternative variant bSep08, mRNA.
<a href="#">Cep68</a>	<a href="#">Cep68.dSep08</a>	<a href="#">289822</a>	1504	566	2	44	centrosomal protein 68 (Cep68) alternative variant dSep08, mRNA.
<a href="#">Cep70</a>	<a href="#">Cep70.bSep08</a>	<a href="#">367153</a>	17795	746	5	141	centrosomal protein 70 (Cep70) alternative variant bSep08, mRNA.
<a href="#">Cep70</a>	<a href="#">Cep70.cSep08</a>	<a href="#">367153</a>	4197	373	4	105	centrosomal protein 70 (Cep70) alternative variant cSep08, mRNA.
<a href="#">Cep70</a>	<a href="#">Cep70.eSep08</a>	<a href="#">367153</a>	14139	749	6	51	centrosomal protein 70 (5.5 kD) (Cep70) alternative variant eSep08, mRNA.
<a href="#">Cep72</a>	<a href="#">Cep72.aSep08</a>	<a href="#">308064</a>	29911	2137	7	622	centrosomal protein 72 (Cep72) alternative variant aSep08, complete mRNA.
<a href="#">Cep72</a>	<a href="#">Cep72.bSep08</a>	<a href="#">308064</a>	12225	2454	7	376	centrosomal protein 72 (Cep72) alternative variant bSep08, mRNA.
<a href="#">Cep76</a>	<a href="#">Cep76.aSep08</a>	<a href="#">291540</a>	21270	1090	8	334	centrosomal protein 76 (Cep76) alternative variant aSep08, mRNA.
<a href="#">Cep76</a>	<a href="#">Cep76.bSep08</a>	<a href="#">291540</a>	16883	1232	8	244	centrosomal protein 76 (27.8 kD) (Cep76) alternative variant bSep08, mRNA.
<a href="#">Cep78</a>	<a href="#">Cep78.bSep08</a>	<a href="#">60347</a>	10635	1342	5	181	centrosomal protein 78 (Cep78) alternative variant bSep08, mRNA.
<a href="#">Cep110</a>	<a href="#">Cep110.bSep08</a>	<a href="#">311886</a>	8694	788	5	262	centrosomal protein 110 (Cep110) alternative variant bSep08, mRNA.
<a href="#">Cep110</a>	<a href="#">Cep110.cSep08</a>	<a href="#">311886</a>	4816	713	5	237	centrosomal protein 110 (Cep110) alternative variant cSep08, mRNA.
<a href="#">Cep110</a>	<a href="#">Cep110.dSep08</a>	<a href="#">311886</a>	4934	502	4	167	centrosomal protein 110 (Cep110) alternative variant dSep08, mRNA.
<a href="#">Cep110</a>	<a href="#">Cep110.eSep08</a>	<a href="#">311886</a>	1423	460	3	109	centrosomal protein 110 (Cep110) alternative variant eSep08, mRNA.
<a href="#">Cep135</a>	<a href="#">Cep135.aSep08</a>	<a href="#">305288</a>	20267	2070		298	centrosomal protein 135 (Cep135) alternative variant aSep08, mRNA.
<a href="#">Cep152</a>	<a href="#">Cep152.aSep08</a>	<a href="#">311391</a>	48731	1800		532	centrosomal protein 152 (Cep152) alternative variant aSep08, mRNA.
<a href="#">Cep152</a>	<a href="#">Cep152.bSep08</a>	<a href="#">311391</a>	12838	558		186	centrosomal protein 152 (Cep152) alternative variant bSep08, mRNA.
<a href="#">Cep164</a>	<a href="#">Cep164.aSep08</a>	<a href="#">363055</a>	34029	752	1	193	centrosomal protein 164 (Cep164) alternative variant aSep08, mRNA.
<a href="#">Cep164</a>	<a href="#">Cep164.bSep08</a>	<a href="#">363055</a>	26042	631	1	117	centrosomal protein 164 (Cep164) alternative variant bSep08, mRNA.
<a href="#">Cep250</a>	<a href="#">Cep250.aSep08</a>	<a href="#">311573</a>	4623	1478	3	227	centrosomal protein 250 (25.3 kD) (Cep250) alternative variant aSep08, mRNA.
<a href="#">Cep250</a>	<a href="#">Cep250.bSep08</a>	<a href="#">311573</a>	1168	703	1	173	centrosomal protein 250 (Cep250) alternative variant bSep08, mRNA.
<a href="#">Cep350</a>	<a href="#">Cep350.aSep08</a>	<a href="#">246304</a>	38018	967		322	centrosomal protein 350 (Cep350) mRNA.

<a href="#">Cept1</a>	<a href="#">Cept1.bSep08</a>	<a href="#">310773</a>	39245	1314	4	390	choline/ethanolamine phosphotransferase 1 (Cept1) alternative variant bSep08, mRNA.
<a href="#">Cercam</a>	<a href="#">Cercam.bSep08</a>	<a href="#">296616</a>	2057	938	3	115	cerebral endothelial cell adhesion molecule (11.9 kD) (Cercam) alternative variant bSep08, mRNA.
<a href="#">Cercam</a>	<a href="#">Cercam.cSep08</a>	<a href="#">296616</a>	3526	427	3	62	cerebral endothelial cell adhesion molecule (Cercam) alternative variant cSep08, mRNA.
<a href="#">Ces3</a>	<a href="#">Ces3.bSep08</a>	<a href="#">113902</a>	13986	695	4	179	carboxylesterase 3 (Ces3) alternative variant bSep08, mRNA.
<a href="#">Ces5</a>	<a href="#">Ces5.aSep08</a>	<a href="#">192257</a>	2048	586		180	carboxylesterase 5 (21.3 kD) (Ces5) mRNA.
<a href="#">Cetn2</a>	<a href="#">Cetn2.aSep08</a>	<a href="#">84593</a>	4094	558	3	185	centrin 2 (Cetn2) alternative variant aSep08, mRNA.
<a href="#">Cetn2</a>	<a href="#">Cetn2.bSep08</a>	<a href="#">84593</a>	5131	1236	4	172	centrin 2 (19.8 kD) (Cetn2) alternative variant bSep08, complete mRNA.
<a href="#">Cetn2</a>	<a href="#">Cetn2.cSep08</a>	<a href="#">84593</a>	3956	438	3	135	centrin 2 (Cetn2) alternative variant cSep08, mRNA.
<a href="#">Cetn2</a>	<a href="#">Cetn2.dSep08</a>	<a href="#">84593</a>	3894	494	3	126	centrin 2 (Cetn2) alternative variant dSep08, mRNA.
<a href="#">Cetn2</a>	<a href="#">Cetn2.eSep08</a>	<a href="#">84593</a>	2888	685	2	99	centrin 2 (11.3 kD) (Cetn2) alternative variant eSep08, mRNA.
<a href="#">Cetn3</a>	<a href="#">Cetn3.aSep08</a>	<a href="#">170895</a>	11501	870	3	174	centrin 3 (Cetn3) alternative variant aSep08, mRNA.
<a href="#">Cetn3</a>	<a href="#">Cetn3.bSep08</a>	<a href="#">170895</a>	12275	799	3	167	centrin 3 (19.5 kD) (Cetn3) alternative variant bSep08, mRNA.
<a href="#">Cetn3</a>	<a href="#">Cetn3.cSep08</a>	<a href="#">170895</a>	9691	820	2	160	centrin 3 (18.8 kD) (Cetn3) alternative variant cSep08, mRNA.
<a href="#">Cetn4</a>	<a href="#">Cetn4.aSep08</a>	<a href="#">688611</a>	1537	964		169	centrin 4 (Cetn4) mRNA.
<a href="#">Cfdp1</a>	<a href="#">Cfdp1.bSep08</a>	<a href="#">292027</a>	76976	890	6	216	craniofacial development protein 1 (Cfdp1) alternative variant bSep08, mRNA.
<a href="#">Cfdp1</a>	<a href="#">Cfdp1.cSep08</a>	<a href="#">292027</a>	79621	821	5	201	craniofacial development protein 1 (Cfdp1) alternative variant cSep08, mRNA.
<a href="#">Cfdp1</a>	<a href="#">Cfdp1.dSep08</a>	<a href="#">292027</a>	2407	367	2	48	craniofacial development protein 1 (Cfdp1) alternative variant dSep08, mRNA.
<a href="#">Cfh</a>	<a href="#">Cfh.bSep08</a>	<a href="#">155012</a>	37239	418		138	complement component factor H (Cfh) alternative variant bSep08, mRNA.
<a href="#">Cfi</a>	<a href="#">Cfi.bSep08</a>	<a href="#">79126</a>	30002	957		312	complement factor I (Cfi) alternative variant bSep08, mRNA.
<a href="#">Cfi</a>	<a href="#">Cfi.cSep08</a>	<a href="#">79126</a>	10448	733		243	complement factor I (Cfi) alternative variant cSep08, mRNA.
<a href="#">Cfi</a>	<a href="#">Cfi.dSep08</a>	<a href="#">79126</a>	8006	821		89	complement factor I (Cfi) alternative variant dSep08, mRNA.
<a href="#">Cfi</a>	<a href="#">Cfi.eSep08</a>	<a href="#">79126</a>	763	455		86	complement factor I (Cfi) alternative variant eSep08, mRNA.
<a href="#">Cfl1</a>	<a href="#">Cfl1.bSep08</a>	<a href="#">29271</a>	2861	628	3	170	cofilin 1, non-muscle (18.9 kD) (Cfl1) alternative variant bSep08, mRNA.
<a href="#">Cfl2</a>	<a href="#">Cfl2.bSep08</a>	<a href="#">366624</a>	826	741	2	94	cofilin 2, muscle (Cfl2) alternative variant bSep08, mRNA.
<a href="#">Cfl2</a>	<a href="#">Cfl2.cSep08</a>	<a href="#">366624</a>	2251	875	4	21	cofilin 2, muscle (Cfl2) alternative variant cSep08, complete mRNA.
<a href="#">Cflar</a>	<a href="#">Cflar.cSep08</a>	<a href="#">117279</a>	2652	622	2	32	CASP8 and FADD-like apoptosis regulator (3.3 kD) (Cflar) alternative variant cSep08, mRNA.

<a href="#">Cflar</a>	<a href="#">Cflar.dSep08</a>	<a href="#">117279</a>	18186	529	3	58	CASP8 and FADD-like apoptosis regulator (Cflar) alternative variant dSep08, mRNA.
<a href="#">Cfp</a>	<a href="#">Cfp.cSep08</a>	<a href="#">299314</a>	1987	482	2	40	complement factor properdin (4.4 kD) (Cfp) alternative variant cSep08, mRNA.
<a href="#">Cga</a>	<a href="#">Cga.bSep08</a>	<a href="#">116700</a>	12132	796	1	120	glycoprotein hormones, alpha subunit (13.5 kD) (Cga) alternative variant bSep08, complete mRNA.
<a href="#">Cga</a>	<a href="#">Cga.cSep08</a>	<a href="#">116700</a>	11897	703	1	117	glycoprotein hormones, alpha subunit (Cga) alternative variant cSep08, mRNA.
<a href="#">Cgi-96</a>	<a href="#">Cgi-96.bSep08</a>	<a href="#">362967</a>	5327	381		125	CGI-96 protein (Cgi-96) alternative variant bSep08, mRNA.
<a href="#">Cgm4</a>	<a href="#">Cgm4.bSep08</a>	<a href="#">24257</a>	1948	446	2	123	carcinoembryonic antigen gene family 4 (Cgm4) alternative variant bSep08, mRNA.
<a href="#">Cgn</a>	<a href="#">Cgn.aSep08</a>	<a href="#">310655</a>	7596	887		295	cingulin (Cgn) mRNA.
<a href="#">Cgnl1</a>	<a href="#">Cgnl1.bSep08</a>	<a href="#">315795</a>	48153	1564	2	491	cingulin-like 1 (Cgnl1) alternative variant bSep08, mRNA.
<a href="#">Cgnl1</a>	<a href="#">Cgnl1.cSep08</a>	<a href="#">315795</a>	4366	1532	5	203	cingulin-like 1 (Cgnl1) alternative variant cSep08, mRNA.
<a href="#">Cgrf1</a>	<a href="#">Cgrf1.bSep08</a>	<a href="#">116679</a>	11128	1043	5	290	cell growth regulator with ring finger domain 1 CRA d (Cgrf1) alternative variant bSep08, mRNA.
<a href="#">Cgrf1</a>	<a href="#">Cgrf1.cSep08</a>	<a href="#">116679</a>	8858	451	2	147	cell growth regulator with ring finger domain 1 CRA c (Cgrf1) alternative variant cSep08, mRNA.
<a href="#">Cgrf1</a>	<a href="#">Cgrf1.dSep08</a>	<a href="#">116679</a>	19763	783	5	135	cell growth regulator with ring finger domain 1 CRA d (Cgrf1) alternative variant dSep08, mRNA.
<a href="#">CH.0</a>	<a href="#">CH.0.aSep08</a>		19827	1392	7	228	calponin-like actin-binding (CH.0) alternative variant aSep08, mRNA.
<a href="#">CH.0</a>	<a href="#">CH.0.bSep08</a>		12202	397	4	132	calponin-like actin-binding (CH.0) alternative variant bSep08, mRNA.
<a href="#">CH.1</a>	<a href="#">CH.1.aSep08</a>		12261	683		227	microtubule-actin crosslinking factor 1 (CH.1) mRNA.
<a href="#">CH.2</a>	<a href="#">CH.2.aSep08</a>		12439	759		252	calponin-like actin-binding (CH.2) mRNA.
<a href="#">CH.3</a>	<a href="#">CH.3.aSep08</a>		3000	518		172	calponin-like actin-binding (CH.3) mRNA.
<a href="#">chabor</a>	<a href="#">chabor.aSep08</a>		14368	977		148	putative protein (15.9 kD) (chabor) mRNA.
<a href="#">Chac1</a>	<a href="#">Chac1.aSep08</a>	<a href="#">362196</a>	3222	1580		222	ChaC, cation transport regulator-like 1 (E. coli) (24.5 kD) (Chac1) mRNA.
<a href="#">Chac2</a>	<a href="#">Chac2.bSep08</a>	<a href="#">360994</a>	6745	395	1	131	ChaC, cation transport regulator homolog 2 (E. coli) (Chac2) alternative variant bSep08, mRNA.
<a href="#">chachy</a>	<a href="#">chachy.aSep08</a>		7681	693		128	adaptor-related protein complex (chachy) mRNA.
<a href="#">Chad</a>	<a href="#">Chad.bSep08</a>	<a href="#">29195</a>	1239	1003	1	133	chondroadherin (Chad) alternative variant bSep08, mRNA.
<a href="#">chadar</a>	<a href="#">chadar.aSep08</a>		19291	550		130	putative cytoplasmic protein (14.7 kD) (chadar) mRNA.
<a href="#">chadoy</a>	<a href="#">chadoy.aSep08</a>		30655	1413		404	protein 130kDa CRA b (44.1 kD) (chadoy) mRNA.
<a href="#">Chaf1a</a>	<a href="#">Chaf1a.aSep08</a>	<a href="#">363333</a>	13495	1147		345	chromatin assembly factor 1, subunit A (p150) (Chaf1a) mRNA.
<a href="#">Chaf1b</a>	<a href="#">Chaf1b.bSep08</a>	<a href="#">288242</a>	20730	1793	7	515	chromatin assembly factor 1, subunit B (p60) (Chaf1b) alternative variant bSep08, mRNA.
<a href="#">Chaf1b</a>	<a href="#">Chaf1b.cSep08</a>	<a href="#">288242</a>	6666	933	6	303	chromatin assembly factor 1, subunit B (p60) (Chaf1b) alternative variant cSep08, mRNA.
<a href="#">chaflu</a>	<a href="#">chaflu.aSep08</a>		13679	758		252	CRA a (chaflu) mRNA.
<a href="#">chafly</a>	<a href="#">chafly.aSep08</a>		6252	394		130	laminin alpha 2 CRA a (chafly) mRNA.
<a href="#">chagar</a>	<a href="#">chagar.aSep08</a>		3356	285		39	putative protein (4.3 kD) (chagar) mRNA.



<a href="#">chaja</a>	<a href="#">chaja.aSep08</a>		5907	638	4	42	putative protein (chaja) alternative variant aSep08, mRNA.
<a href="#">chajey</a>	<a href="#">chajey.aSep08</a>		8151	2408		218	transmembrane anterior posterior transformation 1 (chajey) mRNA.
<a href="#">chakee</a>	<a href="#">chakee.aSep08</a>		1130	676	2	24	putative protein (2.8 kD) (chakee) alternative variant aSep08, mRNA.
<a href="#">chakler</a>	<a href="#">chakler.aSep08</a>		17999	945	4	281	CRA a (chakler) alternative variant aSep08, mRNA.
<a href="#">chakler</a>	<a href="#">chakler.cSep08</a>		21889	633	4	133	CRA b (14.0 kD) (chakler) alternative variant cSep08, mRNA.
<a href="#">chalo</a>	<a href="#">chalo.aSep08</a>		6544	487		119	putative protein (chalo) mRNA.
<a href="#">chamee</a>	<a href="#">chamee.aSep08</a>		3271	1155		51	putative protein of metazoan origin (chamee) mRNA.
<a href="#">chanoy</a>	<a href="#">chanoy.aSep08</a>		3576	239		79	kinesin family member 1A CRA b (chanoy) mRNA.
<a href="#">chapor</a>	<a href="#">chapor.aSep08</a>		3174	727		82	putative mitochondrial protein (9.7 kD) (chapor) mRNA.
<a href="#">charbor</a>	<a href="#">charbor.aSep08</a>		1007	249	2	69	putative protein (charbor) alternative variant aSep08, mRNA.
<a href="#">charchy</a>	<a href="#">charchy.aSep08</a>		14212	700		98	putative protein (10.1 kD) (charchy) mRNA.
<a href="#">chardoy</a>	<a href="#">chardoy.aSep08</a>		3947	599		199	myosin VIIIb (chardoy) mRNA.
<a href="#">charflu</a>	<a href="#">charflu.aSep08</a>		14990	431		53	putative protein (charflu) mRNA.
<a href="#">charfly</a>	<a href="#">charfly.aSep08</a>		95566	479		86	putative protein (charfly) mRNA.
<a href="#">chargar</a>	<a href="#">chargar.aSep08</a>		7568	767	2	93	finger protein 42 (chargar) alternative variant aSep08, mRNA.
<a href="#">chargar</a>	<a href="#">chargar.bSep08</a>		7593	721	1	113	finger protein 42 (chargar) alternative variant bSep08, mRNA.
<a href="#">chargar</a>	<a href="#">chargar.cSep08</a>		7669	717		113	finger protein 42 (chargar) alternative variant cSep08, mRNA.
<a href="#">charja</a>	<a href="#">charja.aSep08</a>		3912	909		55	putative protein (charja) mRNA.
<a href="#">charjey</a>	<a href="#">charjey.aSep08</a>		16942	401		133	putative protein of metazoan origin (charjey) mRNA.
<a href="#">charkee</a>	<a href="#">charkee.aSep08</a>		2552	637		212	transcription factor ELYS (charkee) mRNA.
<a href="#">charkler</a>	<a href="#">charkler.aSep08</a>		13291	309		73	putative protein (charkler) mRNA.
<a href="#">charlo</a>	<a href="#">charlo.aSep08</a>		1881	342		37	putative protein (4.4 kD) (charlo) mRNA.
<a href="#">charmee</a>	<a href="#">charmee.aSep08</a>		2882	424		141	putative protein (charmee) mRNA.
<a href="#">charnoy</a>	<a href="#">charnoy.aSep08</a>		1326	698		53	cytosine deaminase like (6.0 kD) (charnoy) mRNA.
<a href="#">charoy</a>	<a href="#">charoy.bSep08</a>		720	592	2	41	putative protein (charoy) alternative variant bSep08, mRNA.
<a href="#">charpor</a>	<a href="#">charpor.aSep08</a>		5944	820		63	putative protein (6.7 kD) (charpor) mRNA.
<a href="#">charroy</a>	<a href="#">charroy.aSep08</a>		1398	450		49	putative protein (charroy) mRNA.
<a href="#">charsa</a>	<a href="#">charsa.aSep08</a>		16244	477		37	putative protein (charsa) mRNA.
<a href="#">charshee</a>	<a href="#">charshee.aSep08</a>		954	750		57	putative protein (charshee) mRNA.
<a href="#">chartu</a>	<a href="#">chartu.aSep08</a>		4867	575	3	79	putative protein (8.9 kD) (chartu) alternative variant aSep08, mRNA.
<a href="#">charvar</a>	<a href="#">charvar.aSep08</a>		1286	405		56	CRA b like (charvar) mRNA.
<a href="#">charvo</a>	<a href="#">charvo.aSep08</a>		6933	676		51	putative protein (charvo) mRNA.
<a href="#">charwer</a>	<a href="#">charwer.bSep08</a>		5039	354		48	putative protein (charwer) alternative variant bSep08, mRNA.

<a href="#">charwey</a>	<a href="#">charwey.aSep08</a>		6990	454		151	guanylate cyclase 2C (charwey) mRNA.
<a href="#">chasa</a>	<a href="#">chasa.aSep08</a>		10325	648		77	putative protein (chasa) mRNA.
<a href="#">chashee</a>	<a href="#">chashee.aSep08</a>		1363	545			
<a href="#">chatu</a>	<a href="#">chatu.aSep08</a>		42751	458		117	histone deacetylase 9 CRA a (chatu) mRNA.
<a href="#">Chat_predicted</a>	<a href="#">Chat_predicted.aSep08</a>	<a href="#">290567</a>	17862	910		303	choline acetyltransferase (predicted) (Chat_predicted) alternative variant aSep08, mRNA.
<a href="#">Chat_predicted</a>	<a href="#">Chat_predicted.bSep08</a>	<a href="#">290567</a>	6731	902		300	choline acetyltransferase (predicted) (Chat_predicted) alternative variant bSep08, mRNA.
<a href="#">Chat_predicted</a>	<a href="#">Chat_predicted.cSep08</a>	<a href="#">290567</a>	2602	350		116	choline acetyltransferase (predicted) (Chat_predicted) alternative variant cSep08, mRNA.
<a href="#">chavar</a>	<a href="#">chavar.aSep08</a>		2486	1071		78	putative protein (chavar) mRNA.
<a href="#">chavo</a>	<a href="#">chavo.aSep08</a>		41763	801		226	centrosome spindle pole associated protein 1 (chavo) mRNA.
<a href="#">chawbor</a>	<a href="#">chawbor.aSep08</a>		6335	1002	2	55	ring finger protein 8 (6.6 kD) (chawbor) alternative variant aSep08, mRNA.
<a href="#">chawchy</a>	<a href="#">chawchy.aSep08</a>		5044	523		38	putative protein (chawchy) mRNA.
<a href="#">chawdoy</a>	<a href="#">chawdoy.aSep08</a>		22144	513		134	myosin VIIb (chawdoy) mRNA.
<a href="#">chawer</a>	<a href="#">chawer.aSep08</a>		2041	418		138	RNA binding motif protein 33 like (chawer) mRNA.
<a href="#">chawey</a>	<a href="#">chawey.aSep08</a>		1352	367	1	120	putative protein (chawey) alternative variant aSep08, mRNA.
<a href="#">chawey</a>	<a href="#">chawey.bSep08</a>		1614	241	2	68	putative protein (chawey) alternative variant bSep08, mRNA.
<a href="#">chawflu</a>	<a href="#">chawflu.aSep08</a>		1874	408		107	CRA c (chawflu) mRNA.
<a href="#">chawfly</a>	<a href="#">chawfly.aSep08</a>		1355	731		50	putative protein (5.8 kD) (chawfly) mRNA.
<a href="#">chawgar</a>	<a href="#">chawgar.aSep08</a>		22790	765	1	101	putative protein (chawgar) alternative variant aSep08, mRNA.
<a href="#">chawgar</a>	<a href="#">chawgar.bSep08</a>		30586	843	2	94	putative protein (10.3 kD) (chawgar) alternative variant bSep08, mRNA.
<a href="#">chawgar</a>	<a href="#">chawgar.cSep08</a>		6675	728	2	94	putative protein (10.3 kD) (chawgar) alternative variant cSep08, mRNA.
<a href="#">chawja</a>	<a href="#">chawja.aSep08</a>		1672	289		54	putative protein (chawja) mRNA.
<a href="#">chawjey</a>	<a href="#">chawjey.aSep08</a>		8516	399		42	putative protein (chawjey) mRNA.
<a href="#">chawkee</a>	<a href="#">chawkee.aSep08</a>		3756	404		134	transcription factor (chawkee) mRNA.
<a href="#">chawkler</a>	<a href="#">chawkler.aSep08</a>		1425	977		60	putative protein (6.8 kD) (chawkler) mRNA.
<a href="#">chawlo</a>	<a href="#">chawlo.aSep08</a>		4333	346	3	47	putative protein (chawlo) mRNA.
<a href="#">chawmee</a>	<a href="#">chawmee.aSep08</a>		9278	383		127	dynein axonemal heavy (chawmee) mRNA.
<a href="#">chawnoy</a>	<a href="#">chawnoy.aSep08</a>		17161	558		44	putative protein (chawnoy) mRNA.
<a href="#">chawpor</a>	<a href="#">chawpor.aSep08</a>		4300	520		61	putative protein (chawpor) mRNA.
<a href="#">chawroy</a>	<a href="#">chawroy.aSep08</a>		14113	718		59	putative protein (chawroy) mRNA.
<a href="#">chawsa</a>	<a href="#">chawsa.bSep08</a>		2586	404	2	51	putative protein (chawsa) alternative variant bSep08, mRNA.
<a href="#">chawshee</a>	<a href="#">chawshee.aSep08</a>		6425	405		45	putative protein (4.6 kD) (chawshee) mRNA.
<a href="#">chawtu</a>	<a href="#">chawtu.aSep08</a>		42585	833		172	dedicator of cytokinesis 4 CRA b (chawtu) mRNA.
<a href="#">chawvar</a>	<a href="#">chawvar.aSep08</a>		5503	707		134	CRA a (chawvar) mRNA.

<a href="#">chawvo</a>	<a href="#">chawvo.aSep08</a>		3356	411		76	putative cytoplasmic protein (8.4 kD) (chawvo) mRNA.
<a href="#">chawwer</a>	<a href="#">chawwer.aSep08</a>		6169	429	2	79	putative protein (chawwer) alternative variant aSep08, mRNA.
<a href="#">chawwer</a>	<a href="#">chawwer.bSep08</a>		14289	468	3	51	putative protein (chawwer) alternative variant bSep08, mRNA.
<a href="#">chawwey</a>	<a href="#">chawwey.aSep08</a>		2638	444		104	CRA a like (chawwey) mRNA.
<a href="#">CHCH.0</a>	<a href="#">CHCH.0.aSep08</a>		1905	345	2	55	CHCH (CHCH.0) alternative variant aSep08, mRNA.
<a href="#">Chchd1</a>	<a href="#">Chchd1.bSep08</a>	<a href="#">361005</a>	1169	561	3	70	putative protein of bilateral origin (8.1 kD) (Chchd1) alternative variant bSep08, complete mRNA.
<a href="#">Chchd1</a>	<a href="#">Chchd1.cSep08</a>	<a href="#">361005</a>	1557	1027	2	70	putative protein of bilateral origin (8.1 kD) (Chchd1) alternative variant cSep08, mRNA.
<a href="#">Chchd3</a>	<a href="#">Chchd3.bSep08</a>	<a href="#">296966</a>	278737	967	9	236	CHCH (Chchd3) alternative variant bSep08, mRNA.
<a href="#">Chchd3</a>	<a href="#">Chchd3.cSep08</a>	<a href="#">296966</a>	269193	706	7	235	CHCH (Chchd3) alternative variant cSep08, mRNA.
<a href="#">Chchd3</a>	<a href="#">Chchd3.dSep08</a>	<a href="#">296966</a>	278590	726	7	154	CHCH (Chchd3) alternative variant dSep08, mRNA.
<a href="#">Chchd3</a>	<a href="#">Chchd3.eSep08</a>	<a href="#">296966</a>	205780	654	5	132	CHCH (Chchd3) alternative variant eSep08, mRNA.
<a href="#">Chchd5</a>	<a href="#">Chchd5.aSep08</a>	<a href="#">296147</a>	44580	935	4	157	CHCH (Chchd5) alternative variant aSep08, mRNA.
<a href="#">Chchd5</a>	<a href="#">Chchd5.bSep08</a>	<a href="#">296147</a>	1743	685	4	140	CHCH (15.7 kD) (Chchd5) alternative variant bSep08, mRNA.
<a href="#">Chchd5</a>	<a href="#">Chchd5.dSep08</a>	<a href="#">296147</a>	4494	1617	4	110	CHCH (12.3 kD) (Chchd5) alternative variant dSep08, mRNA.
<a href="#">Chchd6</a>	<a href="#">Chchd6.bSep08</a>	<a href="#">297436</a>	87619	555	1	60	putative protein of vertebrate origin (Chchd6) alternative variant bSep08, mRNA.
<a href="#">Chchd8</a>	<a href="#">Chchd8.aSep08</a>	<a href="#">499214</a>	2362	887	2	111	CHCH (12.9 kD) (Chchd8) alternative variant aSep08, mRNA.
<a href="#">Chd1l</a>	<a href="#">Chd1l.aSep08</a>	<a href="#">310707</a>	37477	2263	14	605	chromodomain helicase DNA binding protein 1-like (67.7 kD) (Chd1l) alternative variant aSep08, mRNA.
<a href="#">Chd1l</a>	<a href="#">Chd1l.bSep08</a>	<a href="#">310707</a>	14149	945	7	314	chromodomain helicase DNA binding protein 1-like (Chd1l) alternative variant bSep08, mRNA.
<a href="#">Chd2</a>	<a href="#">Chd2.bSep08</a>	<a href="#">308738</a>	11015	1338	5	242	chromodomain helicase DNA binding protein 2 (Chd2) alternative variant bSep08, mRNA.
<a href="#">Chd2</a>	<a href="#">Chd2.cSep08</a>	<a href="#">308738</a>	8347	2403	3	68	chromodomain helicase DNA binding protein 2 (Chd2) alternative variant cSep08, mRNA.
<a href="#">Chd3</a>	<a href="#">Chd3.aSep08</a>	<a href="#">303241</a>	12940	4255	19	1079	chromodomain helicase DNA binding protein 3 (Chd3) alternative variant aSep08, mRNA.
<a href="#">Chd3</a>	<a href="#">Chd3.bSep08</a>	<a href="#">303241</a>	6736	2287	7	580	chromodomain helicase DNA binding protein 3 (Chd3) alternative variant bSep08, mRNA.
<a href="#">Chd3</a>	<a href="#">Chd3.cSep08</a>	<a href="#">303241</a>	3563	888	2	266	chromodomain helicase DNA binding protein 3 (Chd3) alternative variant cSep08, mRNA.
<a href="#">Chd3</a>	<a href="#">Chd3.dSep08</a>	<a href="#">303241</a>	4503	2204	2	172	chromodomain helicase DNA binding protein 3 (19.4 kD) (Chd3) alternative variant dSep08, mRNA.
<a href="#">Chd4</a>	<a href="#">Chd4.aSep08</a>	<a href="#">117535</a>	20858	3989	25	1131	chromodomain helicase DNA binding protein 4 (Chd4) alternative variant aSep08, mRNA.
<a href="#">Chd4</a>	<a href="#">Chd4.bSep08</a>	<a href="#">117535</a>	7120	1232	9	410	chromodomain helicase DNA binding protein 4 (Chd4) alternative variant bSep08, mRNA.
<a href="#">Chd4</a>	<a href="#">Chd4.cSep08</a>	<a href="#">117535</a>	6026	870	4	182	chromodomain helicase DNA binding protein 4 (Chd4) alternative variant cSep08, mRNA.

<a href="#">Chd4</a>	<a href="#">Chd4.dSep08</a>	<a href="#">117535</a>	7274	914	8	160	chromodomain helicase DNA binding protein 4 (Chd4) alternative variant dSep08, mRNA.
<a href="#">Chd5</a>	<a href="#">Chd5.aSep08</a>	<a href="#">691589</a>	15407	269		89	chromodomain helicase DNA binding protein 5 (Chd5) mRNA.
<a href="#">Chd6</a>	<a href="#">Chd6.bSep08</a>	<a href="#">311607</a>	408	269	2	89	chromodomain helicase DNA binding protein 6 (Chd6) alternative variant bSep08, mRNA.
<a href="#">Chd6</a>	<a href="#">Chd6.cSep08</a>	<a href="#">311607</a>	3961	360	2	75	chromodomain helicase DNA binding protein 6 (Chd6) alternative variant cSep08, mRNA.
<a href="#">Chd6</a>	<a href="#">Chd6.fSep08</a>	<a href="#">311607</a>	9799	503	2	44	chromodomain helicase DNA binding protein 6 (Chd6) alternative variant fSep08, mRNA.
<a href="#">Chd8</a>	<a href="#">Chd8.cSep08</a>	<a href="#">65027</a>	3853	1130	2	230	chromodomain helicase DNA binding protein 8 (Chd8) alternative variant cSep08, mRNA.
<a href="#">CHDNT.0</a>	<a href="#">CHDNT.0.aSep08</a>		3059	789	1	262	chromodomain helicase DNA binding protein 3 like (CHDNT.0) alternative variant aSep08, mRNA.
<a href="#">CHDNT.0</a>	<a href="#">CHDNT.0.bSep08</a>		2319	426	1	141	chromodomain helicase DNA binding protein 3 like (CHDNT.0) alternative variant bSep08, mRNA.
<a href="#">CHDNT.1</a>	<a href="#">CHDNT.1.aSep08</a>		10085	2128	9	655	chromodomain helicase DNA binding protein 4 like (CHDNT.1) alternative variant aSep08, mRNA.
<a href="#">CHDNT.1</a>	<a href="#">CHDNT.1.bSep08</a>		5187	626	1	160	chromodomain helicase DNA binding protein 4 CRA a like (CHDNT.1) alternative variant bSep08, mRNA.
<a href="#">cheebor</a>	<a href="#">cheebor.aSep08</a>		6419	487		132	putative protein (cheebor) mRNA.
<a href="#">cheechy</a>	<a href="#">cheechy.aSep08</a>		4398	501		84	putative protein (cheechy) mRNA.
<a href="#">cheedoy</a>	<a href="#">cheedoy.aSep08</a>		9547	768		48	putative protein (5.2 kD) (cheedoy) mRNA.
<a href="#">cheeflu</a>	<a href="#">cheeflu.aSep08</a>		17733	513		87	caseinolytic peptidase B like (cheeflu) mRNA.
<a href="#">cheefly</a>	<a href="#">cheefly.aSep08</a>		3671	406		108	mediator SUR2 (cheefly) mRNA.
<a href="#">cheegar</a>	<a href="#">cheegar.aSep08</a>		5081	576		191	pericentriolar material 1 (cheegar) mRNA.
<a href="#">cheeja</a>	<a href="#">cheeja.aSep08</a>		5375	736	2	116	putative protein (12.7 kD) (cheeja) alternative variant aSep08, mRNA.
<a href="#">cheejey</a>	<a href="#">cheejey.aSep08</a>		8746	539		64	putative protein (cheejey) mRNA.
<a href="#">cheekee</a>	<a href="#">cheekee.aSep08</a>		3770	348		115	putative protein (cheekee) mRNA.
<a href="#">cheekler</a>	<a href="#">cheekler.aSep08</a>		485	362		10	putative protein (1.1 kD) (cheekler) mRNA.
<a href="#">cheelo</a>	<a href="#">cheelo.aSep08</a>		4863	364		69	putative protein (8.2 kD) (cheelo) mRNA.
<a href="#">cheemee</a>	<a href="#">cheemee.aSep08</a>		3296	356		74	polyprotein (cheemee) mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.aSep08</a>		36908	1652	6	359	histidine acid phosphatase and phosphoglycerate mutase (cheenoy) alternative variant aSep08, mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.bSep08</a>		34818	1099	7	308	putative protein of eukaryotic origin (cheenoy) alternative variant bSep08, mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.cSep08</a>		15939	755	3	169	putative protein (cheenoy) alternative variant cSep08, mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.dSep08</a>		6145	463	3	153	putative protein of eukaryotic origin (cheenoy) alternative variant dSep08, mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.eSep08</a>		6629	1855	1	99	putative protein of mammalian origin (cheenoy) alternative variant eSep08, mRNA.
<a href="#">cheenoy</a>	<a href="#">cheenoy.fSep08</a>		43116	338	2	70	putative protein of vertebrate origin (cheenoy) alternative variant fSep08, mRNA.

<a href="#">cheepor</a>	<a href="#">cheepor.aSep08</a>		19343	784		51	putative protein (5.7 kD) (cheepor) mRNA.
<a href="#">cheeroy</a>	<a href="#">cheeroy.aSep08</a>		4788	380	4	126	zinc finger protein subfamily 1A 4 (cheeroy) alternative variant aSep08, mRNA.
<a href="#">cheeroy</a>	<a href="#">cheeroy.bSep08</a>		21737	630	4	40	CRA c like (4.5 kD) (cheeroy) alternative variant bSep08, mRNA.
<a href="#">cheeroy</a>	<a href="#">cheeroy.dSep08</a>		27579	420	3	62	putative protein (cheeroy) alternative variant dSep08, mRNA.
<a href="#">cheesa</a>	<a href="#">cheesa.aSep08</a>		18443	657	2	64	putative protein (cheesa) alternative variant aSep08, mRNA.
<a href="#">cheesa</a>	<a href="#">cheesa.bSep08</a>		3267	460	1	47	putative protein (cheesa) alternative variant bSep08, mRNA.
<a href="#">cheeshee</a>	<a href="#">cheeshee.aSep08</a>		10858	461	2	61	putative protein (cheeshee) alternative variant aSep08, mRNA.
<a href="#">cheeshee</a>	<a href="#">cheeshee.bSep08</a>		67643	376	1	30	putative protein (3.3 kD) (cheeshee) alternative variant bSep08, mRNA.
<a href="#">cheetu</a>	<a href="#">cheetu.aSep08</a>		12619	3031		263	dedicator of cytokinesis 4 CRA e (cheetu) mRNA.
<a href="#">cheevar</a>	<a href="#">cheevar.aSep08</a>		6669	767	2	84	putative cytoplasmic protein (9.3 kD) (cheevar) mRNA.
<a href="#">cheevo</a>	<a href="#">cheevo.aSep08</a>		53387	406		108	regulator of G-protein signaling 20 like (cheevo) mRNA.
<a href="#">cheewer</a>	<a href="#">cheewer.aSep08</a>		8839	261		86	mlI3 (cheewer) mRNA.
<a href="#">cheewey</a>	<a href="#">cheewey.aSep08</a>		4652	708		83	putative protein (cheewey) mRNA.
<a href="#">Chek1</a>	<a href="#">Chek1.bSep08</a>	<a href="#">140583</a>	13009	497	4	165	checkpoint kinase 1 homolog (S. pombe) (Chek1) alternative variant bSep08, mRNA.
<a href="#">Chek1</a>	<a href="#">Chek1.cSep08</a>	<a href="#">140583</a>	9835	651	3	143	checkpoint kinase 1 homolog (S. pombe) (Chek1) alternative variant cSep08, mRNA.
<a href="#">Chek2</a>	<a href="#">Chek2.bSep08</a>	<a href="#">114212</a>	7981	723	4	204	CHK2 checkpoint homolog (S. pombe) (Chek2) alternative variant bSep08, mRNA.
<a href="#">Chek2</a>	<a href="#">Chek2.cSep08</a>	<a href="#">114212</a>	8419	688	4	204	CHK2 checkpoint homolog (S. pombe) (Chek2) alternative variant cSep08, mRNA.
<a href="#">Chek2</a>	<a href="#">Chek2.eSep08</a>	<a href="#">114212</a>	2368	1255	2	52	CHK2 checkpoint homolog (S. pombe) (5.8 kD) (Chek2) alternative variant eSep08, mRNA.
<a href="#">cherbor</a>	<a href="#">cherbor.aSep08</a>		7598	446		48	putative protein (cherbor) mRNA.
<a href="#">cherchy</a>	<a href="#">cherchy.aSep08</a>		1374	672		44	putative protein (4.8 kD) (cherchy) mRNA.
<a href="#">cherdoy</a>	<a href="#">cherdoy.aSep08</a>		33523	402		51	putative protein (5.5 kD) (cherdoy) mRNA.
<a href="#">cherflu</a>	<a href="#">cherflu.aSep08</a>		8757	344		109	nuclear mitotic apparatus protein 1 (cherflu) mRNA.
<a href="#">cherfly</a>	<a href="#">cherfly.aSep08</a>		763	329		73	putative mitochondrial protein (7.8 kD) (cherfly) mRNA.
<a href="#">chergar</a>	<a href="#">chergar.aSep08</a>		6251	516		171	pericentriolar material 1 (chergar) mRNA.
<a href="#">cherja</a>	<a href="#">cherja.aSep08</a>		17973	799		44	putative protein (cherja) mRNA.
<a href="#">cherjey</a>	<a href="#">cherjey.aSep08</a>		94467	1703		56	putative protein (5.9 kD) (cherjey) mRNA.
<a href="#">cherkee</a>	<a href="#">cherkee.aSep08</a>		1763	671		39	putative protein (cherkee) mRNA.
<a href="#">cherkler</a>	<a href="#">cherkler.aSep08</a>		2143	805	1	124	putative protein, with a coiled coil domain, of mammalian origin (cherkler) alternative variant aSep08, mRNA.
<a href="#">cherkler</a>	<a href="#">cherkler.bSep08</a>		2123	667	2	117	putative protein, with a coiled coil domain, of mammalian origin (cherkler) alternative variant bSep08, mRNA.
<a href="#">cherlo</a>	<a href="#">cherlo.aSep08</a>		5041	312		61	putative protein (cherlo) mRNA.
<a href="#">chermee</a>	<a href="#">chermee.aSep08</a>		5788	503		56	putative protein (chermee) mRNA.

<a href="#">chernoy</a>	<a href="#">chernoy.aSep08</a>		3469	681		70	putative protein (chernoy) mRNA.
<a href="#">Cherp</a>	<a href="#">Cherp.bSep08</a>	<a href="#">290614</a>	1914	1533	4	240	calcium homeostasis endoplasmic reticulum protein CRA a (25.8 kD) (Cherp) alternative variant bSep08, mRNA.
<a href="#">Cherp</a>	<a href="#">Cherp.cSep08</a>	<a href="#">290614</a>	3862	2245	4	192	calcium homeostasis endoplasmic reticulum protein (Cherp) alternative variant cSep08, mRNA.
<a href="#">Cherp</a>	<a href="#">Cherp.dSep08</a>	<a href="#">290614</a>	1168	906	2	133	calcium homeostasis endoplasmic reticulum protein CRA a (Cherp) alternative variant dSep08, mRNA.
<a href="#">Cherp</a>	<a href="#">Cherp.eSep08</a>	<a href="#">290614</a>	922	694	3	80	putative protein (Cherp) alternative variant eSep08, mRNA.
<a href="#">cherpor</a>	<a href="#">cherpor.aSep08</a>		5740	232	1	76	putative protein (cherpor) alternative variant aSep08, mRNA.
<a href="#">cherpor</a>	<a href="#">cherpor.bSep08</a>		62287	583	3	46	putative protein (5.4 kD) (cherpor) alternative variant bSep08, mRNA.
<a href="#">cherroy</a>	<a href="#">cherroy.aSep08</a>		9153	590		27	putative protein (2.9 kD) (cherroy) alternative variant aSep08, mRNA.
<a href="#">cherroy</a>	<a href="#">cherroy.bSep08</a>		1847	291		26	putative protein (cherroy) alternative variant bSep08, mRNA.
<a href="#">chersa</a>	<a href="#">chersa.aSep08</a>		3868	1233	5	101	putative protein (chersa) alternative variant aSep08, mRNA.
<a href="#">chersa</a>	<a href="#">chersa.bSep08</a>		1060	385	3	62	putative protein (chersa) alternative variant bSep08, mRNA.
<a href="#">chershee</a>	<a href="#">chershee.aSep08</a>		3175	650		72	putative protein (8.1 kD) (chershee) mRNA.
<a href="#">chertu</a>	<a href="#">chertu.aSep08</a>		110987	421	2	140	inner mitochondrial membrane -like (chertu) alternative variant aSep08, mRNA.
<a href="#">chertu</a>	<a href="#">chertu.bSep08</a>		57413	382	1	100	putative protein (chertu) alternative variant bSep08, mRNA.
<a href="#">chervar</a>	<a href="#">chervar.aSep08</a>		4250	478	3	94	putative protein of eukaryotic origin (chervar) alternative variant aSep08, mRNA.
<a href="#">chervo</a>	<a href="#">chervo.aSep08</a>		2560	591		49	putative protein (chervo) mRNA.
<a href="#">cherwer</a>	<a href="#">cherwer.aSep08</a>		4695	330		109	mixed-lineage leukemia 3 like (cherwer) mRNA.
<a href="#">cherwey</a>	<a href="#">cherwey.aSep08</a>		2262	359		41	putative protein (cherwey) mRNA.
<a href="#">cheybor</a>	<a href="#">cheybor.aSep08</a>		1364	596		42	putative protein (4.3 kD) (cheybor) mRNA.
<a href="#">cheychy</a>	<a href="#">cheychy.aSep08</a>		57777	552	1	40	putative protein (4.4 kD) (cheychy) alternative variant aSep08, mRNA.
<a href="#">cheychy</a>	<a href="#">cheychy.bSep08</a>		57827	475	2	40	putative protein (4.4 kD) (cheychy) alternative variant bSep08, mRNA.
<a href="#">cheydoy</a>	<a href="#">cheydoy.aSep08</a>		2132	748		45	putative protein (cheydoy) mRNA.
<a href="#">cheyflu</a>	<a href="#">cheyflu.aSep08</a>		2973	606		202	nuclear mitotic apparatus protein 1 (cheyflu) mRNA.
<a href="#">cheyfly</a>	<a href="#">cheyfly.aSep08</a>		3666	751		65	putative protein (7.0 kD) (cheyfly) mRNA.
<a href="#">cheygar</a>	<a href="#">cheygar.aSep08</a>		3476	380		126	pericentriolar material 1 (cheygar) mRNA.
<a href="#">cheyja</a>	<a href="#">cheyja.aSep08</a>		1130	278		50	putative protein (4.4 kD) (cheyja) mRNA.
<a href="#">cheyjey</a>	<a href="#">cheyjey.aSep08</a>		3240	770		52	putative protein (6.2 kD) (cheyjey) mRNA.
<a href="#">cheykee</a>	<a href="#">cheykee.aSep08</a>		1092	612	1	122	putative protein (13.8 kD) (cheykee) alternative variant aSep08, mRNA.
<a href="#">cheykee</a>	<a href="#">cheykee.bSep08</a>		812	329	1	75	putative protein (cheykee) alternative variant bSep08, mRNA.
<a href="#">cheykler</a>	<a href="#">cheykler.aSep08</a>		3513	329		40	putative protein (cheykler) mRNA.

<a href="#">chevlo</a>	<a href="#">chevlo.aSep08</a>		5112	662		86	putative mitochondrial protein (9.1 kD) (chevlo) mRNA.
<a href="#">cheymee</a>	<a href="#">cheymee.aSep08</a>		3137	1081		88	putative protein (9.8 kD) (cheymee) mRNA.
<a href="#">cheynoy</a>	<a href="#">cheynoy.aSep08</a>		16998	255		36	putative protein (3.9 kD) (cheynoy) mRNA.
<a href="#">cheypor</a>	<a href="#">cheypor.aSep08</a>		12749	911		173	putative protein (cheypor) mRNA.
<a href="#">cheyro</a>	<a href="#">cheyro.aSep08</a>		197	123		40	putative protein (cheyro) mRNA.
<a href="#">cheysa</a>	<a href="#">cheysa.aSep08</a>		727	336		112	histone deacetylase 7A CRA b (cheysa) mRNA.
<a href="#">cheyshee</a>	<a href="#">cheyshee.bSep08</a>		36053	699		35	putative protein (4.0 kD) (cheyshee) alternative variant bSep08, mRNA.
<a href="#">cheytu</a>	<a href="#">cheytu.aSep08</a>		29908	467		72	putative cytoplasmic protein (7.5 kD) (cheytu) mRNA.
<a href="#">cheyvar</a>	<a href="#">cheyvar.aSep08</a>		3551	789		262	putative protein of metazoan origin (cheyvar) mRNA.
<a href="#">cheyvo</a>	<a href="#">cheyvo.aSep08</a>		11972	836		84	putative protein (cheyvo) mRNA.
<a href="#">cheywer</a>	<a href="#">cheywer.aSep08</a>		7082	868		61	mixed-lineage leukemia 3 like (cheywer) mRNA.
<a href="#">cheywey</a>	<a href="#">cheywey.aSep08</a>		42140	206		39	putative protein (cheywey) mRNA.
<a href="#">Chfr</a>	<a href="#">Chfr.bSep08</a>	<a href="#">288734</a>	6653	757	5	232	checkpoint with forkhead and ring finger domains (Chfr) alternative variant bSep08, mRNA.
<a href="#">Chga</a>	<a href="#">Chga.bSep08</a>	<a href="#">24258</a>	1228	464	1	133	chromogranin A (Chga) alternative variant bSep08, mRNA.
<a href="#">Chi311</a>	<a href="#">Chi311.bSep08</a>	<a href="#">89824</a>	4659	767	5	176	chitinase 3-like 1 (Chi311) alternative variant bSep08, mRNA.
<a href="#">Chi311</a>	<a href="#">Chi311.cSep08</a>	<a href="#">89824</a>	4382	519	6	172	chitinase 3-like 1 (Chi311) alternative variant cSep08, mRNA.
<a href="#">Chia</a>	<a href="#">Chia.bSep08</a>	<a href="#">113901</a>	9774	618		172	chitinase, acidic (Chia) alternative variant bSep08, mRNA.
<a href="#">Chic1</a>	<a href="#">Chic1.aSep08</a>	<a href="#">363484</a>	19584	392	2	130	cysteine-rich hydrophobic domain 1 (Chic1) alternative variant aSep08, mRNA.
<a href="#">Chic1</a>	<a href="#">Chic1.bSep08</a>	<a href="#">363484</a>	5597	382	1	64	cysteine-rich hydrophobic domain 1 (Chic1) alternative variant bSep08, mRNA.
<a href="#">Chid1</a>	<a href="#">Chid1.bSep08</a>	<a href="#">293628</a>	5572	719	5	66	putative protein (6.6 kD) (Chid1) alternative variant bSep08, mRNA.
<a href="#">Chid1</a>	<a href="#">Chid1.dSep08</a>	<a href="#">293628</a>	1139	612	3	86	putative protein (Chid1) alternative variant dSep08, mRNA.
<a href="#">Chit1</a>	<a href="#">Chit1.aSep08</a>	<a href="#">289032</a>	20420	2177	1	464	chitinase 1 (chitotriosidase) (50.8 kD) (Chit1) alternative variant aSep08, complete mRNA.
<a href="#">Chit1</a>	<a href="#">Chit1.bSep08</a>	<a href="#">289032</a>	20429	1835	2	464	chitinase 1 (chitotriosidase) (50.8 kD) (Chit1) alternative variant bSep08, complete mRNA.
<a href="#">Chit1</a>	<a href="#">Chit1.cSep08</a>	<a href="#">289032</a>	20430	1735	1	464	chitinase 1 (chitotriosidase) (50.8 kD) (Chit1) alternative variant cSep08, complete mRNA.
<a href="#">Chka</a>	<a href="#">Chka.bSep08</a>	<a href="#">29194</a>	41865	1103	6	278	choline kinase alpha (Chka) alternative variant bSep08, mRNA.
<a href="#">Chka</a>	<a href="#">Chka.cSep08</a>	<a href="#">29194</a>	17062	680	7	226	choline kinase alpha (Chka) alternative variant cSep08, mRNA.
<a href="#">Chkb</a>	<a href="#">Chkb.bSep08</a>	<a href="#">29367</a>	2534	1608	7	258	choline kinase beta (29.8 kD) (Chkb) alternative variant bSep08, mRNA.
<a href="#">Chkb</a>	<a href="#">Chkb.cSep08</a>	<a href="#">29367</a>	2399	1247	8	198	choline kinase beta (Chkb) alternative variant cSep08, mRNA.
<a href="#">Chkb</a>	<a href="#">Chkb.dSep08</a>	<a href="#">29367</a>	1216	779	3	160	choline kinase beta (Chkb) alternative variant dSep08, mRNA.

<a href="#">Chkb</a>	<a href="#">Chkb.eSep08</a>	<a href="#">29367</a>	833	475	3	109	choline kinase beta (Chkb) alternative variant eSep08, mRNA.
<a href="#">Chm</a>	<a href="#">Chm.bSep08</a>	<a href="#">24942</a>	45632	912	4	111	choroideremia (Chm) alternative variant bSep08, mRNA.
<a href="#">Chmp1a</a>	<a href="#">Chmp1a.bSep08</a>	<a href="#">365024</a>	7144	881	5	170	chromatin modifying protein 1A (18.7 kD) (Chmp1a) alternative variant bSep08, mRNA.
<a href="#">Chmp1a</a>	<a href="#">Chmp1a.cSep08</a>	<a href="#">365024</a>	2769	406	1	69	chromatin modifying protein 1A (Chmp1a) alternative variant cSep08, mRNA.
<a href="#">Chmp2a</a>	<a href="#">Chmp2a.aSep08</a>	<a href="#">365191</a>	2498	1970	5	418	chromatin modifying protein 2A (Chmp2a) alternative variant aSep08, complete mRNA.
<a href="#">Chmp2a</a>	<a href="#">Chmp2a.bSep08</a>	<a href="#">365191</a>	2501	900	6	222	chromatin modifying protein 2A (25.1 kD) (Chmp2a) alternative variant bSep08, complete mRNA.
<a href="#">Chmp2a</a>	<a href="#">Chmp2a.dSep08</a>	<a href="#">365191</a>	2472	721	6	172	chromatin modifying protein 2A (19.4 kD) (Chmp2a) alternative variant dSep08, mRNA.
<a href="#">Chmp2b</a>	<a href="#">Chmp2b.aSep08</a>	<a href="#">363720</a>	40890	533		177	chromatin modifying protein 2B (Chmp2b) alternative variant aSep08, mRNA.
<a href="#">Chmp2b</a>	<a href="#">Chmp2b.bSep08</a>	<a href="#">363720</a>	6983	1849		107	chromatin modifying protein 2B (Chmp2b) alternative variant bSep08, mRNA.
<a href="#">Chmp6</a>	<a href="#">Chmp6.aSep08</a>	<a href="#">287873</a>	5664	1496	4	240	chromatin modifying protein 6 (27.7 kD) (Chmp6) alternative variant aSep08, complete mRNA.
<a href="#">Chmp6</a>	<a href="#">Chmp6.bSep08</a>	<a href="#">287873</a>	5298	1185	5	200	chromatin modifying protein 6 (23.3 kD) (Chmp6) alternative variant bSep08, mRNA.
<a href="#">Chmp6</a>	<a href="#">Chmp6.cSep08</a>	<a href="#">287873</a>	5340	687	5	114	chromatin modifying protein 6 (12.8 kD) (Chmp6) alternative variant cSep08, mRNA.
<a href="#">Chmp6</a>	<a href="#">Chmp6.dSep08</a>	<a href="#">287873</a>	2298	733	2	112	chromatin modifying protein 6 (Chmp6) alternative variant dSep08, mRNA.
<a href="#">Chmp6</a>	<a href="#">Chmp6.eSep08</a>	<a href="#">287873</a>	2428	778	2	83	chromatin modifying protein 6 (Chmp6) alternative variant eSep08, mRNA.
<a href="#">Chmp7</a>	<a href="#">Chmp7.bSep08</a>	<a href="#">364419</a>	614	389	2	57	CHMP family, member 7 (Chmp7) alternative variant bSep08, mRNA.
<a href="#">Chn1</a>	<a href="#">Chn1.aSep08</a>	<a href="#">84030</a>	224797	2184	14	455	chimerin (chimaerin) 1 (52.8 kD) (Chn1) alternative variant aSep08, mRNA.
<a href="#">Chn1</a>	<a href="#">Chn1.cSep08</a>	<a href="#">84030</a>	41102	1053	7	258	chimerin (chimaerin) 1 (Chn1) alternative variant cSep08, mRNA.
<a href="#">Chn1</a>	<a href="#">Chn1.dSep08</a>	<a href="#">84030</a>	5375	1051	3	128	chimerin (chimaerin) 1 (14.8 kD) (Chn1) alternative variant dSep08, mRNA.
<a href="#">Chn2</a>	<a href="#">Chn2.aSep08</a>	<a href="#">84031</a>	108373	2942	12	458	chimerin (chimaerin) 2 (Chn2) alternative variant aSep08, mRNA.
<a href="#">Chn2</a>	<a href="#">Chn2.cSep08</a>	<a href="#">84031</a>	24692	839	5	240	chimerin (chimaerin) 2 (Chn2) alternative variant cSep08, mRNA.
<a href="#">Chn2</a>	<a href="#">Chn2.dSep08</a>	<a href="#">84031</a>	98331	719	7	217	chimerin (chimaerin) 2 (Chn2) alternative variant dSep08, mRNA.
<a href="#">Chn2</a>	<a href="#">Chn2.eSep08</a>	<a href="#">84031</a>	194264	583	6	194	chimerin (chimaerin) 2 (Chn2) alternative variant eSep08, mRNA.
<a href="#">chobor</a>	<a href="#">chobor.aSep08</a>		3532	529		62	putative protein (7.1 kD) (chobor) mRNA.
<a href="#">chochy</a>	<a href="#">chochy.aSep08</a>		756	460		32	putative protein (3.8 kD) (chochy) mRNA.
<a href="#">Chodl</a>	<a href="#">Chodl.bSep08</a>	<a href="#">288289</a>	6928	1577	2	68	chondrolectin (Chodl) alternative variant bSep08, mRNA.



<a href="#">chodoy</a>	<a href="#">chodoy.aSep08</a>		11750	635		211	myosin VIIb (chodoy) mRNA.
<a href="#">choflu</a>	<a href="#">choflu.aSep08</a>		1413	348	5	115	rho guanine nucleotide exchange factor 17 (choflu) alternative variant aSep08, mRNA.
<a href="#">chofly</a>	<a href="#">chofly.aSep08</a>		5682	429		36	putative protein (chofly) mRNA.
<a href="#">chogar</a>	<a href="#">chogar.aSep08</a>		5597	1776		518	CRA b (chogar) alternative variant aSep08, mRNA.
<a href="#">choja</a>	<a href="#">choja.aSep08</a>		1511	286		67	putative protein (7.2 kD) (choja) alternative variant aSep08, mRNA.
<a href="#">chojey</a>	<a href="#">chojey.aSep08</a>		27410	1643		521	putative protein of bilateral origin (chojey) mRNA.
<a href="#">chokee</a>	<a href="#">chokee.aSep08</a>		3870	598		198	AT hook containing transcription factor 1 (chokee) mRNA.
<a href="#">chokler</a>	<a href="#">chokler.aSep08</a>		7374	419		36	putative protein (4.1 kD) (chokler) mRNA.
<a href="#">cholo</a>	<a href="#">cholo.aSep08</a>		1135	304		45	putative protein (4.9 kD) (cholo) mRNA.
<a href="#">chomee</a>	<a href="#">chomee.aSep08</a>		22373	1812	7	410	rho GTPase-activating protein rich2 (chomee) alternative variant aSep08, mRNA.
<a href="#">chomee</a>	<a href="#">chomee.bSep08</a>		12540	548	4	182	rho gtpase-activating protein rich2 (chomee) alternative variant bSep08, mRNA.
<a href="#">chomee</a>	<a href="#">chomee.cSep08</a>		6605	1819	3	95	rho GTPase-activating protein rich2 like (chomee) alternative variant cSep08, mRNA.
<a href="#">chonoy</a>	<a href="#">chonoy.aSep08</a>		10981	820		62	putative protein (chonoy) mRNA.
<a href="#">chopor</a>	<a href="#">chopor.aSep08</a>		1944	266		34	putative protein (chopor) mRNA.
<a href="#">chorbor</a>	<a href="#">chorbor.aSep08</a>		2002	317		73	putative protein (chorbor) mRNA.
<a href="#">chorchy</a>	<a href="#">chorchy.aSep08</a>		3267	456	2	40	putative protein (chorchy) alternative variant aSep08, mRNA.
<a href="#">Chordc1</a>	<a href="#">Chordc1.bSep08</a>	<a href="#">315447</a>	16335	550	1	62	cysteine and histidine-rich domain (CHORD)-containing, zinc-binding protein 1 (Chordc1) alternative variant bSep08, mRNA.
<a href="#">chordoy</a>	<a href="#">chordoy.aSep08</a>		941	687	2	33	CRA b like (3.8 kD) (chordoy) alternative variant aSep08, mRNA.
<a href="#">chordoy</a>	<a href="#">chordoy.cSep08</a>		1007	500	3	38	putative protein (chordoy) alternative variant cSep08, mRNA.
<a href="#">chorflu</a>	<a href="#">chorflu.bSep08</a>		19044	414		46	putative protein (4.5 kD) (chorflu) alternative variant bSep08, mRNA.
<a href="#">chorfly</a>	<a href="#">chorfly.aSep08</a>		8222	710		44	putative protein (chorfly) mRNA.
<a href="#">chorgar</a>	<a href="#">chorgar.aSep08</a>		1872	208		69	pericentriolar material 1 CRA b (chorgar) mRNA.
<a href="#">chorja</a>	<a href="#">chorja.aSep08</a>		25050	693		145	putative protein of mammalian origin (16.0 kD) (chorja) mRNA.
<a href="#">chorjey</a>	<a href="#">chorjey.aSep08</a>		4092	527		33	putative protein (chorjey) mRNA.
<a href="#">chorkee</a>	<a href="#">chorkee.aSep08</a>		14914	481		53	putative protein (chorkee) mRNA.
<a href="#">chorkler</a>	<a href="#">chorkler.aSep08</a>		36001	663		220	calcium binding atopy-related autoantigen 1 CRA c like (chorkler) mRNA.
<a href="#">chorlo</a>	<a href="#">chorlo.aSep08</a>		11110	440		36	putative protein (4.0 kD) (chorlo) mRNA.
<a href="#">chormee</a>	<a href="#">chormee.aSep08</a>		4058	966		279	myosin heavy chain skeletal muscle adult (chormee) mRNA.
<a href="#">chornoy</a>	<a href="#">chornoy.aSep08</a>		11752	478		38	putative protein (4.4 kD) (chornoy) mRNA.
<a href="#">choroy</a>	<a href="#">choroy.aSep08</a>		7181	511		31	putative protein (3.7 kD) (choroy) mRNA.

<a href="#">chorpor</a>	<a href="#">chorpor.aSep08</a>		2885	751		201	angiotenin like 2 CRA b (chorpor) mRNA.
<a href="#">chorroy</a>	<a href="#">chorroy.aSep08</a>		6003	254		73	putative protein (chorroy) mRNA.
<a href="#">chorsa</a>	<a href="#">chorsa.aSep08</a>		5045	1906		82	specific peptidase 1 CRA c (chorsa) mRNA.
<a href="#">chorshee</a>	<a href="#">chorshee.aSep08</a>		30921	480		88	putative cytoplasmic protein (10.4 kD) (chorshee) mRNA.
<a href="#">chortu</a>	<a href="#">chortu.aSep08</a>		5136	261		50	putative protein (chortu) mRNA.
<a href="#">chorvar</a>	<a href="#">chorvar.aSep08</a>		1702	882		191	CRA a (chorvar) mRNA.
<a href="#">chorvo</a>	<a href="#">chorvo.aSep08</a>		1129	896	2	36	putative protein (4.1 kD) (chorvo) alternative variant aSep08, mRNA.
<a href="#">chorvo</a>	<a href="#">chorvo.bSep08</a>		5088	665	4	18	putative protein (1.9 kD) (chorvo) alternative variant bSep08, mRNA.
<a href="#">chorvo</a>	<a href="#">chorvo.dSep08</a>		5152	565	4		
<a href="#">chorwer</a>	<a href="#">chorwer.aSep08</a>		3296	759		147	putative protein (chorwer) alternative variant aSep08, mRNA.
<a href="#">chorwey</a>	<a href="#">chorwey.aSep08</a>		2429	390		39	putative protein (4.3 kD) (chorwey) mRNA.
<a href="#">chosa</a>	<a href="#">chosa.aSep08</a>		2055	379		44	putative protein (chosa) mRNA.
<a href="#">choshee</a>	<a href="#">choshee.aSep08</a>		1007	460		138	putative protein (choshee) mRNA.
<a href="#">chotu</a>	<a href="#">chotu.aSep08</a>		3357	721		75	putative protein (chotu) mRNA.
<a href="#">chovar</a>	<a href="#">chovar.aSep08</a>		6924	709		44	putative protein (4.5 kD) (chovar) mRNA.
<a href="#">chovo</a>	<a href="#">chovo.aSep08</a>		20819	1788	5	75	putative protein (8.6 kD) (chovo) alternative variant aSep08, mRNA.
<a href="#">chovo</a>	<a href="#">chovo.bSep08</a>		2087	720	3	61	putative protein (chovo) alternative variant bSep08, mRNA.
<a href="#">chovo</a>	<a href="#">chovo.cSep08</a>		1994	703	2	36	putative protein (chovo) alternative variant cSep08, mRNA.
<a href="#">chovo</a>	<a href="#">chovo.dSep08</a>		2071	629	3	33	putative protein (chovo) alternative variant dSep08, mRNA.
<a href="#">chower</a>	<a href="#">chower.aSep08</a>		3306	354		64	polyprotein (chower) mRNA.
<a href="#">chowey</a>	<a href="#">chowey.aSep08</a>		1741	686		65	putative protein (chowey) mRNA.
<a href="#">choybor</a>	<a href="#">choybor.aSep08</a>		1275	253		39	putative protein (4.4 kD) (choybor) mRNA.
<a href="#">choychy</a>	<a href="#">choychy.aSep08</a>		869	416		25	putative protein (2.9 kD) (choychy) mRNA.
<a href="#">choydoy</a>	<a href="#">choydoy.aSep08</a>		9328	715		79	putative protein (8.9 kD) (choydoy) mRNA.
<a href="#">choyflu</a>	<a href="#">choyflu.aSep08</a>		9993	2116	6	383	nucleoporin 98kDa CRA b (choyflu) alternative variant aSep08, mRNA.
<a href="#">choyflu</a>	<a href="#">choyflu.bSep08</a>		384	298	1	46	nucleoporin 98kDa CRA b (choyflu) alternative variant bSep08, mRNA.
<a href="#">choyfly</a>	<a href="#">choyfly.aSep08</a>		719	536		49	putative protein (5.6 kD) (choyfly) mRNA.
<a href="#">choygar</a>	<a href="#">choygar.aSep08</a>		4641	412		137	pericentriolar material 1 CRA b (choygar) mRNA.
<a href="#">choyja</a>	<a href="#">choyja.aSep08</a>		3946	758		72	putative cytoplasmic protein (8.1 kD) (choyja) mRNA.
<a href="#">choyjey</a>	<a href="#">choyjey.aSep08</a>		1227	306		32	putative protein (choyjey) mRNA.
<a href="#">choykee</a>	<a href="#">choykee.aSep08</a>		47119	779		107	putative protein (11.7 kD) (choykee) mRNA.
<a href="#">choykler</a>	<a href="#">choykler.aSep08</a>		35358	663		220	calcium binding atopy-related autoantigen 1 CRA c like (choykler) mRNA.
<a href="#">choylo</a>	<a href="#">choylo.aSep08</a>		1254	304		55	putative protein (choylo) mRNA.
<a href="#">choymee</a>	<a href="#">choymee.aSep08</a>		1076	233		77	myosin heavy chain skeletal muscle (choymee) mRNA.
<a href="#">choynoy</a>	<a href="#">choynoy.aSep08</a>		735	625		28	putative protein (3.0 kD) (choynoy) mRNA.
<a href="#">choypor</a>	<a href="#">choypor.aSep08</a>		14786	483		26	putative protein (3.0 kD) (choypor) mRNA.

<a href="#">choyroy</a>	<a href="#">choyroy.aSep08</a>		13158	340		60	putative protein (choyroy) mRNA.
<a href="#">choysa</a>	<a href="#">choysa.aSep08</a>		2061	407		83	putative protein (choysa) mRNA.
<a href="#">choyshee</a>	<a href="#">choyshee.aSep08</a>		2403	323		107	antigen 17 like (choyshee) mRNA.
<a href="#">choytu</a>	<a href="#">choytu.aSep08</a>		947	723		99	putative protein (choytu) mRNA.
<a href="#">choyvar</a>	<a href="#">choyvar.aSep08</a>		5065	614	4	113	CRA b (choyvar) alternative variant aSep08, mRNA.
<a href="#">choyvo</a>	<a href="#">choyvo.aSep08</a>		2042	757		44	putative protein (4.7 kD) (choyvo) mRNA.
<a href="#">choywer</a>	<a href="#">choywer.aSep08</a>		435	243		61	putative protein (choywer) alternative variant aSep08, mRNA.
<a href="#">choywey</a>	<a href="#">choywey.aSep08</a>		25809	544		26	putative protein (3.1 kD) (choywey) mRNA.
<a href="#">Chp</a>	<a href="#">Chp.aSep08</a>	<a href="#">64152</a>	33769	1035	7	195	calcium binding protein p22 (22.4 kD) (Chp) alternative variant aSep08, mRNA.
<a href="#">Chp</a>	<a href="#">Chp.bSep08</a>	<a href="#">64152</a>	29485	331	5	110	calcium binding protein p22 (Chp) alternative variant bSep08, mRNA.
<a href="#">Chp</a>	<a href="#">Chp.cSep08</a>	<a href="#">64152</a>	13791	2043	3	72	calcium binding protein p22 (8.0 kD) (Chp) alternative variant cSep08, mRNA.
<a href="#">Chp</a>	<a href="#">Chp.dSep08</a>	<a href="#">64152</a>	991	546	2	55	calcium binding protein p22 (6.2 kD) (Chp) alternative variant dSep08, mRNA.
<a href="#">Chpt1</a>	<a href="#">Chpt1.aSep08</a>	<a href="#">362866</a>	26943	1801	6	549	choline phosphotransferase 1 CRA c (Chpt1) alternative variant aSep08, mRNA.
<a href="#">Chpt1</a>	<a href="#">Chpt1.cSep08</a>	<a href="#">362866</a>	8330	655	5	175	choline phosphotransferase 1 (Chpt1) alternative variant cSep08, mRNA.
<a href="#">Chpt1</a>	<a href="#">Chpt1.dSep08</a>	<a href="#">362866</a>	24760	1062	2	86	putative cytoplasmic protein (9.5 kD) (Chpt1) alternative variant dSep08, mRNA.
<a href="#">Chrac1</a>	<a href="#">Chrac1.cSep08</a>	<a href="#">315058</a>	2950	827	3	31	chromatin accessibility complex 1 (3.5 kD) (Chrac1) alternative variant cSep08, mRNA.
<a href="#">Chrd</a>	<a href="#">Chrd.aSep08</a>	<a href="#">117275</a>	812	476		105	chordin (Chrd) mRNA.
<a href="#">Chrd1</a>	<a href="#">Chrd1.bSep08</a>	<a href="#">363455</a>	26338	1115	1	188	kojirin (Chrd1) alternative variant bSep08, mRNA.
<a href="#">Chrna3</a>	<a href="#">Chrna3.bSep08</a>	<a href="#">25101</a>	7976	377	4	29	cholinergic receptor, nicotinic, alpha polypeptide 3 (Chrna3) alternative variant bSep08, mRNA.
<a href="#">Chrna4</a>	<a href="#">Chrna4.bSep08</a>	<a href="#">25590</a>	36692	511	3	97	cholinergic receptor, nicotinic, alpha polypeptide 4 (Chrna4) alternative variant bSep08, mRNA.
<a href="#">Chrb1</a>	<a href="#">Chrb1.bSep08</a>	<a href="#">24261</a>	5736	963	2	229	cholinergic receptor, nicotinic, beta polypeptide 1 (muscle) (26.2 kD) (Chrb1) alternative variant bSep08, mRNA.
<a href="#">Chrne</a>	<a href="#">Chrne.bSep08</a>	<a href="#">29422</a>	1823	828	6	217	cholinergic receptor, nicotinic, epsilon polypeptide (Chrne) alternative variant bSep08, mRNA.
<a href="#">Chromo.0</a>	<a href="#">Chromo.0.aSep08</a>		3580	1213		403	chromodomain helicase DNA binding protein 3 like (Chromo.0) mRNA.
<a href="#">Chst5</a>	<a href="#">Chst5.bSep08</a>	<a href="#">307859</a>	18574	370	2	91	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 (Chst5) alternative variant bSep08, mRNA.
<a href="#">Chst5</a>	<a href="#">Chst5.cSep08</a>	<a href="#">307859</a>	1572	412	1	58	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 (Chst5) alternative variant cSep08, mRNA.
<a href="#">Chst5</a>	<a href="#">Chst5.dSep08</a>	<a href="#">307859</a>	18391	347	2	81	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 (Chst5) alternative variant dSep08, mRNA.
<a href="#">Chst10</a>	<a href="#">Chst10.bSep08</a>	<a href="#">140568</a>	12343	395	1	76	carbohydrate sulfotransferase 10 (Chst10) alternative variant bSep08, mRNA.

<a href="#">Chst14</a>	<a href="#">Chst14.bSep08</a>	<a href="#">691394</a>	1130	560	2	92	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 14 (10.2 kD) (Chst14) alternative variant bSep08, mRNA.
<a href="#">chubor</a>	<a href="#">chubor.aSep08</a>		10373	688	4	184	serine threonine kinase 38 CRA a (chubor) alternative variant aSep08, mRNA.
<a href="#">chuchy</a>	<a href="#">chuchy.aSep08</a>		2679	494		127	activating signal cointegrator 1 complex (chuchy) mRNA.
<a href="#">chudoy</a>	<a href="#">chudoy.aSep08</a>		1974	627		154	myosin VIIIb CRA a (chudoy) mRNA.
<a href="#">chufly</a>	<a href="#">chufly.aSep08</a>		71993	650		71	putative protein (chufly) mRNA.
<a href="#">chugar</a>	<a href="#">chugar.aSep08</a>		5714	752		71	putative mitochondrial protein (8.1 kD) (chugar) mRNA.
<a href="#">chujey</a>	<a href="#">chujey.aSep08</a>		12605	1140		314	putative protein of eukaryotic origin (chujey) mRNA.
<a href="#">Chuk</a>	<a href="#">Chuk.bSep08</a>	<a href="#">309361</a>	35518	3502	21	719	conserved helix-loop-helix ubiquitous kinase (82.2 kD) (Chuk) alternative variant bSep08, mRNA.
<a href="#">Chuk</a>	<a href="#">Chuk.cSep08</a>	<a href="#">309361</a>	3646	956	3	85	conserved helix-loop-helix ubiquitous kinase (9.8 kD) (Chuk) alternative variant cSep08, mRNA.
<a href="#">chukee</a>	<a href="#">chukee.aSep08</a>		536	373	2	82	putative protein of mammalian origin (chukee) alternative variant aSep08, mRNA.
<a href="#">chukler</a>	<a href="#">chukler.aSep08</a>		4421	649		86	uncharacterized protein c3orf1 homolog like (chukler) mRNA.
<a href="#">chulo</a>	<a href="#">chulo.aSep08</a>		818	361		26	putative protein (2.9 kD) (chulo) mRNA.
<a href="#">chumee</a>	<a href="#">chumee.aSep08</a>		5160	668		222	elac homolog 2 (chumee) mRNA.
<a href="#">chunoy</a>	<a href="#">chunoy.aSep08</a>		2522	1270		83	putative nuclear protein (9.3 kD) (chunoy) mRNA.
<a href="#">chupor</a>	<a href="#">chupor.aSep08</a>		455	279		88	putative protein (chupor) mRNA.
<a href="#">Churc1</a>	<a href="#">Churc1.bSep08</a>	<a href="#">299154</a>	14696	258	3	85	churchill (Churc1) alternative variant bSep08, mRNA.
<a href="#">Churc1</a>	<a href="#">Churc1.cSep08</a>	<a href="#">299154</a>	8953	583	3	74	churchill (8.7 kD) (Churc1) alternative variant cSep08, mRNA.
<a href="#">Churc1</a>	<a href="#">Churc1.dSep08</a>	<a href="#">299154</a>	10177	396	4	74	churchill (8.7 kD) (Churc1) alternative variant dSep08, mRNA.
<a href="#">Churc1</a>	<a href="#">Churc1.eSep08</a>	<a href="#">299154</a>	14580	1136	4	74	churchill (8.7 kD) (Churc1) alternative variant eSep08, mRNA.
<a href="#">Churc1</a>	<a href="#">Churc1.fSep08</a>	<a href="#">299154</a>	6701	585	2	35	putative protein of vertebrate origin (4.1 kD) (Churc1) alternative variant fSep08, mRNA.
<a href="#">churoy</a>	<a href="#">churoy.aSep08</a>		17594	693		131	nidogen 2 CRA a (14.1 kD) (churoy) mRNA.
<a href="#">chusa</a>	<a href="#">chusa.aSep08</a>		10123	730		33	putative protein (3.7 kD) (chusa) mRNA.
<a href="#">chushee</a>	<a href="#">chushee.aSep08</a>		6309	665	2	164	putative protein (chushee) alternative variant aSep08, mRNA.
<a href="#">chushee</a>	<a href="#">chushee.bSep08</a>		3278	650	1	159	putative protein (chushee) alternative variant bSep08, mRNA.
<a href="#">chutu</a>	<a href="#">chutu.aSep08</a>		3029	701		60	putative protein (6.9 kD) (chutu) mRNA.
<a href="#">chuvar</a>	<a href="#">chuvar.aSep08</a>		4012	535		41	putative protein (chuvar) mRNA.
<a href="#">chuvo</a>	<a href="#">chuvo.aSep08</a>		810	723		50	chaperonin containing Tcp1 (chuvo) mRNA.
<a href="#">chuwer</a>	<a href="#">chuwer.aSep08</a>		32766	611		121	putative protein (13.3 kD) (chuwer) mRNA.
<a href="#">chuwey</a>	<a href="#">chuwey.aSep08</a>		1830	711		41	putative protein (chuwey) mRNA.

<a href="#">chybor</a>	<a href="#">chybor.aSep08</a>		2606	468		155	putative protein of eukaryotic origin (chybor) mRNA.
<a href="#">chychy</a>	<a href="#">chychy.aSep08</a>		7862	371		36	complex protein 4 like (chychy) mRNA.
<a href="#">chydar</a>	<a href="#">chydar.aSep08</a>		7677	742		78	putative mitochondrial protein (8.7 kD) (chydar) mRNA.
<a href="#">chydoy</a>	<a href="#">chydoy.aSep08</a>		9631	738	2	38	putative protein (4.1 kD) (chydoy) alternative variant aSep08, mRNA.
<a href="#">chyflu</a>	<a href="#">chyflu.aSep08</a>		5810	670		74	putative endoplasmic reticulum protein (8.1 kD) (chyflu) mRNA.
<a href="#">chyfly</a>	<a href="#">chyfly.aSep08</a>		48961	595		198	laminin alpha 2 CRA b (chyfly) mRNA.
<a href="#">chygar</a>	<a href="#">chygar.aSep08</a>		5396	784		55	putative protein (6.1 kD) (chygar) mRNA.
<a href="#">chyja</a>	<a href="#">chyja.aSep08</a>		7987	261		86	putative protein (chyja) alternative variant aSep08, mRNA.
<a href="#">chyjey</a>	<a href="#">chyjey.aSep08</a>		6727	648		177	putative protein of eukaryotic origin (chyjey) mRNA.
<a href="#">chykee</a>	<a href="#">chykee.aSep08</a>		46147	666		222	putative protein of vertebrate origin (chykee) mRNA.
<a href="#">chykler</a>	<a href="#">chykler.aSep08</a>		1205	291		48	putative protein (chykler) mRNA.
<a href="#">chylo</a>	<a href="#">chylo.aSep08</a>		2285	475	2	111	putative protein (chylo) alternative variant aSep08, mRNA.
<a href="#">chymee</a>	<a href="#">chymee.aSep08</a>		661	348		27	putative protein (chymee) mRNA.
<a href="#">chynoy</a>	<a href="#">chynoy.aSep08</a>		3973	458	1	54	putative protein (6.3 kD) (chynoy) alternative variant aSep08, mRNA.
<a href="#">chynoy</a>	<a href="#">chynoy.bSep08</a>		4669	309	1	45	putative protein (chynoy) alternative variant bSep08, mRNA.
<a href="#">chypor</a>	<a href="#">chypor.aSep08</a>		24048	1784		61	putative protein (7.1 kD) (chypor) alternative variant aSep08, mRNA.
<a href="#">chyroy</a>	<a href="#">chyroy.aSep08</a>		28512	777	3	31	putative protein (chyroy) alternative variant aSep08, mRNA.
<a href="#">chyroy</a>	<a href="#">chyroy.bSep08</a>		44112	751	5	31	putative protein (3.3 kD) (chyroy) alternative variant bSep08, mRNA.
<a href="#">chyroy</a>	<a href="#">chyroy.cSep08</a>		42802	747	5	64	putative protein (chyroy) alternative variant cSep08, mRNA.
<a href="#">chyroy</a>	<a href="#">chyroy.dSep08</a>		28530	677	5	43	putative protein (4.4 kD) (chyroy) alternative variant dSep08, complete mRNA.
<a href="#">chyroy</a>	<a href="#">chyroy.eSep08</a>		28597	672	4	59	putative protein (6.3 kD) (chyroy) alternative variant eSep08, mRNA.
<a href="#">chysa</a>	<a href="#">chysa.aSep08</a>		5101	753		79	putative protein (chysa) mRNA.
<a href="#">chyshee</a>	<a href="#">chyshee.aSep08</a>		17003	1797	3	546	putative protein (chyshee) alternative variant aSep08, mRNA.
<a href="#">chytu</a>	<a href="#">chytu.aSep08</a>		11778	732		87	putative protein (chytu) mRNA.
<a href="#">chyvar</a>	<a href="#">chyvar.aSep08</a>		421	236	2	36	putative protein (chyvar) alternative variant aSep08, mRNA.
<a href="#">chyvo</a>	<a href="#">chyvo.aSep08</a>		6538	423		41	putative protein (chyvo) mRNA.
<a href="#">chywer</a>	<a href="#">chywer.aSep08</a>		31864	645		214	CRA a (chywer) mRNA.
<a href="#">chywey</a>	<a href="#">chywey.aSep08</a>		1695	349		71	putative protein (chywey) mRNA.
<a href="#">Ciao1</a>	<a href="#">Ciao1.bSep08</a>	<a href="#">29231</a>	4889	2572	3	194	cytosolic iron-sulfur protein assembly 1 homolog ( <i>S. cerevisiae</i> ) (Ciao1) alternative variant bSep08, mRNA.
<a href="#">Ciapin1</a>	<a href="#">Ciapin1.aSep08</a>	<a href="#">307649</a>	4230	1319	4	316	cytokine induced apoptosis inhibitor 1 (Ciapin1) alternative variant aSep08, mRNA.

<a href="#">Ciapin1</a>	<a href="#">Ciapin1.cSep08</a>	<a href="#">307649</a>	10673	812	7	246	cytokine induced apoptosis inhibitor 1 (Ciapin1) alternative variant cSep08, mRNA.
<a href="#">Ciapin1</a>	<a href="#">Ciapin1.dSep08</a>	<a href="#">307649</a>	10677	813	7	246	cytokine induced apoptosis inhibitor 1 (Ciapin1) alternative variant dSep08, mRNA.
<a href="#">Ciapin1</a>	<a href="#">Ciapin1.eSep08</a>	<a href="#">307649</a>	6667	822	4	151	cytokine induced apoptosis inhibitor 1 (Ciapin1) alternative variant eSep08, mRNA.
<a href="#">Ciapin1</a>	<a href="#">Ciapin1.fSep08</a>	<a href="#">307649</a>	4590	741	2	74	cytokine induced apoptosis inhibitor 1 (7.9 kD) (Ciapin1) alternative variant fSep08, complete mRNA.
<a href="#">Cib1</a>	<a href="#">Cib1.aSep08</a>	<a href="#">81823</a>	13232	1548	1	203	calcium and integrin binding 1 (calmyrin) (23.2 kD) (Cib1) alternative variant aSep08, mRNA.
<a href="#">Cib4</a>	<a href="#">Cib4.aSep08</a>	<a href="#">688819</a>	2580	341		58	calcium and integrin binding family member 4 (Cib4) mRNA.
<a href="#">Cic</a>	<a href="#">Cic.bSep08</a>	<a href="#">308435</a>	1868	1159	5	386	capicua homolog (Drosophila) (Cic) alternative variant bSep08, mRNA.
<a href="#">Cic</a>	<a href="#">Cic.cSep08</a>	<a href="#">308435</a>	1270	848	3	282	capicua homolog (Drosophila) (Cic) alternative variant cSep08, mRNA.
<a href="#">Cic</a>	<a href="#">Cic.dSep08</a>	<a href="#">308435</a>	1074	817	2	135	capicua homolog (Drosophila) (Cic) alternative variant dSep08, mRNA.
<a href="#">Cidea</a>	<a href="#">Cidea.aSep08</a>	<a href="#">291541</a>	25906	1379		215	cell death-inducing DNA fragmentation factor, alpha subunit-like effector A (24.3 kD) (Cidea) mRNA.
<a href="#">Cideb</a>	<a href="#">Cideb.bSep08</a>	<a href="#">364388</a>	3900	819	2	240	cell death-inducing DNA fragmentation factor, alpha subunit-like effector B (Cideb) alternative variant bSep08, mRNA.
<a href="#">Ciita</a>	<a href="#">Ciita.bSep08</a>	<a href="#">85483</a>	2701	2186	2	67	class II transactivator (Ciita) alternative variant bSep08, mRNA.
<a href="#">Cinp</a>	<a href="#">Cinp.bSep08</a>	<a href="#">299334</a>	11931	614	5	158	cyclin-dependent kinase 2-interacting protein (17.7 kD) (Cinp) alternative variant bSep08, complete mRNA.
<a href="#">Cinp</a>	<a href="#">Cinp.cSep08</a>	<a href="#">299334</a>	13279	724	4	118	cyclin-dependent kinase 2-interacting protein (13.0 kD) (Cinp) alternative variant cSep08, mRNA.
<a href="#">Cinp</a>	<a href="#">Cinp.dSep08</a>	<a href="#">299334</a>	3575	648	2	37	cyclin-dependent kinase 2-interacting protein (Cinp) alternative variant dSep08, mRNA.
<a href="#">Cip29</a>	<a href="#">Cip29.bSep08</a>	<a href="#">362819</a>	44177	859	9	175	cytokine induced protein 29 kDa (Cip29) alternative variant bSep08, mRNA.
<a href="#">Cip29</a>	<a href="#">Cip29.cSep08</a>	<a href="#">362819</a>	61376	974	13	127	cytokine induced protein 29 kDa (14.1 kD) (Cip29) alternative variant cSep08, mRNA.
<a href="#">Cip29</a>	<a href="#">Cip29.dSep08</a>	<a href="#">362819</a>	50871	813	10	103	cytokine induced protein 29 kDa (11.4 kD) (Cip29) alternative variant dSep08, mRNA.
<a href="#">Cip29</a>	<a href="#">Cip29.eSep08</a>	<a href="#">362819</a>	5235	575	2	97	cytokine induced protein 29 kDa (11.4 kD) (Cip29) alternative variant eSep08, mRNA.
<a href="#">Cip98</a>	<a href="#">Cip98.bSep08</a>	<a href="#">313255</a>	15672	1120	5	324	CASK-interacting protein CIP98 (Cip98) alternative variant bSep08, mRNA.
<a href="#">Cip98</a>	<a href="#">Cip98.cSep08</a>	<a href="#">313255</a>	38027	1287	3	176	CASK-interacting protein CIP98 (Cip98) alternative variant cSep08, mRNA.
<a href="#">Cip98</a>	<a href="#">Cip98.dSep08</a>	<a href="#">313255</a>	62152	390	3	129	CASK-interacting protein CIP98 (Cip98) alternative variant dSep08, mRNA.
<a href="#">Cip98</a>	<a href="#">Cip98.eSep08</a>	<a href="#">313255</a>	3487	413	3	84	CASK-interacting protein CIP98 (Cip98) alternative variant eSep08, mRNA.

<a href="#">Cipar1</a>	<a href="#">Cipar1.aSep08</a>	<a href="#">286894</a>	32640	1250		282	castration induced prostatic apoptosis-related protein 1 (Cipar1) mRNA.
<a href="#">Cir</a>	<a href="#">Cir.aSep08</a>	<a href="#">362149</a>	30115	1916	8	451	CBF1 interacting corepressor (51.4 kD) (Cir) alternative variant aSep08, complete mRNA.
<a href="#">Cir</a>	<a href="#">Cir.bSep08</a>	<a href="#">362149</a>	29993	1783	7	376	CBF1 interacting corepressor (42.6 kD) (Cir) alternative variant bSep08, complete mRNA.
<a href="#">Cir</a>	<a href="#">Cir.cSep08</a>	<a href="#">362149</a>	3289	781	1	201	CBF1 interacting corepressor (Cir) alternative variant cSep08, mRNA.
<a href="#">Cirbp</a>	<a href="#">Cirbp.bSep08</a>	<a href="#">81825</a>	4943	2518	7	176	cold inducible RNA binding protein (18.8 kD) (Cirbp) alternative variant bSep08, complete mRNA.
<a href="#">Cirbp</a>	<a href="#">Cirbp.cSep08</a>	<a href="#">81825</a>	3036	761	7	174	cold inducible RNA binding protein (Cirbp) alternative variant cSep08, mRNA.
<a href="#">Cirbp</a>	<a href="#">Cirbp.eSep08</a>	<a href="#">81825</a>	4953	909	9	172	cold inducible RNA binding protein (18.6 kD) (Cirbp) alternative variant eSep08, complete mRNA.
<a href="#">Cirh1a</a>	<a href="#">Cirh1a.bSep08</a>	<a href="#">291987</a>	10915	734	1	233	cirrhosis, autosomal recessive 1A (human) (Cirh1a) alternative variant bSep08, mRNA.
<a href="#">Cisd2</a>	<a href="#">Cisd2.aSep08</a>	<a href="#">295457</a>	24810	2883		147	CDGSH iron sulfur domain 2 (Cisd2) mRNA.
<a href="#">Cit</a>	<a href="#">Cit.bSep08</a>	<a href="#">83620</a>	20781	522	4	134	citron (rho-interacting, serine/threonine kinase 21) (Cit) alternative variant bSep08, mRNA.
<a href="#">Cit</a>	<a href="#">Cit.dSep08</a>	<a href="#">83620</a>	29272	672	3	63	citron (rho-interacting, serine/threonine kinase 21) (Cit) alternative variant dSep08, mRNA.
<a href="#">Cit</a>	<a href="#">Cit.eSep08</a>	<a href="#">83620</a>	18323	1182	3	64	citron (rho-interacting, serine/threonine kinase 21) (Cit) alternative variant eSep08, mRNA.
<a href="#">Cited1</a>	<a href="#">Cited1.bSep08</a>	<a href="#">64466</a>	3172	607	1	120	cbp/p300-interacting transactivator with Glu/Asp-rich carboxy-terminal domain 1 (Cited1) alternative variant bSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.bSep08</a>	<a href="#">296639</a>	15030	2590	18	818	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant bSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.cSep08</a>	<a href="#">296639</a>	4065	440	4	146	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant cSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.dSep08</a>	<a href="#">296639</a>	4399	760	5	146	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant dSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.eSep08</a>	<a href="#">296639</a>	3395	737	4	143	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant eSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.fSep08</a>	<a href="#">296639</a>	7113	427	4	142	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant fSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.gSep08</a>	<a href="#">296639</a>	3800	675	4	137	CDKN1A interacting zinc finger protein 1 (Ciz1) alternative variant gSep08, mRNA.
<a href="#">Ciz1</a>	<a href="#">Ciz1.hSep08</a>	<a href="#">296639</a>	1918	816	3	104	CDKN1A interacting zinc finger protein 1 (11.5 kD) (Ciz1) alternative variant hSep08, mRNA.
<a href="#">Ckap2</a>	<a href="#">Ckap2.aSep08</a>	<a href="#">306575</a>	26590	2616	4	675	cytoskeleton associated protein 2 (Ckap2) alternative variant aSep08, mRNA.
<a href="#">Ckap2l</a>	<a href="#">Ckap2l.aSep08</a>	<a href="#">680515</a>	3870	629		157	cytoskeleton associated protein 2-like (Ckap2l) mRNA.
<a href="#">Ckap5</a>	<a href="#">Ckap5.aSep08</a>	<a href="#">311191</a>	40531	3958	3	1166	cytoskeleton associated protein 5 (Ckap5) alternative variant aSep08, mRNA.
<a href="#">Ckap5</a>	<a href="#">Ckap5.bSep08</a>	<a href="#">311191</a>	1686	701	1	189	cytoskeleton associated protein 5 (Ckap5) alternative variant bSep08, mRNA.

<a href="#">Ckb</a>	<a href="#">Ckb.bSep08</a>	<a href="#">24264</a>	1503	652	4	129	creatine kinase, brain (Ckb) alternative variant bSep08, mRNA.
<a href="#">Cklf</a>	<a href="#">Cklf.bSep08</a>	<a href="#">245978</a>	9280	452	3	98	chemokine-like factor (10.8 kD) (Cklf) alternative variant bSep08, mRNA.
<a href="#">Cklf</a>	<a href="#">Cklf.cSep08</a>	<a href="#">245978</a>	2368	701	2	61	chemokine-like factor (6.6 kD) (Cklf) alternative variant cSep08, mRNA.
<a href="#">Ckmt1</a>	<a href="#">Ckmt1.bSep08</a>	<a href="#">29593</a>	2981	780	5	219	creatine kinase, mitochondrial 1, ubiquitous (Ckmt1) alternative variant bSep08, mRNA.
<a href="#">Ckmt2</a>	<a href="#">Ckmt2.aSep08</a>	<a href="#">688698</a>	21055	802		244	creatine kinase, mitochondrial 2, sarcomeric (Ckmt2) alternative variant aSep08, mRNA.
<a href="#">Ckmt2</a>	<a href="#">Ckmt2.bSep08</a>	<a href="#">688698</a>	1593	278		92	creatine kinase, mitochondrial 2, sarcomeric (Ckmt2) alternative variant bSep08, mRNA.
<a href="#">Clasp2</a>	<a href="#">Clasp2.bSep08</a>	<a href="#">114514</a>	14468	434	4	144	CLIP associating protein 2 (Clasp2) alternative variant bSep08, mRNA.
<a href="#">Clasp2</a>	<a href="#">Clasp2.cSep08</a>	<a href="#">114514</a>	2819	1446	2	113	CLIP associating protein 2 (12.9 kD) (Clasp2) alternative variant cSep08, mRNA.
<a href="#">Clca2</a>	<a href="#">Clca2.bSep08</a>	<a href="#">362052</a>	22295	3210		514	chloride channel calcium activated 2 (58.7 kD) (Clca2) alternative variant bSep08, mRNA.
<a href="#">Clca4</a>	<a href="#">Clca4.bSep08</a>	<a href="#">362053</a>	1611	719	2	97	chloride channel, calcium activated, family member 4 (Clca4) alternative variant bSep08, mRNA.
<a href="#">Clca4</a>	<a href="#">Clca4.cSep08</a>	<a href="#">362053</a>	4941	319	3	61	chloride channel, calcium activated, family member 4 (Clca4) alternative variant cSep08, mRNA.
<a href="#">Clca5</a>	<a href="#">Clca5.bSep08</a>	<a href="#">308016</a>	1657	577	2	38	chloride channel calcium activated 5 (Clca5) alternative variant bSep08, mRNA.
<a href="#">Clcc1</a>	<a href="#">Clcc1.bSep08</a>	<a href="#">170927</a>	4234	1293	3	170	chloride channel CLIC-like 1 (Clcc1) alternative variant bSep08, mRNA.
<a href="#">Clcc1</a>	<a href="#">Clcc1.cSep08</a>	<a href="#">170927</a>	23079	603	5	159	chloride channel CLIC-like 1 (Clcc1) alternative variant cSep08, mRNA.
<a href="#">Clcn1</a>	<a href="#">Clcn1.bSep08</a>	<a href="#">25688</a>	7611	1609	6	374	chloride channel 1 (Clcn1) alternative variant bSep08, mRNA.
<a href="#">Clcn2</a>	<a href="#">Clcn2.bSep08</a>	<a href="#">29232</a>	2758	719	6	239	chloride channel 2 (Clcn2) alternative variant bSep08, mRNA.
<a href="#">Clcn2</a>	<a href="#">Clcn2.cSep08</a>	<a href="#">29232</a>	5822	1468	6	176	chloride channel 2 (Clcn2) alternative variant cSep08, mRNA.
<a href="#">Clcn2</a>	<a href="#">Clcn2.dSep08</a>	<a href="#">29232</a>	1737	737	7	118	chloride channel 2 (Clcn2) alternative variant dSep08, mRNA.
<a href="#">Clcn2</a>	<a href="#">Clcn2.eSep08</a>	<a href="#">29232</a>	848	724	2	84	chloride channel 2 (9.1 kD) (Clcn2) alternative variant eSep08, mRNA.
<a href="#">Clcn3</a>	<a href="#">Clcn3.bSep08</a>	<a href="#">84360</a>	14720	1407	4	230	chloride channel 3 (25.2 kD) (Clcn3) alternative variant bSep08, mRNA.
<a href="#">Clcn3</a>	<a href="#">Clcn3.cSep08</a>	<a href="#">84360</a>	21614	440	4	146	chloride channel 3 (Clcn3) alternative variant cSep08, mRNA.
<a href="#">Clcn6</a>	<a href="#">Clcn6.bSep08</a>	<a href="#">295586</a>	2674	973	3	149	chloride channel 6 (Clcn6) alternative variant bSep08, mRNA.
<a href="#">Clcn6</a>	<a href="#">Clcn6.cSep08</a>	<a href="#">295586</a>	2321	696	3	110	chloride channel 6 (Clcn6) alternative variant cSep08, mRNA.



<a href="#">Clcn7</a>	<a href="#">Clcn7.bSep08</a>	<a href="#">29233</a>	6082	2300	7	250	chloride channel 7 (Clcn7) alternative variant bSep08, mRNA.
<a href="#">Clcn7</a>	<a href="#">Clcn7.cSep08</a>	<a href="#">29233</a>	2238	767	5	188	chloride channel 7 (Clcn7) alternative variant cSep08, mRNA.
<a href="#">Clcn7</a>	<a href="#">Clcn7.dSep08</a>	<a href="#">29233</a>	4481	738	7	104	chloride channel 7 (Clcn7) alternative variant dSep08, mRNA.
<a href="#">Clcnka</a>	<a href="#">Clcnka.bSep08</a>	<a href="#">79425</a>	15973	2460	15	687	chloride channel Ka (Clcnka) alternative variant bSep08, mRNA.
<a href="#">Clcnka</a>	<a href="#">Clcnka.cSep08</a>	<a href="#">79425</a>	2995	669	1	183	chloride channel Ka (Clcnka) alternative variant cSep08, mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.aSep08</a>	<a href="#">79430</a>	11751	2229	19	644	chloride channel Kb CRA b (70.4 kD) (Clcnkb) alternative variant aSep08, complete mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.bSep08</a>	<a href="#">79430</a>	11751	2279	19	585	chloride channel Kb CRA b (63.4 kD) (Clcnkb) alternative variant bSep08, complete mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.cSep08</a>	<a href="#">79430</a>	11751	2251	20	583	chloride channel Kb CRA b (63.1 kD) (Clcnkb) alternative variant cSep08, complete mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.dSep08</a>	<a href="#">79430</a>	11751	2301	20	583	chloride channel Kb CRA b (63.1 kD) (Clcnkb) alternative variant dSep08, complete mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.eSep08</a>	<a href="#">79430</a>	11728	2729	18	340	chloride channel Kb CRA b (37.0 kD) (Clcnkb) alternative variant eSep08, complete mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.fSep08</a>	<a href="#">79430</a>	1828	805	4	234	chloride channel Kb CRA b (Clcnkb) alternative variant fSep08, mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.gSep08</a>	<a href="#">79430</a>	4799	661	7	212	chloride channel Kb CRA b (Clcnkb) alternative variant gSep08, mRNA.
<a href="#">Clcnkb</a>	<a href="#">Clcnkb.iSep08</a>	<a href="#">79430</a>	2084	802	3	43	putative protein (Clcnkb) alternative variant iSep08, mRNA.
<a href="#">Cldn6</a>	<a href="#">Cldn6.aSep08</a>	<a href="#">287098</a>	2550	967	2	236	claudin 6 (26.0 kD) (Cldn6) alternative variant aSep08, mRNA.
<a href="#">Cldn10</a>	<a href="#">Cldn10.bSep08</a>	<a href="#">290485</a>	20802	1827	2	196	claudin 10 (Cldn10) alternative variant bSep08, mRNA.
<a href="#">Cldn11</a>	<a href="#">Cldn11.bSep08</a>	<a href="#">84588</a>	7245	342	1	114	claudin 11 (Cldn11) alternative variant bSep08, mRNA.
<a href="#">Cldn19</a>	<a href="#">Cldn19.aSep08</a>	<a href="#">298487</a>	4761	912	5	224	claudin 19 (23.3 kD) (Cldn19) alternative variant aSep08, complete mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.bSep08</a>	<a href="#">288182</a>	5031	875	3	291	putative protein, with at least 3 transmembrane domains (Cldnd1) alternative variant bSep08, mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.cSep08</a>	<a href="#">288182</a>	5070	840	2	213	putative endoplasmic reticulum protein of bilateral origin (23.7 kD) (Cldnd1) alternative variant cSep08, mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.dSep08</a>	<a href="#">288182</a>	4983	748	2	191	putative protein (Cldnd1) alternative variant dSep08, mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.eSep08</a>	<a href="#">288182</a>	4644	594	2	165	putative protein of vertebrate origin (Cldnd1) alternative variant eSep08, mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.fSep08</a>	<a href="#">288182</a>	4641	611	2	159	putative protein of vertebrate origin (Cldnd1) alternative variant fSep08, mRNA.
<a href="#">Cldnd1</a>	<a href="#">Cldnd1.gSep08</a>	<a href="#">288182</a>	3290	408	2	124	putative protein of vertebrate origin (Cldnd1) alternative variant gSep08, mRNA.
<a href="#">Clec1b</a>	<a href="#">Clec1b.aSep08</a>	<a href="#">500336</a>	8099	707	2	197	C-type lectin domain family 1, member b (22.9 kD) (Clec1b) alternative variant aSep08, mRNA.
<a href="#">Clec1b</a>	<a href="#">Clec1b.bSep08</a>	<a href="#">500336</a>	8184	514		59	C-type lectin domain family 1, member b (6.8 kD) (Clec1b) alternative variant bSep08, mRNA.

<a href="#">Clec2d</a>	<a href="#">Clec2d.bSep08</a>	<a href="#">113937</a>	20983	387	3	104	C-type lectin domain family 2, member d (Clec2d) alternative variant bSep08, mRNA.
<a href="#">Clec2h</a>	<a href="#">Clec2h.aSep08</a>	<a href="#">312745</a>	2354	1457		83	C-type lectin domain family 2, member h (Clec2h) mRNA.
<a href="#">Clec4a1</a>	<a href="#">Clec4a1.bSep08</a>	<a href="#">362430</a>	2375	1008	2	81	C-type lectin domain family 4, member a1 (9.3 kD) (Clec4a1) alternative variant bSep08, mRNA.
<a href="#">Clec4b2</a>	<a href="#">Clec4b2.bSep08</a>	<a href="#">450222</a>	24130	421		89	C-type lectin domain family 4, member b2 (10.1 kD) (Clec4b2) alternative variant bSep08, mRNA.
<a href="#">Clec7a</a>	<a href="#">Clec7a.aSep08</a>	<a href="#">502902</a>	11166	2518	2	235	C-type lectin domain family 7, member a (26.5 kD) (Clec7a) alternative variant aSep08, mRNA.
<a href="#">Clec7a</a>	<a href="#">Clec7a.bSep08</a>	<a href="#">502902</a>	9516	733	1	190	C-type lectin domain family 7, member a (21.3 kD) (Clec7a) alternative variant bSep08, mRNA.
<a href="#">Clec16a</a>	<a href="#">Clec16a.aSep08</a>	<a href="#">287044</a>	50072	1785	2	440	C-type lectin domain family 16, member A (Clec16a) alternative variant aSep08, mRNA.
<a href="#">Clec16a</a>	<a href="#">Clec16a.bSep08</a>	<a href="#">287044</a>	31034	666	2	179	C-type lectin domain family 16, member A (Clec16a) alternative variant bSep08, mRNA.
<a href="#">Clecsf6</a>	<a href="#">Clecsf6.bSep08</a>	<a href="#">474143</a>	17711	809	2	63	C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamily member 6 (Clecsf6) alternative variant bSep08, mRNA.
<a href="#">Clgn</a>	<a href="#">Clgn.bSep08</a>	<a href="#">685504</a>	2567	609	3	93	calmegin (Clgn) alternative variant bSep08, mRNA.
<a href="#">Clgn</a>	<a href="#">Clgn.cSep08</a>	<a href="#">685504</a>	7891	257	3	63	calmegin (Clgn) alternative variant cSep08, mRNA.
<a href="#">Clc2</a>	<a href="#">Clc2.bSep08</a>	<a href="#">294141</a>	1405	761	1	114	chloride intracellular channel 2 (Clc2) alternative variant bSep08, mRNA.
<a href="#">Clc5</a>	<a href="#">Clc5.bSep08</a>	<a href="#">94272</a>	37554	5756	2	123	chloride intracellular channel 5 (14.1 kD) (Clc5) alternative variant bSep08, mRNA.
<a href="#">Clc6</a>	<a href="#">Clc6.bSep08</a>	<a href="#">304081</a>	9460	1067	2	163	chloride intracellular channel 6 (Clc6) alternative variant bSep08, mRNA.
<a href="#">Clip1</a>	<a href="#">Clip1.bSep08</a>	<a href="#">65201</a>	40973	2811	11	462	restin CRA c (Clip1) alternative variant bSep08, mRNA.
<a href="#">Clip1</a>	<a href="#">Clip1.cSep08</a>	<a href="#">65201</a>	17950	1337	6	445	restin CRA b (Clip1) alternative variant cSep08, mRNA.
<a href="#">Clip1</a>	<a href="#">Clip1.dSep08</a>	<a href="#">65201</a>	18143	1567	6	340	restin CRA a (35.5 kD) (Clip1) alternative variant dSep08, mRNA.
<a href="#">Clip1</a>	<a href="#">Clip1.eSep08</a>	<a href="#">65201</a>	31688	619	6	197	restin CRA a (Clip1) alternative variant eSep08, mRNA.
<a href="#">Clip1</a>	<a href="#">Clip1.fSep08</a>	<a href="#">65201</a>	29473	781	6	189	restin CRA b (Clip1) alternative variant fSep08, mRNA.
<a href="#">Clip2</a>	<a href="#">Clip2.bSep08</a>	<a href="#">29264</a>	22947	1636	8	451	linker 2 (Clip2) alternative variant bSep08, mRNA.
<a href="#">Clip3</a>	<a href="#">Clip3.bSep08</a>	<a href="#">308493</a>	5042	572	5	190	putative protein of metazoan origin (Clip3) alternative variant bSep08, mRNA.
<a href="#">Clip4</a>	<a href="#">Clip4.bSep08</a>	<a href="#">298801</a>	46490	1429	10	409	restin-like 2 CRA a (Clip4) alternative variant bSep08, mRNA.
<a href="#">Clip4</a>	<a href="#">Clip4.cSep08</a>	<a href="#">298801</a>	15939	753	4	165	restin-like 2 CRA e (Clip4) alternative variant cSep08, mRNA.
<a href="#">Clip4</a>	<a href="#">Clip4.dSep08</a>	<a href="#">298801</a>	1028	330	2	76	restin-like 2 CRA d (Clip4) alternative variant dSep08, mRNA.
<a href="#">Clk1</a>	<a href="#">Clk1.bSep08</a>	<a href="#">301434</a>	11219	3034	11	330	CDC-like kinase 1 (38.6 kD) (Clk1) alternative variant bSep08, complete mRNA.
<a href="#">Clk1</a>	<a href="#">Clk1.cSep08</a>	<a href="#">301434</a>	3821	730	4	179	CDC-like kinase 1 (21.6 kD) (Clk1) alternative variant cSep08, complete mRNA.

<a href="#">Clk1</a>	<a href="#">Clk1.dSep08</a>	<a href="#">301434</a>	1061	975	2	177	CDC-like kinase 1 (Clk1) alternative variant dSep08, mRNA.
<a href="#">Clk1</a>	<a href="#">Clk1.eSep08</a>	<a href="#">301434</a>	6097	717	6	135	CDC-like kinase 1 (16.4 kD) (Clk1) alternative variant eSep08, mRNA.
<a href="#">Clk1</a>	<a href="#">Clk1.fSep08</a>	<a href="#">301434</a>	1115	613	3	95	CDC-like kinase 1 (11.5 kD) (Clk1) alternative variant fSep08, mRNA.
<a href="#">Clk1</a>	<a href="#">Clk1.hSep08</a>	<a href="#">301434</a>	860	487	2	45	CDC-like kinase 1 (5.4 kD) (Clk1) alternative variant hSep08, mRNA.
<a href="#">Clk2</a>	<a href="#">Clk2.bSep08</a>	<a href="#">365842</a>	7828	2858	8	307	CDC-like kinase 2 (35.1 kD) (Clk2) alternative variant bSep08, mRNA.
<a href="#">Clk2</a>	<a href="#">Clk2.cSep08</a>	<a href="#">365842</a>	4491	633	5	211	CDC-like kinase 2 (Clk2) alternative variant cSep08, mRNA.
<a href="#">Clk2</a>	<a href="#">Clk2.dSep08</a>	<a href="#">365842</a>	4467	623	5	100	CDC-like kinase 2 (Clk2) alternative variant dSep08, mRNA.
<a href="#">Clk2</a>	<a href="#">Clk2.fSep08</a>	<a href="#">365842</a>	1685	402	4	81	CDC-like kinase 2 (Clk2) alternative variant fSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.bSep08</a>	<a href="#">171305</a>	11483	939	8	312	CDC-like kinase 3 (Clk3) alternative variant bSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.cSep08</a>	<a href="#">171305</a>	9473	1531	7	272	CDC-like kinase 3 (Clk3) alternative variant cSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.dSep08</a>	<a href="#">171305</a>	5180	893	6	259	CDC-like kinase 3 (Clk3) alternative variant dSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.eSep08</a>	<a href="#">171305</a>	7353	774	6	183	CDC-like kinase 3 (Clk3) alternative variant eSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.fSep08</a>	<a href="#">171305</a>	4718	559	3	151	CDC-like kinase 3 (Clk3) alternative variant fSep08, mRNA.
<a href="#">Clk3</a>	<a href="#">Clk3.gSep08</a>	<a href="#">171305</a>	3804	849	3	117	CDC-like kinase 3 (Clk3) alternative variant gSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.aSep08</a>	<a href="#">287269</a>	12142	794	7	224	CDC like kinase 4 (Clk4) alternative variant aSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.bSep08</a>	<a href="#">287269</a>	13401	625	6	134	CDC like kinase 4 (16.6 kD) (Clk4) alternative variant bSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.cSep08</a>	<a href="#">287269</a>	8590	443	5	122	CDC like kinase 4 (Clk4) alternative variant cSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.dSep08</a>	<a href="#">287269</a>	8742	654	6	113	CDC like kinase 4 (Clk4) alternative variant dSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.eSep08</a>	<a href="#">287269</a>	5248	726	2	113	CDC like kinase 4 (Clk4) alternative variant eSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.fSep08</a>	<a href="#">287269</a>	8621	408	4	102	CDC like kinase 4 (Clk4) alternative variant fSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.gSep08</a>	<a href="#">287269</a>	2697	1157	2	58	CDC like kinase 4 (6.7 kD) (Clk4) alternative variant gSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.hSep08</a>	<a href="#">287269</a>	9922	658	5	59	CDC like kinase 4 (6.6 kD) (Clk4) alternative variant hSep08, mRNA.
<a href="#">Clk4</a>	<a href="#">Clk4.iSep08</a>	<a href="#">287269</a>	11648	562	5	59	CDC like kinase 4 (6.6 kD) (Clk4) alternative variant iSep08, mRNA.
<a href="#">Clmn</a>	<a href="#">Clmn.bSep08</a>	<a href="#">299285</a>	3862	654	1	177	calmin (Clmn) alternative variant bSep08, mRNA.

<a href="#">Cln3</a>	<a href="#">Cln3.bSep08</a>	<a href="#">293485</a>	7343	1244	12	237	ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (Cln3) alternative variant bSep08, mRNA.
<a href="#">Cln3</a>	<a href="#">Cln3.cSep08</a>	<a href="#">293485</a>	5475	730	8	198	ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (Cln3) alternative variant cSep08, mRNA.
<a href="#">Cln3</a>	<a href="#">Cln3.dSep08</a>	<a href="#">293485</a>	5433	801	8	195	ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (Cln3) alternative variant dSep08, mRNA.
<a href="#">Cln3</a>	<a href="#">Cln3.eSep08</a>	<a href="#">293485</a>	3879	811	8	193	ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (Cln3) alternative variant eSep08, mRNA.
<a href="#">Cln3</a>	<a href="#">Cln3.fSep08</a>	<a href="#">293485</a>	3989	755	10	123	ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (Cln3) alternative variant fSep08, mRNA.
<a href="#">Cln5</a>	<a href="#">Cln5.aSep08</a>	<a href="#">306128</a>	9423	2258		339	ceroid-lipofuscinosis, neuronal 5 (Cln5) mRNA.
<a href="#">Cln6andCalml4</a>	<a href="#">Cln6andCalml4.aSep08</a>	<a href="#">315746</a>	15005	2006	7	329	ceroid-lipofuscinosis neuronal 6 (Cln6andCalml4) alternative variant aSep08, mRNA.
<a href="#">Cln6andCalml4</a>	<a href="#">Cln6andCalml4.aSep08</a>	<a href="#">691455</a>	15005	2006	7	329	ceroid-lipofuscinosis neuronal 6 (Cln6andCalml4) alternative variant aSep08, mRNA.
<a href="#">Cln6andCalml4</a>	<a href="#">Cln6andCalml4.cSep08</a>	<a href="#">315746</a>	24314	678	7	165	ceroid-lipofuscinosis neuronal 6 (19.5 kD) (Cln6andCalml4) alternative variant cSep08, mRNA.
<a href="#">Cln6andCalml4</a>	<a href="#">Cln6andCalml4.cSep08</a>	<a href="#">691455</a>	24314	678	7	165	ceroid-lipofuscinosis neuronal 6 (19.5 kD) (Cln6andCalml4) alternative variant cSep08, mRNA.
<a href="#">Clns1a</a>	<a href="#">Clns1a.bSep08</a>	<a href="#">65160</a>	20281	1456	1	241	chloride channel, nucleotide-sensitive, 1A (26.6 kD) (Clns1a) alternative variant bSep08, complete mRNA.
<a href="#">Clp1</a>	<a href="#">Clp1.bSep08</a>	<a href="#">311166</a>	3041	1229	2	346	CLP1, cleavage and polyadenylation factor I subunit, homolog (S. cerevisiae) (Clp1) alternative variant bSep08, mRNA.
<a href="#">Clp1</a>	<a href="#">Clp1.cSep08</a>	<a href="#">311166</a>	2683	343	2	85	CLP1, cleavage and polyadenylation factor I subunit, homolog (S. cerevisiae) (Clp1) alternative variant cSep08, mRNA.
<a href="#">Clp1</a>	<a href="#">Clp1.dSep08</a>	<a href="#">311166</a>	2629	432	2	78	CLP1, cleavage and polyadenylation factor I subunit, homolog (S. cerevisiae) (8.9 kD) (Clp1) alternative variant dSep08, mRNA.
<a href="#">Clpb</a>	<a href="#">Clpb.cSep08</a>	<a href="#">65041</a>	4110	311	2	69	ClpB caseinolytic peptidase B homolog (E. coli) (Clpb) alternative variant cSep08, mRNA.
<a href="#">Clpp</a>	<a href="#">Clpp.aSep08</a>	<a href="#">301117</a>	5679	1024	6	282	caseinolytic peptidase, ATP-dependent, proteolytic subunit homolog (E. coli) (Clpp) alternative variant aSep08, mRNA.
<a href="#">Clpp</a>	<a href="#">Clpp.bSep08</a>	<a href="#">301117</a>	2973	471	3	100	caseinolytic peptidase, ATP-dependent, proteolytic subunit homolog (E. coli) (Clpp) alternative variant bSep08, mRNA.
<a href="#">Clptm1</a>	<a href="#">Clptm1.bSep08</a>	<a href="#">292696</a>	849	770	2	110	cleft lip and palate associated transmembrane protein 1 (Clptm1) alternative variant bSep08, mRNA.
<a href="#">Clptm1</a>	<a href="#">Clptm1.cSep08</a>	<a href="#">292696</a>	11468	327	2	108	cleft lip and palate associated transmembrane protein 1 (Clptm1) alternative variant cSep08, mRNA.
<a href="#">Clptm1l</a>	<a href="#">Clptm1l.bSep08</a>	<a href="#">316916</a>	8379	1114	8	324	CLPTM1-like (Clptm1l) alternative variant bSep08, mRNA.
<a href="#">Clptm1l</a>	<a href="#">Clptm1l.cSep08</a>	<a href="#">316916</a>	9591	884	9	294	CLPTM1-like (Clptm1l) alternative variant cSep08, mRNA.

<a href="#">Clptm1l</a>	<a href="#">Clptm1l.dSep08</a>	<a href="#">316916</a>	7373	1173	7	190	CLPTM1-like (Clptm1l) alternative variant dSep08, mRNA.
<a href="#">Clptm1l</a>	<a href="#">Clptm1l.fSep08</a>	<a href="#">316916</a>	1879	801	2	42	CLPTM1-like (Clptm1l) alternative variant fSep08, mRNA.
<a href="#">Clpx</a>	<a href="#">Clpx.bSep08</a>	<a href="#">300786</a>	20267	728	6	159	caseinolytic peptidase X (E.coli) (Clpx) alternative variant bSep08, mRNA.
<a href="#">Clrn1</a>	<a href="#">Clrn1.bSep08</a>	<a href="#">261738</a>	49925	876	1	115	clarin 1 (13.1 kD) (Clrn1) alternative variant bSep08, mRNA.
<a href="#">Clspn</a>	<a href="#">Clspn.bSep08</a>	<a href="#">298534</a>	2848	1206	3	152	claspin homolog (Xenopus laevis) (Clspn) alternative variant bSep08, mRNA.
<a href="#">Clstn1</a>	<a href="#">Clstn1.aSep08</a>	<a href="#">313717</a>	45258	591	1	196	calsyntenin 1 (Clstn1) alternative variant aSep08, mRNA.
<a href="#">Clstn1</a>	<a href="#">Clstn1.bSep08</a>	<a href="#">313717</a>	45480	783		173	calsyntenin 1 (Clstn1) alternative variant bSep08, mRNA.
<a href="#">Clstn3</a>	<a href="#">Clstn3.bSep08</a>	<a href="#">171393</a>	20607	1800	6	547	calsyntenin 3 (Clstn3) alternative variant bSep08, mRNA.
<a href="#">Clstn3</a>	<a href="#">Clstn3.cSep08</a>	<a href="#">171393</a>	2802	958	5	315	calsyntenin 3 (Clstn3) alternative variant cSep08, mRNA.
<a href="#">Clstn3</a>	<a href="#">Clstn3.dSep08</a>	<a href="#">171393</a>	593	386	2	120	calsyntenin 3 CRA a (Clstn3) alternative variant dSep08, mRNA.
<a href="#">Clstn3</a>	<a href="#">Clstn3.eSep08</a>	<a href="#">171393</a>	830	458	2	72	calsyntenin 3 (Clstn3) alternative variant eSep08, mRNA.
<a href="#">Clstn3</a>	<a href="#">Clstn3.fSep08</a>	<a href="#">171393</a>	11675	741	2	48	calsyntenin 3 CRA a (Clstn3) alternative variant fSep08, mRNA.
<a href="#">Clta</a>	<a href="#">Clta.bSep08</a>	<a href="#">83800</a>	18030	1068	5	265	clathrin, light polypeptide (Lca) (Clta) alternative variant bSep08, mRNA.
<a href="#">Clta</a>	<a href="#">Clta.cSep08</a>	<a href="#">83800</a>	18457	1549	6	236	clathrin, light polypeptide (Lca) (25.6 kD) (Clta) alternative variant cSep08, mRNA.
<a href="#">Cltb</a>	<a href="#">Cltb.bSep08</a>	<a href="#">116561</a>	18146	1511	1	188	clathrin, light polypeptide (Lcb) (Cltb) alternative variant bSep08, mRNA.
<a href="#">Cltc</a>	<a href="#">Cltc.bSep08</a>	<a href="#">54241</a>	10112	2186	7	328	clathrin, heavy polypeptide (Hc) (Cltc) alternative variant bSep08, mRNA.
<a href="#">Cltc</a>	<a href="#">Cltc.dSep08</a>	<a href="#">54241</a>	2643	1111	2	51	clathrin, heavy polypeptide (Hc) (5.6 kD) (Cltc) alternative variant dSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.bSep08</a>	<a href="#">24854</a>	7469	1004	5	275	clusterin CRA a (Clu) alternative variant bSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.cSep08</a>	<a href="#">24854</a>	7176	735	5	221	clusterin CRA a (Clu) alternative variant cSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.dSep08</a>	<a href="#">24854</a>	2218	768	3	219	clusterin CRA a (Clu) alternative variant dSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.eSep08</a>	<a href="#">24854</a>	7346	732	4	208	clusterin CRA a (Clu) alternative variant eSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.fSep08</a>	<a href="#">24854</a>	7113	640	5	193	clusterin CRA a (Clu) alternative variant fSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.gSep08</a>	<a href="#">24854</a>	6747	618	5	167	clusterin (Clu) alternative variant gSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.hSep08</a>	<a href="#">24854</a>	3682	489	3	93	putative secreted or extracellular protein precursor of mammalian origin (10.4 kD) (Clu) alternative variant hSep08, complete mRNA.
<a href="#">Clu</a>	<a href="#">Clu.iSep08</a>	<a href="#">24854</a>	7142	732	5	119	clusterin CRA a (Clu) alternative variant iSep08, mRNA.
<a href="#">Clu</a>	<a href="#">Clu.jSep08</a>	<a href="#">24854</a>	7319	727	5	181	clusterin CRA a (Clu) alternative variant jSep08, mRNA.
<a href="#">Cluap1</a>	<a href="#">Cluap1.bSep08</a>	<a href="#">363544</a>	15992	1243	7	285	clusterin associated protein 1 (Cluap1) alternative variant bSep08, mRNA.
<a href="#">Cluap1</a>	<a href="#">Cluap1.cSep08</a>	<a href="#">363544</a>	15665	804	6	193	clusterin associated protein 1 (Cluap1) alternative variant cSep08, mRNA.
<a href="#">Cluap1</a>	<a href="#">Cluap1.dSep08</a>	<a href="#">363544</a>	16694	1031	5	172	clusterin associated protein 1 (20.3 kD) (Cluap1) alternative variant dSep08, mRNA.

<a href="#">Clybl</a>	<a href="#">Clybl.aSep08</a>	<a href="#">306198</a>	97328	1162	1	322	citrate lyase beta like (Clybl) alternative variant aSep08, mRNA.
<a href="#">Clybl</a>	<a href="#">Clybl.bSep08</a>	<a href="#">306198</a>	70666	379		126	citrate lyase beta like (Clybl) alternative variant bSep08, mRNA.
<a href="#">Cma1</a>	<a href="#">Cma1.aSep08</a>	<a href="#">25627</a>	3256	953		249	chymase 1, mast cell and similar to mast cell protease 1 (Cma1) alternative variant aSep08, mRNA.
<a href="#">Cma1</a>	<a href="#">Cma1.aSep08</a>	<a href="#">691558</a>	3256	953		249	chymase 1, mast cell and similar to mast cell protease 1 (Cma1) alternative variant aSep08, mRNA.
<a href="#">Cma1</a>	<a href="#">Cma1.cSep08</a>	<a href="#">25627</a>	913	602		132	chymase 1, mast cell and similar to mast cell protease 1 (Cma1) alternative variant cSep08, mRNA.
<a href="#">Cma1</a>	<a href="#">Cma1.cSep08</a>	<a href="#">691558</a>	913	602		132	chymase 1, mast cell and similar to mast cell protease 1 (Cma1) alternative variant cSep08, mRNA.
<a href="#">Cmah</a>	<a href="#">Cmah.bSep08</a>	<a href="#">361245</a>	7409	812	5	233	cytidine monophospho-N-acetylneuraminic acid hydroxylase (Cmah) alternative variant bSep08, mRNA.
<a href="#">Cmah</a>	<a href="#">Cmah.cSep08</a>	<a href="#">361245</a>	17435	874	6	187	cytidine monophospho-N-acetylneuraminic acid hydroxylase (Cmah) alternative variant cSep08, mRNA.
<a href="#">Cmas</a>	<a href="#">Cmas.bSep08</a>	<a href="#">312826</a>	3582	769	1	125	cytidine monophospho-N-acetylneuraminic acid synthetase (Cmas) alternative variant bSep08, mRNA.
<a href="#">Cmb1</a>	<a href="#">Cmb1.cSep08</a>	<a href="#">310201</a>	10742	720	2	70	carboxymethylenebutenolidase homolog (Pseudomonas) (Cmb1) alternative variant cSep08, mRNA.
<a href="#">Cmc1.0</a>	<a href="#">Cmc1.0.aSep08</a>		26002	631	1	80	putative protein of vertebrate origin (9.6 kD) (Cmc1.0) alternative variant aSep08, complete mRNA.
<a href="#">Cmc1.0</a>	<a href="#">Cmc1.0.bSep08</a>		26020	618	2	79	putative protein of eukaryotic origin (9.4 kD) (Cmc1.0) alternative variant bSep08, complete mRNA.
<a href="#">Cmip</a>	<a href="#">Cmip.aSep08</a>	<a href="#">292051</a>	81791	4147	20	675	c-Maf-inducing protein (Cmip) alternative variant aSep08, mRNA.
<a href="#">Cmip</a>	<a href="#">Cmip.bSep08</a>	<a href="#">292051</a>	8259	1210	2	134	c-Maf-inducing protein (14.6 kD) (Cmip) alternative variant bSep08, mRNA.
<a href="#">Cmip</a>	<a href="#">Cmip.cSep08</a>	<a href="#">292051</a>	23391	377	5	125	c-Maf-inducing protein (Cmip) alternative variant cSep08, mRNA.
<a href="#">Cmklr1</a>	<a href="#">Cmklr1.bSep08</a>	<a href="#">60669</a>	97696	511	3	38	chemokine-like receptor 1 (Cmklr1) alternative variant bSep08, mRNA.
<a href="#">Cmtm3</a>	<a href="#">Cmtm3.aSep08</a>	<a href="#">291813</a>	6331	982	1	184	marvel (20.3 kD) (Cmtm3) alternative variant aSep08, mRNA.
<a href="#">Cmtm4</a>	<a href="#">Cmtm4.aSep08</a>	<a href="#">498902</a>	43640	4557		208	4-like (22.8 kD) (Cmtm4) mRNA.
<a href="#">Cmtm6</a>	<a href="#">Cmtm6.aSep08</a>	<a href="#">316035</a>	18108	3254		201	marvel (Cmtm6) mRNA.
<a href="#">Cmtm7</a>	<a href="#">Cmtm7.bSep08</a>	<a href="#">501065</a>	2442	1147	2	61	putative protein (Cmtm7) alternative variant bSep08, mRNA.
<a href="#">Cnbp</a>	<a href="#">Cnbp.bSep08</a>	<a href="#">64530</a>	8788	1554	5	227	cellular nucleic acid binding protein (Cnbp) alternative variant bSep08, mRNA.
<a href="#">Cnbp</a>	<a href="#">Cnbp.cSep08</a>	<a href="#">64530</a>	8066	835	5	217	cellular nucleic acid binding protein (Cnbp) alternative variant cSep08, mRNA.
<a href="#">Cnbp</a>	<a href="#">Cnbp.dSep08</a>	<a href="#">64530</a>	8377	1196	5	186	cellular nucleic acid binding protein (Cnbp) alternative variant dSep08, mRNA.
<a href="#">Cnbp</a>	<a href="#">Cnbp.eSep08</a>	<a href="#">64530</a>	8112	864	5	175	cellular nucleic acid binding protein (Cnbp) alternative variant eSep08, mRNA.

<a href="#">Cnbp</a>	<a href="#">Cnbp.gSep08</a>	<a href="#">64530</a>	6388	442	3	117	cellular nucleic acid binding protein (Cnbp) alternative variant gSep08, mRNA.
<a href="#">Cnbp</a>	<a href="#">Cnbp.hSep08</a>	<a href="#">64530</a>	7981	902	4	94	cellular nucleic acid binding protein (Cnbp) alternative variant hSep08, mRNA.
<a href="#">Cndp2</a>	<a href="#">Cndp2.bSep08</a>	<a href="#">291394</a>	8882	825	6	247	CNDP dipeptidase 2 (metallopeptidase M20 family) (Cndp2) alternative variant bSep08, mRNA.
<a href="#">Cndp2</a>	<a href="#">Cndp2.cSep08</a>	<a href="#">291394</a>	3893	1279	5	238	CNDP dipeptidase 2 (metallopeptidase M20 family) (Cndp2) alternative variant cSep08, mRNA.
<a href="#">Cndp2</a>	<a href="#">Cndp2.dSep08</a>	<a href="#">291394</a>	6974	801	5	134	CNDP dipeptidase 2 (metallopeptidase M20 family) (Cndp2) alternative variant dSep08, mRNA.
<a href="#">Cngb1</a>	<a href="#">Cngb1.aSep08</a>	<a href="#">83686</a>	5011	399		133	cyclic nucleotide gated channel beta 1 (Cngb1) mRNA.
<a href="#">CNH.0</a>	<a href="#">CNH.0.aSep08</a>		4244	2422	5	467	misshapen-like kinase 1 (52.2 kD) (CNH.0) alternative variant aSep08, mRNA.
<a href="#">CNH.0</a>	<a href="#">CNH.0.bSep08</a>		2266	1433	3	215	misshapen-like kinase 1 (CNH.0) alternative variant bSep08, mRNA.
<a href="#">CNH.0</a>	<a href="#">CNH.0.cSep08</a>		987	393	2	130	misshapen-like kinase 1 (CNH.0) alternative variant cSep08, mRNA.
<a href="#">CNH.0</a>	<a href="#">CNH.0.dSep08</a>		824	733	2	67	misshapen-like kinase 1 (CNH.0) alternative variant dSep08, mRNA.
<a href="#">Cnksr2</a>	<a href="#">Cnksr2.cSep08</a>	<a href="#">59322</a>	65076	566		188	connector enhancer of kinase suppressor of Ras 2 (Cnksr2) alternative variant cSep08, mRNA.
<a href="#">Cnksr3</a>	<a href="#">Cnksr3.bSep08</a>	<a href="#">308113</a>	10979	3206	4	267	cnksr family member 3 (Cnksr3) alternative variant bSep08, mRNA.
<a href="#">Cnksr3</a>	<a href="#">Cnksr3.cSep08</a>	<a href="#">308113</a>	16561	478	4	158	cnksr family member 3 (Cnksr3) alternative variant cSep08, mRNA.
<a href="#">cNMP_binding.0</a>	<a href="#">cNMP_binding.0.aSep08</a>		816	374		124	cyclic nucleotide-binding (cNMP_binding.0) mRNA.
<a href="#">cNMP_binding.1</a>	<a href="#">cNMP_binding.1.aSep08</a>		6394	812		270	cyclic nucleotide-binding (cNMP_binding.1) mRNA.
<a href="#">Cnn1</a>	<a href="#">Cnn1.bSep08</a>	<a href="#">65204</a>	968	882	2	125	calponin 1 (Cnn1) alternative variant bSep08, mRNA.
<a href="#">Cnn1</a>	<a href="#">Cnn1.cSep08</a>	<a href="#">65204</a>	3285	257	2	85	calponin 1 (Cnn1) alternative variant cSep08, mRNA.
<a href="#">Cnn1</a>	<a href="#">Cnn1.dSep08</a>	<a href="#">65204</a>	3734	228	3	76	calponin 1 (Cnn1) alternative variant dSep08, mRNA.
<a href="#">Cnn3</a>	<a href="#">Cnn3.bSep08</a>	<a href="#">54321</a>	7436	1239	5	288	calponin 3, acidic (Cnn3) alternative variant bSep08, mRNA.
<a href="#">Cnnm1</a>	<a href="#">Cnnm1.bSep08</a>	<a href="#">309387</a>	22338	569	6	189	cyclin M1 (Cnnm1) alternative variant bSep08, mRNA.
<a href="#">Cnnm2</a>	<a href="#">Cnnm2.bSep08</a>	<a href="#">294014</a>	10390	1215	3	161	cyclin M2 (Cnnm2) alternative variant bSep08, mRNA.
<a href="#">Cnnm3</a>	<a href="#">Cnnm3.bSep08</a>	<a href="#">301345</a>	3361	881	3	100	cyclin M3 (Cnnm3) alternative variant bSep08, mRNA.
<a href="#">Cnnm4</a>	<a href="#">Cnnm4.aSep08</a>	<a href="#">363216</a>	9168	2724	4	148	cyclin M4 (Cnnm4) alternative variant aSep08, mRNA.
<a href="#">Cnnm4</a>	<a href="#">Cnnm4.bSep08</a>	<a href="#">363216</a>	9654	511	3	105	cyclin M4 (Cnnm4) alternative variant bSep08, mRNA.
<a href="#">Cnnm4</a>	<a href="#">Cnnm4.cSep08</a>	<a href="#">363216</a>	1598	386	2	79	cyclin M4 (Cnnm4) alternative variant cSep08, mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.aSep08</a>	<a href="#">299805</a>	91144	2643	15	455	transcription complex CRA b (50.2 kD) (Cnot2) alternative variant aSep08, complete mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.cSep08</a>	<a href="#">299805</a>	84556	1426	13	349	transcription complex (38.4 kD) (Cnot2) alternative variant cSep08, mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.dSep08</a>	<a href="#">299805</a>	78739	1299	11	256	transcription complex CRA I (28.0 kD) (Cnot2) alternative variant dSep08, mRNA.

<a href="#">Cnot2</a>	<a href="#">Cnot2.eSep08</a>	<a href="#">299805</a>	48893	776	7	191	transcription complex (Cnot2) alternative variant eSep08, mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.fSep08</a>	<a href="#">299805</a>	71447	821	7	190	transcription complex (20.7 kD) (Cnot2) alternative variant fSep08, mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.gSep08</a>	<a href="#">299805</a>	60345	574	5	69	transcription complex (Cnot2) alternative variant gSep08, mRNA.
<a href="#">Cnot2</a>	<a href="#">Cnot2.hSep08</a>	<a href="#">299805</a>	6553	918	2	56	transcription complex (6.9 kD) (Cnot2) alternative variant hSep08, mRNA.
<a href="#">Cnot3</a>	<a href="#">Cnot3.bSep08</a>	<a href="#">308311</a>	8070	926	8	198	CCR4-NOT transcription complex, subunit 3 (Cnot3) alternative variant bSep08, mRNA.
<a href="#">Cnot3</a>	<a href="#">Cnot3.dSep08</a>	<a href="#">308311</a>	1537	544	3	31	CCR4-NOT transcription complex, subunit 3 (Cnot3) alternative variant dSep08, mRNA.
<a href="#">Cnot4</a>	<a href="#">Cnot4.bSep08</a>	<a href="#">312227</a>	7245	1100	3	173	CCR4-NOT transcription complex, subunit 4 (18.6 kD) (Cnot4) alternative variant bSep08, mRNA.
<a href="#">Cnot6</a>	<a href="#">Cnot6.bSep08</a>	<a href="#">287249</a>	11566	5173	1	373	CCR4-NOT transcription complex, subunit 6 (Cnot6) alternative variant bSep08, mRNA.
<a href="#">Cnot6</a>	<a href="#">Cnot6.cSep08</a>	<a href="#">287249</a>	3908	519	2	97	CCR4-NOT transcription complex, subunit 6 (Cnot6) alternative variant cSep08, mRNA.
<a href="#">Cnot7</a>	<a href="#">Cnot7.bSep08</a>	<a href="#">306492</a>	2732	570	2	116	CCR4-NOT transcription complex, subunit 7 (Cnot7) alternative variant bSep08, mRNA.
<a href="#">Cnot7</a>	<a href="#">Cnot7.cSep08</a>	<a href="#">306492</a>	2737	580	2	71	CCR4-NOT transcription complex, subunit 7 (Cnot7) alternative variant cSep08, mRNA.
<a href="#">Cnot7</a>	<a href="#">Cnot7.dSep08</a>	<a href="#">306492</a>	4998	786	3	50	CCR4-NOT transcription complex, subunit 7 (5.6 kD) (Cnot7) alternative variant dSep08, mRNA.
<a href="#">Cnot8</a>	<a href="#">Cnot8.bSep08</a>	<a href="#">363603</a>	9634	896	6	242	transcription complex (Cnot8) alternative variant bSep08, mRNA.
<a href="#">Cnot8</a>	<a href="#">Cnot8.cSep08</a>	<a href="#">363603</a>	2406	583	3	171	transcription complex (Cnot8) alternative variant cSep08, mRNA.
<a href="#">Cnot8</a>	<a href="#">Cnot8.dSep08</a>	<a href="#">363603</a>	5140	726	3	136	CRA d like (Cnot8) alternative variant dSep08, mRNA.
<a href="#">Cnot8</a>	<a href="#">Cnot8.fSep08</a>	<a href="#">363603</a>	2037	703	2	55	oculocerebrorrenal syndrome of Lowe like (6.2 kD) (Cnot8) alternative variant fSep08, mRNA.
<a href="#">Cnot10</a>	<a href="#">Cnot10.aSep08</a>	<a href="#">316034</a>	34359	2497	13	739	CCR4-NOT transcription complex, subunit 10 (Cnot10) alternative variant aSep08, mRNA.
<a href="#">Cnot10</a>	<a href="#">Cnot10.bSep08</a>	<a href="#">316034</a>	22296	1057	9	251	CCR4-NOT transcription complex, subunit 10 (Cnot10) alternative variant bSep08, mRNA.
<a href="#">Cnot10</a>	<a href="#">Cnot10.cSep08</a>	<a href="#">316034</a>	17502	1408	6	129	CCR4-NOT transcription complex, subunit 10 (Cnot10) alternative variant cSep08, mRNA.
<a href="#">Cnp</a>	<a href="#">Cnp.bSep08</a>	<a href="#">25275</a>	4197	1294	3	266	2',3'-cyclic nucleotide 3' phosphodiesterase (30.3 kD) (Cnp) alternative variant bSep08, mRNA.
<a href="#">Cnp</a>	<a href="#">Cnp.dSep08</a>	<a href="#">25275</a>	866	481	2	160	2',3'-cyclic nucleotide 3' phosphodiesterase (Cnp) alternative variant dSep08, mRNA.
<a href="#">Cnpy2</a>	<a href="#">Cnpy2.aSep08</a>	<a href="#">685814</a>	4168	886	5	274	canopy 2 homolog (zebrafish) (Cnpy2) alternative variant aSep08, mRNA.
<a href="#">Cnpy3</a>	<a href="#">Cnpy3.aSep08</a>	<a href="#">685174</a>	14599	1800	5	276	canopy 3 homolog (zebrafish) (30.6 kD) (Cnpy3) alternative variant aSep08, complete mRNA.
<a href="#">Cnpy3</a>	<a href="#">Cnpy3.bSep08</a>	<a href="#">685174</a>	13483	587	4	102	canopy 3 homolog (zebrafish) (Cnpy3) alternative variant bSep08, mRNA.



<a href="#">Cnpy3</a>	<a href="#">Cnpy3.cSep08</a>	<a href="#">685174</a>	4467	948	3	20	canopy 3 homolog (zebrafish) (Cnpy3) alternative variant cSep08, mRNA.
<a href="#">Cntfr</a>	<a href="#">Cntfr.aSep08</a>	<a href="#">313173</a>	30089	1953	1	373	ciliary neurotrophic factor receptor (41.0 kD) (Cntfr) alternative variant aSep08, mRNA.
<a href="#">Cntn1</a>	<a href="#">Cntn1.bSep08</a>	<a href="#">117258</a>	5821	635	2	51	contactin 1 (5.6 kD) (Cntn1) alternative variant bSep08, mRNA.
<a href="#">Cntn2</a>	<a href="#">Cntn2.bSep08</a>	<a href="#">25356</a>	13777	533	1	97	contactin 2 (Cntn2) alternative variant bSep08, mRNA.
<a href="#">Cntn3</a>	<a href="#">Cntn3.bSep08</a>	<a href="#">54279</a>	23926	419		139	contactin 3 (Cntn3) alternative variant bSep08, mRNA.
<a href="#">Cntn4</a>	<a href="#">Cntn4.bSep08</a>	<a href="#">116658</a>	178048	496	1	91	contactin 4 (10.3 kD) (Cntn4) alternative variant bSep08, mRNA.
<a href="#">Cntnap1</a>	<a href="#">Cntnap1.aSep08</a>	<a href="#">84008</a>	9917	3143		1047	contactin associated protein 1 (Cntnap1) mRNA.
<a href="#">Cntnap4</a>	<a href="#">Cntnap4.bSep08</a>	<a href="#">307865</a>	11478	601		200	contactin associated protein-like 4 (Cntnap4) alternative variant bSep08, mRNA.
<a href="#">Cntrob</a>	<a href="#">Cntrob.bSep08</a>	<a href="#">303240</a>	996	719	2	53	centrobin, centrosomal BRCA2 interacting protein (5.7 kD) (Cntrob) alternative variant bSep08, mRNA.
<a href="#">Coasy</a>	<a href="#">Coasy.bSep08</a>	<a href="#">287711</a>	3838	1779	3	398	coenzyme A synthase (Coasy) alternative variant bSep08, mRNA.
<a href="#">Coasy</a>	<a href="#">Coasy.dSep08</a>	<a href="#">287711</a>	858	755	2	85	coenzyme A synthase (9.6 kD) (Coasy) alternative variant dSep08, mRNA.
<a href="#">Cobl</a>	<a href="#">Cobl.bSep08</a>	<a href="#">305497</a>	89778	2585	4	861	cordons-bleu (Cobl) alternative variant bSep08, mRNA.
<a href="#">Cobl</a>	<a href="#">Cobl.cSep08</a>	<a href="#">305497</a>	90850	761	4	227	cordons-bleu (Cobl) alternative variant cSep08, mRNA.
<a href="#">Cobl</a>	<a href="#">Cobl.dSep08</a>	<a href="#">305497</a>	74204	631	3	150	cordons-bleu (Cobl) alternative variant dSep08, mRNA.
<a href="#">Cobra1andRGD1308019</a>	<a href="#">Cobra1andRGD1308019.bSep08</a>	<a href="#">311795</a>	16356	2672	12	475	similar to hypothetical protein FLJ20245 and cofactor of BRCA1 (53.9 kD) (Cobra1andRGD1308019) alternative variant bSep08, complete mRNA.
<a href="#">Cobra1andRGD1308019</a>	<a href="#">Cobra1andRGD1308019.bSep08</a>	<a href="#">311796</a>	16356	2672	12	475	similar to hypothetical protein FLJ20245 and cofactor of BRCA1 (53.9 kD) (Cobra1andRGD1308019) alternative variant bSep08, complete mRNA.
<a href="#">Cobra1andRGD1308019</a>	<a href="#">Cobra1andRGD1308019.cSep08</a>	<a href="#">311795</a>	2262	1798	3	132	similar to hypothetical protein FLJ20245 and cofactor of BRCA1 (14.4 kD) (Cobra1andRGD1308019) alternative variant cSep08, mRNA.
<a href="#">Cobra1andRGD1308019</a>	<a href="#">Cobra1andRGD1308019.cSep08</a>	<a href="#">311796</a>	2262	1798	3	132	similar to hypothetical protein FLJ20245 and cofactor of BRCA1 (14.4 kD) (Cobra1andRGD1308019) alternative variant cSep08, mRNA.
<a href="#">Coch</a>	<a href="#">Coch.aSep08</a>	<a href="#">362735</a>	13428	2495	7	552	coagulation factor C precursor (Coch) alternative variant aSep08, complete mRNA.
<a href="#">Coch</a>	<a href="#">Coch.bSep08</a>	<a href="#">362735</a>	13350	3319	6	357	coagulation factor C homolog CRA a (39.2 kD) (Coch) alternative variant bSep08, complete mRNA.
<a href="#">Cog1</a>	<a href="#">Cog1.aSep08</a>	<a href="#">303652</a>	13752	3947	14	979	component of oligomeric golgi complex 1 (Cog1) alternative variant aSep08, mRNA.
<a href="#">Cog1</a>	<a href="#">Cog1.bSep08</a>	<a href="#">303652</a>	4113	791	5	263	component of oligomeric golgi complex 1 (Cog1) alternative variant bSep08, mRNA.
<a href="#">Cog1</a>	<a href="#">Cog1.dSep08</a>	<a href="#">303652</a>	2895	543	6	109	component of oligomeric golgi complex 1 (Cog1) alternative variant dSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.bSep08</a>	<a href="#">690961</a>	6322	1786	3	328	putative protein (Cog2) alternative variant bSep08, mRNA.

<a href="#">Cog2</a>	<a href="#">Cog2.cSep08</a>	<a href="#">690961</a>	3316	782	3	175	component of oligomeric golgi complex 2 (Cog2) alternative variant cSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.dSep08</a>	<a href="#">690961</a>	9563	542	5	162	component of oligomeric golgi complex 2 (Cog2) alternative variant dSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.eSep08</a>	<a href="#">690961</a>	1304	744	2	155	component of oligomeric golgi complex 2 (16.9 kD) (Cog2) alternative variant eSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.fSep08</a>	<a href="#">690961</a>	7131	683	6	139	component of oligomeric golgi complex 2 (Cog2) alternative variant fSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.gSep08</a>	<a href="#">690961</a>	6530	423	4	118	component of oligomeric golgi complex 2 (Cog2) alternative variant gSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.hSep08</a>	<a href="#">690961</a>	1661	634	2	74	component of oligomeric golgi complex 2 (Cog2) alternative variant hSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.iSep08</a>	<a href="#">690961</a>	1954	263	2	35	component of oligomeric golgi complex 2 (Cog2) alternative variant iSep08, mRNA.
<a href="#">Cog2</a>	<a href="#">Cog2.jSep08</a>	<a href="#">690961</a>	451	354	2	52	putative protein (Cog2) alternative variant jSep08, mRNA.
<a href="#">Cog3</a>	<a href="#">Cog3.bSep08</a>	<a href="#">361073</a>	31026	2548	15	619	component of oligomeric golgi complex 3 (Cog3) alternative variant bSep08, mRNA.
<a href="#">Cog3</a>	<a href="#">Cog3.cSep08</a>	<a href="#">361073</a>	10238	892	4		
<a href="#">Cog3</a>	<a href="#">Cog3.dSep08</a>	<a href="#">361073</a>	3844	344	3	14	component of oligomeric golgi complex 3 (Cog3) alternative variant dSep08, mRNA.
<a href="#">Cog4</a>	<a href="#">Cog4.bSep08</a>	<a href="#">361407</a>	34460	3038	19	716	component of oligomeric golgi complex 4 CRA d (80.3 kD) (Cog4) alternative variant bSep08, complete mRNA.
<a href="#">Cog4</a>	<a href="#">Cog4.cSep08</a>	<a href="#">361407</a>	11278	1446	6	244	component of oligomeric golgi complex 4 CRA d (Cog4) alternative variant cSep08, mRNA.
<a href="#">Cog4</a>	<a href="#">Cog4.dSep08</a>	<a href="#">361407</a>	1419	838	3	61	component of oligomeric golgi complex 4 CRA d (Cog4) alternative variant dSep08, mRNA.
<a href="#">Cog6</a>	<a href="#">Cog6.bSep08</a>	<a href="#">310411</a>	15845	710	8	121	component of oligomeric golgi complex 6 (Cog6) alternative variant bSep08, mRNA.
<a href="#">Cog8</a>	<a href="#">Cog8.bSep08</a>	<a href="#">291990</a>	2517	466	1	155	component of oligomeric golgi complex 8 (Cog8) alternative variant bSep08, mRNA.
<a href="#">Coil</a>	<a href="#">Coil.bSep08</a>	<a href="#">50998</a>	9265	1700		400	coilin (43.5 kD) (Coil) alternative variant bSep08, complete mRNA.
<a href="#">Col1a1</a>	<a href="#">Col1a1.bSep08</a>	<a href="#">29393</a>	5528	1794	11	517	collagen, type I, alpha 1 (Col1a1) alternative variant bSep08, mRNA.
<a href="#">Col1a1</a>	<a href="#">Col1a1.cSep08</a>	<a href="#">29393</a>	1752	648	6	215	collagen, type I, alpha 1 (Col1a1) alternative variant cSep08, mRNA.
<a href="#">Col1a2</a>	<a href="#">Col1a2.aSep08</a>	<a href="#">84352</a>	35582	5188	5	1395	collagen, type I, alpha 2 (132.0 kD) (Col1a2) alternative variant aSep08, mRNA.
<a href="#">Col1a2</a>	<a href="#">Col1a2.bSep08</a>	<a href="#">84352</a>	19542	3233		789	collagen, type I, alpha 2 (Col1a2) alternative variant bSep08, mRNA.
<a href="#">Col1a2</a>	<a href="#">Col1a2.cSep08</a>	<a href="#">84352</a>	13677	1485		495	collagen, type I, alpha 2 (Col1a2) alternative variant cSep08, mRNA.
<a href="#">Col1a2</a>	<a href="#">Col1a2.dSep08</a>	<a href="#">84352</a>	12488	991		121	collagen, type I, alpha 2 (Col1a2) alternative variant dSep08, mRNA.
<a href="#">Col1a2</a>	<a href="#">Col1a2.eSep08</a>	<a href="#">84352</a>	2542	1496	1	101	collagen, type I, alpha 2 (10.8 kD) (Col1a2) alternative variant eSep08, mRNA.

<a href="#">Col2a1</a>	<a href="#">Col2a1.aSep08</a>	<a href="#">25412</a>	14169	3854	36	1136	collagen, type II, alpha 1 (Col2a1) alternative variant aSep08, mRNA.
<a href="#">Col2a1</a>	<a href="#">Col2a1.bSep08</a>	<a href="#">25412</a>	1466	1020	2	194	collagen, type II, alpha 1 (Col2a1) alternative variant bSep08, mRNA.
<a href="#">Col2a1</a>	<a href="#">Col2a1.dSep08</a>	<a href="#">25412</a>	5768	364	6	121	collagen, type II, alpha 1 (Col2a1) alternative variant dSep08, mRNA.
<a href="#">Col2a1</a>	<a href="#">Col2a1.eSep08</a>	<a href="#">25412</a>	9802	535	3	33	collagen, type II, alpha 1 (Col2a1) alternative variant eSep08, mRNA.
<a href="#">Col3a1</a>	<a href="#">Col3a1.bSep08</a>	<a href="#">84032</a>	4198	870	10	234	collagen, type III, alpha 1 (Col3a1) alternative variant bSep08, mRNA.
<a href="#">Col3a1</a>	<a href="#">Col3a1.cSep08</a>	<a href="#">84032</a>	1041	719	2	141	collagen, type III, alpha 1 (Col3a1) alternative variant cSep08, mRNA.
<a href="#">Col3a1</a>	<a href="#">Col3a1.eSep08</a>	<a href="#">84032</a>	759	599	2	43	collagen, type III, alpha 1 (Col3a1) alternative variant eSep08, mRNA.
<a href="#">Col4a1</a>	<a href="#">Col4a1.aSep08</a>	<a href="#">290905</a>	33953	5063	31	1230	alpha-1 type IV collagen (Col4a1) alternative variant aSep08, mRNA.
<a href="#">Col4a1</a>	<a href="#">Col4a1.bSep08</a>	<a href="#">290905</a>	1697	436	5	145	procollagen type IV alpha 1 (Col4a1) alternative variant bSep08, mRNA.
<a href="#">Col4a1</a>	<a href="#">Col4a1.cSep08</a>	<a href="#">290905</a>	3332	720	3	114	alpha-1 type IV collagen (12.6 kD) (Col4a1) alternative variant cSep08, mRNA.
<a href="#">Col4a1</a>	<a href="#">Col4a1.dSep08</a>	<a href="#">290905</a>	641	514	2	91	collagen triple helix repeat (Col4a1) alternative variant dSep08, mRNA.
<a href="#">Col4a2_predicted</a>	<a href="#">Col4a2_predicted.aSep08</a>	<a href="#">306628</a>	105913	1119		280	procollagen, type IV, alpha 2 (predicted) (Col4a2_predicted) mRNA.
<a href="#">Col4a3bp</a>	<a href="#">Col4a3bp.bSep08</a>	<a href="#">365652</a>	25703	757	1	169	collagen, type IV, alpha 3 (Goodpasture antigen) binding protein (Col4a3bp) alternative variant bSep08, mRNA.
<a href="#">Col4a5</a>	<a href="#">Col4a5.aSep08</a>	<a href="#">363457</a>	42559	3237	18	715	type IV alpha (Col4a5) alternative variant aSep08, mRNA.
<a href="#">Col4a5</a>	<a href="#">Col4a5.bSep08</a>	<a href="#">363457</a>	10169	419	4	129	collagen triple helix repeat (Col4a5) alternative variant bSep08, mRNA.
<a href="#">Col4a5</a>	<a href="#">Col4a5.cSep08</a>	<a href="#">363457</a>	2335	412	2	96	collagen triple helix repeat (Col4a5) alternative variant cSep08, mRNA.
<a href="#">Col5a1</a>	<a href="#">Col5a1.aSep08</a>	<a href="#">85490</a>	45811	5329	1	946	collagen, type V, alpha 1 (Col5a1) alternative variant aSep08, mRNA.
<a href="#">Col5a1</a>	<a href="#">Col5a1.bSep08</a>	<a href="#">85490</a>	17010	1433	1	339	collagen, type V, alpha 1 (Col5a1) alternative variant bSep08, mRNA.
<a href="#">Col5a2</a>	<a href="#">Col5a2.aSep08</a>	<a href="#">85250</a>	29684	4284	30	954	collagen, type V, alpha 2 (Col5a2) alternative variant aSep08, mRNA.
<a href="#">Col5a2</a>	<a href="#">Col5a2.bSep08</a>	<a href="#">85250</a>	27238	1783	15	533	collagen, type V, alpha 2 (Col5a2) alternative variant bSep08, mRNA.
<a href="#">Col5a3</a>	<a href="#">Col5a3.bSep08</a>	<a href="#">60379</a>	6486	474	1	106	collagen, type V, alpha 3 (Col5a3) alternative variant bSep08, mRNA.
<a href="#">Col6a1</a>	<a href="#">Col6a1.aSep08</a>	<a href="#">294337</a>	10713	2932	23	706	collagen, type VI, alpha 1 (Col6a1) alternative variant aSep08, mRNA.
<a href="#">Col6a1</a>	<a href="#">Col6a1.bSep08</a>	<a href="#">294337</a>	1636	747	4	248	collagen, type VI, alpha 1 (Col6a1) alternative variant bSep08, mRNA.
<a href="#">Col6a1</a>	<a href="#">Col6a1.cSep08</a>	<a href="#">294337</a>	4005	804	9	163	collagen, type VI, alpha 1 (Col6a1) alternative variant cSep08, mRNA.

<a href="#">Col6a2</a>	<a href="#">Col6a2.aSep08</a>	<a href="#">361821</a>	19571	3604	13	799	collagen, type VI, alpha 2 (Col6a2) alternative variant aSep08, mRNA.
<a href="#">Col6a2</a>	<a href="#">Col6a2.bSep08</a>	<a href="#">361821</a>	7084	739	2	246	collagen, type VI, alpha 2 (Col6a2) alternative variant bSep08, mRNA.
<a href="#">Col6a2</a>	<a href="#">Col6a2.cSep08</a>	<a href="#">361821</a>	5741	1709	1	158	collagen, type VI, alpha 2 (Col6a2) alternative variant cSep08, mRNA.
<a href="#">Col6a3</a>	<a href="#">Col6a3.bSep08</a>	<a href="#">367313</a>	4991	337	6	112	procollagen, type VI, alpha 3 (Col6a3) alternative variant bSep08, mRNA.
<a href="#">Col7a1</a>	<a href="#">Col7a1.aSep08</a>	<a href="#">301012</a>	1170	665		107	procollagen, type VII, alpha 1 (Col7a1) mRNA.
<a href="#">Col9a1</a>	<a href="#">Col9a1.aSep08</a>	<a href="#">305104</a>	63102	1786		378	collagen, type IX, alpha 1 (Col9a1) alternative variant aSep08, mRNA.
<a href="#">Col9a1</a>	<a href="#">Col9a1.bSep08</a>	<a href="#">305104</a>	41836	1306	1	254	collagen, type IX, alpha 1 (Col9a1) alternative variant bSep08, mRNA.
<a href="#">Col9a1</a>	<a href="#">Col9a1.cSep08</a>	<a href="#">305104</a>	16894	1409	1	201	collagen, type IX, alpha 1 (Col9a1) alternative variant cSep08, mRNA.
<a href="#">Col9a1</a>	<a href="#">Col9a1.dSep08</a>	<a href="#">305104</a>	7384	566	1	110	collagen, type IX, alpha 1 (Col9a1) alternative variant dSep08, mRNA.
<a href="#">Col9a2</a>	<a href="#">Col9a2.aSep08</a>	<a href="#">362584</a>	4487	1774	12	286	collagen, type IX, alpha 2 (32.3 kD) (Col9a2) alternative variant aSep08, mRNA.
<a href="#">Col9a2</a>	<a href="#">Col9a2.bSep08</a>	<a href="#">362584</a>	1489	915	3	138	collagen, type IX, alpha 2 (Col9a2) alternative variant bSep08, mRNA.
<a href="#">Col9a3</a>	<a href="#">Col9a3.aSep08</a>	<a href="#">362285</a>	11320	1194	11	230	procollagen, type IX, alpha 3 (Col9a3) alternative variant aSep08, mRNA.
<a href="#">Col9a3</a>	<a href="#">Col9a3.cSep08</a>	<a href="#">362285</a>	2718	704	5	159	procollagen, type IX, alpha 3 (Col9a3) alternative variant cSep08, mRNA.
<a href="#">Col11a1</a>	<a href="#">Col11a1.aSep08</a>	<a href="#">25654</a>	27124	2766		426	collagen, type XI, alpha 1 (Col11a1) mRNA.
<a href="#">Col11a2</a>	<a href="#">Col11a2.bSep08</a>	<a href="#">294279</a>	3616	547	10	181	procollagen, type XI, alpha 2 (Col11a2) alternative variant bSep08, mRNA.
<a href="#">Col11a2</a>	<a href="#">Col11a2.cSep08</a>	<a href="#">294279</a>	3893	415	6	138	procollagen, type XI, alpha 2 (Col11a2) alternative variant cSep08, mRNA.
<a href="#">Col11a2</a>	<a href="#">Col11a2.dSep08</a>	<a href="#">294279</a>	2542	391	3	130	procollagen, type XI, alpha 2 (Col11a2) alternative variant dSep08, mRNA.
<a href="#">Col11a2</a>	<a href="#">Col11a2.eSep08</a>	<a href="#">294279</a>	1189	608	6	129	procollagen, type XI, alpha 2 (Col11a2) alternative variant eSep08, mRNA.
<a href="#">Col12a1</a>	<a href="#">Col12a1.aSep08</a>	<a href="#">25683</a>	78357	1778	1	473	collagen, type XII, alpha 1 (Col12a1) alternative variant aSep08, mRNA.
<a href="#">Col12a1</a>	<a href="#">Col12a1.bSep08</a>	<a href="#">25683</a>	16586	1691	8	293	collagen, type XII, alpha 1 (Col12a1) alternative variant bSep08, mRNA.
<a href="#">Col12a1</a>	<a href="#">Col12a1.cSep08</a>	<a href="#">25683</a>	2563	1529	1	72	collagen, type XII, alpha 1 (Col12a1) alternative variant cSep08, mRNA.
<a href="#">Col13a1</a>	<a href="#">Col13a1.aSep08</a>	<a href="#">499431</a>	19798	837		121	collagen, type XIII, alpha 1 (Col13a1) mRNA.
<a href="#">Col14a1</a>	<a href="#">Col14a1.bSep08</a>	<a href="#">314981</a>	23796	1385	3	202	collagen, type XIV, alpha 1 (Col14a1) alternative variant bSep08, mRNA.
<a href="#">Col14a1</a>	<a href="#">Col14a1.cSep08</a>	<a href="#">314981</a>	19817	1972	1	119	collagen, type XIV, alpha 1 (Col14a1) alternative variant cSep08, mRNA.
<a href="#">Col15a1</a>	<a href="#">Col15a1.aSep08</a>	<a href="#">298069</a>	43296	3136		755	collagen, type XV, alpha 1 (Col15a1) alternative variant aSep08, mRNA.

<a href="#">Col15a1</a>	<a href="#">Col15a1.bSep08</a>	<a href="#">298069</a>	104727	1772		349	collagen, type XV, alpha 1 (Col15a1) alternative variant bSep08, mRNA.
<a href="#">Col15a1</a>	<a href="#">Col15a1.cSep08</a>	<a href="#">298069</a>	39775	801		266	collagen, type XV, alpha 1 (Col15a1) alternative variant cSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.aSep08</a>	<a href="#">366474</a>	15150	2050	19	415	type XVI alpha 1 (39.9 kD) (Col16a1) alternative variant aSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.bSep08</a>	<a href="#">366474</a>	9388	711	13	236	procollagen type xvi alpha 1 (Col16a1) alternative variant bSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.cSep08</a>	<a href="#">366474</a>	4390	389	7	129	type XVI alpha 1 (Col16a1) alternative variant cSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.eSep08</a>	<a href="#">366474</a>	2708	578	3	102	collagen triple helix repeat (Col16a1) alternative variant eSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.fSep08</a>	<a href="#">366474</a>	1413	726	2	65	putative protein (7.0 kD) (Col16a1) alternative variant fSep08, mRNA.
<a href="#">Col16a1</a>	<a href="#">Col16a1.gSep08</a>	<a href="#">366474</a>	1780	731	4	31	putative protein (Col16a1) alternative variant gSep08, mRNA.
<a href="#">Col17a1</a>	<a href="#">Col17a1.bSep08</a>	<a href="#">294027</a>	2020	766	5	254	collagen, type XVII, alpha 1 (Col17a1) alternative variant bSep08, mRNA.
<a href="#">Col17a1</a>	<a href="#">Col17a1.cSep08</a>	<a href="#">294027</a>	3229	693	7	230	collagen, type XVII, alpha 1 (Col17a1) alternative variant cSep08, mRNA.
<a href="#">Col18a1</a>	<a href="#">Col18a1.aSep08</a>	<a href="#">85251</a>	14312	2672	1	561	collagen, type XVIII, alpha 1 (Col18a1) alternative variant aSep08, mRNA.
<a href="#">Col18a1</a>	<a href="#">Col18a1.bSep08</a>	<a href="#">85251</a>	109979	1926		509	collagen, type XVIII, alpha 1 (Col18a1) alternative variant bSep08, complete mRNA.
<a href="#">Col18a1</a>	<a href="#">Col18a1.cSep08</a>	<a href="#">85251</a>	6836	1832	1	440	collagen, type XVIII, alpha 1 (Col18a1) alternative variant cSep08, mRNA.
<a href="#">Col18a1</a>	<a href="#">Col18a1.dSep08</a>	<a href="#">85251</a>	19240	1319		326	collagen, type XVIII, alpha 1 (Col18a1) alternative variant dSep08, mRNA.
<a href="#">Col20a1</a>	<a href="#">Col20a1.aSep08</a>	<a href="#">311716</a>	3209	441		146	collagen, type XX, alpha 1 (Col20a1) mRNA.
<a href="#">Col22a1</a>	<a href="#">Col22a1.aSep08</a>	<a href="#">315071</a>	37272	766		186	collagen, type XXII, alpha 1 (Col22a1) mRNA.
<a href="#">Col23a1</a>	<a href="#">Col23a1.aSep08</a>	<a href="#">353303</a>	4119	1180	2	57	collagen, type XXIII, alpha 1 (Col23a1) alternative variant aSep08, mRNA.
<a href="#">Col23a1</a>	<a href="#">Col23a1.bSep08</a>	<a href="#">353303</a>	3503	733	1	128	collagen, type XXIII, alpha 1 (14.3 kD) (Col23a1) alternative variant bSep08, mRNA.
<a href="#">Col24a1</a>	<a href="#">Col24a1.aSep08</a>	<a href="#">499723</a>	34251	728	8	242	collagen, type XXIV, alpha 1 (Col24a1) alternative variant aSep08, mRNA.
<a href="#">Col24a1</a>	<a href="#">Col24a1.bSep08</a>	<a href="#">499723</a>	9746	705	2	187	collagen, type XXIV, alpha 1 (Col24a1) alternative variant bSep08, mRNA.
<a href="#">Col27a1</a>	<a href="#">Col27a1.aSep08</a>	<a href="#">298101</a>	17672	1027		342	collagen, type XXVII, alpha 1 (Col27a1) mRNA.
<a href="#">Colec12</a>	<a href="#">Colec12.bSep08</a>	<a href="#">361289</a>	16546	801	4	264	collectin sub-family member 12 (Colec12) alternative variant bSep08, mRNA.
<a href="#">Colec12</a>	<a href="#">Colec12.cSep08</a>	<a href="#">361289</a>	151887	911	3	91	collectin sub-family member 12 (Colec12) alternative variant cSep08, mRNA.
<a href="#">COLFI.0</a>	<a href="#">COLFI.0.aSep08</a>		7490	2413		244	type XXVII alpha 1 (COLFI.0) mRNA.
<a href="#">Collagen.0</a>	<a href="#">Collagen.0.aSep08</a>		11008	388		129	collagen triple helix repeat (Collagen.0) mRNA.
<a href="#">Collagen.1</a>	<a href="#">Collagen.1.aSep08</a>		20567	1010		336	collagen type IV alpha (Collagen.1) mRNA.

<a href="#">Collagen.2</a>	<a href="#">Collagen.2.aSep08</a>		3249	383		127	collagen type IV (Collagen.2) mRNA.
<a href="#">Collagen.3</a>	<a href="#">Collagen.3.aSep08</a>		2930	352		117	collagen (Collagen.3) mRNA.
<a href="#">Collagen.4</a>	<a href="#">Collagen.4.aSep08</a>		1040	968		322	type X alpha 1 (Collagen.4) mRNA.
<a href="#">Collagen.5</a>	<a href="#">Collagen.5.aSep08</a>		25522	1999		666	procollagen type IV alpha 2 (Collagen.5) mRNA.
<a href="#">Collagen.6</a>	<a href="#">Collagen.6.aSep08</a>		73386	1096		314	collagen type IV alpha 1 CRA a (Collagen.6) mRNA.
<a href="#">Collagen.7</a>	<a href="#">Collagen.7.aSep08</a>		10963	542		180	collagen triple helix repeat (Collagen.7) mRNA.
<a href="#">Collagen.8</a>	<a href="#">Collagen.8.aSep08</a>		145002	488		143	collagen (Collagen.8) mRNA.
<a href="#">Collagen.9</a>	<a href="#">Collagen.9.aSep08</a>		1003	292		97	collagen triple helix repeat (Collagen.9) mRNA.
<a href="#">Collagen.10</a>	<a href="#">Collagen.10.aSep08</a>		6284	275		79	alpha 2 type V Collagen (Collagen.10) mRNA.
<a href="#">Collagen.11</a>	<a href="#">Collagen.11.aSep08</a>		9396	752		250	procollagen type IV alpha 4 CRA a (Collagen.11) mRNA.
<a href="#">Collagen.12</a>	<a href="#">Collagen.12.aSep08</a>		5606	577		191	procollagen type IV alpha 4 CRA b like (Collagen.12) mRNA.
<a href="#">Collagen.13</a>	<a href="#">Collagen.13.aSep08</a>		1878	428		142	procollagen type VII alpha 1 (Collagen.13) mRNA.
<a href="#">Collagen.14</a>	<a href="#">Collagen.14.aSep08</a>		4369	517		172	type XXVII alpha 1 (Collagen.14) mRNA.
<a href="#">Collagen.15</a>	<a href="#">Collagen.15.aSep08</a>		11059	1003	15	324	alpha type IX collagen (Collagen.15) alternative variant aSep08, mRNA.
<a href="#">Collagen.15</a>	<a href="#">Collagen.15.bSep08</a>		1377	848	2	93	putative protein (Collagen.15) alternative variant bSep08, mRNA.
<a href="#">Collagen.16</a>	<a href="#">Collagen.16.aSep08</a>		3498	572		190	procollagen type XVI alpha 1 (Collagen.16) mRNA.
<a href="#">Collagen.17</a>	<a href="#">Collagen.17.aSep08</a>		7182	630		209	type XVI alpha 1 (Collagen.17) mRNA.
<a href="#">Collagen.18</a>	<a href="#">Collagen.18.aSep08</a>		34141	1285		428	collagen type V alpha 1 CRA c (Collagen.18) mRNA.
<a href="#">Collagen.19</a>	<a href="#">Collagen.19.aSep08</a>		8727	404		134	collagen triple helix repeat (Collagen.19) mRNA.
<a href="#">Collagen.20</a>	<a href="#">Collagen.20.aSep08</a>		7991	712	14	198	procollagen type IX alpha 3 CRA a like (Collagen.20) alternative variant aSep08, mRNA.
<a href="#">Collagen.21</a>	<a href="#">Collagen.21.aSep08</a>		6178	430		142	collagen type XX alpha 1 CRA a (Collagen.21) mRNA.
<a href="#">Collagen.22</a>	<a href="#">Collagen.22.aSep08</a>		8383	393		130	collagen triple helix repeat (Collagen.22) mRNA.
<a href="#">Collagen.23</a>	<a href="#">Collagen.23.aSep08</a>		7222	439	6	127	collagen triple helix repeat (Collagen.23) alternative variant aSep08, mRNA.
<a href="#">Collagen.23</a>	<a href="#">Collagen.23.bSep08</a>		15782	339	5	112	procollagen type XI alpha 1 CRA b (Collagen.23) alternative variant bSep08, mRNA.
<a href="#">Collagen.24</a>	<a href="#">Collagen.24.aSep08</a>		10023	525		122	collagen chain like (Collagen.24) mRNA.
<a href="#">Commd1</a>	<a href="#">Commd1.bSep08</a>	<a href="#">289831</a>	96215	616		94	putative protein, with a coiled coil domain, of vertebrate origin (10.4 kD) (Commd1) alternative variant bSep08, complete mRNA.
<a href="#">Commd2</a>	<a href="#">Commd2.bSep08</a>	<a href="#">688478</a>	916	820	1	108	putative cytoplasmic protein of eukaryotic origin (11.7 kD) (Commd2) alternative variant bSep08, mRNA.
<a href="#">Commd3</a>	<a href="#">Commd3.bSep08</a>	<a href="#">291339</a>	3454	655	8	82	putative protein (Commd3) alternative variant bSep08, mRNA.
<a href="#">Commd3</a>	<a href="#">Commd3.cSep08</a>	<a href="#">291339</a>	1971	743	2	75	putative protein of vertebrate origin (Commd3) alternative variant cSep08, mRNA.
<a href="#">Commd3</a>	<a href="#">Commd3.dSep08</a>	<a href="#">291339</a>	1617	669	3	52	putative protein of metazoan origin (5.8 kD) (Commd3) alternative variant dSep08, mRNA.
<a href="#">Commd3</a>	<a href="#">Commd3.eSep08</a>	<a href="#">291339</a>	1922	711	2	49	putative protein (Commd3) alternative variant eSep08, mRNA.

<a href="#">Commd4</a>	<a href="#">Commd4.aSep08</a>	<a href="#">363068</a>	3459	837	8	199	HCaRG (21.9 kD) (Commd4) alternative variant aSep08, complete mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.cSep08</a>	<a href="#">363068</a>	2950	729	7	160	HCaRG (Commd4) alternative variant cSep08, mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.dSep08</a>	<a href="#">363068</a>	3451	1245	6	148	HCaRG (16.1 kD) (Commd4) alternative variant dSep08, complete mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.eSep08</a>	<a href="#">363068</a>	1340	689	4	134	HCaRG precursor (14.6 kD) (Commd4) alternative variant eSep08, mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.fSep08</a>	<a href="#">363068</a>	1818	695	2	39	putative protein of metazoan origin (4.4 kD) (Commd4) alternative variant fSep08, mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.gSep08</a>	<a href="#">363068</a>	2141	399	4	37	putative protein (4.1 kD) (Commd4) alternative variant gSep08, mRNA.
<a href="#">Commd4</a>	<a href="#">Commd4.iSep08</a>	<a href="#">363068</a>	1589	241	3		
<a href="#">Commd5</a>	<a href="#">Commd5.cSep08</a>	<a href="#">245974</a>	1150	782	2	220	HCaRG (Commd5) alternative variant cSep08, mRNA.
<a href="#">Commd6</a>	<a href="#">Commd6.bSep08</a>	<a href="#">498559</a>	6251	863	3	74	putative protein of eukaryotic origin (Commd6) alternative variant bSep08, mRNA.
<a href="#">Commd6</a>	<a href="#">Commd6.cSep08</a>	<a href="#">498559</a>	803	714	2	60	putative protein (Commd6) alternative variant cSep08, mRNA.
<a href="#">Commd7</a>	<a href="#">Commd7.aSep08</a>	<a href="#">296285</a>	15204	1525	9	248	HCaRG (Commd7) alternative variant aSep08, mRNA.
<a href="#">Commd7</a>	<a href="#">Commd7.dSep08</a>	<a href="#">296285</a>	4646	585	3	104	putative protein of metazoan origin (Commd7) alternative variant dSep08, mRNA.
<a href="#">Commd7</a>	<a href="#">Commd7.eSep08</a>	<a href="#">296285</a>	3118	734	3	75	putative protein of eukaryotic origin (Commd7) alternative variant eSep08, mRNA.
<a href="#">Commd8</a>	<a href="#">Commd8.bSep08</a>	<a href="#">289603</a>	9720	658	5	90	putative protein of eukaryotic origin (Commd8) alternative variant bSep08, mRNA.
<a href="#">Commd9</a>	<a href="#">Commd9.bSep08</a>	<a href="#">295956</a>	6359	892	6	169	HCaRG (Commd9) alternative variant bSep08, mRNA.
<a href="#">Commd9</a>	<a href="#">Commd9.cSep08</a>	<a href="#">295956</a>	8393	409	2	136	putative protein of vertebrate origin (Commd9) alternative variant cSep08, mRNA.
<a href="#">Commd9</a>	<a href="#">Commd9.dSep08</a>	<a href="#">295956</a>	14435	683	4	99	putative protein of eukaryotic origin (Commd9) alternative variant dSep08, mRNA.
<a href="#">Commd9</a>	<a href="#">Commd9.eSep08</a>	<a href="#">295956</a>	1009	467	2	68	putative protein (Commd9) alternative variant eSep08, mRNA.
<a href="#">Commd9</a>	<a href="#">Commd9.fSep08</a>	<a href="#">295956</a>	2128	1154	2	64	putative protein (Commd9) alternative variant fSep08, mRNA.
<a href="#">Commd10</a>	<a href="#">Commd10.bSep08</a>	<a href="#">361323</a>	134492	597	6	191	HCaRG (Commd10) alternative variant bSep08, mRNA.
<a href="#">Commd10</a>	<a href="#">Commd10.cSep08</a>	<a href="#">361323</a>	136088	1399	6	165	HCaRG (18.8 kD) (Commd10) alternative variant cSep08, complete mRNA.
<a href="#">Complex1_LYR.0</a>	<a href="#">Complex1_LYR.0.aSep08</a>		113693	965		91	LYR motif containing 4 (10.8 kD) (Complex1_LYR.0) mRNA.
<a href="#">Comt</a>	<a href="#">Comt.bSep08</a>	<a href="#">24267</a>	9191	1184	4	200	catechol-O-methyltransferase (22.6 kD) (Comt) alternative variant bSep08, mRNA.
<a href="#">Comtd1</a>	<a href="#">Comtd1.bSep08</a>	<a href="#">305685</a>	1003	698	2	111	O-methyltransferase, family 3 (12.5 kD) (Comtd1) alternative variant bSep08, mRNA.
<a href="#">Copa</a>	<a href="#">Copa.bSep08</a>	<a href="#">304978</a>	4725	733	8	244	coatomer protein complex subunit alpha (Copa) alternative variant bSep08, mRNA.
<a href="#">Copa</a>	<a href="#">Copa.cSep08</a>	<a href="#">304978</a>	2670	260	3	81	coatomer protein complex subunit alpha (Copa) alternative variant cSep08, mRNA.

<a href="#">Copb1</a>	<a href="#">Copb1.bSep08</a>	<a href="#">114023</a>	8684	735	5	202	coatomer protein complex, subunit beta 1 (Copb1) alternative variant bSep08, mRNA.
<a href="#">Copb1</a>	<a href="#">Copb1.cSep08</a>	<a href="#">114023</a>	901	320	2	38	coatomer protein complex, subunit beta 1 (Copb1) alternative variant cSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.aSep08</a>	<a href="#">290659</a>	10787	1253	10	308	coatomer protein complex CRA c (34.7 kD) (Cope) alternative variant aSep08, complete mRNA.
<a href="#">Cope</a>	<a href="#">Cope.bSep08</a>	<a href="#">290659</a>	6065	662	5	217	coatomer protein complex CRA b (Cope) alternative variant bSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.cSep08</a>	<a href="#">290659</a>	6797	697	8	203	coatomer protein complex CRA c (Cope) alternative variant cSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.eSep08</a>	<a href="#">290659</a>	10771	2236	8	166	coatomer protein complex CRA b (18.7 kD) (Cope) alternative variant eSep08, complete mRNA.
<a href="#">Cope</a>	<a href="#">Cope.fSep08</a>	<a href="#">290659</a>	1504	439	2	123	putative protein of eukaryotic origin (Cope) alternative variant fSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.gSep08</a>	<a href="#">290659</a>	3700	445	4	119	coatomer protein complex CRA c (Cope) alternative variant gSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.hSep08</a>	<a href="#">290659</a>	2364	740	2	58	coatomer protein complex CRA c (Cope) alternative variant hSep08, mRNA.
<a href="#">Cope</a>	<a href="#">Cope.iSep08</a>	<a href="#">290659</a>	1009	932	2	36	putative protein (4.1 kD) (Cope) alternative variant iSep08, mRNA.
<a href="#">Copg</a>	<a href="#">Copg.bSep08</a>	<a href="#">297428</a>	24767	1803	11	507	coatomer protein complex (Copg) alternative variant bSep08, mRNA.
<a href="#">Copg</a>	<a href="#">Copg.cSep08</a>	<a href="#">297428</a>	7196	977	3	180	coatomer protein complex (20.4 kD) (Copg) alternative variant cSep08, mRNA.
<a href="#">Copg</a>	<a href="#">Copg.dSep08</a>	<a href="#">297428</a>	2266	961	3	127	coatomer protein complex CRA c (14.5 kD) (Copg) alternative variant dSep08, mRNA.
<a href="#">Copine.0</a>	<a href="#">Copine.0.aSep08</a>		1041	892		212	copine VI (Copine.0) mRNA.
<a href="#">Copine.1</a>	<a href="#">Copine.1.aSep08</a>		6407	1080	8	281	copine I (Copine.1) alternative variant aSep08, mRNA.
<a href="#">Copine.1</a>	<a href="#">Copine.1.bSep08</a>		6322	2422	4	197	copine I (21.2 kD) (Copine.1) alternative variant bSep08, mRNA.
<a href="#">Cops2</a>	<a href="#">Cops2.bSep08</a>	<a href="#">261736</a>	19762	1079	10	354	COP9 (constitutive photomorphogenic) homolog, subunit 2 (Arabidopsis thaliana) (Cops2) alternative variant bSep08, mRNA.
<a href="#">Cops2</a>	<a href="#">Cops2.cSep08</a>	<a href="#">261736</a>	16765	661	8	219	COP9 (constitutive photomorphogenic) homolog, subunit 2 (Arabidopsis thaliana) (Cops2) alternative variant cSep08, mRNA.
<a href="#">Cops2</a>	<a href="#">Cops2.dSep08</a>	<a href="#">261736</a>	2984	925	2	79	COP9 (constitutive photomorphogenic) homolog, subunit 2 (Arabidopsis thaliana) (Cops2) alternative variant dSep08, mRNA.
<a href="#">Cops3</a>	<a href="#">Cops3.bSep08</a>	<a href="#">287367</a>	8704	656	6	102	cop9 (11.6 kD) (Cops3) alternative variant bSep08, mRNA.
<a href="#">Cops3</a>	<a href="#">Cops3.cSep08</a>	<a href="#">287367</a>	9826	735	6	89	cop9 (Cops3) alternative variant cSep08, mRNA.
<a href="#">Cops3</a>	<a href="#">Cops3.dSep08</a>	<a href="#">287367</a>	2994	556	3	67	cop9 constitutive photomorphogenic homolog (7.7 kD) (Cops3) alternative variant dSep08, mRNA.
<a href="#">Cops4</a>	<a href="#">Cops4.bSep08</a>	<a href="#">360915</a>	20663	780	7	260	cop9 constitutive photomorphogenic homolog (Cops4) alternative variant bSep08, mRNA.
<a href="#">Cops4</a>	<a href="#">Cops4.cSep08</a>	<a href="#">360915</a>	1009	874	2	182	cop9 constitutive photomorphogenic homolog (Cops4) alternative variant cSep08, mRNA.



<a href="#">Cops4</a>	<a href="#">Cops4.dSep08</a>	<a href="#">360915</a>	13417	418	3	92	cop9 homolog CRA c (10.6 kD) (Cops4) alternative variant dSep08, mRNA.
<a href="#">Cops5</a>	<a href="#">Cops5.bSep08</a>	<a href="#">312916</a>	15013	723	6	211	COP9 (constitutive photomorphogenic) homolog, subunit 5 (Arabidopsis thaliana) (Cops5) alternative variant bSep08, mRNA.
<a href="#">Cops5</a>	<a href="#">Cops5.cSep08</a>	<a href="#">312916</a>	5727	719	5	199	COP9 (constitutive photomorphogenic) homolog, subunit 5 (Arabidopsis thaliana) (Cops5) alternative variant cSep08, mRNA.
<a href="#">Cops6</a>	<a href="#">Cops6.bSep08</a>	<a href="#">304343</a>	642	470	3	107	COP9 (constitutive photomorphogenic) homolog, subunit 6 (Arabidopsis thaliana) (Cops6) alternative variant bSep08, mRNA.
<a href="#">Cops6</a>	<a href="#">Cops6.cSep08</a>	<a href="#">304343</a>	1201	614	5	89	COP9 (constitutive photomorphogenic) homolog, subunit 6 (Arabidopsis thaliana) (Cops6) alternative variant cSep08, mRNA.
<a href="#">Cops7a</a>	<a href="#">Cops7a.bSep08</a>	<a href="#">312710</a>	2862	826	4	188	COP9 (constitutive photomorphogenic) homolog, subunit 7a (Arabidopsis thaliana) (Cops7a) alternative variant bSep08, mRNA.
<a href="#">Cops7a</a>	<a href="#">Cops7a.cSep08</a>	<a href="#">312710</a>	1486	1324	2	20	COP9 (constitutive photomorphogenic) homolog, subunit 7a (Arabidopsis thaliana) (Cops7a) alternative variant cSep08, mRNA.
<a href="#">Copz1</a>	<a href="#">Copz1.bSep08</a>	<a href="#">315345</a>	9155	617	8	172	copz1 (Copz1) alternative variant bSep08, mRNA.
<a href="#">Copz1</a>	<a href="#">Copz1.cSep08</a>	<a href="#">315345</a>	22521	390	6	110	coatomer protein complex zeta 1 CRA b (Copz1) alternative variant cSep08, mRNA.
<a href="#">Copz1</a>	<a href="#">Copz1.dSep08</a>	<a href="#">315345</a>	25076	759	7	89	copz1 (10.0 kD) (Copz1) alternative variant dSep08, mRNA.
<a href="#">Copz1</a>	<a href="#">Copz1.eSep08</a>	<a href="#">315345</a>	18028	857	3	83	coatomer protein complex zeta 1 CRA b (9.5 kD) (Copz1) alternative variant eSep08, mRNA.
<a href="#">Coq2</a>	<a href="#">Coq2.bSep08</a>	<a href="#">498332</a>	2830	560	2	167	coenzyme Q2 homolog, prenyltransferase (yeast) (Coq2) alternative variant bSep08, mRNA.
<a href="#">Coq4</a>	<a href="#">Coq4.bSep08</a>	<a href="#">366013</a>	2737	815	3	77	coenzyme Q4 homolog (yeast) (Coq4) alternative variant bSep08, mRNA.
<a href="#">Coq6</a>	<a href="#">Coq6.bSep08</a>	<a href="#">299195</a>	2416	681	5	138	coenzyme Q6 homolog CRA a (Coq6) alternative variant bSep08, mRNA.
<a href="#">Coq6</a>	<a href="#">Coq6.cSep08</a>	<a href="#">299195</a>	5647	726	3	138	coenzyme Q6 homolog (Coq6) alternative variant cSep08, mRNA.
<a href="#">Coq6</a>	<a href="#">Coq6.dSep08</a>	<a href="#">299195</a>	1413	526	3	88	coenzyme Q6 homolog CRA a (9.6 kD) (Coq6) alternative variant dSep08, mRNA.
<a href="#">Coq6</a>	<a href="#">Coq6.eSep08</a>	<a href="#">299195</a>	1230	458	2	63	coenzyme Q6 homolog CRA a (7.1 kD) (Coq6) alternative variant eSep08, mRNA.
<a href="#">Coq6</a>	<a href="#">Coq6.fSep08</a>	<a href="#">299195</a>	1024	399	2	25	putative protein (Coq6) alternative variant fSep08, mRNA.
<a href="#">Coq7</a>	<a href="#">Coq7.bSep08</a>	<a href="#">25249</a>	12327	523	1	173	demethyl-Q 7 (Coq7) alternative variant bSep08, mRNA.
<a href="#">Coq10a</a>	<a href="#">Coq10a.bSep08</a>	<a href="#">362810</a>	5319	885	4	185	coenzyme Q10 homolog A (yeast) (Coq10a) alternative variant bSep08, mRNA.
<a href="#">Corin</a>	<a href="#">Corin.bSep08</a>	<a href="#">289596</a>	41553	701	3	128	corin (Corin) alternative variant bSep08, mRNA.
<a href="#">Corin</a>	<a href="#">Corin.cSep08</a>	<a href="#">289596</a>	62272	407	3	111	corin (Corin) alternative variant cSep08, mRNA.
<a href="#">Coro1a</a>	<a href="#">Coro1a.bSep08</a>	<a href="#">155151</a>	3648	1683	6	273	coronin, actin binding protein 1A (29.6 kD) (Coro1a) alternative variant bSep08, mRNA.

<a href="#">Coro1a</a>	<a href="#">Coro1a.dSep08</a>	<a href="#">155151</a>	874	675	3	106	coronin, actin binding protein 1A (Coro1a) alternative variant dSep08, mRNA.
<a href="#">Coro1b</a>	<a href="#">Coro1b.bSep08</a>	<a href="#">29474</a>	3370	1315	5	274	coronin, actin-binding protein, 1B (Coro1b) alternative variant bSep08, mRNA.
<a href="#">Coro1b</a>	<a href="#">Coro1b.cSep08</a>	<a href="#">29474</a>	1337	1257	2	134	coronin, actin-binding protein, 1B (14.8 kD) (Coro1b) alternative variant cSep08, mRNA.
<a href="#">Coro1c</a>	<a href="#">Coro1c.bSep08</a>	<a href="#">501841</a>	67672	1894	4	366	coronin, actin binding protein 1C (Coro1c) alternative variant bSep08, mRNA.
<a href="#">Coro1c</a>	<a href="#">Coro1c.cSep08</a>	<a href="#">501841</a>	63660	424		140	coronin, actin binding protein 1C (Coro1c) alternative variant cSep08, mRNA.
<a href="#">Coro2a</a>	<a href="#">Coro2a.bSep08</a>	<a href="#">313235</a>	6932	1603	7	241	coronin, actin binding protein 2A (27.9 kD) (Coro2a) alternative variant bSep08, mRNA.
<a href="#">Coro2a</a>	<a href="#">Coro2a.cSep08</a>	<a href="#">313235</a>	23357	728	5	189	coronin, actin binding protein 2A (Coro2a) alternative variant cSep08, mRNA.
<a href="#">Coro2a</a>	<a href="#">Coro2a.dSep08</a>	<a href="#">313235</a>	45198	527	4	175	coronin, actin binding protein 2A (Coro2a) alternative variant dSep08, mRNA.
<a href="#">Coro2a</a>	<a href="#">Coro2a.eSep08</a>	<a href="#">313235</a>	4195	642	3	104	coronin, actin binding protein 2A (Coro2a) alternative variant eSep08, mRNA.
<a href="#">Coro2b</a>	<a href="#">Coro2b.aSep08</a>	<a href="#">300768</a>	112271	1133	5	370	coronin, actin binding protein, 2B (Coro2b) alternative variant aSep08, mRNA.
<a href="#">Coro2b</a>	<a href="#">Coro2b.bSep08</a>	<a href="#">300768</a>	67092	568	3	181	coronin, actin binding protein, 2B (Coro2b) alternative variant bSep08, mRNA.
<a href="#">Coro6</a>	<a href="#">Coro6.bSep08</a>	<a href="#">245982</a>	1189	1099	1	70	coronin, actin binding protein 6 (Coro6) alternative variant bSep08, mRNA.
<a href="#">Coro7</a>	<a href="#">Coro7.aSep08</a>	<a href="#">192276</a>	27622	2754	20	688	coronin 7 (75.0 kD) (Coro7) alternative variant aSep08, mRNA.
<a href="#">Coro7</a>	<a href="#">Coro7.bSep08</a>	<a href="#">192276</a>	681	425	4	141	coronin 7 (Coro7) alternative variant bSep08, mRNA.
<a href="#">Coro7</a>	<a href="#">Coro7.dSep08</a>	<a href="#">192276</a>	2084	1092	4	64	coronin 7 (7.3 kD) (Coro7) alternative variant dSep08, mRNA.
<a href="#">Cotl1</a>	<a href="#">Cotl1.bSep08</a>	<a href="#">361422</a>	9012	756	1	159	coactosin-like 1 (Dictyostelium) (Cotl1) alternative variant bSep08, mRNA.
<a href="#">Cotl1</a>	<a href="#">Cotl1.cSep08</a>	<a href="#">361422</a>	21618	411	2	64	coactosin-like 1 (Dictyostelium) (Cotl1) alternative variant cSep08, mRNA.
<a href="#">Cox4i1</a>	<a href="#">Cox4i1.aSep08</a>	<a href="#">29445</a>	6242	717	3	205	cytochrome c oxidase subunit IV isoform 1 (Cox4i1) alternative variant aSep08, mRNA.
<a href="#">Cox4i1</a>	<a href="#">Cox4i1.cSep08</a>	<a href="#">29445</a>	5754	546	3	181	cytochrome c oxidase subunit IV isoform 1 (Cox4i1) alternative variant cSep08, mRNA.
<a href="#">Cox4i1</a>	<a href="#">Cox4i1.dSep08</a>	<a href="#">29445</a>	1856	590	1	163	cytochrome c oxidase subunit IV isoform 1 (Cox4i1) alternative variant dSep08, mRNA.
<a href="#">Cox4i1</a>	<a href="#">Cox4i1.eSep08</a>	<a href="#">29445</a>	5995	474	3	132	cytochrome c oxidase subunit IV isoform 1 (Cox4i1) alternative variant eSep08, mRNA.
<a href="#">Cox4i2</a>	<a href="#">Cox4i2.bSep08</a>	<a href="#">84683</a>	2762	357	3	118	cytochrome c oxidase subunit IV isoform 2 (Cox4i2) alternative variant bSep08, mRNA.
<a href="#">Cox4i2</a>	<a href="#">Cox4i2.cSep08</a>	<a href="#">84683</a>	2441	270	3	89	cytochrome c oxidase subunit IV isoform 2 (Cox4i2) alternative variant cSep08, mRNA.
<a href="#">Cox4nb</a>	<a href="#">Cox4nb.bSep08</a>	<a href="#">361425</a>	781	532	1	118	COX4 neighbor (Cox4nb) alternative variant bSep08, mRNA.

<a href="#">Cox5a</a>	<a href="#">Cox5a.bSep08</a>	<a href="#">252934</a>	3648	572	1	77	cytochrome c oxidase, subunit Va (8.6 kD) (Cox5a) alternative variant bSep08, mRNA.
<a href="#">Cox5b</a>	<a href="#">Cox5b.bSep08</a>	<a href="#">94194</a>	1260	738	1	55	cytochrome c oxidase subunit Vb (5.8 kD) (Cox5b) alternative variant bSep08, complete mRNA.
<a href="#">Cox7a2l</a>	<a href="#">Cox7a2l.bSep08</a>	<a href="#">298762</a>	27143	1070	2	113	cytochrome c oxidase subunit VIIa polypeptide 2 like (12.6 kD) (Cox7a2l) alternative variant bSep08, complete mRNA.
<a href="#">Cox7a2l</a>	<a href="#">Cox7a2l.cSep08</a>	<a href="#">298762</a>	26780	821	3	42	cytochrome c oxidase subunit VIIa polypeptide 2 like (4.6 kD) (Cox7a2l) alternative variant cSep08, mRNA.
<a href="#">Cox7a2l</a>	<a href="#">Cox7a2l.dSep08</a>	<a href="#">298762</a>	26602	766	4	68	cytochrome c oxidase subunit VIIa polypeptide 2 like (Cox7a2l) alternative variant dSep08, mRNA.
<a href="#">Cox7a2l</a>	<a href="#">Cox7a2l.eSep08</a>	<a href="#">298762</a>	26348	696	4	41	cytochrome c oxidase subunit VIIa polypeptide 2 like (Cox7a2l) alternative variant eSep08, mRNA.
<a href="#">Cox15</a>	<a href="#">Cox15.bSep08</a>	<a href="#">309391</a>	5262	827	1	257	COX15 homolog, cytochrome c oxidase assembly protein (yeast) (Cox15) alternative variant bSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.bSep08</a>	<a href="#">89786</a>	27854	1051	4	207	popeye 2 (23.5 kD) (Cox17andPopdc2) alternative variant bSep08, complete mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.bSep08</a>	<a href="#">360718</a>	27854	1051	4	207	popeye 2 (23.5 kD) (Cox17andPopdc2) alternative variant bSep08, complete mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.cSep08</a>	<a href="#">89786</a>	26324	530	2	145	putative protein of vertebrate origin (Cox17andPopdc2) alternative variant cSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.cSep08</a>	<a href="#">360718</a>	26324	530	2	145	putative protein of vertebrate origin (Cox17andPopdc2) alternative variant cSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.eSep08</a>	<a href="#">89786</a>	5768	518	3	63	cytochrome c oxidase (6.9 kD) (Cox17andPopdc2) alternative variant eSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.eSep08</a>	<a href="#">360718</a>	5768	518	3	63	cytochrome c oxidase (6.9 kD) (Cox17andPopdc2) alternative variant eSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.fSep08</a>	<a href="#">89786</a>	5775	425	3	63	cytochrome c oxidase (6.8 kD) (Cox17andPopdc2) alternative variant fSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.fSep08</a>	<a href="#">360718</a>	5775	425	3	63	cytochrome c oxidase (6.8 kD) (Cox17andPopdc2) alternative variant fSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.gSep08</a>	<a href="#">89786</a>	6793	1273	3	166	popeye 2 (18.6 kD) (Cox17andPopdc2) alternative variant gSep08, mRNA.
<a href="#">Cox17andPopdc2</a>	<a href="#">Cox17andPopdc2.gSep08</a>	<a href="#">360718</a>	6793	1273	3	166	popeye 2 (18.6 kD) (Cox17andPopdc2) alternative variant gSep08, mRNA.
<a href="#">Cox18</a>	<a href="#">Cox18.bSep08</a>	<a href="#">289522</a>	11462	1179	3	266	COX18 cytochrome c oxidase assembly homolog (S. cerevisiae) (29.7 kD) (Cox18) alternative variant bSep08, complete mRNA.
<a href="#">Cox18</a>	<a href="#">Cox18.cSep08</a>	<a href="#">289522</a>	6558	714		151	COX18 cytochrome c oxidase assembly homolog (S. cerevisiae) (Cox18) alternative variant cSep08, mRNA.
<a href="#">Cox19</a>	<a href="#">Cox19.aSep08</a>	<a href="#">304330</a>	6707	932	3	111	COX19 cytochrome c oxidase assembly homolog (S. cerevisiae) (13.1 kD) (Cox19) alternative variant aSep08, mRNA.
<a href="#">Cp</a>	<a href="#">Cp.bSep08</a>	<a href="#">24268</a>	20064	1591	5	176	ceruloplasmin (Cp) alternative variant bSep08, mRNA.
<a href="#">Cp</a>	<a href="#">Cp.cSep08</a>	<a href="#">24268</a>	18777	826	5	130	ceruloplasmin (Cp) alternative variant cSep08, mRNA.
<a href="#">Cp110</a>	<a href="#">Cp110.bSep08</a>	<a href="#">361634</a>	7889	2230	6	146	CP110 protein (Cp110) alternative variant bSep08, mRNA.
<a href="#">Cpa1</a>	<a href="#">Cpa1.bSep08</a>	<a href="#">24269</a>	1984	517	1	99	carboxypeptidase A1 (11.1 kD) (Cpa1) alternative variant bSep08, mRNA.

<a href="#">Cpa1</a>	<a href="#">Cpa1.cSep08</a>	<a href="#">24269</a>	1994	405	2	73	carboxypeptidase A1 (8.3 kD) (Cpa1) alternative variant cSep08, mRNA.
<a href="#">Cpa2</a>	<a href="#">Cpa2.bSep08</a>	<a href="#">296959</a>	7459	468	1	156	carboxypeptidase A2 (pancreatic) (Cpa2) alternative variant bSep08, mRNA.
<a href="#">Cpa3</a>	<a href="#">Cpa3.aSep08</a>	<a href="#">54242</a>	18435	727		112	carboxypeptidase A3, mast cell (Cpa3) mRNA.
<a href="#">Cpa6</a>	<a href="#">Cpa6.bSep08</a>	<a href="#">312913</a>	110277	1352	7	314	carboxypeptidase A6 (Cpa6) alternative variant bSep08, mRNA.
<a href="#">Cpa6</a>	<a href="#">Cpa6.cSep08</a>	<a href="#">312913</a>	56381	701	2	96	carboxypeptidase A6 (Cpa6) alternative variant cSep08, mRNA.
<a href="#">Cpa6</a>	<a href="#">Cpa6.dSep08</a>	<a href="#">312913</a>	1898	681	1	55	carboxypeptidase A6 (6.4 kD) (Cpa6) alternative variant dSep08, mRNA.
<a href="#">Cpb1</a>	<a href="#">Cpb1.bSep08</a>	<a href="#">24271</a>	11989	762	2	208	carboxypeptidase B1 (tissue) (Cpb1) alternative variant bSep08, mRNA.
<a href="#">Cpb2</a>	<a href="#">Cpb2.bSep08</a>	<a href="#">113936</a>	34485	674	1	161	carboxypeptidase B2 (plasma) (Cpb2) alternative variant bSep08, mRNA.
<a href="#">Cpd</a>	<a href="#">Cpd.bSep08</a>	<a href="#">25306</a>	4614	1537	4	174	carboxypeptidase D (19.5 kD) (Cpd) alternative variant bSep08, mRNA.
<a href="#">Cpd</a>	<a href="#">Cpd.cSep08</a>	<a href="#">25306</a>	9217	1326	5	132	carboxypeptidase D (Cpd) alternative variant cSep08, mRNA.
<a href="#">Cpd</a>	<a href="#">Cpd.dSep08</a>	<a href="#">25306</a>	2012	391	2	130	carboxypeptidase D (Cpd) alternative variant dSep08, mRNA.
<a href="#">Cpeb1</a>	<a href="#">Cpeb1.bSep08</a>	<a href="#">293056</a>	11165	595	2	98	cytoplasmic polyadenylation element binding protein 1 (Cpeb1) alternative variant bSep08, mRNA.
<a href="#">Cpeb1</a>	<a href="#">Cpeb1.cSep08</a>	<a href="#">293056</a>	14072	1800	3		
<a href="#">Cpeb2</a>	<a href="#">Cpeb2.bSep08</a>	<a href="#">360949</a>	41504	768	8	256	cytoplasmic polyadenylation element binding protein 2 (Cpeb2) alternative variant bSep08, mRNA.
<a href="#">Cpeb2</a>	<a href="#">Cpeb2.cSep08</a>	<a href="#">360949</a>	39127	655	7	218	cytoplasmic polyadenylation element binding protein 2 (Cpeb2) alternative variant cSep08, mRNA.
<a href="#">Cpeb2</a>	<a href="#">Cpeb2.dSep08</a>	<a href="#">360949</a>	31741	575	5	191	cytoplasmic polyadenylation element binding protein 2 (Cpeb2) alternative variant dSep08, mRNA.
<a href="#">Cpeb3</a>	<a href="#">Cpeb3.aSep08</a>	<a href="#">309510</a>	78927	731	6	243	cytoplasmic polyadenylation element binding protein 3 (Cpeb3) alternative variant aSep08, mRNA.
<a href="#">Cpeb4</a>	<a href="#">Cpeb4.bSep08</a>	<a href="#">303010</a>	49713	435	4	145	cytoplasmic polyadenylation element binding protein 4 and hypothetical protein LOC685957 (Cpeb4) alternative variant bSep08, mRNA.
<a href="#">Cpeb4</a>	<a href="#">Cpeb4.bSep08</a>	<a href="#">685957</a>	49713	435	4	145	cytoplasmic polyadenylation element binding protein 4 and hypothetical protein LOC685957 (Cpeb4) alternative variant bSep08, mRNA.
<a href="#">Cplx2</a>	<a href="#">Cplx2.bSep08</a>	<a href="#">116657</a>	70249	304	3	100	complexin 2 (Cplx2) alternative variant bSep08, mRNA.
<a href="#">Cplx4</a>	<a href="#">Cplx4.aSep08</a>	<a href="#">361342</a>	15291	455		151	complexin 4 (Cplx4) mRNA.
<a href="#">Cpm</a>	<a href="#">Cpm.bSep08</a>	<a href="#">314855</a>	745	670	2	48	carboxypeptidase M (Cpm) alternative variant bSep08, mRNA.
<a href="#">Cpn1</a>	<a href="#">Cpn1.bSep08</a>	<a href="#">365466</a>	7368	281	1	75	carboxypeptidase N, polypeptide 1 (Cpn1) alternative variant bSep08, mRNA.
<a href="#">Cpne2</a>	<a href="#">Cpne2.aSep08</a>	<a href="#">291861</a>	11163	689		228	copine II (Cpne2) mRNA.
<a href="#">Cpne3</a>	<a href="#">Cpne3.bSep08</a>	<a href="#">313087</a>	5302	4081	2	85	copine III (Cpne3) alternative variant bSep08, mRNA.

<a href="#">Cpne5</a>	<a href="#">Cpne5.bSep08</a>	<a href="#">309650</a>	16073	727	5	141	copine V (Cpne5) alternative variant bSep08, mRNA.
<a href="#">Cpne5</a>	<a href="#">Cpne5.cSep08</a>	<a href="#">309650</a>	9115	714	2	62	copine V (7.1 kD) (Cpne5) alternative variant cSep08, mRNA.
<a href="#">Cpne6</a>	<a href="#">Cpne6.aSep08</a>	<a href="#">691478</a>	4794	1439	12	327	copine VI (Cpne6) alternative variant aSep08, mRNA.
<a href="#">Cpne6</a>	<a href="#">Cpne6.bSep08</a>	<a href="#">691478</a>	2950	634	6	178	copine VI (Cpne6) alternative variant bSep08, mRNA.
<a href="#">Cpne7</a>	<a href="#">Cpne7.bSep08</a>	<a href="#">361433</a>	7201	584	4	156	copine VII (Cpne7) alternative variant bSep08, mRNA.
<a href="#">Cpne7</a>	<a href="#">Cpne7.cSep08</a>	<a href="#">361433</a>	4420	2145	5	140	copine VII (15.5 kD) (Cpne7) alternative variant cSep08, mRNA.
<a href="#">Cpne8</a>	<a href="#">Cpne8.bSep08</a>	<a href="#">362988</a>	50513	734	8	218	copine VIII (Cpne8) alternative variant bSep08, mRNA.
<a href="#">Cpne9</a>	<a href="#">Cpne9.bSep08</a>	<a href="#">297516</a>	11181	736	5	175	copine family member IX (Cpne9) alternative variant bSep08, mRNA.
<a href="#">Cps1</a>	<a href="#">Cps1.aSep08</a>	<a href="#">497840</a>	110842	4517	37	1469	carbamoyl-phosphate synthetase 1, mitochondrial (Cps1) alternative variant aSep08, mRNA.
<a href="#">Cps1</a>	<a href="#">Cps1.bSep08</a>	<a href="#">497840</a>	32374	664	4	109	carbamoyl-phosphate synthetase 1, mitochondrial (Cps1) alternative variant bSep08, mRNA.
<a href="#">Cps1</a>	<a href="#">Cps1.cSep08</a>	<a href="#">497840</a>	1893	521	2	84	carbamoyl-phosphate synthetase 1, mitochondrial (Cps1) alternative variant cSep08, mRNA.
<a href="#">Cps1</a>	<a href="#">Cps1.dSep08</a>	<a href="#">497840</a>	29113	288	3	66	carbamoyl-phosphate synthetase 1, mitochondrial (Cps1) alternative variant dSep08, mRNA.
<a href="#">Cpsf1</a>	<a href="#">Cpsf1.aSep08</a>	<a href="#">366952</a>	7809	4133	35	1349	cleavage and polyadenylation specific factor 1 (Cpsf1) alternative variant aSep08, mRNA.
<a href="#">Cpsf1</a>	<a href="#">Cpsf1.bSep08</a>	<a href="#">366952</a>	447	366	2	92	cleavage and polyadenylation specific factor 1 (Cpsf1) alternative variant bSep08, mRNA.
<a href="#">Cpsf1</a>	<a href="#">Cpsf1.cSep08</a>	<a href="#">366952</a>	363	286	2	67	cleavage and polyadenylation specific factor 1 (Cpsf1) alternative variant cSep08, mRNA.
<a href="#">Cpsf3</a>	<a href="#">Cpsf3.bSep08</a>	<a href="#">298916</a>	13624	1212	10	190	cleavage and polyadenylation specificity factor 3 (21.8 kD) (Cpsf3) alternative variant bSep08, mRNA.
<a href="#">Cpsf3</a>	<a href="#">Cpsf3.cSep08</a>	<a href="#">298916</a>	2967	968	3	142	cleavage and polyadenylation specificity factor 3 (Cpsf3) alternative variant cSep08, mRNA.
<a href="#">Cpsf3</a>	<a href="#">Cpsf3.eSep08</a>	<a href="#">298916</a>	1498	616	2	39	cleavage and polyadenylation specificity factor 3 (4.4 kD) (Cpsf3) alternative variant eSep08, mRNA.
<a href="#">Cpsf3l</a>	<a href="#">Cpsf3l.bSep08</a>	<a href="#">298688</a>	14610	886	9	211	cleavage polyadenylation specific factor 3-like (Cpsf3l) alternative variant bSep08, mRNA.
<a href="#">Cpsf3l</a>	<a href="#">Cpsf3l.cSep08</a>	<a href="#">298688</a>	15823	806	6	105	cleavage polyadenylation specific factor 3-like (12.2 kD) (Cpsf3l) alternative variant cSep08, mRNA.
<a href="#">Cpsf3l</a>	<a href="#">Cpsf3l.dSep08</a>	<a href="#">298688</a>	815	725	2	69	cleavage polyadenylation specific factor 3-like (7.4 kD) (Cpsf3l) alternative variant dSep08, mRNA.
<a href="#">Cpsf6</a>	<a href="#">Cpsf6.bSep08</a>	<a href="#">299811</a>	41096	2332	11	537	cleavage and polyadenylation specific factor 6 (57.8 kD) (Cpsf6) alternative variant bSep08, mRNA.
<a href="#">Cpsf6</a>	<a href="#">Cpsf6.cSep08</a>	<a href="#">299811</a>	11639	1027	5	184	cleavage and polyadenylation specific factor 6 (Cpsf6) alternative variant cSep08, mRNA.
<a href="#">Cpsf6</a>	<a href="#">Cpsf6.dSep08</a>	<a href="#">299811</a>	27435	755	5	102	cleavage and polyadenylation specific factor 6 (Cpsf6) alternative variant dSep08, mRNA.
<a href="#">Cpt1b</a>	<a href="#">Cpt1b.bSep08</a>	<a href="#">25756</a>	3325	1303	7	277	carnitine palmitoyltransferase 1b, muscle (Cpt1b) alternative variant bSep08, mRNA.

<a href="#">Cpt1b</a>	<a href="#">Cpt1b.cSep08</a>	<a href="#">25756</a>	2355	765	5	187	carnitine palmitoyltransferase 1b, muscle (Cpt1b) alternative variant cSep08, mRNA.
<a href="#">Cpt1b</a>	<a href="#">Cpt1b.dSep08</a>	<a href="#">25756</a>	1970	1140	4	130	carnitine palmitoyltransferase 1b, muscle (Cpt1b) alternative variant dSep08, mRNA.
<a href="#">Cpt1b</a>	<a href="#">Cpt1b.eSep08</a>	<a href="#">25756</a>	1039	413	3	120	carnitine palmitoyltransferase 1b, muscle (Cpt1b) alternative variant eSep08, mRNA.
<a href="#">Cpt1c</a>	<a href="#">Cpt1c.bSep08</a>	<a href="#">308579</a>	4433	775	6	257	carnitine palmitoyltransferase 1c (Cpt1c) alternative variant bSep08, mRNA.
<a href="#">Cpt1c</a>	<a href="#">Cpt1c.cSep08</a>	<a href="#">308579</a>	7427	1417	10	208	carnitine palmitoyltransferase 1c (Cpt1c) alternative variant cSep08, mRNA.
<a href="#">Cpt1c</a>	<a href="#">Cpt1c.dSep08</a>	<a href="#">308579</a>	3308	455	5	151	carnitine palmitoyltransferase 1c (Cpt1c) alternative variant dSep08, mRNA.
<a href="#">Cpt1c</a>	<a href="#">Cpt1c.fSep08</a>	<a href="#">308579</a>	1691	414	3	53	carnitine palmitoyltransferase 1c (6.0 kD) (Cpt1c) alternative variant fSep08, mRNA.
<a href="#">Cpvl</a>	<a href="#">Cpvl.bSep08</a>	<a href="#">502774</a>	19495	523	6	174	carboxypeptidase, vitellogenic-like (Cpvl) alternative variant bSep08, mRNA.
<a href="#">Cpvl</a>	<a href="#">Cpvl.cSep08</a>	<a href="#">502774</a>	10115	353	2	79	carboxypeptidase, vitellogenic-like (Cpvl) alternative variant cSep08, mRNA.
<a href="#">Cpxm2</a>	<a href="#">Cpxm2.bSep08</a>	<a href="#">293566</a>	28442	2457	8	459	carboxypeptidase X 2 (M14 family) (Cpxm2) alternative variant bSep08, mRNA.
<a href="#">Cpxm2</a>	<a href="#">Cpxm2.cSep08</a>	<a href="#">293566</a>	13916	852	4	189	carboxypeptidase X 2 (M14 family) (Cpxm2) alternative variant cSep08, mRNA.
<a href="#">Cr1l</a>	<a href="#">Cr1l.aSep08</a>	<a href="#">54243</a>	43634	1802	9	573	complement component (3b/4b) receptor 1-like (Cr1l) alternative variant aSep08, mRNA.
<a href="#">Cr1l</a>	<a href="#">Cr1l.dSep08</a>	<a href="#">54243</a>	22303	707	6	205	complement component (3b/4b) receptor 1-like (Cr1l) alternative variant dSep08, mRNA.
<a href="#">Cr1l</a>	<a href="#">Cr1l.eSep08</a>	<a href="#">54243</a>	17896	530	6	149	complement component (3b/4b) receptor 1-like (Cr1l) alternative variant eSep08, mRNA.
<a href="#">Cr2</a>	<a href="#">Cr2.bSep08</a>	<a href="#">289395</a>	4167	1851	7	408	complement receptor 2 (Cr2) alternative variant bSep08, mRNA.
<a href="#">Cr2</a>	<a href="#">Cr2.cSep08</a>	<a href="#">289395</a>	8088	830	5	186	complement receptor 2 (Cr2) alternative variant cSep08, mRNA.
<a href="#">Cr2</a>	<a href="#">Cr2.dSep08</a>	<a href="#">289395</a>	12917	643	6	124	complement receptor 2 (Cr2) alternative variant dSep08, mRNA.
<a href="#">Cr2</a>	<a href="#">Cr2.eSep08</a>	<a href="#">289395</a>	5701	685	2	55	complement receptor 2 (6.2 kD) (Cr2) alternative variant eSep08, mRNA.
<a href="#">Crabp2</a>	<a href="#">Crabp2.aSep08</a>	<a href="#">29563</a>	4063	832	3	183	cellular retinoic acid binding protein 2 (20.4 kD) (Crabp2) alternative variant aSep08, mRNA.
<a href="#">Crabp2</a>	<a href="#">Crabp2.dSep08</a>	<a href="#">29563</a>	2015	399	3	86	cellular retinoic acid binding protein 2 (Crabp2) alternative variant dSep08, mRNA.
<a href="#">Cradd</a>	<a href="#">Cradd.aSep08</a>	<a href="#">314756</a>	158845	1177	1	199	adaptor death (22.7 kD) (Cradd) alternative variant aSep08, mRNA.
<a href="#">Cramp1l</a>	<a href="#">Cramp1l.aSep08</a>	<a href="#">287127</a>	11050	793		183	crm, cramped-like (Drosophila) (Cramp1l) mRNA.
<a href="#">Crat</a>	<a href="#">Crat.aSep08</a>	<a href="#">311849</a>	14291	3562	14	626	carnitine acetyltransferase In Complex With (71.0 kD) (Crat) alternative variant aSep08, mRNA.
<a href="#">Crat</a>	<a href="#">Crat.bSep08</a>	<a href="#">311849</a>	8791	2031	6	309	carnitine acetyltransferase (35.4 kD) (Crat) alternative variant bSep08, mRNA.

<a href="#">Crat</a>	<a href="#">Crat.cSep08</a>	<a href="#">311849</a>	2952	659	4	219	carnitine acetyltransferase (Crat) alternative variant cSep08, mRNA.
<a href="#">Crat</a>	<a href="#">Crat.dSep08</a>	<a href="#">311849</a>	1311	527	1	85	carnitine acetyltransferase (Crat) alternative variant dSep08, mRNA.
<a href="#">Crat</a>	<a href="#">Crat.eSep08</a>	<a href="#">311849</a>	529	342	2	58	carnitine acetyltransferase (Crat) alternative variant eSep08, mRNA.
<a href="#">Crb1</a>	<a href="#">Crb1.bSep08</a>	<a href="#">304825</a>	7961	1095	3	364	crumbs homolog 1 (Drosophila) (Crb1) alternative variant bSep08, mRNA.
<a href="#">Crb1</a>	<a href="#">Crb1.dSep08</a>	<a href="#">304825</a>	5609	354	1	85	crumbs homolog 1 (Drosophila) (Crb1) alternative variant dSep08, mRNA.
<a href="#">Crb3</a>	<a href="#">Crb3.aSep08</a>	<a href="#">301112</a>	2832	671	2	118	crumbs homolog 3 (Drosophila) (12.4 kD) (Crb3) alternative variant aSep08, mRNA.
<a href="#">Crbn</a>	<a href="#">Crbn.cSep08</a>	<a href="#">297498</a>	1269	612	2	112	cereblon (Crbn) alternative variant cSep08, mRNA.
<a href="#">Crcp</a>	<a href="#">Crcp.bSep08</a>	<a href="#">114205</a>	34940	691	7	112	calcitonin gene-related peptide-receptor component protein (12.5 kD) (Crcp) alternative variant bSep08, mRNA.
<a href="#">Crcp</a>	<a href="#">Crcp.cSep08</a>	<a href="#">114205</a>	11661	1253	2	83	calcitonin gene-related peptide-receptor component protein (Crcp) alternative variant cSep08, mRNA.
<a href="#">Crcp</a>	<a href="#">Crcp.dSep08</a>	<a href="#">114205</a>	54352	736	5	76	calcitonin gene-related peptide-receptor component protein (8.5 kD) (Crcp) alternative variant dSep08, mRNA.
<a href="#">Creb1</a>	<a href="#">Creb1.cSep08</a>	<a href="#">81646</a>	32434	646	5	90	cAMP responsive element binding protein 1 (9.3 kD) (Creb1) alternative variant cSep08, mRNA.
<a href="#">Creb3</a>	<a href="#">Creb3.aSep08</a>	<a href="#">298400</a>	4609	1529	7	387	cAMP responsive element binding protein 3 (42.9 kD) (Creb3) alternative variant aSep08, complete mRNA.
<a href="#">Creb3</a>	<a href="#">Creb3.cSep08</a>	<a href="#">298400</a>	3603	758	4	112	cAMP responsive element binding protein 3 (Creb3) alternative variant cSep08, mRNA.
<a href="#">Creb311</a>	<a href="#">Creb311.bSep08</a>	<a href="#">362165</a>	7169	775	1	257	cAMP responsive element binding protein 3-like 1 (Creb311) alternative variant bSep08, mRNA.
<a href="#">Creb312</a>	<a href="#">Creb312.bSep08</a>	<a href="#">362339</a>	26873	983	7	325	cAMP responsive element binding protein 3-like 2 (Creb312) alternative variant bSep08, mRNA.
<a href="#">Creb312</a>	<a href="#">Creb312.cSep08</a>	<a href="#">362339</a>	4095	900	2	179	cAMP responsive element binding protein 3-like 2 (19.5 kD) (Creb312) alternative variant cSep08, mRNA.
<a href="#">Creb313</a>	<a href="#">Creb313.bSep08</a>	<a href="#">314638</a>	1679	473	3	106	cAMP responsive element binding protein 3-like 3 (Creb313) alternative variant bSep08, mRNA.
<a href="#">Creb314</a>	<a href="#">Creb314.bSep08</a>	<a href="#">310616</a>	5483	1539	9	222	cAMP responsive element binding protein 3-like 4 (23.7 kD) (Creb314) alternative variant bSep08, mRNA.
<a href="#">Creb314</a>	<a href="#">Creb314.dSep08</a>	<a href="#">310616</a>	873	692	2	67	cAMP responsive element binding protein 3-like 4 (7.3 kD) (Creb314) alternative variant dSep08, mRNA.
<a href="#">Crebbp</a>	<a href="#">Crebbp.bSep08</a>	<a href="#">54244</a>	9374	1792	8	596	CREB binding protein (Crebbp) alternative variant bSep08, mRNA.
<a href="#">Crebl1</a>	<a href="#">Crebl1.bSep08</a>	<a href="#">406169</a>	3283	1341	8	254	cAMP responsive element binding protein-like 1 (Crebl1) alternative variant bSep08, mRNA.
<a href="#">Crebl1</a>	<a href="#">Crebl1.cSep08</a>	<a href="#">406169</a>	2047	720	6	239	cAMP responsive element binding protein-like 1 (Crebl1) alternative variant cSep08, mRNA.
<a href="#">Crebl1</a>	<a href="#">Crebl1.dSep08</a>	<a href="#">406169</a>	1278	792	3	195	cAMP responsive element binding protein-like 1 (21.2 kD) (Crebl1) alternative variant dSep08, mRNA.
<a href="#">Crel1</a>	<a href="#">Crel1.bSep08</a>	<a href="#">312638</a>	7211	1942	5	214	cysteine-rich with EGF-like domains 1 (23.4 kD) (Crel1) alternative variant bSep08, mRNA.

<a href="#">Creld1</a>	<a href="#">Creld1.cSep08</a>	<a href="#">312638</a>	5922	723	5	155	cysteine-rich with EGF-like domains 1 (Creld1) alternative variant cSep08, mRNA.
<a href="#">Creld1</a>	<a href="#">Creld1.dSep08</a>	<a href="#">312638</a>	1085	669	2	86	cysteine-rich with EGF-like domains 1 (9.7 kD) (Creld1) alternative variant dSep08, mRNA.
<a href="#">Creld2</a>	<a href="#">Creld2.bSep08</a>	<a href="#">362978</a>	6854	1174	9	290	cysteine-rich with EGF-like domains 2 (32.0 kD) (Creld2) alternative variant bSep08, complete mRNA.
<a href="#">Creld2</a>	<a href="#">Creld2.cSep08</a>	<a href="#">362978</a>	3309	676	5	225	cysteine-rich with EGF-like domains 2 (Creld2) alternative variant cSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.bSep08</a>	<a href="#">25620</a>	65855	1505	10	357	cAMP responsive element modulator (38.5 kD) (Crem) alternative variant bSep08, complete mRNA.
<a href="#">Crem</a>	<a href="#">Crem.cSep08</a>	<a href="#">25620</a>	14354	1042	5	290	cAMP responsive element modulator (Crem) alternative variant cSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.dSep08</a>	<a href="#">25620</a>	30923	818	7	239	cAMP responsive element modulator (Crem) alternative variant dSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.eSep08</a>	<a href="#">25620</a>	30998	746	6	208	cAMP responsive element modulator (22.1 kD) (Crem) alternative variant eSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.gSep08</a>	<a href="#">25620</a>	14093	698	3	123	cAMP responsive element modulator (Crem) alternative variant gSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.hSep08</a>	<a href="#">25620</a>	13779	780	3	121	cAMP responsive element modulator (Crem) alternative variant hSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.iSep08</a>	<a href="#">25620</a>	12908	710	3	95	cAMP responsive element modulator (10.9 kD) (Crem) alternative variant iSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.jSep08</a>	<a href="#">25620</a>	13640	1046	3	95	cAMP responsive element modulator (10.8 kD) (Crem) alternative variant jSep08, mRNA.
<a href="#">Crem</a>	<a href="#">Crem.lSep08</a>	<a href="#">25620</a>	53484	301	3	59	cAMP responsive element modulator (Crem) alternative variant lSep08, mRNA.
<a href="#">Crhr1</a>	<a href="#">Crhr1.bSep08</a>	<a href="#">58959</a>	42842	2262	13	414	corticotropin releasing hormone receptor 1 (47.7 kD) (Crhr1) alternative variant bSep08, complete mRNA.
<a href="#">Crhr1</a>	<a href="#">Crhr1.cSep08</a>	<a href="#">58959</a>	1209	1038	3	304	corticotropin releasing hormone receptor 1 (Crhr1) alternative variant cSep08, mRNA.
<a href="#">Crhr1</a>	<a href="#">Crhr1.dSep08</a>	<a href="#">58959</a>	41825	1313	13	290	corticotropin releasing hormone receptor 1 (33.7 kD) (Crhr1) alternative variant dSep08, mRNA.
<a href="#">Crhr1</a>	<a href="#">Crhr1.eSep08</a>	<a href="#">58959</a>	41825	1094	12	224	corticotropin releasing hormone receptor 1 (24.9 kD) (Crhr1) alternative variant eSep08, mRNA.
<a href="#">Crim1</a>	<a href="#">Crim1.aSep08</a>	<a href="#">298744</a>	138769	1801	8	589	cysteine rich transmembrane BMP regulator 1 (chordin like) (Crim1) alternative variant aSep08, mRNA.
<a href="#">Crim1</a>	<a href="#">Crim1.bSep08</a>	<a href="#">298744</a>	111968	1106	8	368	cysteine rich transmembrane BMP regulator 1 (chordin like) (Crim1) alternative variant bSep08, mRNA.
<a href="#">Crim1</a>	<a href="#">Crim1.cSep08</a>	<a href="#">298744</a>	23775	3367	6	319	cysteine rich transmembrane BMP regulator 1 (chordin like) (Crim1) alternative variant cSep08, mRNA.
<a href="#">Crim1</a>	<a href="#">Crim1.dSep08</a>	<a href="#">298744</a>	72933	447	4	148	cysteine rich transmembrane BMP regulator 1 (chordin like) (Crim1) alternative variant dSep08, mRNA.
<a href="#">Crim1</a>	<a href="#">Crim1.eSep08</a>	<a href="#">298744</a>	20211	536	3	34	cysteine rich transmembrane BMP regulator 1 (chordin like) (3.8 kD) (Crim1) alternative variant eSep08, mRNA.
<a href="#">Crim2</a>	<a href="#">Crim2.aSep08</a>	<a href="#">296952</a>	5981	843	6	280	cysteine rich BMP regulator 2 (chordin like) (Crim2) alternative variant aSep08, mRNA.



<a href="#">Crim2</a>	<a href="#">Crim2.bSep08</a>	<a href="#">296952</a>	2891	715	3	224	cysteine rich BMP regulator 2 (chordin like) (Crim2) alternative variant bSep08, mRNA.
<a href="#">Crip2</a>	<a href="#">Crip2.bSep08</a>	<a href="#">338401</a>	4157	775	2	174	cysteine-rich protein 2 (18.9 kD) (Crip2) alternative variant bSep08, mRNA.
<a href="#">Crip2</a>	<a href="#">Crip2.cSep08</a>	<a href="#">338401</a>	1517	752	2	151	cysteine-rich protein 2 (Crip2) alternative variant cSep08, mRNA.
<a href="#">Crip2</a>	<a href="#">Crip2.dSep08</a>	<a href="#">338401</a>	4885	1392	3	144	cysteine-rich protein 2 (15.6 kD) (Crip2) alternative variant dSep08, complete mRNA.
<a href="#">Cript</a>	<a href="#">Cript.bSep08</a>	<a href="#">56725</a>	2920	413	1	58	cysteine-rich PDZ-binding protein (Cript) alternative variant bSep08, mRNA.
<a href="#">Crisp1</a>	<a href="#">Crisp1.bSep08</a>	<a href="#">64827</a>	9470	770	2	80	cysteine-rich secretory protein 1 (Crisp1) alternative variant bSep08, mRNA.
<a href="#">Crispld1</a>	<a href="#">Crispld1.aSep08</a>	<a href="#">316482</a>	39181	3334	11	500	allergen V5/Tpx-1 related and LCCL precursor (57.0 kD) (Crispld1) alternative variant aSep08, complete mRNA.
<a href="#">Crispld1</a>	<a href="#">Crispld1.bSep08</a>	<a href="#">316482</a>	4129	558	1	40	putative protein (4.5 kD) (Crispld1) alternative variant bSep08, mRNA.
<a href="#">Crk</a>	<a href="#">Crk.bSep08</a>	<a href="#">54245</a>	22432	583	3	169	v-crk sarcoma virus CT10 oncogene homolog (avian) (Crk) alternative variant bSep08, mRNA.
<a href="#">Crkrs</a>	<a href="#">Crkrs.cSep08</a>	<a href="#">192350</a>	7601	381	1	127	cdc2-related kinase, arginine/serine-rich (Crkrs) alternative variant cSep08, mRNA.
<a href="#">Crif1</a>	<a href="#">Crif1.bSep08</a>	<a href="#">290655</a>	1814	1497	3	131	cytokine receptor-like factor 1 (14.3 kD) (Crif1) alternative variant bSep08, mRNA.
<a href="#">Crmp1</a>	<a href="#">Crmp1.cSep08</a>	<a href="#">25415</a>	28884	274	3	37	collapsin response mediator protein 1 (4.0 kD) (Crmp1) alternative variant cSep08, mRNA.
<a href="#">Crnk1.1</a>	<a href="#">Crnk1.1.aSep08</a>	<a href="#">116481</a>	9422	1535		428	crn, crooked neck-like 1 (Drosophila) (Crnk1.1) mRNA.
<a href="#">Crocc</a>	<a href="#">Crocc.bSep08</a>	<a href="#">313663</a>	2978	978	6	276	ciliary rootlet coiled-coil, rootletin (Crocc) alternative variant bSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.bSep08</a>	<a href="#">360602</a>	7895	1571	9	281	cisplatin resistance-associated overexpressed protein (34.0 kD) (Crop) alternative variant bSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.cSep08</a>	<a href="#">360602</a>	6114	995	4	172	cisplatin resistance-associated overexpressed protein (Crop) alternative variant cSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.dSep08</a>	<a href="#">360602</a>	26968	506	6	168	cisplatin resistance-associated overexpressed protein (Crop) alternative variant dSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.eSep08</a>	<a href="#">360602</a>	4296	668	3	158	cisplatin resistance-associated overexpressed protein (Crop) alternative variant eSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.fSep08</a>	<a href="#">360602</a>	2032	594	3	66	cisplatin resistance-associated overexpressed protein (Crop) alternative variant fSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.gSep08</a>	<a href="#">360602</a>	3331	731	3	75	cisplatin resistance-associated overexpressed protein (8.8 kD) (Crop) alternative variant gSep08, mRNA.
<a href="#">Crop</a>	<a href="#">Crop.hSep08</a>	<a href="#">360602</a>	25747	524	5	45	cisplatin resistance-associated overexpressed protein (5.0 kD) (Crop) alternative variant hSep08, mRNA.
<a href="#">Crot</a>	<a href="#">Crot.bSep08</a>	<a href="#">83842</a>	16777	409	4	136	carnitine O-octanoyltransferase (Crot) alternative variant bSep08, mRNA.
<a href="#">Crtac1</a>	<a href="#">Crtac1.bSep08</a>	<a href="#">171438</a>	9331	536	3	143	cartilage acidic protein 1 (Crtac1) alternative variant bSep08, mRNA.
<a href="#">Crtac1</a>	<a href="#">Crtac1.cSep08</a>	<a href="#">171438</a>	92936	389	3	129	cartilage acidic protein 1 (Crtac1) alternative variant cSep08, mRNA.

<a href="#">Crtam</a>	<a href="#">Crtam.bSep08</a>	<a href="#">300649</a>	18883	602	2	196	cytotoxic and regulatory T cell molecule (Crtam) alternative variant bSep08, mRNA.
<a href="#">Crtap</a>	<a href="#">Crtap.aSep08</a>	<a href="#">363158</a>	19633	1617	5	397	cartilage associated protein (Crtap) alternative variant aSep08, mRNA.
<a href="#">Crtc2</a>	<a href="#">Crtc2.bSep08</a>	<a href="#">310615</a>	2366	1850	4	164	CREB regulated transcription coactivator 2 (17.3 kD) (Crtc2) alternative variant bSep08, mRNA.
<a href="#">Crtc2</a>	<a href="#">Crtc2.dSep08</a>	<a href="#">310615</a>	2026	694	2	93	CREB regulated transcription coactivator 2 (Crtc2) alternative variant dSep08, mRNA.
<a href="#">Crtc2</a>	<a href="#">Crtc2.eSep08</a>	<a href="#">310615</a>	2600	1222	5	61	CREB regulated transcription coactivator 2 (Crtc2) alternative variant eSep08, mRNA.
<a href="#">Crtc3</a>	<a href="#">Crtc3.aSep08</a>	<a href="#">365297</a>	1410	962		80	CREB regulated transcription coactivator 3 (Crtc3) mRNA.
<a href="#">Cry1</a>	<a href="#">Cry1.bSep08</a>	<a href="#">299691</a>	11870	591	5	118	cryptochrome 1 (photolyase-like) (Cry1) alternative variant bSep08, mRNA.
<a href="#">Cry2</a>	<a href="#">Cry2.bSep08</a>	<a href="#">170917</a>	10972	716	5	219	cryptochrome 2 (photolyase-like) (Cry2) alternative variant bSep08, mRNA.
<a href="#">Cry2</a>	<a href="#">Cry2.cSep08</a>	<a href="#">170917</a>	9569	682	4	184	cryptochrome 2 (photolyase-like) (Cry2) alternative variant cSep08, mRNA.
<a href="#">Cryab</a>	<a href="#">Cryab.aSep08</a>	<a href="#">25420</a>	3817	718	2	175	crystallin, alpha B (20.1 kD) (Cryab) alternative variant aSep08, mRNA.
<a href="#">Cryab</a>	<a href="#">Cryab.cSep08</a>	<a href="#">25420</a>	2682	484	1	114	crystallin, alpha B (Cryab) alternative variant cSep08, mRNA.
<a href="#">Cryab</a>	<a href="#">Cryab.dSep08</a>	<a href="#">25420</a>	2593	505	1	108	crystallin, alpha B (12.2 kD) (Cryab) alternative variant dSep08, mRNA.
<a href="#">Cryba1</a>	<a href="#">Cryba1.aSep08</a>	<a href="#">25583</a>	6523	781		215	crystallin, beta A1 (25.3 kD) (Cryba1) complete mRNA.
<a href="#">Cryba2</a>	<a href="#">Cryba2.bSep08</a>	<a href="#">286925</a>	1328	477	1	133	crystallin, beta A2 (Cryba2) alternative variant bSep08, mRNA.
<a href="#">Crygd</a>	<a href="#">Crygd.aSep08</a>	<a href="#">24278</a>	1420	529	1	175	crystallin, gamma D (Crygd) alternative variant aSep08, mRNA.
<a href="#">Crygs</a>	<a href="#">Crygs.bSep08</a>	<a href="#">689897</a>	20143	425		97	crystallin, gamma S (11.3 kD) (Crygs) alternative variant bSep08, complete mRNA.
<a href="#">Cryl1</a>	<a href="#">Cryl1.bSep08</a>	<a href="#">290277</a>	50985	847	1	254	crystallin, lambda 1 (Cryl1) alternative variant bSep08, mRNA.
<a href="#">Crystall.0</a>	<a href="#">Crystall.0.aSep08</a>		15849	2118		286	in melanoma 1 like (Crystall.0) mRNA.
<a href="#">Cryzl1</a>	<a href="#">Cryzl1.bSep08</a>	<a href="#">288256</a>	10995	278	3	92	crystallin, zeta (quinone reductase)-like 1 (Cryzl1) alternative variant bSep08, mRNA.
<a href="#">Cryzl1</a>	<a href="#">Cryzl1.cSep08</a>	<a href="#">288256</a>	2539	1699	2	42	crystallin, zeta (quinone reductase)-like 1 (4.9 kD) (Cryzl1) alternative variant cSep08, mRNA.
<a href="#">Cs</a>	<a href="#">Cs.bSep08</a>	<a href="#">170587</a>	2344	983	4	145	citrate synthase (Cs) alternative variant bSep08, mRNA.
<a href="#">Cs</a>	<a href="#">Cs.cSep08</a>	<a href="#">170587</a>	24583	850	1	130	citrate synthase (Cs) alternative variant cSep08, mRNA.
<a href="#">Cs</a>	<a href="#">Cs.dSep08</a>	<a href="#">170587</a>	490	409	2	70	citrate synthase (7.7 kD) (Cs) alternative variant dSep08, mRNA.
<a href="#">CS.1</a>	<a href="#">CS.1.aSep08</a>		10200	1231		146	CS (CS.1) mRNA.
<a href="#">Csad</a>	<a href="#">Csad.cSep08</a>	<a href="#">60356</a>	19081	731	7	243	cysteine sulfinic acid decarboxylase (Csad) alternative variant cSep08, mRNA.
<a href="#">Csad</a>	<a href="#">Csad.dSep08</a>	<a href="#">60356</a>	1804	745	5	172	cysteine sulfinic acid decarboxylase (Csad) alternative variant dSep08, mRNA.

<a href="#">Csda</a>	<a href="#">Csda.aSep08</a>	<a href="#">83807</a>	23336	1663	9	365	cold shock domain protein A (Csda) alternative variant aSep08, mRNA.
<a href="#">Csda</a>	<a href="#">Csda.bSep08</a>	<a href="#">83807</a>	19510	1406	9	272	cold shock domain protein A (Csda) alternative variant bSep08, mRNA.
<a href="#">Csda</a>	<a href="#">Csda.cSep08</a>	<a href="#">83807</a>	7521	580	4	191	cold shock domain protein A (Csda) alternative variant cSep08, mRNA.
<a href="#">Csdc2</a>	<a href="#">Csdc2.aSep08</a>	<a href="#">266600</a>	32794	1768		232	PIPPin protein (24.8 kD) (Csdc2) mRNA.
<a href="#">Csde1</a>	<a href="#">Csde1.bSep08</a>	<a href="#">117180</a>	10239	2596	10	448	cold-shock protein, DNA-binding (Csde1) alternative variant bSep08, mRNA.
<a href="#">Csde1</a>	<a href="#">Csde1.cSep08</a>	<a href="#">117180</a>	15396	1440	7	262	upstream of NRAS CRA a (29.6 kD) (Csde1) alternative variant cSep08, mRNA.
<a href="#">Csde1</a>	<a href="#">Csde1.dSep08</a>	<a href="#">117180</a>	4715	590	5	143	cold-shock protein, DNA-binding (Csde1) alternative variant dSep08, mRNA.
<a href="#">Csde1</a>	<a href="#">Csde1.eSep08</a>	<a href="#">117180</a>	3481	388	3	129	cold-shock protein, DNA-binding (Csde1) alternative variant eSep08, mRNA.
<a href="#">Csde1</a>	<a href="#">Csde1.fSep08</a>	<a href="#">117180</a>	9019	365	3	81	putative protein of vertebrate origin (Csde1) alternative variant fSep08, mRNA.
<a href="#">Csf1r</a>	<a href="#">Csf1r.bSep08</a>	<a href="#">307403</a>	4673	598	3	177	colony stimulating factor 1 receptor (Csf1r) alternative variant bSep08, mRNA.
<a href="#">Csf1r</a>	<a href="#">Csf1r.cSep08</a>	<a href="#">307403</a>	25372	701	4	139	colony stimulating factor 1 receptor (Csf1r) alternative variant cSep08, mRNA.
<a href="#">Csf2rb</a>	<a href="#">Csf2rb.bSep08</a>	<a href="#">171081</a>	447	333	2	97	colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage) (Csf2rb) alternative variant bSep08, mRNA.
<a href="#">Csf3r</a>	<a href="#">Csf3r.bSep08</a>	<a href="#">298518</a>	1714	349	2	51	colony stimulating factor 3 receptor (granulocyte) (5.6 kD) (Csf3r) alternative variant bSep08, mRNA.
<a href="#">Csgalnact2</a>	<a href="#">Csgalnact2.bSep08</a>	<a href="#">297554</a>	22980	1785	1	376	chondroitin sulfate N-acetylgalactosaminyltransferase 2 (Csgalnact2) alternative variant bSep08, mRNA.
<a href="#">Csgalnact2</a>	<a href="#">Csgalnact2.cSep08</a>	<a href="#">297554</a>	1465	225	1	51	chondroitin sulfate N-acetylgalactosaminyltransferase 2 (Csgalnact2) alternative variant cSep08, mRNA.
<a href="#">Csk</a>	<a href="#">Csk.bSep08</a>	<a href="#">315707</a>	1427	777	6	199	c-src tyrosine kinase (Csk) alternative variant bSep08, mRNA.
<a href="#">Csk</a>	<a href="#">Csk.dSep08</a>	<a href="#">315707</a>	1660	1560	2	108	c-src tyrosine kinase (Csk) alternative variant dSep08, mRNA.
<a href="#">Csmd1</a>	<a href="#">Csmd1.aSep08</a>	<a href="#">364634</a>	7794	859		286	CUB and Sushi multiple domains 1 (Csmd1) mRNA.
<a href="#">Csn1s1</a>	<a href="#">Csn1s1.aSep08</a>	<a href="#">24284</a>	16178	1779		487	casein alpha s1 (Csn1s1) alternative variant aSep08, mRNA.
<a href="#">Csn1s1</a>	<a href="#">Csn1s1.bSep08</a>	<a href="#">24284</a>	1740	493		35	casein alpha s1 (Csn1s1) alternative variant bSep08, mRNA.
<a href="#">Csn3</a>	<a href="#">Csn3.aSep08</a>	<a href="#">29188</a>	2465	427		71	casein kappa (Csn3) mRNA.
<a href="#">Csnk1a1</a>	<a href="#">Csnk1a1.bSep08</a>	<a href="#">113927</a>	3274	1515	2	86	casein kinase 1, alpha 1 (Csnk1a1) alternative variant bSep08, mRNA.
<a href="#">Csnk1a1</a>	<a href="#">Csnk1a1.cSep08</a>	<a href="#">113927</a>	7860	1000	2	50	casein kinase 1, alpha 1 (Csnk1a1) alternative variant cSep08, mRNA.
<a href="#">Csnk1d</a>	<a href="#">Csnk1d.bSep08</a>	<a href="#">64462</a>	34613	1901	11	211	casein kinase 1, delta (24.3 kD) (Csnk1d) alternative variant bSep08, complete mRNA.

<a href="#">Csnk1d</a>	<a href="#">Csnk1d.cSep08</a>	<a href="#">64462</a>	8863	1407	5	149	casein kinase 1, delta (16.4 kD) (Csnk1d) alternative variant cSep08, mRNA.
<a href="#">Csnk1e</a>	<a href="#">Csnk1e.bSep08</a>	<a href="#">58822</a>	2968	1813	4	135	casein kinase 1, epsilon (Csnk1e) alternative variant bSep08, mRNA.
<a href="#">Csnk1e</a>	<a href="#">Csnk1e.cSep08</a>	<a href="#">58822</a>	3096	1408	4	114	casein kinase 1, epsilon (Csnk1e) alternative variant cSep08, mRNA.
<a href="#">Csnk1e</a>	<a href="#">Csnk1e.eSep08</a>	<a href="#">58822</a>	2661	1204	4	107	casein kinase 1, epsilon (Csnk1e) alternative variant eSep08, mRNA.
<a href="#">Csnk1e</a>	<a href="#">Csnk1e.fSep08</a>	<a href="#">58822</a>	2628	586	4	98	casein kinase 1, epsilon (10.6 kD) (Csnk1e) alternative variant fSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.aSep08</a>	<a href="#">65278</a>	18314	2325	12	441	casein kinase 1 gamma 2 CRA a (50.1 kD) (Csnk1g2) alternative variant aSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.bSep08</a>	<a href="#">65278</a>	6213	1494	10	400	casein kinase 1 gamma 2 (46.0 kD) (Csnk1g2) alternative variant bSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.cSep08</a>	<a href="#">65278</a>	3215	1795	11	299	casein kinase 1 gamma 2 CRA a (34.7 kD) (Csnk1g2) alternative variant cSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.dSep08</a>	<a href="#">65278</a>	8505	753	5	155	casein kinase 1 gamma 2 (Csnk1g2) alternative variant dSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.eSep08</a>	<a href="#">65278</a>	3522	326	5	108	casein kinase 1 gamma 2 (Csnk1g2) alternative variant eSep08, mRNA.
<a href="#">Csnk1g2</a>	<a href="#">Csnk1g2.fSep08</a>	<a href="#">65278</a>	1305	314	2	45	casein kinase 1 gamma 2 (Csnk1g2) alternative variant fSep08, mRNA.
<a href="#">Csnk1g3</a>	<a href="#">Csnk1g3.bSep08</a>	<a href="#">64823</a>	25696	599	6	199	casein kinase 1, gamma 3 (Csnk1g3) alternative variant bSep08, mRNA.
<a href="#">Csnk1g3</a>	<a href="#">Csnk1g3.cSep08</a>	<a href="#">64823</a>	24404	1023	2	119	casein kinase 1, gamma 3 (Csnk1g3) alternative variant cSep08, mRNA.
<a href="#">Csnk1g3</a>	<a href="#">Csnk1g3.dSep08</a>	<a href="#">64823</a>	23760	403	3	109	casein kinase 1, gamma 3 (Csnk1g3) alternative variant dSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.aSep08</a>	<a href="#">116549</a>	47172	4156	13	391	casein kinase II (45.1 kD) (Csnk2a1) alternative variant aSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.cSep08</a>	<a href="#">116549</a>	33930	633	7	158	casein kinase II (Csnk2a1) alternative variant cSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.dSep08</a>	<a href="#">116549</a>	4052	504	4	145	casein kinase II (15.9 kD) (Csnk2a1) alternative variant dSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.eSep08</a>	<a href="#">116549</a>	30300	335	4	78	casein kinase II alpha (Csnk2a1) alternative variant eSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.fSep08</a>	<a href="#">116549</a>	3071	602	2	78	casein kinase II (Csnk2a1) alternative variant fSep08, mRNA.
<a href="#">Csnk2a1</a>	<a href="#">Csnk2a1.gSep08</a>	<a href="#">116549</a>	1635	442	2	54	casein kinase II (Csnk2a1) alternative variant gSep08, mRNA.
<a href="#">Csnk2a2</a>	<a href="#">Csnk2a2.bSep08</a>	<a href="#">307641</a>	35630	2687	9	313	casein kinase 2, alpha prime polypeptide (Csnk2a2) alternative variant bSep08, mRNA.
<a href="#">Csnk2a2</a>	<a href="#">Csnk2a2.cSep08</a>	<a href="#">307641</a>	7731	809	3	31	casein kinase 2, alpha prime polypeptide (3.5 kD) (Csnk2a2) alternative variant cSep08, mRNA.
<a href="#">Csnk2b</a>	<a href="#">Csnk2b.cSep08</a>	<a href="#">81650</a>	4180	735	6	198	casein kinase 2, beta subunit (Csnk2b) alternative variant cSep08, mRNA.

<a href="#">Csnk2b</a>	<a href="#">Csnk2b.dSep08</a>	<a href="#">81650</a>	730	594	2	140	casein kinase 2, beta subunit (Csnk2b) alternative variant dSep08, mRNA.
<a href="#">Cspg4</a>	<a href="#">Cspg4.bSep08</a>	<a href="#">81651</a>	7375	594	2	60	chondroitin sulfate proteoglycan 4 (Cspg4) alternative variant bSep08, mRNA.
<a href="#">Csrnp3</a>	<a href="#">Csrnp3.aSep08</a>	<a href="#">311093</a>	135560	402		106	cysteine-serine-rich nuclear protein 3 (Csrnp3) mRNA.
<a href="#">Csrp1</a>	<a href="#">Csrp1.bSep08</a>	<a href="#">29276</a>	23765	1694	6	181	cysteine and glycine-rich protein 1 (19.4 kD) (Csrp1) alternative variant bSep08, mRNA.
<a href="#">Csrp1</a>	<a href="#">Csrp1.cSep08</a>	<a href="#">29276</a>	24791	744	5	112	cysteine and glycine-rich protein 1 (11.8 kD) (Csrp1) alternative variant cSep08, mRNA.
<a href="#">Csrp2</a>	<a href="#">Csrp2.bSep08</a>	<a href="#">29317</a>	19116	1187	4	184	cysteine and glycine-rich protein 2 (20.1 kD) (Csrp2) alternative variant bSep08, mRNA.
<a href="#">Csrp2</a>	<a href="#">Csrp2.cSep08</a>	<a href="#">29317</a>	4025	575	3	120	cysteine and glycine-rich protein 2 (Csrp2) alternative variant cSep08, mRNA.
<a href="#">Csrp2bp</a>	<a href="#">Csrp2bp.aSep08</a>	<a href="#">362224</a>	6917	1430	2	186	cysteine and glycine-rich protein 2 binding protein (Csrp2bp) alternative variant aSep08, mRNA.
<a href="#">Csrp2bp</a>	<a href="#">Csrp2bp.bSep08</a>	<a href="#">362224</a>	5100	1423	1	129	cysteine and glycine-rich protein 2 binding protein (15.3 kD) (Csrp2bp) alternative variant bSep08, mRNA.
<a href="#">Csrp3</a>	<a href="#">Csrp3.bSep08</a>	<a href="#">117505</a>	5329	708	1	94	cysteine and glycine-rich protein 3 (10.2 kD) (Csrp3) alternative variant bSep08, mRNA.
<a href="#">Cst3</a>	<a href="#">Cst3.bSep08</a>	<a href="#">25307</a>	3787	233	2	76	cystatin C (Cst3) alternative variant bSep08, mRNA.
<a href="#">Cst7</a>	<a href="#">Cst7.aSep08</a>	<a href="#">296257</a>	10548	1073	3	172	cystatin F (leukocystatin) (Cst7) alternative variant aSep08, mRNA.
<a href="#">Cst7</a>	<a href="#">Cst7.cSep08</a>	<a href="#">296257</a>	8080	390	2	86	cystatin F (leukocystatin) (Cst7) alternative variant cSep08, mRNA.
<a href="#">Cst10</a>	<a href="#">Cst10.bSep08</a>	<a href="#">366219</a>	8338	369	1	49	cystatin 10 (chondrocytes) (5.7 kD) (Cst10) alternative variant bSep08, mRNA.
<a href="#">Cst12</a>	<a href="#">Cst12.bSep08</a>	<a href="#">266776</a>	3549	341	1	52	cystatin 12 (5.9 kD) (Cst12) alternative variant bSep08, mRNA.
<a href="#">Cst13</a>	<a href="#">Cst13.bSep08</a>	<a href="#">502679</a>	1389	268	1	30	cystatin 13 (Cst13) alternative variant bSep08, mRNA.
<a href="#">Cstb</a>	<a href="#">Cstb.bSep08</a>	<a href="#">25308</a>	1039	554	1	86	cystatin B (10.0 kD) (Cstb) alternative variant bSep08, mRNA.
<a href="#">Cstf1</a>	<a href="#">Cstf1.bSep08</a>	<a href="#">311670</a>	7044	877	3	291	cleavage stimulation factor, 3' pre-RNA, subunit 1 (Cstf1) alternative variant bSep08, mRNA.
<a href="#">Cstf1</a>	<a href="#">Cstf1.cSep08</a>	<a href="#">311670</a>	6898	904	3	269	cleavage stimulation factor, 3' pre-RNA, subunit 1 (Cstf1) alternative variant cSep08, mRNA.
<a href="#">Cstf1</a>	<a href="#">Cstf1.dSep08</a>	<a href="#">311670</a>	5446	725	2	241	cleavage stimulation factor, 3' pre-RNA, subunit 1 (Cstf1) alternative variant dSep08, mRNA.
<a href="#">Cstf1</a>	<a href="#">Cstf1.eSep08</a>	<a href="#">311670</a>	5220	641	2	197	cleavage stimulation factor, 3' pre-RNA, subunit 1 (Cstf1) alternative variant eSep08, mRNA.
<a href="#">Cstf3</a>	<a href="#">Cstf3.aSep08</a>	<a href="#">362178</a>	18740	2194	14	563	cleavage stimulation factor, 3' pre-RNA, subunit 3 (Cstf3) alternative variant aSep08, mRNA.
<a href="#">Cstf3</a>	<a href="#">Cstf3.cSep08</a>	<a href="#">362178</a>	18699	1417	2	44	cleavage stimulation factor, 3' pre-RNA, subunit 3 (4.9 kD) (Cstf3) alternative variant cSep08, complete mRNA.
<a href="#">Ctage5</a>	<a href="#">Ctage5.aSep08</a>	<a href="#">299078</a>	52425	3142	20	636	ctage family member 5 CRA e (71.8 kD) (Ctage5) alternative variant aSep08, mRNA.
<a href="#">Ctage5</a>	<a href="#">Ctage5.bSep08</a>	<a href="#">299078</a>	54044	1486	16	378	ctage family member 5 (Ctage5) alternative variant bSep08, mRNA.

<a href="#">Ctage5</a>	<a href="#">Ctage5.cSep08</a>	<a href="#">299078</a>	24008	999	10	333	ctage family member 5 CRA d (Ctage5) alternative variant cSep08, mRNA.
<a href="#">Ctage5</a>	<a href="#">Ctage5.dSep08</a>	<a href="#">299078</a>	2400	380	2	108	meningioma expressed antigen 6 CRA b like (Ctage5) alternative variant dSep08, mRNA.
<a href="#">Ctage5</a>	<a href="#">Ctage5.eSep08</a>	<a href="#">299078</a>	4018	652	2	93	meningioma expressed antigen 6 CRA b like (Ctage5) alternative variant eSep08, mRNA.
<a href="#">Ctage5</a>	<a href="#">Ctage5.fSep08</a>	<a href="#">299078</a>	3666	509	3	82	meningioma expressed antigen 6 CRA b like (Ctage5) alternative variant fSep08, mRNA.
<a href="#">Ctbp1</a>	<a href="#">Ctbp1.bSep08</a>	<a href="#">29382</a>	3706	1928	5	267	C-terminal binding protein 1 (28.9 kD) (Ctbp1) alternative variant bSep08, mRNA.
<a href="#">Ctbp1</a>	<a href="#">Ctbp1.cSep08</a>	<a href="#">29382</a>	14665	620	5	140	C-terminal binding protein 1 (Ctbp1) alternative variant cSep08, mRNA.
<a href="#">Ctbp1</a>	<a href="#">Ctbp1.dSep08</a>	<a href="#">29382</a>	2142	642	5	118	C-terminal binding protein 1 (Ctbp1) alternative variant dSep08, mRNA.
<a href="#">Ctbp2</a>	<a href="#">Ctbp2.bSep08</a>	<a href="#">81717</a>	55805	1034	7	287	C-terminal binding protein 2 (Ctbp2) alternative variant bSep08, mRNA.
<a href="#">Ctbp2</a>	<a href="#">Ctbp2.cSep08</a>	<a href="#">81717</a>	21098	740	5	246	C-terminal binding protein 2 (Ctbp2) alternative variant cSep08, mRNA.
<a href="#">Ctbp2</a>	<a href="#">Ctbp2.dSep08</a>	<a href="#">81717</a>	2946	818	3	127	C-terminal binding protein 2 (14.0 kD) (Ctbp2) alternative variant dSep08, mRNA.
<a href="#">Ctbp2</a>	<a href="#">Ctbp2.eSep08</a>	<a href="#">81717</a>	82482	373	3	92	C-terminal binding protein 2 (Ctbp2) alternative variant eSep08, mRNA.
<a href="#">Ctbp2</a>	<a href="#">Ctbp2.gSep08</a>	<a href="#">81717</a>	96292	440	3	84	C-terminal binding protein 2 (Ctbp2) alternative variant gSep08, mRNA.
<a href="#">Ctcf</a>	<a href="#">Ctcf.bSep08</a>	<a href="#">83726</a>	20102	3525	10	740	CCCTC-binding factor (Ctcf) alternative variant bSep08, mRNA.
<a href="#">Ctcf</a>	<a href="#">Ctcf.cSep08</a>	<a href="#">83726</a>	12263	2383	7	357	CCCTC-binding factor (Ctcf) alternative variant cSep08, mRNA.
<a href="#">Ctcf</a>	<a href="#">Ctcf.dSep08</a>	<a href="#">83726</a>	5475	753	4	192	CCCTC-binding factor (Ctcf) alternative variant dSep08, mRNA.
<a href="#">Ctdp1</a>	<a href="#">Ctdp1.bSep08</a>	<a href="#">291414</a>	40588	1789	4	422	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1 (Ctdp1) alternative variant bSep08, mRNA.
<a href="#">Ctdp1</a>	<a href="#">Ctdp1.cSep08</a>	<a href="#">291414</a>	10963	560	4	186	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1 (Ctdp1) alternative variant cSep08, mRNA.
<a href="#">Ctdsp1</a>	<a href="#">Ctdsp1.bSep08</a>	<a href="#">363249</a>	1637	437	5	145	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 1 (Ctdsp1) alternative variant bSep08, mRNA.
<a href="#">Ctdspl</a>	<a href="#">Ctdspl.bSep08</a>	<a href="#">301056</a>	8045	868	3	95	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase-like (10.5 kD) (Ctdspl) alternative variant bSep08, mRNA.
<a href="#">Ctdspl2</a>	<a href="#">Ctdspl2.bSep08</a>	<a href="#">311368</a>	26201	2692	8	236	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2 (Ctdspl2) alternative variant bSep08, mRNA.

<a href="#">Ctdspl2</a>	<a href="#">Ctdspl2.cSep08</a>	<a href="#">311368</a>	21341	772	3	72	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2 (Ctdspl2) alternative variant cSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.bSep08</a>	<a href="#">364996</a>	1120	455	3	132	putative protein of eukaryotic origin (Ctf8) alternative variant bSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.bSep08</a>	<a href="#">690176</a>	1120	455	3	132	putative protein of eukaryotic origin (Ctf8) alternative variant bSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.cSep08</a>	<a href="#">364996</a>	5920	952	3	129	putative protein (14.3 kD) (Ctf8) alternative variant cSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.cSep08</a>	<a href="#">690176</a>	5920	952	3	129	putative protein (14.3 kD) (Ctf8) alternative variant cSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.dSep08</a>	<a href="#">364996</a>	35399	396	4	63	putative protein (Ctf8) alternative variant dSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.dSep08</a>	<a href="#">690176</a>	35399	396	4	63	putative protein (Ctf8) alternative variant dSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.eSep08</a>	<a href="#">364996</a>	6837	479	2	32	putative protein (Ctf8) alternative variant eSep08, mRNA.
<a href="#">Ctf8</a>	<a href="#">Ctf8.eSep08</a>	<a href="#">690176</a>	6837	479	2	32	putative protein (Ctf8) alternative variant eSep08, mRNA.
<a href="#">Cth</a>	<a href="#">Cth.bSep08</a>	<a href="#">24962</a>	10818	772	1	178	cystathionase (cystathionine gamma-lyase) (Cth) alternative variant bSep08, mRNA.
<a href="#">Ctla2a</a>	<a href="#">Ctla2a.bSep08</a>	<a href="#">498690</a>	1284	591	3	135	cytotoxic T lymphocyte-associated protein 2 alpha (Ctla2a) alternative variant bSep08, mRNA.
<a href="#">Ctla2a</a>	<a href="#">Ctla2a.cSep08</a>	<a href="#">498690</a>	917	816	1	48	cytotoxic T lymphocyte-associated protein 2 alpha (Ctla2a) alternative variant cSep08, mRNA.
<a href="#">Ctnna1</a>	<a href="#">Ctnna1.cSep08</a>	<a href="#">307505</a>	4574	267	3	41	catenin (cadherin associated protein), alpha 1 (Ctnna1) alternative variant cSep08, mRNA.
<a href="#">Ctnnal1</a>	<a href="#">Ctnnal1.bSep08</a>	<a href="#">298019</a>	22147	764	6	254	catenin (cadherin associated protein), alpha-like 1 (Ctnnal1) alternative variant bSep08, mRNA.
<a href="#">Ctnnal1</a>	<a href="#">Ctnnal1.cSep08</a>	<a href="#">298019</a>	3219	409	3	108	catenin (cadherin associated protein), alpha-like 1 (Ctnnal1) alternative variant cSep08, mRNA.
<a href="#">Ctnnb1</a>	<a href="#">Ctnnb1.bSep08</a>	<a href="#">84353</a>	8296	2858	8	417	catenin (cadherin associated protein), beta 1 (45.7 kD) (Ctnnb1) alternative variant bSep08, mRNA.
<a href="#">Ctnnb1</a>	<a href="#">Ctnnb1.cSep08</a>	<a href="#">84353</a>	2845	1037	5	143	catenin (cadherin associated protein), beta 1 (Ctnnb1) alternative variant cSep08, mRNA.
<a href="#">Ctnnbip1</a>	<a href="#">Ctnnbip1.aSep08</a>	<a href="#">503000</a>	47174	630	3	128	catenin, beta-interacting protein 1 (Ctnnbip1) alternative variant aSep08, mRNA.
<a href="#">Ctnnbip1</a>	<a href="#">Ctnnbip1.bSep08</a>	<a href="#">503000</a>	47465	709	1	114	catenin, beta-interacting protein 1 (Ctnnbip1) alternative variant bSep08, mRNA.
<a href="#">Ctnnbip1</a>	<a href="#">Ctnnbip1.cSep08</a>	<a href="#">503000</a>	49010	2551	4	81	catenin, beta-interacting protein 1 (9.2 kD) (Ctnnbip1) alternative variant cSep08, complete mRNA.
<a href="#">Ctnnd1</a>	<a href="#">Ctnnd1.bSep08</a>	<a href="#">311163</a>	15970	1784	4	524	catenin (cadherin associated protein), delta 1 (Ctnnd1) alternative variant bSep08, mRNA.
<a href="#">Ctnnd1</a>	<a href="#">Ctnnd1.cSep08</a>	<a href="#">311163</a>	9625	1390	9	322	catenin (cadherin associated protein), delta 1 (Ctnnd1) alternative variant cSep08, mRNA.
<a href="#">Ctnnd1</a>	<a href="#">Ctnnd1.dSep08</a>	<a href="#">311163</a>	3991	1771	4	85	catenin (cadherin associated protein), delta 1 (Ctnnd1) alternative variant dSep08, mRNA.
<a href="#">Ctnnd2</a>	<a href="#">Ctnnd2.aSep08</a>	<a href="#">114028</a>	123985	1782		404	catenin (cadherin-associated protein), delta 2 (Ctnnd2) alternative variant aSep08, mRNA.
<a href="#">Ctnnd2</a>	<a href="#">Ctnnd2.bSep08</a>	<a href="#">114028</a>	17758	1447		164	catenin (cadherin-associated protein), delta 2 (Ctnnd2) alternative variant bSep08, mRNA.

<a href="#">Ctns</a>	<a href="#">Ctns.aSep08</a>	<a href="#">287478</a>	15143	2054		367	cystinosis, nephropathic (41.9 kD) (Ctns) mRNA.
<a href="#">Ctps</a>	<a href="#">Ctps.bSep08</a>	<a href="#">313560</a>	4785	369	3	123	cytidine 5'-triphosphate synthase (Ctps) alternative variant bSep08, mRNA.
<a href="#">Ctps</a>	<a href="#">Ctps.cSep08</a>	<a href="#">313560</a>	1194	254	3	78	cytidine 5'-triphosphate synthase (Ctps) alternative variant cSep08, mRNA.
<a href="#">Ctps2</a>	<a href="#">Ctps2.bSep08</a>	<a href="#">619580</a>	34806	716	6	213	cytidine 5'-triphosphate synthase 2 (Ctps2) alternative variant bSep08, mRNA.
<a href="#">Ctps2</a>	<a href="#">Ctps2.cSep08</a>	<a href="#">619580</a>	28440	729	5	178	cytidine 5'-triphosphate synthase 2 (Ctps2) alternative variant cSep08, mRNA.
<a href="#">Ctps2</a>	<a href="#">Ctps2.dSep08</a>	<a href="#">619580</a>	27968	759	5	167	cytidine 5'-triphosphate synthase 2 (Ctps2) alternative variant dSep08, mRNA.
<a href="#">Ctr9</a>	<a href="#">Ctr9.aSep08</a>	<a href="#">293184</a>	30595	4234	25	1214	ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (Ctr9) alternative variant aSep08, mRNA.
<a href="#">Ctr9</a>	<a href="#">Ctr9.bSep08</a>	<a href="#">293184</a>	527	371	2	75	ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (Ctr9) alternative variant bSep08, mRNA.
<a href="#">Ctr9</a>	<a href="#">Ctr9.eSep08</a>	<a href="#">293184</a>	6098	408	3	13	ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (1.6 kD) (Ctr9) alternative variant eSep08, mRNA.
<a href="#">Ctr9</a>	<a href="#">Ctr9.fSep08</a>	<a href="#">293184</a>	2128	301	3	59	ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (Ctr9) alternative variant fSep08, mRNA.
<a href="#">Ctr9</a>	<a href="#">Ctr9.gSep08</a>	<a href="#">293184</a>	1457	263	3	48	ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (Ctr9) alternative variant gSep08, mRNA.
<a href="#">Ctrl</a>	<a href="#">Ctrl.cSep08</a>	<a href="#">117184</a>	820	663	3	147	chymotrypsin-like (15.6 kD) (Ctrl) alternative variant cSep08, mRNA.
<a href="#">Ctsb</a>	<a href="#">Ctsb.bSep08</a>	<a href="#">64529</a>	2890	597	1	67	cathepsin B (Ctsb) alternative variant bSep08, mRNA.
<a href="#">Ctsc</a>	<a href="#">Ctsc.bSep08</a>	<a href="#">25423</a>	2058	836	2	198	cathepsin C (22.1 kD) (Ctsc) alternative variant bSep08, mRNA.
<a href="#">Ctsc</a>	<a href="#">Ctsc.cSep08</a>	<a href="#">25423</a>	3123	477	2	107	cathepsin C CRA b (Ctsc) alternative variant cSep08, mRNA.
<a href="#">Ctsd</a>	<a href="#">Ctsd.bSep08</a>	<a href="#">171293</a>	10386	809	6	168	cathepsin D (Ctsd) alternative variant bSep08, mRNA.
<a href="#">Ctsd</a>	<a href="#">Ctsd.cSep08</a>	<a href="#">171293</a>	1800	1254	5	167	cathepsin D (18.4 kD) (Ctsd) alternative variant cSep08, mRNA.
<a href="#">Ctse</a>	<a href="#">Ctse.bSep08</a>	<a href="#">25424</a>	9586	1456	7	304	cathepsin E (Ctse) alternative variant bSep08, mRNA.
<a href="#">Ctse</a>	<a href="#">Ctse.cSep08</a>	<a href="#">25424</a>	680	582	2	46	cathepsin E (Ctse) alternative variant cSep08, mRNA.
<a href="#">Ctsf</a>	<a href="#">Ctsf.bSep08</a>	<a href="#">361704</a>	993	387	1	128	cathepsin F (Ctsf) alternative variant bSep08, mRNA.
<a href="#">Ctsl</a>	<a href="#">Ctsl.bSep08</a>	<a href="#">25697</a>	6773	2093	3	279	cathepsin L (31.1 kD) (Ctsl) alternative variant bSep08, complete mRNA.
<a href="#">Ctsl</a>	<a href="#">Ctsl.cSep08</a>	<a href="#">25697</a>	2273	780	1	80	cathepsin L (Ctsl) alternative variant cSep08, mRNA.
<a href="#">Ctsm</a>	<a href="#">Ctsm.bSep08</a>	<a href="#">306720</a>	2629	511	1	170	cathepsin M (Ctsm) alternative variant bSep08, mRNA.
<a href="#">Ctsq</a>	<a href="#">Ctsq.bSep08</a>	<a href="#">246147</a>	3638	785		236	cathepsin Q (Ctsq) alternative variant bSep08, mRNA.
<a href="#">Ctsql2</a>	<a href="#">Ctsql2.bSep08</a>	<a href="#">408201</a>	1899	712	3	144	cathepsin Q-like 2 (Ctsql2) alternative variant bSep08, mRNA.



<a href="#">Ctsql2</a>	<a href="#">Ctsql2.cSep08</a>	<a href="#">408201</a>	1592	383	3	127	cathepsin Q-like 2 (Ctsql2) alternative variant cSep08, mRNA.
<a href="#">Ctss</a>	<a href="#">Ctss.aSep08</a>	<a href="#">50654</a>	44298	1393	1	342	cathepsin S (Ctss) alternative variant aSep08, complete mRNA.
<a href="#">Ctss</a>	<a href="#">Ctss.cSep08</a>	<a href="#">50654</a>	12211	914		200	cathepsin S (Ctss) alternative variant cSep08, mRNA.
<a href="#">Ctss</a>	<a href="#">Ctss.dSep08</a>	<a href="#">50654</a>	10248	390	1	95	cathepsin S (Ctss) alternative variant dSep08, mRNA.
<a href="#">Cttn</a>	<a href="#">Cttn.bSep08</a>	<a href="#">60465</a>	17231	1013	10	337	cortactin (Cttn) alternative variant bSep08, mRNA.
<a href="#">Cttn</a>	<a href="#">Cttn.cSep08</a>	<a href="#">60465</a>	15531	640	7	213	cortactin (Cttn) alternative variant cSep08, mRNA.
<a href="#">Cttn</a>	<a href="#">Cttn.eSep08</a>	<a href="#">60465</a>	1535	741	2	79	cortactin (9.1 kD) (Cttn) alternative variant eSep08, mRNA.
<a href="#">Ctxn3</a>	<a href="#">Ctxn3.bSep08</a>	<a href="#">100188934</a>	10393	1245	3	37	cortixin 3 (4.0 kD) (Ctxn3) alternative variant bSep08, mRNA.
<a href="#">Ctxn3</a>	<a href="#">Ctxn3.cSep08</a>	<a href="#">100188934</a>	4492	415	1	50	cortixin 3 (Ctxn3) alternative variant cSep08, mRNA.
<a href="#">CUB.0</a>	<a href="#">CUB.0.aSep08</a>		2674	449		149	deleted in malignant brain tumors 1 like (CUB.0) mRNA.
<a href="#">Cubn</a>	<a href="#">Cubn.aSep08</a>	<a href="#">80848</a>	15332	1594	7	440	cubilin (intrinsic factor-cobalamin receptor) (Cubn) alternative variant aSep08, mRNA.
<a href="#">Cubn</a>	<a href="#">Cubn.bSep08</a>	<a href="#">80848</a>	2248	629	2	79	cubilin (intrinsic factor-cobalamin receptor) (Cubn) alternative variant bSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.aSep08</a>	<a href="#">294009</a>	8269	1012	9	270	putative cytoplasmic protein of bilateral origin (30.7 kD) (Cuedc2) alternative variant aSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.cSep08</a>	<a href="#">294009</a>	8098	829	9	250	putative protein of bilateral origin (Cuedc2) alternative variant cSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.dSep08</a>	<a href="#">294009</a>	8220	941	7	146	putative cytoplasmic protein (16.3 kD) (Cuedc2) alternative variant dSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.eSep08</a>	<a href="#">294009</a>	1860	911	5	121	putative protein of vertebrate origin (13.4 kD) (Cuedc2) alternative variant eSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.gSep08</a>	<a href="#">294009</a>	427	370	2	73	putative protein (Cuedc2) alternative variant gSep08, mRNA.
<a href="#">Cuedc2</a>	<a href="#">Cuedc2.hSep08</a>	<a href="#">294009</a>	7020	769	3	32	putative protein of mammalian origin (Cuedc2) alternative variant hSep08, mRNA.
<a href="#">Cugbp1</a>	<a href="#">Cugbp1.bSep08</a>	<a href="#">362160</a>	75256	1804	7	374	CUG triplet repeat, RNA binding protein 1 (Cugbp1) alternative variant bSep08, mRNA.
<a href="#">Cugbp1</a>	<a href="#">Cugbp1.cSep08</a>	<a href="#">362160</a>	8556	686	4	132	CUG triplet repeat, RNA binding protein 1 (Cugbp1) alternative variant cSep08, mRNA.
<a href="#">Cugbp1</a>	<a href="#">Cugbp1.dSep08</a>	<a href="#">362160</a>	14286	399	3	132	CUG triplet repeat, RNA binding protein 1 (Cugbp1) alternative variant dSep08, mRNA.
<a href="#">Cugbp1</a>	<a href="#">Cugbp1.eSep08</a>	<a href="#">362160</a>	4456	2135	3	104	CUG triplet repeat, RNA binding protein 1 (Cugbp1) alternative variant eSep08, mRNA.
<a href="#">Cugbp2</a>	<a href="#">Cugbp2.aSep08</a>	<a href="#">29428</a>	264960	1791	5	582	CUG triplet repeat RNA binding protein 2 like (Cugbp2) alternative variant aSep08, mRNA.
<a href="#">Cugbp2</a>	<a href="#">Cugbp2.dSep08</a>	<a href="#">29428</a>	44202	5958	5	235	CUG triplet repeat RNA binding protein 2 like (Cugbp2) alternative variant dSep08, mRNA.
<a href="#">Cugbp2</a>	<a href="#">Cugbp2.eSep08</a>	<a href="#">29428</a>	40846	1219	6	196	CUG triplet repeat RNA binding protein 2 like (Cugbp2) alternative variant eSep08, mRNA.
<a href="#">Cugbp2</a>	<a href="#">Cugbp2.fSep08</a>	<a href="#">29428</a>	15311	766	5	177	CUG triplet repeat RNA binding protein 2 like (18.6 kD) (Cugbp2) alternative variant fSep08, mRNA.

<a href="#">Cugbp2</a>	<a href="#">Cugbp2.gSep08</a>	<a href="#">29428</a>	228298	382	4	127	CUG triplet repeat RNA binding protein 2 like (Cugbp2) alternative variant gSep08, mRNA.
<a href="#">Cul1</a>	<a href="#">Cul1.bSep08</a>	<a href="#">362356</a>	9202	1609	10	327	cullin 1 (38.1 kD) (Cul1) alternative variant bSep08, mRNA.
<a href="#">Cul1</a>	<a href="#">Cul1.cSep08</a>	<a href="#">362356</a>	38157	678	4	145	cullin 1 (Cul1) alternative variant cSep08, mRNA.
<a href="#">Cul1</a>	<a href="#">Cul1.eSep08</a>	<a href="#">362356</a>	36427	635	3	95	cullin 1 (Cul1) alternative variant eSep08, mRNA.
<a href="#">Cul2</a>	<a href="#">Cul2.bSep08</a>	<a href="#">361258</a>	2862	339	4	112	cullin 2 (Cul2) alternative variant bSep08, mRNA.
<a href="#">Cul3</a>	<a href="#">Cul3.aSep08</a>	<a href="#">301555</a>	77797	2743	16	768	cullin 3 (88.9 kD) (Cul3) alternative variant aSep08, complete mRNA.
<a href="#">Cul3</a>	<a href="#">Cul3.cSep08</a>	<a href="#">301555</a>	77514	1780	8	503	cullin 3 (Cul3) alternative variant cSep08, mRNA.
<a href="#">Cul3</a>	<a href="#">Cul3.dSep08</a>	<a href="#">301555</a>	37420	549	6	131	cullin 3 (Cul3) alternative variant dSep08, mRNA.
<a href="#">Cul3</a>	<a href="#">Cul3.eSep08</a>	<a href="#">301555</a>	5231	787	4	129	cullin 3 (Cul3) alternative variant eSep08, mRNA.
<a href="#">Cul3</a>	<a href="#">Cul3.fSep08</a>	<a href="#">301555</a>	2750	739	2	58	cullin 3 (6.9 kD) (Cul3) alternative variant fSep08, mRNA.
<a href="#">Cul4b</a>	<a href="#">Cul4b.bSep08</a>	<a href="#">302502</a>	766	606	2	201	cullin 4B (Cul4b) alternative variant bSep08, mRNA.
<a href="#">Cul5</a>	<a href="#">Cul5.bSep08</a>	<a href="#">64624</a>	29862	1785	9	404	cullin 5 (Cul5) alternative variant bSep08, mRNA.
<a href="#">Cul5</a>	<a href="#">Cul5.cSep08</a>	<a href="#">64624</a>	20919	753	6	251	cullin 5 (Cul5) alternative variant cSep08, mRNA.
<a href="#">Cul5</a>	<a href="#">Cul5.dSep08</a>	<a href="#">64624</a>	2374	1062	3	95	cullin 5 (Cul5) alternative variant dSep08, mRNA.
<a href="#">CUT.0</a>	<a href="#">CUT.0.aSep08</a>		4405	751		250	cut-like 1 CRA a (CUT.0) mRNA.
<a href="#">CUT.1</a>	<a href="#">CUT.1.aSep08</a>		31212	2174	4	460	cut-like 1 CRA a (49.1 kD) (CUT.1) alternative variant aSep08, complete mRNA.
<a href="#">CUT.1</a>	<a href="#">CUT.1.bSep08</a>		4286	623	1	174	cut-like 1 CRA a (CUT.1) alternative variant bSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.aSep08</a>	<a href="#">294288</a>	1576	664	6	189	cutA (Cuta) alternative variant aSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.bSep08</a>	<a href="#">294288</a>	1421	644	6	183	cutA (Cuta) alternative variant bSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.cSep08</a>	<a href="#">294288</a>	1500	938	5	154	cutA (16.4 kD) (Cuta) alternative variant cSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.dSep08</a>	<a href="#">294288</a>	710	627	2	108	putative protein (Cuta) alternative variant dSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.eSep08</a>	<a href="#">294288</a>	648	406	2	58	divalent cation cutA (6.3 kD) (Cuta) alternative variant eSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.fSep08</a>	<a href="#">294288</a>	1447	789	5	126	putative secreted or extracellular protein precursor of mammalian origin (12.9 kD) (Cuta) alternative variant fSep08, mRNA.
<a href="#">Cuta</a>	<a href="#">Cuta.gSep08</a>	<a href="#">294288</a>	1508	715	5	59	divalent cation cutA (6.5 kD) (Cuta) alternative variant gSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.bSep08</a>	<a href="#">361760</a>	8871	922	6	195	cutC copper transporter homolog (E.coli) (Cutc) alternative variant bSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.cSep08</a>	<a href="#">361760</a>	14813	2171	7	187	cutC copper transporter homolog (E.coli) (20.0 kD) (Cutc) alternative variant cSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.dSep08</a>	<a href="#">361760</a>	8561	572	5	149	cutC copper transporter homolog (E.coli) (Cutc) alternative variant dSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.eSep08</a>	<a href="#">361760</a>	8764	779	5	147	cutC copper transporter homolog (E.coli) (Cutc) alternative variant eSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.fSep08</a>	<a href="#">361760</a>	8058	751	5	119	cutC copper transporter homolog (E.coli) (12.6 kD) (Cutc) alternative variant fSep08, mRNA.
<a href="#">Cutc</a>	<a href="#">Cutc.gSep08</a>	<a href="#">361760</a>	5728	947	3	45	cutC copper transporter homolog (E.coli) (4.8 kD) (Cutc) alternative variant gSep08, mRNA.

<a href="#">Cux1</a>	<a href="#">Cux1.aSep08</a>	<a href="#">116639</a>	317072	3100	1	773	cut-like homeobox 1 (Cux1) alternative variant aSep08, mRNA.
<a href="#">Cux1</a>	<a href="#">Cux1.bSep08</a>	<a href="#">116639</a>	47210	590	1	196	cut-like homeobox 1 (Cux1) alternative variant bSep08, mRNA.
<a href="#">Cuzd1</a>	<a href="#">Cuzd1.bSep08</a>	<a href="#">117179</a>	33813	1410	8	194	CUB and zona pellucida-like domains 1 (22.0 kD) (Cuzd1) alternative variant bSep08, mRNA.
<a href="#">Cuzd1</a>	<a href="#">Cuzd1.cSep08</a>	<a href="#">117179</a>	29313	678	5	148	CUB and zona pellucida-like domains 1 (Cuzd1) alternative variant cSep08, mRNA.
<a href="#">Cuzd1</a>	<a href="#">Cuzd1.dSep08</a>	<a href="#">117179</a>	29360	620	4	123	CUB and zona pellucida-like domains 1 (Cuzd1) alternative variant dSep08, mRNA.
<a href="#">Cuzd1</a>	<a href="#">Cuzd1.eSep08</a>	<a href="#">117179</a>	1277	743	2	32	CUB and zona pellucida-like domains 1 (3.8 kD) (Cuzd1) alternative variant eSep08, mRNA.
<a href="#">Cwc15</a>	<a href="#">Cwc15.aSep08</a>	<a href="#">300361</a>	10404	856	6	229	CWC15 homolog (S. cerevisiae) (26.6 kD) (Cwc15) alternative variant aSep08, complete mRNA.
<a href="#">Cwc15</a>	<a href="#">Cwc15.cSep08</a>	<a href="#">300361</a>	1809	661	1	73	CWC15 homolog (S. cerevisiae) (Cwc15) alternative variant cSep08, mRNA.
<a href="#">Cwf191</a>	<a href="#">Cwf191.cSep08</a>	<a href="#">365465</a>	3859	1349	3	82	CWF19-like 1, cell cycle control (S. pombe) (9.7 kD) (Cwf191) alternative variant cSep08, mRNA.
<a href="#">Cwf192</a>	<a href="#">Cwf192.aSep08</a>	<a href="#">362804</a>	8117	1805	1	148	CWF19-like 2, cell cycle control (S. pombe) (Cwf192) alternative variant aSep08, mRNA.
<a href="#">Cwf192</a>	<a href="#">Cwf192.bSep08</a>	<a href="#">362804</a>	7561	1172	1	116	CWF19-like 2, cell cycle control (S. pombe) (Cwf192) alternative variant bSep08, mRNA.
<a href="#">cwf21.0</a>	<a href="#">cwf21.0.aSep08</a>		4957	867	8	288	serine arginine repetitive matrix 2 CRA b (cwf21.0) alternative variant aSep08, mRNA.
<a href="#">Cxcl3</a>	<a href="#">Cxcl3.aSep08</a>	<a href="#">171551</a>	12753	1524	3	129	chemokine (C-X-C motif) ligand 3 (14.8 kD) (Cxcl3) alternative variant aSep08, mRNA.
<a href="#">Cxcl3</a>	<a href="#">Cxcl3.bSep08</a>	<a href="#">171551</a>	12123	1164	2	128	chemokine (C-X-C motif) ligand 3 (14.8 kD) (Cxcl3) alternative variant bSep08, complete mRNA.
<a href="#">Cxcl13</a>	<a href="#">Cxcl13.aSep08</a>	<a href="#">498335</a>	5071	1126	2	109	chemokine (C-X-C motif) ligand 13 (12.1 kD) (Cxcl13) alternative variant aSep08, mRNA.
<a href="#">Cxcl13</a>	<a href="#">Cxcl13.cSep08</a>	<a href="#">498335</a>	4945	914	2	75	chemokine (C-X-C motif) ligand 13 (8.5 kD) (Cxcl13) alternative variant cSep08, mRNA.
<a href="#">Cxcl13</a>	<a href="#">Cxcl13.dSep08</a>	<a href="#">498335</a>	4819	741	1	41	chemokine (C-X-C motif) ligand 13 (Cxcl13) alternative variant dSep08, mRNA.
<a href="#">Cxcl14</a>	<a href="#">Cxcl14.bSep08</a>	<a href="#">306748</a>	4840	2085	2	64	chemokine (C-X-C motif) ligand 14 (6.5 kD) (Cxcl14) alternative variant bSep08, mRNA.
<a href="#">Cxcl17</a>	<a href="#">Cxcl17.bSep08</a>	<a href="#">308436</a>	11716	563		58	chemokine (C-X-C motif) ligand 17 (6.6 kD) (Cxcl17) alternative variant bSep08, mRNA.
<a href="#">Cxcr7</a>	<a href="#">Cxcr7.bSep08</a>	<a href="#">84348</a>	10504	1062	3	277	chemokine (C-X-C motif) receptor 7 (Cxcr7) alternative variant bSep08, mRNA.
<a href="#">Cxcr7</a>	<a href="#">Cxcr7.cSep08</a>	<a href="#">84348</a>	10432	901	3	272	chemokine (C-X-C motif) receptor 7 (Cxcr7) alternative variant cSep08, mRNA.
<a href="#">Cxxc1</a>	<a href="#">Cxxc1.aSep08</a>	<a href="#">291440</a>	5237	2693	14	698	CXXC finger 1 (79.7 kD) (Cxxc1) alternative variant aSep08, mRNA.
<a href="#">Cxxc1</a>	<a href="#">Cxxc1.bSep08</a>	<a href="#">291440</a>	3023	1845	9	350	CXXC finger 1 (41.2 kD) (Cxxc1) alternative variant bSep08, mRNA.

<a href="#">Cxxc1</a>	<a href="#">Cxxc1.cSep08</a>	<a href="#">291440</a>	2473	977	6	285	CXXC finger 1 (31.5 kD) (Cxxc1) alternative variant cSep08, mRNA.
<a href="#">Cxxc1</a>	<a href="#">Cxxc1.dSep08</a>	<a href="#">291440</a>	1845	856	6	285	CXXC finger 1 (Cxxc1) alternative variant dSep08, mRNA.
<a href="#">Cxxc1</a>	<a href="#">Cxxc1.eSep08</a>	<a href="#">291440</a>	1130	1002	2	167	CXXC finger 1 (Cxxc1) alternative variant eSep08, mRNA.
<a href="#">Cxxc1</a>	<a href="#">Cxxc1.fSep08</a>	<a href="#">291440</a>	1807	753	4	162	CXXC finger 1 (17.9 kD) (Cxxc1) alternative variant fSep08, mRNA.
<a href="#">Cxxc5</a>	<a href="#">Cxxc5.bSep08</a>	<a href="#">291670</a>	30499	2367	3	306	CXXC finger 5 (31.4 kD) (Cxxc5) alternative variant bSep08, mRNA.
<a href="#">Cyb5</a>	<a href="#">Cyb5.bSep08</a>	<a href="#">64001</a>	36496	571	4	96	cytochrome b-5 (Cyb5) alternative variant bSep08, mRNA.
<a href="#">Cyb5</a>	<a href="#">Cyb5.cSep08</a>	<a href="#">64001</a>	4390	1770	2	53	cytochrome b-5 (6.1 kD) (Cyb5) alternative variant cSep08, mRNA.
<a href="#">Cyb5</a>	<a href="#">Cyb5.dSep08</a>	<a href="#">64001</a>	17626	462	5	17	cytochrome b-5 (Cyb5) alternative variant dSep08, mRNA.
<a href="#">Cyb5b</a>	<a href="#">Cyb5b.bSep08</a>	<a href="#">80773</a>	3642	775	2	94	cytochrome b5 type B (10.4 kD) (Cyb5b) alternative variant bSep08, mRNA.
<a href="#">Cyb5r1</a>	<a href="#">Cyb5r1.bSep08</a>	<a href="#">304805</a>	3645	1155	5	125	cytochrome b5 reductase 1 CRA d (14.3 kD) (Cyb5r1) alternative variant bSep08, mRNA.
<a href="#">Cyb5r1</a>	<a href="#">Cyb5r1.cSep08</a>	<a href="#">304805</a>	815	420	3	83	cytochrome b5 reductase 1 CRA b (Cyb5r1) alternative variant cSep08, mRNA.
<a href="#">Cyb5r2</a>	<a href="#">Cyb5r2.bSep08</a>	<a href="#">365345</a>	5806	798	7	170	cytochrome b5 reductase 2 (Cyb5r2) alternative variant bSep08, mRNA.
<a href="#">Cyb5r3</a>	<a href="#">Cyb5r3.bSep08</a>	<a href="#">25035</a>	10702	739	1	246	cytochrome b5 reductase 3 (Cyb5r3) alternative variant bSep08, mRNA.
<a href="#">Cyb5r4</a>	<a href="#">Cyb5r4.aSep08</a>	<a href="#">171015</a>	65381	2551	3	564	cytochrome b5 reductase 4 CRA a (Cyb5r4) alternative variant aSep08, mRNA.
<a href="#">Cyb5r4</a>	<a href="#">Cyb5r4.bSep08</a>	<a href="#">171015</a>	49288	855	1	169	flavoheмоprotein b5 b5r (Cyb5r4) alternative variant bSep08, mRNA.
<a href="#">Cyb5r4</a>	<a href="#">Cyb5r4.cSep08</a>	<a href="#">171015</a>	70596	1796	2	97	putative protein (Cyb5r4) alternative variant cSep08, mRNA.
<a href="#">Cyb561</a>	<a href="#">Cyb561.bSep08</a>	<a href="#">303601</a>	2652	2144		151	cytochrome b-561 (Cyb561) alternative variant bSep08, mRNA.
<a href="#">Cyb561d2</a>	<a href="#">Cyb561d2.aSep08</a>	<a href="#">363137</a>	2190	745	2	248	cytochrome b561 (Cyb561d2) alternative variant aSep08, mRNA.
<a href="#">Cyba</a>	<a href="#">Cyba.bSep08</a>	<a href="#">79129</a>	1505	386	2	98	cytochrome b-245, alpha polypeptide (Cyba) alternative variant bSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.aSep08</a>	<a href="#">361729</a>	11635	2131	6	275	cytochrome b, ascorbate dependent 3 (Cybasc3) alternative variant aSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.bSep08</a>	<a href="#">361729</a>	5905	912	5	261	cytochrome b, ascorbate dependent 3 (Cybasc3) alternative variant bSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.dSep08</a>	<a href="#">361729</a>	5212	794	3	235	cytochrome b, ascorbate dependent 3 (Cybasc3) alternative variant dSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.eSep08</a>	<a href="#">361729</a>	6166	974	4	191	cytochrome b, ascorbate dependent 3 (21.3 kD) (Cybasc3) alternative variant eSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.fSep08</a>	<a href="#">361729</a>	5962	835	3	135	cytochrome b, ascorbate dependent 3 (15.0 kD) (Cybasc3) alternative variant fSep08, mRNA.
<a href="#">Cybasc3</a>	<a href="#">Cybasc3.gSep08</a>	<a href="#">361729</a>	8414	1074	4	130	cytochrome b, ascorbate dependent 3 (Cybasc3) alternative variant gSep08, mRNA.

<a href="#">Cybasc3</a>	<a href="#">Cybasc3.hSep08</a>	<a href="#">361729</a>	7885	712	4	78	cytochrome b, ascorbate dependent 3 (Cybasc3) alternative variant hSep08, mRNA.
<a href="#">Cyc1</a>	<a href="#">Cyc1.aSep08</a>	<a href="#">300047</a>	2001	900	6	291	cytochrome c-1 (Cyc1) alternative variant aSep08, mRNA.
<a href="#">Cyc1</a>	<a href="#">Cyc1.bSep08</a>	<a href="#">300047</a>	2361	1752	6	266	cytochrome c-1 (29.5 kD) (Cyc1) alternative variant bSep08, complete mRNA.
<a href="#">Cyc1</a>	<a href="#">Cyc1.eSep08</a>	<a href="#">300047</a>	1269	556	4	107	cytochrome c-1 (Cyc1) alternative variant eSep08, mRNA.
<a href="#">Cyc1</a>	<a href="#">Cyc1.gSep08</a>	<a href="#">300047</a>	1007	892	2	63	cytochrome c-1 (6.4 kD) (Cyc1) alternative variant gSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.bSep08</a>	<a href="#">308666</a>	7964	675	6	225	cytoplasmic FMR1 interacting protein 1 CRA a (Cyfip1) alternative variant bSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.bSep08</a>	<a href="#">691443</a>	7964	675	6	225	cytoplasmic FMR1 interacting protein 1 CRA a (Cyfip1) alternative variant bSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.cSep08</a>	<a href="#">308666</a>	11886	739	7	158	cytoplasmic fmr1 interacting protein 1 CRA b (Cyfip1) alternative variant cSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.cSep08</a>	<a href="#">691443</a>	11886	739	7	158	cytoplasmic fmr1 interacting protein 1 CRA b (Cyfip1) alternative variant cSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.dSep08</a>	<a href="#">308666</a>	31893	420	3	84	cytoplasmic fmr1 interacting protein 1 CRA a (9.4 kD) (Cyfip1) alternative variant dSep08, mRNA.
<a href="#">Cyfip1</a>	<a href="#">Cyfip1.dSep08</a>	<a href="#">691443</a>	31893	420	3	84	cytoplasmic fmr1 interacting protein 1 CRA a (9.4 kD) (Cyfip1) alternative variant dSep08, mRNA.
<a href="#">Cyhr1</a>	<a href="#">Cyhr1.aSep08</a>	<a href="#">315097</a>	1659	923	2	192	cysteine and histidine rich 1 (20.2 kD) (Cyhr1) alternative variant aSep08, mRNA.
<a href="#">Cyld</a>	<a href="#">Cyld.aSep08</a>	<a href="#">312937</a>	25580	2424		383	cyliodromatosis (turban tumor syndrome) (Cyld) mRNA.
<a href="#">Cyp2a1</a>	<a href="#">Cyp2a1.bSep08</a>	<a href="#">24894</a>	8343	821	5	273	cytochrome P450 IIA1 (hepatic steroid hydroxylase IIA1) gene (Cyp2a1) alternative variant bSep08, mRNA.
<a href="#">Cyp2a3a</a>	<a href="#">Cyp2a3a.bSep08</a>	<a href="#">24299</a>	1611	428		75	cytochrome P450, family 2, subfamily A, polypeptide 3a (Cyp2a3a) alternative variant bSep08, mRNA.
<a href="#">Cyp2b2</a>	<a href="#">Cyp2b2.aSep08</a>	<a href="#">361523</a>	13797	1691	8	491	cytochrome P450, family 2, subfamily b, polypeptide 2 (Cyp2b2) alternative variant aSep08, mRNA.
<a href="#">Cyp2b2</a>	<a href="#">Cyp2b2.bSep08</a>	<a href="#">361523</a>	738	300	1	99	cytochrome P450, family 2, subfamily b, polypeptide 2 (Cyp2b2) alternative variant bSep08, mRNA.
<a href="#">Cyp2b3</a>	<a href="#">Cyp2b3.bSep08</a>	<a href="#">286953</a>	31045	739	1	216	cytochrome P450IIB3 (24.6 kD) (Cyp2b3) alternative variant bSep08, complete mRNA.
<a href="#">Cyp2b21</a>	<a href="#">Cyp2b21.aSep08</a>	<a href="#">292728</a>	28979	1784		575	cytochrome P450, family 2, subfamily b, polypeptide 21 (Cyp2b21) alternative variant aSep08, mRNA.
<a href="#">Cyp2c</a>	<a href="#">Cyp2c.bSep08</a>	<a href="#">29277</a>	1289	547	2	118	cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase) (Cyp2c) alternative variant bSep08, mRNA.
<a href="#">Cyp2c</a>	<a href="#">Cyp2c.cSep08</a>	<a href="#">29277</a>	1038	254	2	84	cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase) (Cyp2c) alternative variant cSep08, mRNA.
<a href="#">Cyp2c6</a>	<a href="#">Cyp2c6.aSep08</a>	<a href="#">246070</a>	3562	794		177	cytochrome P450, subfamily IIC6 (Cyp2c6) mRNA.
<a href="#">Cyp2c7</a>	<a href="#">Cyp2c7.aSep08</a>	<a href="#">29298</a>	19751	906	2	199	cytochrome P450, family 2, subfamily c, polypeptide 7 (22.6 kD) (Cyp2c7) alternative variant aSep08, mRNA.
<a href="#">Cyp2c7</a>	<a href="#">Cyp2c7.bSep08</a>	<a href="#">29298</a>	615727	913	2	199	cytochrome P450, family 2, subfamily c, polypeptide 7 (22.7 kD) (Cyp2c7) alternative variant bSep08, complete mRNA.
<a href="#">Cyp2c12</a>	<a href="#">Cyp2c12.bSep08</a>	<a href="#">25011</a>	47998	1066	3	287	cytochrome P450, family 2, subfamily c, polypeptide 12 (Cyp2c12) alternative variant bSep08, mRNA.

<a href="#">Cyp2c12</a>	<a href="#">Cyp2c12.cSep08</a>	<a href="#">25011</a>	9599	1111	1	119	cytochrome P450, family 2, subfamily c, polypeptide 12 (Cyp2c12) alternative variant cSep08, mRNA.
<a href="#">Cyp2c13</a>	<a href="#">Cyp2c13.bSep08</a>	<a href="#">171521</a>	5776	296	3	98	cytochrome P450 2c13 (Cyp2c13) alternative variant bSep08, mRNA.
<a href="#">Cyp2c13</a>	<a href="#">Cyp2c13.cSep08</a>	<a href="#">171521</a>	18124	290	3	96	cytochrome P450 2c13 (Cyp2c13) alternative variant cSep08, mRNA.
<a href="#">Cyp2c22</a>	<a href="#">Cyp2c22.bSep08</a>	<a href="#">171518</a>	17136	732	2	244	cytochrome P450, family 2, subfamily c, polypeptide 22 (Cyp2c22) alternative variant bSep08, mRNA.
<a href="#">Cyp2c22</a>	<a href="#">Cyp2c22.cSep08</a>	<a href="#">171518</a>	24548	914	3	231	cytochrome P450, family 2, subfamily c, polypeptide 22 (Cyp2c22) alternative variant cSep08, mRNA.
<a href="#">Cyp2c23</a>	<a href="#">Cyp2c23.bSep08</a>	<a href="#">83790</a>	5527	843	2	133	cytochrome P-450 S-mephenytoin (Cyp2c23) alternative variant bSep08, mRNA.
<a href="#">Cyp2c23</a>	<a href="#">Cyp2c23.cSep08</a>	<a href="#">83790</a>	6548	645	4	106	cytochrome 2C78 (Cyp2c23) alternative variant cSep08, mRNA.
<a href="#">Cyp2c23</a>	<a href="#">Cyp2c23.eSep08</a>	<a href="#">83790</a>	2637	919	3	101	cytochrome 2C78 (11.5 kD) (Cyp2c23) alternative variant eSep08, mRNA.
<a href="#">Cyp2c23</a>	<a href="#">Cyp2c23.fSep08</a>	<a href="#">83790</a>	2656	1047	2	101	cytochrome 2C78 (11.5 kD) (Cyp2c23) alternative variant fSep08, mRNA.
<a href="#">Cyp2c23</a>	<a href="#">Cyp2c23.gSep08</a>	<a href="#">83790</a>	1045	280	2	93	putative protein (Cyp2c23) alternative variant gSep08, mRNA.
<a href="#">Cyp2c24</a>	<a href="#">Cyp2c24.aSep08</a>	<a href="#">499353</a>	3385	454		151	cytochrome P450, family 2, subfamily c, polypeptide 24 (Cyp2c24) mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.aSep08</a>	<a href="#">29296</a>	47247	1753	9	286	cytochrome P450, 2c37 (32.9 kD) (Cyp2c37) alternative variant aSep08, mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.bSep08</a>	<a href="#">29296</a>	21982	727	4	147	cytochrome P450, 2c37 (17.3 kD) (Cyp2c37) alternative variant bSep08, mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.cSep08</a>	<a href="#">29296</a>	1952	341	2	113	cytochrome P450, 2c37 (Cyp2c37) alternative variant cSep08, mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.dSep08</a>	<a href="#">29296</a>	1846	237	3	79	cytochrome P450, 2c37 (Cyp2c37) alternative variant dSep08, mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.eSep08</a>	<a href="#">29296</a>	1970	207	2	68	cytochrome P450, 2c37 (Cyp2c37) alternative variant eSep08, mRNA.
<a href="#">Cyp2c37</a>	<a href="#">Cyp2c37.hSep08</a>	<a href="#">29296</a>	309	217	2	19	cytochrome P450, 2c37 (2.4 kD) (Cyp2c37) alternative variant hSep08, mRNA.
<a href="#">Cyp2c77-ps</a>	<a href="#">Cyp2c77-ps.aSep08</a>	<a href="#">686022</a>	51260	1000	7	142	cytochrome P450, family 2, subfamily c, polypeptide 77, pseudogene (Cyp2c77-ps) alternative variant aSep08, mRNA.
<a href="#">Cyp2c77-ps</a>	<a href="#">Cyp2c77-ps.bSep08</a>	<a href="#">686022</a>	3264	522	3	135	cytochrome P450, family 2, subfamily c, polypeptide 77, pseudogene (Cyp2c77-ps) alternative variant bSep08, mRNA.
<a href="#">Cyp2c77-ps</a>	<a href="#">Cyp2c77-ps.cSep08</a>	<a href="#">686022</a>	2085	343	2	110	cytochrome P450, family 2, subfamily c, polypeptide 77, pseudogene (Cyp2c77-ps) alternative variant cSep08, mRNA.
<a href="#">Cyp2c77-ps</a>	<a href="#">Cyp2c77-ps.dSep08</a>	<a href="#">686022</a>	20859	351	3		
<a href="#">Cyp2c80</a>	<a href="#">Cyp2c80.aSep08</a>	<a href="#">292330</a>	18733	1590		341	cytochrome P450, family 2, subfamily c, polypeptide 80 (Cyp2c80) mRNA.

<a href="#">Cyp2d2</a>	<a href="#">Cyp2d2.bSep08</a>	<a href="#">25053</a>	1730	877	3	195	p450 cytochrome 2d2 (Cyp2d2) alternative variant bSep08, mRNA.
<a href="#">Cyp2d2</a>	<a href="#">Cyp2d2.cSep08</a>	<a href="#">25053</a>	1524	1308	3	130	p450 cytochrome 2d2 (14.9 kD) (Cyp2d2) alternative variant cSep08, mRNA.
<a href="#">Cyp2d3</a>	<a href="#">Cyp2d3.bSep08</a>	<a href="#">24303</a>	1684	828	1	187	cytochrome P450, family 2, subfamily d, polypeptide 3 (21.0 kD) (Cyp2d3) alternative variant bSep08, mRNA.
<a href="#">Cyp2d4v1</a>	<a href="#">Cyp2d4v1.bSep08</a>	<a href="#">171522</a>	1111	539	4	149	cytochrome p450 (Cyp2d4v1) alternative variant bSep08, mRNA.
<a href="#">Cyp2d4v1</a>	<a href="#">Cyp2d4v1.cSep08</a>	<a href="#">171522</a>	787	687	2	115	cytochrome p450 (12.9 kD) (Cyp2d4v1) alternative variant cSep08, mRNA.
<a href="#">Cyp2d5</a>	<a href="#">Cyp2d5.bSep08</a>	<a href="#">286963</a>	1908	777	5	259	cytochrome P450, family 2, subfamily d, polypeptide 5 (Cyp2d5) alternative variant bSep08, mRNA.
<a href="#">Cyp2e1</a>	<a href="#">Cyp2e1.bSep08</a>	<a href="#">25086</a>	5709	863	1	285	cytochrome P450, family 2, subfamily e, polypeptide 1 (Cyp2e1) alternative variant bSep08, mRNA.
<a href="#">Cyp2f4</a>	<a href="#">Cyp2f4.bSep08</a>	<a href="#">54246</a>	5448	751	3	221	cytochrome P450, family 2, subfamily f, polypeptide 4 (Cyp2f4) alternative variant bSep08, mRNA.
<a href="#">Cyp2f4</a>	<a href="#">Cyp2f4.cSep08</a>	<a href="#">54246</a>	10103	660	5	184	cytochrome P450, family 2, subfamily f, polypeptide 4 (Cyp2f4) alternative variant cSep08, mRNA.
<a href="#">Cyp2f4</a>	<a href="#">Cyp2f4.dSep08</a>	<a href="#">54246</a>	5239	615	3	106	cytochrome P450, family 2, subfamily f, polypeptide 4 (Cyp2f4) alternative variant dSep08, mRNA.
<a href="#">Cyp2j10</a>	<a href="#">Cyp2j10.aSep08</a>	<a href="#">313373</a>	36914	1926	8	502	cytochrome P450, family 2, subfamily j, polypeptide 10 (57.7 kD) (Cyp2j10) alternative variant aSep08, mRNA.
<a href="#">Cyp2j10</a>	<a href="#">Cyp2j10.bSep08</a>	<a href="#">313373</a>	2524	754	1	82	cytochrome P450, family 2, subfamily j, polypeptide 10 (9.0 kD) (Cyp2j10) alternative variant bSep08, mRNA.
<a href="#">Cyp2j13</a>	<a href="#">Cyp2j13.aSep08</a>	<a href="#">313372</a>	18073	1018		167	cytochrome P450, family 2, subfamily j, polypeptide 13 (Cyp2j13) mRNA.
<a href="#">Cyp2r1</a>	<a href="#">Cyp2r1.aSep08</a>	<a href="#">361631</a>	13791	2813	3	431	cytochrome P450, family 2, subfamily r, polypeptide 1 (Cyp2r1) alternative variant aSep08, mRNA.
<a href="#">Cyp2r1</a>	<a href="#">Cyp2r1.cSep08</a>	<a href="#">361631</a>	9247	383	2	127	cytochrome P450, family 2, subfamily r, polypeptide 1 (Cyp2r1) alternative variant cSep08, mRNA.
<a href="#">Cyp2t1</a>	<a href="#">Cyp2t1.bSep08</a>	<a href="#">171380</a>	1703	701	5	197	cytochrome p450 monooxygenase CYP2T1 CRA a (Cyp2t1) alternative variant bSep08, mRNA.
<a href="#">Cyp2t1</a>	<a href="#">Cyp2t1.cSep08</a>	<a href="#">171380</a>	1534	392	2	97	cytochrome P450 monooxygenase CYP2T1 like (10.8 kD) (Cyp2t1) alternative variant cSep08, mRNA.
<a href="#">Cyp2t1</a>	<a href="#">Cyp2t1.dSep08</a>	<a href="#">171380</a>	961	740	2	90	cytochrome p450 monooxygenase CYP2T1 CRA a (Cyp2t1) alternative variant dSep08, mRNA.
<a href="#">Cyp2w1</a>	<a href="#">Cyp2w1.aSep08</a>	<a href="#">288517</a>	1680	457		151	cytochrome P450, family 2, subfamily W, polypeptide 1 (Cyp2w1) mRNA.
<a href="#">Cyp3a2</a>	<a href="#">Cyp3a2.bSep08</a>	<a href="#">266682</a>	3389	629	1	64	cytochrome P450, family 3, subfamily a, polypeptide 2 (Cyp3a2) alternative variant bSep08, mRNA.
<a href="#">Cyp3a2</a>	<a href="#">Cyp3a2.cSep08</a>	<a href="#">266682</a>	873	220	1	30	cytochrome P450, family 3, subfamily a, polypeptide 2 (3.5 kD) (Cyp3a2) alternative variant cSep08, mRNA.
<a href="#">Cyp3a9</a>	<a href="#">Cyp3a9.bSep08</a>	<a href="#">171352</a>	14088	774	1	101	cytochrome P450, family 3, subfamily a, polypeptide 9 (11.4 kD) (Cyp3a9) alternative variant bSep08, mRNA.
<a href="#">Cyp3a23/3a1</a>	<a href="#">Cyp3a23/3a1.bSep08</a>	<a href="#">25642</a>	5341	927	6	220	cytochrome p450 (Cyp3a23/3a1) alternative variant bSep08, mRNA.

<a href="#">Cyp3a23/3a1</a>	<a href="#">Cyp3a23/3a1.bSep08</a>	<a href="#">286929</a>	5341	927	6	220	cytochrome p450 (Cyp3a23/3a1) alternative variant bSep08, mRNA.
<a href="#">Cyp4a3</a>	<a href="#">Cyp4a3.aSep08</a>	<a href="#">298423</a>	8458	2143	9	386	cytochrome P450, family 4, subfamily a, polypeptide 3 (Cyp4a3) alternative variant aSep08, mRNA.
<a href="#">Cyp4a3</a>	<a href="#">Cyp4a3.bSep08</a>	<a href="#">298423</a>	11808	967	7	317	cytochrome P450, family 4, subfamily a, polypeptide 3 (Cyp4a3) alternative variant bSep08, mRNA.
<a href="#">Cyp4a10.1</a>	<a href="#">Cyp4a10.1.bSep08</a>	<a href="#">50549</a>	5877	847	2	257	cytochrome P450, family 4, subfamily a, polypeptide 10 and cytochrome P450, family 4, subfamily a, polypeptide 1 (Cyp4a10.1) alternative variant bSep08, mRNA.
<a href="#">Cyp4a10.1</a>	<a href="#">Cyp4a10.1.bSep08</a>	<a href="#">170544</a>	5877	847	2	257	cytochrome P450, family 4, subfamily a, polypeptide 10 and cytochrome P450, family 4, subfamily a, polypeptide 1 (Cyp4a10.1) alternative variant bSep08, mRNA.
<a href="#">Cyp4a10.1</a>	<a href="#">Cyp4a10.1.cSep08</a>	<a href="#">50549</a>	12131	1145	6	254	cytochrome P450, family 4, subfamily a, polypeptide 10 and cytochrome P450, family 4, subfamily a, polypeptide 1 (28.9 kD) (Cyp4a10.1) alternative variant cSep08, mRNA.
<a href="#">Cyp4a10.1</a>	<a href="#">Cyp4a10.1.cSep08</a>	<a href="#">170544</a>	12131	1145	6	254	cytochrome P450, family 4, subfamily a, polypeptide 10 and cytochrome P450, family 4, subfamily a, polypeptide 1 (28.9 kD) (Cyp4a10.1) alternative variant cSep08, mRNA.
<a href="#">Cyp4b1</a>	<a href="#">Cyp4b1.bSep08</a>	<a href="#">24307</a>	5640	709	1	236	cytochrome P450, family 4, subfamily b, polypeptide 1 (Cyp4b1) alternative variant bSep08, mRNA.
<a href="#">Cyp4f1</a>	<a href="#">Cyp4f1.bSep08</a>	<a href="#">56266</a>	5127	911	1	220	cytochrome P450, family 4, subfamily f, polypeptide 1 (Cyp4f1) alternative variant bSep08, mRNA.
<a href="#">Cyp4f1</a>	<a href="#">Cyp4f1.cSep08</a>	<a href="#">56266</a>	1418	256		84	cytochrome P450, family 4, subfamily f, polypeptide 1 (Cyp4f1) alternative variant cSep08, mRNA.
<a href="#">Cyp4f6</a>	<a href="#">Cyp4f6.bSep08</a>	<a href="#">266689</a>	8926	1166	7	263	cytochrome p450 4F6 (30.3 kD) (Cyp4f6) alternative variant bSep08, mRNA.
<a href="#">Cyp4f6</a>	<a href="#">Cyp4f6.cSep08</a>	<a href="#">266689</a>	9068	1443	7	212	cytochrome P450 4F6 (24.2 kD) (Cyp4f6) alternative variant cSep08, mRNA.
<a href="#">Cyp4f6</a>	<a href="#">Cyp4f6.dSep08</a>	<a href="#">266689</a>	20479	839	6	181	cytochrome P450 4F6 (Cyp4f6) alternative variant dSep08, mRNA.
<a href="#">Cyp4f17</a>	<a href="#">Cyp4f17.aSep08</a>	<a href="#">500801</a>	6167	1590	2	335	cytochrome P450, family 4, subfamily f, polypeptide 17 (Cyp4f17) alternative variant aSep08, mRNA.
<a href="#">Cyp4f17</a>	<a href="#">Cyp4f17.bSep08</a>	<a href="#">500801</a>	1769	853	1	140	cytochrome P450, family 4, subfamily f, polypeptide 17 (Cyp4f17) alternative variant bSep08, mRNA.
<a href="#">Cyp4f37</a>	<a href="#">Cyp4f37.aSep08</a>	<a href="#">691312</a>	3969	745		183	cytochrome P450, family 4, subfamily f, polypeptide 37 (Cyp4f37) mRNA.
<a href="#">Cyp4f39</a>	<a href="#">Cyp4f39.aSep08</a>	<a href="#">299566</a>	16320	756		251	cytochrome P450, family 4, subfamily f, polypeptide 39 (Cyp4f39) mRNA.
<a href="#">Cyp4v3</a>	<a href="#">Cyp4v3.aSep08</a>	<a href="#">266761</a>	24938	3191		525	cytochrome P450, family 4, subfamily v, polypeptide 3 (60.6 kD) (Cyp4v3) mRNA.
<a href="#">Cyp11a1</a>	<a href="#">Cyp11a1.bSep08</a>	<a href="#">29680</a>	4647	700	5	233	cytochrome P450, family 11, subfamily a, polypeptide 1 (Cyp11a1) alternative variant bSep08, mRNA.
<a href="#">Cyp11a1</a>	<a href="#">Cyp11a1.cSep08</a>	<a href="#">29680</a>	2514	1355	4	197	cytochrome P450, family 11, subfamily a, polypeptide 1 (22.9 kD) (Cyp11a1) alternative variant cSep08, mRNA.
<a href="#">Cyp11a1</a>	<a href="#">Cyp11a1.dSep08</a>	<a href="#">29680</a>	4554	708	4	150	cytochrome P450, family 11, subfamily a, polypeptide 1 (Cyp11a1) alternative variant dSep08, mRNA.



<a href="#">Cyp11a1</a>	<a href="#">Cyp11a1.eSep08</a>	<a href="#">29680</a>	6128	760	6	148	cytochrome P450, family 11, subfamily a, polypeptide 1 (Cyp11a1) alternative variant eSep08, mRNA.
<a href="#">Cyp11a1</a>	<a href="#">Cyp11a1.fSep08</a>	<a href="#">29680</a>	5388	763	3		
<a href="#">Cyp17a1</a>	<a href="#">Cyp17a1.bSep08</a>	<a href="#">25146</a>	2508	855	5	285	cytochrome P450, family 17, subfamily a, polypeptide 1 (Cyp17a1) alternative variant bSep08, mRNA.
<a href="#">Cyp17a1</a>	<a href="#">Cyp17a1.cSep08</a>	<a href="#">25146</a>	4085	870	4	215	cytochrome P450, family 17, subfamily a, polypeptide 1 (Cyp17a1) alternative variant cSep08, mRNA.
<a href="#">Cyp20a1</a>	<a href="#">Cyp20a1.bSep08</a>	<a href="#">316435</a>	40702	725		148	cytochrome P450, family 20, subfamily A, polypeptide 1 (Cyp20a1) alternative variant bSep08, mRNA.
<a href="#">Cyp24a1</a>	<a href="#">Cyp24a1.bSep08</a>	<a href="#">25279</a>	8580	780	7	223	cytochrome P450 family 24 subfamily a polypeptide 1 (Cyp24a1) alternative variant bSep08, mRNA.
<a href="#">Cyp24a1</a>	<a href="#">Cyp24a1.cSep08</a>	<a href="#">25279</a>	1640	441	3	124	putative protein (Cyp24a1) alternative variant cSep08, mRNA.
<a href="#">Cyp27a1</a>	<a href="#">Cyp27a1.bSep08</a>	<a href="#">301517</a>	2083	693	2	121	cytochrome p450 family 27 subfamily a polypeptide 1 (Cyp27a1) alternative variant bSep08, mRNA.
<a href="#">Cyp27a1</a>	<a href="#">Cyp27a1.cSep08</a>	<a href="#">301517</a>	1981	739	1	110	cytochrome p450 family 27 subfamily a polypeptide 1 (Cyp27a1) alternative variant cSep08, mRNA.
<a href="#">Cyp46a1</a>	<a href="#">Cyp46a1.aSep08</a>	<a href="#">362782</a>	27169	2160	15	500	cytochrome P450, family 46, subfamily a, polypeptide 1 (56.7 kD) (Cyp46a1) alternative variant aSep08, mRNA.
<a href="#">Cystatin.0</a>	<a href="#">Cystatin.0.aSep08</a>		1961	437		104	kininogen 2 (Cystatin.0) mRNA.
<a href="#">Cystatin.1</a>	<a href="#">Cystatin.1.aSep08</a>		3930	404		134	histidine-rich glycoprotein (Cystatin.1) mRNA.
<a href="#">Cyt11</a>	<a href="#">Cyt11.bSep08</a>	<a href="#">498392</a>	2255	727	2	110	cytokine like 1 (Cyt11) alternative variant bSep08, mRNA.
<a href="#">D2hgdh</a>	<a href="#">D2hgdh.aSep08</a>	<a href="#">301624</a>	18588	2811	9	542	D-2-hydroxyglutarate dehydrogenase (D2hgdh) alternative variant aSep08, mRNA.
<a href="#">D2hgdh</a>	<a href="#">D2hgdh.cSep08</a>	<a href="#">301624</a>	8968	833	5	277	D-2-hydroxyglutarate dehydrogenase (D2hgdh) alternative variant cSep08, mRNA.
<a href="#">D2hgdh</a>	<a href="#">D2hgdh.dSep08</a>	<a href="#">301624</a>	1055	314	2	104	D-2-hydroxyglutarate dehydrogenase (D2hgdh) alternative variant dSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.bSep08</a>	<a href="#">314212</a>	4673	1034	3	181	activator of morphogenesis 1 (21.2 kD) (Daam1) alternative variant bSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.cSep08</a>	<a href="#">314212</a>	8788	526	4	149	activator of morphogenesis 1 (Daam1) alternative variant cSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.dSep08</a>	<a href="#">314212</a>	112987	610	5	136	activator of morphogenesis 1 (Daam1) alternative variant dSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.fSep08</a>	<a href="#">314212</a>	22046	452	3	106	activator of morphogenesis 1 (Daam1) alternative variant fSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.gSep08</a>	<a href="#">314212</a>	84338	384	3	75	activator of morphogenesis 1 (Daam1) alternative variant gSep08, mRNA.
<a href="#">Daam1</a>	<a href="#">Daam1.hSep08</a>	<a href="#">314212</a>	63711	426	3	71	putative protein (Daam1) alternative variant hSep08, mRNA.
<a href="#">Daam2</a>	<a href="#">Daam2.aSep08</a>	<a href="#">316201</a>	7620	1800	5	509	dishevelled associated activator of morphogenesis 2 (Daam2) alternative variant aSep08, mRNA.
<a href="#">Daam2</a>	<a href="#">Daam2.bSep08</a>	<a href="#">316201</a>	19190	1299	7	290	dishevelled associated activator of morphogenesis 2 (Daam2) alternative variant bSep08, mRNA.
<a href="#">Daam2</a>	<a href="#">Daam2.cSep08</a>	<a href="#">316201</a>	2167	553	3	154	dishevelled associated activator of morphogenesis 2 (Daam2) alternative variant cSep08, mRNA.

<a href="#">Daam2</a>	<a href="#">Daam2.dSep08</a>	<a href="#">316201</a>	3899	383	1	113	dishevelled associated activator of morphogenesis 2 (Daam2) alternative variant dSep08, mRNA.
<a href="#">Daam2</a>	<a href="#">Daam2.fSep08</a>	<a href="#">316201</a>	4902	542	5	48	dishevelled associated activator of morphogenesis 2 (Daam2) alternative variant fSep08, mRNA.
<a href="#">Dab1</a>	<a href="#">Dab1.bSep08</a>	<a href="#">266729</a>	309892	1783	5	535	disabled homolog 1 (Drosophila) (Dab1) alternative variant bSep08, mRNA.
<a href="#">Dab1</a>	<a href="#">Dab1.cSep08</a>	<a href="#">266729</a>	14307	1855	4	132	disabled homolog 1 (Drosophila) (Dab1) alternative variant cSep08, mRNA.
<a href="#">Dab1</a>	<a href="#">Dab1.dSep08</a>	<a href="#">266729</a>	1679	571	3	115	disabled homolog 1 (Drosophila) (Dab1) alternative variant dSep08, mRNA.
<a href="#">Dab2</a>	<a href="#">Dab2.bSep08</a>	<a href="#">79128</a>	6126	2007	2	182	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) (Dab2) alternative variant bSep08, mRNA.
<a href="#">Dab2</a>	<a href="#">Dab2.cSep08</a>	<a href="#">79128</a>	2474	570	5	127	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) (Dab2) alternative variant cSep08, mRNA.
<a href="#">Dab2ip</a>	<a href="#">Dab2ip.bSep08</a>	<a href="#">192126</a>	18173	674	4	224	disabled homolog 2 (Drosophila) interacting protein (Dab2ip) alternative variant bSep08, mRNA.
<a href="#">Dab2ip</a>	<a href="#">Dab2ip.cSep08</a>	<a href="#">192126</a>	1544	515	2	171	disabled homolog 2 (Drosophila) interacting protein (Dab2ip) alternative variant cSep08, mRNA.
<a href="#">Dab2ip</a>	<a href="#">Dab2ip.dSep08</a>	<a href="#">192126</a>	5223	750	3	151	disabled homolog 2 (Drosophila) interacting protein (Dab2ip) alternative variant dSep08, mRNA.
<a href="#">Dab2ip</a>	<a href="#">Dab2ip.eSep08</a>	<a href="#">192126</a>	933	444	2	47	disabled homolog 2 (Drosophila) interacting protein (Dab2ip) alternative variant eSep08, mRNA.
<a href="#">daby</a>	<a href="#">daby.aSep08</a>		7530	684		114	dedicator of cytokinesis 11 (daby) mRNA.
<a href="#">dachy</a>	<a href="#">dachy.aSep08</a>		563	320		97	putative protein (dachy) mRNA.
<a href="#">Dact1</a>	<a href="#">Dact1.aSep08</a>	<a href="#">500666</a>	10251	2306	4	656	dapper homolog 1, antagonist of beta-catenin (xenopus) (Dact1) alternative variant aSep08, mRNA.
<a href="#">Dact1</a>	<a href="#">Dact1.cSep08</a>	<a href="#">500666</a>	8321	648	1	48	dapper homolog 1, antagonist of beta-catenin (xenopus) (Dact1) alternative variant cSep08, mRNA.
<a href="#">Dact2</a>	<a href="#">Dact2.bSep08</a>	<a href="#">308212</a>	7197	462	1	154	dapper homolog 2, antagonist of beta-catenin (xenopus) (Dact2) alternative variant bSep08, mRNA.
<a href="#">dafer</a>	<a href="#">dafer.aSep08</a>		29072	320		54	putative protein (dafer) mRNA.
<a href="#">daflo</a>	<a href="#">daflo.aSep08</a>		23697	498		69	putative protein (daflo) mRNA.
<a href="#">daflu</a>	<a href="#">daflu.aSep08</a>		604	410		94	putative mitochondrial protein (10.0 kD) (daflu) mRNA.
<a href="#">Dag1</a>	<a href="#">Dag1.aSep08</a>	<a href="#">114489</a>	12684	5171		893	dystroglycan 1 (96.7 kD) (Dag1) mRNA.
<a href="#">DAGK_acc.0</a>	<a href="#">DAGK_acc.0.aSep08</a>		8172	337		112	diacylglycerol kinase eta CRA d (DAGK_acc.0) mRNA.
<a href="#">DAGK_acc.1</a>	<a href="#">DAGK_acc.1.aSep08</a>		2764	2192		184	diacylglycerol kinase theta CRA a (DAGK_acc.1) mRNA.
<a href="#">DAGK_cat.0</a>	<a href="#">DAGK_cat.0.aSep08</a>		1047	709	2	149	diacylglycerol kinase theta (DAGK_cat.0) alternative variant aSep08, mRNA.
<a href="#">DAGK_cat.0</a>	<a href="#">DAGK_cat.0.bSep08</a>		495	285	1	94	diacylglycerol kinase theta (DAGK_cat.0) alternative variant bSep08, mRNA.
<a href="#">DAGK_cat.1</a>	<a href="#">DAGK_cat.1.aSep08</a>		7276	678	5	226	diacylglycerol kinase delta (DAGK_cat.1) alternative variant aSep08, mRNA.
<a href="#">DAGK_cat.1</a>	<a href="#">DAGK_cat.1.bSep08</a>		2833	537	1	127	diacylglycerol kinase delta (DAGK_cat.1) alternative variant bSep08, mRNA.
<a href="#">Dagla</a>	<a href="#">Dagla.bSep08</a>	<a href="#">309207</a>	1582	522	2	132	diacylglycerol lipase, alpha (Dagla) alternative variant bSep08, mRNA.

<a href="#">Daglb</a>	<a href="#">Daglb.bSep08</a>	<a href="#">304289</a>	8736	808	2	261	diacylglycerol lipase, beta (Daglb) alternative variant bSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.bSep08</a>	<a href="#">361730</a>	13855	2990	18	365	dihydroxyacetone kinase 2 (37.9 kD) (Dak) alternative variant bSep08, complete mRNA.
<a href="#">Dak</a>	<a href="#">Dak.cSep08</a>	<a href="#">361730</a>	10688	1161	11	308	dihydroxyacetone kinase 2 (Dak) alternative variant cSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.dSep08</a>	<a href="#">361730</a>	7579	802	6	267	dihydroxyacetone kinase 2 (Dak) alternative variant dSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.eSep08</a>	<a href="#">361730</a>	8694	768	7	255	dihydroxyacetone kinase 2 (Dak) alternative variant eSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.fSep08</a>	<a href="#">361730</a>	3184	794	9	241	dihydroxyacetone kinase 2 (Dak) alternative variant fSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.gSep08</a>	<a href="#">361730</a>	13235	448	4	122	dihydroxyacetone kinase 2 (Dak) alternative variant gSep08, mRNA.
<a href="#">Dak</a>	<a href="#">Dak.hSep08</a>	<a href="#">361730</a>	3229	650	4	95	putative protein (Dak) alternative variant hSep08, mRNA.
<a href="#">dakee</a>	<a href="#">dakee.aSep08</a>		28285	1001		148	phosphatidylinositol glycan anchor biosynthesis class N (dakee) mRNA.
<a href="#">daloy</a>	<a href="#">daloy.aSep08</a>		16889	742	2	162	leukemia nup98 fusion partner 1 like (daloy) alternative variant aSep08, mRNA.
<a href="#">daloy</a>	<a href="#">daloy.bSep08</a>		20637	537	3	68	putative protein of mammalian origin (7.8 kD) (daloy) alternative variant bSep08, mRNA.
<a href="#">Dalrd3</a>	<a href="#">Dalrd3.bSep08</a>	<a href="#">363146</a>	1549	1435	2	192	putative protein of vertebrate origin (21.0 kD) (Dalrd3) alternative variant bSep08, mRNA.
<a href="#">Dalrd3</a>	<a href="#">Dalrd3.cSep08</a>	<a href="#">363146</a>	1194	1009	3	63	putative protein of metazoan origin (7.3 kD) (Dalrd3) alternative variant cSep08, mRNA.
<a href="#">damer</a>	<a href="#">damer.aSep08</a>		1589	938		43	putative protein human specific (4.7 kD) (damer) mRNA.
<a href="#">danoy</a>	<a href="#">danoy.aSep08</a>		39244	1035		51	putative protein (6.0 kD) (danoy) mRNA.
<a href="#">Dao1</a>	<a href="#">Dao1.aSep08</a>	<a href="#">114027</a>	20136	1100	6	329	D-amino acid oxidase 1 (Dao1) alternative variant aSep08, mRNA.
<a href="#">Dao1</a>	<a href="#">Dao1.bSep08</a>	<a href="#">114027</a>	12901	622	1	207	D-amino acid oxidase 1 (Dao1) alternative variant bSep08, mRNA.
<a href="#">Dap</a>	<a href="#">Dap.bSep08</a>	<a href="#">64322</a>	13335	1728	3	97	death-associated protein (Dap) alternative variant bSep08, mRNA.
<a href="#">Dap</a>	<a href="#">Dap.dSep08</a>	<a href="#">64322</a>	38362	438	3	56	death-associated protein (6.2 kD) (Dap) alternative variant dSep08, complete mRNA.
<a href="#">Dap3</a>	<a href="#">Dap3.bSep08</a>	<a href="#">295238</a>	26914	1109	8	197	death associated protein 3 (22.4 kD) (Dap3) alternative variant bSep08, mRNA.
<a href="#">Dap3</a>	<a href="#">Dap3.cSep08</a>	<a href="#">295238</a>	21317	740	7	194	death associated protein 3 (Dap3) alternative variant cSep08, mRNA.
<a href="#">Dap3</a>	<a href="#">Dap3.dSep08</a>	<a href="#">295238</a>	396	321	2	106	death associated protein 3 (Dap3) alternative variant dSep08, mRNA.
<a href="#">Dap3</a>	<a href="#">Dap3.eSep08</a>	<a href="#">295238</a>	2174	835	2	74	death associated protein 3 (8.8 kD) (Dap3) alternative variant eSep08, mRNA.
<a href="#">Dap3</a>	<a href="#">Dap3.fSep08</a>	<a href="#">295238</a>	2967	702	3	42	death associated protein 3 (Dap3) alternative variant fSep08, mRNA.
<a href="#">Dapk1</a>	<a href="#">Dapk1.aSep08</a>	<a href="#">306722</a>	52840	4937	19	1207	death associated protein kinase 1 (Dapk1) alternative variant aSep08, mRNA.

<a href="#">Dapk1</a>	<a href="#">Dapk1.cSep08</a>	<a href="#">306722</a>	105779	519	4	120	death associated protein kinase 1 (Dapk1) alternative variant cSep08, mRNA.
<a href="#">Dapk2</a>	<a href="#">Dapk2.aSep08</a>	<a href="#">300799</a>	101528	849	8	282	death-associated kinase 2 (Dapk2) alternative variant aSep08, mRNA.
<a href="#">Dapk2</a>	<a href="#">Dapk2.bSep08</a>	<a href="#">300799</a>	94347	729	6	243	death-associated kinase 2 (Dapk2) alternative variant bSep08, mRNA.
<a href="#">Dapk2</a>	<a href="#">Dapk2.dSep08</a>	<a href="#">300799</a>	18796	1108	6	162	death-associated kinase 2 (18.9 kD) (Dapk2) alternative variant dSep08, mRNA.
<a href="#">Dapk2</a>	<a href="#">Dapk2.eSep08</a>	<a href="#">300799</a>	4151	784	3	38	death-associated kinase 2 (4.5 kD) (Dapk2) alternative variant eSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.bSep08</a>	<a href="#">64391</a>	6189	1408	4	286	death-associated protein kinase 3 (32.3 kD) (Dapk3) alternative variant bSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.cSep08</a>	<a href="#">64391</a>	6001	1043	3	259	death-associated protein kinase 3 (29.1 kD) (Dapk3) alternative variant cSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.dSep08</a>	<a href="#">64391</a>	6166	795	6	209	death-associated protein kinase 3 (Dapk3) alternative variant dSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.eSep08</a>	<a href="#">64391</a>	6343	753	6	209	death-associated protein kinase 3 (Dapk3) alternative variant eSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.fSep08</a>	<a href="#">64391</a>	5918	791	5	184	death-associated protein kinase 3 (Dapk3) alternative variant fSep08, mRNA.
<a href="#">Dapk3</a>	<a href="#">Dapk3.gSep08</a>	<a href="#">64391</a>	5353	728	3	61	death-associated protein kinase 3 (6.4 kD) (Dapk3) alternative variant gSep08, mRNA.
<a href="#">dapor</a>	<a href="#">dapor.aSep08</a>		1939	357		119	pleckstrin homology-like domain family B member 1 CRA a (dapor) mRNA.
<a href="#">darby</a>	<a href="#">darby.aSep08</a>		9291	617		205	WD repeat domain 44 (darby) mRNA.
<a href="#">darchy</a>	<a href="#">darchy.aSep08</a>		11997	388	1	78	putative protein (darchy) alternative variant aSep08, mRNA.
<a href="#">darchy</a>	<a href="#">darchy.bSep08</a>		8320	245	1	14	putative protein (darchy) alternative variant bSep08, mRNA.
<a href="#">darfer</a>	<a href="#">darfer.aSep08</a>		7443	1783		22	putative protein (darfer) alternative variant aSep08, mRNA.
<a href="#">darfer</a>	<a href="#">darfer.bSep08</a>		4275	485		36	putative protein (darfer) alternative variant bSep08, mRNA.
<a href="#">darflo</a>	<a href="#">darflo.aSep08</a>		26034	506		168	bnip2 motif-containing molecule at the c-terminal region 1 (darflo) mRNA.
<a href="#">darflu</a>	<a href="#">darflu.aSep08</a>		1921	691		230	n-methyltransferase mll4 like (darflu) mRNA.
<a href="#">darkee</a>	<a href="#">darkee.aSep08</a>		16098	483		104	putative protein (darkee) mRNA.
<a href="#">darloy</a>	<a href="#">darloy.aSep08</a>		10980	758		252	specific 7 (darloy) mRNA.
<a href="#">darmer</a>	<a href="#">darmer.aSep08</a>		1764	259		85	rap guanine nucleotide exchange factor -like 1 (darmer) mRNA.
<a href="#">darnoy</a>	<a href="#">darnoy.aSep08</a>		1999	838		120	putative nuclear protein (12.9 kD) (darnoy) mRNA.
<a href="#">darpor</a>	<a href="#">darpor.aSep08</a>		11855	625		208	CRA a (darpor) mRNA.
<a href="#">darsa</a>	<a href="#">darsa.aSep08</a>		1459	431		109	putative protein (darsa) mRNA.
<a href="#">darshee</a>	<a href="#">darshee.aSep08</a>		542	312		38	putative protein (4.3 kD) (darshee) mRNA.
<a href="#">darto</a>	<a href="#">darto.aSep08</a>		3452	855		285	necrosis factor alpha-induced protein 2 like (darto) alternative variant aSep08, mRNA.
<a href="#">darto</a>	<a href="#">darto.bSep08</a>		2249	513		170	necrosis factor alpha-induced protein 2 like (darto) alternative variant bSep08, mRNA.

<a href="#">darvar</a>	<a href="#">darvar.aSep08</a>		477	361		41	putative protein (darvar) mRNA.
<a href="#">darwey</a>	<a href="#">darwey.aSep08</a>		1309	546	3	86	putative protein (9.1 kD) (darwey) alternative variant aSep08, mRNA.
<a href="#">dasa</a>	<a href="#">dasa.aSep08</a>		2549	650		216	regulator of G-protein 22 (dasa) mRNA.
<a href="#">dashee</a>	<a href="#">dashee.aSep08</a>		47712	666		50	putative protein (dashee) mRNA.
<a href="#">dato</a>	<a href="#">dato.aSep08</a>		8298	431		21	putative protein (2.3 kD) (dato) mRNA.
<a href="#">davar</a>	<a href="#">davar.aSep08</a>		2811	387		41	putative protein (4.5 kD) (davar) mRNA.
<a href="#">dawby</a>	<a href="#">dawby.aSep08</a>		27177	365		69	putative protein (dawby) mRNA.
<a href="#">dawchy</a>	<a href="#">dawchy.bSep08</a>		7953	752	4	122	putative protein human specific (dawchy) alternative variant bSep08, mRNA.
<a href="#">dawchy</a>	<a href="#">dawchy.cSep08</a>		3314	709	2	94	putative mitochondrial protein human specific (10.2 kD) (dawchy) alternative variant cSep08, mRNA.
<a href="#">dawey</a>	<a href="#">dawey.aSep08</a>		19055	487		35	putative protein (4.3 kD) (dawey) mRNA.
<a href="#">dawfer</a>	<a href="#">dawfer.aSep08</a>		10795	589		88	putative protein (dawfer) mRNA.
<a href="#">dawflo</a>	<a href="#">dawflo.aSep08</a>		980	678		121	putative protein (13.4 kD) (dawflo) mRNA.
<a href="#">dawflu</a>	<a href="#">dawflu.aSep08</a>		4630	1060		275	uncharacterized protein (dawflu) mRNA.
<a href="#">dawkee</a>	<a href="#">dawkee.aSep08</a>		3011	1083		156	putative protein of mammalian origin (dawkee) mRNA.
<a href="#">dawloy</a>	<a href="#">dawloy.aSep08</a>		599	253		59	putative protein (dawloy) mRNA.
<a href="#">dawmer</a>	<a href="#">dawmer.aSep08</a>		29961	1793			
<a href="#">dawnoy</a>	<a href="#">dawnoy.aSep08</a>		1681	1132		113	protein tyrosine phosphatase non-receptor type 18 CRA e (dawnoy) mRNA.
<a href="#">dawpor</a>	<a href="#">dawpor.aSep08</a>		1983	762		239	CRA c (dawpor) mRNA.
<a href="#">dawsa</a>	<a href="#">dawsa.aSep08</a>		7991	1768		60	putative protein (6.9 kD) (dawsa) mRNA.
<a href="#">dawshee</a>	<a href="#">dawshee.aSep08</a>		108556	355		25	putative protein (2.8 kD) (dawshee) mRNA.
<a href="#">dawto</a>	<a href="#">dawto.aSep08</a>		56148	757	2	60	putative protein (dawto) mRNA.
<a href="#">dawvar</a>	<a href="#">dawvar.aSep08</a>		2301	386		128	apolipoprotein E receptor 2 (dawvar) mRNA.
<a href="#">dawwey</a>	<a href="#">dawwey.aSep08</a>		2268	1248		64	putative protein (dawwey) mRNA.
<a href="#">Daxx</a>	<a href="#">Daxx.cSep08</a>	<a href="#">140926</a>	1853	501	3	104	fas death domain-associated protein (Daxx) alternative variant cSep08, mRNA.
<a href="#">Daxx</a>	<a href="#">Daxx.dSep08</a>	<a href="#">140926</a>	1852	600	2	87	fas death domain-associated protein (9.2 kD) (Daxx) alternative variant dSep08, mRNA.
<a href="#">Dazap1</a>	<a href="#">Dazap1.bSep08</a>	<a href="#">362836</a>	8030	2228	6	234	DAZ associated protein 1 (24.0 kD) (Dazap1) alternative variant bSep08, mRNA.
<a href="#">Dazap1</a>	<a href="#">Dazap1.cSep08</a>	<a href="#">362836</a>	16101	695	7	156	DAZ associated protein 1 (17.3 kD) (Dazap1) alternative variant cSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.aSep08</a>	<a href="#">300235</a>	4644	1171	5	234	DAZ associated protein 2 (24.6 kD) (Dazap2) alternative variant aSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.bSep08</a>	<a href="#">300235</a>	4440	623	5	207	DAZ associated protein 2 (Dazap2) alternative variant bSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.cSep08</a>	<a href="#">300235</a>	4442	862	5	168	DAZ associated protein 2 (17.3 kD) (Dazap2) alternative variant cSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.dSep08</a>	<a href="#">300235</a>	4263	914	4	168	DAZ associated protein 2 (17.3 kD) (Dazap2) alternative variant dSep08, mRNA.

<a href="#">Dazap2</a>	<a href="#">Dazap2.eSep08</a>	<a href="#">300235</a>	2912	1079	3	167	DAZ associated protein 2 (17.4 kD) (Dazap2) alternative variant eSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.gSep08</a>	<a href="#">300235</a>	5513	1673	4	139	DAZ associated protein 2 (Dazap2) alternative variant gSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.hSep08</a>	<a href="#">300235</a>	4387	736	4	108	DAZ associated protein 2 (10.8 kD) (Dazap2) alternative variant hSep08, mRNA.
<a href="#">Dazap2</a>	<a href="#">Dazap2.iSep08</a>	<a href="#">300235</a>	4337	742	3	86	DAZ associated protein 2 (9.2 kD) (Dazap2) alternative variant iSep08, mRNA.
<a href="#">Dazl</a>	<a href="#">Dazl.bSep08</a>	<a href="#">680486</a>	7551	700	1	126	deleted in azoospermia-like (Dazl) alternative variant bSep08, mRNA.
<a href="#">Dbf4</a>	<a href="#">Dbf4.aSep08</a>	<a href="#">312046</a>	15861	1667		543	DBF4 homolog (S. cerevisiae) (Dbf4) mRNA.
<a href="#">Dbh</a>	<a href="#">Dbh.bSep08</a>	<a href="#">25699</a>	5663	970	1	200	dopamine beta hydroxylase (Dbh) alternative variant bSep08, mRNA.
<a href="#">Dbi</a>	<a href="#">Dbi.aSep08</a>	<a href="#">25045</a>	5014	761	2	115	diazepam binding inhibitor (Dbi) alternative variant aSep08, mRNA.
<a href="#">Dbi</a>	<a href="#">Dbi.cSep08</a>	<a href="#">25045</a>	7626	454	3	63	diazepam binding inhibitor (7.3 kD) (Dbi) alternative variant cSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.aSep08</a>	<a href="#">81653</a>	14343	2907	14	702	drebrin E2 (Dbn1) alternative variant aSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.bSep08</a>	<a href="#">81653</a>	8355	2197	10	535	drebrin E2 (Dbn1) alternative variant bSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.cSep08</a>	<a href="#">81653</a>	6287	1382	5	218	drebrin E2 (Dbn1) alternative variant cSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.dSep08</a>	<a href="#">81653</a>	5876	598	5	199	drebrin 1 CRA b (Dbn1) alternative variant dSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.eSep08</a>	<a href="#">81653</a>	11595	585	7	178	drebrin E2 (Dbn1) alternative variant eSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.fSep08</a>	<a href="#">81653</a>	4867	739	6	110	drebrin E2 (Dbn1) alternative variant fSep08, mRNA.
<a href="#">Dbn1</a>	<a href="#">Dbn1.gSep08</a>	<a href="#">81653</a>	4608	414	2	63	drebrin 1 CRA a (Dbn1) alternative variant gSep08, mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.bSep08</a>	<a href="#">83527</a>	14344	1597	13	432	drebrin-like (48.3 kD) (Dbnl) alternative variant bSep08, mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.cSep08</a>	<a href="#">83527</a>	13446	1175	12	390	drebrin-like (Dbnl) alternative variant cSep08, mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.dSep08</a>	<a href="#">83527</a>	10321	976	5	178	drebrin-like (19.1 kD) (Dbnl) alternative variant dSep08, complete mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.eSep08</a>	<a href="#">83527</a>	5027	748	4	122	drebrin-like (Dbnl) alternative variant eSep08, mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.fSep08</a>	<a href="#">83527</a>	4753	406	2	72	drebrin-like (8.0 kD) (Dbnl) alternative variant fSep08, mRNA.
<a href="#">Dbnl</a>	<a href="#">Dbnl.gSep08</a>	<a href="#">83527</a>	543	343	2	43	putative protein (Dbnl) alternative variant gSep08, mRNA.
<a href="#">Dbp</a>	<a href="#">Dbp.bSep08</a>	<a href="#">24309</a>	1511	669	2	221	D site albumin promoter binding protein (Dbp) alternative variant bSep08, mRNA.
<a href="#">Dbp</a>	<a href="#">Dbp.cSep08</a>	<a href="#">24309</a>	2210	722	3	154	D site albumin promoter binding protein (Dbp) alternative variant cSep08, mRNA.
<a href="#">Dbt</a>	<a href="#">Dbt.bSep08</a>	<a href="#">29611</a>	16241	845	6	203	dihydrolipoamide branched chain transacylase E2 (22.9 kD) (Dbt) alternative variant bSep08, complete mRNA.
<a href="#">Dcakd</a>	<a href="#">Dcakd.bSep08</a>	<a href="#">360639</a>	25233	584	5	147	dephospho-CoA kinase (Dcakd) alternative variant bSep08, mRNA.
<a href="#">Dcakd</a>	<a href="#">Dcakd.cSep08</a>	<a href="#">360639</a>	26447	941	4	119	dephospho-CoA kinase (13.2 kD) (Dcakd) alternative variant cSep08, mRNA.
<a href="#">Dcakd</a>	<a href="#">Dcakd.dSep08</a>	<a href="#">360639</a>	16421	354	2	63	putative protein of ancient origin (Dcakd) alternative variant dSep08, mRNA.

<a href="#">Dcbl1</a>	<a href="#">Dcbl1.aSep08</a>	<a href="#">309773</a>	15505	1690	6	318	putative protein of eukaryotic origin (Dcbl1) alternative variant aSep08, mRNA.
<a href="#">Dcdc5</a>	<a href="#">Dcdc5.bSep08</a>	<a href="#">295980</a>	29913	872	5	53	putative protein (Dcdc5) alternative variant bSep08, mRNA.
<a href="#">Dcdc5</a>	<a href="#">Dcdc5.cSep08</a>	<a href="#">295980</a>	3686	480	2	36	putative protein (4.1 kD) (Dcdc5) alternative variant cSep08, mRNA.
<a href="#">Dchs1</a>	<a href="#">Dchs1.bSep08</a>	<a href="#">308912</a>	426	334	1	111	dachsous 1 (Drosophila) (Dchs1) alternative variant bSep08, mRNA.
<a href="#">Dci</a>	<a href="#">Dci.bSep08</a>	<a href="#">29740</a>	12617	613	4	203	dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase) (Dci) alternative variant bSep08, mRNA.
<a href="#">Dclk1</a>	<a href="#">Dclk1.aSep08</a>	<a href="#">83825</a>	48636	932	7	310	doublecortin-like kinase 1 (Dclk1) alternative variant aSep08, mRNA.
<a href="#">Dclk1</a>	<a href="#">Dclk1.bSep08</a>	<a href="#">83825</a>	29357	406		112	doublecortin-like kinase 1 (Dclk1) alternative variant bSep08, mRNA.
<a href="#">Dclk2</a>	<a href="#">Dclk2.bSep08</a>	<a href="#">310698</a>	23676	522	7	174	doublecortin-like kinase 2 (Dclk2) alternative variant bSep08, mRNA.
<a href="#">Dclk2</a>	<a href="#">Dclk2.dSep08</a>	<a href="#">310698</a>	6317	692	4	71	doublecortin-like kinase 2 (Dclk2) alternative variant dSep08, mRNA.
<a href="#">Dclk2</a>	<a href="#">Dclk2.eSep08</a>	<a href="#">310698</a>	3025	1125	2	83	doublecortin-like kinase 2 (8.7 kD) (Dclk2) alternative variant eSep08, mRNA.
<a href="#">Dclk3</a>	<a href="#">Dclk3.aSep08</a>	<a href="#">316023</a>	30137	637		155	doublecortin-like kinase 3 (Dclk3) mRNA.
<a href="#">Dclre1a</a>	<a href="#">Dclre1a.cSep08</a>	<a href="#">292127</a>	816	343	2	15	DNA cross-link repair 1A, PSO2 homolog (S. cerevisiae) (Dclre1a) alternative variant cSep08, mRNA.
<a href="#">Dclre1b</a>	<a href="#">Dclre1b.bSep08</a>	<a href="#">310745</a>	5284	530	2	166	DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae) (Dclre1b) alternative variant bSep08, mRNA.
<a href="#">Dclre1b</a>	<a href="#">Dclre1b.cSep08</a>	<a href="#">310745</a>	1742	576	3	128	DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae) (Dclre1b) alternative variant cSep08, mRNA.
<a href="#">Dclre1b</a>	<a href="#">Dclre1b.dSep08</a>	<a href="#">310745</a>	4504	920	2	73	DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae) (Dclre1b) alternative variant dSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.bSep08</a>	<a href="#">259171</a>	9583	2558	4	347	DNA cross-link repair 1C PSO2 homolog (37.8 kD) (Dclre1c) alternative variant bSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.cSep08</a>	<a href="#">259171</a>	7185	982	5	299	artemis protein (Dclre1c) alternative variant cSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.dSep08</a>	<a href="#">259171</a>	13564	914	8	174	artemis protein (Dclre1c) alternative variant dSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.eSep08</a>	<a href="#">259171</a>	5175	793	3	111	artemis protein (Dclre1c) alternative variant eSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.fSep08</a>	<a href="#">259171</a>	971	528	2	58	artemis protein (6.6 kD) (Dclre1c) alternative variant fSep08, mRNA.
<a href="#">Dclre1c</a>	<a href="#">Dclre1c.gSep08</a>	<a href="#">259171</a>	2292	415	2	42	artemis protein (Dclre1c) alternative variant gSep08, mRNA.
<a href="#">Dcn</a>	<a href="#">Dcn.bSep08</a>	<a href="#">29139</a>	37290	1787	1	354	decorin (39.8 kD) (Dcn) alternative variant bSep08, mRNA.
<a href="#">DCP1.0</a>	<a href="#">DCP1.0.aSep08</a>		32242	744	7	248	decapping enzyme Dcp1b (DCP1.0) alternative variant aSep08, mRNA.
<a href="#">DCP1.0</a>	<a href="#">DCP1.0.bSep08</a>		27622	734	7	197	decapping enzyme Dcp1b (DCP1.0) alternative variant bSep08, mRNA.

<a href="#">Dcp1a</a>	<a href="#">Dcp1a.aSep08</a>	<a href="#">361109</a>	21298	394	4	118	DCP1 decapping enzyme homolog A ( <i>S. cerevisiae</i> ) (Dcp1a) mRNA.
<a href="#">Dcp1b</a>	<a href="#">Dcp1b.aSep08</a>	<a href="#">500305</a>	4766	486	2	118	DCP1 decapping enzyme homolog b ( <i>S. cerevisiae</i> ) (Dcp1b) alternative variant aSep08, mRNA.
<a href="#">Dcp1b</a>	<a href="#">Dcp1b.bSep08</a>	<a href="#">500305</a>	2628	560	1	72	DCP1 decapping enzyme homolog b ( <i>S. cerevisiae</i> ) (7.9 kD) (Dcp1b) alternative variant bSep08, mRNA.
<a href="#">Dcps</a>	<a href="#">Dcps.bSep08</a>	<a href="#">266605</a>	4720	354	3	68	decapping enzyme, scavenger (Dcps) alternative variant bSep08, mRNA.
<a href="#">Dcst1andAdam15</a>	<a href="#">Dcst1andAdam15.bSep08</a>	<a href="#">57025</a>	2695	744	6	158	a disintegrin and metallopeptidase domain 15 (metargidin) (Dcst1andAdam15) alternative variant bSep08, mRNA.
<a href="#">Dcst1andAdam15</a>	<a href="#">Dcst1andAdam15.bSep08</a>	<a href="#">295246</a>	2695	744	6	158	a disintegrin and metallopeptidase domain 15 (metargidin) (Dcst1andAdam15) alternative variant bSep08, mRNA.
<a href="#">Dcst1andAdam15</a>	<a href="#">Dcst1andAdam15.cSep08</a>	<a href="#">57025</a>	2326	467	4	65	a disintegrin and metallopeptidase domain 15 (metargidin) (Dcst1andAdam15) alternative variant cSep08, mRNA.
<a href="#">Dcst1andAdam15</a>	<a href="#">Dcst1andAdam15.cSep08</a>	<a href="#">295246</a>	2326	467	4	65	a disintegrin and metallopeptidase domain 15 (metargidin) (Dcst1andAdam15) alternative variant cSep08, mRNA.
<a href="#">Dcst2</a>	<a href="#">Dcst2.aSep08</a>	<a href="#">295247</a>	7304	709		169	putative protein of bilateral origin (Dcst2) mRNA.
<a href="#">Dct</a>	<a href="#">Dct.aSep08</a>	<a href="#">290484</a>	33529	1817	2	462	dopachrome tautomerase (Dct) alternative variant aSep08, mRNA.
<a href="#">Dct</a>	<a href="#">Dct.bSep08</a>	<a href="#">290484</a>	12153	608	2	136	dopachrome tautomerase (Dct) alternative variant bSep08, mRNA.
<a href="#">Dctd</a>	<a href="#">Dctd.bSep08</a>	<a href="#">290741</a>	19715	779	5	160	dCMP deaminase (Dctd) alternative variant bSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.bSep08</a>	<a href="#">29167</a>	4570	1254	9	400	dynactin 1 (Dctn1) alternative variant bSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.cSep08</a>	<a href="#">29167</a>	1599	964	5	320	dynactin 1 (Dctn1) alternative variant cSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.dSep08</a>	<a href="#">29167</a>	3233	425	4	141	dynactin 1 (Dctn1) alternative variant dSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.eSep08</a>	<a href="#">29167</a>	22407	463	8	141	dynactin 1 (Dctn1) alternative variant eSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.fSep08</a>	<a href="#">29167</a>	6566	419	4	139	dynactin 1 (Dctn1) alternative variant fSep08, mRNA.
<a href="#">Dctn1</a>	<a href="#">Dctn1.gSep08</a>	<a href="#">29167</a>	2865	353	3	117	dynactin 1 (Dctn1) alternative variant gSep08, mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.aSep08</a>	<a href="#">299850</a>	15220	1689	15	415	dynactin 2 (Dctn2) alternative variant aSep08, mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.cSep08</a>	<a href="#">299850</a>	15424	1836	12	300	dynactin 2 (32.4 kD) (Dctn2) alternative variant cSep08, complete mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.dSep08</a>	<a href="#">299850</a>	8515	710	5	164	dynactin 2 (Dctn2) alternative variant dSep08, mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.eSep08</a>	<a href="#">299850</a>	10257	789	5	154	dynactin 2 (17.4 kD) (Dctn2) alternative variant eSep08, complete mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.fSep08</a>	<a href="#">299850</a>	865	439	4	107	dynactin 2 (Dctn2) alternative variant fSep08, mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.gSep08</a>	<a href="#">299850</a>	15009	457	4	89	dynactin 2 (Dctn2) alternative variant gSep08, mRNA.
<a href="#">Dctn2</a>	<a href="#">Dctn2.hSep08</a>	<a href="#">299850</a>	2782	1943	3	79	dynactin 2 (Dctn2) alternative variant hSep08, mRNA.
<a href="#">Dctn3</a>	<a href="#">Dctn3.bSep08</a>	<a href="#">362504</a>	744	594	2	71	dynactin 3 (8.1 kD) (Dctn3) alternative variant bSep08, mRNA.
<a href="#">Dctn4</a>	<a href="#">Dctn4.bSep08</a>	<a href="#">84428</a>	27007	3801	11	460	dynactin 4 (52.3 kD) (Dctn4) alternative variant bSep08, complete mRNA.
<a href="#">Dctn4</a>	<a href="#">Dctn4.cSep08</a>	<a href="#">84428</a>	7347	558	1	138	dynactin 4 (Dctn4) alternative variant cSep08, mRNA.
<a href="#">Dctn5</a>	<a href="#">Dctn5.bSep08</a>	<a href="#">308961</a>	15564	2058	1	163	dynactin 5 (Dctn5) alternative variant bSep08, mRNA.
<a href="#">Dctn6</a>	<a href="#">Dctn6.bSep08</a>	<a href="#">290798</a>	7121	531	5	167	dynactin 6 (Dctn6) alternative variant bSep08, mRNA.



<a href="#">Dctn6</a>	<a href="#">Dctn6.dSep08</a>	<a href="#">290798</a>	7290	952	3	32	dynactin 6 (3.4 kD) (Dctn6) alternative variant dSep08, mRNA.
<a href="#">Dcun1d1</a>	<a href="#">Dcun1d1.aSep08</a>	<a href="#">310324</a>	44132	2404	7	275	putative protein of eukaryotic origin (Dcun1d1) alternative variant aSep08, mRNA.
<a href="#">Dcun1d1</a>	<a href="#">Dcun1d1.cSep08</a>	<a href="#">310324</a>	28756	2562	5	100	putative protein of eukaryotic origin (Dcun1d1) alternative variant cSep08, mRNA.
<a href="#">Dcun1d4</a>	<a href="#">Dcun1d4.bSep08</a>	<a href="#">360928</a>	8959	1158	2	57	putative protein (6.2 kD) (Dcun1d4) alternative variant bSep08, mRNA.
<a href="#">Dcx</a>	<a href="#">Dcx.aSep08</a>	<a href="#">84394</a>	9684	521		173	doublecortin (Dcx) mRNA.
<a href="#">DCX.1</a>	<a href="#">DCX.1.aSep08</a>		4837	477		158	doublecortin-like kinase 1 (DCX.1) mRNA.
<a href="#">Dd5</a>	<a href="#">Dd5.aSep08</a>	<a href="#">117060</a>	24949	3354	21	1079	progesterone induced protein (Dd5) alternative variant aSep08, mRNA.
<a href="#">Dd5</a>	<a href="#">Dd5.bSep08</a>	<a href="#">117060</a>	6221	1233	4	270	progesterone induced protein (Dd5) alternative variant bSep08, mRNA.
<a href="#">Dd5</a>	<a href="#">Dd5.cSep08</a>	<a href="#">117060</a>	4077	380	4	126	progesterone induced protein (Dd5) alternative variant cSep08, mRNA.
<a href="#">Dd5</a>	<a href="#">Dd5.dSep08</a>	<a href="#">117060</a>	1753	324	1	81	progesterone induced protein (Dd5) alternative variant dSep08, mRNA.
<a href="#">Dd25</a>	<a href="#">Dd25.bSep08</a>	<a href="#">360863</a>	17039	3540	7	670	hypothetical protein Dd25 (Dd25) alternative variant bSep08, mRNA.
<a href="#">Dd25</a>	<a href="#">Dd25.cSep08</a>	<a href="#">360863</a>	8271	649	6	111	hypothetical protein Dd25 (Dd25) alternative variant cSep08, mRNA.
<a href="#">Dd25</a>	<a href="#">Dd25.dSep08</a>	<a href="#">360863</a>	14316	485	3	80	hypothetical protein Dd25 (Dd25) alternative variant dSep08, mRNA.
<a href="#">Dda1</a>	<a href="#">Dda1.aSep08</a>	<a href="#">688813</a>	6626	908	5	132	DET1 and DDB1 associated 1 (Dda1) alternative variant aSep08, mRNA.
<a href="#">Ddah1</a>	<a href="#">Ddah1.bSep08</a>	<a href="#">64157</a>	100257	319	2	106	dimethylarginine dimethylaminohydrolase 1 (Ddah1) alternative variant bSep08, mRNA.
<a href="#">Ddah2</a>	<a href="#">Ddah2.bSep08</a>	<a href="#">294239</a>	721	585	2	169	dimethylarginine dimethylaminohydrolase 2 (Ddah2) alternative variant bSep08, mRNA.
<a href="#">Ddah2</a>	<a href="#">Ddah2.cSep08</a>	<a href="#">294239</a>	915	514	3	108	dimethylarginine dimethylaminohydrolase 2 (Ddah2) alternative variant cSep08, mRNA.
<a href="#">Ddah2</a>	<a href="#">Ddah2.dSep08</a>	<a href="#">294239</a>	1033	424	3	86	dimethylarginine dimethylaminohydrolase 2 (Ddah2) alternative variant dSep08, mRNA.
<a href="#">Ddb1</a>	<a href="#">Ddb1.aSep08</a>	<a href="#">64470</a>	25800	4249		1140	damage-specific DNA binding protein 1 (126.9 kD) (Ddb1) mRNA.
<a href="#">Ddc</a>	<a href="#">Ddc.bSep08</a>	<a href="#">24311</a>	40440	1253	8	296	dopa decarboxylase CRA b (Ddc) alternative variant bSep08, mRNA.
<a href="#">Ddc</a>	<a href="#">Ddc.cSep08</a>	<a href="#">24311</a>	49215	701	6	216	dopa decarboxylase CRA b (Ddc) alternative variant cSep08, mRNA.
<a href="#">Ddc</a>	<a href="#">Ddc.dSep08</a>	<a href="#">24311</a>	18211	422	4	123	dopa decarboxylase CRA b (Ddc) alternative variant dSep08, mRNA.
<a href="#">Ddc</a>	<a href="#">Ddc.eSep08</a>	<a href="#">24311</a>	7217	569	3	112	putative protein of ancient origin (Ddc) alternative variant eSep08, mRNA.
<a href="#">Ddc</a>	<a href="#">Ddc.fSep08</a>	<a href="#">24311</a>	4343	444	2	97	dopa decarboxylase CRA b (11.3 kD) (Ddc) alternative variant fSep08, mRNA.

<a href="#">Ddc8</a>	<a href="#">Ddc8.aSep08</a>	<a href="#">498028</a>	7937	1934	2	591	differential display clone 8 (68.2 kD) (Ddc8) alternative variant aSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.cSep08</a>	<a href="#">498028</a>	7843	711	2	195	differential display clone 8 (Ddc8) alternative variant cSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.dSep08</a>	<a href="#">498028</a>	8313	709	3	72	differential display clone 8 (Ddc8) alternative variant dSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.eSep08</a>	<a href="#">498028</a>	8082	707	3	58	differential display clone 8 (6.3 kD) (Ddc8) alternative variant eSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.fSep08</a>	<a href="#">498028</a>	7649	687	2	58	differential display clone 8 (6.3 kD) (Ddc8) alternative variant fSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.gSep08</a>	<a href="#">498028</a>	8393	583	3	97	differential display clone 8 (Ddc8) alternative variant gSep08, mRNA.
<a href="#">Ddc8</a>	<a href="#">Ddc8.hSep08</a>	<a href="#">498028</a>	8083	443	3	33	differential display clone 8 (3.7 kD) (Ddc8) alternative variant hSep08, mRNA.
<a href="#">Ddef1</a>	<a href="#">Ddef1.bSep08</a>	<a href="#">314961</a>	30618	536	7	178	development and differentiation enhancing (Ddef1) alternative variant bSep08, mRNA.
<a href="#">Ddef1</a>	<a href="#">Ddef1.cSep08</a>	<a href="#">314961</a>	18145	276	4	73	development and differentiation enhancing (Ddef1) alternative variant cSep08, mRNA.
<a href="#">Ddef1</a>	<a href="#">Ddef1.dSep08</a>	<a href="#">314961</a>	855	409	2	61	development and differentiation enhancing (Ddef1) alternative variant dSep08, mRNA.
<a href="#">Ddef1</a>	<a href="#">Ddef1.eSep08</a>	<a href="#">314961</a>	1532	735	3	94	development and differentiation enhancing (11.1 kD) (Ddef1) alternative variant eSep08, mRNA.
<a href="#">Ddef2</a>	<a href="#">Ddef2.bSep08</a>	<a href="#">362719</a>	3150	577	3	151	arf GTPase activating protein (Ddef2) alternative variant bSep08, mRNA.
<a href="#">Ddef2</a>	<a href="#">Ddef2.dSep08</a>	<a href="#">362719</a>	4201	2559	2	96	putative protein (11.2 kD) (Ddef2) alternative variant dSep08, mRNA.
<a href="#">dDENN.0</a>	<a href="#">dDENN.0.aSep08</a>		3801	1677	13	558	SET binding factor 1 like (dDENN.0) alternative variant aSep08, mRNA.
<a href="#">dDENN.1</a>	<a href="#">dDENN.1.aSep08</a>		5095	1523		507	denn 4B (dDENN.1) mRNA.
<a href="#">Ddhd1</a>	<a href="#">Ddhd1.bSep08</a>	<a href="#">305816</a>	7342	390	1	129	putative protein of eukaryotic origin (Ddhd1) alternative variant bSep08, mRNA.
<a href="#">Ddhd2</a>	<a href="#">Ddhd2.aSep08</a>	<a href="#">680971</a>	11582	1790	4	158	ddhd 2 (Ddhd2) alternative variant aSep08, mRNA.
<a href="#">Ddhd2</a>	<a href="#">Ddhd2.bSep08</a>	<a href="#">680971</a>	10898	1041	4	117	ddhd 2 (Ddhd2) alternative variant bSep08, mRNA.
<a href="#">Ddit4</a>	<a href="#">Ddit4.bSep08</a>	<a href="#">140942</a>	2065	1941	1	117	DNA-damage-inducible transcript 4 (Ddit4) alternative variant bSep08, mRNA.
<a href="#">Ddit4l</a>	<a href="#">Ddit4l.bSep08</a>	<a href="#">140582</a>	743	488	1	51	DNA-damage-inducible transcript 4-like and hypothetical protein LOC680872 (Ddit4l) alternative variant bSep08, mRNA.
<a href="#">Ddit4l</a>	<a href="#">Ddit4l.bSep08</a>	<a href="#">680872</a>	743	488	1	51	DNA-damage-inducible transcript 4-like and hypothetical protein LOC680872 (Ddit4l) alternative variant bSep08, mRNA.
<a href="#">Ddo</a>	<a href="#">Ddo.bSep08</a>	<a href="#">685325</a>	10814	1027	1	194	D-aspartate oxidase (Ddo) alternative variant bSep08, mRNA.
<a href="#">Ddost</a>	<a href="#">Ddost.bSep08</a>	<a href="#">313648</a>	1107	1018	2	127	dolichyl-di-phosphooligosaccharide-protein glycotransferase (Ddost) alternative variant bSep08, mRNA.

<a href="#">Ddost</a>	<a href="#">Ddost.cSep08</a>	<a href="#">313648</a>	1509	543	2	96	dolichyl-di-phosphooligosaccharide-protein glycotransferase (Ddost) alternative variant cSep08, mRNA.
<a href="#">Ddr1</a>	<a href="#">Ddr1.aSep08</a>	<a href="#">25678</a>	18005	2875	18	932	discoidin domain receptor family, member 1 (Ddr1) alternative variant aSep08, mRNA.
<a href="#">Ddr1</a>	<a href="#">Ddr1.cSep08</a>	<a href="#">25678</a>	10625	555	5	184	discoidin domain receptor family, member 1 (Ddr1) alternative variant cSep08, mRNA.
<a href="#">Ddr1</a>	<a href="#">Ddr1.dSep08</a>	<a href="#">25678</a>	6465	873	5	155	discoidin domain receptor family, member 1 (Ddr1) alternative variant dSep08, mRNA.
<a href="#">Ddr1</a>	<a href="#">Ddr1.eSep08</a>	<a href="#">25678</a>	3345	408	3	135	discoidin domain receptor family, member 1 (Ddr1) alternative variant eSep08, mRNA.
<a href="#">Ddr1</a>	<a href="#">Ddr1.gSep08</a>	<a href="#">25678</a>	10709	356	4	118	discoidin domain receptor family, member 1 (Ddr1) alternative variant gSep08, mRNA.
<a href="#">Ddt</a>	<a href="#">Ddt.bSep08</a>	<a href="#">29318</a>	2950	1392	2	73	D-dopachrome tautomerase (8.3 kD) (Ddt) alternative variant bSep08, mRNA.
<a href="#">Ddt</a>	<a href="#">Ddt.dSep08</a>	<a href="#">29318</a>	3929	534	2	56	D-dopachrome tautomerase (Ddt) alternative variant dSep08, mRNA.
<a href="#">DDT.2</a>	<a href="#">DDT.2.aSep08</a>		8015	704		234	bromodomain adjacent zinc finger domain 1A (DDT.2) mRNA.
<a href="#">Ddx1</a>	<a href="#">Ddx1.aSep08</a>	<a href="#">84474</a>	30881	2472	26	740	DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 (82.5 kD) (Ddx1) alternative variant aSep08, complete mRNA.
<a href="#">Ddx1</a>	<a href="#">Ddx1.bSep08</a>	<a href="#">84474</a>	11259	675	10	215	DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 (Ddx1) alternative variant bSep08, mRNA.
<a href="#">Ddx1</a>	<a href="#">Ddx1.cSep08</a>	<a href="#">84474</a>	1582	653	3	85	DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 (9.5 kD) (Ddx1) alternative variant cSep08, mRNA.
<a href="#">Ddx1</a>	<a href="#">Ddx1.dSep08</a>	<a href="#">84474</a>	1468	385	3	76	DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 (Ddx1) alternative variant dSep08, mRNA.
<a href="#">Ddx3x</a>	<a href="#">Ddx3x.aSep08</a>	<a href="#">317335</a>	11487	2265	17	683	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked (Ddx3x) alternative variant aSep08, mRNA.
<a href="#">Ddx3x</a>	<a href="#">Ddx3x.cSep08</a>	<a href="#">317335</a>	6932	775	6	175	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked (Ddx3x) alternative variant cSep08, mRNA.
<a href="#">Ddx3x</a>	<a href="#">Ddx3x.dSep08</a>	<a href="#">317335</a>	6881	721	6	173	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked (Ddx3x) alternative variant dSep08, mRNA.
<a href="#">Ddx4</a>	<a href="#">Ddx4.aSep08</a>	<a href="#">310090</a>	12282	900		247	DEAD (Asp-Glu-Ala-Asp) box polypeptide 4 (Ddx4) mRNA.
<a href="#">Ddx5</a>	<a href="#">Ddx5.bSep08</a>	<a href="#">287765</a>	8158	4794	11	406	dead box polypeptide 5 (46.1 kD) (Ddx5) alternative variant bSep08, mRNA.
<a href="#">Ddx5</a>	<a href="#">Ddx5.cSep08</a>	<a href="#">287765</a>	1876	1776	2	127	dead box polypeptide 5 (13.9 kD) (Ddx5) alternative variant cSep08, mRNA.
<a href="#">Ddx5</a>	<a href="#">Ddx5.dSep08</a>	<a href="#">287765</a>	1655	426	2	79	putative protein of ancient origin (Ddx5) alternative variant dSep08, mRNA.
<a href="#">Ddx11</a>	<a href="#">Ddx11.aSep08</a>	<a href="#">316767</a>	1819	1583		145	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, <i>S. cerevisiae</i> ) (Ddx11) mRNA.
<a href="#">Ddx17</a>	<a href="#">Ddx17.aSep08</a>	<a href="#">315133</a>	20210	4575	13	501	dead box polypeptide 17 CRA b (Ddx17) alternative variant aSep08, mRNA.
<a href="#">Ddx17</a>	<a href="#">Ddx17.cSep08</a>	<a href="#">315133</a>	7949	742	5	246	dead box polypeptide 17 CRA h (Ddx17) alternative variant cSep08, mRNA.

<a href="#">Ddx17</a>	<a href="#">Ddx17.dSep08</a>	<a href="#">315133</a>	2883	1637	2	202	dead box polypeptide 17 (Ddx17) alternative variant dSep08, mRNA.
<a href="#">Ddx17</a>	<a href="#">Ddx17.eSep08</a>	<a href="#">315133</a>	3326	773	5	168	dead box polypeptide 17 (Ddx17) alternative variant eSep08, mRNA.
<a href="#">Ddx17</a>	<a href="#">Ddx17.gSep08</a>	<a href="#">315133</a>	3297	447	2	32	putative protein (Ddx17) alternative variant gSep08, mRNA.
<a href="#">Ddx18</a>	<a href="#">Ddx18.cSep08</a>	<a href="#">308490</a>	1626	400	2	63	DEAD (Asp-Glu-Ala-Asp) box polypeptide 18 (Ddx18) alternative variant cSep08, mRNA.
<a href="#">Ddx20</a>	<a href="#">Ddx20.aSep08</a>	<a href="#">84473</a>	7988	2748	11	821	DEAD (Asp-Glu-Ala-Asp) box polypeptide 20 (Ddx20) alternative variant aSep08, mRNA.
<a href="#">Ddx21</a>	<a href="#">Ddx21.aSep08</a>	<a href="#">317399</a>	9719	3072	8	360	DEAD (Asp-Glu-Ala-Asp) box polypeptide 21 (Ddx21) alternative variant aSep08, mRNA.
<a href="#">Ddx21</a>	<a href="#">Ddx21.bSep08</a>	<a href="#">317399</a>	21858	1809	1	358	DEAD (Asp-Glu-Ala-Asp) box polypeptide 21 (Ddx21) alternative variant bSep08, complete mRNA.
<a href="#">Ddx21</a>	<a href="#">Ddx21.cSep08</a>	<a href="#">317399</a>	1758	748		104	DEAD (Asp-Glu-Ala-Asp) box polypeptide 21 (Ddx21) alternative variant cSep08, mRNA.
<a href="#">Ddx23</a>	<a href="#">Ddx23.aSep08</a>	<a href="#">300208</a>	17231	3109	17	819	DEAD box polypeptide 23 (95.5 kD) (Ddx23) alternative variant aSep08, mRNA.
<a href="#">Ddx23</a>	<a href="#">Ddx23.bSep08</a>	<a href="#">300208</a>	11201	915	8	284	DEAD box polypeptide 23 (Ddx23) alternative variant bSep08, mRNA.
<a href="#">Ddx23</a>	<a href="#">Ddx23.dSep08</a>	<a href="#">300208</a>	2787	742	4	246	DEAD box polypeptide 23 (Ddx23) alternative variant dSep08, mRNA.
<a href="#">Ddx23</a>	<a href="#">Ddx23.eSep08</a>	<a href="#">300208</a>	3412	755	5	135	DEAD box polypeptide 23 (Ddx23) alternative variant eSep08, mRNA.
<a href="#">Ddx23</a>	<a href="#">Ddx23.fSep08</a>	<a href="#">300208</a>	1115	750	2	121	DEAD box polypeptide 23 (Ddx23) alternative variant fSep08, mRNA.
<a href="#">Ddx24</a>	<a href="#">Ddx24.bSep08</a>	<a href="#">373065</a>	6271	766	3	254	dead box polypeptide 24 (Ddx24) alternative variant bSep08, mRNA.
<a href="#">Ddx24</a>	<a href="#">Ddx24.cSep08</a>	<a href="#">373065</a>	3502	1136	3	169	dead box polypeptide 24 CRA a (19.9 kD) (Ddx24) alternative variant cSep08, mRNA.
<a href="#">Ddx24</a>	<a href="#">Ddx24.dSep08</a>	<a href="#">373065</a>	1782	706	2	115	putative mitochondrial protein (12.5 kD) (Ddx24) alternative variant dSep08, mRNA.
<a href="#">Ddx24</a>	<a href="#">Ddx24.eSep08</a>	<a href="#">373065</a>	1529	612	2	108	putative mitochondrial protein (11.6 kD) (Ddx24) alternative variant eSep08, mRNA.
<a href="#">Ddx24</a>	<a href="#">Ddx24.fSep08</a>	<a href="#">373065</a>	1667	384	2	69	putative protein (7.4 kD) (Ddx24) alternative variant fSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.bSep08</a>	<a href="#">58856</a>	9048	1634	4	313	hydrolethalus syndrome 1 like (35.7 kD) (Ddx25) alternative variant bSep08, complete mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.bSep08</a>	<a href="#">680262</a>	9048	1634	4	313	hydrolethalus syndrome 1 like (35.7 kD) (Ddx25) alternative variant bSep08, complete mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.cSep08</a>	<a href="#">58856</a>	16770	705	8	146	dead box polypeptide 25 CRA a (Ddx25) alternative variant cSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.cSep08</a>	<a href="#">680262</a>	16770	705	8	146	dead box polypeptide 25 CRA a (Ddx25) alternative variant cSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.dSep08</a>	<a href="#">58856</a>	7826	741	4	129	hydrolethalus syndrome 1 like (Ddx25) alternative variant dSep08, mRNA.

<a href="#">Ddx25</a>	<a href="#">Ddx25.dSep08</a>	<a href="#">680262</a>	7826	741	4	129	hydrolethalus syndrome 1 like (Ddx25) alternative variant dSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.eSep08</a>	<a href="#">58856</a>	1007	755	2	118	DEAD box polypeptide 25 (12.6 kD) (Ddx25) alternative variant eSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.eSep08</a>	<a href="#">680262</a>	1007	755	2	118	DEAD box polypeptide 25 (12.6 kD) (Ddx25) alternative variant eSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.fSep08</a>	<a href="#">58856</a>	2330	392	2	103	DEAD box polypeptide (Ddx25) alternative variant fSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.fSep08</a>	<a href="#">680262</a>	2330	392	2	103	DEAD box polypeptide (Ddx25) alternative variant fSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.hSep08</a>	<a href="#">58856</a>	1741	261	2	47	putative protein (Ddx25) alternative variant hSep08, mRNA.
<a href="#">Ddx25</a>	<a href="#">Ddx25.hSep08</a>	<a href="#">680262</a>	1741	261	2	47	putative protein (Ddx25) alternative variant hSep08, mRNA.
<a href="#">Ddx27</a>	<a href="#">Ddx27.aSep08</a>	<a href="#">362274</a>	19831	2595	21	768	dead box polypeptide 27 (Ddx27) alternative variant aSep08, mRNA.
<a href="#">Ddx27</a>	<a href="#">Ddx27.bSep08</a>	<a href="#">362274</a>	2313	1410	3	146	dead box polypeptide 27 (Ddx27) alternative variant bSep08, mRNA.
<a href="#">Ddx27</a>	<a href="#">Ddx27.cSep08</a>	<a href="#">362274</a>	3370	1929	3	134	dead box polypeptide 27 (15.4 kD) (Ddx27) alternative variant cSep08, mRNA.
<a href="#">Ddx31</a>	<a href="#">Ddx31.bSep08</a>	<a href="#">311835</a>	17729	905	4	139	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31 (Ddx31) alternative variant bSep08, mRNA.
<a href="#">Ddx39</a>	<a href="#">Ddx39.bSep08</a>	<a href="#">89827</a>	8042	1480	9	345	DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 (Ddx39) alternative variant bSep08, mRNA.
<a href="#">Ddx39</a>	<a href="#">Ddx39.cSep08</a>	<a href="#">89827</a>	7564	1532	8	208	DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 (23.2 kD) (Ddx39) alternative variant cSep08, mRNA.
<a href="#">Ddx39</a>	<a href="#">Ddx39.eSep08</a>	<a href="#">89827</a>	1626	742	3	114	DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 (Ddx39) alternative variant eSep08, mRNA.
<a href="#">Ddx41</a>	<a href="#">Ddx41.bSep08</a>	<a href="#">314336</a>	976	806	3	115	DEAD (Asp-Glu-Ala-Asp) box polypeptide 41 (12.9 kD) (Ddx41) alternative variant bSep08, mRNA.
<a href="#">Ddx41</a>	<a href="#">Ddx41.cSep08</a>	<a href="#">314336</a>	1062	716	5	107	DEAD (Asp-Glu-Ala-Asp) box polypeptide 41 (Ddx41) alternative variant cSep08, mRNA.
<a href="#">Ddx41</a>	<a href="#">Ddx41.eSep08</a>	<a href="#">314336</a>	470	368	2	73	DEAD (Asp-Glu-Ala-Asp) box polypeptide 41 (8.3 kD) (Ddx41) alternative variant eSep08, mRNA.
<a href="#">Ddx42</a>	<a href="#">Ddx42.bSep08</a>	<a href="#">303607</a>	3344	1489	3	283	DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (Ddx42) alternative variant bSep08, mRNA.
<a href="#">Ddx42</a>	<a href="#">Ddx42.cSep08</a>	<a href="#">303607</a>	13039	615	5	204	DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (Ddx42) alternative variant cSep08, mRNA.
<a href="#">Ddx42</a>	<a href="#">Ddx42.eSep08</a>	<a href="#">303607</a>	3557	541	3	75	DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (Ddx42) alternative variant eSep08, mRNA.
<a href="#">Ddx47</a>	<a href="#">Ddx47.bSep08</a>	<a href="#">297685</a>	2050	726	4	125	DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 (Ddx47) alternative variant bSep08, mRNA.
<a href="#">Ddx47</a>	<a href="#">Ddx47.cSep08</a>	<a href="#">297685</a>	4586	1778	3	61	DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 (Ddx47) alternative variant cSep08, mRNA.
<a href="#">Ddx49</a>	<a href="#">Ddx49.bSep08</a>	<a href="#">290660</a>	3103	672	8	161	DEAD (Asp-Glu-Ala-Asp) box polypeptide 49 (Ddx49) alternative variant bSep08, mRNA.
<a href="#">Ddx50</a>	<a href="#">Ddx50.bSep08</a>	<a href="#">361848</a>	13294	735	6	228	CRA a (Ddx50) alternative variant bSep08, mRNA.

<a href="#">Ddx50</a>	<a href="#">Ddx50.cSep08</a>	<a href="#">361848</a>	8472	1577	5	160	DEAD box polypeptide 50 (Ddx50) alternative variant cSep08, mRNA.
<a href="#">Ddx50</a>	<a href="#">Ddx50.eSep08</a>	<a href="#">361848</a>	903	316	2	46	DEAD box polypeptide 50 like (5.3 kD) (Ddx50) alternative variant eSep08, mRNA.
<a href="#">Ddx51</a>	<a href="#">Ddx51.bSep08</a>	<a href="#">304570</a>	934	495	4	93	DEAD (Asp-Glu-Ala-Asp) box polypeptide 51 (Ddx51) alternative variant bSep08, mRNA.
<a href="#">Ddx51</a>	<a href="#">Ddx51.cSep08</a>	<a href="#">304570</a>	514	282	3	64	DEAD (Asp-Glu-Ala-Asp) box polypeptide 51 (Ddx51) alternative variant cSep08, mRNA.
<a href="#">Ddx54</a>	<a href="#">Ddx54.aSep08</a>	<a href="#">360815</a>	15247	2166	12	613	DEAD box polypeptide 54 (Ddx54) alternative variant aSep08, mRNA.
<a href="#">Ddx54</a>	<a href="#">Ddx54.bSep08</a>	<a href="#">360815</a>	4184	1669	8	402	DEAD box polypeptide 54 (Ddx54) alternative variant bSep08, mRNA.
<a href="#">Ddx54</a>	<a href="#">Ddx54.cSep08</a>	<a href="#">360815</a>	5355	1879	7	369	DEAD box polypeptide 54 (Ddx54) alternative variant cSep08, mRNA.
<a href="#">Ddx54</a>	<a href="#">Ddx54.dSep08</a>	<a href="#">360815</a>	1698	1587	3	348	putative protein (Ddx54) alternative variant dSep08, mRNA.
<a href="#">Ddx56</a>	<a href="#">Ddx56.bSep08</a>	<a href="#">289780</a>	3714	1140	8	374	DEAD (Asp-Glu-Ala-Asp) box polypeptide 56 (Ddx56) alternative variant bSep08, mRNA.
<a href="#">Ddx56</a>	<a href="#">Ddx56.cSep08</a>	<a href="#">289780</a>	332	252	2	79	DEAD (Asp-Glu-Ala-Asp) box polypeptide 56 (Ddx56) alternative variant cSep08, mRNA.
<a href="#">Ddx58</a>	<a href="#">Ddx58.bSep08</a>	<a href="#">297989</a>	18545	1033	1	115	DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 (13.6 kD) (Ddx58) alternative variant bSep08, mRNA.
<a href="#">Ddx59</a>	<a href="#">Ddx59.bSep08</a>	<a href="#">289402</a>	15813	1215	1	404	DEAD (Asp-Glu-Ala-Asp) box polypeptide 59 (Ddx59) alternative variant bSep08, mRNA.
<a href="#">Ddx60</a>	<a href="#">Ddx60.aSep08</a>	<a href="#">306409</a>	7325	571		189	DEAD (Asp-Glu-Ala-Asp) box polypeptide 60 (Ddx60) mRNA.
<a href="#">DEAD.0</a>	<a href="#">DEAD.0.aSep08</a>		13638	1378	12	444	dead box polypeptide 55 CRA a (DEAD.0) alternative variant aSep08, mRNA.
<a href="#">DEAD.0</a>	<a href="#">DEAD.0.bSep08</a>		4756	746	5	248	dead box polypeptide 55 (DEAD.0) alternative variant bSep08, mRNA.
<a href="#">DEAD.0</a>	<a href="#">DEAD.0.cSep08</a>		5506	698	4	163	dead box polypeptide 55 (DEAD.0) alternative variant cSep08, mRNA.
<a href="#">DEAD.0</a>	<a href="#">DEAD.0.dSep08</a>		1932	415	3	79	dead box polypeptide 55 (DEAD.0) alternative variant dSep08, mRNA.
<a href="#">DEAD.1</a>	<a href="#">DEAD.1.aSep08</a>		6841	1080		360	DEAH box polypeptide 29 (DEAD.1) mRNA.
<a href="#">Deadc1</a>	<a href="#">Deadc1.aSep08</a>	<a href="#">361453</a>	14491	545	1	181	CMP/dCMP deaminase, zinc-binding (Deadc1) alternative variant aSep08, mRNA.
<a href="#">Deadc1</a>	<a href="#">Deadc1.bSep08</a>	<a href="#">361453</a>	14404	429	1	124	CMP/dCMP deaminase, zinc-binding (Deadc1) alternative variant bSep08, mRNA.
<a href="#">Deaf1</a>	<a href="#">Deaf1.bSep08</a>	<a href="#">83632</a>	30546	2250	1	207	deformed epidermal autoregulatory factor 1 (Drosophila) (21.8 kD) (Deaf1) alternative variant bSep08, mRNA.
<a href="#">Death.0</a>	<a href="#">Death.0.aSep08</a>		5626	1122	3	373	ankyrin 2 (Death.0) alternative variant aSep08, mRNA.
<a href="#">Death.0</a>	<a href="#">Death.0.bSep08</a>		2767	401	2	133	ankyrin 2 (Death.0) alternative variant bSep08, mRNA.
<a href="#">Decr1</a>	<a href="#">Decr1.bSep08</a>	<a href="#">117543</a>	22312	750	5	100	2,4-dienoyl CoA reductase 1, mitochondrial (10.5 kD) (Decr1) alternative variant bSep08, mRNA.
<a href="#">Decr1</a>	<a href="#">Decr1.cSep08</a>	<a href="#">117543</a>	10000	682	6	82	2,4-dienoyl CoA reductase 1, mitochondrial (Decr1) alternative variant cSep08, mRNA.

<a href="#">Decr2</a>	<a href="#">Decr2.bSep08</a>	<a href="#">64461</a>	7547	881	7	252	2-4-dienoyl-Coenzyme A reductase 2, peroxisomal (Decr2) alternative variant bSep08, mRNA.
<a href="#">Decr2</a>	<a href="#">Decr2.cSep08</a>	<a href="#">64461</a>	5594	677	8	225	2-4-dienoyl-Coenzyme A reductase 2, peroxisomal (Decr2) alternative variant cSep08, mRNA.
<a href="#">Decr2</a>	<a href="#">Decr2.dSep08</a>	<a href="#">64461</a>	3279	1434	3	202	2-4-dienoyl-Coenzyme A reductase 2, peroxisomal (Decr2) alternative variant dSep08, mRNA.
<a href="#">Dedd</a>	<a href="#">Dedd.bSep08</a>	<a href="#">83631</a>	10242	725	4	170	death effector domain-containing (Dedd) alternative variant bSep08, mRNA.
<a href="#">Dedd</a>	<a href="#">Dedd.cSep08</a>	<a href="#">83631</a>	11980	679	5	160	death effector domain-containing (Dedd) alternative variant cSep08, mRNA.
<a href="#">Dedd</a>	<a href="#">Dedd.eSep08</a>	<a href="#">83631</a>	9769	405	3	35	putative protein (Dedd) alternative variant eSep08, mRNA.
<a href="#">Dedd</a>	<a href="#">Dedd.fSep08</a>	<a href="#">83631</a>	1546	360	2	30	putative protein (3.6 kD) (Dedd) alternative variant fSep08, mRNA.
<a href="#">deeby</a>	<a href="#">deeby.bSep08</a>		1310	751	2	38	putative protein (4.6 kD) (deeby) alternative variant bSep08, mRNA.
<a href="#">deechy</a>	<a href="#">deechy.aSep08</a>		8195	653		46	putative protein (deechy) mRNA.
<a href="#">deefer</a>	<a href="#">deefer.aSep08</a>		2021	1065		38	putative protein (4.2 kD) (deefer) mRNA.
<a href="#">deeflo</a>	<a href="#">deeflo.aSep08</a>		19582	1645		317	proprotein convertase subtilisin kexin type 5 (deeflo) mRNA.
<a href="#">deeflu</a>	<a href="#">deeflu.aSep08</a>		3260	362	2	120	CRA a like (deeflu) alternative variant aSep08, mRNA.
<a href="#">deeflu</a>	<a href="#">deeflu.bSep08</a>		5763	1083	6	118	uncharacterized protein like (13.6 kD) (deeflu) alternative variant bSep08, mRNA.
<a href="#">deekee</a>	<a href="#">deekee.aSep08</a>		2639	753		121	putative protein of mammalian origin (deekee) mRNA.
<a href="#">deeloy</a>	<a href="#">deeloy.aSep08</a>		6366	755	1	54	putative protein (deeloy) alternative variant aSep08, mRNA.
<a href="#">deeloy</a>	<a href="#">deeloy.bSep08</a>		1450	239	1	45	putative protein (deeloy) alternative variant bSep08, mRNA.
<a href="#">deemer</a>	<a href="#">deemer.aSep08</a>		2724	395		98	dna topoisomerase (deemer) mRNA.
<a href="#">deenoy</a>	<a href="#">deenoy.aSep08</a>		512	453		52	putative protein (6.2 kD) (deenoy) mRNA.
<a href="#">deepor</a>	<a href="#">deepor.aSep08</a>		6983	810		65	putative protein of mammalian origin (7.1 kD) (deepor) mRNA.
<a href="#">deesa</a>	<a href="#">deesa.aSep08</a>		24646	695		76	putative cytoplasmic protein (8.8 kD) (deesa) mRNA.
<a href="#">deeshee</a>	<a href="#">deeshee.aSep08</a>		25553	3401	7	353	family with sequence similarity 105 member B (40.4 kD) (deeshee) alternative variant aSep08, complete mRNA.
<a href="#">deeshee</a>	<a href="#">deeshee.bSep08</a>		19507	633	5	186	sertolin (deeshee) alternative variant bSep08, mRNA.
<a href="#">deeto</a>	<a href="#">deeto.aSep08</a>		841	522		71	putative cytoplasmic protein (7.8 kD) (deeto) mRNA.
<a href="#">deever</a>	<a href="#">deever.aSep08</a>		2062	394		104	putative protein (deever) mRNA.
<a href="#">deewey</a>	<a href="#">deewey.aSep08</a>		4079	355		56	mob1 Mps One Binder kinase activator-like 1A CRA a (deewey) mRNA.
<a href="#">Def6</a>	<a href="#">Def6.aSep08</a>	<a href="#">309642</a>	16235	984		323	differentially expressed in FDCP 6 (Def6) mRNA.
<a href="#">Def8</a>	<a href="#">Def8.bSep08</a>	<a href="#">307973</a>	14517	945	8	247	differentially expressed in FDCP 8 (Def8) alternative variant bSep08, mRNA.
<a href="#">Def8</a>	<a href="#">Def8.cSep08</a>	<a href="#">307973</a>	2133	289	2	65	differentially expressed in FDCP 8 (Def8) alternative variant cSep08, mRNA.

<a href="#">Def8</a>	<a href="#">Def8.dSep08</a>	<a href="#">307973</a>	950	450	2	63	differentially expressed in FDCP 8 (Def8) alternative variant dSep08, mRNA.
<a href="#">Defa</a>	<a href="#">Defa.aSep08</a>	<a href="#">286995</a>	1149	632	1	135	defensin, alpha 5, Paneth cell-specific (15.3 kD) (Defa) alternative variant aSep08, mRNA.
<a href="#">Defb11</a>	<a href="#">Defb11.bSep08</a>	<a href="#">641630</a>	1053	300	2	64	defensin beta 11 (Defb11) alternative variant bSep08, mRNA.
<a href="#">Defb17andDefb16-ps</a>	<a href="#">Defb17andDefb16-ps.bSep08</a>	<a href="#">641622</a>	8530	189	1	35	putative protein (Defb17andDefb16-ps) alternative variant bSep08, mRNA.
<a href="#">Defb17andDefb16-ps</a>	<a href="#">Defb17andDefb16-ps.bSep08</a>	<a href="#">641658</a>	8530	189	1	35	putative protein (Defb17andDefb16-ps) alternative variant bSep08, mRNA.
<a href="#">Degs1</a>	<a href="#">Degs1.bSep08</a>	<a href="#">58970</a>	2650	695	3	115	degenerative spermatocyte homolog 1 (Drosophila) (Degs1) alternative variant bSep08, mRNA.
<a href="#">Dek</a>	<a href="#">Dek.bSep08</a>	<a href="#">306817</a>	13598	735	6	148	DEK oncogene (DNA binding) (Dek) alternative variant bSep08, mRNA.
<a href="#">Dek</a>	<a href="#">Dek.cSep08</a>	<a href="#">306817</a>	4304	408	3	119	DEK oncogene (DNA binding) (Dek) alternative variant cSep08, mRNA.
<a href="#">DENN.0</a>	<a href="#">DENN.0.aSep08</a>		773	361		120	DENN (DENN.0) mRNA.
<a href="#">DENN.1</a>	<a href="#">DENN.1.aSep08</a>		12759	1074	3	358	rab6 interacting protein 1 (DENN.1) alternative variant aSep08, mRNA.
<a href="#">Dennd1a</a>	<a href="#">Dennd1a.aSep08</a>	<a href="#">311913</a>	158468	3853	9	767	dDENN (Dennd1a) alternative variant aSep08, mRNA.
<a href="#">Dennd1a</a>	<a href="#">Dennd1a.bSep08</a>	<a href="#">311913</a>	14921	740	1	182	putative protein of vertebrate origin (Dennd1a) alternative variant bSep08, mRNA.
<a href="#">Dennd1a</a>	<a href="#">Dennd1a.cSep08</a>	<a href="#">311913</a>	33972	592	6	128	putative protein of bilateral origin (Dennd1a) alternative variant cSep08, mRNA.
<a href="#">Dennd1b</a>	<a href="#">Dennd1b.aSep08</a>	<a href="#">289051</a>	45266	400	5	133	uDENN (Dennd1b) alternative variant aSep08, mRNA.
<a href="#">Dennd1b</a>	<a href="#">Dennd1b.bSep08</a>	<a href="#">289051</a>	36843	294	3	97	putative protein of eukaryotic origin (Dennd1b) alternative variant bSep08, mRNA.
<a href="#">Dennd2a</a>	<a href="#">Dennd2a.aSep08</a>	<a href="#">312257</a>	18913	1226	2	298	denn 2A (Dennd2a) alternative variant aSep08, mRNA.
<a href="#">Dennd2a</a>	<a href="#">Dennd2a.bSep08</a>	<a href="#">312257</a>	2251	875	1	148	denn 2A (Dennd2a) alternative variant bSep08, mRNA.
<a href="#">Dennd2c</a>	<a href="#">Dennd2c.aSep08</a>	<a href="#">295333</a>	13477	1154		336	DENN (Dennd2c) mRNA.
<a href="#">Dennd3</a>	<a href="#">Dennd3.aSep08</a>	<a href="#">315055</a>	7849	2063		242	WD-40 repeat (Dennd3) mRNA.
<a href="#">Dennd4b</a>	<a href="#">Dennd4b.aSep08</a>	<a href="#">361987</a>	3097	1768		339	putative protein of eukaryotic origin (Dennd4b) mRNA.
<a href="#">DEP.0</a>	<a href="#">DEP.0.aSep08</a>		369	298		72	regulator of G-protein 11 (DEP.0) mRNA.
<a href="#">DEP.1</a>	<a href="#">DEP.1.aSep08</a>		4048	2025	10	490	dishevelled homolog (51.8 kD) (DEP.1) alternative variant aSep08, complete mRNA.
<a href="#">DEP.1</a>	<a href="#">DEP.1.bSep08</a>		1376	877	4	194	dishevelled homolog (21.1 kD) (DEP.1) alternative variant bSep08, mRNA.
<a href="#">DEP.1</a>	<a href="#">DEP.1.cSep08</a>		948	860	2	138	dishevelled homolog precursor (15.0 kD) (DEP.1) alternative variant cSep08, mRNA.
<a href="#">DEP.1</a>	<a href="#">DEP.1.dSep08</a>		660	458	3	75	dishevelled homolog (DEP.1) alternative variant dSep08, mRNA.
<a href="#">DEP.2</a>	<a href="#">DEP.2.aSep08</a>		12266	745		248	phosphatidylinositol 3 4 5-trisphosphate-dependent Rac exchanger 1 (DEP.2) mRNA.
<a href="#">Depdc1a</a>	<a href="#">Depdc1a.aSep08</a>	<a href="#">295538</a>	2901	139		45	putative protein of bilateral origin (Depdc1a) mRNA.
<a href="#">Depdc1b</a>	<a href="#">Depdc1b.bSep08</a>	<a href="#">310074</a>	13714	747	1	81	DEP domain-containing protein 1B (Depdc1b) alternative variant bSep08, mRNA.



<a href="#">Depdc2</a>	<a href="#">Depdc2.bSep08</a>	<a href="#">312912</a>	28294	827	8	275	ptdins-dependent rac exchanger 2a (Depdc2) alternative variant bSep08, mRNA.
<a href="#">Depdc2</a>	<a href="#">Depdc2.dSep08</a>	<a href="#">312912</a>	34148	1311	4	84	putative protein of metazoan origin (Depdc2) alternative variant dSep08, mRNA.
<a href="#">Depdc5</a>	<a href="#">Depdc5.bSep08</a>	<a href="#">305464</a>	16902	755	5	154	putative protein of metazoan origin (Depdc5) alternative variant bSep08, mRNA.
<a href="#">Depdc5</a>	<a href="#">Depdc5.cSep08</a>	<a href="#">305464</a>	4483	425	4	68	putative protein (Depdc5) alternative variant cSep08, mRNA.
<a href="#">Depdc7</a>	<a href="#">Depdc7.bSep08</a>	<a href="#">295971</a>	2556	1230	2	109	putative protein of vertebrate origin (Depdc7) alternative variant bSep08, mRNA.
<a href="#">derby</a>	<a href="#">derby.aSep08</a>		3558	685		49	putative protein (5.8 kD) (derby) mRNA.
<a href="#">derchy</a>	<a href="#">derchy.aSep08</a>		3982	292		34	putative protein (derchy) mRNA.
<a href="#">derfer</a>	<a href="#">derfer.aSep08</a>		26258	553		69	putative protein (derfer) mRNA.
<a href="#">derflo</a>	<a href="#">derflo.aSep08</a>		14231	374	2	85	putative mitochondrial protein (9.7 kD) (derflo) alternative variant aSep08, mRNA.
<a href="#">derflo</a>	<a href="#">derflo.bSep08</a>		16980	466	3	24	putative protein (2.6 kD) (derflo) alternative variant bSep08, mRNA.
<a href="#">derflo</a>	<a href="#">derflo.cSep08</a>		17344	398	3	40	putative protein (derflo) alternative variant cSep08, mRNA.
<a href="#">derflu</a>	<a href="#">derflu.aSep08</a>		1161	380		34	putative protein (derflu) mRNA.
<a href="#">derkee</a>	<a href="#">derkee.aSep08</a>		24134	475		42	putative protein (4.8 kD) (derkee) mRNA.
<a href="#">Der1</a>	<a href="#">Der1.bSep08</a>	<a href="#">362912</a>	4245	524	2	65	der1-like domain family, member 1 (7.2 kD) (Der1) alternative variant bSep08, mRNA.
<a href="#">Der1</a>	<a href="#">Der1.cSep08</a>	<a href="#">362912</a>	8786	260	2	34	der1-like domain family, member 1 (4.0 kD) (Der1) alternative variant cSep08, mRNA.
<a href="#">Der2</a>	<a href="#">Der2.aSep08</a>	<a href="#">691956</a>	9142	1146	7	246	der1-like domain family, member 2 (Der2) alternative variant aSep08, mRNA.
<a href="#">Der2</a>	<a href="#">Der2.bSep08</a>	<a href="#">691956</a>	6399	792	5	61	der1-like domain family, member 2 (Der2) alternative variant bSep08, mRNA.
<a href="#">Der2</a>	<a href="#">Der2.cSep08</a>	<a href="#">691956</a>	5218	437	3	60	der1-like domain family, member 2 (Der2) alternative variant cSep08, mRNA.
<a href="#">Der3</a>	<a href="#">Der3.bSep08</a>	<a href="#">690315</a>	3442	1117	6	222	der1-like domain family, member 3 (Der3) alternative variant bSep08, mRNA.
<a href="#">Der3</a>	<a href="#">Der3.cSep08</a>	<a href="#">690315</a>	12336	774	5	154	der1-like domain family, member 3 (17.7 kD) (Der3) alternative variant cSep08, mRNA.
<a href="#">Der3</a>	<a href="#">Der3.dSep08</a>	<a href="#">690315</a>	1787	1083	2	79	der1-like domain family, member 3 (Der3) alternative variant dSep08, mRNA.
<a href="#">derloy</a>	<a href="#">derloy.aSep08</a>		882	456	2	25	putative protein (derloy) alternative variant aSep08, mRNA.
<a href="#">dermer</a>	<a href="#">dermer.aSep08</a>		1793	928		74	putative cytoplasmic protein (8.5 kD) (dermer) mRNA.
<a href="#">dernoy</a>	<a href="#">dernoy.bSep08</a>		25627	447	3	19	putative protein (dernoy) alternative variant bSep08, mRNA.
<a href="#">derpor</a>	<a href="#">derpor.aSep08</a>		5136	599		26	putative protein (2.8 kD) (derpor) mRNA.
<a href="#">dersa</a>	<a href="#">dersa.aSep08</a>		6426	1247		415	putative protein of metazoan origin (dersa) alternative variant aSep08, mRNA.
<a href="#">dershee</a>	<a href="#">dershee.aSep08</a>		20897	527		118	putative protein (dershee) mRNA.
<a href="#">derto</a>	<a href="#">derto.aSep08</a>		4711	545		47	putative protein (derto) mRNA.

<a href="#">dervar</a>	<a href="#">dervar.aSep08</a>		12988	794	4	108	putative protein (dervar) alternative variant aSep08, mRNA.
<a href="#">dervar</a>	<a href="#">dervar.bSep08</a>		3744	219	1	73	putative protein (dervar) alternative variant bSep08, mRNA.
<a href="#">derwey</a>	<a href="#">derwey.aSep08</a>		3650	507		81	putative protein of eukaryotic origin (derwey) mRNA.
<a href="#">Det1</a>	<a href="#">Det1.bSep08</a>	<a href="#">308775</a>	10350	1207	4	392	de-etiolated homolog 1 (Arabidopsis) (Det1) alternative variant bSep08, mRNA.
<a href="#">Dexi</a>	<a href="#">Dexi.cSep08</a>	<a href="#">497857</a>	4344	1463	2	50	dexamethasone-induced transcript (Dexi) alternative variant cSep08, mRNA.
<a href="#">deyby</a>	<a href="#">deyby.aSep08</a>		4113	405		134	putative protein of metazoan origin (deyby) mRNA.
<a href="#">deychy</a>	<a href="#">deychy.aSep08</a>		20247	347		27	putative protein (deychy) mRNA.
<a href="#">deyfer</a>	<a href="#">deyfer.aSep08</a>		2748	546	1	66	CRA c like (7.9 kD) (deyfer) alternative variant aSep08, mRNA.
<a href="#">deyfer</a>	<a href="#">deyfer.bSep08</a>		17160	663	2	61	CRA b like (7.4 kD) (deyfer) alternative variant bSep08, mRNA.
<a href="#">deyflo</a>	<a href="#">deyflo.aSep08</a>		4325	666		36	putative protein (3.9 kD) (deyflo) mRNA.
<a href="#">deyflu</a>	<a href="#">deyflu.aSep08</a>		957	774		32	putative protein (3.4 kD) (deyflu) mRNA.
<a href="#">deykee</a>	<a href="#">deykee.aSep08</a>		2702	1033		83	putative protein (9.3 kD) (deykee) mRNA.
<a href="#">deyloy</a>	<a href="#">deyloy.aSep08</a>		6375	590		57	putative protein (deyloy) mRNA.
<a href="#">deymer</a>	<a href="#">deymer.aSep08</a>		5661	427	3	72	truncated type I keratin KA21 (deymer) alternative variant aSep08, mRNA.
<a href="#">deynoy</a>	<a href="#">deynoy.aSep08</a>		3135	343		84	putative protein (deynoy) mRNA.
<a href="#">deypor</a>	<a href="#">deypor.aSep08</a>		18263	570		40	putative protein (4.5 kD) (deypor) alternative variant aSep08, mRNA.
<a href="#">deypor</a>	<a href="#">deypor.bSep08</a>		18037	493	1	40	putative protein (4.5 kD) (deypor) alternative variant bSep08, mRNA.
<a href="#">deysa</a>	<a href="#">deysa.aSep08</a>		2041	242		80	ubiquitin protein ligase E3 component n-recognin 5 (deysa) mRNA.
<a href="#">deyshee</a>	<a href="#">deyshee.aSep08</a>		312424	510		107	catenin delta 2 (11.3 kD) (deyshee) mRNA.
<a href="#">deyto</a>	<a href="#">deyto.aSep08</a>		1247	213		71	atp-dependent rna helicase tdr9 (deyto) mRNA.
<a href="#">deyvar</a>	<a href="#">deyvar.aSep08</a>		14301	451		149	putative protein of vertebrate origin (deyvar) mRNA.
<a href="#">deywey</a>	<a href="#">deywey.aSep08</a>		2538	754		69	activator-like MOB1 Mps One Binder kinase 1 (deywey) mRNA.
<a href="#">Dffb</a>	<a href="#">Dffb.bSep08</a>	<a href="#">84359</a>	5006	713	6	197	DNA fragmentation factor, beta subunit (Dffb) alternative variant bSep08, mRNA.
<a href="#">Dfna5h</a>	<a href="#">Dfna5h.aSep08</a>	<a href="#">353316</a>	15057	1308	6	278	deafness, autosomal dominant 5 homolog (human) (Dfna5h) alternative variant aSep08, mRNA.
<a href="#">Dfnb59</a>	<a href="#">Dfnb59.aSep08</a>	<a href="#">679552</a>	5898	639	2	206	deafness, autosomal recessive 59 (human) (Dfnb59) alternative variant aSep08, mRNA.
<a href="#">Dfnb59</a>	<a href="#">Dfnb59.bSep08</a>	<a href="#">679552</a>	2008	460	2	118	deafness, autosomal recessive 59 (human) (Dfnb59) alternative variant bSep08, mRNA.
<a href="#">Dgat1</a>	<a href="#">Dgat1.bSep08</a>	<a href="#">84497</a>	1590	744	10	238	diacylglycerol O-acyltransferase 1 (Dgat1) alternative variant bSep08, mRNA.
<a href="#">Dgat1</a>	<a href="#">Dgat1.cSep08</a>	<a href="#">84497</a>	4686	1574	14	214	diacylglycerol O-acyltransferase 1 (Dgat1) alternative variant cSep08, mRNA.
<a href="#">Dgat1</a>	<a href="#">Dgat1.dSep08</a>	<a href="#">84497</a>	1043	662	5	149	diacylglycerol O-acyltransferase 1 (Dgat1) alternative variant dSep08, mRNA.

<a href="#">Dgat1</a>	<a href="#">Dgat1.eSep08</a>	<a href="#">84497</a>	1164	819	5	107	diacylglycerol O-acyltransferase 1 (12.4 kD) (Dgat1) alternative variant eSep08, mRNA.
<a href="#">Dgat2</a>	<a href="#">Dgat2.bSep08</a>	<a href="#">252900</a>	25783	747	3	171	diacylglycerol O-acyltransferase 2 (Dgat2) alternative variant bSep08, mRNA.
<a href="#">Dgat2</a>	<a href="#">Dgat2.cSep08</a>	<a href="#">252900</a>	19837	728	2	84	diacylglycerol O-acyltransferase 2 (9.2 kD) (Dgat2) alternative variant cSep08, mRNA.
<a href="#">Dgcr2</a>	<a href="#">Dgcr2.aSep08</a>	<a href="#">360742</a>	33000	730		150	DiGeorge syndrome critical region gene 2 (Dgcr2) mRNA.
<a href="#">Dgcr6</a>	<a href="#">Dgcr6.bSep08</a>	<a href="#">303794</a>	4659	607	4	197	DiGeorge syndrome critical region gene 6 (Dgcr6) alternative variant bSep08, mRNA.
<a href="#">Dgcr6</a>	<a href="#">Dgcr6.cSep08</a>	<a href="#">303794</a>	1383	775	1	62	DiGeorge syndrome critical region gene 6 (7.0 kD) (Dgcr6) alternative variant cSep08, mRNA.
<a href="#">Dgcr14</a>	<a href="#">Dgcr14.bSep08</a>	<a href="#">360741</a>	5406	2138	2	283	DiGeorge syndrome critical region gene 14 (Dgcr14) alternative variant bSep08, mRNA.
<a href="#">Dgka</a>	<a href="#">Dgka.bSep08</a>	<a href="#">140866</a>	13915	2012	12	226	diacylglycerol kinase alpha (25.4 kD) (Dgka) alternative variant bSep08, mRNA.
<a href="#">Dgka</a>	<a href="#">Dgka.cSep08</a>	<a href="#">140866</a>	2729	595	6	198	diacylglycerol kinase alpha (Dgka) alternative variant cSep08, mRNA.
<a href="#">Dgka</a>	<a href="#">Dgka.dSep08</a>	<a href="#">140866</a>	1614	645	5	193	diacylglycerol kinase alpha (Dgka) alternative variant dSep08, mRNA.
<a href="#">Dgka</a>	<a href="#">Dgka.eSep08</a>	<a href="#">140866</a>	2228	538	3	127	diacylglycerol kinase alpha (13.8 kD) (Dgka) alternative variant eSep08, mRNA.
<a href="#">Dgkb</a>	<a href="#">Dgkb.bSep08</a>	<a href="#">54248</a>	38185	3953	3	90	diacylglycerol kinase, beta (Dgkb) alternative variant bSep08, mRNA.
<a href="#">Dgkb</a>	<a href="#">Dgkb.cSep08</a>	<a href="#">54248</a>	242553	1785	5	75	diacylglycerol kinase, beta (Dgkb) alternative variant cSep08, mRNA.
<a href="#">Dgkg</a>	<a href="#">Dgkg.aSep08</a>	<a href="#">25666</a>	105378	1795	7	572	diacylglycerol kinase gamma (Dgkg) alternative variant aSep08, mRNA.
<a href="#">Dgkg</a>	<a href="#">Dgkg.bSep08</a>	<a href="#">25666</a>	88371	1801	10	428	diacylglycerol kinase gamma (Dgkg) alternative variant bSep08, mRNA.
<a href="#">Dgkg</a>	<a href="#">Dgkg.dSep08</a>	<a href="#">25666</a>	1164	327	2	101	putative protein (Dgkg) alternative variant dSep08, mRNA.
<a href="#">Dgki</a>	<a href="#">Dgki.bSep08</a>	<a href="#">688705</a>	209956	1060	1	150	diacylglycerol kinase, iota (Dgki) alternative variant bSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.bSep08</a>	<a href="#">81821</a>	9161	1454	9	459	diacylglycerol kinase zeta 104kDa (Dgkz) alternative variant bSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.cSep08</a>	<a href="#">81821</a>	4906	1882	13	250	diacylglycerol kinase zeta (27.6 kD) (Dgkz) alternative variant cSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.dSep08</a>	<a href="#">81821</a>	2092	669	7	222	diacylglycerol kinase zeta CRA a (Dgkz) alternative variant dSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.fSep08</a>	<a href="#">81821</a>	4147	1415	10	164	diacylglycerol kinase zeta CRA a (Dgkz) alternative variant fSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.gSep08</a>	<a href="#">81821</a>	31271	397	4	131	diacylglycerol kinase zeta (Dgkz) alternative variant gSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.hSep08</a>	<a href="#">81821</a>	5176	2261	12	126	diacylglycerol kinase zeta 104kDa (13.8 kD) (Dgkz) alternative variant hSep08, mRNA.
<a href="#">Dgkz</a>	<a href="#">Dgkz.iSep08</a>	<a href="#">81821</a>	946	397	3	74	diacylglycerol kinase zeta (Dgkz) alternative variant iSep08, mRNA.

<a href="#">Dguok</a>	<a href="#">Dguok.bSep08</a>	<a href="#">297389</a>	17480	1676	6	259	deoxyguanosine kinase (30.0 kD) (Dguok) alternative variant bSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.cSep08</a>	<a href="#">297389</a>	6460	742	4	247	deoxyguanosine kinase (Dguok) alternative variant cSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.dSep08</a>	<a href="#">297389</a>	21230	733	5	244	deoxyguanosine kinase (Dguok) alternative variant dSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.eSep08</a>	<a href="#">297389</a>	26937	739	5	201	deoxyguanosine kinase (Dguok) alternative variant eSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.fSep08</a>	<a href="#">297389</a>	7458	1198	3	159	deoxyguanosine kinase (Dguok) alternative variant fSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.gSep08</a>	<a href="#">297389</a>	6450	597	4	132	deoxyguanosine kinase (Dguok) alternative variant gSep08, mRNA.
<a href="#">Dguok</a>	<a href="#">Dguok.hSep08</a>	<a href="#">297389</a>	6773	508	3	121	deoxyguanosine kinase (Dguok) alternative variant hSep08, mRNA.
<a href="#">Dhdds</a>	<a href="#">Dhdds.bSep08</a>	<a href="#">298541</a>	23600	2802	7	229	dehydrodolichyl diphosphate synthase (26.4 kD) (Dhdds) alternative variant bSep08, mRNA.
<a href="#">Dhdds</a>	<a href="#">Dhdds.cSep08</a>	<a href="#">298541</a>	21545	662	6	220	dehydrodolichyl diphosphate synthase (Dhdds) alternative variant cSep08, mRNA.
<a href="#">Dhdds</a>	<a href="#">Dhdds.dSep08</a>	<a href="#">298541</a>	7226	766	6	191	dehydrodolichyl diphosphate synthase (Dhdds) alternative variant dSep08, mRNA.
<a href="#">Dhdds</a>	<a href="#">Dhdds.eSep08</a>	<a href="#">298541</a>	21422	653	7	186	dehydrodolichyl diphosphate synthase (Dhdds) alternative variant eSep08, mRNA.
<a href="#">Dhh</a>	<a href="#">Dhh.bSep08</a>	<a href="#">84380</a>	3356	606	2	126	desert hedgehog (Dhh) alternative variant bSep08, mRNA.
<a href="#">Dhodh</a>	<a href="#">Dhodh.bSep08</a>	<a href="#">65156</a>	12875	861	7	262	dihydroorotate dehydrogenase (Dhodh) alternative variant bSep08, mRNA.
<a href="#">Dhps</a>	<a href="#">Dhps.bSep08</a>	<a href="#">288923</a>	1629	787	4	171	deoxyhypusine synthase (Dhps) alternative variant bSep08, mRNA.
<a href="#">Dhps</a>	<a href="#">Dhps.cSep08</a>	<a href="#">288923</a>	1505	1104	2	128	deoxyhypusine synthase (13.9 kD) (Dhps) alternative variant cSep08, complete mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.bSep08</a>	<a href="#">290234</a>	5690	909	6	226	dehydrogenase reductase member 1 (24.3 kD) (Dhrs1) alternative variant bSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.cSep08</a>	<a href="#">290234</a>	5263	815	6	208	dehydrogenase reductase member 1 (Dhrs1) alternative variant cSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.dSep08</a>	<a href="#">290234</a>	5508	773	7	196	dehydrogenase reductase member 1 (Dhrs1) alternative variant dSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.eSep08</a>	<a href="#">290234</a>	2822	698	3	101	dehydrogenase reductase member 1 (10.8 kD) (Dhrs1) alternative variant eSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.fSep08</a>	<a href="#">290234</a>	3136	898	4	101	dehydrogenase reductase member 1 (10.8 kD) (Dhrs1) alternative variant fSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.gSep08</a>	<a href="#">290234</a>	1264	833	3	95	dehydrogenase reductase member 1 CRA a (10.5 kD) (Dhrs1) alternative variant gSep08, mRNA.
<a href="#">Dhrs1</a>	<a href="#">Dhrs1.iSep08</a>	<a href="#">290234</a>	3197	615	4	54	dehydrogenase reductase member 1 (Dhrs1) alternative variant iSep08, mRNA.
<a href="#">Dhrs3</a>	<a href="#">Dhrs3.bSep08</a>	<a href="#">313689</a>	26299	737	5	245	dehydrogenase/reductase (SDR family) member 3 (Dhrs3) alternative variant bSep08, mRNA.
<a href="#">Dhrs3</a>	<a href="#">Dhrs3.cSep08</a>	<a href="#">313689</a>	26439	752	4	147	dehydrogenase/reductase (SDR family) member 3 (Dhrs3) alternative variant cSep08, mRNA.

<a href="#">Dhrs3</a>	<a href="#">Dhrs3.dSep08</a>	<a href="#">313689</a>	23992	757	4	129	dehydrogenase/reductase (SDR family) member 3 (Dhrs3) alternative variant dSep08, mRNA.
<a href="#">Dhrs4</a>	<a href="#">Dhrs4.bSep08</a>	<a href="#">266686</a>	9036	737	6	183	dehydrogenase/reductase (SDR family) member 4 (19.4 kD) (Dhrs4) alternative variant bSep08, mRNA.
<a href="#">Dhrs4</a>	<a href="#">Dhrs4.cSep08</a>	<a href="#">266686</a>	9502	721	7	156	dehydrogenase/reductase (SDR family) member 4 (16.6 kD) (Dhrs4) alternative variant cSep08, mRNA.
<a href="#">Dhrs4</a>	<a href="#">Dhrs4.dSep08</a>	<a href="#">266686</a>	1942	955	2	128	dehydrogenase/reductase (SDR family) member 4 (13.7 kD) (Dhrs4) alternative variant dSep08, mRNA.
<a href="#">Dhrs4</a>	<a href="#">Dhrs4.eSep08</a>	<a href="#">266686</a>	7721	531	4	112	dehydrogenase/reductase (SDR family) member 4 (Dhrs4) alternative variant eSep08, mRNA.
<a href="#">Dhrs4</a>	<a href="#">Dhrs4.gSep08</a>	<a href="#">266686</a>	2463	711	3	47	dehydrogenase/reductase (SDR family) member 4 (5.1 kD) (Dhrs4) alternative variant gSep08, mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.bSep08</a>	<a href="#">287380</a>	32536	1325	6	316	dehydrogenase/reductase (SDR family) member 7B (34.4 kD) (Dhrs7b) alternative variant bSep08, complete mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.cSep08</a>	<a href="#">287380</a>	32375	1071	5	308	dehydrogenase/reductase (SDR family) member 7B (Dhrs7b) alternative variant cSep08, mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.dSep08</a>	<a href="#">287380</a>	31868	1343	5	286	dehydrogenase/reductase (SDR family) member 7B (30.8 kD) (Dhrs7b) alternative variant dSep08, mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.eSep08</a>	<a href="#">287380</a>	17497	849	5	256	dehydrogenase/reductase (SDR family) member 7B (Dhrs7b) alternative variant eSep08, mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.fSep08</a>	<a href="#">287380</a>	28830	1781	4	223	dehydrogenase/reductase (SDR family) member 7B (24.2 kD) (Dhrs7b) alternative variant fSep08, complete mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.gSep08</a>	<a href="#">287380</a>	27296	747	4	174	dehydrogenase/reductase (SDR family) member 7B (Dhrs7b) alternative variant gSep08, mRNA.
<a href="#">Dhrs7b</a>	<a href="#">Dhrs7b.hSep08</a>	<a href="#">287380</a>	9118	766	2	163	dehydrogenase/reductase (SDR family) member 7B (Dhrs7b) alternative variant hSep08, mRNA.
<a href="#">Dhrs7c</a>	<a href="#">Dhrs7c.aSep08</a>	<a href="#">287411</a>	18047	1107		311	dehydrogenase/reductase (SDR family) member 7C (34.4 kD) (Dhrs7c) mRNA.
<a href="#">Dhrsx</a>	<a href="#">Dhrsx.aSep08</a>	<a href="#">288525</a>	1482	636	3	202	dhrsx protein (Dhrsx) alternative variant aSep08, mRNA.
<a href="#">Dhrsx</a>	<a href="#">Dhrsx.bSep08</a>	<a href="#">288525</a>	3190	1026	5	201	dhrsx protein (21.4 kD) (Dhrsx) alternative variant bSep08, mRNA.
<a href="#">Dhrsx</a>	<a href="#">Dhrsx.cSep08</a>	<a href="#">288525</a>	4559	1057	6	201	dhrsx protein (21.4 kD) (Dhrsx) alternative variant cSep08, complete mRNA.
<a href="#">Dhrsx</a>	<a href="#">Dhrsx.eSep08</a>	<a href="#">288525</a>	2377	2136	2	150	putative protein of ancient origin (15.8 kD) (Dhrsx) alternative variant eSep08, mRNA.
<a href="#">Dhrsx</a>	<a href="#">Dhrsx.fSep08</a>	<a href="#">288525</a>	1266	296	2	83	putative protein (Dhrsx) alternative variant fSep08, mRNA.
<a href="#">Dhtkd1</a>	<a href="#">Dhtkd1.bSep08</a>	<a href="#">361272</a>	4138	250		61	putative protein of ancient origin (Dhtkd1) alternative variant bSep08, mRNA.
<a href="#">Dhtkd1</a>	<a href="#">Dhtkd1.cSep08</a>	<a href="#">361272</a>	2724	1613		61	putative protein of ancient origin (Dhtkd1) alternative variant cSep08, mRNA.
<a href="#">Dhx8</a>	<a href="#">Dhx8.bSep08</a>	<a href="#">287727</a>	14943	2399	11	630	DEAH (Asp-Glu-Ala-His) box polypeptide 8 (Dhx8) alternative variant bSep08, mRNA.
<a href="#">Dhx8</a>	<a href="#">Dhx8.cSep08</a>	<a href="#">287727</a>	10835	824	6	274	DEAH (Asp-Glu-Ala-His) box polypeptide 8 (Dhx8) alternative variant cSep08, mRNA.
<a href="#">Dhx8</a>	<a href="#">Dhx8.dSep08</a>	<a href="#">287727</a>	2812	726	4	227	DEAH (Asp-Glu-Ala-His) box polypeptide 8 (Dhx8) alternative variant dSep08, mRNA.

<a href="#">Dhx8</a>	<a href="#">Dhx8.eSep08</a>	<a href="#">287727</a>	10331	760	6	226	DEAH (Asp-Glu-Ala-His) box polypeptide 8 (Dhx8) alternative variant eSep08, mRNA.
<a href="#">Dhx9</a>	<a href="#">Dhx9.bSep08</a>	<a href="#">304859</a>	10721	406	1	134	DEAH (Asp-Glu-Ala-His) box polypeptide 9 (Dhx9) alternative variant bSep08, mRNA.
<a href="#">Dhx15</a>	<a href="#">Dhx15.aSep08</a>	<a href="#">289693</a>	37924	2981	9	795	DEAH (Asp-Glu-Ala-His) box polypeptide 15 (91.0 kD) (Dhx15) alternative variant aSep08, mRNA.
<a href="#">Dhx15</a>	<a href="#">Dhx15.bSep08</a>	<a href="#">289693</a>	18530	1175	2	391	DEAH (Asp-Glu-Ala-His) box polypeptide 15 (Dhx15) alternative variant bSep08, mRNA.
<a href="#">Dhx15</a>	<a href="#">Dhx15.cSep08</a>	<a href="#">289693</a>	5296	1134	1	256	DEAH (Asp-Glu-Ala-His) box polypeptide 15 (Dhx15) alternative variant cSep08, mRNA.
<a href="#">Dhx15</a>	<a href="#">Dhx15.dSep08</a>	<a href="#">289693</a>	7692	1609	4	218	DEAH (Asp-Glu-Ala-His) box polypeptide 15 (25.9 kD) (Dhx15) alternative variant dSep08, mRNA.
<a href="#">Dhx16</a>	<a href="#">Dhx16.bSep08</a>	<a href="#">294232</a>	9588	1958	14	415	deah box polypeptide 16 (47.2 kD) (Dhx16) alternative variant bSep08, mRNA.
<a href="#">Dhx16</a>	<a href="#">Dhx16.cSep08</a>	<a href="#">294232</a>	2212	732	4	133	deah box polypeptide 16 (15.3 kD) (Dhx16) alternative variant cSep08, mRNA.
<a href="#">Dhx16</a>	<a href="#">Dhx16.dSep08</a>	<a href="#">294232</a>	2648	730	6	127	deah box polypeptide 16 (Dhx16) alternative variant dSep08, mRNA.
<a href="#">Dhx16</a>	<a href="#">Dhx16.eSep08</a>	<a href="#">294232</a>	2335	653	5	99	deah box polypeptide 16 (11.4 kD) (Dhx16) alternative variant eSep08, mRNA.
<a href="#">Dhx16</a>	<a href="#">Dhx16.gSep08</a>	<a href="#">294232</a>	1877	677	2	45	deah box polypeptide 16 (5.3 kD) (Dhx16) alternative variant gSep08, mRNA.
<a href="#">Dhx30</a>	<a href="#">Dhx30.bSep08</a>	<a href="#">367172</a>	1207	743	5	247	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (Dhx30) alternative variant bSep08, mRNA.
<a href="#">Dhx30</a>	<a href="#">Dhx30.cSep08</a>	<a href="#">367172</a>	1089	809	3	159	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (18.0 kD) (Dhx30) alternative variant cSep08, mRNA.
<a href="#">Dhx30</a>	<a href="#">Dhx30.dSep08</a>	<a href="#">367172</a>	6465	462	4	153	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (Dhx30) alternative variant dSep08, mRNA.
<a href="#">Dhx30</a>	<a href="#">Dhx30.eSep08</a>	<a href="#">367172</a>	8689	747	4	143	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (Dhx30) alternative variant eSep08, mRNA.
<a href="#">Dhx30</a>	<a href="#">Dhx30.fSep08</a>	<a href="#">367172</a>	9047	431	4	54	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (6.1 kD) (Dhx30) alternative variant fSep08, mRNA.
<a href="#">Dhx32</a>	<a href="#">Dhx32.bSep08</a>	<a href="#">361667</a>	1697	960	2	104	DEAH (Asp-Glu-Ala-His) box polypeptide 32 (11.9 kD) (Dhx32) alternative variant bSep08, mRNA.
<a href="#">Dhx35</a>	<a href="#">Dhx35.bSep08</a>	<a href="#">362260</a>	7575	350	4	116	DEAH (Asp-Glu-Ala-His) box polypeptide 35 (Dhx35) alternative variant bSep08, mRNA.
<a href="#">Dhx35</a>	<a href="#">Dhx35.cSep08</a>	<a href="#">362260</a>	2824	504	2	62	DEAH (Asp-Glu-Ala-His) box polypeptide 35 (Dhx35) alternative variant cSep08, mRNA.
<a href="#">Dhx36</a>	<a href="#">Dhx36.bSep08</a>	<a href="#">310461</a>	2391	931	2	37	DEAH (Asp-Glu-Ala-His) box polypeptide 36 (4.2 kD) (Dhx36) alternative variant bSep08, mRNA.
<a href="#">Dhx36</a>	<a href="#">Dhx36.cSep08</a>	<a href="#">310461</a>	1883	889	2	33	DEAH (Asp-Glu-Ala-His) box polypeptide 36 (3.6 kD) (Dhx36) alternative variant cSep08, mRNA.
<a href="#">Dhx37</a>	<a href="#">Dhx37.bSep08</a>	<a href="#">288647</a>	2171	1132	7	180	DEAH (Asp-Glu-Ala-His) box polypeptide 37 (Dhx37) alternative variant bSep08, mRNA.
<a href="#">Dhx37</a>	<a href="#">Dhx37.dSep08</a>	<a href="#">288647</a>	1804	468	4	95	DEAH (Asp-Glu-Ala-His) box polypeptide 37 (Dhx37) alternative variant dSep08, mRNA.

<a href="#">Dhx38</a>	<a href="#">Dhx38.bSep08</a>	<a href="#">292007</a>	2025	686	6	141	DEAH box polypeptide 38 (Dhx38) alternative variant bSep08, mRNA.
<a href="#">Dhx38</a>	<a href="#">Dhx38.cSep08</a>	<a href="#">292007</a>	3344	493	3	122	DEAH box polypeptide 38 (Dhx38) alternative variant cSep08, mRNA.
<a href="#">Dhx38</a>	<a href="#">Dhx38.dSep08</a>	<a href="#">292007</a>	682	415	3	76	DEAH box polypeptide 38 (Dhx38) alternative variant dSep08, mRNA.
<a href="#">Dhx40</a>	<a href="#">Dhx40.bSep08</a>	<a href="#">287595</a>	7438	1905	2	137	DEAH (Asp-Glu-Ala-His) box polypeptide 40 (16.2 kD) (Dhx40) alternative variant bSep08, mRNA.
<a href="#">Dhx57</a>	<a href="#">Dhx57.aSep08</a>	<a href="#">366532</a>	23417	2297	1	600	DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 (Dhx57) alternative variant aSep08, mRNA.
<a href="#">Dhx57</a>	<a href="#">Dhx57.bSep08</a>	<a href="#">366532</a>	3118	757	1	88	DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 (9.8 kD) (Dhx57) alternative variant bSep08, mRNA.
<a href="#">Dhx58</a>	<a href="#">Dhx58.bSep08</a>	<a href="#">303538</a>	624	385	1	127	DEXH (Asp-Glu-X-His) box polypeptide 58 (Dhx58) alternative variant bSep08, mRNA.
<a href="#">Diablo</a>	<a href="#">Diablo.bSep08</a>	<a href="#">288753</a>	2568	1178	2	117	diablo homolog (Drosophila) (Diablo) alternative variant bSep08, mRNA.
<a href="#">Diap1</a>	<a href="#">Diap1.aSep08</a>	<a href="#">307483</a>	43290	1800	12	600	diaphanous homolog 1 (Drosophila) (Diap1) alternative variant aSep08, mRNA.
<a href="#">Diap1</a>	<a href="#">Diap1.cSep08</a>	<a href="#">307483</a>	56248	1785	3	368	diaphanous homolog 1 (Drosophila) (Diap1) alternative variant cSep08, mRNA.
<a href="#">Diap1</a>	<a href="#">Diap1.dSep08</a>	<a href="#">307483</a>	10198	2206	4	160	diaphanous homolog 1 (Drosophila) (Diap1) alternative variant dSep08, mRNA.
<a href="#">Diap1</a>	<a href="#">Diap1.fSep08</a>	<a href="#">307483</a>	633	486	2	84	diaphanous homolog 1 (Drosophila) (Diap1) alternative variant fSep08, mRNA.
<a href="#">Diap3</a>	<a href="#">Diap3.aSep08</a>	<a href="#">290396</a>	66568	628		209	diaphanous homolog 3 (Drosophila) (Diap3) mRNA.
<a href="#">Dicer1</a>	<a href="#">Dicer1.aSep08</a>	<a href="#">299284</a>	13196	2650	7	725	dicer1, Dcr-1 homolog (Drosophila) (Dicer1) alternative variant aSep08, mRNA.
<a href="#">Dido1</a>	<a href="#">Dido1.bSep08</a>	<a href="#">362286</a>	15215	1312	1	75	death inducer-obliterator 1 (8.5 kD) (Dido1) alternative variant bSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.bSep08</a>	<a href="#">294718</a>	4983	671	7	190	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (Dimt1) alternative variant bSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.cSep08</a>	<a href="#">294718</a>	4946	428	5	142	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (Dimt1) alternative variant cSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.dSep08</a>	<a href="#">294718</a>	2426	653	3	72	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (Dimt1) alternative variant dSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.eSep08</a>	<a href="#">294718</a>	3552	1975	2	79	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (9.2 kD) (Dimt1) alternative variant eSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.fSep08</a>	<a href="#">294718</a>	13339	1005	6	86	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (9.5 kD) (Dimt1) alternative variant fSep08, mRNA.
<a href="#">Dimt1</a>	<a href="#">Dimt1.gSep08</a>	<a href="#">294718</a>	2002	851	3	73	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (8.5 kD) (Dimt1) alternative variant gSep08, mRNA.
<a href="#">Dio1</a>	<a href="#">Dio1.aSep08</a>	<a href="#">25430</a>	18219	710	3	141	deiodinase, iodothyronine, type I (15.9 kD) (Dio1) alternative variant aSep08, mRNA.
<a href="#">Dio1</a>	<a href="#">Dio1.bSep08</a>	<a href="#">25430</a>	17631	751	3	125	deiodinase, iodothyronine, type I (14.5 kD) (Dio1) alternative variant bSep08, mRNA.
<a href="#">Dip2a</a>	<a href="#">Dip2a.aSep08</a>	<a href="#">690211</a>	15152	940		313	DIP2 disco-interacting protein 2 homolog A (Drosophila) (Dip2a) mRNA.

<a href="#">Dip2b</a>	<a href="#">Dip2b.aSep08</a>	<a href="#">300231</a>	22544	1970		547	DIP2 disco-interacting protein 2 homolog B (Drosophila) (Dip2b) alternative variant aSep08, mRNA.
<a href="#">Dis3l</a>	<a href="#">Dis3l.bSep08</a>	<a href="#">363077</a>	2767	844	3	169	DIS3 mitotic control homolog (S. cerevisiae)-like (Dis3l) alternative variant bSep08, mRNA.
<a href="#">Dis3l</a>	<a href="#">Dis3l.cSep08</a>	<a href="#">363077</a>	4051	445	2	148	DIS3 mitotic control homolog (S. cerevisiae)-like (Dis3l) alternative variant cSep08, mRNA.
<a href="#">Dis3l2</a>	<a href="#">Dis3l2.aSep08</a>	<a href="#">367307</a>	379823	2001	13	626	DIS3 mitotic control homolog (S. cerevisiae)-like 2 (Dis3l2) alternative variant aSep08, mRNA.
<a href="#">Dis3l2</a>	<a href="#">Dis3l2.bSep08</a>	<a href="#">367307</a>	55081	223	2	67	DIS3 mitotic control homolog (S. cerevisiae)-like 2 (Dis3l2) alternative variant bSep08, mRNA.
<a href="#">Disp1</a>	<a href="#">Disp1.bSep08</a>	<a href="#">289338</a>	6568	356	1	118	dispatched homolog 1 (Drosophila) (Disp1) alternative variant bSep08, mRNA.
<a href="#">Disp2</a>	<a href="#">Disp2.aSep08</a>	<a href="#">311324</a>	8393	6176	3	1260	dispatched homolog 2 (Drosophila) (139.0 kD) (Disp2) alternative variant aSep08, mRNA.
<a href="#">DIX.0</a>	<a href="#">DIX.0.aSep08</a>		5342	931		244	dishevelled 2 (DIX.0) mRNA.
<a href="#">Dixdc1</a>	<a href="#">Dixdc1.bSep08</a>	<a href="#">363062</a>	3477	607	3	82	DIX (Dixdc1) alternative variant bSep08, mRNA.
<a href="#">Dixdc1</a>	<a href="#">Dixdc1.cSep08</a>	<a href="#">363062</a>	701	248	2	55	putative protein, with a coiled coil domain, of vertebrate origin (Dixdc1) alternative variant cSep08, mRNA.
<a href="#">Dkk2</a>	<a href="#">Dkk2.bSep08</a>	<a href="#">295445</a>	6469	2810	2	159	dickkopf homolog 2 (Xenopus laevis) (18.1 kD) (Dkk2) alternative variant bSep08, mRNA.
<a href="#">Dkk3</a>	<a href="#">Dkk3.bSep08</a>	<a href="#">171548</a>	29908	1010	2	176	dickkopf homolog 3 (Xenopus laevis) (18.3 kD) (Dkk3) alternative variant bSep08, mRNA.
<a href="#">Dkk3</a>	<a href="#">Dkk3.cSep08</a>	<a href="#">171548</a>	6440	386	2	128	dickkopf homolog 3 (Xenopus laevis) (Dkk3) alternative variant cSep08, mRNA.
<a href="#">Dlc1</a>	<a href="#">Dlc1.bSep08</a>	<a href="#">58834</a>	1768	653	2	87	deleted in liver cancer 1 (Dlc1) alternative variant bSep08, mRNA.
<a href="#">Dld</a>	<a href="#">Dld.bSep08</a>	<a href="#">298942</a>	8354	2552	6	255	dihydrolipoamide dehydrogenase (27.3 kD) (Dld) alternative variant bSep08, mRNA.
<a href="#">Dld</a>	<a href="#">Dld.cSep08</a>	<a href="#">298942</a>	10604	772	8	226	dihydrolipoamide dehydrogenase (Dld) alternative variant cSep08, mRNA.
<a href="#">Dld</a>	<a href="#">Dld.dSep08</a>	<a href="#">298942</a>	13665	847	9	139	dihydrolipoamide dehydrogenase (Dld) alternative variant dSep08, mRNA.
<a href="#">Dld</a>	<a href="#">Dld.eSep08</a>	<a href="#">298942</a>	711	616	2	65	dihydrolipoamide dehydrogenase (Dld) alternative variant eSep08, mRNA.
<a href="#">Dlec1</a>	<a href="#">Dlec1.aSep08</a>	<a href="#">501070</a>	7736	694		230	deleted in lung and esophageal cancer 1 isoform DLEC1-N1 (Dlec1) mRNA.
<a href="#">Dleu7</a>	<a href="#">Dleu7.bSep08</a>	<a href="#">290308</a>	16186	1584		209	deleted in lymphocytic leukemia, 7 (22.6 kD) (Dleu7) alternative variant bSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.bSep08</a>	<a href="#">25252</a>	196138	2882	13	467	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant bSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.cSep08</a>	<a href="#">25252</a>	28300	730	9	242	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant cSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.dSep08</a>	<a href="#">25252</a>	39323	725	8	241	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant dSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.eSep08</a>	<a href="#">25252</a>	18010	725	8	241	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant eSep08, mRNA.



<a href="#">Dlg1</a>	<a href="#">Dlg1.fSep08</a>	<a href="#">25252</a>	9945	755	5	208	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant fSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.gSep08</a>	<a href="#">25252</a>	14610	1138	4	136	discs, large homolog 1 (Drosophila) (15.8 kD) (Dlg1) alternative variant gSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.hSep08</a>	<a href="#">25252</a>	39640	539	4	134	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant hSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.iSep08</a>	<a href="#">25252</a>	24229	731	9	132	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant iSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.jSep08</a>	<a href="#">25252</a>	21938	397	6	131	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant jSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.kSep08</a>	<a href="#">25252</a>	11484	626	4	100	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant kSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.mSep08</a>	<a href="#">25252</a>	11365	312	4	69	discs, large homolog 1 (Drosophila) (Dlg1) alternative variant mSep08, mRNA.
<a href="#">Dlg1</a>	<a href="#">Dlg1.nSep08</a>	<a href="#">25252</a>	196198	1645	7	133	discs, large homolog 1 (Drosophila) (14.7 kD) (Dlg1) alternative variant nSep08, mRNA.
<a href="#">Dlg2</a>	<a href="#">Dlg2.bSep08</a>	<a href="#">64053</a>	183445	456	4	99	discs large homolog 2 CRA b (Dlg2) alternative variant bSep08, mRNA.
<a href="#">Dlg2</a>	<a href="#">Dlg2.cSep08</a>	<a href="#">64053</a>	21512	404	2	66	putative protein (Dlg2) alternative variant cSep08, mRNA.
<a href="#">Dlg2</a>	<a href="#">Dlg2.dSep08</a>	<a href="#">64053</a>	2704	1009	2	64	discs large homolog 2 CRA b (7.6 kD) (Dlg2) alternative variant dSep08, mRNA.
<a href="#">Dlg2</a>	<a href="#">Dlg2.eSep08</a>	<a href="#">64053</a>	331628	458	4	62	putative protein of vertebrate origin (7.2 kD) (Dlg2) alternative variant eSep08, mRNA.
<a href="#">Dlg2</a>	<a href="#">Dlg2.fSep08</a>	<a href="#">64053</a>	93309	393	2	64	putative protein (Dlg2) alternative variant fSep08, mRNA.
<a href="#">Dlg3</a>	<a href="#">Dlg3.bSep08</a>	<a href="#">58948</a>	37122	638	8	212	discs, large homolog 3 (Drosophila) (Dlg3) alternative variant bSep08, mRNA.
<a href="#">Dlg3</a>	<a href="#">Dlg3.cSep08</a>	<a href="#">58948</a>	23461	637	6	154	discs, large homolog 3 (Drosophila) (Dlg3) alternative variant cSep08, mRNA.
<a href="#">Dlg3</a>	<a href="#">Dlg3.dSep08</a>	<a href="#">58948</a>	4173	2873	3	66	discs, large homolog 3 (Drosophila) (7.9 kD) (Dlg3) alternative variant dSep08, mRNA.
<a href="#">Dlg4</a>	<a href="#">Dlg4.aSep08</a>	<a href="#">29495</a>	5100	1110	10	249	discs, large homolog 4 (Drosophila) (Dlg4) alternative variant aSep08, mRNA.
<a href="#">Dlg4</a>	<a href="#">Dlg4.cSep08</a>	<a href="#">29495</a>	831	681	2	125	discs, large homolog 4 (Drosophila) (Dlg4) alternative variant cSep08, mRNA.
<a href="#">Dlg4</a>	<a href="#">Dlg4.dSep08</a>	<a href="#">29495</a>	1854	1141	4	101	discs, large homolog 4 (Drosophila) (Dlg4) alternative variant dSep08, mRNA.
<a href="#">Dlg5</a>	<a href="#">Dlg5.cSep08</a>	<a href="#">305645</a>	10096	572	5	190	discs, large homolog 5 (Drosophila) (Dlg5) alternative variant cSep08, mRNA.
<a href="#">Dlg5</a>	<a href="#">Dlg5.dSep08</a>	<a href="#">305645</a>	9065	358	3	118	discs, large homolog 5 (Drosophila) (Dlg5) alternative variant dSep08, mRNA.
<a href="#">Dlg5</a>	<a href="#">Dlg5.eSep08</a>	<a href="#">305645</a>	2415	897	3	109	discs, large homolog 5 (Drosophila) (Dlg5) alternative variant eSep08, mRNA.
<a href="#">Dlg7</a>	<a href="#">Dlg7.aSep08</a>	<a href="#">289997</a>	34803	2739	19	815	discs, large homolog 7 (Drosophila) (90.5 kD) (Dlg7) alternative variant aSep08, mRNA.
<a href="#">Dlg7</a>	<a href="#">Dlg7.bSep08</a>	<a href="#">289997</a>	5096	408	2	63	discs, large homolog 7 (Drosophila) (Dlg7) alternative variant bSep08, mRNA.

<a href="#">Dlgap1</a>	<a href="#">Dlgap1.bSep08</a>	<a href="#">65040</a>	141481	748	4	249	discs, large (Drosophila) homolog-associated protein 1 (Dlgap1) alternative variant bSep08, mRNA.
<a href="#">Dlgap1</a>	<a href="#">Dlgap1.cSep08</a>	<a href="#">65040</a>	5627	388	3	63	discs, large (Drosophila) homolog-associated protein 1 (Dlgap1) alternative variant cSep08, mRNA.
<a href="#">Dlgap4</a>	<a href="#">Dlgap4.bSep08</a>	<a href="#">286930</a>	27821	842	4	280	discs, large homolog-associated protein 4 (Drosophila) (Dlgap4) alternative variant bSep08, mRNA.
<a href="#">Dlgap4</a>	<a href="#">Dlgap4.cSep08</a>	<a href="#">286930</a>	27787	817	4	271	discs, large homolog-associated protein 4 (Drosophila) (Dlgap4) alternative variant cSep08, mRNA.
<a href="#">Dlgap4</a>	<a href="#">Dlgap4.dSep08</a>	<a href="#">286930</a>	7051	787	4	194	discs, large homolog-associated protein 4 (Drosophila) (Dlgap4) alternative variant dSep08, mRNA.
<a href="#">Dlgap4</a>	<a href="#">Dlgap4.fSep08</a>	<a href="#">286930</a>	783	477	2	49	discs, large homolog-associated protein 4 (Drosophila) (Dlgap4) alternative variant fSep08, mRNA.
<a href="#">Dlk1</a>	<a href="#">Dlk1.bSep08</a>	<a href="#">114587</a>	5835	1400	1	383	delta-like 1 homolog (Drosophila) (Dlk1) alternative variant bSep08, mRNA.
<a href="#">Dlk2</a>	<a href="#">Dlk2.bSep08</a>	<a href="#">316232</a>	4344	1781	2	331	delta-like 2 homolog (Drosophila) (Dlk2) alternative variant bSep08, mRNA.
<a href="#">Dlst</a>	<a href="#">Dlst.bSep08</a>	<a href="#">299201</a>	13620	753	9	250	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) (Dlst) alternative variant bSep08, mRNA.
<a href="#">Dlst</a>	<a href="#">Dlst.cSep08</a>	<a href="#">299201</a>	8697	3385	6	164	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) (Dlst) alternative variant cSep08, mRNA.
<a href="#">Dlst</a>	<a href="#">Dlst.dSep08</a>	<a href="#">299201</a>	8379	453	7	102	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) (Dlst) alternative variant dSep08, mRNA.
<a href="#">Dlst</a>	<a href="#">Dlst.eSep08</a>	<a href="#">299201</a>	7565	432	6	48	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) (Dlst) alternative variant eSep08, mRNA.
<a href="#">Dlx1</a>	<a href="#">Dlx1.aSep08</a>	<a href="#">296500</a>	2825	2171	2	189	distal-less homeobox 1 (20.2 kD) (Dlx1) alternative variant aSep08, mRNA.
<a href="#">Dlx1</a>	<a href="#">Dlx1.bSep08</a>	<a href="#">296500</a>	1583	403	2	89	distal-less homeobox 1 (Dlx1) alternative variant bSep08, mRNA.
<a href="#">Dlx2</a>	<a href="#">Dlx2.aSep08</a>	<a href="#">296499</a>	1112	631		165	distal-less homeobox 2 (Dlx2) mRNA.
<a href="#">Dmap1</a>	<a href="#">Dmap1.aSep08</a>	<a href="#">298447</a>	7676	1460	10	449	DNA methyltransferase 1 associated protein (Dmap1) alternative variant aSep08, mRNA.
<a href="#">Dmap1</a>	<a href="#">Dmap1.bSep08</a>	<a href="#">298447</a>	2039	882	6	275	DNA methyltransferase 1 associated protein (Dmap1) alternative variant bSep08, mRNA.
<a href="#">Dmap1</a>	<a href="#">Dmap1.cSep08</a>	<a href="#">298447</a>	958	624	3	142	DNA methyltransferase protein 1 (15.2 kD) (Dmap1) alternative variant cSep08, mRNA.
<a href="#">Dmap1</a>	<a href="#">Dmap1.dSep08</a>	<a href="#">298447</a>	1164	498	3	95	CRA a like (10.4 kD) (Dmap1) alternative variant dSep08, mRNA.
<a href="#">DMAP_binding_0</a>	<a href="#">DMAP_binding_0.aSep08</a>		28972	580		151	dip2 Disco-interacting protein 2 homolog A (16.8 kD) (DMAP_binding_0) complete mRNA.
<a href="#">Dmbt1</a>	<a href="#">Dmbt1.aSep08</a>	<a href="#">170568</a>	19319	1316		309	deleted in malignant brain tumors 1 (Dmbt1) mRNA.
<a href="#">Dmc1</a>	<a href="#">Dmc1.bSep08</a>	<a href="#">362960</a>	5399	797	6	60	DMC1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast) (7.0 kD) (Dmc1) alternative variant bSep08, mRNA.

<a href="#">Dmkn</a>	<a href="#">Dmkn.aSep08</a>	<a href="#">361548</a>	17396	2033	14	569	dermokine (Dmkn) alternative variant aSep08, mRNA.
<a href="#">Dmkn</a>	<a href="#">Dmkn.bSep08</a>	<a href="#">361548</a>	4668	644	1	107	dermokine (Dmkn) alternative variant bSep08, mRNA.
<a href="#">Dmn</a>	<a href="#">Dmn.bSep08</a>	<a href="#">308709</a>	23516	3317	3	1032	desmuslin (Dmn) alternative variant bSep08, mRNA.
<a href="#">Dmn</a>	<a href="#">Dmn.cSep08</a>	<a href="#">308709</a>	1424	518	2	172	desmuslin (Dmn) alternative variant cSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.aSep08</a>	<a href="#">308405</a>	11001	2910	14	687	myotonic dystrophy kinase (76.1 kD) (Dmpk) alternative variant aSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.bSep08</a>	<a href="#">308405</a>	6904	1360	9	387	myotonic dystrophy kinase (Dmpk) alternative variant bSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.dSep08</a>	<a href="#">308405</a>	1612	730	5	173	myotonic dystrophy kinase (Dmpk) alternative variant dSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.eSep08</a>	<a href="#">308405</a>	1443	1348	2	164	myotonic dystrophy kinase (18.0 kD) (Dmpk) alternative variant eSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.fSep08</a>	<a href="#">308405</a>	927	383	4	127	myotonic dystrophy kinase (Dmpk) alternative variant fSep08, mRNA.
<a href="#">Dmpk</a>	<a href="#">Dmpk.hSep08</a>	<a href="#">308405</a>	528	358	2	96	myotonic dystrophy kinase (Dmpk) alternative variant hSep08, mRNA.
<a href="#">Dmrt2</a>	<a href="#">Dmrt2.bSep08</a>	<a href="#">309430</a>	4973	982	4	123	doublesex and mab-3 related transcription factor 2 (13.6 kD) (Dmrt2) alternative variant bSep08, complete mRNA.
<a href="#">Dmrta2</a>	<a href="#">Dmrta2.bSep08</a>	<a href="#">313471</a>	2010	731	1	243	doublesex and mab-3 related transcription factor like family A2 (Dmrta2) alternative variant bSep08, mRNA.
<a href="#">Dmrtb1</a>	<a href="#">Dmrtb1.aSep08</a>	<a href="#">313484</a>	6951	1280	4	189	DMRT-like family B with proline-rich C-terminal, 1 (Dmrtb1) alternative variant aSep08, mRNA.
<a href="#">Dmrta1a</a>	<a href="#">Dmrta1a.bSep08</a>	<a href="#">501668</a>	2894	680	1	125	DMRT-like family C1a (13.5 kD) (Dmrta1a) alternative variant bSep08, mRNA.
<a href="#">Dmrta1b</a>	<a href="#">Dmrta1b.aSep08</a>	<a href="#">680068</a>	33763	856	3	251	DMRT-like family C1b (Dmrta1b) alternative variant aSep08, mRNA.
<a href="#">Dmrta1b</a>	<a href="#">Dmrta1b.bSep08</a>	<a href="#">680068</a>	7884	431	1	131	DMRT-like family C1b (Dmrta1b) alternative variant bSep08, mRNA.
<a href="#">Dmrta1c</a>	<a href="#">Dmrta1c.bSep08</a>	<a href="#">363483</a>	5927	1318	8	224	DMRT-like family C1c (24.4 kD) (Dmrta1c) alternative variant bSep08, mRNA.
<a href="#">Dmrta2</a>	<a href="#">Dmrta2.bSep08</a>	<a href="#">499100</a>	1701	458	1	139	doublesex and mab-3 related transcription factor like family C2 (Dmrta2) alternative variant bSep08, mRNA.
<a href="#">Dmtf1</a>	<a href="#">Dmtf1.bSep08</a>	<a href="#">114485</a>	19133	782	5	110	cyclin D binding myb-like transcription factor 1 (11.9 kD) (Dmtf1) alternative variant bSep08, mRNA.
<a href="#">Dmxl1</a>	<a href="#">Dmxl1.aSep08</a>	<a href="#">307429</a>	28434	974		324	dmx-like 1 (Dmxl1) mRNA.
<a href="#">Dmxl2</a>	<a href="#">Dmxl2.aSep08</a>	<a href="#">315676</a>	11621	742	8	247	dmx-like 2 (Dmxl2) alternative variant aSep08, mRNA.
<a href="#">Dmxl2</a>	<a href="#">Dmxl2.bSep08</a>	<a href="#">315676</a>	9175	745	7	247	dmx-like 2 (Dmxl2) alternative variant bSep08, mRNA.
<a href="#">Dmxl2</a>	<a href="#">Dmxl2.cSep08</a>	<a href="#">315676</a>	9486	707	6	140	dmx-like 2 (Dmxl2) alternative variant cSep08, mRNA.
<a href="#">Dmxl2</a>	<a href="#">Dmxl2.dSep08</a>	<a href="#">315676</a>	4704	488	5	120	dmx-like 2 (Dmxl2) alternative variant dSep08, mRNA.
<a href="#">Dna2</a>	<a href="#">Dna2.aSep08</a>	<a href="#">309762</a>	5650	966	3	322	DNA replication helicase 2 homolog (yeast) (Dna2) alternative variant aSep08, mRNA.
<a href="#">Dna2</a>	<a href="#">Dna2.bSep08</a>	<a href="#">309762</a>	1909	1091	1	129	DNA replication helicase 2 homolog (yeast) (Dna2) alternative variant bSep08, mRNA.
<a href="#">Dna2.1</a>	<a href="#">Dna2.1.aSep08</a>		8324	780		260	putative protein of eukaryotic origin (Dna2.1) mRNA.
<a href="#">Dnah3_predicted</a>	<a href="#">Dnah3_predicted.aSep08</a>	<a href="#">293539</a>	2827	502		149	dynein, axonemal, heavy polypeptide 3 (predicted) (Dnah3_predicted) mRNA.

<a href="#">Dnah9</a>	<a href="#">Dnah9.aSep08</a>	<a href="#">117251</a>	8320	923		233	dynein, axonemal, heavy polypeptide 9 (Dnah9) mRNA.
<a href="#">Dnah10</a>	<a href="#">Dnah10.aSep08</a>	<a href="#">117252</a>	1719	743		247	dynein, axonemal, heavy polypeptide 10 (Dnah10) mRNA.
<a href="#">Dnah12</a>	<a href="#">Dnah12.aSep08</a>	<a href="#">252959</a>	13903	1366	3	454	dynein, axonemal, heavy polypeptide 12 (Dnah12) alternative variant aSep08, mRNA.
<a href="#">Dnah12</a>	<a href="#">Dnah12.bSep08</a>	<a href="#">252959</a>	5038	655	1	163	dynein, axonemal, heavy polypeptide 12 (Dnah12) alternative variant bSep08, mRNA.
<a href="#">Dnahc2</a>	<a href="#">Dnahc2.aSep08</a>	<a href="#">303242</a>	1762	749		249	dynein, axonemal, heavy chain 2 (Dnahc2) mRNA.
<a href="#">Dnahc5</a>	<a href="#">Dnahc5.aSep08</a>	<a href="#">294854</a>	12545	1320		118	dynein, axonemal, heavy chain 5 (Dnahc5) mRNA.
<a href="#">Dnahc6</a>	<a href="#">Dnahc6.aSep08</a>	<a href="#">117250</a>	8994	767		255	dynein, axonemal, heavy chain 6 (Dnahc6) mRNA.
<a href="#">Dnahc8</a>	<a href="#">Dnahc8.aSep08</a>	<a href="#">207117</a>	23161	716		146	dynein, axonemal, heavy chain 8 (16.5 kD) (Dnahc8) mRNA.
<a href="#">Dnahc11</a>	<a href="#">Dnahc11.aSep08</a>	<a href="#">117253</a>	5235	977	4	194	dynein, axonemal, heavy chain 11 (Dnahc11) alternative variant aSep08, mRNA.
<a href="#">Dnahc17</a>	<a href="#">Dnahc17.bSep08</a>	<a href="#">287845</a>	3304	361	1	107	dynein heavy axonemal (Dnahc17) alternative variant bSep08, mRNA.
<a href="#">DnaJ.0</a>	<a href="#">DnaJ.0.aSep08</a>		156795	2088	7	565	DnaJ homolog subfamily C member 1 (65.4 kD) (DnaJ.0) alternative variant aSep08, mRNA.
<a href="#">DnaJ.0</a>	<a href="#">DnaJ.0.bSep08</a>		36455	583	1	39	putative protein (4.7 kD) (DnaJ.0) alternative variant bSep08, mRNA.
<a href="#">DnaJ.1</a>	<a href="#">DnaJ.1.bSep08</a>		1052	305	2	70	DnaJ homolog subfamily B member 4 (DnaJ.1) alternative variant bSep08, mRNA.
<a href="#">Dnaja1</a>	<a href="#">Dnaja1.bSep08</a>	<a href="#">65028</a>	11883	3506	8	336	DnaJ homolog subfamily A member 1 (37.6 kD) (Dnaja1) alternative variant bSep08, mRNA.
<a href="#">Dnaja1</a>	<a href="#">Dnaja1.cSep08</a>	<a href="#">65028</a>	6448	1228	5	217	DnaJ homolog subfamily A member 1 (24.2 kD) (Dnaja1) alternative variant cSep08, mRNA.
<a href="#">Dnaja1</a>	<a href="#">Dnaja1.dSep08</a>	<a href="#">65028</a>	4560	2872	4	182	DnaJ homolog subfamily A member 1 (20.9 kD) (Dnaja1) alternative variant dSep08, mRNA.
<a href="#">Dnaja1</a>	<a href="#">Dnaja1.eSep08</a>	<a href="#">65028</a>	9447	686	6	103	DnaJ homolog subfamily A member 1 (Dnaja1) alternative variant eSep08, mRNA.
<a href="#">Dnaja2</a>	<a href="#">Dnaja2.bSep08</a>	<a href="#">84026</a>	6404	408	4	135	DnaJ (Hsp40) homolog, subfamily A, member 2 (Dnaja2) alternative variant bSep08, mRNA.
<a href="#">Dnaja3</a>	<a href="#">Dnaja3.bSep08</a>	<a href="#">360481</a>	21442	1388	2	342	DnaJ (Hsp40) homolog, subfamily A, member 3 (Dnaja3) alternative variant bSep08, mRNA.
<a href="#">Dnaja3</a>	<a href="#">Dnaja3.cSep08</a>	<a href="#">360481</a>	15162	1233	1	271	DnaJ (Hsp40) homolog, subfamily A, member 3 (Dnaja3) alternative variant cSep08, mRNA.
<a href="#">Dnaja4</a>	<a href="#">Dnaja4.bSep08</a>	<a href="#">300721</a>	10449	1505	4	220	DnaJ homolog subfamily A member (24.5 kD) (Dnaja4) alternative variant bSep08, mRNA.
<a href="#">Dnaja4</a>	<a href="#">Dnaja4.cSep08</a>	<a href="#">300721</a>	5912	2172	3	181	DnaJ homolog subfamily A member 4 (20.8 kD) (Dnaja4) alternative variant cSep08, mRNA.
<a href="#">Dnaja4</a>	<a href="#">Dnaja4.dSep08</a>	<a href="#">300721</a>	1227	1127	2	89	DnaJ homolog subfamily A member 4 (10.5 kD) (Dnaja4) alternative variant dSep08, mRNA.
<a href="#">Dnajb1</a>	<a href="#">Dnajb1.aSep08</a>	<a href="#">361384</a>	3674	2176	3	340	DnaJ (Hsp40) homolog, subfamily B, member 1 (38.1 kD) (Dnajb1) alternative variant aSep08, mRNA.
<a href="#">Dnajb1</a>	<a href="#">Dnajb1.bSep08</a>	<a href="#">361384</a>	1829	726	2	242	DnaJ (Hsp40) homolog, subfamily B, member 1 (Dnajb1) alternative variant bSep08, mRNA.

<a href="#">Dnajb5</a>	<a href="#">Dnajb5.bSep08</a>	<a href="#">313811</a>	9219	2960	2	402	DnaJ (Hsp40) homolog, subfamily B, member 5 (Dnajb5) alternative variant bSep08, mRNA.
<a href="#">Dnajb5</a>	<a href="#">Dnajb5.cSep08</a>	<a href="#">313811</a>	5724	614	1	188	DnaJ (Hsp40) homolog, subfamily B, member 5 (Dnajb5) alternative variant cSep08, mRNA.
<a href="#">Dnajb6</a>	<a href="#">Dnajb6.aSep08</a>	<a href="#">362293</a>	59175	1371	3	364	DnaJ (Hsp40) homolog, subfamily B, member 6 (39.3 kD) (Dnajb6) alternative variant aSep08, mRNA.
<a href="#">Dnajb6</a>	<a href="#">Dnajb6.cSep08</a>	<a href="#">362293</a>	11526	985	1	91	DnaJ (Hsp40) homolog, subfamily B, member 6 (Dnajb6) alternative variant cSep08, mRNA.
<a href="#">Dnajb11</a>	<a href="#">Dnajb11.bSep08</a>	<a href="#">360734</a>	12694	809	5	262	DnaJ (Hsp40) homolog, subfamily B, member 11 (Dnajb11) alternative variant bSep08, mRNA.
<a href="#">Dnajb11</a>	<a href="#">Dnajb11.cSep08</a>	<a href="#">360734</a>	14749	819	8	260	DnaJ (Hsp40) homolog, subfamily B, member 11 (Dnajb11) alternative variant cSep08, mRNA.
<a href="#">Dnajb11</a>	<a href="#">Dnajb11.dSep08</a>	<a href="#">360734</a>	2436	317	3	105	DnaJ (Hsp40) homolog, subfamily B, member 11 (Dnajb11) alternative variant dSep08, mRNA.
<a href="#">Dnajb12</a>	<a href="#">Dnajb12.bSep08</a>	<a href="#">294513</a>	6699	1269	7	238	DnaJ homolog subfamily B member 12 (Dnajb12) alternative variant bSep08, mRNA.
<a href="#">Dnajb12</a>	<a href="#">Dnajb12.cSep08</a>	<a href="#">294513</a>	2085	535	3	161	DnaJ homolog subfamily B member 12 (Dnajb12) alternative variant cSep08, mRNA.
<a href="#">Dnajb12</a>	<a href="#">Dnajb12.dSep08</a>	<a href="#">294513</a>	4441	409	4	131	DnaJ homolog subfamily B member 12 (Dnajb12) alternative variant dSep08, mRNA.
<a href="#">Dnajb12</a>	<a href="#">Dnajb12.fSep08</a>	<a href="#">294513</a>	1949	811	2	114	putative protein (12.0 kD) (Dnajb12) alternative variant fSep08, mRNA.
<a href="#">Dnajb12.1</a>	<a href="#">Dnajb12.1.bSep08</a>	<a href="#">294513</a>	6699	1269	7	238	DnaJ homolog subfamily B member 12 (Dnajb12.1) alternative variant bSep08, mRNA.
<a href="#">Dnajb12.1</a>	<a href="#">Dnajb12.1.cSep08</a>	<a href="#">294513</a>	2085	535	3	161	DnaJ homolog subfamily B member 12 (Dnajb12.1) alternative variant cSep08, mRNA.
<a href="#">Dnajb12.1</a>	<a href="#">Dnajb12.1.dSep08</a>	<a href="#">294513</a>	4441	409	4	131	DnaJ homolog subfamily B member 12 (Dnajb12.1) alternative variant dSep08, mRNA.
<a href="#">Dnajb12.1</a>	<a href="#">Dnajb12.1.fSep08</a>	<a href="#">294513</a>	1949	811	2	114	putative protein (12.0 kD) (Dnajb12.1) alternative variant fSep08, mRNA.
<a href="#">Dnajb13</a>	<a href="#">Dnajb13.bSep08</a>	<a href="#">308857</a>	13190	649	3	167	DnaJ (Hsp40) related, subfamily B, member 13 (Dnajb13) alternative variant bSep08, mRNA.
<a href="#">Dnajb13</a>	<a href="#">Dnajb13.cSep08</a>	<a href="#">308857</a>	4322	528	2	127	DnaJ (Hsp40) related, subfamily B, member 13 (Dnajb13) alternative variant cSep08, mRNA.
<a href="#">Dnajb14</a>	<a href="#">Dnajb14.aSep08</a>	<a href="#">499716</a>	43277	1726	8	338	DnaJ (Hsp40) homolog, subfamily B, member 14 (37.9 kD) (Dnajb14) alternative variant aSep08, mRNA.
<a href="#">Dnajb14</a>	<a href="#">Dnajb14.bSep08</a>	<a href="#">499716</a>	24094	451	3	140	DnaJ (Hsp40) homolog, subfamily B, member 14 (Dnajb14) alternative variant bSep08, mRNA.
<a href="#">Dnajb14</a>	<a href="#">Dnajb14.cSep08</a>	<a href="#">499716</a>	15503	610	2	102	DnaJ (Hsp40) homolog, subfamily B, member 14 (Dnajb14) alternative variant cSep08, complete mRNA.
<a href="#">Dnajb14</a>	<a href="#">Dnajb14.dSep08</a>	<a href="#">499716</a>	16083	1189	3	102	DnaJ (Hsp40) homolog, subfamily B, member 14 (10.9 kD) (Dnajb14) alternative variant dSep08, mRNA.
<a href="#">Dnajc2</a>	<a href="#">Dnajc2.bSep08</a>	<a href="#">116456</a>	6197	1171	1	233	DnaJ (Hsp40) homolog, subfamily C, member 2 (Dnajc2) alternative variant bSep08, mRNA.
<a href="#">Dnajc4</a>	<a href="#">Dnajc4.bSep08</a>	<a href="#">361717</a>	2323	1236	4	188	DnaJ (Hsp40) homolog, subfamily C, member 4 (21.8 kD) (Dnajc4) alternative variant bSep08, mRNA.

<a href="#">Dnajc4</a>	<a href="#">Dnajc4.cSep08</a>	<a href="#">361717</a>	1553	457	4	122	DnaJ (Hsp40) homolog, subfamily C, member 4 (Dnajc4) alternative variant cSep08, mRNA.
<a href="#">Dnajc4</a>	<a href="#">Dnajc4.dSep08</a>	<a href="#">361717</a>	1826	892	3	77	DnaJ (Hsp40) homolog, subfamily C, member 4 (9.0 kD) (Dnajc4) alternative variant dSep08, mRNA.
<a href="#">Dnajc4</a>	<a href="#">Dnajc4.eSep08</a>	<a href="#">361717</a>	846	767	2	55	DnaJ (Hsp40) homolog, subfamily C, member 4 (6.4 kD) (Dnajc4) alternative variant eSep08, mRNA.
<a href="#">Dnajc5</a>	<a href="#">Dnajc5.bSep08</a>	<a href="#">79130</a>	34608	4879	2	198	DnaJ (Hsp40) homolog, subfamily C, member 5 (22.1 kD) (Dnajc5) alternative variant bSep08, mRNA.
<a href="#">Dnajc5</a>	<a href="#">Dnajc5.cSep08</a>	<a href="#">79130</a>	3413	1114	2	145	DnaJ (Hsp40) homolog, subfamily C, member 5 (Dnajc5) alternative variant cSep08, mRNA.
<a href="#">Dnajc5g</a>	<a href="#">Dnajc5g.bSep08</a>	<a href="#">366567</a>	3730	1073		164	DnaJ (Hsp40) homolog, subfamily C, member 5 gamma (18.4 kD) (Dnajc5g) alternative variant bSep08, mRNA.
<a href="#">Dnajc6</a>	<a href="#">Dnajc6.bSep08</a>	<a href="#">313409</a>	2044	518	3	172	DnaJ (Hsp40) homolog, subfamily C, member 6 (Dnajc6) alternative variant bSep08, mRNA.
<a href="#">Dnajc7</a>	<a href="#">Dnajc7.bSep08</a>	<a href="#">303536</a>	36847	2404	12	253	DnaJ (Hsp40) homolog, subfamily C, member 7 (28.7 kD) (Dnajc7) alternative variant bSep08, complete mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.bSep08</a>	<a href="#">313035</a>	16078	1001	7	216	DnaJ (Hsp40) homolog, subfamily C, member 8 (25.2 kD) (Dnajc8) alternative variant bSep08, mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.cSep08</a>	<a href="#">313035</a>	15737	603	6	196	DnaJ (Hsp40) homolog, subfamily C, member 8 (Dnajc8) alternative variant cSep08, mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.dSep08</a>	<a href="#">313035</a>	18858	739	8	184	DnaJ (Hsp40) homolog, subfamily C, member 8 (Dnajc8) alternative variant dSep08, mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.eSep08</a>	<a href="#">313035</a>	15696	640	7	141	DnaJ (Hsp40) homolog, subfamily C, member 8 (Dnajc8) alternative variant eSep08, mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.fSep08</a>	<a href="#">313035</a>	14636	1093	5	109	DnaJ (Hsp40) homolog, subfamily C, member 8 (12.2 kD) (Dnajc8) alternative variant fSep08, complete mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.gSep08</a>	<a href="#">313035</a>	6759	1074	2	47	DnaJ (Hsp40) homolog, subfamily C, member 8 (Dnajc8) alternative variant gSep08, mRNA.
<a href="#">Dnajc8</a>	<a href="#">Dnajc8.hSep08</a>	<a href="#">313035</a>	2651	716	2	30	DnaJ (Hsp40) homolog, subfamily C, member 8 (Dnajc8) alternative variant hSep08, mRNA.
<a href="#">Dnajc9</a>	<a href="#">Dnajc9.cSep08</a>	<a href="#">364240</a>	26918	363	3	35	DnaJ (Hsp40) homolog, subfamily C, member 9 (Dnajc9) alternative variant cSep08, mRNA.
<a href="#">Dnajc10</a>	<a href="#">Dnajc10.aSep08</a>	<a href="#">295690</a>	41041	4998	23	793	DnaJ homolog subfamily C member 10 (90.7 kD) (Dnajc10) alternative variant aSep08, mRNA.
<a href="#">Dnajc10</a>	<a href="#">Dnajc10.bSep08</a>	<a href="#">295690</a>	5375	1292	6	226	DnaJ homolog subfamily C member 10 (Dnajc10) alternative variant bSep08, mRNA.
<a href="#">Dnajc10</a>	<a href="#">Dnajc10.cSep08</a>	<a href="#">295690</a>	8004	1313	6	179	DnaJ homolog subfamily C member 10 (20.9 kD) (Dnajc10) alternative variant cSep08, mRNA.
<a href="#">Dnajc10</a>	<a href="#">Dnajc10.dSep08</a>	<a href="#">295690</a>	5097	373	2	90	DnaJ homolog subfamily C member 10 (Dnajc10) alternative variant dSep08, mRNA.
<a href="#">Dnajc11</a>	<a href="#">Dnajc11.bSep08</a>	<a href="#">362666</a>	3781	1664	2	122	DnaJ (Hsp40) homolog, subfamily C, member 11 (13.6 kD) (Dnajc11) alternative variant bSep08, mRNA.
<a href="#">Dnajc13</a>	<a href="#">Dnajc13.bSep08</a>	<a href="#">363127</a>	6467	707	7	235	DnaJ (Hsp40) homolog, subfamily C, member 13 (Dnajc13) alternative variant bSep08, mRNA.
<a href="#">Dnajc13</a>	<a href="#">Dnajc13.cSep08</a>	<a href="#">363127</a>	10703	1088	6	171	DnaJ (Hsp40) homolog, subfamily C, member 13 (Dnajc13) alternative variant cSep08, mRNA.

<a href="#">Dnajc13</a>	<a href="#">Dnajc13.dSep08</a>	<a href="#">363127</a>	2748	556	3	104	DnaJ (Hsp40) homolog, subfamily C, member 13 (Dnajc13) alternative variant dSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.bSep08</a>	<a href="#">114481</a>	6575	735	6	245	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant bSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.cSep08</a>	<a href="#">114481</a>	6735	701	6	233	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant cSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.dSep08</a>	<a href="#">114481</a>	6587	838	7	179	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant dSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.eSep08</a>	<a href="#">114481</a>	6288	741	5	172	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant eSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.fSep08</a>	<a href="#">114481</a>	6539	1057	6	168	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant fSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.gSep08</a>	<a href="#">114481</a>	6483	731	5	148	DnaJ (Hsp40) homolog, subfamily C, member 14 (Dnajc14) alternative variant gSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.hSep08</a>	<a href="#">114481</a>	3125	706	3	108	DnaJ (Hsp40) homolog, subfamily C, member 14 (12.6 kD) (Dnajc14) alternative variant hSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.iSep08</a>	<a href="#">114481</a>	6136	772	5	88	DnaJ (Hsp40) homolog, subfamily C, member 14 (10.3 kD) (Dnajc14) alternative variant iSep08, mRNA.
<a href="#">Dnajc14</a>	<a href="#">Dnajc14.jSep08</a>	<a href="#">114481</a>	6095	727	5	57	DnaJ (Hsp40) homolog, subfamily C, member 14 (6.5 kD) (Dnajc14) alternative variant jSep08, mRNA.
<a href="#">Dnajc15</a>	<a href="#">Dnajc15.bSep08</a>	<a href="#">290370</a>	66369	390	2	129	DnaJ (Hsp40) homolog, subfamily C, member 15 (Dnajc15) alternative variant bSep08, mRNA.
<a href="#">Dnajc15</a>	<a href="#">Dnajc15.cSep08</a>	<a href="#">290370</a>	24444	386	1	40	DnaJ (Hsp40) homolog, subfamily C, member 15 (Dnajc15) alternative variant cSep08, mRNA.
<a href="#">Dnajc17</a>	<a href="#">Dnajc17.aSep08</a>	<a href="#">311329</a>	34465	980	10	304	DnaJ (Hsp40) homolog, subfamily C, member 17 (Dnajc17) alternative variant aSep08, mRNA.
<a href="#">Dnajc17</a>	<a href="#">Dnajc17.bSep08</a>	<a href="#">311329</a>	3091	363	1	33	DnaJ (Hsp40) homolog, subfamily C, member 17 (Dnajc17) alternative variant bSep08, mRNA.
<a href="#">Dnajc18</a>	<a href="#">Dnajc18.aSep08</a>	<a href="#">291677</a>	23154	1207	1	339	DnaJ (Hsp40) homolog, subfamily C, member 18 (39.0 kD) (Dnajc18) alternative variant aSep08, mRNA.
<a href="#">Dnalc1</a>	<a href="#">Dnalc1.bSep08</a>	<a href="#">685664</a>	21650	2857	2	77	dynein, axonemal, light chain 1 (8.9 kD) (Dnalc1) alternative variant bSep08, mRNA.
<a href="#">Dnase1</a>	<a href="#">Dnase1.bSep08</a>	<a href="#">25633</a>	2066	745	8	200	deoxyribonuclease I (Dnase1) alternative variant bSep08, mRNA.
<a href="#">Dnase111</a>	<a href="#">Dnase111.bSep08</a>	<a href="#">363522</a>	3248	1230	6	262	deoxyribonuclease 1-like 1 (Dnase111) alternative variant bSep08, mRNA.
<a href="#">Dnase111</a>	<a href="#">Dnase111.cSep08</a>	<a href="#">363522</a>	7830	868	6	237	deoxyribonuclease 1-like 1 (Dnase111) alternative variant cSep08, mRNA.
<a href="#">Dnase111</a>	<a href="#">Dnase111.dSep08</a>	<a href="#">363522</a>	8482	1379	8	197	deoxyribonuclease 1-like 1 (22.4 kD) (Dnase111) alternative variant dSep08, complete mRNA.
<a href="#">Dnase112</a>	<a href="#">Dnase112.aSep08</a>	<a href="#">681124</a>	1439	1053		226	deoxyribonuclease 1-like 2 (Dnase112) mRNA.
<a href="#">Dnase2b</a>	<a href="#">Dnase2b.bSep08</a>	<a href="#">59296</a>	21177	447	2	81	putative protein (Dnase2b) alternative variant bSep08, mRNA.
<a href="#">Dnase2b</a>	<a href="#">Dnase2b.cSep08</a>	<a href="#">59296</a>	5961	436	2	68	putative protein (Dnase2b) alternative variant cSep08, mRNA.
<a href="#">DNA_pol_E_B.0</a>	<a href="#">DNA_pol_E_B.0.aSep08</a>		5610	583	6	163	polymerase epsilon (DNA_pol_E_B.0) alternative variant aSep08, mRNA.

<a href="#">DNA_pol_E_B.0</a>	<a href="#">DNA_pol_E_B.0.bSep08</a>		4251	648	4	105	polymerase epsilon (11.9 kD) (DNA_pol_E_B.0) alternative variant bSep08, mRNA.
<a href="#">DNA_pol_E_B.0</a>	<a href="#">DNA_pol_E_B.0.cSep08</a>		2276	914	2	93	polymerase epsilon (10.7 kD) (DNA_pol_E_B.0) alternative variant cSep08, mRNA.
<a href="#">DNA_RNApol_7kD.0</a>	<a href="#">DNA_RNApol_7kD.0.aSep08</a>		3371	528	4	58	putative protein (DNA_RNApol_7kD.0) alternative variant aSep08, mRNA.
<a href="#">DNA_RNApol_7kD.0</a>	<a href="#">DNA_RNApol_7kD.0.cSep08</a>		1199	288	3	42	polymerase II (DNA_RNApol_7kD.0) alternative variant cSep08, mRNA.
<a href="#">Dnd1</a>	<a href="#">Dnd1.aSep08</a>	<a href="#">307492</a>	7038	1599	10	404	dead end homolog 1 (zebrafish) (Dnd1) alternative variant aSep08, mRNA.
<a href="#">Dnd1</a>	<a href="#">Dnd1.cSep08</a>	<a href="#">307492</a>	822	618	2	102	dead end homolog 1 (zebrafish) (Dnd1) alternative variant cSep08, mRNA.
<a href="#">Dnd1</a>	<a href="#">Dnd1.dSep08</a>	<a href="#">307492</a>	54119	296	3	98	dead end homolog 1 (zebrafish) (Dnd1) alternative variant dSep08, mRNA.
<a href="#">Dnd1</a>	<a href="#">Dnd1.eSep08</a>	<a href="#">307492</a>	8872	845	3	53	dead end homolog 1 (zebrafish) (5.8 kD) (Dnd1) alternative variant eSep08, complete mRNA.
<a href="#">Dner</a>	<a href="#">Dner.aSep08</a>	<a href="#">316573</a>	105476	2129	6	371	delta/notch-like EGF-related receptor (40.6 kD) (Dner) alternative variant aSep08, mRNA.
<a href="#">Dner</a>	<a href="#">Dner.bSep08</a>	<a href="#">316573</a>	30451	564	1	120	delta/notch-like EGF-related receptor (Dner) alternative variant bSep08, mRNA.
<a href="#">Dnhd1</a>	<a href="#">Dnhd1.aSep08</a>	<a href="#">690115</a>	925	730		242	dynein heavy chain domain 1 (Dnhd1) mRNA.
<a href="#">Dnm1</a>	<a href="#">Dnm1.bSep08</a>	<a href="#">140694</a>	24023	1285	14	427	dynamin 1 (Dnm1) alternative variant bSep08, mRNA.
<a href="#">Dnm1</a>	<a href="#">Dnm1.cSep08</a>	<a href="#">140694</a>	6626	1127	4	199	dynamin 1 (Dnm1) alternative variant cSep08, mRNA.
<a href="#">Dnm1</a>	<a href="#">Dnm1.dSep08</a>	<a href="#">140694</a>	9080	526	4	146	dynamin 1 (Dnm1) alternative variant dSep08, mRNA.
<a href="#">Dnm1</a>	<a href="#">Dnm1.eSep08</a>	<a href="#">140694</a>	11431	1040	5	117	dynamin 1 (Dnm1) alternative variant eSep08, mRNA.
<a href="#">Dnm1l</a>	<a href="#">Dnm1l.bSep08</a>	<a href="#">114114</a>	18960	3827	10	391	dynamin 1-like (Dnm1l) alternative variant bSep08, mRNA.
<a href="#">Dnm1l</a>	<a href="#">Dnm1l.cSep08</a>	<a href="#">114114</a>	23001	647	6	215	dynamin 1-like (Dnm1l) alternative variant cSep08, mRNA.
<a href="#">Dnm1l</a>	<a href="#">Dnm1l.dSep08</a>	<a href="#">114114</a>	11455	630	7	209	dynamin 1-like (Dnm1l) alternative variant dSep08, mRNA.
<a href="#">Dnm1l</a>	<a href="#">Dnm1l.eSep08</a>	<a href="#">114114</a>	930	646	2	66	dynamin 1-like (7.6 kD) (Dnm1l) alternative variant eSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.aSep08</a>	<a href="#">25751</a>	83642	3470	20	909	dynamin 2 (Dnm2) alternative variant aSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.bSep08</a>	<a href="#">25751</a>	28094	1645	12	430	dynamin 2 (Dnm2) alternative variant bSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.cSep08</a>	<a href="#">25751</a>	15543	526	7	175	dynamin 2 (Dnm2) alternative variant cSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.dSep08</a>	<a href="#">25751</a>	3238	800	3	159	dynamin 2 (Dnm2) alternative variant dSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.eSep08</a>	<a href="#">25751</a>	1616	714	2	126	dynamin 2 (Dnm2) alternative variant eSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.fSep08</a>	<a href="#">25751</a>	43959	373	3	124	dynamin 2 (Dnm2) alternative variant fSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.hSep08</a>	<a href="#">25751</a>	13918	530	5	117	dynamin 2 (Dnm2) alternative variant hSep08, mRNA.
<a href="#">Dnm2</a>	<a href="#">Dnm2.iSep08</a>	<a href="#">25751</a>	4022	433	2	36	dynamin 2 (Dnm2) alternative variant iSep08, mRNA.
<a href="#">Dnm3</a>	<a href="#">Dnm3.bSep08</a>	<a href="#">171574</a>	464697	2951	20	894	dynamin 3 (Dnm3) alternative variant bSep08, mRNA.
<a href="#">Dnm3</a>	<a href="#">Dnm3.cSep08</a>	<a href="#">171574</a>	479861	1748	4	430	dynamin 3 (Dnm3) alternative variant cSep08, mRNA.
<a href="#">Dnm3</a>	<a href="#">Dnm3.eSep08</a>	<a href="#">171574</a>	34883	468	5	145	dynamin 3 (Dnm3) alternative variant eSep08, mRNA.
<a href="#">Dnmbp</a>	<a href="#">Dnmbp.aSep08</a>	<a href="#">309362</a>	32442	747	4	249	dynamin binding protein (Dnmbp) alternative variant aSep08, mRNA.



<a href="#">Dnmt1</a>	<a href="#">Dnmt1.bSep08</a>	<a href="#">84350</a>	2576	1265	4	319	DNA methyltransferase (cytosine-5) 1 (Dnmt1) alternative variant bSep08, mRNA.
<a href="#">Dnmt1</a>	<a href="#">Dnmt1.cSep08</a>	<a href="#">84350</a>	5664	939	5	313	DNA methyltransferase (cytosine-5) 1 (Dnmt1) alternative variant cSep08, mRNA.
<a href="#">Dnmt1</a>	<a href="#">Dnmt1.dSep08</a>	<a href="#">84350</a>	3814	988	4	169	DNA methyltransferase (cytosine-5) 1 (Dnmt1) alternative variant dSep08, mRNA.
<a href="#">Dnmt1</a>	<a href="#">Dnmt1.eSep08</a>	<a href="#">84350</a>	1009	732	1	144	DNA methyltransferase (cytosine-5) 1 (Dnmt1) alternative variant eSep08, mRNA.
<a href="#">Dnmt3a</a>	<a href="#">Dnmt3a.dSep08</a>	<a href="#">444984</a>	15183	639	3	69	DNA methyltransferase 3A (7.8 kD) (Dnmt3a) alternative variant dSep08, mRNA.
<a href="#">Dnmt3a</a>	<a href="#">Dnmt3a.eSep08</a>	<a href="#">444984</a>	1568	398	2	44	DNA methyltransferase 3A (4.6 kD) (Dnmt3a) alternative variant eSep08, mRNA.
<a href="#">Dnmt3b</a>	<a href="#">Dnmt3b.cSep08</a>	<a href="#">444985</a>	7135	418	2	72	DNA methyltransferase 3B (Dnmt3b) alternative variant cSep08, mRNA.
<a href="#">Dnpep</a>	<a href="#">Dnpep.bSep08</a>	<a href="#">301529</a>	1798	752	3	219	aspartyl aminopeptidase CRA a (Dnpep) alternative variant bSep08, mRNA.
<a href="#">Dnpep</a>	<a href="#">Dnpep.cSep08</a>	<a href="#">301529</a>	2110	928	2	190	aspartyl aminopeptidase CRA e (Dnpep) alternative variant cSep08, mRNA.
<a href="#">Dnpep</a>	<a href="#">Dnpep.dSep08</a>	<a href="#">301529</a>	4084	1264	4	177	aspartyl aminopeptidase (Dnpep) alternative variant dSep08, mRNA.
<a href="#">Dnpep</a>	<a href="#">Dnpep.eSep08</a>	<a href="#">301529</a>	1276	750	2	150	aspartyl aminopeptidase CRA e (16.7 kD) (Dnpep) alternative variant eSep08, mRNA.
<a href="#">Dnttip1</a>	<a href="#">Dnttip1.bSep08</a>	<a href="#">171437</a>	3079	886	2	52	deoxynucleotidyltransferase, terminal, interacting protein 1 (5.7 kD) (Dnttip1) alternative variant bSep08, mRNA.
<a href="#">doby</a>	<a href="#">doby.aSep08</a>		48942	441		146	dedicator of cytokinesis 11 (doby) mRNA.
<a href="#">Doc2a</a>	<a href="#">Doc2a.bSep08</a>	<a href="#">65031</a>	1352	306	2	101	double C2, alpha (Doc2a) alternative variant bSep08, mRNA.
<a href="#">Doc2g</a>	<a href="#">Doc2g.bSep08</a>	<a href="#">293654</a>	2266	555	1	92	double C2, gamma (Doc2g) alternative variant bSep08, mRNA.
<a href="#">dochy</a>	<a href="#">dochy.aSep08</a>	<a href="#">499795</a>	133010	550	1	171	rho GTPase activating protein 15 (dochy) alternative variant aSep08, mRNA.
<a href="#">dochy</a>	<a href="#">dochy.bSep08</a>	<a href="#">499795</a>	132965	436		145	rho GTPase activating protein 15 (dochy) alternative variant bSep08, mRNA.
<a href="#">Dock1</a>	<a href="#">Dock1.aSep08</a>	<a href="#">309081</a>	33729	2653	2	408	dedicator of cyto-kinesis 1 (45.6 kD) (Dock1) alternative variant aSep08, mRNA.
<a href="#">Dock2</a>	<a href="#">Dock2.aSep08</a>	<a href="#">360509</a>	44778	2007		668	dedicator of cytokinesis 2 (Dock2) mRNA.
<a href="#">Dock3</a>	<a href="#">Dock3.bSep08</a>	<a href="#">315992</a>	10092	497	5	165	dedicator of cyto-kinesis 3 (Dock3) alternative variant bSep08, mRNA.
<a href="#">Dock5</a>	<a href="#">Dock5.bSep08</a>	<a href="#">305987</a>	12633	833	3	277	dedicator of cytokinesis 5 (Dock5) alternative variant bSep08, mRNA.
<a href="#">Dock6</a>	<a href="#">Dock6.aSep08</a>	<a href="#">367039</a>	38966	1785		528	dedicator of cytokinesis 6 (Dock6) alternative variant aSep08, mRNA.
<a href="#">Dock6</a>	<a href="#">Dock6.bSep08</a>	<a href="#">367039</a>	2713	1186		139	dedicator of cytokinesis 6 (16.1 kD) (Dock6) alternative variant bSep08, mRNA.
<a href="#">Dock7</a>	<a href="#">Dock7.aSep08</a>	<a href="#">313388</a>	37278	2407	16	802	dedicator of cytokinesis 7 (Dock7) alternative variant aSep08, mRNA.

<a href="#">Dock7</a>	<a href="#">Dock7.bSep08</a>	<a href="#">313388</a>	15994	1510	7	303	dedicator of cytokinesis 7 (35.6 kD) (Dock7) alternative variant bSep08, complete mRNA.
<a href="#">Dock7</a>	<a href="#">Dock7.cSep08</a>	<a href="#">313388</a>	22339	1116	8	301	dedicator of cytokinesis 7 (34.0 kD) (Dock7) alternative variant cSep08, mRNA.
<a href="#">Dock7</a>	<a href="#">Dock7.dSep08</a>	<a href="#">313388</a>	8835	587	4	195	dedicator of cytokinesis 7 (Dock7) alternative variant dSep08, mRNA.
<a href="#">Dock7.1</a>	<a href="#">Dock7.1.aSep08</a>	<a href="#">313388</a>	60964	1437		479	dedicator of cytokinesis 7 (Dock7.1) mRNA.
<a href="#">Dock8</a>	<a href="#">Dock8.bSep08</a>	<a href="#">499337</a>	11728	734	6	244	dedicator of cytokinesis 8 (Dock8) alternative variant bSep08, mRNA.
<a href="#">Dock8</a>	<a href="#">Dock8.cSep08</a>	<a href="#">499337</a>	26735	1001	8	205	dedicator of cytokinesis 8 CRA a (23.4 kD) (Dock8) alternative variant cSep08, mRNA.
<a href="#">Dock8</a>	<a href="#">Dock8.dSep08</a>	<a href="#">499337</a>	8656	1880	4	148	dedicator of cytokinesis 8 (17.3 kD) (Dock8) alternative variant dSep08, mRNA.
<a href="#">Dock8</a>	<a href="#">Dock8.fSep08</a>	<a href="#">499337</a>	975	264	2	35	dedicator of cytokinesis 8 (Dock8) alternative variant fSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.bSep08</a>	<a href="#">259237</a>	20566	889	10	296	dedicator of cytokinesis 9 (Dock9) alternative variant bSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.cSep08</a>	<a href="#">259237</a>	22573	823	9	274	dedicator of cytokinesis 9 (Dock9) alternative variant cSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.dSep08</a>	<a href="#">259237</a>	17137	576	7	192	dedicator of cytokinesis 9 CRA i (Dock9) alternative variant dSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.eSep08</a>	<a href="#">259237</a>	4807	785	4	121	dedicator of cytokinesis 9 (13.6 kD) (Dock9) alternative variant eSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.fSep08</a>	<a href="#">259237</a>	2812	447	2	88	dedicator of cytokinesis 9 (Dock9) alternative variant fSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.gSep08</a>	<a href="#">259237</a>	5447	351	2	86	dedicator of cytokinesis 9 (Dock9) alternative variant gSep08, mRNA.
<a href="#">Dock9</a>	<a href="#">Dock9.hSep08</a>	<a href="#">259237</a>	4081	382	3	63	dedicator of cytokinesis 9 (Dock9) alternative variant hSep08, mRNA.
<a href="#">Dock11</a>	<a href="#">Dock11.aSep08</a>	<a href="#">313438</a>	25714	1291		430	dedicator of cytokinesis 11 (Dock11) mRNA.
<a href="#">dofer</a>	<a href="#">dofer.aSep08</a>		96944	823		72	putative cytoplasmic protein (8.1 kD) (dofer) mRNA.
<a href="#">doflo</a>	<a href="#">doflo.aSep08</a>		3385	458		152	vacuolar protein sorting 13A (doflo) mRNA.
<a href="#">doflu</a>	<a href="#">doflu.aSep08</a>		686	383		127	myeloid lymphoid leukemia like (doflu) mRNA.
<a href="#">Dohh</a>	<a href="#">Dohh.bSep08</a>	<a href="#">314644</a>	3743	700	3	185	deoxyhypusine hydroxylase/monooxygenase (Dohh) alternative variant bSep08, mRNA.
<a href="#">Dohh</a>	<a href="#">Dohh.cSep08</a>	<a href="#">314644</a>	4771	538	4	178	deoxyhypusine hydroxylase/monooxygenase (Dohh) alternative variant cSep08, mRNA.
<a href="#">Dohh</a>	<a href="#">Dohh.dSep08</a>	<a href="#">314644</a>	2872	618	3	108	deoxyhypusine hydroxylase/monooxygenase (Dohh) alternative variant dSep08, mRNA.
<a href="#">Dok4</a>	<a href="#">Dok4.bSep08</a>	<a href="#">361364</a>	3571	386	2	58	docking protein 4 (Dok4) alternative variant bSep08, mRNA.
<a href="#">dokee</a>	<a href="#">dokee.aSep08</a>		766	505		33	putative protein (3.8 kD) (dokee) mRNA.
<a href="#">doloy</a>	<a href="#">doloy.aSep08</a>		4167	377		125	specific 7 (doloy) mRNA.
<a href="#">Dolpp1</a>	<a href="#">Dolpp1.aSep08</a>	<a href="#">296624</a>	8486	2276	8	238	dolichyl pyrophosphate phosphatase 1 (27.1 kD) (Dolpp1) alternative variant aSep08, mRNA.

<a href="#">Dolpp1</a>	<a href="#">Dolpp1.dSep08</a>	<a href="#">296624</a>	2692	333	2	60	dolichyl pyrophosphate phosphatase 1 (Dolpp1) alternative variant dSep08, mRNA.
<a href="#">Dom3z</a>	<a href="#">Dom3z.bSep08</a>	<a href="#">361799</a>	2091	1455	6	392	RAI1 like (Dom3z) alternative variant bSep08, mRNA.
<a href="#">domer</a>	<a href="#">domer.aSep08</a>		810	431		66	putative protein (domer) mRNA.
<a href="#">donoy</a>	<a href="#">donoy.aSep08</a>		415	317		50	putative protein (5.3 kD) (donoy) mRNA.
<a href="#">Donson</a>	<a href="#">Donson.aSep08</a>	<a href="#">288257</a>	9651	2385	9	546	downstream neighbor of SON (60.5 kD) (Donson) alternative variant aSep08, mRNA.
<a href="#">Donson</a>	<a href="#">Donson.cSep08</a>	<a href="#">288257</a>	1989	758	2	42	downstream neighbor of SON (Donson) alternative variant cSep08, mRNA.
<a href="#">Dopey2</a>	<a href="#">Dopey2.aSep08</a>	<a href="#">304077</a>	39015	1931	13	490	dopey family member 2 (55.7 kD) (Dopey2) alternative variant aSep08, mRNA.
<a href="#">Dopey_N.0</a>	<a href="#">Dopey_N.0.aSep08</a>		37100	989		329	dopey family member 2 CRA b (Dopey_N.0) mRNA.
<a href="#">Dopey_N.1</a>	<a href="#">Dopey_N.1.aSep08</a>		36380	603		140	dopey family member 1 (Dopey_N.1) mRNA.
<a href="#">dopor</a>	<a href="#">dopor.aSep08</a>		1307	689		229	centrosomal protein 164kDa CRA d (dopor) mRNA.
<a href="#">dorby</a>	<a href="#">dorby.aSep08</a>		6240	818		272	putative protein of vertebrate origin (dorby) mRNA.
<a href="#">dorchy</a>	<a href="#">dorchy.aSep08</a>		5455	630		209	methyl-CpG binding domain protein 5 like (dorchy) mRNA.
<a href="#">dorfer</a>	<a href="#">dorfer.aSep08</a>		1768	298		76	putative protein (dorfer) mRNA.
<a href="#">dorflo</a>	<a href="#">dorflo.aSep08</a>		9091	489		163	transient receptor potential cation channel subfamily M member 6 (dorflo) mRNA.
<a href="#">dorflu</a>	<a href="#">dorflu.aSep08</a>		547	374		43	putative protein (dorflu) mRNA.
<a href="#">dorkee</a>	<a href="#">dorkee.bSep08</a>		523	266	2	38	putative protein (dorkee) alternative variant bSep08, mRNA.
<a href="#">dorloy</a>	<a href="#">dorloy.aSep08</a>		1053	623		81	CRA a like (9.4 kD) (dorloy) mRNA.
<a href="#">dormer</a>	<a href="#">dormer.aSep08</a>		1578	290		69	keratin 2 (dormer) mRNA.
<a href="#">dornoy</a>	<a href="#">dornoy.aSep08</a>		4715	1366		91	CRA a like (dornoy) mRNA.
<a href="#">dorpor</a>	<a href="#">dorpor.aSep08</a>		1049	626	2	165	putative protein (dorpor) alternative variant aSep08, mRNA.
<a href="#">dorsa</a>	<a href="#">dorsa.aSep08</a>		2515	452		46	putative protein (5.0 kD) (dorsa) mRNA.
<a href="#">dorshee</a>	<a href="#">dorshee.aSep08</a>		649	401		41	putative protein (4.4 kD) (dorshee) mRNA.
<a href="#">dorto</a>	<a href="#">dorto.aSep08</a>		3689	780		70	atp-dependent rna helicase tdrd9 (dorto) mRNA.
<a href="#">dorvar</a>	<a href="#">dorvar.aSep08</a>		4388	705	2	165	zyg-11 homolog B (dorvar) alternative variant aSep08, mRNA.
<a href="#">dorvar</a>	<a href="#">dorvar.bSep08</a>		4269	710	1	121	zyg-11 homolog B (14.0 kD) (dorvar) alternative variant bSep08, mRNA.
<a href="#">dorwey</a>	<a href="#">dorwey.aSep08</a>		1149	551	2	69	putative protein (7.3 kD) (dorwey) alternative variant aSep08, mRNA.
<a href="#">Dos</a>	<a href="#">Dos.bSep08</a>	<a href="#">314622</a>	1947	1349	4	179	putative protein of vertebrate origin (Dos) alternative variant bSep08, mRNA.
<a href="#">Dos</a>	<a href="#">Dos.eSep08</a>	<a href="#">314622</a>	1974	296	2	98	putative protein (Dos) alternative variant eSep08, mRNA.
<a href="#">Dos</a>	<a href="#">Dos.fSep08</a>	<a href="#">314622</a>	1449	399	5	92	putative protein (Dos) alternative variant fSep08, mRNA.
<a href="#">dosa</a>	<a href="#">dosa.aSep08</a>		1614	1366	3	74	putative protein (dosa) alternative variant aSep08, mRNA.
<a href="#">dosa</a>	<a href="#">dosa.bSep08</a>		20327	715	3	111	putative protein (dosa) alternative variant bSep08, mRNA.
<a href="#">doshee</a>	<a href="#">doshee.aSep08</a>		2058	632		67	putative protein (7.6 kD) (doshee) mRNA.

<a href="#">Dot1l</a>	<a href="#">Dot1l.bSep08</a>	<a href="#">362831</a>	26817	737	6	240	DOT1-like, histone H3 methyltransferase ( <i>S. cerevisiae</i> ) (Dot1l) alternative variant bSep08, mRNA.
<a href="#">Dot1l</a>	<a href="#">Dot1l.dSep08</a>	<a href="#">362831</a>	13729	442	5	141	DOT1-like, histone H3 methyltransferase ( <i>S. cerevisiae</i> ) (Dot1l) alternative variant dSep08, mRNA.
<a href="#">doto</a>	<a href="#">doto.aSep08</a>		3312	738		178	putative protein of mammalian origin (doto) mRNA.
<a href="#">dovar</a>	<a href="#">dovar.aSep08</a>		460	362		51	putative protein (5.6 kD) (dovar) mRNA.
<a href="#">dowey</a>	<a href="#">dowey.aSep08</a>		3897	753		87	putative mitochondrial protein (9.7 kD) (dowey) mRNA.
<a href="#">doyby</a>	<a href="#">doyby.aSep08</a>		3398	353		73	polyprotein (doyby) mRNA.
<a href="#">doychy</a>	<a href="#">doychy.aSep08</a>		10065	641		40	putative protein (4.7 kD) (doychy) mRNA.
<a href="#">doyfer</a>	<a href="#">doyfer.aSep08</a>		1802	725		60	putative protein (6.8 kD) (doyfer) mRNA.
<a href="#">doyflo</a>	<a href="#">doyflo.aSep08</a>		3911	472	3	47	putative protein (5.2 kD) (doyflo) alternative variant aSep08, mRNA.
<a href="#">doyflu</a>	<a href="#">doyflu.aSep08</a>		26811	476		63	putative protein (6.6 kD) (doyflu) mRNA.
<a href="#">doykee</a>	<a href="#">doykee.aSep08</a>		119126	1240		117	CLIP-associating protein 1 (13.2 kD) (doykee) mRNA.
<a href="#">doyloy</a>	<a href="#">doyloy.aSep08</a>		5401	1191	2	114	CRA a like (13.7 kD) (doyloy) alternative variant aSep08, mRNA.
<a href="#">doyloy</a>	<a href="#">doyloy.cSep08</a>		4102	386	2	39	putative protein (4.3 kD) (doyloy) alternative variant cSep08, mRNA.
<a href="#">doymer</a>	<a href="#">doymer.aSep08</a>		3453	661		150	putative protein (16.8 kD) (doymer) mRNA.
<a href="#">doynoy</a>	<a href="#">doynoy.aSep08</a>		1506	360		63	putative protein (doynoy) mRNA.
<a href="#">doypor</a>	<a href="#">doypor.aSep08</a>		5726	844	2	70	ab2-143 like (doypor) alternative variant aSep08, mRNA.
<a href="#">doysa</a>	<a href="#">doysa.aSep08</a>		4729	723		69	putative protein (7.6 kD) (doysa) mRNA.
<a href="#">doyshee</a>	<a href="#">doyshee.aSep08</a>		34088	588		196	catenin delta 2 (doyshee) mRNA.
<a href="#">doyto</a>	<a href="#">doyto.bSep08</a>		22506	329	4	69	putative protein (7.6 kD) (doyto) alternative variant bSep08, mRNA.
<a href="#">doyvar</a>	<a href="#">doyvar.aSep08</a>		1303	627		53	putative protein (doyvar) mRNA.
<a href="#">doywey</a>	<a href="#">doywey.aSep08</a>		4262	610		93	putative protein (doywey) mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.bSep08</a>	<a href="#">300668</a>	6651	2393	8	301	transferase (34.5 kD) (Dpagt1) alternative variant bSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.cSep08</a>	<a href="#">300668</a>	4915	953	5	198	transferase (Dpagt1) alternative variant cSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.dSep08</a>	<a href="#">300668</a>	1730	560	4	152	transferase (16.9 kD) (Dpagt1) alternative variant dSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.eSep08</a>	<a href="#">300668</a>	7083	701	4	119	transferase (Dpagt1) alternative variant eSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.fSep08</a>	<a href="#">300668</a>	2974	774	4	118	transferase (13.0 kD) (Dpagt1) alternative variant fSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.gSep08</a>	<a href="#">300668</a>	1947	739	3	93	transferase (Dpagt1) alternative variant gSep08, mRNA.
<a href="#">Dpagt1</a>	<a href="#">Dpagt1.hSep08</a>	<a href="#">300668</a>	648	524	2	63	putative protein (Dpagt1) alternative variant hSep08, mRNA.
<a href="#">Dpep1</a>	<a href="#">Dpep1.bSep08</a>	<a href="#">94199</a>	23477	720	7	240	dipeptidase 1 (renal) (Dpep1) alternative variant bSep08, mRNA.
<a href="#">Dpep2</a>	<a href="#">Dpep2.bSep08</a>	<a href="#">291984</a>	6661	1633	8	310	dipeptidase 2 (34.2 kD) (Dpep2) alternative variant bSep08, mRNA.
<a href="#">Dpep2</a>	<a href="#">Dpep2.cSep08</a>	<a href="#">291984</a>	2683	1107	4	251	dipeptidase 2 (Dpep2) alternative variant cSep08, mRNA.

<a href="#">Dpf1</a>	<a href="#">Dpf1.bSep08</a>	<a href="#">50545</a>	3838	630	6	189	d4 zinc double phd fingers family 1 CRA a (Dpf1) alternative variant bSep08, mRNA.
<a href="#">Dpf1</a>	<a href="#">Dpf1.cSep08</a>	<a href="#">50545</a>	4794	561	5	186	d4 zinc double phd fingers family 1 CRA c (Dpf1) alternative variant cSep08, mRNA.
<a href="#">Dpf1</a>	<a href="#">Dpf1.dSep08</a>	<a href="#">50545</a>	3349	1616	4	77	d4 zinc double PHD fingers family 1 (8.8 kD) (Dpf1) alternative variant dSep08, mRNA.
<a href="#">Dpf2</a>	<a href="#">Dpf2.bSep08</a>	<a href="#">361711</a>	14006	2552	7	384	d4, zinc and double PHD fingers family 2 (43.6 kD) (Dpf2) alternative variant bSep08, mRNA.
<a href="#">Dpf2</a>	<a href="#">Dpf2.cSep08</a>	<a href="#">361711</a>	4918	943	6	313	d4, zinc and double PHD fingers family 2 (Dpf2) alternative variant cSep08, mRNA.
<a href="#">Dpf2</a>	<a href="#">Dpf2.dSep08</a>	<a href="#">361711</a>	4336	751	2	143	d4, zinc and double PHD fingers family 2 (Dpf2) alternative variant dSep08, mRNA.
<a href="#">Dpf2</a>	<a href="#">Dpf2.eSep08</a>	<a href="#">361711</a>	2422	813	3	136	d4, zinc and double PHD fingers family 2 (Dpf2) alternative variant eSep08, mRNA.
<a href="#">Dpf3</a>	<a href="#">Dpf3.aSep08</a>	<a href="#">299186</a>	224699	1677	1	367	d4, zinc and double PHD fingers, family 3 (Dpf3) alternative variant aSep08, mRNA.
<a href="#">Dpf3</a>	<a href="#">Dpf3.bSep08</a>	<a href="#">299186</a>	59722	519	1	172	d4, zinc and double PHD fingers, family 3 (Dpf3) alternative variant bSep08, mRNA.
<a href="#">Dph1andOvca2</a>	<a href="#">Dph1andOvca2.cSep08</a>	<a href="#">287523</a>	8306	770	5	253	diphtheria toxin resistance protein required for diphthamide biosynthesis-like 1 (Dph1andOvca2) alternative variant cSep08, mRNA.
<a href="#">Dph1andOvca2</a>	<a href="#">Dph1andOvca2.cSep08</a>	<a href="#">497954</a>	8306	770	5	253	diphtheria toxin resistance protein required for diphthamide biosynthesis-like 1 (Dph1andOvca2) alternative variant cSep08, mRNA.
<a href="#">Dph1andOvca2</a>	<a href="#">Dph1andOvca2.eSep08</a>	<a href="#">287523</a>	2505	800	6	139	tumor suppressor like (Dph1andOvca2) alternative variant eSep08, mRNA.
<a href="#">Dph1andOvca2</a>	<a href="#">Dph1andOvca2.eSep08</a>	<a href="#">497954</a>	2505	800	6	139	tumor suppressor like (Dph1andOvca2) alternative variant eSep08, mRNA.
<a href="#">Dph3</a>	<a href="#">Dph3.bSep08</a>	<a href="#">680594</a>	2832	2425	2	73	DPH3, KTI11 homolog (S. cerevisiae) (8.2 kD) (Dph3) alternative variant bSep08, mRNA.
<a href="#">Dph3</a>	<a href="#">Dph3.cSep08</a>	<a href="#">680594</a>	2792	542	2	57	DPH3, KTI11 homolog (S. cerevisiae) (6.5 kD) (Dph3) alternative variant cSep08, mRNA.
<a href="#">Dph3</a>	<a href="#">Dph3.dSep08</a>	<a href="#">680594</a>	2524	756	2	54	DPH3, KTI11 homolog (S. cerevisiae) (Dph3) alternative variant dSep08, mRNA.
<a href="#">Dph4</a>	<a href="#">Dph4.aSep08</a>	<a href="#">362184</a>	34328	534	3	128	DPH4 homolog (JJJ3, S. cerevisiae) (Dph4) alternative variant aSep08, mRNA.
<a href="#">Dph4</a>	<a href="#">Dph4.bSep08</a>	<a href="#">362184</a>	14937	614	3	72	DPH4 homolog (JJJ3, S. cerevisiae) (Dph4) alternative variant bSep08, mRNA.
<a href="#">Dph4</a>	<a href="#">Dph4.cSep08</a>	<a href="#">362184</a>	35372	773	3	48	DPH4 homolog (JJJ3, S. cerevisiae) (5.4 kD) (Dph4) alternative variant cSep08, mRNA.
<a href="#">Dph5</a>	<a href="#">Dph5.aSep08</a>	<a href="#">295394</a>	33460	1776	2	468	DPH5 homolog (S. cerevisiae) (Dph5) alternative variant aSep08, mRNA.
<a href="#">Dph5</a>	<a href="#">Dph5.cSep08</a>	<a href="#">295394</a>	31099	729	1	211	DPH5 homolog (S. cerevisiae) (Dph5) alternative variant cSep08, mRNA.
<a href="#">Dpm1</a>	<a href="#">Dpm1.bSep08</a>	<a href="#">296394</a>	6082	400	5	127	dolichol-phosphate (beta-D) mannosyltransferase 1 (Dpm1) alternative variant bSep08, mRNA.

<a href="#">Dpm1</a>	<a href="#">Dpm1.cSep08</a>	<a href="#">296394</a>	9903	332	4	110	dolichol-phosphate (beta-D) mannosyltransferase 1 (Dpm1) alternative variant cSep08, mRNA.
<a href="#">Dpm1</a>	<a href="#">Dpm1.dSep08</a>	<a href="#">296394</a>	10159	982	4	102	dolichol-phosphate (beta-D) mannosyltransferase 1 (Dpm1) alternative variant dSep08, mRNA.
<a href="#">Dpp4</a>	<a href="#">Dpp4.aSep08</a>	<a href="#">25253</a>	52079	2378		703	dipeptidylpeptidase 4 (Dpp4) alternative variant aSep08, mRNA.
<a href="#">Dpp4</a>	<a href="#">Dpp4.bSep08</a>	<a href="#">25253</a>	23512	1327		323	dipeptidylpeptidase 4 (Dpp4) alternative variant bSep08, mRNA.
<a href="#">Dpp6</a>	<a href="#">Dpp6.bSep08</a>	<a href="#">29272</a>	562440	1791	4	544	dipeptidylpeptidase 6 CRA a (Dpp6) alternative variant bSep08, mRNA.
<a href="#">Dpp6</a>	<a href="#">Dpp6.cSep08</a>	<a href="#">29272</a>	471784	641	8	192	dipeptidylpeptidase 6 CRA c (Dpp6) alternative variant cSep08, mRNA.
<a href="#">Dpp6</a>	<a href="#">Dpp6.dSep08</a>	<a href="#">29272</a>	6502	540	4	116	dipeptidylpeptidase 6 CRA c (Dpp6) alternative variant dSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.bSep08</a>	<a href="#">83799</a>	1351	764	4	152	dipeptidylpeptidase 7 CRA c (15.9 kD) (Dpp7) alternative variant bSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.cSep08</a>	<a href="#">83799</a>	1431	729	4	150	dipeptidylpeptidase 7 CRA a (Dpp7) alternative variant cSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.dSep08</a>	<a href="#">83799</a>	861	773	2	122	dipeptidylpeptidase 7 (Dpp7) alternative variant dSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.eSep08</a>	<a href="#">83799</a>	861	755	2	116	dipeptidylpeptidase 7 CRA a (Dpp7) alternative variant eSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.fSep08</a>	<a href="#">83799</a>	877	721	2	103	dipeptidylpeptidase 7 CRA a (Dpp7) alternative variant fSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.gSep08</a>	<a href="#">83799</a>	712	220	3	73	dipeptidyl-peptidase 7 (Dpp7) alternative variant gSep08, mRNA.
<a href="#">Dpp7</a>	<a href="#">Dpp7.iSep08</a>	<a href="#">83799</a>	396	287	2	58	dipeptidyl-peptidase 7 (Dpp7) alternative variant iSep08, mRNA.
<a href="#">Dpp8</a>	<a href="#">Dpp8.bSep08</a>	<a href="#">315758</a>	26397	1777	12	494	dipeptidylpeptidase 8 (Dpp8) alternative variant bSep08, mRNA.
<a href="#">Dpp8</a>	<a href="#">Dpp8.cSep08</a>	<a href="#">315758</a>	21947	1264	10	359	dipeptidylpeptidase 8 (Dpp8) alternative variant cSep08, mRNA.
<a href="#">Dpp8</a>	<a href="#">Dpp8.dSep08</a>	<a href="#">315758</a>	9226	373	3	124	dipeptidylpeptidase 8 (Dpp8) alternative variant dSep08, mRNA.
<a href="#">Dpp8</a>	<a href="#">Dpp8.hSep08</a>	<a href="#">315758</a>	3531	721	5	58	dipeptidylpeptidase 8 (Dpp8) alternative variant hSep08, mRNA.
<a href="#">Dpp9</a>	<a href="#">Dpp9.aSep08</a>	<a href="#">301130</a>	15502	769		255	dipeptidyl peptidase 9 (Dpp9) mRNA.
<a href="#">DPPIV_N.0</a>	<a href="#">DPPIV_N.0.aSep08</a>		12891	1028		342	dipeptidylpeptidase 9 (DPPIV_N.0) mRNA.
<a href="#">Dpt</a>	<a href="#">Dpt.aSep08</a>	<a href="#">289178</a>	28723	1975	3	202	dermatopontin (24.2 kD) (Dpt) alternative variant aSep08, mRNA.
<a href="#">Dpt</a>	<a href="#">Dpt.cSep08</a>	<a href="#">289178</a>	27608	734	2	160	dermatopontin (19.0 kD) (Dpt) alternative variant cSep08, complete mRNA.
<a href="#">Dpy1911</a>	<a href="#">Dpy1911.aSep08</a>	<a href="#">315496</a>	65262	4203	19	592	CRA a (Dpy1911) alternative variant aSep08, mRNA.
<a href="#">Dpy1911</a>	<a href="#">Dpy1911.bSep08</a>	<a href="#">315496</a>	7507	786	3	85	CRA a (Dpy1911) alternative variant bSep08, mRNA.
<a href="#">Dpy1911</a>	<a href="#">Dpy1911.cSep08</a>	<a href="#">315496</a>	6117	635	2	46	putative protein (5.1 kD) (Dpy1911) alternative variant cSep08, mRNA.

<a href="#">Dpy1914</a>	<a href="#">Dpy1914.aSep08</a>	<a href="#">297824</a>	20341	2037		249	CRA a (Dpy1914) alternative variant aSep08, mRNA.
<a href="#">Dpy1914</a>	<a href="#">Dpy1914.bSep08</a>	<a href="#">297824</a>	7946	1067		141	CRA a (Dpy1914) alternative variant bSep08, mRNA.
<a href="#">Dpy30</a>	<a href="#">Dpy30.bSep08</a>	<a href="#">286897</a>	14753	673	4	99	protein CRA b (11.2 kD) (Dpy30) alternative variant bSep08, mRNA.
<a href="#">Dpy30</a>	<a href="#">Dpy30.cSep08</a>	<a href="#">286897</a>	14711	488	3	69	dpy-30 homolog like (Dpy30) alternative variant cSep08, mRNA.
<a href="#">Dpy30</a>	<a href="#">Dpy30.dSep08</a>	<a href="#">286897</a>	5711	598	1	51	dosage compensation-related protein DPY30 like (6.3 kD) (Dpy30) alternative variant dSep08, mRNA.
<a href="#">Dpyd</a>	<a href="#">Dpyd.aSep08</a>	<a href="#">81656</a>	322931	2380		386	dihydropyrimidine dehydrogenase (Dpyd) mRNA.
<a href="#">Dpys</a>	<a href="#">Dpys.bSep08</a>	<a href="#">65135</a>	40448	351	3	60	dihydropyrimidinase (Dpys) alternative variant bSep08, mRNA.
<a href="#">Dpys</a>	<a href="#">Dpys.cSep08</a>	<a href="#">65135</a>	40533	290	2	60	dihydropyrimidinase (Dpys) alternative variant cSep08, mRNA.
<a href="#">Dpysl2</a>	<a href="#">Dpysl2.bSep08</a>	<a href="#">25416</a>	44696	602	3	200	dihydropyrimidinase-like 2 (Dpysl2) alternative variant bSep08, mRNA.
<a href="#">Dpysl3</a>	<a href="#">Dpysl3.bSep08</a>	<a href="#">25418</a>	19269	710	6	204	dihydropyrimidinase-like 3 (Dpysl3) alternative variant bSep08, mRNA.
<a href="#">Dpysl3</a>	<a href="#">Dpysl3.cSep08</a>	<a href="#">25418</a>	76149	593	4	197	dihydropyrimidinase-like 3 (Dpysl3) alternative variant cSep08, mRNA.
<a href="#">Dr1</a>	<a href="#">Dr1.bSep08</a>	<a href="#">289881</a>	4657	2207	1	116	down-regulator of transcription 1 (Dr1) alternative variant bSep08, mRNA.
<a href="#">Drap1</a>	<a href="#">Drap1.aSep08</a>	<a href="#">293674</a>	2004	793	7	229	dr1 associated protein 1 (negative cofactor 2 alpha) (Drap1) alternative variant aSep08, mRNA.
<a href="#">DREV.0</a>	<a href="#">DREV.0.aSep08</a>		45235	1649	3	318	methyltransferase like 9 precursor (36.4 kD) (DREV.0) alternative variant aSep08, mRNA.
<a href="#">DREV.0</a>	<a href="#">DREV.0.bSep08</a>		44979	1390	3	296	methyltransferase like 9 (DREV.0) alternative variant bSep08, mRNA.
<a href="#">DREV.0</a>	<a href="#">DREV.0.cSep08</a>		28534	870	3	238	methyltransferase like 9 (27.4 kD) (DREV.0) alternative variant cSep08, mRNA.
<a href="#">Drf_DAD.0</a>	<a href="#">Drf_DAD.0.aSep08</a>		64096	610		200	diaphanous homolog 3 CRA a (Drf_DAD.0) mRNA.
<a href="#">Drf_FH3.0</a>	<a href="#">Drf_FH3.0.aSep08</a>		11975	881		240	dishevelled associated activator of morphogenesis 2 CRA a (Drf_FH3.0) mRNA.
<a href="#">Drf_FH3.1</a>	<a href="#">Drf_FH3.1.aSep08</a>		2498	652		217	formin-like 3 CRA c (Drf_FH3.1) mRNA.
<a href="#">Drg1</a>	<a href="#">Drg1.bSep08</a>	<a href="#">305470</a>	6533	634	4	178	developmentally regulated GTP binding protein 1 (Drg1) alternative variant bSep08, mRNA.
<a href="#">Drg2</a>	<a href="#">Drg2.aSep08</a>	<a href="#">497915</a>	14046	1784	3	430	developmentally regulated GTP binding protein 2 (Drg2) alternative variant aSep08, complete mRNA.
<a href="#">Drg2</a>	<a href="#">Drg2.bSep08</a>	<a href="#">497915</a>	8387	1067	5	269	developmentally regulated GTP binding protein 2 (29.7 kD) (Drg2) alternative variant bSep08, complete mRNA.
<a href="#">Drg2</a>	<a href="#">Drg2.cSep08</a>	<a href="#">497915</a>	13808	1459	6	188	developmentally regulated GTP binding protein 2 (20.9 kD) (Drg2) alternative variant cSep08, mRNA.
<a href="#">Drg2</a>	<a href="#">Drg2.dSep08</a>	<a href="#">497915</a>	2797	503	3	135	developmentally regulated GTP binding protein 2 (Drg2) alternative variant dSep08, mRNA.
<a href="#">Drg2</a>	<a href="#">Drg2.eSep08</a>	<a href="#">497915</a>	1057	726	1	46	developmentally regulated GTP binding protein 2 (Drg2) alternative variant eSep08, mRNA.

<a href="#">Drp2</a>	<a href="#">Drp2.bSep08</a>	<a href="#">66027</a>	6652	595	5	198	dystrophin related protein 2 (Drp2) alternative variant bSep08, mRNA.
<a href="#">Drp2</a>	<a href="#">Drp2.cSep08</a>	<a href="#">66027</a>	645	385	2	33	dystrophin related protein 2 (Drp2) alternative variant cSep08, mRNA.
<a href="#">Dsc1</a>	<a href="#">Dsc1.bSep08</a>	<a href="#">291759</a>	7068	748	2	249	desmocollin 1 (Dsc1) alternative variant bSep08, mRNA.
<a href="#">Dsc2</a>	<a href="#">Dsc2.aSep08</a>	<a href="#">291760</a>	9236	1042		347	desmocollin 2 (Dsc2) mRNA.
<a href="#">Dsc3</a>	<a href="#">Dsc3.bSep08</a>	<a href="#">307563</a>	4102	1135	2	119	desmocollin 3 (Dsc3) alternative variant bSep08, mRNA.
<a href="#">Dscam</a>	<a href="#">Dscam.aSep08</a>	<a href="#">171119</a>	18652	1431		284	down syndrome cell adhesion molecule (Dscam) mRNA.
<a href="#">Dsccl1</a>	<a href="#">Dsccl1.aSep08</a>	<a href="#">299933</a>	15467	1369		413	defective in sister chromatid cohesion 1 homolog (S. cerevisiae) (Dsccl1) mRNA.
<a href="#">Dscr3</a>	<a href="#">Dscr3.aSep08</a>	<a href="#">360703</a>	25162	1112	6	219	down syndrome critical region gene 3 (Dscr3) alternative variant aSep08, mRNA.
<a href="#">Dscr3</a>	<a href="#">Dscr3.bSep08</a>	<a href="#">360703</a>	22674	665	5	177	down syndrome critical region gene 3 (Dscr3) alternative variant bSep08, mRNA.
<a href="#">Dscr3</a>	<a href="#">Dscr3.eSep08</a>	<a href="#">360703</a>	14804	641	2	62	down syndrome critical region gene 3 (6.5 kD) (Dscr3) alternative variant eSep08, mRNA.
<a href="#">Dsg1b</a>	<a href="#">Dsg1b.aSep08</a>	<a href="#">291755</a>	3427	547	3	158	desmoglein 1 beta (Dsg1b) alternative variant aSep08, mRNA.
<a href="#">Dsg2</a>	<a href="#">Dsg2.aSep08</a>	<a href="#">307562</a>	2707	346		115	desmoglein 2 (Dsg2) mRNA.
<a href="#">Dsg3</a>	<a href="#">Dsg3.aSep08</a>	<a href="#">291752</a>	3794	424		141	desmoglein 3 (Dsg3) mRNA.
<a href="#">DSHCT.0</a>	<a href="#">DSHCT.0.aSep08</a>		11981	761	6	211	superkiller viralicidic activity 2-like 2 (DSHCT.0) alternative variant aSep08, mRNA.
<a href="#">DSHCT.0</a>	<a href="#">DSHCT.0.bSep08</a>		6466	692	4	210	superkiller viralicidic activity 2-like 2 (DSHCT.0) alternative variant bSep08, mRNA.
<a href="#">DSHCT.0</a>	<a href="#">DSHCT.0.cSep08</a>		15882	879	6	196	superkiller viralicidic activity 2-like 2 (DSHCT.0) alternative variant cSep08, mRNA.
<a href="#">DSHCT.0</a>	<a href="#">DSHCT.0.dSep08</a>		12029	688	5	161	superkiller viralicidic activity 2-like 2 (DSHCT.0) alternative variant dSep08, mRNA.
<a href="#">DSHCT.0</a>	<a href="#">DSHCT.0.eSep08</a>		10789	732	5	150	superkiller viralicidic activity 2-like 2 (17.2 kD) (DSHCT.0) alternative variant eSep08, mRNA.
<a href="#">DSL.0</a>	<a href="#">DSL.0.aSep08</a>		13744	1244		414	jagged 2 (DSL.0) mRNA.
<a href="#">Dsn1</a>	<a href="#">Dsn1.bSep08</a>	<a href="#">499933</a>	4618	449	2	128	DSN1, MIND kinetochore complex component, homolog (S. cerevisiae) and similar to 40S ribosomal protein S2 (Dsn1) alternative variant bSep08, mRNA.
<a href="#">Dsn1</a>	<a href="#">Dsn1.bSep08</a>	<a href="#">679915</a>	4618	449	2	128	DSN1, MIND kinetochore complex component, homolog (S. cerevisiae) and similar to 40S ribosomal protein S2 (Dsn1) alternative variant bSep08, mRNA.
<a href="#">Dsn1</a>	<a href="#">Dsn1.cSep08</a>	<a href="#">499933</a>	3412	362	2	33	DSN1, MIND kinetochore complex component, homolog (S. cerevisiae) and similar to 40S ribosomal protein S2 (Dsn1) alternative variant cSep08, mRNA.
<a href="#">Dsn1</a>	<a href="#">Dsn1.cSep08</a>	<a href="#">679915</a>	3412	362	2	33	DSN1, MIND kinetochore complex component, homolog (S. cerevisiae) and similar to 40S ribosomal protein S2 (Dsn1) alternative variant cSep08, mRNA.
<a href="#">dsRNA_bind.0</a>	<a href="#">dsRNA_bind.0.aSep08</a>		5141	423		140	dicer1 (dsRNA_bind.0) mRNA.
<a href="#">Dst</a>	<a href="#">Dst.bSep08</a>	<a href="#">316313</a>	13766	1986	9	398	dystonin (Dst) alternative variant bSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.cSep08</a>	<a href="#">316313</a>	12039	766	7	254	dystonin (Dst) alternative variant cSep08, mRNA.



<a href="#">Dst</a>	<a href="#">Dst.dSep08</a>	<a href="#">316313</a>	3804	717	6	238	dystonin (Dst) alternative variant dSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.eSep08</a>	<a href="#">316313</a>	12331	665	4	221	dystonin (Dst) alternative variant eSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.fSep08</a>	<a href="#">316313</a>	11235	565	7	188	dystonin (Dst) alternative variant fSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.gSep08</a>	<a href="#">316313</a>	1474	856	2	142	dystonin (15.0 kD) (Dst) alternative variant gSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.hSep08</a>	<a href="#">316313</a>	5234	413	3	137	dystonin CRA f (Dst) alternative variant hSep08, mRNA.
<a href="#">Dst</a>	<a href="#">Dst.iSep08</a>	<a href="#">316313</a>	4207	738	3	127	dystonin (Dst) alternative variant iSep08, mRNA.
<a href="#">Dstn</a>	<a href="#">Dstn.bSep08</a>	<a href="#">502674</a>	3033	796	2	51	destrin (5.5 kD) (Dstn) alternative variant bSep08, mRNA.
<a href="#">Dtd1</a>	<a href="#">Dtd1.aSep08</a>	<a href="#">362227</a>	169273	1245	1	209	D-tyrosyl-tRNA deacylase 1 homolog (S. cerevisiae) (23.4 kD) (Dtd1) alternative variant aSep08, complete mRNA.
<a href="#">Dtl</a>	<a href="#">Dtl.aSep08</a>	<a href="#">305073</a>	24218	1063	2	353	denticleless homolog (Drosophila) (Dtl) alternative variant aSep08, mRNA.
<a href="#">Dtl</a>	<a href="#">Dtl.bSep08</a>	<a href="#">305073</a>	28747	948	4	315	denticleless homolog (Drosophila) (Dtl) alternative variant bSep08, mRNA.
<a href="#">Dtna</a>	<a href="#">Dtna.aSep08</a>	<a href="#">307548</a>	11894	761	5	174	dystrobrevin alpha (Dtna) alternative variant aSep08, mRNA.
<a href="#">Dtna</a>	<a href="#">Dtna.bSep08</a>	<a href="#">307548</a>	20621	733	6	172	dystrobrevin alpha (Dtna) alternative variant bSep08, mRNA.
<a href="#">Dtna</a>	<a href="#">Dtna.cSep08</a>	<a href="#">307548</a>	28783	785	6	168	dystrobrevin alpha (Dtna) alternative variant cSep08, mRNA.
<a href="#">Dtna</a>	<a href="#">Dtna.dSep08</a>	<a href="#">307548</a>	20993	403	4	134	dystrobrevin alpha (Dtna) alternative variant dSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.bSep08</a>	<a href="#">362715</a>	200332	1797	7	572	dystrobrevin beta CRA a (Dtnb) alternative variant bSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.cSep08</a>	<a href="#">362715</a>	71007	768	7	255	dystrobrevin beta CRA a (Dtnb) alternative variant cSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.dSep08</a>	<a href="#">362715</a>	49921	1192	9	190	dystrobrevin beta (21.1 kD) (Dtnb) alternative variant dSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.eSep08</a>	<a href="#">362715</a>	8365	668	6	135	CRA a like (Dtnb) alternative variant eSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.fSep08</a>	<a href="#">362715</a>	30380	350	3	116	dystrobrevin beta CRA a (Dtnb) alternative variant fSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.gSep08</a>	<a href="#">362715</a>	8332	455	4	116	CRA a like (Dtnb) alternative variant gSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.iSep08</a>	<a href="#">362715</a>	1896	414	2	98	putative protein (Dtnb) alternative variant iSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.jSep08</a>	<a href="#">362715</a>	26706	601	5	91	dystrobrevin beta (Dtnb) alternative variant jSep08, mRNA.
<a href="#">Dtnb</a>	<a href="#">Dtnb.kSep08</a>	<a href="#">362715</a>	8131	295	4	76	putative protein (Dtnb) alternative variant kSep08, mRNA.
<a href="#">Dtwd1</a>	<a href="#">Dtwd1.cSep08</a>	<a href="#">296119</a>	2914	365	2	56	putative protein of mammalian origin (Dtwd1) alternative variant cSep08, mRNA.
<a href="#">Dtwd2</a>	<a href="#">Dtwd2.bSep08</a>	<a href="#">361326</a>	90241	2044	1	184	CRA c (20.3 kD) (Dtwd2) alternative variant bSep08, mRNA.
<a href="#">Dtx2</a>	<a href="#">Dtx2.bSep08</a>	<a href="#">304591</a>	39604	2589	6	619	deltex 2 homolog (Drosophila) (67.2 kD) (Dtx2) alternative variant bSep08, mRNA.
<a href="#">Dtx2</a>	<a href="#">Dtx2.cSep08</a>	<a href="#">304591</a>	11876	858	1	271	deltex 2 homolog (Drosophila) (Dtx2) alternative variant cSep08, mRNA.
<a href="#">Dtx3</a>	<a href="#">Dtx3.aSep08</a>	<a href="#">500847</a>	1210	898	2	97	deltex 3 homolog (Drosophila) (Dtx3) alternative variant aSep08, mRNA.

<a href="#">Dtymk</a>	<a href="#">Dtymk.bSep08</a>	<a href="#">301622</a>	8613	1169	6	195	deoxythymidylate kinase (Dtymk) alternative variant bSep08, mRNA.
<a href="#">Dtymk</a>	<a href="#">Dtymk.cSep08</a>	<a href="#">301622</a>	1334	418	2	130	deoxythymidylate kinase (Dtymk) alternative variant cSep08, mRNA.
<a href="#">Dtymk</a>	<a href="#">Dtymk.dSep08</a>	<a href="#">301622</a>	7456	778	4	121	deoxythymidylate kinase (Dtymk) alternative variant dSep08, mRNA.
<a href="#">Dtymk</a>	<a href="#">Dtymk.fSep08</a>	<a href="#">301622</a>	4982	578	3	94	deoxythymidylate kinase CRA d (10.6 kD) (Dtymk) alternative variant fSep08, mRNA.
<a href="#">Dtymk</a>	<a href="#">Dtymk.iSep08</a>	<a href="#">301622</a>	3611	707	3	45	putative protein (6.4 kD) (Dtymk) alternative variant iSep08, mRNA.
<a href="#">duby</a>	<a href="#">duby.aSep08</a>		13712	1791		597	dedicator of cytokinesis 11 (duby) alternative variant aSep08, mRNA.
<a href="#">duby</a>	<a href="#">duby.bSep08</a>		11116	563		187	dedicator of cytokinesis 11 (duby) alternative variant bSep08, mRNA.
<a href="#">duchy</a>	<a href="#">duchy.aSep08</a>		65997	783		49	putative protein (5.5 kD) (duchy) mRNA.
<a href="#">DUF590.0</a>	<a href="#">DUF590.0.aSep08</a>		585	396		132	transmembrane protein 16h (DUF590.0) mRNA.
<a href="#">DUF618.0</a>	<a href="#">DUF618.0.aSep08</a>		5784	668	3	222	cleavage polyadenylation factor homolog (DUF618.0) alternative variant aSep08, mRNA.
<a href="#">DUF622.0</a>	<a href="#">DUF622.0.aSep08</a>		10020	1092		240	putative protein, with a coiled coil domain, of vertebrate origin (DUF622.0) mRNA.
<a href="#">DUF622.1</a>	<a href="#">DUF622.1.aSep08</a>	<a href="#">501248</a>	13465	709		154	putative protein, with 2 coiled coil domains, of mammalian origin (DUF622.1) mRNA.
<a href="#">DUF622.2</a>	<a href="#">DUF622.2.aSep08</a>		1861	545		83	putative protein, with 2 coiled coil domains, of mammalian origin (DUF622.2) mRNA.
<a href="#">DUF622.3</a>	<a href="#">DUF622.3.bSep08</a>		37707	751	3	121	putative protein, with a coiled coil domain, of mammalian origin (14.5 kD) (DUF622.3) alternative variant bSep08, mRNA.
<a href="#">DUF622.4</a>	<a href="#">DUF622.4.aSep08</a>		2485	437		145	putative protein, with a coiled coil domain, of mammalian origin (DUF622.4) mRNA.
<a href="#">DUF622.6</a>	<a href="#">DUF622.6.aSep08</a>		8366	779	3	206	putative protein, with 2 coiled coil domains, of mammalian origin (DUF622.6) alternative variant aSep08, mRNA.
<a href="#">DUF622.6</a>	<a href="#">DUF622.6.bSep08</a>		3293	735	1	122	putative protein, with 2 coiled coil domains, of mammalian origin (DUF622.6) alternative variant bSep08, mRNA.
<a href="#">DUF663.0</a>	<a href="#">DUF663.0.aSep08</a>		7669	2212	10	444	tsr1 20S rRNA accumulation (DUF663.0) alternative variant aSep08, mRNA.
<a href="#">DUF663.0</a>	<a href="#">DUF663.0.bSep08</a>		2378	783	5	214	tsr1 20S rRNA accumulation homolog (24.6 kD) (DUF663.0) alternative variant bSep08, mRNA.
<a href="#">DUF663.0</a>	<a href="#">DUF663.0.cSep08</a>		5342	737	6	171	tsr1 20S rRNA accumulation (DUF663.0) alternative variant cSep08, mRNA.
<a href="#">DUF663.0</a>	<a href="#">DUF663.0.dSep08</a>		1084	837	2	64	tsr1 20S rRNA accumulation (DUF663.0) alternative variant dSep08, mRNA.
<a href="#">DUF729.0</a>	<a href="#">DUF729.0.aSep08</a>		15705	1930	8	404	putative protein, with a coiled coil domain, of eukaryotic origin (DUF729.0) alternative variant aSep08, mRNA.
<a href="#">DUF729.0</a>	<a href="#">DUF729.0.bSep08</a>		4222	1018	3	175	aa2-041 (19.2 kD) (DUF729.0) alternative variant bSep08, mRNA.
<a href="#">DUF729.0</a>	<a href="#">DUF729.0.dSep08</a>		6661	315	3	49	putative protein (5.4 kD) (DUF729.0) alternative variant dSep08, mRNA.

<a href="#">DUF814.0</a>	<a href="#">DUF814.0.aSep08</a>		18165	1394		463	serologically defined colon cancer antigen 1 like (DUF814.0) alternative variant aSep08, mRNA.
<a href="#">DUF814.0</a>	<a href="#">DUF814.0.bSep08</a>		3515	542		180	serologically defined colon cancer antigen 1 CRA f like (DUF814.0) alternative variant bSep08, mRNA.
<a href="#">DUF908.0</a>	<a href="#">DUF908.0.aSep08</a>		5859	771		256	putative protein of eukaryotic origin (DUF908.0) mRNA.
<a href="#">DUF913.0</a>	<a href="#">DUF913.0.aSep08</a>		10665	864	4	287	putative protein of metazoan origin (DUF913.0) alternative variant aSep08, mRNA.
<a href="#">DUF913.0</a>	<a href="#">DUF913.0.bSep08</a>		8038	1119	3	240	putative protein of eukaryotic origin (DUF913.0) alternative variant bSep08, mRNA.
<a href="#">DUF947.0</a>	<a href="#">DUF947.0.aSep08</a>		7901	1095	7	254	CRA a (DUF947.0) alternative variant aSep08, mRNA.
<a href="#">DUF947.0</a>	<a href="#">DUF947.0.bSep08</a>		7870	1061	7	244	CRA a (28.5 kD) (DUF947.0) alternative variant bSep08, mRNA.
<a href="#">DUF947.0</a>	<a href="#">DUF947.0.cSep08</a>		6148	795	6	207	CRA a (DUF947.0) alternative variant cSep08, mRNA.
<a href="#">DUF947.0</a>	<a href="#">DUF947.0.dSep08</a>		2385	752	4	196	CRA b (DUF947.0) alternative variant dSep08, mRNA.
<a href="#">DUF1041.0</a>	<a href="#">DUF1041.0.aSep08</a>		1541	384		128	unc-13 homolog (DUF1041.0) mRNA.
<a href="#">DUF1041.1</a>	<a href="#">DUF1041.1.aSep08</a>		22281	281		74	activator protein for secretion 2 (DUF1041.1) mRNA.
<a href="#">DUF1227.0</a>	<a href="#">DUF1227.0.aSep08</a>		1721	698		232	excision repair cross-complementing rodent deficiency complementation group 2 (DUF1227.0) mRNA.
<a href="#">DUF1242.1</a>	<a href="#">DUF1242.1.aSep08</a>		18181	3273	4	72	transmembrane protein 167 (8.1 kD) (DUF1242.1) alternative variant aSep08, mRNA.
<a href="#">DUF1242.1</a>	<a href="#">DUF1242.1.bSep08</a>		14470	875	4	39	putative protein (4.5 kD) (DUF1242.1) alternative variant bSep08, mRNA.
<a href="#">DUF1242.1</a>	<a href="#">DUF1242.1.cSep08</a>		6488	595	3	39	putative protein (4.5 kD) (DUF1242.1) alternative variant cSep08, mRNA.
<a href="#">DUF1358.0</a>	<a href="#">DUF1358.0.aSep08</a>		26747	967	3	144	putative protein of metazoan origin (DUF1358.0) alternative variant aSep08, mRNA.
<a href="#">DUF1358.0</a>	<a href="#">DUF1358.0.bSep08</a>		12744	562	3	142	putative cytoplasmic protein of bilaterial origin (14.9 kD) (DUF1358.0) alternative variant bSep08, complete mRNA.
<a href="#">DUF1358.0</a>	<a href="#">DUF1358.0.cSep08</a>		1543	1003	1	65	putative protein of vertebrate origin (6.6 kD) (DUF1358.0) alternative variant cSep08, mRNA.
<a href="#">DUF1387.0</a>	<a href="#">DUF1387.0.bSep08</a>	<a href="#">501207</a>	88105	473	3	123	CRA b (DUF1387.0) alternative variant bSep08, mRNA.
<a href="#">DUF1387.0</a>	<a href="#">DUF1387.0.cSep08</a>	<a href="#">501207</a>	88142	425	2	116	CRA b (DUF1387.0) alternative variant cSep08, mRNA.
<a href="#">DUF1387.0</a>	<a href="#">DUF1387.0.dSep08</a>	<a href="#">501207</a>	85018	363	2	34	putative protein (4.0 kD) (DUF1387.0) alternative variant dSep08, mRNA.
<a href="#">DUF1725.115</a>	<a href="#">DUF1725.115.aSep08</a>		143333	676		73	endonuclease reverse transcriptase like (8.7 kD) (DUF1725.115) mRNA.
<a href="#">DUF1725.254</a>	<a href="#">DUF1725.254.aSep08</a>		29459	722		228	reverse transcriptase (DUF1725.254) mRNA.
<a href="#">DUF1725.282</a>	<a href="#">DUF1725.282.aSep08</a>		1545	750		61	endonuclease reverse transcriptase like (7.7 kD) (DUF1725.282) mRNA.
<a href="#">DUF1725.308</a>	<a href="#">DUF1725.308.bSep08</a>		98844	313	1	52	novel protein containing a Zinc carboxypeptidase domain (DUF1725.308) alternative variant bSep08, mRNA.
<a href="#">DUF1781.0</a>	<a href="#">DUF1781.0.aSep08</a>		892	709		178	CRA b (DUF1781.0) mRNA.
<a href="#">DUF1855.0</a>	<a href="#">DUF1855.0.aSep08</a>		17987	628		155	axin dorsalization (DUF1855.0) mRNA.
<a href="#">DUF2039.0</a>	<a href="#">DUF2039.0.aSep08</a>		1271	867		168	uncharacterized protein c9orf85 like (DUF2039.0) mRNA.
<a href="#">DUF2346.0</a>	<a href="#">DUF2346.0.aSep08</a>		1657	320	4	82	CRA a (DUF2346.0) alternative variant aSep08, mRNA.

<a href="#">DUF2346.0</a>	<a href="#">DUF2346.0.bSep08</a>		1657	441	4	67	CRA a (8.4 kD) (DUF2346.0) alternative variant bSep08, mRNA.
<a href="#">DUF2346.0</a>	<a href="#">DUF2346.0.cSep08</a>		1493	748	2	41	CRA a like (DUF2346.0) alternative variant cSep08, mRNA.
<a href="#">DUF2356.0</a>	<a href="#">DUF2356.0.aSep08</a>		11999	788		262	integrator complex CRA a (DUF2356.0) mRNA.
<a href="#">DUF2372.0</a>	<a href="#">DUF2372.0.aSep08</a>		2890	773	3	122	proteasome chaperone 3 (13.3 kD) (DUF2372.0) alternative variant aSep08, complete mRNA.
<a href="#">DUF2372.0</a>	<a href="#">DUF2372.0.bSep08</a>		2424	395	3	85	proteasome chaperone 3 (DUF2372.0) alternative variant bSep08, mRNA.
<a href="#">DUF2372.0</a>	<a href="#">DUF2372.0.cSep08</a>		2347	587	2	56	proteasome chaperone 3 (DUF2372.0) alternative variant cSep08, mRNA.
<a href="#">DUF2372.0</a>	<a href="#">DUF2372.0.dSep08</a>		903	457	2	96	proteasome chaperone 3 (10.6 kD) (DUF2372.0) alternative variant dSep08, mRNA.
<a href="#">DUF2450.0</a>	<a href="#">DUF2450.0.aSep08</a>		56815	1224		407	putative protein, with 2 coiled coil domains, of eukaryotic origin (DUF2450.0) mRNA.
<a href="#">dufer</a>	<a href="#">dufer.aSep08</a>		13534	468		53	putative protein (dufer) mRNA.
<a href="#">duflo</a>	<a href="#">duflo.aSep08</a>		7609	712		167	putative protein (18.3 kD) (duflo) mRNA.
<a href="#">duflu</a>	<a href="#">duflu.aSep08</a>		572	325		108	WW domain binding protein 7 like (duflu) mRNA.
<a href="#">dukee</a>	<a href="#">dukee.aSep08</a>		5436	428		36	putative protein (4.1 kD) (dukee) mRNA.
<a href="#">Dullard</a>	<a href="#">Dullard.aSep08</a>	<a href="#">287447</a>	9640	1603	7	309	dullard homolog (Xenopus laevis) (Dullard) alternative variant aSep08, mRNA.
<a href="#">Dullard</a>	<a href="#">Dullard.bSep08</a>	<a href="#">287447</a>	1542	957	1	40	dullard homolog (Xenopus laevis) (Dullard) alternative variant bSep08, mRNA.
<a href="#">duloy</a>	<a href="#">duloy.aSep08</a>		6166	373		124	putative protein of mammalian origin (duloy) mRNA.
<a href="#">dumer</a>	<a href="#">dumer.aSep08</a>		1551	788		40	putative protein (dumer) mRNA.
<a href="#">dunoy</a>	<a href="#">dunoy.aSep08</a>		3443	1001	2	83	putative protein (dunoy) alternative variant aSep08, mRNA.
<a href="#">dunoy</a>	<a href="#">dunoy.bSep08</a>		619	230	1	35	putative protein (4.1 kD) (dunoy) alternative variant bSep08, mRNA.
<a href="#">Duox2</a>	<a href="#">Duox2.aSep08</a>	<a href="#">79107</a>	882	743		80	dual oxidase 2 (Duox2) mRNA.
<a href="#">dupor</a>	<a href="#">dupor.aSep08</a>		3634	1801		169	CRA a (dupor) alternative variant aSep08, mRNA.
<a href="#">Dus3l</a>	<a href="#">Dus3l.bSep08</a>	<a href="#">301122</a>	1975	1447	2	298	dihydrouridine synthase 3-like (33.1 kD) (Dus3l) alternative variant bSep08, mRNA.
<a href="#">Dus3l</a>	<a href="#">Dus3l.cSep08</a>	<a href="#">301122</a>	1788	850	5	174	dihydrouridine synthase 3-like (19.9 kD) (Dus3l) alternative variant cSep08, mRNA.
<a href="#">Dus3l</a>	<a href="#">Dus3l.dSep08</a>	<a href="#">301122</a>	1058	787	3	141	dihydrouridine synthase 3-like (Dus3l) alternative variant dSep08, mRNA.
<a href="#">dusa</a>	<a href="#">dusa.bSep08</a>		421	293	2	59	putative protein (dusa) alternative variant bSep08, mRNA.
<a href="#">dushee</a>	<a href="#">dushee.aSep08</a>		1450	310		34	putative protein (dushee) mRNA.
<a href="#">Dusp3</a>	<a href="#">Dusp3.aSep08</a>	<a href="#">498003</a>	11824	4232	2	193	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) (Dusp3) alternative variant aSep08, mRNA.
<a href="#">Dusp3</a>	<a href="#">Dusp3.bSep08</a>	<a href="#">498003</a>	1409	389	1	26	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) (2.8 kD) (Dusp3) alternative variant bSep08, mRNA.
<a href="#">Dusp5</a>	<a href="#">Dusp5.cSep08</a>	<a href="#">171109</a>	7752	323	2	107	dual specificity phosphatase 5 (Dusp5) alternative variant cSep08, mRNA.

<a href="#">Dusp7</a>	<a href="#">Dusp7.aSep08</a>	<a href="#">300980</a>	1979	565		188	dual specificity phosphatase 7 (Dusp7) mRNA.
<a href="#">Dusp8</a>	<a href="#">Dusp8.bSep08</a>	<a href="#">361679</a>	5012	3940	4	514	dual specificity phosphatase 8 (Dusp8) alternative variant bSep08, mRNA.
<a href="#">Dusp11</a>	<a href="#">Dusp11.bSep08</a>	<a href="#">297412</a>	3818	638	1	75	dual specificity phosphatase 11 (RNA/RNP complex 1-interacting) (Dusp11) alternative variant bSep08, mRNA.
<a href="#">Dusp12</a>	<a href="#">Dusp12.bSep08</a>	<a href="#">64014</a>	4998	387	1	23	dual specificity phosphatase 12 (Dusp12) alternative variant bSep08, mRNA.
<a href="#">Dusp13</a>	<a href="#">Dusp13.aSep08</a>	<a href="#">361002</a>	15369	1747	8	326	dual specificity phosphatase 13 CRA b (36.4 kD) (Dusp13) alternative variant aSep08, complete mRNA.
<a href="#">Dusp13</a>	<a href="#">Dusp13.cSep08</a>	<a href="#">361002</a>	1732	751	3	99	dual specificity phosphatase 13 like (10.7 kD) (Dusp13) alternative variant cSep08, complete mRNA.
<a href="#">Dusp14</a>	<a href="#">Dusp14.aSep08</a>	<a href="#">360580</a>	18395	1363	2	258	dual specificity phosphatase 14 (Dusp14) alternative variant aSep08, mRNA.
<a href="#">Dusp15</a>	<a href="#">Dusp15.bSep08</a>	<a href="#">362238</a>	10505	3109	1	236	dual specificity phosphatase-like 15 (26.3 kD) (Dusp15) alternative variant bSep08, mRNA.
<a href="#">Dusp16</a>	<a href="#">Dusp16.bSep08</a>	<a href="#">297682</a>	40174	782	3	215	dual specificity phosphatase 16 (Dusp16) alternative variant bSep08, mRNA.
<a href="#">Dusp16</a>	<a href="#">Dusp16.cSep08</a>	<a href="#">297682</a>	22769	694	2	189	dual specificity phosphatase 16 (Dusp16) alternative variant cSep08, mRNA.
<a href="#">Dusp19</a>	<a href="#">Dusp19.bSep08</a>	<a href="#">311151</a>	140026	525	4	92	dual specificity phosphatase 19 (Dusp19) alternative variant bSep08, mRNA.
<a href="#">Dusp19</a>	<a href="#">Dusp19.cSep08</a>	<a href="#">311151</a>	5409	433	2	91	dual specificity phosphatase 19 (Dusp19) alternative variant cSep08, mRNA.
<a href="#">Dusp19</a>	<a href="#">Dusp19.dSep08</a>	<a href="#">311151</a>	12515	810	4	56	dual specificity phosphatase 19 (6.0 kD) (Dusp19) alternative variant dSep08, mRNA.
<a href="#">Dusp22</a>	<a href="#">Dusp22.bSep08</a>	<a href="#">361242</a>	51513	963	7	143	dual specificity phosphatase 22 (16.4 kD) (Dusp22) alternative variant bSep08, mRNA.
<a href="#">Dusp22</a>	<a href="#">Dusp22.dSep08</a>	<a href="#">361242</a>	4118	893	3	142	dual specificity phosphatase 22 (Dusp22) alternative variant dSep08, mRNA.
<a href="#">Dusp23</a>	<a href="#">Dusp23.aSep08</a>	<a href="#">360881</a>	1607	867	2	181	dual specificity phosphatase 23 (Dusp23) alternative variant aSep08, mRNA.
<a href="#">Dusp26</a>	<a href="#">Dusp26.aSep08</a>	<a href="#">306527</a>	5722	1952	1	211	dual specificity phosphatase 26 (putative) (23.9 kD) (Dusp26) alternative variant aSep08, mRNA.
<a href="#">Dusp26</a>	<a href="#">Dusp26.cSep08</a>	<a href="#">306527</a>	6762	595	2	151	dual specificity phosphatase 26 (putative) (Dusp26) alternative variant cSep08, mRNA.
<a href="#">Dusp27</a>	<a href="#">Dusp27.aSep08</a>	<a href="#">498267</a>	19246	760		237	dual specificity phosphatase 27 (putative) (26.6 kD) (Dusp27) mRNA.
<a href="#">Dut-ps</a>	<a href="#">Dut-ps.cSep08</a>	<a href="#">94200</a>	11679	1480	6	129	deoxyuridine triphosphatase pseudogene (Dut-ps) alternative variant cSep08, mRNA.
<a href="#">Dut-ps</a>	<a href="#">Dut-ps.dSep08</a>	<a href="#">94200</a>	8036	930	5	102	deoxyuridine triphosphatase pseudogene (Dut-ps) alternative variant dSep08, mRNA.
<a href="#">Dut-ps</a>	<a href="#">Dut-ps.eSep08</a>	<a href="#">94200</a>	2183	1815	3	52	deoxyuridine triphosphatase pseudogene (5.7 kD) (Dut-ps) alternative variant eSep08, mRNA.
<a href="#">Dut-ps</a>	<a href="#">Dut-ps.hSep08</a>	<a href="#">94200</a>	869	430	2	39	deoxyuridine triphosphatase pseudogene (4.2 kD) (Dut-ps) alternative variant hSep08, mRNA.
<a href="#">duto</a>	<a href="#">duto.aSep08</a>		3025	389		90	rest corepressor 1 (duto) mRNA.

<a href="#">duvar</a>	<a href="#">duvar.aSep08</a>		21135	1511	8	345	transmembrane protein 59 (duvar) alternative variant aSep08, mRNA.
<a href="#">duvar</a>	<a href="#">duvar.bSep08</a>		8623	862	3	154	transmembrane protein 59 (17.2 kD) (duvar) alternative variant bSep08, mRNA.
<a href="#">duvar</a>	<a href="#">duvar.cSep08</a>		3258	497	2	81	transmembrane protein 59 (duvar) alternative variant cSep08, mRNA.
<a href="#">duwey</a>	<a href="#">duwey.aSep08</a>		2859	1780		33	putative protein (3.6 kD) (duwey) mRNA.
<a href="#">Duxbl</a>	<a href="#">Duxbl.aSep08</a>	<a href="#">306226</a>	9305	1782	4	263	double homeobox B-like (Duxbl) alternative variant aSep08, mRNA.
<a href="#">Duxbl</a>	<a href="#">Duxbl.cSep08</a>	<a href="#">306226</a>	3792	447	2	104	double homeobox B-like (Duxbl) alternative variant cSep08, mRNA.
<a href="#">Dvl3</a>	<a href="#">Dvl3.bSep08</a>	<a href="#">303811</a>	489	295	1	97	dishevelled 3, dsh homolog (Drosophila) (Dvl3) alternative variant bSep08, mRNA.
<a href="#">DWNN.0</a>	<a href="#">DWNN.0.aSep08</a>		8501	1109		123	retinoblastoma binding protein 6 CRA g like (13.8 kD) (DWNN.0) mRNA.
<a href="#">dyby</a>	<a href="#">dyby.aSep08</a>		10330	514		171	dedicator of cytokinesis (dyby) mRNA.
<a href="#">dychy</a>	<a href="#">dychy.aSep08</a>		6392	429		83	putative protein (dychy) mRNA.
<a href="#">Dydc2</a>	<a href="#">Dydc2.aSep08</a>	<a href="#">688422</a>	13669	735	1	139	dpy-30 (15.7 kD) (Dydc2) alternative variant aSep08, mRNA.
<a href="#">dyfer</a>	<a href="#">dyfer.aSep08</a>		7957	377		58	putative protein (dyfer) mRNA.
<a href="#">dyflo</a>	<a href="#">dyflo.aSep08</a>		8589	309		103	putative protein (dyflo) mRNA.
<a href="#">dyflu</a>	<a href="#">dyflu.aSep08</a>		3702	773		246	sorting nexin 26 (dyflu) mRNA.
<a href="#">dykee</a>	<a href="#">dykee.aSep08</a>		2797	733		66	phosphatidylinositol glycan anchor biosynthesis class N (dykee) mRNA.
<a href="#">dyloy</a>	<a href="#">dyloy.aSep08</a>		59815	606		201	ABI gene family member 3 binding protein like (dyloy) mRNA.
<a href="#">Dym</a>	<a href="#">Dym.aSep08</a>	<a href="#">291433</a>	299222	2538	17	674	dymeclin (76.5 kD) (Dym) alternative variant aSep08, complete mRNA.
<a href="#">Dym</a>	<a href="#">Dym.cSep08</a>	<a href="#">291433</a>	48950	712	6	236	dymeclin (Dym) alternative variant cSep08, mRNA.
<a href="#">Dym</a>	<a href="#">Dym.dSep08</a>	<a href="#">291433</a>	72139	733	6	98	dymeclin (10.9 kD) (Dym) alternative variant dSep08, mRNA.
<a href="#">Dym</a>	<a href="#">Dym.eSep08</a>	<a href="#">291433</a>	49462	675	5	96	dymeclin (Dym) alternative variant eSep08, mRNA.
<a href="#">Dym</a>	<a href="#">Dym.fSep08</a>	<a href="#">291433</a>	108923	1777	4	53	dymeclin (Dym) alternative variant fSep08, mRNA.
<a href="#">dymer</a>	<a href="#">dymer.aSep08</a>		951	784	2	93	putative protein (dymer) mRNA.
<a href="#">Dync1i1</a>	<a href="#">Dync1i1.bSep08</a>	<a href="#">29564</a>	191899	690	4	182	dynein cytoplasmic 1 intermediate chain 1 (Dync1i1) alternative variant bSep08, mRNA.
<a href="#">Dync1i1</a>	<a href="#">Dync1i1.cSep08</a>	<a href="#">29564</a>	64996	578	3	146	dynein cytoplasmic 1 intermediate chain 1 (Dync1i1) alternative variant cSep08, mRNA.
<a href="#">Dync1i1</a>	<a href="#">Dync1i1.dSep08</a>	<a href="#">29564</a>	84135	542	2	141	dynein cytoplasmic 1 intermediate chain 1 (Dync1i1) alternative variant dSep08, mRNA.
<a href="#">Dync1i1</a>	<a href="#">Dync1i1.eSep08</a>	<a href="#">29564</a>	84042	500	2	137	dynein cytoplasmic 1 intermediate chain 1 (Dync1i1) alternative variant eSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.bSep08</a>	<a href="#">116659</a>	52471	2675	17	632	dynein cytoplasmic intermediate (70.6 kD) (Dync1i2) alternative variant bSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.cSep08</a>	<a href="#">116659</a>	34072	939	9	313	dynein cytoplasmic intermediate CRA c (Dync1i2) alternative variant cSep08, mRNA.

<a href="#">Dync1i2</a>	<a href="#">Dync1i2.dSep08</a>	<a href="#">116659</a>	35516	1045	9	289	dynein cytoplasmic intermediate CRA c (Dync1i2) alternative variant dSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.eSep08</a>	<a href="#">116659</a>	33293	874	8	255	dynein cytoplasmic intermediate CRA c (Dync1i2) alternative variant eSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.fSep08</a>	<a href="#">116659</a>	32810	669	7	180	cytoplasmic dynein intermediate (Dync1i2) alternative variant fSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.gSep08</a>	<a href="#">116659</a>	23383	503	7	167	dynein cytoplasmic intermediate (Dync1i2) alternative variant gSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.hSep08</a>	<a href="#">116659</a>	4878	1206	3	108	dynein cytoplasmic intermediate CRA a (11.8 kD) (Dync1i2) alternative variant hSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.iSep08</a>	<a href="#">116659</a>	2019	614	2	91	cytoplasmic dynein intermediate (Dync1i2) alternative variant iSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.jSep08</a>	<a href="#">116659</a>	2431	729	2	61	cytoplasmic dynein intermediate (Dync1i2) alternative variant jSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.kSep08</a>	<a href="#">116659</a>	32788	814	6	52	putative protein (6.4 kD) (Dync1i2) alternative variant kSep08, mRNA.
<a href="#">Dync1i2</a>	<a href="#">Dync1i2.mSep08</a>	<a href="#">116659</a>	4584	635	3	47	putative protein (Dync1i2) alternative variant mSep08, mRNA.
<a href="#">Dync1li1</a>	<a href="#">Dync1li1.cSep08</a>	<a href="#">252902</a>	1168	614	3	84	dynein cytoplasmic 1 light intermediate chain 1 (Dync1li1) alternative variant cSep08, mRNA.
<a href="#">Dync1li1</a>	<a href="#">Dync1li1.dSep08</a>	<a href="#">252902</a>	4695	1801	5	53	dynein cytoplasmic 1 light intermediate chain 1 (Dync1li1) alternative variant dSep08, mRNA.
<a href="#">Dync1li2</a>	<a href="#">Dync1li2.aSep08</a>	<a href="#">81655</a>	38981	1783	13	520	dynein, cytoplasmic 1 light intermediate chain 2 (Dync1li2) alternative variant aSep08, mRNA.
<a href="#">Dync1li2</a>	<a href="#">Dync1li2.cSep08</a>	<a href="#">81655</a>	4784	372	4	55	dynein, cytoplasmic 1 light intermediate chain 2 (Dync1li2) alternative variant cSep08, mRNA.
<a href="#">Dync1li2</a>	<a href="#">Dync1li2.dSep08</a>	<a href="#">81655</a>	2675	1086	2	62	dynein, cytoplasmic 1 light intermediate chain 2 (Dync1li2) alternative variant dSep08, mRNA.
<a href="#">Dync1li2</a>	<a href="#">Dync1li2.eSep08</a>	<a href="#">81655</a>	2702	415	3	13	dynein, cytoplasmic 1 light intermediate chain 2 (Dync1li2) alternative variant eSep08, mRNA.
<a href="#">Dync2h1</a>	<a href="#">Dync2h1.aSep08</a>	<a href="#">65209</a>	24101	1109		369	dynein cytoplasmic 2 heavy chain 1 (Dync2h1) mRNA.
<a href="#">Dync2li1</a>	<a href="#">Dync2li1.bSep08</a>	<a href="#">298767</a>	14353	815	8	197	dynein cytoplasmic 2 light intermediate chain 1 (Dync2li1) alternative variant bSep08, mRNA.
<a href="#">Dync2li1</a>	<a href="#">Dync2li1.dSep08</a>	<a href="#">298767</a>	1236	420	2	68	dynein cytoplasmic 2 light intermediate chain 1 (Dync2li1) alternative variant dSep08, mRNA.
<a href="#">Dyntl1</a>	<a href="#">Dyntl1.aSep08</a>	<a href="#">58945</a>	2291	611	2	89	dynein light chain LC8-type 1 (10.4 kD) (Dyntl1) alternative variant aSep08, complete mRNA.
<a href="#">Dyntl2</a>	<a href="#">Dyntl2.bSep08</a>	<a href="#">140734</a>	14371	157	1	31	dynein light chain LC8-type 2 (Dyntl2) alternative variant bSep08, mRNA.
<a href="#">Dynlrb1</a>	<a href="#">Dynlrb1.bSep08</a>	<a href="#">170714</a>	11241	647	3	63	dynein light chain roadblock-type 1 (7.4 kD) (Dynlrb1) alternative variant bSep08, mRNA.
<a href="#">Dynlrb1</a>	<a href="#">Dynlrb1.cSep08</a>	<a href="#">170714</a>	10987	593	4	63	dynein light chain roadblock-type 1 (7.4 kD) (Dynlrb1) alternative variant cSep08, mRNA.
<a href="#">Dynlrb2</a>	<a href="#">Dynlrb2.bSep08</a>	<a href="#">361415</a>	3139	419	2	55	dynein light chain roadblock-type 2 (6.3 kD) (Dynlrb2) alternative variant bSep08, mRNA.
<a href="#">Dyntl1</a>	<a href="#">Dyntl1.aSep08</a>	<a href="#">83462</a>	6186	466	4	119	dynein light chain Tctex-type 1 (13.0 kD) (Dyntl1) alternative variant aSep08, mRNA.

<a href="#">Dyntl3</a>	<a href="#">Dyntl3.bSep08</a>	<a href="#">363448</a>	2845	2231	1	48	dynein light chain Tctex-type 3 (Dyntl3) alternative variant bSep08, mRNA.
<a href="#">dynoy</a>	<a href="#">dynoy.aSep08</a>		1239	586		29	putative protein (3.1 kD) (dynoy) mRNA.
<a href="#">dypor</a>	<a href="#">dypor.aSep08</a>		28977	570		82	putative protein (dypor) mRNA.
<a href="#">Dyrk1a</a>	<a href="#">Dyrk1a.bSep08</a>	<a href="#">25255</a>	50260	392	1	105	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A (Dyrk1a) alternative variant bSep08, mRNA.
<a href="#">Dyrk1b</a>	<a href="#">Dyrk1b.bSep08</a>	<a href="#">308468</a>	852	425	3	141	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1b (Dyrk1b) alternative variant bSep08, mRNA.
<a href="#">Dyrk1b</a>	<a href="#">Dyrk1b.cSep08</a>	<a href="#">308468</a>	1345	568	3	111	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1b (Dyrk1b) alternative variant cSep08, mRNA.
<a href="#">Dyrk3</a>	<a href="#">Dyrk3.aSep08</a>	<a href="#">304775</a>	7654	1012		337	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3 (Dyrk3) mRNA.
<a href="#">dysa</a>	<a href="#">dysa.aSep08</a>		1814	505		36	putative protein (dysa) mRNA.
<a href="#">Dysf</a>	<a href="#">Dysf.aSep08</a>	<a href="#">312492</a>	60139	2223	16	665	dysferlin (Dysf) alternative variant aSep08, mRNA.
<a href="#">Dysf</a>	<a href="#">Dysf.bSep08</a>	<a href="#">312492</a>	1383	290	1	75	dysferlin (Dysf) alternative variant bSep08, mRNA.
<a href="#">dyshee</a>	<a href="#">dyshee.aSep08</a>		15294	422		34	putative protein (dyshee) mRNA.
<a href="#">Dyt1</a>	<a href="#">Dyt1.bSep08</a>	<a href="#">266606</a>	5989	712	4	139	dystonia 1 (Dyt1) alternative variant bSep08, mRNA.
<a href="#">dyto</a>	<a href="#">dyto.aSep08</a>		2159	688	4	112	putative protein (12.7 kD) (dyto) alternative variant aSep08, mRNA.
<a href="#">dyvar</a>	<a href="#">dyvar.aSep08</a>		5137	754		37	putative protein (4.5 kD) (dyvar) mRNA.
<a href="#">dywey</a>	<a href="#">dywey.aSep08</a>		3166	750		69	putative protein (dywey) mRNA.
<a href="#">Dzip1</a>	<a href="#">Dzip1.aSep08</a>	<a href="#">364475</a>	68419	3803	8	875	zinc finger protein dzip1 (100.2 kD) (Dzip1) alternative variant aSep08, complete mRNA.
<a href="#">Dzip1</a>	<a href="#">Dzip1.bSep08</a>	<a href="#">364475</a>	2974	1165	5	186	putative protein (Dzip1) alternative variant bSep08, mRNA.
<a href="#">Dzip1</a>	<a href="#">Dzip1.cSep08</a>	<a href="#">364475</a>	7375	668	2	169	DAZ interacting protein 1 (Dzip1) alternative variant cSep08, mRNA.
<a href="#">Dzip1</a>	<a href="#">Dzip1.dSep08</a>	<a href="#">364475</a>	591	192	1	51	DAZ interacting protein 1 (Dzip1) alternative variant dSep08, mRNA.
<a href="#">Dzip1l</a>	<a href="#">Dzip1l.bSep08</a>	<a href="#">315952</a>	5817	683	1	227	DAZ interacting protein 1-like (Dzip1l) alternative variant bSep08, mRNA.
<a href="#">E2f1</a>	<a href="#">E2f1.bSep08</a>	<a href="#">399489</a>	3793	1039	5	346	E2F transcription factor 1 (E2f1) alternative variant bSep08, mRNA.
<a href="#">E2f1</a>	<a href="#">E2f1.cSep08</a>	<a href="#">399489</a>	3313	654	4	217	E2F transcription factor 1 (E2f1) alternative variant cSep08, mRNA.
<a href="#">E2f1</a>	<a href="#">E2f1.eSep08</a>	<a href="#">399489</a>	1564	382	2	127	E2F transcription factor 1 (E2f1) alternative variant eSep08, mRNA.
<a href="#">E2f6</a>	<a href="#">E2f6.aSep08</a>	<a href="#">313978</a>	15587	1395	7	203	E2F transcription factor 6 (E2f6) alternative variant aSep08, mRNA.
<a href="#">E2f6</a>	<a href="#">E2f6.bSep08</a>	<a href="#">313978</a>	2213	1539	2	117	E2F transcription factor 6 (13.1 kD) (E2f6) alternative variant bSep08, mRNA.
<a href="#">E2f6</a>	<a href="#">E2f6.cSep08</a>	<a href="#">313978</a>	16352	338	3	50	E2F transcription factor 6 (5.6 kD) (E2f6) alternative variant cSep08, mRNA.
<a href="#">E2f7</a>	<a href="#">E2f7.bSep08</a>	<a href="#">314818</a>	8236	674	3	142	E2F transcription factor 7 (16.1 kD) (E2f7) alternative variant bSep08, mRNA.
<a href="#">E2f8</a>	<a href="#">E2f8.aSep08</a>	<a href="#">308607</a>	3354	772		257	E2F transcription factor 8 (E2f8) mRNA.



<a href="#">E2F_TDP.0</a>	<a href="#">E2F_TDP.0.aSep08</a>		7400	2023	1	412	E2F transcription factor 4 (43.9 kD) (E2F_TDP.0) alternative variant aSep08, mRNA.
<a href="#">E2F_TDP.0</a>	<a href="#">E2F_TDP.0.bSep08</a>		5853	904	1	301	E2F transcription factor 4 (E2F_TDP.0) alternative variant bSep08, mRNA.
<a href="#">E2F_TDP.1</a>	<a href="#">E2F_TDP.1.aSep08</a>		4195	764		254	e2f transcription factor 8 (E2F_TDP.1) mRNA.
<a href="#">E4f1</a>	<a href="#">E4f1.aSep08</a>	<a href="#">681359</a>	11400	2519	13	401	transcription factor (E4f1) alternative variant aSep08, mRNA.
<a href="#">E4f1</a>	<a href="#">E4f1.bSep08</a>	<a href="#">681359</a>	3497	2016	10	393	transcription factor (42.8 kD) (E4f1) alternative variant bSep08, mRNA.
<a href="#">E4f1</a>	<a href="#">E4f1.cSep08</a>	<a href="#">681359</a>	2344	874	6	291	transcription factor (E4f1) alternative variant cSep08, mRNA.
<a href="#">E4f1</a>	<a href="#">E4f1.dSep08</a>	<a href="#">681359</a>	958	813	3	181	transcription factor (18.9 kD) (E4f1) alternative variant dSep08, mRNA.
<a href="#">E4f1</a>	<a href="#">E4f1.eSep08</a>	<a href="#">681359</a>	874	703	2	137	transcription factor (E4f1) alternative variant eSep08, mRNA.
<a href="#">E030032D13Rik</a>	<a href="#">E030032D13Rik.aSep08</a>	<a href="#">415052</a>	734	367	3	82	E030032D13Rik gene (E030032D13Rik) alternative variant aSep08, mRNA.
<a href="#">E230034O05Rik</a>	<a href="#">E230034O05Rik.aSep08</a>	<a href="#">415065</a>	2158	1470		63	E230034O05Rik gene (E230034O05Rik) mRNA.
<a href="#">Eapa2</a>	<a href="#">Eapa2.bSep08</a>	<a href="#">494321</a>	1596	314	2	104	experimental autoimmune prostatitis antigen 2 (Eapa2) alternative variant bSep08, mRNA.
<a href="#">Eapa2</a>	<a href="#">Eapa2.cSep08</a>	<a href="#">494321</a>	2649	757	2	103	experimental autoimmune prostatitis antigen 2 (Eapa2) alternative variant cSep08, mRNA.
<a href="#">Eapp</a>	<a href="#">Eapp.aSep08</a>	<a href="#">299043</a>	27216	1418	7	280	E2F-associated phosphoprotein (32.2 kD) (Eapp) alternative variant aSep08, mRNA.
<a href="#">Eapp</a>	<a href="#">Eapp.bSep08</a>	<a href="#">299043</a>	14318	2450	4	190	E2F-associated phosphoprotein (Eapp) alternative variant bSep08, mRNA.
<a href="#">Eapp</a>	<a href="#">Eapp.dSep08</a>	<a href="#">299043</a>	32361	705	3	113	E2F-associated phosphoprotein (Eapp) alternative variant dSep08, mRNA.
<a href="#">Ears2</a>	<a href="#">Ears2.aSep08</a>	<a href="#">361641</a>	21688	1785		550	glutamyl-tRNA synthetase 2 mitochondrial (putative) (Ears2) mRNA.
<a href="#">Ebag9</a>	<a href="#">Ebag9.aSep08</a>	<a href="#">299864</a>	17625	1783	2	361	estrogen receptor-binding fragment-associated gene 9 (Ebag9) alternative variant aSep08, mRNA.
<a href="#">Ebag9</a>	<a href="#">Ebag9.cSep08</a>	<a href="#">299864</a>	9989	508	1	143	estrogen receptor-binding fragment-associated gene 9 (Ebag9) alternative variant cSep08, mRNA.
<a href="#">Ebf1</a>	<a href="#">Ebf1.bSep08</a>	<a href="#">116543</a>	379021	1530	14	448	early B-cell factor 1 (Ebf1) alternative variant bSep08, mRNA.
<a href="#">Ebf1</a>	<a href="#">Ebf1.cSep08</a>	<a href="#">116543</a>	9020	1077	2	68	early B-cell factor 1 (6.9 kD) (Ebf1) alternative variant cSep08, mRNA.
<a href="#">Ebf3</a>	<a href="#">Ebf3.bSep08</a>	<a href="#">361668</a>	2236	1089	4	152	early B-cell factor 3 (17.4 kD) (Ebf3) alternative variant bSep08, mRNA.
<a href="#">Ebf4</a>	<a href="#">Ebf4.aSep08</a>	<a href="#">680751</a>	5512	1393	1	210	early B-cell factor 4 (10.9 kD) (Ebf4) alternative variant aSep08, mRNA.
<a href="#">Ebf4</a>	<a href="#">Ebf4.bSep08</a>	<a href="#">680751</a>	8195	860	3	124	early B-cell factor 4 (Ebf4) alternative variant bSep08, mRNA.
<a href="#">Ebf4</a>	<a href="#">Ebf4.cSep08</a>	<a href="#">680751</a>	7534	400	2	121	early B-cell factor 4 (Ebf4) alternative variant cSep08, mRNA.

<a href="#">Ebp</a>	<a href="#">Ebp.bSep08</a>	<a href="#">117278</a>	5889	563	5	187	phenylalkylamine Ca2+ antagonist (emopamil) binding protein (Ebp) alternative variant bSep08, mRNA.
<a href="#">Ebp</a>	<a href="#">Ebp.cSep08</a>	<a href="#">117278</a>	6339	428	2	54	phenylalkylamine Ca2+ antagonist (emopamil) binding protein (Ebp) alternative variant cSep08, mRNA.
<a href="#">Ebpl</a>	<a href="#">Ebpl.aSep08</a>	<a href="#">361054</a>	21988	1118	3	211	emopamil binding protein-like (Ebpl) alternative variant aSep08, mRNA.
<a href="#">Ebpl</a>	<a href="#">Ebpl.bSep08</a>	<a href="#">361054</a>	1140	319	1	45	emopamil binding protein-like (Ebpl) alternative variant bSep08, mRNA.
<a href="#">Ece1</a>	<a href="#">Ece1.bSep08</a>	<a href="#">94204</a>	5980	372	2	123	endothelin converting enzyme 1 (Ece1) alternative variant bSep08, mRNA.
<a href="#">Ece2</a>	<a href="#">Ece2.bSep08</a>	<a href="#">408243</a>	15717	2021	6	258	endothelin-converting enzyme 2 (29.3 kD) (Ece2) alternative variant bSep08, mRNA.
<a href="#">Ece2</a>	<a href="#">Ece2.cSep08</a>	<a href="#">408243</a>	4899	742	7	247	endothelin-converting enzyme 2 (Ece2) alternative variant cSep08, mRNA.
<a href="#">Ecel1</a>	<a href="#">Ecel1.bSep08</a>	<a href="#">60417</a>	6221	1796	5	598	endothelin converting enzyme-like 1 (Ecel1) alternative variant bSep08, mRNA.
<a href="#">ECH.0</a>	<a href="#">ECH.0.aSep08</a>	<a href="#">500621</a>	17657	713	6	225	mitochondrial trifunctional protein (ECH.0) alternative variant aSep08, mRNA.
<a href="#">ECH.0</a>	<a href="#">ECH.0.bSep08</a>	<a href="#">500621</a>	3708	349	1	48	putative protein (ECH.0) alternative variant bSep08, mRNA.
<a href="#">Ech1</a>	<a href="#">Ech1.bSep08</a>	<a href="#">64526</a>	4385	701	4	226	enoyl coenzyme A hydratase 1, peroxisomal (25.1 kD) (Ech1) alternative variant bSep08, mRNA.
<a href="#">Ech1</a>	<a href="#">Ech1.cSep08</a>	<a href="#">64526</a>	4891	742	5	194	enoyl coenzyme A hydratase 1, peroxisomal (21.8 kD) (Ech1) alternative variant cSep08, mRNA.
<a href="#">Ech1</a>	<a href="#">Ech1.dSep08</a>	<a href="#">64526</a>	4328	835	3	170	enoyl coenzyme A hydratase 1, peroxisomal (19.0 kD) (Ech1) alternative variant dSep08, mRNA.
<a href="#">Ech1</a>	<a href="#">Ech1.eSep08</a>	<a href="#">64526</a>	3790	483	1	92	enoyl coenzyme A hydratase 1, peroxisomal (10.6 kD) (Ech1) alternative variant eSep08, mRNA.
<a href="#">Echdc1</a>	<a href="#">Echdc1.bSep08</a>	<a href="#">361465</a>	29437	669	2	223	enoyl-CoA hydratase/isomerase (Echdc1) alternative variant bSep08, mRNA.
<a href="#">Echdc1</a>	<a href="#">Echdc1.cSep08</a>	<a href="#">361465</a>	29174	741	2	218	enoyl-CoA hydratase/isomerase (Echdc1) alternative variant cSep08, mRNA.
<a href="#">Echdc2</a>	<a href="#">Echdc2.bSep08</a>	<a href="#">298381</a>	15108	946	2	191	enoyl-CoA hydratase/isomerase (20.1 kD) (Echdc2) alternative variant bSep08, mRNA.
<a href="#">Ecm1</a>	<a href="#">Ecm1.bSep08</a>	<a href="#">116662</a>	5163	1506	8	436	extracellular matrix protein 1 (48.7 kD) (Ecm1) alternative variant bSep08, complete mRNA.
<a href="#">Ecm1</a>	<a href="#">Ecm1.cSep08</a>	<a href="#">116662</a>	2701	731	7	243	extracellular matrix protein 1 (Ecm1) alternative variant cSep08, mRNA.
<a href="#">Ecm1</a>	<a href="#">Ecm1.dSep08</a>	<a href="#">116662</a>	1130	514	1	66	extracellular matrix protein 1 (Ecm1) alternative variant dSep08, mRNA.
<a href="#">Ecop</a>	<a href="#">Ecop.bSep08</a>	<a href="#">362374</a>	68778	919		96	EGFR-coamplified and overexpressed protein (Ecop) alternative variant bSep08, mRNA.
<a href="#">Ecsit</a>	<a href="#">Ecsit.bSep08</a>	<a href="#">300447</a>	3709	724	5	240	ECSIT homolog (Drosophila) (Ecsit) alternative variant bSep08, mRNA.
<a href="#">Ecsit</a>	<a href="#">Ecsit.cSep08</a>	<a href="#">300447</a>	880	469	2	106	ECSIT homolog (Drosophila) (11.8 kD) (Ecsit) alternative variant cSep08, mRNA.
<a href="#">Ect2</a>	<a href="#">Ect2.bSep08</a>	<a href="#">361921</a>	30506	1275	5	241	ect2 oncogene (Ect2) alternative variant bSep08, mRNA.

<a href="#">Ect2</a>	<a href="#">Ect2.cSep08</a>	<a href="#">361921</a>	31589	1757	4	196	ect2 oncogene (Ect2) alternative variant cSep08, mRNA.
<a href="#">Ect2</a>	<a href="#">Ect2.dSep08</a>	<a href="#">361921</a>	5167	412	2	88	ect2 oncogene (Ect2) alternative variant dSep08, mRNA.
<a href="#">Edaradd</a>	<a href="#">Edaradd.aSep08</a>	<a href="#">498769</a>	1032	522	1	108	EDAR (ectodysplasin-A receptor)-associated death domain (Edaradd) alternative variant aSep08, mRNA.
<a href="#">Edaradd</a>	<a href="#">Edaradd.bSep08</a>	<a href="#">498769</a>	660	472	1	74	EDAR (ectodysplasin-A receptor)-associated death domain (8.7 kD) (Edaradd) alternative variant bSep08, mRNA.
<a href="#">Edem2</a>	<a href="#">Edem2.bSep08</a>	<a href="#">296304</a>	9401	747	5	184	ER degradation enhancer mannosidase alpha-like 2 (Edem2) alternative variant bSep08, mRNA.
<a href="#">Edem2</a>	<a href="#">Edem2.cSep08</a>	<a href="#">296304</a>	9977	436	4	89	ER degradation enhancer mannosidase alpha-like 2 (Edem2) alternative variant cSep08, mRNA.
<a href="#">Edem2</a>	<a href="#">Edem2.dSep08</a>	<a href="#">296304</a>	10292	416	4	87	ER degradation enhancer mannosidase alpha-like 2 precursor (9.8 kD) (Edem2) alternative variant dSep08, mRNA.
<a href="#">Edem3</a>	<a href="#">Edem3.aSep08</a>	<a href="#">289085</a>	28747	2007		483	ER degradation enhancer, mannosidase alpha-like 3 (Edem3) alternative variant aSep08, mRNA.
<a href="#">Edem3</a>	<a href="#">Edem3.bSep08</a>	<a href="#">289085</a>	2744	373		124	ER degradation enhancer, mannosidase alpha-like 3 (Edem3) alternative variant bSep08, mRNA.
<a href="#">Edf1</a>	<a href="#">Edf1.cSep08</a>	<a href="#">296570</a>	1085	698	2	71	endothelial differentiation-related factor 1 (Edf1) alternative variant cSep08, mRNA.
<a href="#">Edn3</a>	<a href="#">Edn3.bSep08</a>	<a href="#">366270</a>	4921	326	1	89	endothelin 3 (Edn3) alternative variant bSep08, mRNA.
<a href="#">Ednra</a>	<a href="#">Ednra.bSep08</a>	<a href="#">24326</a>	60855	3684	3	399	endothelin receptor type A (45.3 kD) (Ednra) alternative variant bSep08, complete mRNA.
<a href="#">Ednra</a>	<a href="#">Ednra.cSep08</a>	<a href="#">24326</a>	58930	1487	4	267	endothelin receptor type A (30.9 kD) (Ednra) alternative variant cSep08, mRNA.
<a href="#">Ednra</a>	<a href="#">Ednra.dSep08</a>	<a href="#">24326</a>	58930	1429	4	259	endothelin receptor type A (28.5 kD) (Ednra) alternative variant dSep08, mRNA.
<a href="#">Ednra</a>	<a href="#">Ednra.eSep08</a>	<a href="#">24326</a>	59130	1427	2	247	endothelin receptor type A (28.8 kD) (Ednra) alternative variant eSep08, mRNA.
<a href="#">Ednra</a>	<a href="#">Ednra.fSep08</a>	<a href="#">24326</a>	26100	697	2	195	endothelin receptor type A (Ednra) alternative variant fSep08, mRNA.
<a href="#">Eea1</a>	<a href="#">Eea1.bSep08</a>	<a href="#">314764</a>	15496	457	4	123	early endosome antigen 1 162kD CRA b like (Eea1) alternative variant bSep08, mRNA.
<a href="#">Eea1</a>	<a href="#">Eea1.cSep08</a>	<a href="#">314764</a>	8603	468	2	81	early endosome antigen 1 CRA a like (Eea1) alternative variant cSep08, mRNA.
<a href="#">Eed</a>	<a href="#">Eed.bSep08</a>	<a href="#">293104</a>	1489	429	3	66	embryonic ectoderm development (7.5 kD) (Eed) alternative variant bSep08, mRNA.
<a href="#">Eed</a>	<a href="#">Eed.cSep08</a>	<a href="#">293104</a>	1335	978	2	66	embryonic ectoderm development (7.5 kD) (Eed) alternative variant cSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.aSep08</a>	<a href="#">29652</a>	5893	4400	8	462	eukaryotic translation elongation factor 1 alpha 1 (50.1 kD) (Eef1a1.1) alternative variant aSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.aSep08</a>	<a href="#">171361</a>	5893	4400	8	462	eukaryotic translation elongation factor 1 alpha 1 (50.1 kD) (Eef1a1.1) alternative variant aSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.bSep08</a>	<a href="#">29652</a>	2289	740	5	246	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant bSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.bSep08</a>	<a href="#">171361</a>	2289	740	5	246	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant bSep08, mRNA.

<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.cSep08</a>	<a href="#">29652</a>	2426	1313	4	232	eukaryotic translation elongation factor 1 alpha 1 (25.6 kD) (Eef1a1.1) alternative variant cSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.cSep08</a>	<a href="#">171361</a>	2426	1313	4	232	eukaryotic translation elongation factor 1 alpha 1 (25.6 kD) (Eef1a1.1) alternative variant cSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.dSep08</a>	<a href="#">29652</a>	1769	679	4	197	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant dSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.dSep08</a>	<a href="#">171361</a>	1769	679	4	197	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant dSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.fSep08</a>	<a href="#">29652</a>	1644	1172	4	156	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant fSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.fSep08</a>	<a href="#">171361</a>	1644	1172	4	156	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant fSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.gSep08</a>	<a href="#">29652</a>	760	658	2	124	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant gSep08, mRNA.
<a href="#">Eef1a1.1</a>	<a href="#">Eef1a1.1.gSep08</a>	<a href="#">171361</a>	760	658	2	124	eukaryotic translation elongation factor 1 alpha 1 (Eef1a1.1) alternative variant gSep08, mRNA.
<a href="#">Eef1a2</a>	<a href="#">Eef1a2.bSep08</a>	<a href="#">24799</a>	7646	444	3	133	eukaryotic translation elongation factor 1 alpha 2 (Eef1a2) alternative variant bSep08, mRNA.
<a href="#">Eef1a2</a>	<a href="#">Eef1a2.cSep08</a>	<a href="#">24799</a>	1818	384	2	127	eukaryotic translation elongation factor 1 alpha 2 (Eef1a2) alternative variant cSep08, mRNA.
<a href="#">Eef1b2</a>	<a href="#">Eef1b2.bSep08</a>	<a href="#">363241</a>	2461	1266	3	157	eukaryotic translation elongation factor 1 beta 2 (17.2 kD) (Eef1b2) alternative variant bSep08, complete mRNA.
<a href="#">Eef1b2</a>	<a href="#">Eef1b2.cSep08</a>	<a href="#">363241</a>	2382	1224	2	141	eukaryotic translation elongation factor 1 beta 2 (Eef1b2) alternative variant cSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.bSep08</a>	<a href="#">300033</a>	2622	1162	2	387	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant bSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.cSep08</a>	<a href="#">300033</a>	15024	1142	8	314	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant cSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.dSep08</a>	<a href="#">300033</a>	13864	716	7	223	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant dSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.eSep08</a>	<a href="#">300033</a>	14150	730	6	204	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant eSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.fSep08</a>	<a href="#">300033</a>	14448	648	7	195	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant fSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.gSep08</a>	<a href="#">300033</a>	14529	589	5	195	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant gSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.hSep08</a>	<a href="#">300033</a>	13810	590	6	177	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant hSep08, mRNA.

<a href="#">Eef1d</a>	<a href="#">Eef1d.iSep08</a>	<a href="#">300033</a>	1635	879	4	147	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (16.6 kD) (Eef1d) alternative variant iSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.jSep08</a>	<a href="#">300033</a>	13951	459	5	125	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant jSep08, mRNA.
<a href="#">Eef1d</a>	<a href="#">Eef1d.kSep08</a>	<a href="#">300033</a>	13679	402	5	118	eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (Eef1d) alternative variant kSep08, mRNA.
<a href="#">Eef1g</a>	<a href="#">Eef1g.bSep08</a>	<a href="#">293725</a>	1631	1355	2	136	eukaryotic translation elongation factor 1 gamma (Eef1g) alternative variant bSep08, mRNA.
<a href="#">Eef1g</a>	<a href="#">Eef1g.cSep08</a>	<a href="#">293725</a>	1105	766	2	82	elongation factor 1 gamma (Eef1g) alternative variant cSep08, mRNA.
<a href="#">Eef2</a>	<a href="#">Eef2.bSep08</a>	<a href="#">29565</a>	3702	2864	8	703	eukaryotic translation elongation factor 2 (78.4 kD) (Eef2) alternative variant bSep08, mRNA.
<a href="#">Eef2</a>	<a href="#">Eef2.cSep08</a>	<a href="#">29565</a>	4122	3231	6	316	eukaryotic translation elongation factor 2 (35.2 kD) (Eef2) alternative variant cSep08, mRNA.
<a href="#">Eef2</a>	<a href="#">Eef2.dSep08</a>	<a href="#">29565</a>	3972	768	6	253	eukaryotic translation elongation factor 2 (Eef2) alternative variant dSep08, mRNA.
<a href="#">Eef2k</a>	<a href="#">Eef2k.bSep08</a>	<a href="#">25435</a>	1701	1303	2	84	eukaryotic elongation factor-2 kinase (9.0 kD) (Eef2k) alternative variant bSep08, mRNA.
<a href="#">Eefsec</a>	<a href="#">Eefsec.aSep08</a>	<a href="#">500255</a>	198640	2116	7	589	eukaryotic elongation factor, selenocysteine-tRNA-specific (Eefsec) alternative variant aSep08, mRNA.
<a href="#">Efcab2</a>	<a href="#">Efcab2.bSep08</a>	<a href="#">289280</a>	1538	472	4	63	EF-hand calcium binding domain 2 (Efcab2) alternative variant bSep08, mRNA.
<a href="#">Efcab3</a>	<a href="#">Efcab3.bSep08</a>	<a href="#">303589</a>	14662	626		50	EF-hand calcium binding domain 3 (Efcab3) alternative variant bSep08, mRNA.
<a href="#">Efcab4a</a>	<a href="#">Efcab4a.bSep08</a>	<a href="#">309112</a>	1886	1083	4	261	EF-hand calcium binding domain 4A (Efcab4a) alternative variant bSep08, mRNA.
<a href="#">Efcab4a</a>	<a href="#">Efcab4a.cSep08</a>	<a href="#">309112</a>	1608	1509	2	236	EF-hand calcium binding domain 4A (25.7 kD) (Efcab4a) alternative variant cSep08, mRNA.
<a href="#">Efcab4a</a>	<a href="#">Efcab4a.dSep08</a>	<a href="#">309112</a>	3981	750	6	224	EF-hand calcium binding domain 4A (Efcab4a) alternative variant dSep08, mRNA.
<a href="#">Efcab4a</a>	<a href="#">Efcab4a.eSep08</a>	<a href="#">309112</a>	3966	748	6	224	EF-hand calcium binding domain 4A (Efcab4a) alternative variant eSep08, mRNA.
<a href="#">Efcab5</a>	<a href="#">Efcab5.aSep08</a>	<a href="#">363653</a>	13637	648		215	EF-hand calcium binding domain 5 (Efcab5) mRNA.
<a href="#">Efcab6</a>	<a href="#">Efcab6.aSep08</a>	<a href="#">315179</a>	42686	1406	9	468	EF-hand calcium binding domain 6 (Efcab6) alternative variant aSep08, mRNA.
<a href="#">Efcab6</a>	<a href="#">Efcab6.bSep08</a>	<a href="#">315179</a>	22452	763	6	254	EF-hand calcium binding domain 6 (Efcab6) alternative variant bSep08, mRNA.
<a href="#">Efcab6</a>	<a href="#">Efcab6.cSep08</a>	<a href="#">315179</a>	22155	759	6	253	EF-hand calcium binding domain 6 (Efcab6) alternative variant cSep08, mRNA.
<a href="#">Efcab6</a>	<a href="#">Efcab6.dSep08</a>	<a href="#">315179</a>	13120	734	4	244	EF-hand calcium binding domain 6 (Efcab6) alternative variant dSep08, mRNA.
<a href="#">Efcab6</a>	<a href="#">Efcab6.eSep08</a>	<a href="#">315179</a>	4627	697	3	77	EF-hand calcium binding domain 6 (Efcab6) alternative variant eSep08, mRNA.
<a href="#">Efcab7</a>	<a href="#">Efcab7.aSep08</a>	<a href="#">362549</a>	15666	730		211	EF-hand calcium binding domain 7 (Efcab7) mRNA.

<a href="#">Efemp1</a>	<a href="#">Efemp1.bSep08</a>	<a href="#">305604</a>	84964	1784	8	451	epidermal growth factor-containing fibulin-like extracellular matrix protein 1 (Efemp1) alternative variant bSep08, mRNA.
<a href="#">Efemp1</a>	<a href="#">Efemp1.cSep08</a>	<a href="#">305604</a>	75238	779	5	259	epidermal growth factor-containing fibulin-like extracellular matrix protein 1 (Efemp1) alternative variant cSep08, mRNA.
<a href="#">Efemp1</a>	<a href="#">Efemp1.dSep08</a>	<a href="#">305604</a>	19339	645	6	187	epidermal growth factor-containing fibulin-like extracellular matrix protein 1 (Efemp1) alternative variant dSep08, mRNA.
<a href="#">Efemp1</a>	<a href="#">Efemp1.eSep08</a>	<a href="#">305604</a>	343	257	2	56	epidermal growth factor-containing fibulin-like extracellular matrix protein 1 (Efemp1) alternative variant eSep08, mRNA.
<a href="#">Efemp1</a>	<a href="#">Efemp1.fSep08</a>	<a href="#">305604</a>	10472	699	4	172	epidermal growth factor-containing fibulin-like extracellular matrix protein 1 (Efemp1) alternative variant fSep08, mRNA.
<a href="#">Efemp2</a>	<a href="#">Efemp2.bSep08</a>	<a href="#">293677</a>	5875	869	8	275	EGF-containing fibulin-like extracellular matrix protein 2 (Efemp2) alternative variant bSep08, mRNA.
<a href="#">Efemp2</a>	<a href="#">Efemp2.cSep08</a>	<a href="#">293677</a>	6752	1325	12	255	EGF-containing fibulin-like extracellular matrix protein 2 (Efemp2) alternative variant cSep08, mRNA.
<a href="#">Efemp2</a>	<a href="#">Efemp2.dSep08</a>	<a href="#">293677</a>	1207	688	3	215	EGF-containing fibulin-like extracellular matrix protein 2 (Efemp2) alternative variant dSep08, mRNA.
<a href="#">Efha1</a>	<a href="#">Efha1.bSep08</a>	<a href="#">171433</a>	70876	623	8	207	ef-hand domain family member A1 (Efha1) alternative variant bSep08, mRNA.
<a href="#">Efha1</a>	<a href="#">Efha1.cSep08</a>	<a href="#">171433</a>	39724	742	8	188	ef-hand domain family member A1 (Efha1) alternative variant cSep08, mRNA.
<a href="#">Efha1</a>	<a href="#">Efha1.dSep08</a>	<a href="#">171433</a>	28114	863	5	103	ef-hand domain family member A1 (Efha1) alternative variant dSep08, mRNA.
<a href="#">Efha1</a>	<a href="#">Efha1.eSep08</a>	<a href="#">171433</a>	1048	672	2	48	putative protein of bilateral origin (Efha1) alternative variant eSep08, mRNA.
<a href="#">Efha2</a>	<a href="#">Efha2.aSep08</a>	<a href="#">364601</a>	34030	3716	8	261	EF hand domain family, member A2 (Efha2) alternative variant aSep08, mRNA.
<a href="#">efhand.0</a>	<a href="#">efhand.0.aSep08</a>		21021	1104		111	calcium binding atopy-related autoantigen 1 like (efhand.0) mRNA.
<a href="#">efhand.1</a>	<a href="#">efhand.1.aSep08</a>		8556	652		122	lysophosphatidylcholine acyltransferase 2 (efhand.1) mRNA.
<a href="#">efhand.2</a>	<a href="#">efhand.2.aSep08</a>		30699	1372		454	intersectin 1 (efhand.2) mRNA.
<a href="#">efhand.3</a>	<a href="#">efhand.3.aSep08</a>		14247	542		94	interacting protein 3 (efhand.3) mRNA.
<a href="#">efhand.4</a>	<a href="#">efhand.4.aSep08</a>		7733	1022	4	230	myosin light chain regulatory B-like (efhand.4) alternative variant aSep08, mRNA.
<a href="#">efhand.4</a>	<a href="#">efhand.4.bSep08</a>		6181	1202	3	119	myosin light (13.5 kD) (efhand.4) alternative variant bSep08, mRNA.
<a href="#">efhand.4</a>	<a href="#">efhand.4.cSep08</a>		4807	356	2	118	myosin light (efhand.4) alternative variant cSep08, mRNA.
<a href="#">efhand.6</a>	<a href="#">efhand.6.aSep08</a>		11335	688		218	intersectin 2 (efhand.6) mRNA.
<a href="#">efhand_1.0</a>	<a href="#">efhand_1.0.aSep08</a>		125412	630	6	164	dystrobrevin (efhand_1.0) alternative variant aSep08, mRNA.
<a href="#">efhand_2.0</a>	<a href="#">efhand_2.0.aSep08</a>		48866	3136	12	326	utrophin CRA a (37.1 kD) (efhand_2.0) alternative variant aSep08, mRNA.

<a href="#">Efhb</a>	<a href="#">Efhb.bSep08</a>	<a href="#">301168</a>	71886	3163	1	768	EF hand domain family, member B (86.0 kD) (Efhb) alternative variant bSep08, complete mRNA.
<a href="#">Efhc1</a>	<a href="#">Efhc1.bSep08</a>	<a href="#">301295</a>	6586	762	3	193	EF-hand domain (C-terminal) containing 1 (Efhc1) alternative variant bSep08, mRNA.
<a href="#">Efhd2</a>	<a href="#">Efhd2.bSep08</a>	<a href="#">298609</a>	1493	1405	2	102	putative protein (Efhd2) alternative variant bSep08, mRNA.
<a href="#">Efna2</a>	<a href="#">Efna2.aSep08</a>	<a href="#">84358</a>	687	495		61	ephrin A2 (Efna2) mRNA.
<a href="#">Efna4</a>	<a href="#">Efna4.bSep08</a>	<a href="#">310643</a>	1131	784	1	146	ephrin A4 (Efna4) alternative variant bSep08, mRNA.
<a href="#">Efnb1</a>	<a href="#">Efnb1.aSep08</a>	<a href="#">25186</a>	12844	3296		501	ephrin B1 (54.0 kD) (Efnb1) mRNA.
<a href="#">Efr3a</a>	<a href="#">Efr3a.aSep08</a>	<a href="#">362923</a>	18609	1362		453	EFR3 homolog A ( <i>S. cerevisiae</i> ) (Efr3a) mRNA.
<a href="#">Efs</a>	<a href="#">Efs.bSep08</a>	<a href="#">290212</a>	872	518	2	172	embryonal Fyn-associated substrate (Efs) alternative variant bSep08, mRNA.
<a href="#">Eftud1</a>	<a href="#">Eftud1.bSep08</a>	<a href="#">308789</a>	14479	1434	3	423	elongation factor G, C-terminal (Eftud1) alternative variant bSep08, mRNA.
<a href="#">Eftud1</a>	<a href="#">Eftud1.cSep08</a>	<a href="#">308789</a>	7530	1567	3	101	putative protein of ancient origin (11.2 kD) (Eftud1) alternative variant cSep08, mRNA.
<a href="#">Eftud1</a>	<a href="#">Eftud1.eSep08</a>	<a href="#">308789</a>	3481	387	3	58	putative protein of ancient origin (6.3 kD) (Eftud1) alternative variant eSep08, mRNA.
<a href="#">Eftud2</a>	<a href="#">Eftud2.aSep08</a>	<a href="#">287739</a>	24939	2521	19	738	elongation factor Tu, domain 2 containing protein and elongation factor G, domain IV containing protein and elongation factor G, C-terminal (Eftud2) alternative variant aSep08, mRNA.
<a href="#">Eftud2</a>	<a href="#">Eftud2.bSep08</a>	<a href="#">287739</a>	5969	1249	7	213	putative cytoplasmic protein of eukaryotic origin (23.6 kD) (Eftud2) alternative variant bSep08, mRNA.
<a href="#">Eftud2</a>	<a href="#">Eftud2.cSep08</a>	<a href="#">287739</a>	5118	712	5	187	putative protein of eukaryotic origin (Eftud2) alternative variant cSep08, mRNA.
<a href="#">Egf</a>	<a href="#">Egf.cSep08</a>	<a href="#">25313</a>	3965	1299	2	67	epidermal growth factor (7.5 kD) (Egf) alternative variant cSep08, mRNA.
<a href="#">Egf</a>	<a href="#">Egf.dSep08</a>	<a href="#">25313</a>	4870	373	3	58	epidermal growth factor (6.1 kD) (Egf) alternative variant dSep08, mRNA.
<a href="#">EGF.1</a>	<a href="#">EGF.1.aSep08</a>		784	341		113	stabilin 1 (EGF.1) mRNA.
<a href="#">EGF.2</a>	<a href="#">EGF.2.aSep08</a>		7470	443		147	sushi nidogen egf-like 1 (EGF.2) mRNA.
<a href="#">EGF.4</a>	<a href="#">EGF.4.aSep08</a>		1628	547	5	181	jagged 2 (EGF.4) alternative variant aSep08, mRNA.
<a href="#">EGF.5</a>	<a href="#">EGF.5.aSep08</a>		809	536		178	cadherin egf lag seven-pass g-type receptor 2 (EGF.5) mRNA.
<a href="#">EGF.6</a>	<a href="#">EGF.6.aSep08</a>		1318	465		154	cadherin egf lag seven-pass g-type receptor 2 (EGF.6) mRNA.
<a href="#">EGF.7</a>	<a href="#">EGF.7.aSep08</a>		3861	535		178	latent transforming growth factor beta binding protein 4 like (EGF.7) mRNA.
<a href="#">EGF.8</a>	<a href="#">EGF.8.aSep08</a>		3139	382		127	latent transforming growth factor beta binding protein 3 like (EGF.8) mRNA.
<a href="#">Egfl6</a>	<a href="#">Egfl6.bSep08</a>	<a href="#">317470</a>	15162	1078	5	247	EGF-like-domain, multiple 6 (Egfl6) alternative variant bSep08, mRNA.
<a href="#">Egfl6</a>	<a href="#">Egfl6.cSep08</a>	<a href="#">317470</a>	14956	609	4	150	EGF-like-domain, multiple 6 (Egfl6) alternative variant cSep08, mRNA.
<a href="#">Egfl7</a>	<a href="#">Egfl7.bSep08</a>	<a href="#">245963</a>	7176	858	7	171	EGF-like domain 7 (Egfl7) alternative variant bSep08, mRNA.

<a href="#">Egfl7</a>	<a href="#">Egfl7.cSep08</a>	<a href="#">245963</a>	6874	675	6	125	EGF-like domain 7 (Egfl7) alternative variant cSep08, mRNA.
<a href="#">Egfl7</a>	<a href="#">Egfl7.dSep08</a>	<a href="#">245963</a>	1313	488	4	97	EGF-like domain 7 (10.7 kD) (Egfl7) alternative variant dSep08, mRNA.
<a href="#">Egfl7</a>	<a href="#">Egfl7.eSep08</a>	<a href="#">245963</a>	6554	548	5	74	EGF-like domain 7 (Egfl7) alternative variant eSep08, mRNA.
<a href="#">Egfl7</a>	<a href="#">Egfl7.fSep08</a>	<a href="#">245963</a>	2360	2244	2	80	EGF-like domain 7 (8.6 kD) (Egfl7) alternative variant fSep08, mRNA.
<a href="#">Egfl7</a>	<a href="#">Egfl7.gSep08</a>	<a href="#">245963</a>	911	342	2	44	EGF-like domain 7 (Egfl7) alternative variant gSep08, mRNA.
<a href="#">Egfl8</a>	<a href="#">Egfl8.bSep08</a>	<a href="#">406166</a>	2356	1377	7	334	EGF-like domain 8 (Egfl8) alternative variant bSep08, mRNA.
<a href="#">Egfl8</a>	<a href="#">Egfl8.cSep08</a>	<a href="#">406166</a>	1421	633	6	138	EGF-like domain 8 (Egfl8) alternative variant cSep08, mRNA.
<a href="#">Egfr</a>	<a href="#">Egfr.aSep08</a>	<a href="#">24329</a>	20568	2099		428	epidermal growth factor receptor (Egfr) mRNA.
<a href="#">EGF_2.0</a>	<a href="#">EGF_2.0.aSep08</a>		676	301		100	tenascin XB (EGF_2.0) mRNA.
<a href="#">EGF_2.1</a>	<a href="#">EGF_2.1.aSep08</a>		9766	1235		126	tenascin-X precursor (13.0 kD) (EGF_2.1) mRNA.
<a href="#">EGF_2.2</a>	<a href="#">EGF_2.2.aSep08</a>		6673	626		208	odd Oz ten-m homolog 3 (EGF_2.2) mRNA.
<a href="#">EGF_2.3</a>	<a href="#">EGF_2.3.aSep08</a>		32147	542	1	175	A disintegrin metallopeptidase domain 32 (EGF_2.3) alternative variant aSep08, mRNA.
<a href="#">EGF_2.3</a>	<a href="#">EGF_2.3.bSep08</a>		28622	568	2	76	putative secreted or extracellular protein precursor of mammalian origin (8.4 kD) (EGF_2.3) alternative variant bSep08, mRNA.
<a href="#">EGF_CA.0</a>	<a href="#">EGF_CA.0.aSep08</a>		18256	712		236	nidogen (EGF_CA.0) mRNA.
<a href="#">EGF_CA.1</a>	<a href="#">EGF_CA.1.aSep08</a>		46946	1312		436	latent transforming growth factor beta binding protein 1 like (EGF_CA.1) mRNA.
<a href="#">EGF_CA.2</a>	<a href="#">EGF_CA.2.aSep08</a>		1355	1104		158	CRA b (EGF_CA.2) mRNA.
<a href="#">EGF_CA.3</a>	<a href="#">EGF_CA.3.aSep08</a>		1698	606	5	202	latent transforming growth factor beta binding protein 3 CRA e like (EGF_CA.3) alternative variant aSep08, mRNA.
<a href="#">EGF_CA.3</a>	<a href="#">EGF_CA.3.bSep08</a>		600	423	1	85	latent transforming growth factor binding protein like (EGF_CA.3) alternative variant bSep08, mRNA.
<a href="#">Egr3</a>	<a href="#">Egr3.bSep08</a>	<a href="#">25148</a>	1895	655	1	71	early growth response 3 (7.7 kD) (Egr3) alternative variant bSep08, mRNA.
<a href="#">Ehbp1</a>	<a href="#">Ehbp1.aSep08</a>	<a href="#">305556</a>	122934	3197	5	753	EH domain binding protein 1 (Ehbp1) alternative variant aSep08, mRNA.
<a href="#">Ehbp1</a>	<a href="#">Ehbp1.bSep08</a>	<a href="#">305556</a>	58454	965	2	321	EH domain binding protein 1 (Ehbp1) alternative variant bSep08, mRNA.
<a href="#">Ehbp111</a>	<a href="#">Ehbp111.bSep08</a>	<a href="#">309169</a>	18128	3171	16	844	calponin-like actin-binding (90.9 kD) (Ehbp111) alternative variant bSep08, complete mRNA.
<a href="#">Ehbp111</a>	<a href="#">Ehbp111.cSep08</a>	<a href="#">309169</a>	10567	3470	5	813	CRA d (Ehbp111) alternative variant cSep08, mRNA.
<a href="#">Ehbp111</a>	<a href="#">Ehbp111.eSep08</a>	<a href="#">309169</a>	18091	2603	15	667	tangerin (72.5 kD) (Ehbp111) alternative variant eSep08, complete mRNA.
<a href="#">Ehbp111</a>	<a href="#">Ehbp111.fSep08</a>	<a href="#">309169</a>	7227	1392	2	247	tangerin (25.5 kD) (Ehbp111) alternative variant fSep08, mRNA.
<a href="#">Ehd1</a>	<a href="#">Ehd1.aSep08</a>	<a href="#">293692</a>	22381	3240		558	hpast (Ehd1) mRNA.



<a href="#">Ehmt1</a>	<a href="#">Ehmt1.bSep08</a>	<a href="#">362078</a>	24025	627	6	208	euchromatic histone methyltransferase 1 (Ehmt1) alternative variant bSep08, mRNA.
<a href="#">Ehmt1</a>	<a href="#">Ehmt1.cSep08</a>	<a href="#">362078</a>	58757	398	3	132	euchromatic histone methyltransferase 1 (Ehmt1) alternative variant cSep08, mRNA.
<a href="#">Ehmt2</a>	<a href="#">Ehmt2.bSep08</a>	<a href="#">361798</a>	9303	2987	19	582	euchromatic histone lysine N-methyltransferase 2 (Ehmt2) alternative variant bSep08, mRNA.
<a href="#">Ehmt2</a>	<a href="#">Ehmt2.dSep08</a>	<a href="#">361798</a>	538	370	2	123	euchromatic histone lysine N-methyltransferase 2 (Ehmt2) alternative variant dSep08, mRNA.
<a href="#">Ei24</a>	<a href="#">Ei24.bSep08</a>	<a href="#">300514</a>	12400	909	8	222	etoposide induced 2.4 mRNA (Ei24) alternative variant bSep08, mRNA.
<a href="#">Ei24</a>	<a href="#">Ei24.cSep08</a>	<a href="#">300514</a>	1711	542	2	56	etoposide induced 2.4 mRNA (Ei24) alternative variant cSep08, mRNA.
<a href="#">Eid1</a>	<a href="#">Eid1.bSep08</a>	<a href="#">499882</a>	1426	1256	2	67	EP300 interacting inhibitor of differentiation 1 (7.8 kD) (Eid1) alternative variant bSep08, mRNA.
<a href="#">Eif1</a>	<a href="#">Eif1.aSep08</a>	<a href="#">287703</a>	2304	1594	3	196	eukaryotic translation initiation factor 1 (Eif1) alternative variant aSep08, mRNA.
<a href="#">Eif1</a>	<a href="#">Eif1.cSep08</a>	<a href="#">287703</a>	1673	1251	3	164	eukaryotic translation initiation factor 1 (18.5 kD) (Eif1) alternative variant cSep08, complete mRNA.
<a href="#">Eif1a</a>	<a href="#">Eif1a.aSep08</a>	<a href="#">317163</a>	9271	589	2	101	eukaryotic translation initiation factor 1A (11.1 kD) (Eif1a) alternative variant aSep08, mRNA.
<a href="#">Eif1ad</a>	<a href="#">Eif1ad.bSep08</a>	<a href="#">293673</a>	3487	520	5	86	putative protein of eukaryotic origin (Eif1ad) alternative variant bSep08, mRNA.
<a href="#">Eif1ad</a>	<a href="#">Eif1ad.eSep08</a>	<a href="#">293673</a>	421	275	2	51	eukaryotic initiation factor 1A (Eif1ad) alternative variant eSep08, mRNA.
<a href="#">Eif2a</a>	<a href="#">Eif2a.bSep08</a>	<a href="#">502531</a>	3134	985	4	299	eukaryotic translation initiation factor 2A (Eif2a) alternative variant bSep08, mRNA.
<a href="#">Eif2a</a>	<a href="#">Eif2a.cSep08</a>	<a href="#">502531</a>	23449	883	10	294	eukaryotic translation initiation factor 2A (Eif2a) alternative variant cSep08, mRNA.
<a href="#">Eif2ak1</a>	<a href="#">Eif2ak1.bSep08</a>	<a href="#">27137</a>	13636	830	7	276	eukaryotic translation initiation factor 2 alpha kinase 1 (Eif2ak1) alternative variant bSep08, mRNA.
<a href="#">Eif2ak1</a>	<a href="#">Eif2ak1.cSep08</a>	<a href="#">27137</a>	2040	429	2	78	eukaryotic translation initiation factor 2 alpha kinase 1 (Eif2ak1) alternative variant cSep08, mRNA.
<a href="#">Eif2ak2</a>	<a href="#">Eif2ak2.bSep08</a>	<a href="#">54287</a>	3240	735	1	195	eukaryotic translation initiation factor 2-alpha kinase 2 (Eif2ak2) alternative variant bSep08, mRNA.
<a href="#">Eif2ak3</a>	<a href="#">Eif2ak3.bSep08</a>	<a href="#">29702</a>	3369	360		119	eukaryotic translation initiation factor 2 alpha kinase 3 (Eif2ak3) alternative variant bSep08, mRNA.
<a href="#">Eif2ak4</a>	<a href="#">Eif2ak4.aSep08</a>	<a href="#">114859</a>	20724	1723	15	512	eukaryotic translation initiation factor 2 alpha kinase 4 (Eif2ak4) alternative variant aSep08, mRNA.
<a href="#">Eif2ak4</a>	<a href="#">Eif2ak4.bSep08</a>	<a href="#">114859</a>	2719	648	3	81	eukaryotic translation initiation factor 2 alpha kinase 4 (9.9 kD) (Eif2ak4) alternative variant bSep08, mRNA.
<a href="#">Eif2ak4</a>	<a href="#">Eif2ak4.cSep08</a>	<a href="#">114859</a>	1563	740	2	44	eukaryotic translation initiation factor 2 alpha kinase 4 (5.3 kD) (Eif2ak4) alternative variant cSep08, mRNA.
<a href="#">Eif2b4</a>	<a href="#">Eif2b4.bSep08</a>	<a href="#">117019</a>	4052	2143	9	289	eukaryotic translation initiation factor 2B, subunit 4 delta (Eif2b4) alternative variant bSep08, mRNA.
<a href="#">Eif2b4</a>	<a href="#">Eif2b4.cSep08</a>	<a href="#">117019</a>	3187	1799	3	127	eukaryotic translation initiation factor 2B, subunit 4 delta (Eif2b4) alternative variant cSep08, mRNA.
<a href="#">Eif2c1</a>	<a href="#">Eif2c1.aSep08</a>	<a href="#">313594</a>	20612	5166		511	eukaryotic translation initiation factor 2C, 1 (Eif2c1) mRNA.

<a href="#">Eif2c3</a>	<a href="#">Eif2c3.aSep08</a>	<a href="#">313593</a>	18371	806		205	eukaryotic translation initiation factor 2C, 3 (Eif2c3) mRNA.
<a href="#">Eif2s2</a>	<a href="#">Eif2s2.bSep08</a>	<a href="#">296302</a>	14660	773	5	188	eukaryotic translation initiation factor 2 beta (21.9 kD) (Eif2s2) alternative variant bSep08, complete mRNA.
<a href="#">Eif2s2</a>	<a href="#">Eif2s2.cSep08</a>	<a href="#">296302</a>	5769	2109	3	90	eukaryotic translation initiation factor 2 beta 38kDa (10.2 kD) (Eif2s2) alternative variant cSep08, mRNA.
<a href="#">Eif2s2</a>	<a href="#">Eif2s2.dSep08</a>	<a href="#">296302</a>	2382	1709	2	88	CRA b like (10.1 kD) (Eif2s2) alternative variant dSep08, mRNA.
<a href="#">Eif2s2</a>	<a href="#">Eif2s2.eSep08</a>	<a href="#">296302</a>	9185	521	4	44	putative protein (5.5 kD) (Eif2s2) alternative variant eSep08, complete mRNA.
<a href="#">Eif2s2</a>	<a href="#">Eif2s2.fSep08</a>	<a href="#">296302</a>	6272	457	3	44	putative protein (5.5 kD) (Eif2s2) alternative variant fSep08, complete mRNA.
<a href="#">Eif2s3x</a>	<a href="#">Eif2s3x.bSep08</a>	<a href="#">299027</a>	12505	886	5	286	eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked (Eif2s3x) alternative variant bSep08, mRNA.
<a href="#">Eif2s3x</a>	<a href="#">Eif2s3x.cSep08</a>	<a href="#">299027</a>	6536	707	2	196	eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked (Eif2s3x) alternative variant cSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.bSep08</a>	<a href="#">293484</a>	8097	744	4	242	eukaryotic translation initiation factor C-like (Eif3c) alternative variant bSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.cSep08</a>	<a href="#">293484</a>	7966	702	3	223	eukaryotic translation initiation factor C-like (Eif3c) alternative variant cSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.dSep08</a>	<a href="#">293484</a>	2366	655	6	218	eukaryotic translation initiation factor C-like (Eif3c) alternative variant dSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.eSep08</a>	<a href="#">293484</a>	1521	761	4	211	eukaryotic translation initiation factor 3 (Eif3c) alternative variant eSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.fSep08</a>	<a href="#">293484</a>	895	775	2	116	eukaryotic translation initiation factor 3 110kDa CRA a (Eif3c) alternative variant fSep08, mRNA.
<a href="#">Eif3c</a>	<a href="#">Eif3c.gSep08</a>	<a href="#">293484</a>	629	528	2	74	putative protein (Eif3c) alternative variant gSep08, mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.bSep08</a>	<a href="#">362952</a>	8292	1194	9	298	eukaryotic translation initiation factor 3 CRA c (35.2 kD) (Eif3d) alternative variant bSep08, complete mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.cSep08</a>	<a href="#">362952</a>	6818	736	8	237	eukaryotic translation initiation factor 3 CRA c (Eif3d) alternative variant cSep08, mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.dSep08</a>	<a href="#">362952</a>	7125	900	8	235	eukaryotic translation initiation factor 3 CRA c (Eif3d) alternative variant dSep08, mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.eSep08</a>	<a href="#">362952</a>	7179	790	8	134	eukaryotic translation initiation factor 3 CRA c (Eif3d) alternative variant eSep08, mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.fSep08</a>	<a href="#">362952</a>	3517	458	4	93	eukaryotic translation initiation factor 3 CRA c (Eif3d) alternative variant fSep08, mRNA.
<a href="#">Eif3d</a>	<a href="#">Eif3d.gSep08</a>	<a href="#">362952</a>	2954	416	3	76	eukaryotic translation initiation factor 3 CRA c (8.7 kD) (Eif3d) alternative variant gSep08, complete mRNA.
<a href="#">Eif3e</a>	<a href="#">Eif3e.bSep08</a>	<a href="#">299872</a>	14882	726	6	133	eukaryotic translation initiation factor 3, subunit E (Eif3e) alternative variant bSep08, mRNA.
<a href="#">Eif3f</a>	<a href="#">Eif3f.aSep08</a>	<a href="#">293427</a>	9026	1251	8	363	eukaryotic translation initiation factor 3, subunit F (Eif3f) alternative variant aSep08, mRNA.
<a href="#">Eif3g</a>	<a href="#">Eif3g.bSep08</a>	<a href="#">298700</a>	861	769	2	222	eukaryotic translation initiation factor 3, subunit G (Eif3g) alternative variant bSep08, mRNA.
<a href="#">Eif3h</a>	<a href="#">Eif3h.bSep08</a>	<a href="#">299899</a>	72148	620	1	91	eukaryotic translation initiation factor 3, subunit H (Eif3h) alternative variant bSep08, mRNA.

<a href="#">Eif3j</a>	<a href="#">Eif3j.cSep08</a>	<a href="#">691947</a>	12241	302	2	36	eukaryotic translation initiation factor 3, subunit J (Eif3j) alternative variant cSep08, mRNA.
<a href="#">Eif3k</a>	<a href="#">Eif3k.aSep08</a>	<a href="#">292762</a>	9301	1523	7	264	eukaryotic translation initiation factor 3 (30.1 kD) (Eif3k) alternative variant aSep08, mRNA.
<a href="#">Eif3k</a>	<a href="#">Eif3k.bSep08</a>	<a href="#">292762</a>	10318	1212	8	218	eukaryotic translation initiation factor 3 (25.1 kD) (Eif3k) alternative variant bSep08, mRNA.
<a href="#">Eif3k</a>	<a href="#">Eif3k.cSep08</a>	<a href="#">292762</a>	5335	1063	5	119	eukaryotic translation initiation factor 3 CRA f (13.9 kD) (Eif3k) alternative variant cSep08, mRNA.
<a href="#">Eif3k</a>	<a href="#">Eif3k.dSep08</a>	<a href="#">292762</a>	3584	598	4	109	eukaryotic translation initiation factor 3 (12.3 kD) (Eif3k) alternative variant dSep08, mRNA.
<a href="#">Eif3k</a>	<a href="#">Eif3k.fSep08</a>	<a href="#">292762</a>	2952	690	2	41	putative protein (4.9 kD) (Eif3k) alternative variant fSep08, mRNA.
<a href="#">Eif3s6ip</a>	<a href="#">Eif3s6ip.aSep08</a>	<a href="#">300069</a>	20809	1913	1	583	eukaryotic translation initiation factor 3, subunit 6 interacting protein (Eif3s6ip) alternative variant aSep08, mRNA.
<a href="#">Eif3s9</a>	<a href="#">Eif3s9.bSep08</a>	<a href="#">288516</a>	9665	757	8	191	eukaryotic translation initiation factor 3, subunit 9 (eta) (Eif3s9) alternative variant bSep08, mRNA.
<a href="#">Eif3s10</a>	<a href="#">Eif3s10.bSep08</a>	<a href="#">292148</a>	3537	443	4	141	eukaryotic translation initiation factor 3, subunit 10 (theta) (Eif3s10) alternative variant bSep08, mRNA.
<a href="#">Eif4a1</a>	<a href="#">Eif4a1.bSep08</a>	<a href="#">287436</a>	5275	1853	10	332	eukaryotic translation initiation factor 4A1 (37.9 kD) (Eif4a1) alternative variant bSep08, mRNA.
<a href="#">Eif4a1</a>	<a href="#">Eif4a1.cSep08</a>	<a href="#">287436</a>	1034	859	3	128	eukaryotic translation initiation factor 4A1 (Eif4a1) alternative variant cSep08, mRNA.
<a href="#">Eif4a1</a>	<a href="#">Eif4a1.gSep08</a>	<a href="#">287436</a>	475	315	2	58	eukaryotic translation initiation factor 4A1 (Eif4a1) alternative variant gSep08, mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.bSep08</a>	<a href="#">303831</a>	6611	2011	12	363	factor Eukaryotic initiation 4 (41.4 kD) (Eif4a2) alternative variant bSep08, complete mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.cSep08</a>	<a href="#">303831</a>	6597	2560	10	235	eukaryotic initiation factor 4 (27.4 kD) (Eif4a2) alternative variant cSep08, complete mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.dSep08</a>	<a href="#">303831</a>	5830	2265	9	233	eukaryotic initiation factor 4 (27.1 kD) (Eif4a2) alternative variant dSep08, mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.eSep08</a>	<a href="#">303831</a>	4027	1558	7	115	factor eukaryotic initiation 4 (12.6 kD) (Eif4a2) alternative variant eSep08, mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.fSep08</a>	<a href="#">303831</a>	3332	869	6	146	factor eukaryotic initiation 4 (17.3 kD) (Eif4a2) alternative variant fSep08, mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.gSep08</a>	<a href="#">303831</a>	2323	768	4	86	eukaryotic initiation factor 4 (9.5 kD) (Eif4a2) alternative variant gSep08, mRNA.
<a href="#">Eif4a2</a>	<a href="#">Eif4a2.hSep08</a>	<a href="#">303831</a>	1205	744	3	68	initiation factor eukaryotic 4 (7.9 kD) (Eif4a2) alternative variant hSep08, mRNA.
<a href="#">Eif4b</a>	<a href="#">Eif4b.bSep08</a>	<a href="#">300253</a>	2937	795	3	142	eukaryotic translation initiation factor 4B (Eif4b) alternative variant bSep08, mRNA.
<a href="#">Eif4e</a>	<a href="#">Eif4e.bSep08</a>	<a href="#">117045</a>	30641	1178	8	212	eukaryotic translation initiation factor 4E (24.6 kD) (Eif4e) alternative variant bSep08, mRNA.
<a href="#">Eif4e2</a>	<a href="#">Eif4e2.aSep08</a>	<a href="#">363275</a>	16220	1286	7	240	eukaryotic translation initiation factor 4E member 2 (27.6 kD) (Eif4e2) alternative variant aSep08, mRNA.
<a href="#">Eif4e2</a>	<a href="#">Eif4e2.cSep08</a>	<a href="#">363275</a>	12240	466	5	154	eukaryotic translation initiation factor 4E member 2 (Eif4e2) alternative variant cSep08, mRNA.

<a href="#">Eif4e2</a>	<a href="#">Eif4e2.dSep08</a>	<a href="#">363275</a>	8925	287	3	85	eukaryotic translation initiation factor 4E member 2 (Eif4e2) alternative variant dSep08, mRNA.
<a href="#">Eif4e3</a>	<a href="#">Eif4e3.bSep08</a>	<a href="#">297481</a>	39155	582	1	101	eukaryotic translation initiation factor 4E member 3 (Eif4e3) alternative variant bSep08, mRNA.
<a href="#">Eif4ebp1</a>	<a href="#">Eif4ebp1.cSep08</a>	<a href="#">116636</a>	13748	770	3	126	eukaryotic translation initiation factor 4E binding protein 1 (13.6 kD) (Eif4ebp1) alternative variant cSep08, mRNA.
<a href="#">Eif4g1</a>	<a href="#">Eif4g1.aSep08</a>	<a href="#">287986</a>	20174	5413	32	1591	eukaryotic translation initiation factor 4 gamma, 1 (175.0 kD) (Eif4g1) alternative variant aSep08, mRNA.
<a href="#">Eif4g1</a>	<a href="#">Eif4g1.bSep08</a>	<a href="#">287986</a>	3104	1121	8	373	eukaryotic translation initiation factor 4 gamma, 1 (Eif4g1) alternative variant bSep08, mRNA.
<a href="#">Eif4g1</a>	<a href="#">Eif4g1.cSep08</a>	<a href="#">287986</a>	4974	670	7	223	eukaryotic translation initiation factor 4 gamma, 1 (Eif4g1) alternative variant cSep08, mRNA.
<a href="#">Eif4g1</a>	<a href="#">Eif4g1.dSep08</a>	<a href="#">287986</a>	5311	607	6	119	eukaryotic translation initiation factor 4 gamma, 1 (Eif4g1) alternative variant dSep08, mRNA.
<a href="#">Eif4g1</a>	<a href="#">Eif4g1.eSep08</a>	<a href="#">287986</a>	744	339	3	112	eukaryotic translation initiation factor 4 gamma, 1 (Eif4g1) alternative variant eSep08, mRNA.
<a href="#">Eif4g2_predicted</a>	<a href="#">Eif4g2_predicted.bSep08</a>	<a href="#">361628</a>	7073	1723	14	473	eukaryotic translation initiation factor 4, gamma 2 (predicted) (Eif4g2_predicted) alternative variant bSep08, mRNA.
<a href="#">Eif4g2_predicted</a>	<a href="#">Eif4g2_predicted.cSep08</a>	<a href="#">361628</a>	5070	3340	8	409	eukaryotic translation initiation factor 4, gamma 2 (predicted) (Eif4g2_predicted) alternative variant cSep08, mRNA.
<a href="#">Eif4g2_predicted</a>	<a href="#">Eif4g2_predicted.dSep08</a>	<a href="#">361628</a>	5832	1221	12	381	eukaryotic translation initiation factor 4, gamma 2 (predicted) (Eif4g2_predicted) alternative variant dSep08, mRNA.
<a href="#">Eif4g2_predicted</a>	<a href="#">Eif4g2_predicted.eSep08</a>	<a href="#">361628</a>	2033	802	6	267	eukaryotic translation initiation factor 4, gamma 2 (predicted) (Eif4g2_predicted) alternative variant eSep08, mRNA.
<a href="#">Eif4g2_predicted</a>	<a href="#">Eif4g2_predicted.fSep08</a>	<a href="#">361628</a>	1301	261	3	8	eukaryotic translation initiation factor 4, gamma 2 (predicted) (Eif4g2_predicted) alternative variant fSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.bSep08</a>	<a href="#">298573</a>	139868	1196	11	398	eukaryotic translation initiation factor 4 gamma 3 (Eif4g3) alternative variant bSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.cSep08</a>	<a href="#">298573</a>	17216	1112	9	370	eukaryotic translation initiation factor 4 gamma 3 CRA a (Eif4g3) alternative variant cSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.dSep08</a>	<a href="#">298573</a>	66544	706	6	235	eukaryotic translation initiation factor 4 gamma 3 CRA c (Eif4g3) alternative variant dSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.eSep08</a>	<a href="#">298573</a>	64529	687	7	229	eukaryotic translation initiation factor 4 gamma 3 (Eif4g3) alternative variant eSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.fSep08</a>	<a href="#">298573</a>	14181	1066	6	214	eukaryotic translation initiation factor 4 gamma 3 CRA a (Eif4g3) alternative variant fSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.gSep08</a>	<a href="#">298573</a>	13408	1534	5	208	eukaryotic translation initiation factor 4 gamma 3 CRA b (Eif4g3) alternative variant gSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.hSep08</a>	<a href="#">298573</a>	45028	456	5	152	eukaryotic translation initiation factor 4 gamma 3 (Eif4g3) alternative variant hSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.iSep08</a>	<a href="#">298573</a>	21759	1414	8	151	eukaryotic translation initiation factor 4 gamma 3 CRA b (17.2 kD) (Eif4g3) alternative variant iSep08, mRNA.

<a href="#">Eif4g3</a>	<a href="#">Eif4g3.jSep08</a>	<a href="#">298573</a>	96884	497	4	139	eukaryotic translation initiation factor 4 gamma 3 like (Eif4g3) alternative variant jSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.kSep08</a>	<a href="#">298573</a>	58025	408	4	136	eukaryotic translation initiation factor 4 gamma 3 (Eif4g3) alternative variant kSep08, mRNA.
<a href="#">Eif4g3</a>	<a href="#">Eif4g3.mSep08</a>	<a href="#">298573</a>	1159	390	2	38	eukaryotic translation initiation factor 4 gamma 3 CRA b (Eif4g3) alternative variant mSep08, mRNA.
<a href="#">Eif4h</a>	<a href="#">Eif4h.aSep08</a>	<a href="#">288599</a>	3753	3309	2	69	putative protein (31.7 kD) (Eif4h) alternative variant aSep08, mRNA.
<a href="#">Eif4h</a>	<a href="#">Eif4h.cSep08</a>	<a href="#">288599</a>	15908	1611	6	228	eukaryotic translation initiation factor 4H (25.2 kD) (Eif4h) alternative variant cSep08, mRNA.
<a href="#">Eif4h</a>	<a href="#">Eif4h.eSep08</a>	<a href="#">288599</a>	3059	1020	4	84	eukaryotic translation initiation factor 4H (Eif4h) alternative variant eSep08, mRNA.
<a href="#">Eif5</a>	<a href="#">Eif5.bSep08</a>	<a href="#">56783</a>	3639	949	7	163	eukaryotic translation initiation factor 5 (Eif5) alternative variant bSep08, mRNA.
<a href="#">Eif5</a>	<a href="#">Eif5.dSep08</a>	<a href="#">56783</a>	1771	1232	4	59	eukaryotic translation initiation factor 5 (Eif5) alternative variant dSep08, mRNA.
<a href="#">Eif5</a>	<a href="#">Eif5.eSep08</a>	<a href="#">56783</a>	788	518	2	68	eukaryotic translation initiation factor 5 (Eif5) alternative variant eSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.aSep08</a>	<a href="#">287444</a>	3640	820	6	154	eukaryotic translation initiation factor 5A (16.8 kD) (Eif5a) alternative variant aSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.bSep08</a>	<a href="#">287444</a>	4365	1525	6	154	eukaryotic translation initiation factor 5A (16.8 kD) (Eif5a) alternative variant bSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.cSep08</a>	<a href="#">287444</a>	3759	1381	6	154	eukaryotic translation initiation factor 5A (16.8 kD) (Eif5a) alternative variant cSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.dSep08</a>	<a href="#">287444</a>	3986	878	7	154	eukaryotic translation initiation factor 5A (16.8 kD) (Eif5a) alternative variant dSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.fSep08</a>	<a href="#">287444</a>	3698	854	3	140	eukaryotic translation initiation factor 5A (Eif5a) alternative variant fSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.gSep08</a>	<a href="#">287444</a>	3855	739	5	130	eukaryotic translation initiation factor 5A (Eif5a) alternative variant gSep08, mRNA.
<a href="#">Eif5a</a>	<a href="#">Eif5a.hSep08</a>	<a href="#">287444</a>	4121	733	5	116	eukaryotic translation initiation factor 5A (Eif5a) alternative variant hSep08, mRNA.
<a href="#">Eif5a2</a>	<a href="#">Eif5a2.aSep08</a>	<a href="#">310261</a>	13815	1920		188	eukaryotic translation initiation factor 5A2 (Eif5a2) mRNA.
<a href="#">Eif5b</a>	<a href="#">Eif5b.aSep08</a>	<a href="#">308306</a>	27742	1308		383	eukaryotic translation initiation factor 5B (Eif5b) mRNA.
<a href="#">Ela1</a>	<a href="#">Ela1.bSep08</a>	<a href="#">24331</a>	6821	627	2	113	elastase 1, pancreatic (Ela1) alternative variant bSep08, mRNA.
<a href="#">Elac2</a>	<a href="#">Elac2.aSep08</a>	<a href="#">282826</a>	18334	2957	18	611	elaC homolog 2 (E. coli) (Elac2) alternative variant aSep08, mRNA.
<a href="#">Elac2</a>	<a href="#">Elac2.bSep08</a>	<a href="#">282826</a>	1453	722	4	109	elaC homolog 2 (E. coli) (Elac2) alternative variant bSep08, mRNA.
<a href="#">Elac2</a>	<a href="#">Elac2.eSep08</a>	<a href="#">282826</a>	4234	508	3	50	elaC homolog 2 (E. coli) (Elac2) alternative variant eSep08, mRNA.
<a href="#">Elavl1</a>	<a href="#">Elavl1.bSep08</a>	<a href="#">363854</a>	5967	1210	4	111	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R) (12.2 kD) (Elavl1) alternative variant bSep08, mRNA.
<a href="#">Elavl2</a>	<a href="#">Elavl2.bSep08</a>	<a href="#">286973</a>	7243	440	3	146	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B) (Elavl2) alternative variant bSep08, mRNA.

<a href="#">Elavl2</a>	<a href="#">Elavl2.cSep08</a>	<a href="#">286973</a>	94868	410	3	105	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B) (Elavl2) alternative variant cSep08, mRNA.
<a href="#">Elavl3</a>	<a href="#">Elavl3.bSep08</a>	<a href="#">282824</a>	14866	536	2	178	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu antigen C) (Elavl3) alternative variant bSep08, mRNA.
<a href="#">Elavl3</a>	<a href="#">Elavl3.cSep08</a>	<a href="#">282824</a>	8538	1205	3	160	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu antigen C) (Elavl3) alternative variant cSep08, mRNA.
<a href="#">Elavl3</a>	<a href="#">Elavl3.dSep08</a>	<a href="#">282824</a>	1891	521	3	108	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu antigen C) (Elavl3) alternative variant dSep08, mRNA.
<a href="#">Elavl4</a>	<a href="#">Elavl4.aSep08</a>	<a href="#">432358</a>	98375	3880	7	385	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D) (42.4 kD) (Elavl4) alternative variant aSep08, mRNA.
<a href="#">Elavl4</a>	<a href="#">Elavl4.bSep08</a>	<a href="#">432358</a>	21380	1785	4	13	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D) (1.4 kD) (Elavl4) alternative variant bSep08, mRNA.
<a href="#">Elavl4</a>	<a href="#">Elavl4.cSep08</a>	<a href="#">432358</a>	35951	387	2	37	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D) (Elavl4) alternative variant cSep08, mRNA.
<a href="#">Elf1</a>	<a href="#">Elf1.bSep08</a>	<a href="#">85424</a>	32300	568	3	81	e74-like factor 1 (Elf1) alternative variant bSep08, mRNA.
<a href="#">Elf1</a>	<a href="#">Elf1.cSep08</a>	<a href="#">85424</a>	13412	403	2	45	e74-like factor 1 (5.3 kD) (Elf1) alternative variant cSep08, mRNA.
<a href="#">Elf2</a>	<a href="#">Elf2.cSep08</a>	<a href="#">361944</a>	10956	630	2	100	e74-like factor 2 (10.7 kD) (Elf2) alternative variant cSep08, mRNA.
<a href="#">Elf2</a>	<a href="#">Elf2.dSep08</a>	<a href="#">361944</a>	42976	541	2	99	e74-like factor 2 (Elf2) alternative variant dSep08, mRNA.
<a href="#">Elf2</a>	<a href="#">Elf2.eSep08</a>	<a href="#">361944</a>	43788	418	3	33	e74-like factor 2 (4.0 kD) (Elf2) alternative variant eSep08, mRNA.
<a href="#">Elf3</a>	<a href="#">Elf3.bSep08</a>	<a href="#">304815</a>	2057	751	6	250	e74-like factor 3 (Elf3) alternative variant bSep08, mRNA.
<a href="#">Elf3</a>	<a href="#">Elf3.cSep08</a>	<a href="#">304815</a>	1400	851	2	79	e74-like factor 3 (9.6 kD) (Elf3) alternative variant cSep08, mRNA.
<a href="#">Elf4</a>	<a href="#">Elf4.aSep08</a>	<a href="#">302811</a>	2885	853		283	e74-like factor 4 (ets domain transcription factor) (Elf4) mRNA.
<a href="#">Elk3</a>	<a href="#">Elk3.aSep08</a>	<a href="#">362871</a>	62241	3952	6	462	ELK3, member of ETS oncogene family (Elk3) alternative variant aSep08, mRNA.
<a href="#">Elk3</a>	<a href="#">Elk3.bSep08</a>	<a href="#">362871</a>	1759	478	3	103	ELK3, member of ETS oncogene family (Elk3) alternative variant bSep08, mRNA.
<a href="#">Elk3</a>	<a href="#">Elk3.cSep08</a>	<a href="#">362871</a>	5379	975	2	63	ELK3, member of ETS oncogene family (Elk3) alternative variant cSep08, mRNA.
<a href="#">Elk4</a>	<a href="#">Elk4.aSep08</a>	<a href="#">304786</a>	14340	2168	5	533	ELK4, member of ETS oncogene family (Elk4) alternative variant aSep08, mRNA.
<a href="#">Elk4</a>	<a href="#">Elk4.dSep08</a>	<a href="#">304786</a>	3417	438	2	52	ELK4, member of ETS oncogene family (5.6 kD) (Elk4) alternative variant dSep08, mRNA.
<a href="#">EII2</a>	<a href="#">EII2.aSep08</a>	<a href="#">309918</a>	68739	3542	12	640	elongation factor RNA polymerase II 2 (72.3 kD) (EII2) alternative variant aSep08, complete mRNA.
<a href="#">EII2</a>	<a href="#">EII2.bSep08</a>	<a href="#">309918</a>	7823	1192	6	263	elongation factor RNA polymerase II 2 (EII2) alternative variant bSep08, mRNA.
<a href="#">EII2</a>	<a href="#">EII2.cSep08</a>	<a href="#">309918</a>	21114	367	4	95	elongation factor RNA polymerase II 2 (EII2) alternative variant cSep08, mRNA.
<a href="#">EII2</a>	<a href="#">EII2.dSep08</a>	<a href="#">309918</a>	881	619	2	80	elongation factor RNA polymerase II 2 (EII2) alternative variant dSep08, mRNA.

<a href="#">EII2</a>	<a href="#">EII2.eSep08</a>	<a href="#">309918</a>	604	341	2	72	elongation factor RNA polymerase II 2 (7.9 kD) (EII2) alternative variant eSep08, mRNA.
<a href="#">ELM2.0</a>	<a href="#">ELM2.0.aSep08</a>		2324	470	5	156	ELM2 (ELM2.0) alternative variant aSep08, mRNA.
<a href="#">Elmo1</a>	<a href="#">Elmo1.bSep08</a>	<a href="#">361251</a>	266306	2331	9	365	engulfment cell motility 1 (Elmo1) alternative variant bSep08, mRNA.
<a href="#">Elmo1</a>	<a href="#">Elmo1.cSep08</a>	<a href="#">361251</a>	101158	548	4	116	engulfment cell motility 1 (Elmo1) alternative variant cSep08, mRNA.
<a href="#">Elmo1</a>	<a href="#">Elmo1.dSep08</a>	<a href="#">361251</a>	161084	479	6	105	engulfment cell motility (Elmo1) alternative variant dSep08, mRNA.
<a href="#">Elmo1</a>	<a href="#">Elmo1.eSep08</a>	<a href="#">361251</a>	161194	593	6	105	engulfment cell motility (Elmo1) alternative variant eSep08, mRNA.
<a href="#">Elmo2</a>	<a href="#">Elmo2.aSep08</a>	<a href="#">362271</a>	35744	2791	21	746	engulfment cell motility 2 (Elmo2) alternative variant aSep08, mRNA.
<a href="#">Elmo2</a>	<a href="#">Elmo2.bSep08</a>	<a href="#">362271</a>	20733	816	9	260	engulfment cell motility 2 (Elmo2) alternative variant bSep08, mRNA.
<a href="#">Elmo2</a>	<a href="#">Elmo2.cSep08</a>	<a href="#">362271</a>	9894	749	5	243	engulfment cell motility 2 (Elmo2) alternative variant cSep08, mRNA.
<a href="#">Elmo2</a>	<a href="#">Elmo2.dSep08</a>	<a href="#">362271</a>	6381	2633	5	181	engulfment cell motility 2 (Elmo2) alternative variant dSep08, mRNA.
<a href="#">Elmo2</a>	<a href="#">Elmo2.eSep08</a>	<a href="#">362271</a>	2607	683	2	65	putative protein (7.0 kD) (Elmo2) alternative variant eSep08, mRNA.
<a href="#">Elmo3</a>	<a href="#">Elmo3.aSep08</a>	<a href="#">291962</a>	2534	1310	12	384	engulfment cell motility 3 (Elmo3) alternative variant aSep08, mRNA.
<a href="#">Elmo3</a>	<a href="#">Elmo3.bSep08</a>	<a href="#">291962</a>	821	415	4	92	engulfment cell motility 3 (Elmo3) alternative variant bSep08, mRNA.
<a href="#">Elmo3</a>	<a href="#">Elmo3.cSep08</a>	<a href="#">291962</a>	496	413	2	46	engulfment cell motility 3 (Elmo3) alternative variant cSep08, mRNA.
<a href="#">Elmod1</a>	<a href="#">Elmod1.aSep08</a>	<a href="#">315670</a>	44301	1793	2	504	putative protein of eukaryotic origin (Elmod1) alternative variant aSep08, mRNA.
<a href="#">Elmod1</a>	<a href="#">Elmod1.bSep08</a>	<a href="#">315670</a>	43823	825	1	207	engulfment and cell motility, ELM (Elmod1) alternative variant bSep08, mRNA.
<a href="#">ELMO_CED12.0</a>	<a href="#">ELMO_CED12.0.aSep08</a>		1319	867	6	252	engulfment cell motility 3 (ELMO_CED12.0) alternative variant aSep08, mRNA.
<a href="#">ELMO_CED12.0</a>	<a href="#">ELMO_CED12.0.bSep08</a>		1177	779	5	178	engulfment cell motility 3 (ELMO_CED12.0) alternative variant bSep08, mRNA.
<a href="#">Eln</a>	<a href="#">Eln.aSep08</a>	<a href="#">25043</a>	43387	3593	37	877	elastin (Eln) alternative variant aSep08, mRNA.
<a href="#">Elof1</a>	<a href="#">Elof1.bSep08</a>	<a href="#">691193</a>	5709	725	3	83	elongation factor 1 homolog (S. cerevisiae) (9.5 kD) (Elof1) alternative variant bSep08, mRNA.
<a href="#">Elof1</a>	<a href="#">Elof1.cSep08</a>	<a href="#">691193</a>	4601	177	1	46	elongation factor 1 homolog (S. cerevisiae) (Elof1) alternative variant cSep08, mRNA.
<a href="#">Elov13</a>	<a href="#">Elov13.bSep08</a>	<a href="#">309449</a>	891	675	2	207	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 3 (Elov13) alternative variant bSep08, mRNA.
<a href="#">Elov14</a>	<a href="#">Elov14.aSep08</a>	<a href="#">315851</a>	17136	375	1	124	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 4 (Elov14) alternative variant aSep08, mRNA.

<a href="#">Elov4</a>	<a href="#">Elov4.bSep08</a>	<a href="#">315851</a>	16635	401	1	62	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 4 (Elov4) alternative variant bSep08, mRNA.
<a href="#">Elov6</a>	<a href="#">Elov6.aSep08</a>	<a href="#">171402</a>	106329	1190	4	267	ELOVL family member 6, elongation of long chain fatty acids (yeast) (31.6 kD) (Elov6) alternative variant aSep08, mRNA.
<a href="#">Elov7</a>	<a href="#">Elov7.aSep08</a>	<a href="#">361895</a>	68081	958	3	265	ELOVL family member 7, elongation of long chain fatty acids (yeast) (Elov7) alternative variant aSep08, mRNA.
<a href="#">Elov7</a>	<a href="#">Elov7.bSep08</a>	<a href="#">361895</a>	58780	486	1	71	ELOVL family member 7, elongation of long chain fatty acids (yeast) (8.3 kD) (Elov7) alternative variant bSep08, mRNA.
<a href="#">Elp4</a>	<a href="#">Elp4.aSep08</a>	<a href="#">691133</a>	4426	267		84	elongation protein 4 homolog ( <i>S. cerevisiae</i> ) (Elp4) mRNA.
<a href="#">Elt1</a>	<a href="#">Elt1.bSep08</a>	<a href="#">64124</a>	60589	725	6	241	EGF calcium-binding (Elt1) alternative variant bSep08, mRNA.
<a href="#">Emcn</a>	<a href="#">Emcn.bSep08</a>	<a href="#">295490</a>	67535	644	8	214	endomucin (Emcn) alternative variant bSep08, mRNA.
<a href="#">Eme1</a>	<a href="#">Eme1.bSep08</a>	<a href="#">287634</a>	3206	765	1	246	essential meiotic endonuclease 1 homolog 1 ( <i>S. pombe</i> ) (Eme1) alternative variant bSep08, mRNA.
<a href="#">EMI.0</a>	<a href="#">EMI.0.aSep08</a>		45785	508		169	MEGF11 (EMI.0) mRNA.
<a href="#">Emid1</a>	<a href="#">Emid1.bSep08</a>	<a href="#">685462</a>	18217	483	5	161	EMI (Emid1) alternative variant bSep08, mRNA.
<a href="#">Emilin1</a>	<a href="#">Emilin1.bSep08</a>	<a href="#">298845</a>	5226	2628	3	867	elastin microfibril interfacier 1 (Emilin1) alternative variant bSep08, mRNA.
<a href="#">Emilin1</a>	<a href="#">Emilin1.cSep08</a>	<a href="#">298845</a>	897	391	1	38	elastin microfibril interfacier 1 (Emilin1) alternative variant cSep08, mRNA.
<a href="#">Emilin2</a>	<a href="#">Emilin2.aSep08</a>	<a href="#">316736</a>	18000	1093		363	elastin microfibril interfacier 2 (Emilin2) mRNA.
<a href="#">Eml1</a>	<a href="#">Eml1.aSep08</a>	<a href="#">362783</a>	70972	760	2	187	echinoderm microtubule associated protein like 1 (20.0 kD) (Eml1) alternative variant aSep08, mRNA.
<a href="#">Eml1</a>	<a href="#">Eml1.cSep08</a>	<a href="#">362783</a>	96789	535	2	150	echinoderm microtubule associated protein like 1 (Eml1) alternative variant cSep08, mRNA.
<a href="#">Eml2</a>	<a href="#">Eml2.aSep08</a>	<a href="#">192360</a>	26315	2247	19	656	echinoderm microtubule associated protein like 2 (Eml2) alternative variant aSep08, mRNA.
<a href="#">Eml2</a>	<a href="#">Eml2.bSep08</a>	<a href="#">192360</a>	5818	561	6	186	echinoderm microtubule associated protein like 2 (Eml2) alternative variant bSep08, mRNA.
<a href="#">Eml2</a>	<a href="#">Eml2.cSep08</a>	<a href="#">192360</a>	1591	547	4	149	echinoderm microtubule associated protein like 2 (Eml2) alternative variant cSep08, mRNA.
<a href="#">Eml2</a>	<a href="#">Eml2.dSep08</a>	<a href="#">192360</a>	600	426	2	136	echinoderm microtubule associated protein like 2 (Eml2) alternative variant dSep08, mRNA.
<a href="#">Eml2</a>	<a href="#">Eml2.eSep08</a>	<a href="#">192360</a>	1803	368	3	107	echinoderm microtubule associated protein like 2 (Eml2) alternative variant eSep08, mRNA.
<a href="#">Eml4</a>	<a href="#">Eml4.bSep08</a>	<a href="#">313861</a>	6494	440	3	146	echinoderm microtubule associated protein like 4 (Eml4) alternative variant bSep08, mRNA.
<a href="#">Eml4</a>	<a href="#">Eml4.cSep08</a>	<a href="#">313861</a>	2970	727	3	38	echinoderm microtubule associated protein like 4 (Eml4) alternative variant cSep08, mRNA.
<a href="#">Eml5</a>	<a href="#">Eml5.aSep08</a>	<a href="#">444982</a>	4130	384		127	echinoderm microtubule associated protein like 5 (Eml5) mRNA.
<a href="#">Emr1</a>	<a href="#">Emr1.bSep08</a>	<a href="#">316137</a>	3995	593	1	62	EGF-like module containing, mucin-like, hormone receptor-like sequence 1 (7.0 kD) (Emr1) alternative variant bSep08, mRNA.



<a href="#">Emx2</a>	<a href="#">Emx2.bSep08</a>	<a href="#">499380</a>	979	412	1	67	empty spiracles homeobox 2 (Emx2) alternative variant bSep08, mRNA.
<a href="#">Enah</a>	<a href="#">Enah.bSep08</a>	<a href="#">360891</a>	96802	1359	7	375	enabled homolog (Drosophila) (Enah) alternative variant bSep08, mRNA.
<a href="#">Enah</a>	<a href="#">Enah.cSep08</a>	<a href="#">360891</a>	1442	733	2	132	enabled homolog (Drosophila) (Enah) alternative variant cSep08, mRNA.
<a href="#">Enah</a>	<a href="#">Enah.dSep08</a>	<a href="#">360891</a>	28417	328	4	108	enabled homolog (Drosophila) (Enah) alternative variant dSep08, mRNA.
<a href="#">Enah</a>	<a href="#">Enah.eSep08</a>	<a href="#">360891</a>	1889	291	2	96	enabled homolog (Drosophila) (Enah) alternative variant eSep08, mRNA.
<a href="#">Enah</a>	<a href="#">Enah.fSep08</a>	<a href="#">360891</a>	7447	255	2	77	enabled homolog (Drosophila) (Enah) alternative variant fSep08, mRNA.
<a href="#">Eno1</a>	<a href="#">Eno1.cSep08</a>	<a href="#">24333</a>	7981	933	7	238	enolase 1, alpha non-neuron (25.6 kD) (Eno1) alternative variant cSep08, mRNA.
<a href="#">Eno1</a>	<a href="#">Eno1.dSep08</a>	<a href="#">24333</a>	4096	389	3	129	enolase 1, alpha non-neuron (Eno1) alternative variant dSep08, mRNA.
<a href="#">Eno1</a>	<a href="#">Eno1.eSep08</a>	<a href="#">24333</a>	3105	806	4	121	enolase 1, alpha non-neuron (Eno1) alternative variant eSep08, mRNA.
<a href="#">Eno2</a>	<a href="#">Eno2.bSep08</a>	<a href="#">24334</a>	8840	1318	11	404	enolase 2 (43.6 kD) (Eno2) alternative variant bSep08, complete mRNA.
<a href="#">Eno2</a>	<a href="#">Eno2.cSep08</a>	<a href="#">24334</a>	2451	883	4	146	enolase 2 (16.2 kD) (Eno2) alternative variant cSep08, mRNA.
<a href="#">Eno2</a>	<a href="#">Eno2.dSep08</a>	<a href="#">24334</a>	2222	398	2	128	enolase 2 (Eno2) alternative variant dSep08, mRNA.
<a href="#">Eno2</a>	<a href="#">Eno2.eSep08</a>	<a href="#">24334</a>	2489	581	5	100	enolase 2 (Eno2) alternative variant eSep08, mRNA.
<a href="#">Eno2</a>	<a href="#">Eno2.fSep08</a>	<a href="#">24334</a>	454	286	2	89	putative protein (Eno2) alternative variant fSep08, mRNA.
<a href="#">Eno3</a>	<a href="#">Eno3.bSep08</a>	<a href="#">25438</a>	3677	715	7	218	enolase 3, beta, muscle (Eno3) alternative variant bSep08, mRNA.
<a href="#">Eno3</a>	<a href="#">Eno3.cSep08</a>	<a href="#">25438</a>	575	380	2	126	enolase 3, beta, muscle (Eno3) alternative variant cSep08, mRNA.
<a href="#">Eno3</a>	<a href="#">Eno3.dSep08</a>	<a href="#">25438</a>	2998	747	5	115	enolase 3, beta, muscle (Eno3) alternative variant dSep08, mRNA.
<a href="#">Eno3</a>	<a href="#">Eno3.eSep08</a>	<a href="#">25438</a>	1219	712	4	102	enolase 3, beta, muscle (Eno3) alternative variant eSep08, mRNA.
<a href="#">Eno3</a>	<a href="#">Eno3.fSep08</a>	<a href="#">25438</a>	1147	476	4	78	enolase 3, beta, muscle (Eno3) alternative variant fSep08, mRNA.
<a href="#">Enoph1</a>	<a href="#">Enoph1.bSep08</a>	<a href="#">305177</a>	19813	597	1	84	enolase-phosphatase 1 (Enoph1) alternative variant bSep08, mRNA.
<a href="#">Enox1</a>	<a href="#">Enox1.aSep08</a>	<a href="#">306038</a>	49365	822		126	ecto-NOX disulfide-thiol exchanger 1 (Enox1) mRNA.
<a href="#">Enpp1</a>	<a href="#">Enpp1.bSep08</a>	<a href="#">85496</a>	13060	2165	2	309	ectonucleotide pyrophosphatase/phosphodiesterase 1 (Enpp1) alternative variant bSep08, mRNA.
<a href="#">Enpp2</a>	<a href="#">Enpp2.aSep08</a>	<a href="#">84050</a>	125394	3350	25	887	pyrophosphatase phosphodiesterase (Enpp2) alternative variant aSep08, mRNA.
<a href="#">Enpp2</a>	<a href="#">Enpp2.cSep08</a>	<a href="#">84050</a>	26360	1697	10	408	pyrophosphatase phosphodiesterase (Enpp2) alternative variant cSep08, mRNA.
<a href="#">Enpp2</a>	<a href="#">Enpp2.dSep08</a>	<a href="#">84050</a>	30453	1208	9	380	ectonucleotide pyrophosphatase phosphodiesterase 2 CRA a (Enpp2) alternative variant dSep08, mRNA.

<a href="#">Enpp2</a>	<a href="#">Enpp2.eSep08</a>	<a href="#">84050</a>	81239	1203	11	356	ectonucleotide pyrophosphatase phosphodiesterase 2 (Enpp2) alternative variant eSep08, mRNA.
<a href="#">Enpp3</a>	<a href="#">Enpp3.bSep08</a>	<a href="#">54410</a>	3556	713	3	124	ectonucleotide pyrophosphatase/phosphodiesterase 3 (Enpp3) alternative variant bSep08, mRNA.
<a href="#">Enpp4</a>	<a href="#">Enpp4.bSep08</a>	<a href="#">301261</a>	1840	918	1	305	ectonucleotide pyrophosphatase/phosphodiesterase 4 (Enpp4) alternative variant bSep08, mRNA.
<a href="#">Enpp4</a>	<a href="#">Enpp4.cSep08</a>	<a href="#">301261</a>	3448	483	1	160	ectonucleotide pyrophosphatase/phosphodiesterase 4 (Enpp4) alternative variant cSep08, mRNA.
<a href="#">Enpp5</a>	<a href="#">Enpp5.cSep08</a>	<a href="#">316249</a>	1779	385	2	99	ectonucleotide pyrophosphatase/phosphodiesterase 5 (Enpp5) alternative variant cSep08, mRNA.
<a href="#">Enpp6</a>	<a href="#">Enpp6.aSep08</a>	<a href="#">306460</a>	124226	1503	8	440	ectonucleotide pyrophosphatase/phosphodiesterase 6 (50.7 kD) (Enpp6) alternative variant aSep08, complete mRNA.
<a href="#">ENT.0</a>	<a href="#">ENT.0.aSep08</a>		10573	381		91	putative protein (10.4 kD) (ENT.0) mRNA.
<a href="#">Enth</a>	<a href="#">Enth.bSep08</a>	<a href="#">360515</a>	14413	1362	6	384	clathrin interactor 1 (Enth) alternative variant bSep08, mRNA.
<a href="#">Enth</a>	<a href="#">Enth.cSep08</a>	<a href="#">360515</a>	15583	2478	6	384	clathrin interactor 1 (Enth) alternative variant cSep08, mRNA.
<a href="#">Enth</a>	<a href="#">Enth.dSep08</a>	<a href="#">360515</a>	3181	750	3	244	clathrin interactor 1 (Enth) alternative variant dSep08, mRNA.
<a href="#">Enth</a>	<a href="#">Enth.eSep08</a>	<a href="#">360515</a>	28661	373	3	63	enthoprotin (Enth) alternative variant eSep08, mRNA.
<a href="#">Enth</a>	<a href="#">Enth.gSep08</a>	<a href="#">360515</a>	1776	769	2	62	CRA b (Enth) alternative variant gSep08, mRNA.
<a href="#">ENTH.1</a>	<a href="#">ENTH.1.aSep08</a>		36411	553	4	162	synaptosomal-associated protein 91kDa homolog (ENTH.1) alternative variant aSep08, mRNA.
<a href="#">ENTH.1</a>	<a href="#">ENTH.1.bSep08</a>		34472	450	2	73	synaptosomal-associated protein 91kDa homolog CRA d (ENTH.1) alternative variant bSep08, mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.aSep08</a>	<a href="#">685815</a>	114318	2672	7	616	hypothetical protein LOC685815 and hypothetical protein LOC685869 (67.9 kD) (Enthd1) alternative variant aSep08, complete mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.aSep08</a>	<a href="#">685869</a>	114318	2672	7	616	hypothetical protein LOC685815 and hypothetical protein LOC685869 (67.9 kD) (Enthd1) alternative variant aSep08, complete mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.aSep08</a>	<a href="#">685900</a>	114318	2672	7	616	hypothetical protein LOC685815 and hypothetical protein LOC685869 (67.9 kD) (Enthd1) alternative variant aSep08, complete mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.bSep08</a>	<a href="#">685815</a>	19291	345	3	62	hypothetical protein LOC685815 and hypothetical protein LOC685869 (Enthd1) alternative variant bSep08, mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.bSep08</a>	<a href="#">685869</a>	19291	345	3	62	hypothetical protein LOC685815 and hypothetical protein LOC685869 (Enthd1) alternative variant bSep08, mRNA.
<a href="#">Enthd1</a>	<a href="#">Enthd1.bSep08</a>	<a href="#">685900</a>	19291	345	3	62	hypothetical protein LOC685815 and hypothetical protein LOC685869 (Enthd1) alternative variant bSep08, mRNA.
<a href="#">Entpd2</a>	<a href="#">Entpd2.aSep08</a>	<a href="#">64467</a>	2702	1783	5	592	ectonucleoside triphosphate diphosphohydrolase 2 (Entpd2) alternative variant aSep08, mRNA.
<a href="#">Entpd2</a>	<a href="#">Entpd2.cSep08</a>	<a href="#">64467</a>	2400	1108	6	350	ectonucleoside triphosphate diphosphohydrolase 2 (Entpd2) alternative variant cSep08, mRNA.
<a href="#">Entpd2</a>	<a href="#">Entpd2.dSep08</a>	<a href="#">64467</a>	1949	594	3	82	ectonucleoside triphosphate diphosphohydrolase 2 (12.4 kD) (Entpd2) alternative variant dSep08, mRNA.

<a href="#">Entpd4</a>	<a href="#">Entpd4.aSep08</a>	<a href="#">361063</a>	11026	1560	8	410	ectonucleoside triphosphate diphosphohydrolase 4 CRA b (Entpd4) alternative variant aSep08, mRNA.
<a href="#">Entpd4</a>	<a href="#">Entpd4.cSep08</a>	<a href="#">361063</a>	17958	1278	8	309	ectonucleoside triphosphate diphosphohydrolase 4 CRA b (34.4 kD) (Entpd4) alternative variant cSep08, mRNA.
<a href="#">Entpd4</a>	<a href="#">Entpd4.dSep08</a>	<a href="#">361063</a>	2484	785	3	133	ectonucleoside triphosphate diphosphohydrolase 4 CRA b (Entpd4) alternative variant dSep08, mRNA.
<a href="#">Entpd4</a>	<a href="#">Entpd4.eSep08</a>	<a href="#">361063</a>	1329	769	2	90	ectonucleoside triphosphate diphosphohydrolase 4 CRA b (Entpd4) alternative variant eSep08, mRNA.
<a href="#">Entpd4</a>	<a href="#">Entpd4.iSep08</a>	<a href="#">361063</a>	9724	374	3	79	putative protein (Entpd4) alternative variant iSep08, mRNA.
<a href="#">Entpd6</a>	<a href="#">Entpd6.bSep08</a>	<a href="#">85260</a>	7784	510	4	153	ectonucleoside triphosphate diphosphohydrolase 6 (Entpd6) alternative variant bSep08, mRNA.
<a href="#">Entpd6</a>	<a href="#">Entpd6.cSep08</a>	<a href="#">85260</a>	4177	747	5	144	ectonucleoside triphosphate diphosphohydrolase 6 (16.2 kD) (Entpd6) alternative variant cSep08, mRNA.
<a href="#">Entpd7</a>	<a href="#">Entpd7.bSep08</a>	<a href="#">309390</a>	4793	712	3	101	ectonucleoside triphosphate diphosphohydrolase 7 (Entpd7) alternative variant bSep08, mRNA.
<a href="#">Entpd8</a>	<a href="#">Entpd8.bSep08</a>	<a href="#">613267</a>	802	488	1	162	ectonucleoside triphosphate diphosphohydrolase 8 (Entpd8) alternative variant bSep08, mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.bSep08</a>	<a href="#">685258</a>	7348	525	5	96	enhancer of yellow 2 homolog (Drosophila) (11.0 kD) (Eny2) alternative variant bSep08, complete mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.cSep08</a>	<a href="#">685258</a>	7298	472	4	93	enhancer of yellow 2 homolog (Drosophila) (Eny2) alternative variant cSep08, mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.eSep08</a>	<a href="#">685258</a>	5987	551	4	54	enhancer of yellow 2 homolog (Drosophila) (Eny2) alternative variant eSep08, mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.fSep08</a>	<a href="#">685258</a>	4520	248	4	43	enhancer of yellow 2 homolog (Drosophila) (Eny2) alternative variant fSep08, mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.gSep08</a>	<a href="#">685258</a>	2522	1071	3	36	enhancer of yellow 2 homolog (Drosophila) (4.4 kD) (Eny2) alternative variant gSep08, mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.iSep08</a>	<a href="#">685258</a>	7384	559	5	48	enhancer of yellow 2 homolog (Drosophila) (5.6 kD) (Eny2) alternative variant iSep08, complete mRNA.
<a href="#">Eny2</a>	<a href="#">Eny2.jSep08</a>	<a href="#">685258</a>	7339	511	4	9	enhancer of yellow 2 homolog (Drosophila) (1.0 kD) (Eny2) alternative variant jSep08, complete mRNA.
<a href="#">Ep300</a>	<a href="#">Ep300.aSep08</a>	<a href="#">170915</a>	3591	3409		775	E1A binding protein p300 (Ep300) mRNA.
<a href="#">Ep400</a>	<a href="#">Ep400.bSep08</a>	<a href="#">304569</a>	12622	1615	2	434	E1A binding protein p400 (Ep400) alternative variant bSep08, mRNA.
<a href="#">Ep400</a>	<a href="#">Ep400.cSep08</a>	<a href="#">304569</a>	6442	725	5	241	E1A binding protein p400 (Ep400) alternative variant cSep08, mRNA.
<a href="#">Ep400</a>	<a href="#">Ep400.dSep08</a>	<a href="#">304569</a>	1184	684	2	160	E1A binding protein p400 (Ep400) alternative variant dSep08, mRNA.
<a href="#">Ep400</a>	<a href="#">Ep400.eSep08</a>	<a href="#">304569</a>	686	523	2	111	E1A binding protein p400 (Ep400) alternative variant eSep08, mRNA.
<a href="#">Epb4.1</a>	<a href="#">Epb4.1.aSep08</a>	<a href="#">313052</a>	154264	5007	18	838	erythrocyte membrane protein band 4.1 (Epb4.1) alternative variant aSep08, mRNA.
<a href="#">Epb4.1</a>	<a href="#">Epb4.1.bSep08</a>	<a href="#">313052</a>	48022	749	3	213	erythrocyte membrane protein band 4.1 (Epb4.1) alternative variant bSep08, mRNA.
<a href="#">Epb4.1</a>	<a href="#">Epb4.1.cSep08</a>	<a href="#">313052</a>	13004	1333	6	201	erythrocyte membrane protein band 4.1 (Epb4.1) alternative variant cSep08, mRNA.

<a href="#">Epb4.111</a>	<a href="#">Epb4.111.cSep08</a>	<a href="#">59317</a>	35668	1771	5	590	erythrocyte protein band 4.1-like 1 (Epb4.111) alternative variant cSep08, mRNA.
<a href="#">Epb4.111</a>	<a href="#">Epb4.111.dSep08</a>	<a href="#">59317</a>	30663	1145	11	381	erythrocyte protein band 4.1-like 1 (Epb4.111) alternative variant dSep08, mRNA.
<a href="#">Epb4.112</a>	<a href="#">Epb4.112.aSep08</a>	<a href="#">309557</a>	43753	2777	10	512	erythrocyte membrane protein band 4.1-like 2 (Epb4.112) alternative variant aSep08, mRNA.
<a href="#">Epb4.112</a>	<a href="#">Epb4.112.bSep08</a>	<a href="#">309557</a>	20342	897	5	268	erythrocyte membrane protein band 4.1-like 2 (Epb4.112) alternative variant bSep08, mRNA.
<a href="#">Epb4.112</a>	<a href="#">Epb4.112.cSep08</a>	<a href="#">309557</a>	27475	534	4	177	erythrocyte membrane protein band 4.1-like 2 (Epb4.112) alternative variant cSep08, mRNA.
<a href="#">Epb4.112</a>	<a href="#">Epb4.112.dSep08</a>	<a href="#">309557</a>	5041	422	1	138	erythrocyte membrane protein band 4.1-like 2 (Epb4.112) alternative variant dSep08, mRNA.
<a href="#">Epb4.112</a>	<a href="#">Epb4.112.eSep08</a>	<a href="#">309557</a>	4103	369	3	123	erythrocyte membrane protein band 4.1-like 2 (Epb4.112) alternative variant eSep08, mRNA.
<a href="#">Epb4.113</a>	<a href="#">Epb4.113.bSep08</a>	<a href="#">116724</a>	19523	978	7	325	erythrocyte protein band 4.1-like 3 (Epb4.113) alternative variant bSep08, mRNA.
<a href="#">Epb4.113</a>	<a href="#">Epb4.113.cSep08</a>	<a href="#">116724</a>	5271	1249	5	214	erythrocyte protein band 4.1-like 3 (Epb4.113) alternative variant cSep08, mRNA.
<a href="#">Epb4.113</a>	<a href="#">Epb4.113.dSep08</a>	<a href="#">116724</a>	20555	610	5	202	erythrocyte protein band 4.1-like 3 (Epb4.113) alternative variant dSep08, mRNA.
<a href="#">Epb4.113</a>	<a href="#">Epb4.113.eSep08</a>	<a href="#">116724</a>	154092	804	5	190	erythrocyte protein band 4.1-like 3 (Epb4.113) alternative variant eSep08, mRNA.
<a href="#">Epb4.113</a>	<a href="#">Epb4.113.fSep08</a>	<a href="#">116724</a>	71432	574	4	164	erythrocyte protein band 4.1-like 3 (Epb4.113) alternative variant fSep08, mRNA.
<a href="#">Epb4.114a</a>	<a href="#">Epb4.114a.bSep08</a>	<a href="#">307514</a>	8314	544	5	122	erythrocyte protein band 4.1-like 4a (Epb4.114a) alternative variant bSep08, mRNA.
<a href="#">Epb4.114b</a>	<a href="#">Epb4.114b.aSep08</a>	<a href="#">366376</a>	16816	386		128	erythrocyte protein band 4.1-like 4b (Epb4.114b) mRNA.
<a href="#">Epb4.115</a>	<a href="#">Epb4.115.bSep08</a>	<a href="#">304733</a>	26166	445	2	148	erythrocyte protein band 4.1-like 5 (Epb4.115) alternative variant bSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.aSep08</a>	<a href="#">361069</a>	24638	3390	14	354	erythrocyte protein band 4.9 (39.5 kD) (Epb4.9) alternative variant aSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.bSep08</a>	<a href="#">361069</a>	26873	1858	16	343	erythrocyte protein band 4.9 (37.9 kD) (Epb4.9) alternative variant bSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.dSep08</a>	<a href="#">361069</a>	13801	715	7	173	erythrocyte protein band 4.9 (Epb4.9) alternative variant dSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.eSep08</a>	<a href="#">361069</a>	18092	423	4	140	erythrocyte protein band 4.9 (Epb4.9) alternative variant eSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.fSep08</a>	<a href="#">361069</a>	8853	341	3	65	erythrocyte protein band 4.9 (Epb4.9) alternative variant fSep08, mRNA.
<a href="#">Epb4.9</a>	<a href="#">Epb4.9.gSep08</a>	<a href="#">361069</a>	10679	318	3	55	erythrocyte protein band 4.9 (Epb4.9) alternative variant gSep08, mRNA.
<a href="#">Epc1</a>	<a href="#">Epc1.aSep08</a>	<a href="#">307042</a>	3203	794		146	enhancer of polycomb homolog 1 (Drosophila) (Epc1) mRNA.
<a href="#">Epc2</a>	<a href="#">Epc2.bSep08</a>	<a href="#">362132</a>	4399	1223	4	282	enhancer of polycomb homolog 2 (Drosophila) (Epc2) alternative variant bSep08, mRNA.
<a href="#">Epc2</a>	<a href="#">Epc2.cSep08</a>	<a href="#">362132</a>	2355	751	2	164	enhancer of polycomb homolog 2 (Drosophila) (18.2 kD) (Epc2) alternative variant cSep08, mRNA.

<a href="#">Epgn</a>	<a href="#">Epgn.aSep08</a>	<a href="#">289515</a>	5415	591		163	epithelial mitogen (Epgn) mRNA.
<a href="#">Epha2</a>	<a href="#">Epha2.bSep08</a>	<a href="#">366492</a>	8521	1732	6	287	eph receptor A2 (Epha2) alternative variant bSep08, mRNA.
<a href="#">Epha2</a>	<a href="#">Epha2.cSep08</a>	<a href="#">366492</a>	973	353	4	117	eph receptor A2 (Epha2) alternative variant cSep08, mRNA.
<a href="#">Epha4</a>	<a href="#">Epha4.aSep08</a>	<a href="#">316539</a>	74579	1784		404	eph receptor A4 (Epha4) alternative variant aSep08, mRNA.
<a href="#">Epha4</a>	<a href="#">Epha4.bSep08</a>	<a href="#">316539</a>	5861	2700		86	eph receptor A4 (10.0 kD) (Epha4) alternative variant bSep08, mRNA.
<a href="#">Epha5</a>	<a href="#">Epha5.aSep08</a>	<a href="#">79208</a>	34040	1783		558	EphA5 (Epha5) alternative variant aSep08, mRNA.
<a href="#">Epha5</a>	<a href="#">Epha5.bSep08</a>	<a href="#">79208</a>	10445	396		131	EphA5 (Epha5) alternative variant bSep08, mRNA.
<a href="#">Epha6</a>	<a href="#">Epha6.aSep08</a>	<a href="#">29202</a>	33050	1049		281	eph receptor A6 (Epha6) mRNA.
<a href="#">Epha7</a>	<a href="#">Epha7.bSep08</a>	<a href="#">171287</a>	13016	865	8	277	eph receptor A7 (Epha7) alternative variant bSep08, mRNA.
<a href="#">Epha7</a>	<a href="#">Epha7.cSep08</a>	<a href="#">171287</a>	17402	756	7	187	eph receptor A7 (Epha7) alternative variant cSep08, mRNA.
<a href="#">Epha8</a>	<a href="#">Epha8.aSep08</a>	<a href="#">60589</a>	7486	487		161	eph receptor A8 (Epha8) mRNA.
<a href="#">Ephb3</a>	<a href="#">Ephb3.bSep08</a>	<a href="#">287989</a>	575	369	3	101	eph receptor B3 (Ephb3) alternative variant bSep08, mRNA.
<a href="#">Ephb6</a>	<a href="#">Ephb6.aSep08</a>	<a href="#">312275</a>	2525	1032	8	277	eph receptor B6 (Ephb6) alternative variant aSep08, mRNA.
<a href="#">Ephb6</a>	<a href="#">Ephb6.bSep08</a>	<a href="#">312275</a>	1720	740	5	246	eph receptor B6 (Ephb6) alternative variant bSep08, mRNA.
<a href="#">Ephb6</a>	<a href="#">Ephb6.cSep08</a>	<a href="#">312275</a>	823	470	2	81	eph receptor B6 (Ephb6) alternative variant cSep08, mRNA.
<a href="#">Ephrin_lbd.0</a>	<a href="#">Ephrin_lbd.0.aSep08</a>		25210	735		158	EPH receptor A5 CRA c (Ephrin_lbd.0) mRNA.
<a href="#">Ephrin_lbd.1</a>	<a href="#">Ephrin_lbd.1.aSep08</a>		8046	405		124	epha4 (Ephrin_lbd.1) mRNA.
<a href="#">Ephx1</a>	<a href="#">Ephx1.cSep08</a>	<a href="#">25315</a>	23739	747	1	152	epoxide hydrolase 1, microsomal (Ephx1) alternative variant cSep08, mRNA.
<a href="#">Ephx2</a>	<a href="#">Ephx2.bSep08</a>	<a href="#">65030</a>	4771	864	7	166	epoxide hydrolase (Ephx2) alternative variant bSep08, mRNA.
<a href="#">Ephx2</a>	<a href="#">Ephx2.cSep08</a>	<a href="#">65030</a>	3771	1263	6	131	epoxide hydrolase (15.3 kD) (Ephx2) alternative variant cSep08, mRNA.
<a href="#">EPL1.0</a>	<a href="#">EPL1.0.aSep08</a>		9958	265		88	enhancer of polycomb 1 (EPL1.0) mRNA.
<a href="#">EPL1.1</a>	<a href="#">EPL1.1.bSep08</a>		1606	477	2	63	enhancer of polycomb homolog 1 (7.1 kD) (EPL1.1) alternative variant bSep08, mRNA.
<a href="#">EPL1.2</a>	<a href="#">EPL1.2.aSep08</a>		4118	762		254	bromodomain PHD finger containing 1 CRA d (EPL1.2) mRNA.
<a href="#">Epn1</a>	<a href="#">Epn1.bSep08</a>	<a href="#">117277</a>	6014	734	2	244	epsin 1 (Epn1) alternative variant bSep08, mRNA.
<a href="#">Epn1</a>	<a href="#">Epn1.cSep08</a>	<a href="#">117277</a>	4509	733	4	244	epsin 1 (Epn1) alternative variant cSep08, mRNA.
<a href="#">Epn2</a>	<a href="#">Epn2.cSep08</a>	<a href="#">60443</a>	23651	852	6	283	epsin 2 (Epn2) alternative variant cSep08, mRNA.
<a href="#">Epn3</a>	<a href="#">Epn3.bSep08</a>	<a href="#">360605</a>	1430	1248	3	358	epsin 3 (Epn3) alternative variant bSep08, mRNA.
<a href="#">Epn3</a>	<a href="#">Epn3.cSep08</a>	<a href="#">360605</a>	3763	784	2	162	epsin 3 (Epn3) alternative variant cSep08, mRNA.
<a href="#">Epor</a>	<a href="#">Epor.bSep08</a>	<a href="#">24336</a>	1757	1075	2	292	erythropoietin receptor (31.7 kD) (Epor) alternative variant bSep08, mRNA.

<a href="#">Epor</a>	<a href="#">Epor.cSep08</a>	<a href="#">24336</a>	1755	682	4	171	erythropoietin receptor (Epor) alternative variant cSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.bSep08</a>	<a href="#">289352</a>	33418	1988	16	622	glutaryl-prolyl-tRNA synthetase (Eprs) alternative variant bSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.cSep08</a>	<a href="#">289352</a>	6579	1114	3	268	glutaryl-prolyl-tRNA synthetase (Eprs) alternative variant cSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.dSep08</a>	<a href="#">289352</a>	3238	830	5	159	glutaryl-prolyl-tRNA synthetase (Eprs) alternative variant dSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.eSep08</a>	<a href="#">289352</a>	8591	329	4	109	glutaryl-prolyl-tRNA synthetase (Eprs) alternative variant eSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.gSep08</a>	<a href="#">289352</a>	2842	574	2	55	glutaryl-prolyl-tRNA synthetase (Eprs) alternative variant gSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.hSep08</a>	<a href="#">289352</a>	1430	544	2	49	putative protein (5.3 kD) (Eprs) alternative variant hSep08, mRNA.
<a href="#">Eprs</a>	<a href="#">Eprs.iSep08</a>	<a href="#">289352</a>	9494	406	4	47	putative protein (5.4 kD) (Eprs) alternative variant iSep08, mRNA.
<a href="#">Eps8</a>	<a href="#">Eps8.aSep08</a>	<a href="#">312812</a>	50546	2211		619	epidermal growth factor receptor pathway substrate 8 (Eps8) alternative variant aSep08, mRNA.
<a href="#">Eps8</a>	<a href="#">Eps8.bSep08</a>	<a href="#">312812</a>	13815	1968		213	epidermal growth factor receptor pathway substrate 8 (Eps8) alternative variant bSep08, mRNA.
<a href="#">Eps8l2</a>	<a href="#">Eps8l2.aSep08</a>	<a href="#">361674</a>	25481	3282	21	729	epidermal growth factor receptor pathway substrate 8-like protein 2 (82.3 kD) (Eps8l2) alternative variant aSep08, mRNA.
<a href="#">Eps8l2</a>	<a href="#">Eps8l2.cSep08</a>	<a href="#">361674</a>	24829	1186	9	320	epidermal growth factor receptor pathway substrate 8-like protein 2 (36.3 kD) (Eps8l2) alternative variant cSep08, mRNA.
<a href="#">Eps8l2</a>	<a href="#">Eps8l2.dSep08</a>	<a href="#">361674</a>	1609	765	5	195	epidermal growth factor receptor pathway substrate 8-like protein 2 (Eps8l2) alternative variant dSep08, mRNA.
<a href="#">Eps8l2</a>	<a href="#">Eps8l2.eSep08</a>	<a href="#">361674</a>	507	421	2	111	epidermal growth factor receptor pathway substrate 8-like protein 2 (Eps8l2) alternative variant eSep08, mRNA.
<a href="#">Eps8l2</a>	<a href="#">Eps8l2.fSep08</a>	<a href="#">361674</a>	17425	415	4	43	2 CRA b (4.4 kD) (Eps8l2) alternative variant fSep08, mRNA.
<a href="#">Eps15</a>	<a href="#">Eps15.bSep08</a>	<a href="#">313474</a>	101245	5067	25	693	epidermal growth factor receptor pathway substrate 15 (Eps15) alternative variant bSep08, mRNA.
<a href="#">Eps15</a>	<a href="#">Eps15.cSep08</a>	<a href="#">313474</a>	11129	516	3	132	epidermal growth factor receptor pathway substrate 15 like (Eps15) alternative variant cSep08, mRNA.
<a href="#">Eps15</a>	<a href="#">Eps15.dSep08</a>	<a href="#">313474</a>	15561	738	5	128	epidermal growth factor receptor pathway substrate 15 (Eps15) alternative variant dSep08, mRNA.
<a href="#">Eps15</a>	<a href="#">Eps15.eSep08</a>	<a href="#">313474</a>	1835	515	2	82	epidermal growth factor receptor pathway substrate 15 (Eps15) alternative variant eSep08, mRNA.
<a href="#">Eps15</a>	<a href="#">Eps15.fSep08</a>	<a href="#">313474</a>	3720	755	3	74	epidermal growth factor receptor pathway substrate 15 (Eps15) alternative variant fSep08, mRNA.
<a href="#">Eps15l1</a>	<a href="#">Eps15l1.bSep08</a>	<a href="#">361120</a>	43396	1785	8	464	epidermal growth factor receptor pathway substrate 15-like 1 CRA a (Eps15l1) alternative variant bSep08, mRNA.
<a href="#">Eps15l1</a>	<a href="#">Eps15l1.cSep08</a>	<a href="#">361120</a>	3877	868	5	220	epidermal growth factor receptor pathway substrate 15-like 1 (Eps15l1) alternative variant cSep08, mRNA.

<a href="#">Eps151</a>	<a href="#">Eps151.dSep08</a>	<a href="#">361120</a>	7203	470	4	156	epidermal growth factor receptor pathway substrate 15-like 1 (Eps151) alternative variant dSep08, mRNA.
<a href="#">Eps151</a>	<a href="#">Eps151.eSep08</a>	<a href="#">361120</a>	10876	821	6	129	putative protein (14.3 kD) (Eps151) alternative variant eSep08, mRNA.
<a href="#">Eps151</a>	<a href="#">Eps151.fSep08</a>	<a href="#">361120</a>	30285	980	4	81	epidermal growth factor receptor pathway substrate 15-like 1 CRA c (8.9 kD) (Eps151) alternative variant fSep08, mRNA.
<a href="#">Eps151</a>	<a href="#">Eps151.gSep08</a>	<a href="#">361120</a>	4482	603	2	38	putative protein (Eps151) alternative variant gSep08, mRNA.
<a href="#">Eps151</a>	<a href="#">Eps151.hSep08</a>	<a href="#">361120</a>	64712	358	2	22	putative protein (Eps151) alternative variant hSep08, mRNA.
<a href="#">EPTP.0</a>	<a href="#">EPTP.0.aSep08</a>		21852	1727		458	leucine-rich glioma inactivated 1 CRA a (EPTP.0) mRNA.
<a href="#">Epyc</a>	<a href="#">Epyc.bSep08</a>	<a href="#">314772</a>	34394	2055	2	322	epiphyccan (36.8 kD) (Epyc) alternative variant bSep08, mRNA.
<a href="#">Eral1</a>	<a href="#">Eral1.bSep08</a>	<a href="#">363646</a>	2543	1514	1	259	era (G-protein)-like 1 (E. coli) (Eral1) alternative variant bSep08, mRNA.
<a href="#">Erap1</a>	<a href="#">Erap1.aSep08</a>	<a href="#">80897</a>	38836	2953	19	930	endoplasmic reticulum aminopeptidase 1 (106.4 kD) (Erap1) alternative variant aSep08, complete mRNA.
<a href="#">Erap1</a>	<a href="#">Erap1.bSep08</a>	<a href="#">80897</a>	16837	784	6	245	endoplasmic reticulum aminopeptidase 1 (Erap1) alternative variant bSep08, mRNA.
<a href="#">Erap1</a>	<a href="#">Erap1.cSep08</a>	<a href="#">80897</a>	5280	819	6	223	endoplasmic reticulum aminopeptidase 1 (Erap1) alternative variant cSep08, mRNA.
<a href="#">Erb2</a>	<a href="#">Erb2.bSep08</a>	<a href="#">24337</a>	11181	1055	7	351	v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian) (Erb2) alternative variant bSep08, mRNA.
<a href="#">Erb2</a>	<a href="#">Erb2.cSep08</a>	<a href="#">24337</a>	514	386	2	44	v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian) (Erb2) alternative variant cSep08, mRNA.
<a href="#">Erc1</a>	<a href="#">Erc1.cSep08</a>	<a href="#">266806</a>	27732	600	3	98	rab6 interacting protein 2 CRA b (Erc1) alternative variant cSep08, mRNA.
<a href="#">Erc1</a>	<a href="#">Erc1.dSep08</a>	<a href="#">266806</a>	4907	424	2	73	rab6 interacting protein 2 (Erc1) alternative variant dSep08, mRNA.
<a href="#">Ercc1</a>	<a href="#">Ercc1.aSep08</a>	<a href="#">292673</a>	10870	1524	7	471	excision repair cross-complementing rodent repair deficiency, complementation group 1 (Ercc1) alternative variant aSep08, mRNA.
<a href="#">Ercc1</a>	<a href="#">Ercc1.cSep08</a>	<a href="#">292673</a>	4098	395	1	131	excision repair cross-complementing rodent repair deficiency, complementation group 1 (Ercc1) alternative variant cSep08, mRNA.
<a href="#">Ercc3</a>	<a href="#">Ercc3.bSep08</a>	<a href="#">291703</a>	8890	737	5	195	excision repair cross-complementing rodent repair deficiency, complementation group 3 (Ercc3) alternative variant bSep08, mRNA.
<a href="#">Ercc4</a>	<a href="#">Ercc4.aSep08</a>	<a href="#">304719</a>	10178	906	5	302	excision repair cross-complementing rodent repair deficiency, complementation group 4 (Ercc4) alternative variant aSep08, mRNA.
<a href="#">ERCC4.1</a>	<a href="#">ERCC4.1.aSep08</a>		9828	1744		304	excision repair cross-complementing rodent deficiency complementation group 4 CRA b (ERCC4.1) mRNA.

<a href="#">Ercc6</a>	<a href="#">Ercc6.bSep08</a>	<a href="#">306274</a>	7576	1070	4	356	excision repair cross-complementing rodent repair deficiency, complementation group 6 (Ercc6) alternative variant bSep08, mRNA.
<a href="#">Ercc6</a>	<a href="#">Ercc6.cSep08</a>	<a href="#">306274</a>	13373	994	8	331	excision repair cross-complementing rodent repair deficiency, complementation group 6 (Ercc6) alternative variant cSep08, mRNA.
<a href="#">Ercc6</a>	<a href="#">Ercc6.dSep08</a>	<a href="#">306274</a>	3416	793	4	247	excision repair cross-complementing rodent repair deficiency, complementation group 6 (26.5 kD) (Ercc6) alternative variant dSep08, mRNA.
<a href="#">Ercc8</a>	<a href="#">Ercc8.aSep08</a>	<a href="#">310071</a>	36904	1606	13	201	excision repair cross-complementing rodent repair deficiency, complementation group 8 (Ercc8) alternative variant aSep08, mRNA.
<a href="#">Ercc8</a>	<a href="#">Ercc8.bSep08</a>	<a href="#">310071</a>	18710	874	6	194	excision repair cross-complementing rodent repair deficiency, complementation group 8 (Ercc8) alternative variant bSep08, mRNA.
<a href="#">Ercc8</a>	<a href="#">Ercc8.cSep08</a>	<a href="#">310071</a>	37825	2308	11	184	excision repair cross-complementing rodent repair deficiency, complementation group 8 (20.3 kD) (Ercc8) alternative variant cSep08, complete mRNA.
<a href="#">Ercc8</a>	<a href="#">Ercc8.dSep08</a>	<a href="#">310071</a>	7323	864	2	53	excision repair cross-complementing rodent repair deficiency, complementation group 8 (5.7 kD) (Ercc8) alternative variant dSep08, mRNA.
<a href="#">Erf</a>	<a href="#">Erf.aSep08</a>	<a href="#">292721</a>	2289	1832	3	519	ets2 repressor factor (Erf) alternative variant aSep08, mRNA.
<a href="#">Erg</a>	<a href="#">Erg.bSep08</a>	<a href="#">170909</a>	19383	1144	1	230	avian erythroblastosis virus E-26 (v-ets) oncogene related (Erg) alternative variant bSep08, mRNA.
<a href="#">Ergic1</a>	<a href="#">Ergic1.aSep08</a>	<a href="#">287177</a>	26888	2415		178	endoplasmic reticulum-golgi intermediate compartment (ERGIC) 1 (Ergic1) mRNA.
<a href="#">Ergic3</a>	<a href="#">Ergic3.bSep08</a>	<a href="#">296306</a>	8532	958	10	217	ERGIC and golgi 3 (24.7 kD) (Ergic3) alternative variant bSep08, mRNA.
<a href="#">Ergic3</a>	<a href="#">Ergic3.cSep08</a>	<a href="#">296306</a>	7609	550	7	183	ERGIC and golgi 3 (Ergic3) alternative variant cSep08, mRNA.
<a href="#">Ergic3</a>	<a href="#">Ergic3.dSep08</a>	<a href="#">296306</a>	1347	543	4	134	ERGIC and golgi 3 (Ergic3) alternative variant dSep08, mRNA.
<a href="#">Ergic3</a>	<a href="#">Ergic3.eSep08</a>	<a href="#">296306</a>	1010	374	2	36	ERGIC and golgi 3 (3.9 kD) (Ergic3) alternative variant eSep08, mRNA.
<a href="#">Erh</a>	<a href="#">Erh.bSep08</a>	<a href="#">681415</a>	10921	1468	2	104	enhancer of rudimentary homolog (Drosophila) (12.3 kD) (Erh) alternative variant bSep08, mRNA.
<a href="#">Erh</a>	<a href="#">Erh.cSep08</a>	<a href="#">681415</a>	5806	785	1	70	enhancer of rudimentary homolog (Drosophila) (8.4 kD) (Erh) alternative variant cSep08, mRNA.
<a href="#">Erich1</a>	<a href="#">Erich1.bSep08</a>	<a href="#">306622</a>	62012	402		133	glutamate-rich 1 (Erich1) alternative variant bSep08, mRNA.
<a href="#">Erlin1</a>	<a href="#">Erlin1.bSep08</a>	<a href="#">293939</a>	14375	432	5	143	ER lipid raft associated 1 (Erlin1) alternative variant bSep08, mRNA.
<a href="#">Erlin2</a>	<a href="#">Erlin2.aSep08</a>	<a href="#">290823</a>	12873	1785	9	530	ER lipid raft associated 2 (Erlin2) alternative variant aSep08, mRNA.
<a href="#">Ermap</a>	<a href="#">Ermap.aSep08</a>	<a href="#">298485</a>	2017	215		71	erythroblast membrane-associated protein (Ermap) mRNA.



<a href="#">Ernm</a>	<a href="#">Ernm.bSep08</a>	<a href="#">295619</a>	759	342	2	98	ermin, ERM-like protein (Ernm) alternative variant bSep08, mRNA.
<a href="#">Ern1</a>	<a href="#">Ern1.aSep08</a>	<a href="#">498013</a>	5166	505		148	endoplasmic reticulum (ER) to nucleus signalling 1 (Ern1) mRNA.
<a href="#">Ero1l</a>	<a href="#">Ero1l.aSep08</a>	<a href="#">171562</a>	35608	1834		476	ERO1-like (S. cerevisiae) (Ero1l) mRNA.
<a href="#">Ero1lb</a>	<a href="#">Ero1lb.aSep08</a>	<a href="#">364755</a>	34530	991		329	ERO1-like beta (S. cerevisiae) (Ero1lb) alternative variant aSep08, mRNA.
<a href="#">Ero1lb</a>	<a href="#">Ero1lb.bSep08</a>	<a href="#">364755</a>	11125	1069		179	ERO1-like beta (S. cerevisiae) (Ero1lb) alternative variant bSep08, mRNA.
<a href="#">Ero1lb</a>	<a href="#">Ero1lb.cSep08</a>	<a href="#">364755</a>	3216	887		65	ERO1-like beta (S. cerevisiae) (Ero1lb) alternative variant cSep08, mRNA.
<a href="#">Erp29</a>	<a href="#">Erp29.bSep08</a>	<a href="#">117030</a>	2824	2436	2	238	endoplasmic reticulum protein 29 (26.7 kD) (Erp29) alternative variant bSep08, mRNA.
<a href="#">Erp29</a>	<a href="#">Erp29.cSep08</a>	<a href="#">117030</a>	6293	978	2	209	endoplasmic reticulum protein 29 precursor (22.9 kD) (Erp29) alternative variant cSep08, mRNA.
<a href="#">Erp29</a>	<a href="#">Erp29.eSep08</a>	<a href="#">117030</a>	434	235	2	78	putative protein (Erp29) alternative variant eSep08, mRNA.
<a href="#">Erp29</a>	<a href="#">Erp29.fSep08</a>	<a href="#">117030</a>	4866	715	3	78	putative secreted or extracellular protein precursor (Erp29) alternative variant fSep08, mRNA.
<a href="#">Es1</a>	<a href="#">Es1.bSep08</a>	<a href="#">24346</a>	8350	545		126	esterase 1 (Es1) alternative variant bSep08, mRNA.
<a href="#">Es22</a>	<a href="#">Es22.bSep08</a>	<a href="#">29225</a>	14073	822	1	118	esterase 22 (13.4 kD) (Es22) alternative variant bSep08, mRNA.
<a href="#">Esam</a>	<a href="#">Esam.bSep08</a>	<a href="#">300519</a>	1622	1100	1	192	endothelial cell adhesion molecule (Esam) alternative variant bSep08, mRNA.
<a href="#">Esco1</a>	<a href="#">Esco1.aSep08</a>	<a href="#">680014</a>	39163	3884	9	840	establishment of cohesion 1 homolog 1 (S. cerevisiae) (94.6 kD) (Esco1) alternative variant aSep08, mRNA.
<a href="#">Esd</a>	<a href="#">Esd.bSep08</a>	<a href="#">290401</a>	14275	601	5	152	esterase D/formylglutathione hydrolase (Esd) alternative variant bSep08, mRNA.
<a href="#">Esd</a>	<a href="#">Esd.dSep08</a>	<a href="#">290401</a>	4511	550	2	55	esterase D/formylglutathione hydrolase (5.9 kD) (Esd) alternative variant dSep08, mRNA.
<a href="#">Esf1</a>	<a href="#">Esf1.aSep08</a>	<a href="#">366203</a>	70302	3336	14	861	ESF1, nucleolar pre-rRNA processing protein, homolog (S. cerevisiae) (Esf1) alternative variant aSep08, mRNA.
<a href="#">Esf1</a>	<a href="#">Esf1.bSep08</a>	<a href="#">366203</a>	2660	829	2	207	ESF1, nucleolar pre-rRNA processing protein, homolog (S. cerevisiae) (Esf1) alternative variant bSep08, mRNA.
<a href="#">Esp1l</a>	<a href="#">Esp1l.aSep08</a>	<a href="#">315330</a>	7668	1785	6	543	extra spindle poles like 1 (S. cerevisiae) (Esp1l) alternative variant aSep08, mRNA.
<a href="#">Esp1l</a>	<a href="#">Esp1l.bSep08</a>	<a href="#">315330</a>	4696	1938	10	443	extra spindle poles like 1 (S. cerevisiae) (Esp1l) alternative variant bSep08, mRNA.
<a href="#">Esp1l</a>	<a href="#">Esp1l.cSep08</a>	<a href="#">315330</a>	6225	585	3	146	extra spindle poles like 1 (S. cerevisiae) (Esp1l) alternative variant cSep08, mRNA.
<a href="#">Espn</a>	<a href="#">Espn.aSep08</a>	<a href="#">56227</a>	5731	423		140	espin (Espn) mRNA.
<a href="#">Esr1</a>	<a href="#">Esr1.bSep08</a>	<a href="#">24890</a>	730	424	1	70	estrogen receptor 1 (alpha) (Esr1) alternative variant bSep08, mRNA.
<a href="#">Esrra</a>	<a href="#">Esrra.bSep08</a>	<a href="#">293701</a>	8345	699	1	117	estrogen related receptor, alpha (12.4 kD) (Esrra) alternative variant bSep08, mRNA.
<a href="#">EST1.0</a>	<a href="#">EST1.0.aSep08</a>		39793	572		161	smg-7 homolog (EST1.0) mRNA.

<a href="#">Etaa1</a>	<a href="#">Etaa1.bSep08</a>	<a href="#">498420</a>	7583	2533	5	669	ewing's tumor-associated antigen 1 (Etaa1) alternative variant bSep08, mRNA.
<a href="#">Etaa1</a>	<a href="#">Etaa1.dSep08</a>	<a href="#">498420</a>	5666	561	5	93	ewing's tumor-associated antigen 1 (10.1 kD) (Etaa1) alternative variant dSep08, mRNA.
<a href="#">Etf1</a>	<a href="#">Etf1.bSep08</a>	<a href="#">307503</a>	21469	754	5	251	eukaryotic translation termination factor 1 (Etf1) alternative variant bSep08, mRNA.
<a href="#">Etfb</a>	<a href="#">Etfb.bSep08</a>	<a href="#">292845</a>	19941	899	3	221	electron transferring flavoprotein, beta polypeptide (24.1 kD) (Etfb) alternative variant bSep08, mRNA.
<a href="#">Etfb</a>	<a href="#">Etfb.cSep08</a>	<a href="#">292845</a>	12995	614	1	198	electron transferring flavoprotein, beta polypeptide (Etfb) alternative variant cSep08, mRNA.
<a href="#">Etnk2</a>	<a href="#">Etnk2.aSep08</a>	<a href="#">360843</a>	17730	1754	3	506	ethanolamine kinase 2 (Etnk2) alternative variant aSep08, mRNA.
<a href="#">Etnk2</a>	<a href="#">Etnk2.cSep08</a>	<a href="#">360843</a>	9071	500	2	166	ethanolamine kinase 2 (Etnk2) alternative variant cSep08, mRNA.
<a href="#">Ets.0</a>	<a href="#">Ets.0.aSep08</a>		3159	644		214	spi-B transcription factor (Ets.0) mRNA.
<a href="#">Ets2</a>	<a href="#">Ets2.bSep08</a>	<a href="#">304063</a>	1295	376	2	118	e26 avian leukemia oncogene 2, 3' domain (Ets2) alternative variant bSep08, mRNA.
<a href="#">Etv1</a>	<a href="#">Etv1.aSep08</a>	<a href="#">362733</a>	78264	1204	11	339	ets variant gene 1 (Etv1) alternative variant aSep08, mRNA.
<a href="#">Etv2</a>	<a href="#">Etv2.aSep08</a>	<a href="#">361544</a>	1139	459		146	ets variant gene 2 (Etv2) mRNA.
<a href="#">Etv3</a>	<a href="#">Etv3.bSep08</a>	<a href="#">295297</a>	4093	2342	3	147	ets variant gene 3 (17.5 kD) (Etv3) alternative variant bSep08, complete mRNA.
<a href="#">Etv3</a>	<a href="#">Etv3.cSep08</a>	<a href="#">295297</a>	1850	635	2	47	ets variant gene 3 (Etv3) alternative variant cSep08, mRNA.
<a href="#">Etv3l</a>	<a href="#">Etv3l.bSep08</a>	<a href="#">499651</a>	11824	1035	1	290	ets variant gene 3-like (Etv3l) alternative variant bSep08, mRNA.
<a href="#">Etv3l</a>	<a href="#">Etv3l.cSep08</a>	<a href="#">499651</a>	6449	703	1	108	ets variant gene 3-like (Etv3l) alternative variant cSep08, mRNA.
<a href="#">Etv4</a>	<a href="#">Etv4.bSep08</a>	<a href="#">360635</a>	15133	2202	13	487	ets gene 4 CRA b (54.1 kD) (Etv4) alternative variant bSep08, mRNA.
<a href="#">Etv4</a>	<a href="#">Etv4.cSep08</a>	<a href="#">360635</a>	11374	1390	6	267	ets gene 4 (29.3 kD) (Etv4) alternative variant cSep08, mRNA.
<a href="#">Etv4</a>	<a href="#">Etv4.dSep08</a>	<a href="#">360635</a>	1111	802	3	111	ets gene 4 CRA b (Etv4) alternative variant dSep08, mRNA.
<a href="#">Etv5</a>	<a href="#">Etv5.bSep08</a>	<a href="#">303828</a>	56309	1800	1	474	ets variant gene 5 (Etv5) alternative variant bSep08, mRNA.
<a href="#">Etv6</a>	<a href="#">Etv6.bSep08</a>	<a href="#">312777</a>	1132	720	2	115	putative protein, with a transmembrane domain (Etv6) alternative variant bSep08, mRNA.
<a href="#">Evc</a>	<a href="#">Evc.aSep08</a>	<a href="#">289712</a>	11618	1186		266	ellis van Creveld gene homolog (human) (Evc) mRNA.
<a href="#">Evc2</a>	<a href="#">Evc2.bSep08</a>	<a href="#">289711</a>	3259	418	1	63	ellis van Creveld syndrome 2 homolog (human) (Evc2) alternative variant bSep08, mRNA.
<a href="#">Evl</a>	<a href="#">Evl.aSep08</a>	<a href="#">79115</a>	50400	2922	2	420	ena-vasodilator stimulated phosphoprotein (45.1 kD) (Evl) alternative variant aSep08, mRNA.
<a href="#">Evl</a>	<a href="#">Evl.cSep08</a>	<a href="#">79115</a>	7000	1218	3	118	ena-vasodilator stimulated phosphoprotein (Evl) alternative variant cSep08, mRNA.
<a href="#">Ewsr1</a>	<a href="#">Ewsr1.bSep08</a>	<a href="#">289752</a>	11405	935	8	304	ewing sarcoma breakpoint region 1 (Ewsr1) alternative variant bSep08, mRNA.

<a href="#">Ewsr1</a>	<a href="#">Ewsr1.cSep08</a>	<a href="#">289752</a>	11314	718	6	239	ewing sarcoma breakpoint region 1 (Ewsr1) alternative variant cSep08, mRNA.
<a href="#">Ewsr1</a>	<a href="#">Ewsr1.dSep08</a>	<a href="#">289752</a>	7795	1712	5	189	ewing sarcoma breakpoint region 1 (Ewsr1) alternative variant dSep08, mRNA.
<a href="#">Ewsr1</a>	<a href="#">Ewsr1.eSep08</a>	<a href="#">289752</a>	6493	400	5	132	ewing sarcoma breakpoint region 1 (Ewsr1) alternative variant eSep08, mRNA.
<a href="#">Ewsr1</a>	<a href="#">Ewsr1.fSep08</a>	<a href="#">289752</a>	1637	724	3	65	ewing sarcoma breakpoint region 1 (7.3 kD) (Ewsr1) alternative variant fSep08, mRNA.
<a href="#">Exdl2</a>	<a href="#">Exdl2.bSep08</a>	<a href="#">362759</a>	20238	798	1	234	exonuclease 3"-5" domain-like 2 (Exdl2) alternative variant bSep08, mRNA.
<a href="#">Exo1</a>	<a href="#">Exo1.bSep08</a>	<a href="#">305000</a>	1279	612	1	21	exonuclease 1 (Exo1) alternative variant bSep08, mRNA.
<a href="#">Exoc3</a>	<a href="#">Exoc3.bSep08</a>	<a href="#">252881</a>	30746	1798	5	408	exocyst complex component 3 (Exoc3) alternative variant bSep08, complete mRNA.
<a href="#">Exoc3l</a>	<a href="#">Exoc3l.bSep08</a>	<a href="#">291961</a>	2307	1151	3	326	exocyst complex component 3-like (Exoc3l) alternative variant bSep08, mRNA.
<a href="#">Exoc3l</a>	<a href="#">Exoc3l.cSep08</a>	<a href="#">291961</a>	1218	858	5	191	exocyst complex component 3-like (Exoc3l) alternative variant cSep08, mRNA.
<a href="#">Exoc4</a>	<a href="#">Exoc4.aSep08</a>	<a href="#">116654</a>	260412	688	1	174	exocyst complex component 4 (Exoc4) alternative variant aSep08, mRNA.
<a href="#">Exoc4</a>	<a href="#">Exoc4.bSep08</a>	<a href="#">116654</a>	538972	753	1	168	exocyst complex component 4 (Exoc4) alternative variant bSep08, mRNA.
<a href="#">Exoc4</a>	<a href="#">Exoc4.cSep08</a>	<a href="#">116654</a>	163612	591	2	74	exocyst complex component 4 (Exoc4) alternative variant cSep08, mRNA.
<a href="#">Exoc5</a>	<a href="#">Exoc5.bSep08</a>	<a href="#">60627</a>	10139	769	7	110	exocyst complex component 5 (Exoc5) alternative variant bSep08, mRNA.
<a href="#">Exoc7</a>	<a href="#">Exoc7.bSep08</a>	<a href="#">64632</a>	8866	645	7	214	exocyst complex component 7 (Exoc7) alternative variant bSep08, mRNA.
<a href="#">Exoc7</a>	<a href="#">Exoc7.cSep08</a>	<a href="#">64632</a>	1488	478	3	146	exocyst complex component 7 (Exoc7) alternative variant cSep08, mRNA.
<a href="#">Exosc2</a>	<a href="#">Exosc2.bSep08</a>	<a href="#">366017</a>	3853	409	5	67	exosome component 2 (Exosc2) alternative variant bSep08, mRNA.
<a href="#">Exosc3</a>	<a href="#">Exosc3.aSep08</a>	<a href="#">313243</a>	5170	1038		274	exosome component 3 (Exosc3) mRNA.
<a href="#">Exosc5</a>	<a href="#">Exosc5.aSep08</a>	<a href="#">308441</a>	9151	1026	6	235	exosome component 5 (25.2 kD) (Exosc5) alternative variant aSep08, mRNA.
<a href="#">Exosc5</a>	<a href="#">Exosc5.bSep08</a>	<a href="#">308441</a>	5585	551	3	183	exosome component 5 (Exosc5) alternative variant bSep08, mRNA.
<a href="#">Exosc7</a>	<a href="#">Exosc7.aSep08</a>	<a href="#">316098</a>	25359	1035		298	exosome component 7 (Exosc7) mRNA.
<a href="#">Exosc8</a>	<a href="#">Exosc8.aSep08</a>	<a href="#">295050</a>	6520	947	11	281	exosome component 8 (Exosc8) alternative variant aSep08, mRNA.
<a href="#">Exosc9</a>	<a href="#">Exosc9.bSep08</a>	<a href="#">294975</a>	5279	1919	6	167	exosome component 9 (Exosc9) alternative variant bSep08, mRNA.
<a href="#">Exosc9</a>	<a href="#">Exosc9.cSep08</a>	<a href="#">294975</a>	1148	504	2	129	exosome component 9 (Exosc9) alternative variant cSep08, mRNA.
<a href="#">Exosc9</a>	<a href="#">Exosc9.dSep08</a>	<a href="#">294975</a>	3828	1752	3	64	exosome component 9 (7.0 kD) (Exosc9) alternative variant dSep08, mRNA.
<a href="#">Exo_endo_phos.11</a>	<a href="#">Exo_endo_phos.11.aSep08</a>		6664	745		247	synaptojanin 1 (Exo_endo_phos.11) mRNA.

<a href="#">Exo_endo_phos.21</a>	<a href="#">Exo_endo_phos.21.aSep08</a>		4078	2299	5	341	inositol e (Exo_endo_phos.21) alternative variant aSep08, mRNA.
<a href="#">Exo_endo_phos.21</a>	<a href="#">Exo_endo_phos.21.bSep08</a>		2410	1198	4	208	inositol e (Exo_endo_phos.21) alternative variant bSep08, mRNA.
<a href="#">Exo_endo_phos.21</a>	<a href="#">Exo_endo_phos.21.cSep08</a>		2259	1271	3	81	CRA c like (Exo_endo_phos.21) alternative variant cSep08, mRNA.
<a href="#">Ext1</a>	<a href="#">Ext1.aSep08</a>	<a href="#">299907</a>	277872	3366	11	749	exostoses (multiple) 1 (Ext1) alternative variant aSep08, mRNA.
<a href="#">Ext2</a>	<a href="#">Ext2.aSep08</a>	<a href="#">311215</a>	133015	2882	13	787	exostoses (multiple) 2 (Ext2) alternative variant aSep08, mRNA.
<a href="#">Ext2</a>	<a href="#">Ext2.cSep08</a>	<a href="#">311215</a>	66300	419	2	99	exostoses (multiple) 2 (Ext2) alternative variant cSep08, mRNA.
<a href="#">Extl2</a>	<a href="#">Extl2.cSep08</a>	<a href="#">310803</a>	14964	460	3	35	exostoses (multiple)-like 2 (4.0 kD) (Extl2) alternative variant cSep08, mRNA.
<a href="#">Extl3</a>	<a href="#">Extl3.bSep08</a>	<a href="#">56819</a>	22935	1604	1	457	exostoses (multiple)-like 3 (Extl3) alternative variant bSep08, mRNA.
<a href="#">Eya1</a>	<a href="#">Eya1.aSep08</a>	<a href="#">502935</a>	22884	394		42	eyes absent 1 homolog (Drosophila) (4.4 kD) (Eya1) mRNA.
<a href="#">Eya2</a>	<a href="#">Eya2.bSep08</a>	<a href="#">156826</a>	51275	359	4	119	eyes absent 2 homolog (Drosophila) (Eya2) alternative variant bSep08, mRNA.
<a href="#">Eya4</a>	<a href="#">Eya4.aSep08</a>	<a href="#">292172</a>	13383	783		228	eyes absent 4 homolog (Drosophila) (Eya4) mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.aSep08</a>	<a href="#">312299</a>	63760	3233	20	790	enhancer of zeste homolog 2 (Drosophila) (Ezh2) alternative variant aSep08, mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.bSep08</a>	<a href="#">312299</a>	46424	2198	18	404	enhancer of zeste homolog 2 (Drosophila) (45.9 kD) (Ezh2) alternative variant bSep08, mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.cSep08</a>	<a href="#">312299</a>	40439	431	5	143	enhancer of zeste homolog 2 (Drosophila) (Ezh2) alternative variant cSep08, mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.dSep08</a>	<a href="#">312299</a>	5411	463	3	117	enhancer of zeste homolog 2 (Drosophila) (Ezh2) alternative variant dSep08, mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.eSep08</a>	<a href="#">312299</a>	793	410	2	95	enhancer of zeste homolog 2 (Drosophila) (Ezh2) alternative variant eSep08, mRNA.
<a href="#">Ezh2</a>	<a href="#">Ezh2.fSep08</a>	<a href="#">312299</a>	5765	719	3	45	enhancer of zeste homolog 2 (Drosophila) (Ezh2) alternative variant fSep08, mRNA.
<a href="#">Ezr</a>	<a href="#">Ezr.bSep08</a>	<a href="#">54319</a>	1693	717	3	131	ezrin (Ezr) alternative variant bSep08, mRNA.
<a href="#">F-box.0</a>	<a href="#">F-box.0.aSep08</a>		10672	2243		380	F-box leucine-rich repeat protein 10 (F-box.0) mRNA.
<a href="#">F-box.1</a>	<a href="#">F-box.1.aSep08</a>		31380	1038	3	227	F-box wd-40 domain protein 8 (F-box.1) alternative variant aSep08, mRNA.
<a href="#">F-box.1</a>	<a href="#">F-box.1.bSep08</a>		26157	553	1	184	F-box wd-40 domain protein 8 (F-box.1) alternative variant bSep08, mRNA.
<a href="#">F-box.2</a>	<a href="#">F-box.2.aSep08</a>		156498	1401		466	F-box leucine-rich repeat protein 13 (F-box.2) mRNA.
<a href="#">F2</a>	<a href="#">F2.bSep08</a>	<a href="#">29251</a>	2216	1200	5	216	coagulation factor II (F2) alternative variant bSep08, mRNA.
<a href="#">F2</a>	<a href="#">F2.cSep08</a>	<a href="#">29251</a>	328	249	2	82	coagulation factor II (F2) alternative variant cSep08, mRNA.
<a href="#">F5</a>	<a href="#">F5.bSep08</a>	<a href="#">304929</a>	8355	2877	3	959	coagulation factor V (F5) alternative variant bSep08, mRNA.

<a href="#">F5</a>	<a href="#">F5.cSep08</a>	<a href="#">304929</a>	22927	708	5	235	coagulation factor V (F5) alternative variant cSep08, mRNA.
<a href="#">F5</a>	<a href="#">F5.dSep08</a>	<a href="#">304929</a>	9785	764	5	233	coagulation factor V (F5) alternative variant dSep08, mRNA.
<a href="#">F5</a>	<a href="#">F5.eSep08</a>	<a href="#">304929</a>	3090	559	3		
<a href="#">F5_F8_type_C.0</a>	<a href="#">F5_F8_type_C.0.aSep08</a>		1477	642	4	214	contactin associated protein 1 CRA a (F5_F8_type_C.0) alternative variant aSep08, mRNA.
<a href="#">F5_F8_type_C.1</a>	<a href="#">F5_F8_type_C.1.aSep08</a>		1994	286	1	95	EGF-like repeats discoidin I-like domains 3 (F5_F8_type_C.1) alternative variant aSep08, mRNA.
<a href="#">F5_F8_type_C.1</a>	<a href="#">F5_F8_type_C.1.bSep08</a>		560	253	1	46	EGF-like repeats discoidin I-like domains 3 (F5_F8_type_C.1) alternative variant bSep08, mRNA.
<a href="#">F5_F8_type_C.2</a>	<a href="#">F5_F8_type_C.2.aSep08</a>		178862	537		144	EGF-like repeats discoidin I-like domains 3 (F5_F8_type_C.2) mRNA.
<a href="#">F8</a>	<a href="#">F8.bSep08</a>	<a href="#">302470</a>	6799	3007	2	1002	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant bSep08, mRNA.
<a href="#">F8</a>	<a href="#">F8.bSep08</a>	<a href="#">679968</a>	6799	3007	2	1002	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant bSep08, mRNA.
<a href="#">F8</a>	<a href="#">F8.cSep08</a>	<a href="#">302470</a>	3748	1094	5	236	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant cSep08, mRNA.
<a href="#">F8</a>	<a href="#">F8.cSep08</a>	<a href="#">679968</a>	3748	1094	5	236	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant cSep08, mRNA.
<a href="#">F8</a>	<a href="#">F8.dSep08</a>	<a href="#">302470</a>	3989	325	3	75	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant dSep08, mRNA.
<a href="#">F8</a>	<a href="#">F8.dSep08</a>	<a href="#">679968</a>	3989	325	3	75	coagulation factor VIII and hypothetical protein LOC679968 (F8) alternative variant dSep08, mRNA.
<a href="#">F9</a>	<a href="#">F9.aSep08</a>	<a href="#">24946</a>	43629	1857		153	coagulation factor IX (F9) mRNA.
<a href="#">F11</a>	<a href="#">F11.bSep08</a>	<a href="#">290757</a>	8817	1209	5	402	coagulation factor XI (F11) alternative variant bSep08, mRNA.
<a href="#">F11r</a>	<a href="#">F11r.aSep08</a>	<a href="#">116479</a>	21526	971		267	f11 receptor (F11r) mRNA.
<a href="#">F13b</a>	<a href="#">F13b.aSep08</a>	<a href="#">289055</a>	21570	2352	11	669	coagulation factor XIII, beta subunit (F13b) alternative variant aSep08, mRNA.
<a href="#">F13b</a>	<a href="#">F13b.cSep08</a>	<a href="#">289055</a>	5542	689	4	229	coagulation factor XIII, beta subunit (F13b) alternative variant cSep08, mRNA.
<a href="#">F13b</a>	<a href="#">F13b.dSep08</a>	<a href="#">289055</a>	5314	817	3	119	coagulation factor XIII, beta subunit (F13b) alternative variant dSep08, mRNA.
<a href="#">Faah</a>	<a href="#">Faah.bSep08</a>	<a href="#">29347</a>	1781	1060	3	227	fatty acid amide hydrolase (25.2 kD) (Faah) alternative variant bSep08, mRNA.
<a href="#">Fabp3</a>	<a href="#">Fabp3.aSep08</a>	<a href="#">79131</a>	6300	1355	3	136	fatty acid binding protein 3, muscle and heart (Fabp3) alternative variant aSep08, mRNA.
<a href="#">Fabp4</a>	<a href="#">Fabp4.aSep08</a>	<a href="#">79451</a>	3556	610	3	142	fatty acid binding protein 4, adipocyte (15.7 kD) (Fabp4) alternative variant aSep08, mRNA.
<a href="#">Fabp4</a>	<a href="#">Fabp4.cSep08</a>	<a href="#">79451</a>	4672	818	4	97	fatty acid binding protein 4, adipocyte (10.9 kD) (Fabp4) alternative variant cSep08, complete mRNA.
<a href="#">Fabp7</a>	<a href="#">Fabp7.aSep08</a>	<a href="#">80841</a>	2242	1152	4	224	fatty acid binding protein 7, brain (Fabp7) alternative variant aSep08, complete mRNA.
<a href="#">Fabp7</a>	<a href="#">Fabp7.cSep08</a>	<a href="#">80841</a>	1549	458	3	130	fatty acid binding protein 7, brain (Fabp7) alternative variant cSep08, mRNA.

<a href="#">faby</a>	<a href="#">faby.aSep08</a>		46795	1605	3	86	putative secreted or extracellular protein precursor (10.1 kD) (faby) alternative variant aSep08, mRNA.
<a href="#">faby</a>	<a href="#">faby.bSep08</a>		5384	748	3	43	putative protein (faby) alternative variant bSep08, mRNA.
<a href="#">faby</a>	<a href="#">faby.cSep08</a>		5338	717	3	52	putative protein (faby) alternative variant cSep08, mRNA.
<a href="#">faby</a>	<a href="#">faby.dSep08</a>		5395	554	4	77	putative protein (faby) alternative variant dSep08, mRNA.
<a href="#">fachy</a>	<a href="#">fachy.aSep08</a>		1149	229		76	nebulin (fachy) mRNA.
<a href="#">Fads1</a>	<a href="#">Fads1.bSep08</a>	<a href="#">84575</a>	1312	463	1	116	fatty acid desaturase 1 (Fads1) alternative variant bSep08, mRNA.
<a href="#">Fads3</a>	<a href="#">Fads3.bSep08</a>	<a href="#">286922</a>	18750	3091	12	219	fatty acid desaturase 3 (25.6 kD) (Fads3) alternative variant bSep08, mRNA.
<a href="#">Faf1</a>	<a href="#">Faf1.bSep08</a>	<a href="#">140657</a>	117612	671	6	169	fas-associated factor 1 (Faf1) alternative variant bSep08, mRNA.
<a href="#">Faf1</a>	<a href="#">Faf1.cSep08</a>	<a href="#">140657</a>	66289	418	3	56	fas-associated factor 1 (Faf1) alternative variant cSep08, mRNA.
<a href="#">fafer</a>	<a href="#">fafer.aSep08</a>		51685	325		28	putative protein (3.0 kD) (fafer) mRNA.
<a href="#">faflo</a>	<a href="#">faflo.aSep08</a>		1427	1274		58	putative protein (6.6 kD) (faflo) alternative variant aSep08, mRNA.
<a href="#">faflu</a>	<a href="#">faflu.aSep08</a>		3144	2629		129	CRA a (faflu) mRNA.
<a href="#">Fahd2a</a>	<a href="#">Fahd2a.bSep08</a>	<a href="#">296131</a>	7926	959	6	216	putative protein of ancient origin (Fahd2a) alternative variant bSep08, mRNA.
<a href="#">Fahd2a</a>	<a href="#">Fahd2a.cSep08</a>	<a href="#">296131</a>	827	726	2	93	putative protein of ancient origin (10.3 kD) (Fahd2a) alternative variant cSep08, mRNA.
<a href="#">Fahd2a</a>	<a href="#">Fahd2a.dSep08</a>	<a href="#">296131</a>	7553	417	3	85	putative protein of metazoan origin (Fahd2a) alternative variant dSep08, mRNA.
<a href="#">Faim</a>	<a href="#">Faim.aSep08</a>	<a href="#">140930</a>	16013	1004	6	201	fas apoptotic inhibitory molecule (22.7 kD) (Faim) alternative variant aSep08, complete mRNA.
<a href="#">Faim.1</a>	<a href="#">Faim.1.aSep08</a>	<a href="#">140930</a>	16738	1004	6	201	fas apoptotic inhibitory molecule (22.7 kD) (Faim.1) alternative variant aSep08, complete mRNA.
<a href="#">Faim2</a>	<a href="#">Faim2.bSep08</a>	<a href="#">246274</a>	10653	796	9	167	fas apoptotic inhibitory molecule 2 (Faim2) alternative variant bSep08, mRNA.
<a href="#">Faim2</a>	<a href="#">Faim2.cSep08</a>	<a href="#">246274</a>	6783	440	5	120	fas apoptotic inhibitory molecule 2 (Faim2) alternative variant cSep08, mRNA.
<a href="#">Faim2</a>	<a href="#">Faim2.dSep08</a>	<a href="#">246274</a>	1276	410	2	59	fas apoptotic inhibitory molecule 2 (Faim2) alternative variant dSep08, mRNA.
<a href="#">Faim3</a>	<a href="#">Faim3.bSep08</a>	<a href="#">548326</a>	5569	1558	2	161	fas apoptotic inhibitory molecule 3 (18.3 kD) (Faim3) alternative variant bSep08, mRNA.
<a href="#">fakee</a>	<a href="#">fakee.aSep08</a>		25942	895		297	CLIP-associating protein 1 (fakee) mRNA.
<a href="#">faloy</a>	<a href="#">faloy.aSep08</a>		1451	933		64	putative protein of eukaryotic origin (7.1 kD) (faloy) mRNA.
<a href="#">Fam3a</a>	<a href="#">Fam3a.bSep08</a>	<a href="#">501664</a>	2314	1480	4	143	family with sequence similarity 3, member A (15.8 kD) (Fam3a) alternative variant bSep08, mRNA.
<a href="#">Fam3a</a>	<a href="#">Fam3a.cSep08</a>	<a href="#">501664</a>	2159	595	2	118	family with sequence similarity 3, member A (Fam3a) alternative variant cSep08, mRNA.
<a href="#">Fam3a</a>	<a href="#">Fam3a.dSep08</a>	<a href="#">501664</a>	2021	1230	4	103	family with sequence similarity 3, member A (Fam3a) alternative variant dSep08, mRNA.
<a href="#">Fam8a1</a>	<a href="#">Fam8a1.aSep08</a>	<a href="#">291031</a>	7110	3025		162	family with sequence similarity 8, member A1 (Fam8a1) mRNA.

<a href="#">Fam13c1</a>	<a href="#">Fam13c1.aSep08</a>	<a href="#">294565</a>	88269	821	6	190	CRA a (Fam13c1) alternative variant aSep08, mRNA.
<a href="#">Fam13c1</a>	<a href="#">Fam13c1.bSep08</a>	<a href="#">294565</a>	24433	701	3	116	CRA a (Fam13c1) alternative variant bSep08, mRNA.
<a href="#">Fam13c1</a>	<a href="#">Fam13c1.cSep08</a>	<a href="#">294565</a>	7729	348	2	67	CRA b (Fam13c1) alternative variant cSep08, mRNA.
<a href="#">Fam19a2</a>	<a href="#">Fam19a2.aSep08</a>	<a href="#">680647</a>	47301	2259		44	family with sequence similarity 19 (chemokine (C-C motif)-like), member A2 (Fam19a2) mRNA.
<a href="#">Fam19a5</a>	<a href="#">Fam19a5.aSep08</a>	<a href="#">500915</a>	139624	1134	2	129	family with sequence similarity 19 (chemokine (C-C motif)-like), member A5 (Fam19a5) alternative variant aSep08, mRNA.
<a href="#">Fam19a5</a>	<a href="#">Fam19a5.bSep08</a>	<a href="#">500915</a>	43032	1668	1	83	family with sequence similarity 19 (chemokine (C-C motif)-like), member A5 (Fam19a5) alternative variant bSep08, mRNA.
<a href="#">Fam20a</a>	<a href="#">Fam20a.bSep08</a>	<a href="#">303635</a>	932	328	1	66	family with sequence similarity 20, member A (Fam20a) alternative variant bSep08, mRNA.
<a href="#">Fam20b</a>	<a href="#">Fam20b.bSep08</a>	<a href="#">304885</a>	1978	749	2	117	family with sequence similarity 20, member B (Fam20b) alternative variant bSep08, mRNA.
<a href="#">Fam20b</a>	<a href="#">Fam20b.cSep08</a>	<a href="#">304885</a>	2149	1965	2	67	family with sequence similarity 20, member B (7.8 kD) (Fam20b) alternative variant cSep08, mRNA.
<a href="#">Fam20c</a>	<a href="#">Fam20c.bSep08</a>	<a href="#">304334</a>	48252	776	4	199	family with sequence similarity 20, member C (Fam20c) alternative variant bSep08, mRNA.
<a href="#">Fam20c</a>	<a href="#">Fam20c.cSep08</a>	<a href="#">304334</a>	15526	694	4	122	family with sequence similarity 20, member C (Fam20c) alternative variant cSep08, mRNA.
<a href="#">Fam21c</a>	<a href="#">Fam21c.bSep08</a>	<a href="#">297530</a>	37160	3293	22	1035	DNA segment Chr 6 Wayne State University 116 expressed (Fam21c) alternative variant bSep08, mRNA.
<a href="#">Fam21c</a>	<a href="#">Fam21c.cSep08</a>	<a href="#">297530</a>	17403	614	5	204	DNA segment Chr 6 Wayne State University 116 expressed (Fam21c) alternative variant cSep08, mRNA.
<a href="#">Fam21c</a>	<a href="#">Fam21c.dSep08</a>	<a href="#">297530</a>	3595	1092	3	200	DNA segment Chr 6 Wayne State University 116 expressed (Fam21c) alternative variant dSep08, mRNA.
<a href="#">Fam21c</a>	<a href="#">Fam21c.eSep08</a>	<a href="#">297530</a>	1843	1204	3	118	DNA segment Chr 6 Wayne State University 116 expressed (Fam21c) alternative variant eSep08, mRNA.
<a href="#">Fam24a</a>	<a href="#">Fam24a.aSep08</a>	<a href="#">499278</a>	2265	1455	2	98	family with sequence similarity 24, member A (10.7 kD) (Fam24a) alternative variant aSep08, mRNA.
<a href="#">Fam24a</a>	<a href="#">Fam24a.cSep08</a>	<a href="#">499278</a>	4240	1850	3	98	family with sequence similarity 24, member A (10.7 kD) (Fam24a) alternative variant cSep08, mRNA.
<a href="#">Fam26f</a>	<a href="#">Fam26f.bSep08</a>	<a href="#">294430</a>	1658	834	1	242	family with sequence similarity 26, member F (27.0 kD) (Fam26f) alternative variant bSep08, complete mRNA.
<a href="#">Fam26f</a>	<a href="#">Fam26f.cSep08</a>	<a href="#">294430</a>	17728	681	1	184	family with sequence similarity 26, member F (21.4 kD) (Fam26f) alternative variant cSep08, mRNA.
<a href="#">Fam26f</a>	<a href="#">Fam26f.dSep08</a>	<a href="#">294430</a>	5255	585	1	170	family with sequence similarity 26, member F (Fam26f) alternative variant dSep08, mRNA.
<a href="#">Fam29a</a>	<a href="#">Fam29a.aSep08</a>	<a href="#">366403</a>	4361	1771	1	410	family with sequence similarity 29, member A (Fam29a) alternative variant aSep08, mRNA.
<a href="#">Fam29a</a>	<a href="#">Fam29a.bSep08</a>	<a href="#">366403</a>	2866	591	1	58	family with sequence similarity 29, member A (6.4 kD) (Fam29a) alternative variant bSep08, mRNA.
<a href="#">Fam32a</a>	<a href="#">Fam32a.aSep08</a>	<a href="#">498600</a>	2838	678	3	176	family with sequence similarity 32, member A (Fam32a) alternative variant aSep08, mRNA.
<a href="#">Fam35a</a>	<a href="#">Fam35a.aSep08</a>	<a href="#">364514</a>	89131	1006	6	243	family with sequence similarity 35, member A (Fam35a) alternative variant aSep08, mRNA.

<a href="#">Fam35a</a>	<a href="#">Fam35a.bSep08</a>	<a href="#">364514</a>	23992	355	4	118	family with sequence similarity 35, member A (Fam35a) alternative variant bSep08, mRNA.
<a href="#">Fam35a</a>	<a href="#">Fam35a.cSep08</a>	<a href="#">364514</a>	60764	1157	4	130	family with sequence similarity 35, member A (Fam35a) alternative variant cSep08, mRNA.
<a href="#">Fam36a</a>	<a href="#">Fam36a.bSep08</a>	<a href="#">289278</a>	5147	1825	2	86	family with sequence similarity 36, member A (10.0 kD) (Fam36a) alternative variant bSep08, mRNA.
<a href="#">Fam36a</a>	<a href="#">Fam36a.cSep08</a>	<a href="#">289278</a>	1562	1009	3	84	family with sequence similarity 36, member A (9.5 kD) (Fam36a) alternative variant cSep08, mRNA.
<a href="#">Fam36a</a>	<a href="#">Fam36a.dSep08</a>	<a href="#">289278</a>	3567	488	4	82	family with sequence similarity 36, member A (Fam36a) alternative variant dSep08, mRNA.
<a href="#">Fam36a</a>	<a href="#">Fam36a.eSep08</a>	<a href="#">289278</a>	3580	373	3	73	family with sequence similarity 36, member A (Fam36a) alternative variant eSep08, mRNA.
<a href="#">Fam38b</a>	<a href="#">Fam38b.aSep08</a>	<a href="#">688934</a>	8299	1986		230	family with sequence similarity 38, member B (Fam38b) mRNA.
<a href="#">Fam40a</a>	<a href="#">Fam40a.aSep08</a>	<a href="#">362012</a>	20395	3200		839	family with sequence similarity 40, member A (Fam40a) mRNA.
<a href="#">Fam43a</a>	<a href="#">Fam43a.aSep08</a>	<a href="#">288031</a>	3132	2539	2	430	family with sequence similarity 43, member A (47.0 kD) (Fam43a) alternative variant aSep08, complete mRNA.
<a href="#">Fam45a</a>	<a href="#">Fam45a.bSep08</a>	<a href="#">308009</a>	13974	621	1	200	family with sequence similarity 45, member A (Fam45a) alternative variant bSep08, mRNA.
<a href="#">Fam46a</a>	<a href="#">Fam46a.bSep08</a>	<a href="#">300870</a>	2655	1385	3	427	family with sequence similarity 46, member A (48.5 kD) (Fam46a) alternative variant bSep08, complete mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.bSep08</a>	<a href="#">361946</a>	13750	1519	9	407	family with sequence similarity 48, member A (Fam48a) alternative variant bSep08, mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.cSep08</a>	<a href="#">361946</a>	8608	941	11	313	family with sequence similarity 48, member A (Fam48a) alternative variant cSep08, mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.dSep08</a>	<a href="#">361946</a>	11064	1005	5	223	family with sequence similarity 48, member A (Fam48a) alternative variant dSep08, mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.eSep08</a>	<a href="#">361946</a>	13531	894	10	221	family with sequence similarity 48, member A (Fam48a) alternative variant eSep08, mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.fSep08</a>	<a href="#">361946</a>	12517	820	10	190	family with sequence similarity 48, member A (Fam48a) alternative variant fSep08, mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.gSep08</a>	<a href="#">361946</a>	12474	673	9	99	family with sequence similarity 48, member A (11.4 kD) (Fam48a) alternative variant gSep08, complete mRNA.
<a href="#">Fam48a</a>	<a href="#">Fam48a.hSep08</a>	<a href="#">361946</a>	13387	755	9	98	family with sequence similarity 48, member A (11.3 kD) (Fam48a) alternative variant hSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.bSep08</a>	<a href="#">299909</a>	116430	700	7	232	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant bSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.bSep08</a>	<a href="#">690541</a>	116430	700	7	232	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant bSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.cSep08</a>	<a href="#">299909</a>	9551	764	5	179	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant cSep08, mRNA.



<a href="#">Fam49b</a>	<a href="#">Fam49b.cSep08</a>	<a href="#">690541</a>	9551	764	5	179	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant cSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.dSep08</a>	<a href="#">299909</a>	51165	725	7	172	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant dSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.dSep08</a>	<a href="#">690541</a>	51165	725	7	172	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant dSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.eSep08</a>	<a href="#">299909</a>	114310	738	7	155	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant eSep08, mRNA.
<a href="#">Fam49b</a>	<a href="#">Fam49b.eSep08</a>	<a href="#">690541</a>	114310	738	7	155	family with sequence similarity 49, member B and hypothetical protein LOC690541 (Fam49b) alternative variant eSep08, mRNA.
<a href="#">FAM50A</a>	<a href="#">FAM50A.bSep08</a>	<a href="#">293862</a>	5499	1134	11	241	family with sequence similarity 50, member A (FAM50A) alternative variant bSep08, mRNA.
<a href="#">Fam53a</a>	<a href="#">Fam53a.bSep08</a>	<a href="#">305461</a>	24268	726	3	151	family with sequence similarity 53, member A (16.2 kD) (Fam53a) alternative variant bSep08, mRNA.
<a href="#">Fam53a</a>	<a href="#">Fam53a.cSep08</a>	<a href="#">305461</a>	5295	415	2	61	family with sequence similarity 53, member A (6.8 kD) (Fam53a) alternative variant cSep08, mRNA.
<a href="#">Fam53a</a>	<a href="#">Fam53a.dSep08</a>	<a href="#">305461</a>	34267	1777	4	31	family with sequence similarity 53, member A (Fam53a) alternative variant dSep08, mRNA.
<a href="#">Fam53a</a>	<a href="#">Fam53a.fSep08</a>	<a href="#">305461</a>	17151	803	4	83	family with sequence similarity 53, member A (8.9 kD) (Fam53a) alternative variant fSep08, mRNA.
<a href="#">Fam54b</a>	<a href="#">Fam54b.aSep08</a>	<a href="#">298549</a>	9630	1799	4	468	family with sequence similarity 54, member B (Fam54b) alternative variant aSep08, complete mRNA.
<a href="#">Fam54b</a>	<a href="#">Fam54b.cSep08</a>	<a href="#">298549</a>	6309	836	5	224	family with sequence similarity 54, member B (Fam54b) alternative variant cSep08, mRNA.
<a href="#">Fam54b</a>	<a href="#">Fam54b.dSep08</a>	<a href="#">298549</a>	5890	761	5	163	family with sequence similarity 54, member B (Fam54b) alternative variant dSep08, mRNA.
<a href="#">Fam54b</a>	<a href="#">Fam54b.eSep08</a>	<a href="#">298549</a>	4653	660	3	77	family with sequence similarity 54, member B (8.5 kD) (Fam54b) alternative variant eSep08, mRNA.
<a href="#">Fam55c</a>	<a href="#">Fam55c.bSep08</a>	<a href="#">681096</a>	30537	627	2	116	family with sequence similarity 55, member C (Fam55c) alternative variant bSep08, mRNA.
<a href="#">Fam62c</a>	<a href="#">Fam62c.aSep08</a>	<a href="#">363120</a>	4625	934		281	DNA segment chr 9 ERATO Doi 280 expressed (Fam62c) mRNA.
<a href="#">Fam63a</a>	<a href="#">Fam63a.bSep08</a>	<a href="#">310665</a>	1000	714	1	84	family with sequence similarity 63, member A (Fam63a) alternative variant bSep08, mRNA.
<a href="#">Fam64a</a>	<a href="#">Fam64a.bSep08</a>	<a href="#">360559</a>	3871	774	1	250	family with sequence similarity 64, member A (Fam64a) alternative variant bSep08, mRNA.
<a href="#">Fam65a</a>	<a href="#">Fam65a.aSep08</a>	<a href="#">291974</a>	7812	4135	21	1211	family with sequence similarity 65, member A (Fam65a) alternative variant aSep08, mRNA.
<a href="#">Fam65a</a>	<a href="#">Fam65a.bSep08</a>	<a href="#">291974</a>	1122	798	4	149	family with sequence similarity 65, member A (Fam65a) alternative variant bSep08, mRNA.
<a href="#">Fam65a</a>	<a href="#">Fam65a.cSep08</a>	<a href="#">291974</a>	437	325	2	80	family with sequence similarity 65, member A (Fam65a) alternative variant cSep08, mRNA.

<a href="#">Fam65a</a>	<a href="#">Fam65a.dSep08</a>	<a href="#">291974</a>	785	686	2	72	family with sequence similarity 65, member A (Fam65a) alternative variant dSep08, mRNA.
<a href="#">Fam69a</a>	<a href="#">Fam69a.aSep08</a>	<a href="#">360906</a>	71325	2754	5	426	family with sequence similarity 69, member A (Fam69a) alternative variant aSep08, mRNA.
<a href="#">Fam69a</a>	<a href="#">Fam69a.bSep08</a>	<a href="#">360906</a>	1757	502	2	167	family with sequence similarity 69, member A (Fam69a) alternative variant bSep08, mRNA.
<a href="#">Fam69b</a>	<a href="#">Fam69b.cSep08</a>	<a href="#">362090</a>	547	298	2	92	family with sequence similarity 69, member B (Fam69b) alternative variant cSep08, mRNA.
<a href="#">Fam70a</a>	<a href="#">Fam70a.bSep08</a>	<a href="#">313453</a>	6795	393	2	93	family with sequence similarity 70, member A (Fam70a) alternative variant bSep08, mRNA.
<a href="#">Fam70b</a>	<a href="#">Fam70b.bSep08</a>	<a href="#">290877</a>	11465	1157	1	66	family with sequence similarity 70, member B (Fam70b) alternative variant bSep08, mRNA.
<a href="#">Fam71d</a>	<a href="#">Fam71d.bSep08</a>	<a href="#">366671</a>	19192	757	1	154	family with sequence similarity 71, member D (Fam71d) alternative variant bSep08, mRNA.
<a href="#">Fam71f1</a>	<a href="#">Fam71f1.bSep08</a>	<a href="#">500061</a>	8151	741	2	134	family with sequence similarity 71, member F1 (Fam71f1) alternative variant bSep08, mRNA.
<a href="#">Fam71f1</a>	<a href="#">Fam71f1.cSep08</a>	<a href="#">500061</a>	8159	561	2	74	family with sequence similarity 71, member F1 (Fam71f1) alternative variant cSep08, mRNA.
<a href="#">Fam72a</a>	<a href="#">Fam72a.aSep08</a>	<a href="#">681249</a>	15898	632		77	family with sequence similarity 72, member A (8.7 kD) (Fam72a) mRNA.
<a href="#">Fam73b</a>	<a href="#">Fam73b.bSep08</a>	<a href="#">296623</a>	8187	1764	10	334	family with sequence similarity 73, member B (37.6 kD) (Fam73b) alternative variant bSep08, mRNA.
<a href="#">Fam73b</a>	<a href="#">Fam73b.cSep08</a>	<a href="#">296623</a>	22651	284	3	79	family with sequence similarity 73, member B (Fam73b) alternative variant cSep08, mRNA.
<a href="#">Fam76a</a>	<a href="#">Fam76a.bSep08</a>	<a href="#">362618</a>	20476	737	4	209	family with sequence similarity 76, member A (Fam76a) alternative variant bSep08, mRNA.
<a href="#">Fam76a</a>	<a href="#">Fam76a.cSep08</a>	<a href="#">362618</a>	27785	1794	3	119	family with sequence similarity 76, member A (13.7 kD) (Fam76a) alternative variant cSep08, mRNA.
<a href="#">Fam78a</a>	<a href="#">Fam78a.aSep08</a>	<a href="#">499776</a>	14489	1803		283	family with sequence similarity 78, member A (32.1 kD) (Fam78a) mRNA.
<a href="#">Fam80b</a>	<a href="#">Fam80b.aSep08</a>	<a href="#">362428</a>	47353	3387	6	387	family with sequence similarity 80, member B (42.5 kD) (Fam80b) alternative variant aSep08, mRNA.
<a href="#">Fam80b</a>	<a href="#">Fam80b.bSep08</a>	<a href="#">362428</a>	4387	411	3	87	family with sequence similarity 80, member B (Fam80b) alternative variant bSep08, mRNA.
<a href="#">Fam80b</a>	<a href="#">Fam80b.cSep08</a>	<a href="#">362428</a>	15384	624	2	30	family with sequence similarity 80, member B (Fam80b) alternative variant cSep08, mRNA.
<a href="#">Fam81a</a>	<a href="#">Fam81a.bSep08</a>	<a href="#">315789</a>	46053	798	6	204	family with sequence similarity 81, member A (Fam81a) alternative variant bSep08, mRNA.
<a href="#">Fam81a</a>	<a href="#">Fam81a.cSep08</a>	<a href="#">315789</a>	10316	752	4	105	family with sequence similarity 81, member A (Fam81a) alternative variant cSep08, mRNA.
<a href="#">Fam81b</a>	<a href="#">Fam81b.aSep08</a>	<a href="#">309925</a>	91390	762	4	227	family with sequence similarity 81, member B (Fam81b) alternative variant aSep08, mRNA.
<a href="#">Fam81b</a>	<a href="#">Fam81b.bSep08</a>	<a href="#">309925</a>	91477	730	3	118	family with sequence similarity 81, member B (Fam81b) alternative variant bSep08, mRNA.
<a href="#">Fam81b</a>	<a href="#">Fam81b.cSep08</a>	<a href="#">309925</a>	96488	814	3	98	family with sequence similarity 81, member B (11.6 kD) (Fam81b) alternative variant cSep08, mRNA.

<a href="#">Fam82a</a>	<a href="#">Fam82a.aSep08</a>	<a href="#">313840</a>	71786	2267	8	615	of microtubule dynamics protein 2 (Fam82a) alternative variant aSep08, mRNA.
<a href="#">Fam82a</a>	<a href="#">Fam82a.cSep08</a>	<a href="#">313840</a>	46314	721	3	240	of microtubule dynamics protein 2 (Fam82a) alternative variant cSep08, mRNA.
<a href="#">Fam82a</a>	<a href="#">Fam82a.dSep08</a>	<a href="#">313840</a>	22694	716	2	238	putative protein of mammalian origin (Fam82a) alternative variant dSep08, mRNA.
<a href="#">Fam82a</a>	<a href="#">Fam82a.eSep08</a>	<a href="#">313840</a>	4884	1763	2	206	of microtubule dynamics protein 2 (23.5 kD) (Fam82a) alternative variant eSep08, mRNA.
<a href="#">Fam82a</a>	<a href="#">Fam82a.fSep08</a>	<a href="#">313840</a>	17625	677	8	189	of microtubule dynamics protein 2 (Fam82a) alternative variant fSep08, mRNA.
<a href="#">Fam82a</a>	<a href="#">Fam82a.gSep08</a>	<a href="#">313840</a>	29657	781	4	107	of microtubule dynamics protein 2 (Fam82a) alternative variant gSep08, mRNA.
<a href="#">Fam82c</a>	<a href="#">Fam82c.bSep08</a>	<a href="#">311328</a>	15010	1090	5	239	family with sequence similarity 82, member C (Fam82c) alternative variant bSep08, mRNA.
<a href="#">Fam83e</a>	<a href="#">Fam83e.aSep08</a>	<a href="#">292913</a>	2639	1072		229	family with sequence similarity 83, member E (Fam83e) mRNA.
<a href="#">Fam84a</a>	<a href="#">Fam84a.bSep08</a>	<a href="#">313969</a>	4663	3651	1	94	family with sequence similarity 84, member A (10.6 kD) (Fam84a) alternative variant bSep08, mRNA.
<a href="#">Fam86a</a>	<a href="#">Fam86a.bSep08</a>	<a href="#">302931</a>	10287	1247	1	93	family with sequence similarity 86, member A (Fam86a) alternative variant bSep08, mRNA.
<a href="#">Fam92a1</a>	<a href="#">Fam92a1.aSep08</a>	<a href="#">297903</a>	18109	1801	7	587	family with sequence similarity 92, member A1 (Fam92a1) alternative variant aSep08, mRNA.
<a href="#">Fam92a1</a>	<a href="#">Fam92a1.bSep08</a>	<a href="#">297903</a>	11526	612	4	140	family with sequence similarity 92, member A1 (Fam92a1) alternative variant bSep08, mRNA.
<a href="#">Fam92a1</a>	<a href="#">Fam92a1.cSep08</a>	<a href="#">297903</a>	14900	978	6	103	family with sequence similarity 92, member A1 (Fam92a1) alternative variant cSep08, mRNA.
<a href="#">Fam92a1</a>	<a href="#">Fam92a1.dSep08</a>	<a href="#">297903</a>	2690	249	3	83	family with sequence similarity 92, member A1 (Fam92a1) alternative variant dSep08, mRNA.
<a href="#">Fam92a1</a>	<a href="#">Fam92a1.eSep08</a>	<a href="#">297903</a>	13257	847	5	44	family with sequence similarity 92, member A1 (Fam92a1) alternative variant eSep08, mRNA.
<a href="#">Fam92b</a>	<a href="#">Fam92b.aSep08</a>	<a href="#">361423</a>	5238	488		162	family with sequence similarity 92, member B (Fam92b) mRNA.
<a href="#">Fam96b</a>	<a href="#">Fam96b.aSep08</a>	<a href="#">680987</a>	1890	717	5	165	family with sequence similarity 96, member B (17.8 kD) (Fam96b) alternative variant aSep08, mRNA.
<a href="#">Fam96b</a>	<a href="#">Fam96b.bSep08</a>	<a href="#">680987</a>	1814	1039	4	135	family with sequence similarity 96, member B (Fam96b) alternative variant bSep08, mRNA.
<a href="#">Fam96b</a>	<a href="#">Fam96b.cSep08</a>	<a href="#">680987</a>	1810	454	3	100	family with sequence similarity 96, member B (Fam96b) alternative variant cSep08, mRNA.
<a href="#">Fam96b</a>	<a href="#">Fam96b.dSep08</a>	<a href="#">680987</a>	1615	599	4	57	family with sequence similarity 96, member B (Fam96b) alternative variant dSep08, mRNA.
<a href="#">Fam98b</a>	<a href="#">Fam98b.aSep08</a>	<a href="#">499866</a>	17065	438		137	family with sequence similarity 98, member B (Fam98b) mRNA.
<a href="#">Fam100b</a>	<a href="#">Fam100b.aSep08</a>	<a href="#">287840</a>	4593	1293		163	family with sequence similarity 100, member B (17.6 kD) (Fam100b) complete mRNA.
<a href="#">Fam103a1</a>	<a href="#">Fam103a1.bSep08</a>	<a href="#">293058</a>	5879	1445	1	109	family with sequence similarity 103, member A1 (13.4 kD) (Fam103a1) alternative variant bSep08, complete mRNA.

<a href="#">Fam107b</a>	<a href="#">Fam107b.aSep08</a>	<a href="#">498796</a>	61865	3059	3	193	family with sequence similarity 107, member B (Fam107b) alternative variant aSep08, mRNA.
<a href="#">Fam107b</a>	<a href="#">Fam107b.bSep08</a>	<a href="#">498796</a>	32676	751	3	131	family with sequence similarity 107, member B (15.6 kD) (Fam107b) alternative variant bSep08, mRNA.
<a href="#">Fam107b</a>	<a href="#">Fam107b.cSep08</a>	<a href="#">498796</a>	32413	651	3	131	family with sequence similarity 107, member B (15.6 kD) (Fam107b) alternative variant cSep08, mRNA.
<a href="#">Fam107b</a>	<a href="#">Fam107b.eSep08</a>	<a href="#">498796</a>	3770	491	2	99	family with sequence similarity 107, member B (11.7 kD) (Fam107b) alternative variant eSep08, complete mRNA.
<a href="#">Fam108a1</a>	<a href="#">Fam108a1.bSep08</a>	<a href="#">299617</a>	1588	723	3	148	family with sequence similarity 108, member A1 (Fam108a1) alternative variant bSep08, mRNA.
<a href="#">Fam108b1</a>	<a href="#">Fam108b1.aSep08</a>	<a href="#">309399</a>	34477	2379	4	288	family with sequence similarity 108, member B1 (32.2 kD) (Fam108b1) alternative variant aSep08, mRNA.
<a href="#">Fam108b1</a>	<a href="#">Fam108b1.cSep08</a>	<a href="#">309399</a>	4942	754	2	121	family with sequence similarity 108, member B1 (Fam108b1) alternative variant cSep08, mRNA.
<a href="#">Fam108b1</a>	<a href="#">Fam108b1.dSep08</a>	<a href="#">309399</a>	32895	571	4	94	family with sequence similarity 108, member B1 (Fam108b1) alternative variant dSep08, mRNA.
<a href="#">Fam108c1</a>	<a href="#">Fam108c1.bSep08</a>	<a href="#">361601</a>	57719	770	3	160	family with sequence similarity 108, member C1 (18.0 kD) (Fam108c1) alternative variant bSep08, mRNA.
<a href="#">Fam108c1</a>	<a href="#">Fam108c1.cSep08</a>	<a href="#">361601</a>	9015	360	2	119	family with sequence similarity 108, member C1 (Fam108c1) alternative variant cSep08, mRNA.
<a href="#">Fam108c1</a>	<a href="#">Fam108c1.dSep08</a>	<a href="#">361601</a>	39306	772	2	94	family with sequence similarity 108, member C1 (10.7 kD) (Fam108c1) alternative variant dSep08, mRNA.
<a href="#">Fam108c1</a>	<a href="#">Fam108c1.eSep08</a>	<a href="#">361601</a>	57637	602	2	69	family with sequence similarity 108, member C1 (Fam108c1) alternative variant eSep08, mRNA.
<a href="#">Fam110a</a>	<a href="#">Fam110a.aSep08</a>	<a href="#">311535</a>	10594	1728	2	386	family with sequence similarity 110, member A (Fam110a) alternative variant aSep08, mRNA.
<a href="#">Fam110a</a>	<a href="#">Fam110a.cSep08</a>	<a href="#">311535</a>	3415	425	2	141	family with sequence similarity 110, member A (Fam110a) alternative variant cSep08, mRNA.
<a href="#">Fam110a</a>	<a href="#">Fam110a.eSep08</a>	<a href="#">311535</a>	3449	752	2	136	family with sequence similarity 110, member A (Fam110a) alternative variant eSep08, mRNA.
<a href="#">Fam110a</a>	<a href="#">Fam110a.fSep08</a>	<a href="#">311535</a>	3031	805	3	98	family with sequence similarity 110, member A (Fam110a) alternative variant fSep08, mRNA.
<a href="#">Fam110b</a>	<a href="#">Fam110b.bSep08</a>	<a href="#">500400</a>	139849	502	3	37	family with sequence similarity 110, member B (Fam110b) alternative variant bSep08, mRNA.
<a href="#">Fam110b</a>	<a href="#">Fam110b.cSep08</a>	<a href="#">500400</a>	139737	477	4	37	family with sequence similarity 110, member B (Fam110b) alternative variant cSep08, mRNA.
<a href="#">Fam110b</a>	<a href="#">Fam110b.dSep08</a>	<a href="#">500400</a>	23712	386	2	50	family with sequence similarity 110, member B (Fam110b) alternative variant dSep08, mRNA.
<a href="#">Fam113a</a>	<a href="#">Fam113a.aSep08</a>	<a href="#">296158</a>	6059	1735	8	318	family with sequence similarity 113, member A (36.5 kD) (Fam113a) alternative variant aSep08, mRNA.
<a href="#">Fam113a</a>	<a href="#">Fam113a.cSep08</a>	<a href="#">296158</a>	5082	1502	4	123	family with sequence similarity 113, member A (Fam113a) alternative variant cSep08, mRNA.
<a href="#">Fam113b</a>	<a href="#">Fam113b.aSep08</a>	<a href="#">315283</a>	23542	1202	2	219	putative protein of metazoan origin (Fam113b) alternative variant aSep08, mRNA.
<a href="#">Fam113b</a>	<a href="#">Fam113b.bSep08</a>	<a href="#">315283</a>	7668	898	2	200	putative protein of metazoan origin (Fam113b) alternative variant bSep08, mRNA.

<a href="#">Fam113b</a>	<a href="#">Fam113b.dSep08</a>	<a href="#">315283</a>	7297	562	2	101	putative protein (Fam113b) alternative variant dSep08, mRNA.
<a href="#">Fam114a1</a>	<a href="#">Fam114a1.aSep08</a>	<a href="#">498366</a>	56076	935		311	family with sequence similarity 114, member A1 (Fam114a1) mRNA.
<a href="#">Fam116a</a>	<a href="#">Fam116a.bSep08</a>	<a href="#">306229</a>	2895	1574	2	115	family with sequence similarity 116, member A (Fam116a) alternative variant bSep08, mRNA.
<a href="#">Fam116a</a>	<a href="#">Fam116a.cSep08</a>	<a href="#">306229</a>	8557	380	2	84	family with sequence similarity 116, member A (Fam116a) alternative variant cSep08, mRNA.
<a href="#">Fam116a</a>	<a href="#">Fam116a.dSep08</a>	<a href="#">306229</a>	550	479	2	38	family with sequence similarity 116, member A (4.0 kD) (Fam116a) alternative variant dSep08, mRNA.
<a href="#">Fam116a</a>	<a href="#">Fam116a.eSep08</a>	<a href="#">306229</a>	1228	311	2	14	family with sequence similarity 116, member A (Fam116a) alternative variant eSep08, mRNA.
<a href="#">Fam116b</a>	<a href="#">Fam116b.bSep08</a>	<a href="#">362983</a>	1929	1065	5	170	family with sequence similarity 116, member B (19.9 kD) (Fam116b) alternative variant bSep08, mRNA.
<a href="#">Fam116b</a>	<a href="#">Fam116b.cSep08</a>	<a href="#">362983</a>	1179	408	4	73	family with sequence similarity 116, member B (Fam116b) alternative variant cSep08, mRNA.
<a href="#">Fam117a</a>	<a href="#">Fam117a.bSep08</a>	<a href="#">497983</a>	4348	1286	3	181	family with sequence similarity 117, member A (19.5 kD) (Fam117a) alternative variant bSep08, mRNA.
<a href="#">Fam117a</a>	<a href="#">Fam117a.cSep08</a>	<a href="#">497983</a>	4080	712	3	61	family with sequence similarity 117, member A (Fam117a) alternative variant cSep08, mRNA.
<a href="#">Fam117a</a>	<a href="#">Fam117a.dSep08</a>	<a href="#">497983</a>	10506	734	4	61	family with sequence similarity 117, member A (Fam117a) alternative variant dSep08, mRNA.
<a href="#">Fam118a</a>	<a href="#">Fam118a.aSep08</a>	<a href="#">300120</a>	26526	2444	8	390	family with sequence similarity 118, member A (Fam118a) alternative variant aSep08, mRNA.
<a href="#">Fam118a</a>	<a href="#">Fam118a.bSep08</a>	<a href="#">300120</a>	9950	753	2	81	family with sequence similarity 118, member A (Fam118a) alternative variant bSep08, mRNA.
<a href="#">Fam118a</a>	<a href="#">Fam118a.cSep08</a>	<a href="#">300120</a>	9132	734	2	80	family with sequence similarity 118, member A (Fam118a) alternative variant cSep08, mRNA.
<a href="#">Fam119a</a>	<a href="#">Fam119a.aSep08</a>	<a href="#">301466</a>	9269	1260	5	177	family with sequence similarity 119, member A (Fam119a) alternative variant aSep08, mRNA.
<a href="#">Fam120a</a>	<a href="#">Fam120a.aSep08</a>	<a href="#">291019</a>	91021	4953	9	908	family with sequence similarity 120A (Fam120a) alternative variant aSep08, mRNA.
<a href="#">Fam120b</a>	<a href="#">Fam120b.bSep08</a>	<a href="#">308218</a>	5265	1334	2	319	family with sequence similarity 120B (Fam120b) alternative variant bSep08, mRNA.
<a href="#">Fam120b</a>	<a href="#">Fam120b.cSep08</a>	<a href="#">308218</a>	8276	771	2	104	family with sequence similarity 120B (Fam120b) alternative variant cSep08, mRNA.
<a href="#">Fam120b</a>	<a href="#">Fam120b.eSep08</a>	<a href="#">308218</a>	8048	739	4	93	family with sequence similarity 120B (Fam120b) alternative variant eSep08, mRNA.
<a href="#">FAM120C</a>	<a href="#">FAM120C.aSep08</a>	<a href="#">317423</a>	27739	992		300	family with sequence similarity 120C (FAM120C) mRNA.
<a href="#">Fam122b</a>	<a href="#">Fam122b.aSep08</a>	<a href="#">501647</a>	9408	1784		59	family with sequence similarity 122B (Fam122b) alternative variant aSep08, mRNA.
<a href="#">Fam122c</a>	<a href="#">Fam122c.aSep08</a>	<a href="#">685701</a>	42301	365		45	family with sequence similarity 122C (Fam122c) mRNA.
<a href="#">Fam124a</a>	<a href="#">Fam124a.aSep08</a>	<a href="#">691938</a>	34831	1636		301	family with sequence similarity 124A (34.5 kD) (Fam124a) mRNA.
<a href="#">Fam125a</a>	<a href="#">Fam125a.bSep08</a>	<a href="#">290635</a>	2964	761	6	253	family with sequence similarity 125, member A (Fam125a) alternative variant bSep08, mRNA.

<a href="#">Fam125a</a>	<a href="#">Fam125a.cSep08</a>	<a href="#">290635</a>	2505	767	5	137	family with sequence similarity 125, member A (14.0 kD) (Fam125a) alternative variant cSep08, mRNA.
<a href="#">Fam125a</a>	<a href="#">Fam125a.dSep08</a>	<a href="#">290635</a>	2637	684	6	137	family with sequence similarity 125, member A (14.0 kD) (Fam125a) alternative variant dSep08, mRNA.
<a href="#">Fam125a</a>	<a href="#">Fam125a.eSep08</a>	<a href="#">290635</a>	933	724	3	117	family with sequence similarity 125, member A (12.1 kD) (Fam125a) alternative variant eSep08, mRNA.
<a href="#">Fam125b</a>	<a href="#">Fam125b.aSep08</a>	<a href="#">362118</a>	60535	622	6	206	family with sequence similarity 125, member B (Fam125b) alternative variant aSep08, mRNA.
<a href="#">Fam125b</a>	<a href="#">Fam125b.bSep08</a>	<a href="#">362118</a>	45442	437	4	145	family with sequence similarity 125, member B (Fam125b) alternative variant bSep08, mRNA.
<a href="#">Fam126a</a>	<a href="#">Fam126a.aSep08</a>	<a href="#">499975</a>	37976	857		285	family with sequence similarity 126, member A (Fam126a) mRNA.
<a href="#">Fam126b</a>	<a href="#">Fam126b.bSep08</a>	<a href="#">316415</a>	56943	747	8	192	family with sequence similarity 126, member B (Fam126b) alternative variant bSep08, mRNA.
<a href="#">Fam126b</a>	<a href="#">Fam126b.cSep08</a>	<a href="#">316415</a>	5640	401	3	104	family with sequence similarity 126, member B (Fam126b) alternative variant cSep08, mRNA.
<a href="#">Fam128b</a>	<a href="#">Fam128b.aSep08</a>	<a href="#">287929</a>	5817	680	4	179	family with sequence similarity 128, member B (Fam128b) alternative variant aSep08, mRNA.
<a href="#">Fam128b</a>	<a href="#">Fam128b.bSep08</a>	<a href="#">287929</a>	5790	774	4	158	family with sequence similarity 128, member B (16.4 kD) (Fam128b) alternative variant bSep08, mRNA.
<a href="#">Fam128b</a>	<a href="#">Fam128b.dSep08</a>	<a href="#">287929</a>	7217	2463	3	133	family with sequence similarity 128, member B (Fam128b) alternative variant dSep08, mRNA.
<a href="#">Fam128b</a>	<a href="#">Fam128b.eSep08</a>	<a href="#">287929</a>	5431	761	3	102	family with sequence similarity 128, member B (Fam128b) alternative variant eSep08, mRNA.
<a href="#">Fam129b</a>	<a href="#">Fam129b.bSep08</a>	<a href="#">362115</a>	1054	726	1	240	family with sequence similarity 129, member B (Fam129b) alternative variant bSep08, mRNA.
<a href="#">Fam131c</a>	<a href="#">Fam131c.aSep08</a>	<a href="#">690880</a>	1138	249		82	family with sequence similarity 131, member C (Fam131c) mRNA.
<a href="#">Fam132b</a>	<a href="#">Fam132b.aSep08</a>	<a href="#">681056</a>	1697	780		91	family with sequence similarity 132, member B (10.0 kD) (Fam132b) mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.bSep08</a>	<a href="#">362320</a>	25101	748	11	138	CRA a like (Fam133b) alternative variant bSep08, mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.cSep08</a>	<a href="#">362320</a>	11509	986	7	121	CRA a like (Fam133b) alternative variant cSep08, mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.dSep08</a>	<a href="#">362320</a>	16662	968	7	98	protein CRA f (11.6 kD) (Fam133b) alternative variant dSep08, mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.eSep08</a>	<a href="#">362320</a>	4688	814	3	63	CRA a like (7.2 kD) (Fam133b) alternative variant eSep08, mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.fSep08</a>	<a href="#">362320</a>	3000	1623	2	51	CRA a like (5.9 kD) (Fam133b) alternative variant fSep08, mRNA.
<a href="#">Fam133b</a>	<a href="#">Fam133b.gSep08</a>	<a href="#">362320</a>	1619	727	4	21	putative protein (2.6 kD) (Fam133b) alternative variant gSep08, mRNA.
<a href="#">Fam134a</a>	<a href="#">Fam134a.aSep08</a>	<a href="#">363252</a>	6239	2931	9	558	family with sequence similarity 134, member A (Fam134a) alternative variant aSep08, mRNA.
<a href="#">Fam134a</a>	<a href="#">Fam134a.bSep08</a>	<a href="#">363252</a>	3670	731	8	243	family with sequence similarity 134, member A (Fam134a) alternative variant bSep08, mRNA.
<a href="#">Fam134a</a>	<a href="#">Fam134a.cSep08</a>	<a href="#">363252</a>	2829	523	6	124	family with sequence similarity 134, member A (Fam134a) alternative variant cSep08, mRNA.

<a href="#">Fam134b</a>	<a href="#">Fam134b.bSep08</a>	<a href="#">619558</a>	35521	650	6	184	family with sequence similarity 134, member B (Fam134b) alternative variant bSep08, mRNA.
<a href="#">Fam134b</a>	<a href="#">Fam134b.dSep08</a>	<a href="#">619558</a>	2122	435	2	69	family with sequence similarity 134, member B (7.7 kD) (Fam134b) alternative variant dSep08, mRNA.
<a href="#">Fam134c</a>	<a href="#">Fam134c.aSep08</a>	<a href="#">360632</a>	24149	3079	9	476	family with sequence similarity 134, member C (Fam134c) alternative variant aSep08, mRNA.
<a href="#">Fam134c</a>	<a href="#">Fam134c.bSep08</a>	<a href="#">360632</a>	1154	752	2	162	family with sequence similarity 134, member C (Fam134c) alternative variant bSep08, mRNA.
<a href="#">Fam134c</a>	<a href="#">Fam134c.cSep08</a>	<a href="#">360632</a>	16755	388	3	129	family with sequence similarity 134, member C (Fam134c) alternative variant cSep08, mRNA.
<a href="#">Fam139a</a>	<a href="#">Fam139a.aSep08</a>	<a href="#">680422</a>	4904	2411		318	family with sequence similarity 139, member A (Fam139a) mRNA.
<a href="#">Fam148b</a>	<a href="#">Fam148b.aSep08</a>	<a href="#">501015</a>	1496	1327		340	family with sequence similarity 148, member B (37.2 kD) (Fam148b) mRNA.
<a href="#">Fam149a</a>	<a href="#">Fam149a.aSep08</a>	<a href="#">361153</a>	6046	508		80	family with sequence similarity 149, member A (9.0 kD) (Fam149a) mRNA.
<a href="#">Fam149b1</a>	<a href="#">Fam149b1.bSep08</a>	<a href="#">289900</a>	10409	652	4	211	family with sequence similarity 149, member B1 (Fam149b1) alternative variant bSep08, mRNA.
<a href="#">Fam149b1</a>	<a href="#">Fam149b1.cSep08</a>	<a href="#">289900</a>	4908	668	3	143	family with sequence similarity 149, member B1 (Fam149b1) alternative variant cSep08, mRNA.
<a href="#">Fam149b1</a>	<a href="#">Fam149b1.dSep08</a>	<a href="#">289900</a>	2238	588	2	84	family with sequence similarity 149, member B1 (Fam149b1) alternative variant dSep08, mRNA.
<a href="#">Fam151b</a>	<a href="#">Fam151b.bSep08</a>	<a href="#">499507</a>	10331	756	4	202	family with sequence similarity 151, member B (Fam151b) alternative variant bSep08, mRNA.
<a href="#">Fam151b</a>	<a href="#">Fam151b.cSep08</a>	<a href="#">499507</a>	15663	786	3	107	family with sequence similarity 151, member B (11.7 kD) (Fam151b) alternative variant cSep08, complete mRNA.
<a href="#">Fam151b</a>	<a href="#">Fam151b.dSep08</a>	<a href="#">499507</a>	35558	346	2	93	family with sequence similarity 151, member B (Fam151b) alternative variant dSep08, mRNA.
<a href="#">Fam151b</a>	<a href="#">Fam151b.eSep08</a>	<a href="#">499507</a>	4945	897	1	59	family with sequence similarity 151, member B (6.5 kD) (Fam151b) alternative variant eSep08, mRNA.
<a href="#">Fam152b</a>	<a href="#">Fam152b.aSep08</a>	<a href="#">315160</a>	22766	2929	2	214	family with sequence similarity 152, member B (Fam152b) alternative variant aSep08, mRNA.
<a href="#">Fam152b</a>	<a href="#">Fam152b.bSep08</a>	<a href="#">315160</a>	21909	2106	2	213	family with sequence similarity 152, member B (Fam152b) alternative variant bSep08, mRNA.
<a href="#">Fam152b</a>	<a href="#">Fam152b.dSep08</a>	<a href="#">315160</a>	16866	701	3	138	family with sequence similarity 152, member B (Fam152b) alternative variant dSep08, mRNA.
<a href="#">Fam166a</a>	<a href="#">Fam166a.bSep08</a>	<a href="#">311797</a>	5460	1405	8	128	family with sequence similarity 166, member A (14.4 kD) (Fam166a) alternative variant bSep08, mRNA.
<a href="#">Fam166a</a>	<a href="#">Fam166a.cSep08</a>	<a href="#">311797</a>	5291	685	7	66	family with sequence similarity 166, member A (Fam166a) alternative variant cSep08, mRNA.
<a href="#">famer</a>	<a href="#">famer.aSep08</a>		1406	517		56	keratin 12 (famer) mRNA.
<a href="#">Fanca</a>	<a href="#">Fanca.aSep08</a>	<a href="#">361435</a>	14902	1667		473	fanconi anemia, complementation group A (Fanca) mRNA.
<a href="#">Fancc</a>	<a href="#">Fancc.bSep08</a>	<a href="#">24361</a>	8147	382	1	115	fanconi anemia, complementation group C (Fancc) alternative variant bSep08, mRNA.
<a href="#">Fandc2</a>	<a href="#">Fandc2.bSep08</a>	<a href="#">312641</a>	28890	1784	12	485	fanconi anemia, complementation group D2 (Fandc2) alternative variant bSep08, mRNA.

<a href="#">Fancd2</a>	<a href="#">Fancd2.cSep08</a>	<a href="#">312641</a>	13907	1237	10	302	fanconi anemia, complementation group D2 (Fancd2) alternative variant cSep08, mRNA.
<a href="#">Fancd2</a>	<a href="#">Fancd2.dSep08</a>	<a href="#">312641</a>	4423	667	3	106	fanconi anemia, complementation group D2 (Fancd2) alternative variant dSep08, mRNA.
<a href="#">Fancd2</a>	<a href="#">Fancd2.eSep08</a>	<a href="#">312641</a>	1650	730	2	78	fanconi anemia, complementation group D2 (Fancd2) alternative variant eSep08, mRNA.
<a href="#">Fance</a>	<a href="#">Fance.aSep08</a>	<a href="#">309643</a>	11115	1938	10	531	fanconi anemia, complementation group E (57.7 kD) (Fance) alternative variant aSep08, mRNA.
<a href="#">Fance</a>	<a href="#">Fance.bSep08</a>	<a href="#">309643</a>	2702	386	4	128	fanconi anemia, complementation group E (Fance) alternative variant bSep08, mRNA.
<a href="#">Fance</a>	<a href="#">Fance.cSep08</a>	<a href="#">309643</a>	4987	500	3	123	fanconi anemia, complementation group E (Fance) alternative variant cSep08, mRNA.
<a href="#">Fance</a>	<a href="#">Fance.dSep08</a>	<a href="#">309643</a>	3304	733	2	112	fanconi anemia, complementation group E (11.5 kD) (Fance) alternative variant dSep08, mRNA.
<a href="#">Fancg</a>	<a href="#">Fancg.aSep08</a>	<a href="#">691105</a>	2638	790		263	fanconi anemia, complementation group G (Fancg) mRNA.
<a href="#">Fancl</a>	<a href="#">Fancl.aSep08</a>	<a href="#">305600</a>	65356	1787		435	fanconi anemia, complementation group L (Fancl) alternative variant aSep08, mRNA.
<a href="#">Fancl</a>	<a href="#">Fancl.bSep08</a>	<a href="#">305600</a>	65386	1658	5	377	fanconi anemia, complementation group L (Fancl) alternative variant bSep08, mRNA.
<a href="#">Fancl</a>	<a href="#">Fancl.cSep08</a>	<a href="#">305600</a>	61655	773	4	179	fanconi anemia, complementation group L (Fancl) alternative variant cSep08, mRNA.
<a href="#">fanoy</a>	<a href="#">fanoy.cSep08</a>		1763	768	2	109	putative protein (fanoy) alternative variant cSep08, mRNA.
<a href="#">Fap</a>	<a href="#">Fap.bSep08</a>	<a href="#">192203</a>	23127	703	9	234	fibroblast activation protein (Fap) alternative variant bSep08, mRNA.
<a href="#">Fap</a>	<a href="#">Fap.cSep08</a>	<a href="#">192203</a>	16415	725	1	163	fibroblast activation protein (Fap) alternative variant cSep08, mRNA.
<a href="#">fapor</a>	<a href="#">fapor.aSep08</a>		6926	596		17	putative protein (2.1 kD) (fapor) mRNA.
<a href="#">farby</a>	<a href="#">farby.aSep08</a>		9521	401		133	cofactor required for Sp1 transcriptional activation 150kDa CRA b (farby) mRNA.
<a href="#">farchy</a>	<a href="#">farchy.aSep08</a>		568	300	2	63	putative protein (farchy) alternative variant aSep08, mRNA.
<a href="#">farchy</a>	<a href="#">farchy.bSep08</a>		7346	1057	2	61	putative protein (farchy) alternative variant bSep08, mRNA.
<a href="#">farfer</a>	<a href="#">farfer.aSep08</a>		10527	958		319	putative protein of metazoan origin (farfer) mRNA.
<a href="#">farflo</a>	<a href="#">farflo.aSep08</a>		17516	587		195	transient receptor potential cation channel subfamily M member 3 (farflo) mRNA.
<a href="#">farflu</a>	<a href="#">farflu.aSep08</a>		1260	433		35	putative protein (farflu) mRNA.
<a href="#">farkee</a>	<a href="#">farkee.aSep08</a>		983	413		64	putative protein (farkee) mRNA.
<a href="#">farloy</a>	<a href="#">farloy.aSep08</a>		8887	241		33	putative protein (farloy) mRNA.
<a href="#">farmer</a>	<a href="#">farmer.aSep08</a>		1503	641		75	putative cytoplasmic protein (8.4 kD) (farmer) mRNA.
<a href="#">farnoy</a>	<a href="#">farnoy.aSep08</a>		1807	471		150	CRA a (farnoy) mRNA.
<a href="#">Farp2</a>	<a href="#">Farp2.bSep08</a>	<a href="#">316639</a>	2662	722	4	156	FERM, RhoGEF and pleckstrin domain protein 2 (18.2 kD) (Farp2) alternative variant bSep08, mRNA.
<a href="#">Farp2</a>	<a href="#">Farp2.cSep08</a>	<a href="#">316639</a>	52500	481	5	132	FERM, RhoGEF and pleckstrin domain protein 2 (Farp2) alternative variant cSep08, mRNA.
<a href="#">farpor</a>	<a href="#">farpor.aSep08</a>		1257	514		50	nicotinamide N-methyltransferase (farpor) mRNA.



<a href="#">Fars2</a>	<a href="#">Fars2.bSep08</a>	<a href="#">306879</a>	375166	1366	5	403	phenylalanine-tRNA synthetase 2 (mitochondrial) (Fars2) alternative variant bSep08, mRNA.
<a href="#">Fars2</a>	<a href="#">Fars2.cSep08</a>	<a href="#">306879</a>	20256	727	2	118	phenylalanine-tRNA synthetase 2 (mitochondrial) (13.1 kD) (Fars2) alternative variant cSep08, mRNA.
<a href="#">Farsa</a>	<a href="#">Farsa.bSep08</a>	<a href="#">288917</a>	1052	612	2	117	phenylalanyl-tRNA synthetase, alpha subunit (Farsa) alternative variant bSep08, mRNA.
<a href="#">Farsa</a>	<a href="#">Farsa.cSep08</a>	<a href="#">288917</a>	3666	713	3	113	phenylalanyl-tRNA synthetase, alpha subunit (Farsa) alternative variant cSep08, mRNA.
<a href="#">farshee</a>	<a href="#">farshee.aSep08</a>		10921	540		179	membrane-associated ring finger 6 (farshee) mRNA.
<a href="#">farto</a>	<a href="#">farto.aSep08</a>		2366	1304		117	putative protein (farto) mRNA.
<a href="#">farvar</a>	<a href="#">farvar.aSep08</a>		12547	648		47	putative protein (5.5 kD) (farvar) mRNA.
<a href="#">farwey</a>	<a href="#">farwey.aSep08</a>		5062	633		86	putative protein of eukaryotic origin (9.4 kD) (farwey) mRNA.
<a href="#">Fas</a>	<a href="#">Fas.aSep08</a>	<a href="#">246097</a>	14538	1705		304	fas (TNF receptor superfamily, member 6) (34.6 kD) (Fas) mRNA.
<a href="#">fasa</a>	<a href="#">fasa.aSep08</a>		1314	733	1	89	putative protein (fasa) alternative variant aSep08, mRNA.
<a href="#">fasa</a>	<a href="#">fasa.bSep08</a>		6386	692	2	38	putative protein (4.3 kD) (fasa) alternative variant bSep08, mRNA.
<a href="#">Fasciclin.0</a>	<a href="#">Fasciclin.0.aSep08</a>		1036	382		127	stabilin 1 (Fasciclin.0) mRNA.
<a href="#">fashee</a>	<a href="#">fashee.aSep08</a>		17657	750		87	putative protein (9.6 kD) (fashee) mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.aSep08</a>	<a href="#">296741</a>	4028	1824	10	545	fas-activated serine/threonine kinase (61.3 kD) (Fastk) alternative variant aSep08, complete mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.cSep08</a>	<a href="#">296741</a>	4027	2289	7	255	fas-activated serine/threonine kinase (28.9 kD) (Fastk) alternative variant cSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.dSep08</a>	<a href="#">296741</a>	1875	1427	2	177	fas-activated serine/threonine kinase (Fastk) alternative variant dSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.eSep08</a>	<a href="#">296741</a>	1848	551	4	166	fas-activated serine/threonine kinase (Fastk) alternative variant eSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.fSep08</a>	<a href="#">296741</a>	1353	694	4	137	fas-activated serine/threonine kinase (Fastk) alternative variant fSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.gSep08</a>	<a href="#">296741</a>	565	488	2	77	fas-activated serine/threonine kinase (Fastk) alternative variant gSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.hSep08</a>	<a href="#">296741</a>	983	694	2	48	fas-activated serine/threonine kinase (Fastk) alternative variant hSep08, mRNA.
<a href="#">Fastk</a>	<a href="#">Fastk.iSep08</a>	<a href="#">296741</a>	880	679	2	63	fas-activated serine/threonine kinase (Fastk) alternative variant iSep08, mRNA.
<a href="#">Fastkd1</a>	<a href="#">Fastkd1.aSep08</a>	<a href="#">311112</a>	7874	878		292	FAST kinase domains 1 (Fastkd1) mRNA.
<a href="#">Fastkd2</a>	<a href="#">Fastkd2.bSep08</a>	<a href="#">301463</a>	1270	325	2	82	FAST kinase domains 2 (Fastkd2) alternative variant bSep08, mRNA.
<a href="#">Fastkd3</a>	<a href="#">Fastkd3.bSep08</a>	<a href="#">290946</a>	7307	2057	5	515	fast kinase domains 3 CRA a (Fastkd3) alternative variant bSep08, mRNA.
<a href="#">Fastkd3</a>	<a href="#">Fastkd3.dSep08</a>	<a href="#">290946</a>	3089	1005	3	102	kinase 3 (11.9 kD) (Fastkd3) alternative variant dSep08, mRNA.
<a href="#">Fastkd3</a>	<a href="#">Fastkd3.eSep08</a>	<a href="#">290946</a>	5249	708	4	95	kinase 3 (10.8 kD) (Fastkd3) alternative variant eSep08, mRNA.

<a href="#">Fat1</a>	<a href="#">Fat1.bSep08</a>	<a href="#">83720</a>	8346	2117	4	437	FAT tumor suppressor homolog 1 (Drosophila) (Fat1) alternative variant bSep08, mRNA.
<a href="#">Fat1</a>	<a href="#">Fat1.cSep08</a>	<a href="#">83720</a>	7150	938	4	126	FAT tumor suppressor homolog 1 (Drosophila) (Fat1) alternative variant cSep08, mRNA.
<a href="#">Fat1</a>	<a href="#">Fat1.dSep08</a>	<a href="#">83720</a>	3314	1206	2	98	FAT tumor suppressor homolog 1 (Drosophila) (11.2 kD) (Fat1) alternative variant dSep08, mRNA.
<a href="#">Fat1</a>	<a href="#">Fat1.eSep08</a>	<a href="#">83720</a>	6021	456	4	26	FAT tumor suppressor homolog 1 (Drosophila) (Fat1) alternative variant eSep08, mRNA.
<a href="#">Fat2</a>	<a href="#">Fat2.bSep08</a>	<a href="#">65048</a>	3259	2234	2	284	FAT tumor suppressor homolog 2 (Drosophila) (Fat2) alternative variant bSep08, mRNA.
<a href="#">Fat2</a>	<a href="#">Fat2.cSep08</a>	<a href="#">65048</a>	4133	347	3	115	FAT tumor suppressor homolog 2 (Drosophila) (Fat2) alternative variant cSep08, mRNA.
<a href="#">Fat4</a>	<a href="#">Fat4.aSep08</a>	<a href="#">310341</a>	64424	2139		712	FAT tumor suppressor homolog 4 (Drosophila) (Fat4) mRNA.
<a href="#">FATC.0</a>	<a href="#">FATC.0.aSep08</a>		5627	410		127	ataxia telangiectasia Rad3 related (FATC.0) mRNA.
<a href="#">fato</a>	<a href="#">fato.aSep08</a>		5186	346		93	putative secreted or extracellular protein precursor (10.1 kD) (fato) alternative variant aSep08, mRNA.
<a href="#">Fau</a>	<a href="#">Fau.aSep08</a>	<a href="#">29752</a>	1509	504	2	133	finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (14.4 kD) (Fau) alternative variant aSep08, complete mRNA.
<a href="#">Fau</a>	<a href="#">Fau.bSep08</a>	<a href="#">29752</a>	1488	485	2	133	finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (14.4 kD) (Fau) alternative variant bSep08, complete mRNA.
<a href="#">favar</a>	<a href="#">favar.aSep08</a>		6133	852		255	putative protein, with 2 coiled coil domains, of vertebrate origin (favar) mRNA.
<a href="#">fawby</a>	<a href="#">fawby.aSep08</a>		5019	2434	2	72	cofactor required for Sp1 transcriptional activation CRA a (fawby) alternative variant aSep08, mRNA.
<a href="#">fawchy</a>	<a href="#">fawchy.aSep08</a>		2306	289		54	putative protein (6.3 kD) (fawchy) mRNA.
<a href="#">fawey</a>	<a href="#">fawey.aSep08</a>		13820	793		118	putative protein (fawey) mRNA.
<a href="#">fawfer</a>	<a href="#">fawfer.aSep08</a>		13217	585		107	putative protein (fawfer) mRNA.
<a href="#">fawflo</a>	<a href="#">fawflo.aSep08</a>		5793	261	2	36	putative protein (4.0 kD) (fawflo) alternative variant aSep08, mRNA.
<a href="#">fawflo</a>	<a href="#">fawflo.bSep08</a>		6486	823	3	35	putative protein (3.9 kD) (fawflo) alternative variant bSep08, mRNA.
<a href="#">fawflu</a>	<a href="#">fawflu.aSep08</a>		786	694		92	putative protein (10.4 kD) (fawflu) mRNA.
<a href="#">fawkee</a>	<a href="#">fawkee.aSep08</a>		7674	1108	2	87	putative protein (9.6 kD) (fawkee) alternative variant aSep08, mRNA.
<a href="#">fawkee</a>	<a href="#">fawkee.bSep08</a>		7088	412		32	putative protein (fawkee) alternative variant bSep08, mRNA.
<a href="#">fawloy</a>	<a href="#">fawloy.aSep08</a>		14814	640		13	putative protein (1.5 kD) (fawloy) mRNA.
<a href="#">fawmer</a>	<a href="#">fawmer.aSep08</a>		5800	634		64	putative protein (fawmer) mRNA.
<a href="#">fawnoy</a>	<a href="#">fawnoy.aSep08</a>		9291	621		51	CRA a like (fawnoy) mRNA.
<a href="#">fawpor</a>	<a href="#">fawpor.aSep08</a>		1262	293		50	putative protein (fawpor) mRNA.
<a href="#">fawsa</a>	<a href="#">fawsa.aSep08</a>		2346	251		38	putative protein (fawsa) mRNA.
<a href="#">fawshee</a>	<a href="#">fawshee.aSep08</a>		79754	725		60	putative protein (6.7 kD) (fawshee) mRNA.

<a href="#">fawto</a>	<a href="#">fawto.aSep08</a>		398	271		41	putative protein (fawto) mRNA.
<a href="#">fawvar</a>	<a href="#">fawvar.aSep08</a>		22064	1380		24	putative protein (fawvar) mRNA.
<a href="#">fawwey</a>	<a href="#">fawwey.bSep08</a>		4720	362	2	29	putative protein (3.5 kD) (fawwey) alternative variant bSep08, mRNA.
<a href="#">FBA.0</a>	<a href="#">FBA.0.bSep08</a>	<a href="#">685932</a>	4671	1009	2	257	F-box protein 27 (FBA.0) alternative variant bSep08, mRNA.
<a href="#">FBA.0</a>	<a href="#">FBA.0.cSep08</a>	<a href="#">685932</a>	2612	1071	1	80	f-box protein 27 (FBA.0) alternative variant cSep08, mRNA.
<a href="#">Fbf1</a>	<a href="#">Fbf1.aSep08</a>	<a href="#">287836</a>	8009	2035	15	632	fas (TNFRSF6) binding factor 1 (Fbf1) alternative variant aSep08, mRNA.
<a href="#">Fbf1</a>	<a href="#">Fbf1.cSep08</a>	<a href="#">287836</a>	2900	750	4	205	fas (TNFRSF6) binding factor 1 (Fbf1) alternative variant cSep08, mRNA.
<a href="#">Fbf1</a>	<a href="#">Fbf1.dSep08</a>	<a href="#">287836</a>	3815	629	2	56	fas (TNFRSF6) binding factor 1 (Fbf1) alternative variant dSep08, mRNA.
<a href="#">Fblim1</a>	<a href="#">Fblim1.bSep08</a>	<a href="#">362650</a>	11887	2215	4	181	filamin binding LIM protein 1 (Fblim1) alternative variant bSep08, mRNA.
<a href="#">Fblim1</a>	<a href="#">Fblim1.cSep08</a>	<a href="#">362650</a>	14585	440	4	146	filamin binding LIM protein 1 (Fblim1) alternative variant cSep08, mRNA.
<a href="#">Fblim1</a>	<a href="#">Fblim1.dSep08</a>	<a href="#">362650</a>	4488	401	3	133	filamin binding LIM protein 1 (Fblim1) alternative variant dSep08, mRNA.
<a href="#">Fblim1</a>	<a href="#">Fblim1.eSep08</a>	<a href="#">362650</a>	5826	856	2	56	filamin binding LIM protein 1 (Fblim1) alternative variant eSep08, mRNA.
<a href="#">Fbln1</a>	<a href="#">Fbln1.bSep08</a>	<a href="#">315191</a>	44517	2232	15	685	fibulin 1 (75.4 kD) (Fbln1) alternative variant bSep08, complete mRNA.
<a href="#">Fbln2</a>	<a href="#">Fbln2.aSep08</a>	<a href="#">282583</a>	60530	4193	14	1174	fibulin 2 (126.0 kD) (Fbln2) alternative variant aSep08, mRNA.
<a href="#">Fbln2</a>	<a href="#">Fbln2.bSep08</a>	<a href="#">282583</a>	9026	738	3	245	fibulin 2 (Fbln2) alternative variant bSep08, mRNA.
<a href="#">Fbln5</a>	<a href="#">Fbln5.bSep08</a>	<a href="#">29158</a>	59504	686	1	228	fibulin 5 (Fbln5) alternative variant bSep08, mRNA.
<a href="#">Fbln7</a>	<a href="#">Fbln7.aSep08</a>	<a href="#">296145</a>	15723	407		135	fibulin 7 (Fbln7) mRNA.
<a href="#">Fbn1</a>	<a href="#">Fbn1.bSep08</a>	<a href="#">83727</a>	4537	430	3	95	fibrillin 1 (Fbn1) alternative variant bSep08, mRNA.
<a href="#">Fbn1</a>	<a href="#">Fbn1.cSep08</a>	<a href="#">83727</a>	8567	499	5	36	fibrillin 1 (3.7 kD) (Fbn1) alternative variant cSep08, mRNA.
<a href="#">Fbn2</a>	<a href="#">Fbn2.aSep08</a>	<a href="#">689008</a>	32564	4192	6	861	fibrillin 2 (Fbn2) alternative variant aSep08, mRNA.
<a href="#">Fbn2</a>	<a href="#">Fbn2.bSep08</a>	<a href="#">689008</a>	2235	728	1	202	fibrillin 2 (Fbn2) alternative variant bSep08, mRNA.
<a href="#">Fbp1</a>	<a href="#">Fbp1.bSep08</a>	<a href="#">24362</a>	17884	966	4	282	fructose-1,6- biphosphatase 1 (Fbp1) alternative variant bSep08, mRNA.
<a href="#">Fbp1</a>	<a href="#">Fbp1.cSep08</a>	<a href="#">24362</a>	8140	909	5	252	fructose-1,6- biphosphatase 1 (Fbp1) alternative variant cSep08, mRNA.
<a href="#">Fbp1</a>	<a href="#">Fbp1.dSep08</a>	<a href="#">24362</a>	4703	753	3	201	fructose-1,6- biphosphatase 1 (Fbp1) alternative variant dSep08, mRNA.
<a href="#">Fbp2</a>	<a href="#">Fbp2.bSep08</a>	<a href="#">114508</a>	3821	673	2	127	fructose-1,6-bisphosphatase 2 (13.9 kD) (Fbp2) alternative variant bSep08, mRNA.
<a href="#">Fbp2</a>	<a href="#">Fbp2.cSep08</a>	<a href="#">114508</a>	2960	658	1	91	fructose-1,6-bisphosphatase 2 (10.0 kD) (Fbp2) alternative variant cSep08, mRNA.
<a href="#">Fbrs</a>	<a href="#">Fbrs.aSep08</a>	<a href="#">691899</a>	6125	2645	11	482	fibrosin (Fbrs) alternative variant aSep08, mRNA.
<a href="#">Fbxl2</a>	<a href="#">Fbxl2.aSep08</a>	<a href="#">363156</a>	26413	659	1	207	F-box and leucine-rich repeat protein 2 (Fbxl2) alternative variant aSep08, mRNA.

<a href="#">Fbxl2</a>	<a href="#">Fbxl2.bSep08</a>	<a href="#">363156</a>	29652	748	2	162	F-box and leucine-rich repeat protein 2 (Fbxl2) alternative variant bSep08, mRNA.
<a href="#">Fbxl3</a>	<a href="#">Fbxl3.aSep08</a>	<a href="#">306129</a>	18210	1623	1	478	F-box and leucine-rich repeat protein 3 (Fbxl3) alternative variant aSep08, mRNA.
<a href="#">Fbxl3</a>	<a href="#">Fbxl3.bSep08</a>	<a href="#">306129</a>	10375	593	1	197	F-box and leucine-rich repeat protein 3 (Fbxl3) alternative variant bSep08, mRNA.
<a href="#">Fbxl4</a>	<a href="#">Fbxl4.aSep08</a>	<a href="#">313101</a>	73323	2487	9	621	F-box and leucine-rich repeat protein 4 (70.2 kD) (Fbxl4) alternative variant aSep08, mRNA.
<a href="#">Fbxl5</a>	<a href="#">Fbxl5.bSep08</a>	<a href="#">305424</a>	10554	1240	6	412	F-box and leucine-rich repeat protein 5 (Fbxl5) alternative variant bSep08, mRNA.
<a href="#">Fbxl6</a>	<a href="#">Fbxl6.bSep08</a>	<a href="#">362941</a>	2878	2111	6	324	F-box and leucine-rich repeat protein 6 (34.8 kD) (Fbxl6) alternative variant bSep08, mRNA.
<a href="#">Fbxl6</a>	<a href="#">Fbxl6.cSep08</a>	<a href="#">362941</a>	1296	730	5	212	F-box and leucine-rich repeat protein 6 (Fbxl6) alternative variant cSep08, mRNA.
<a href="#">Fbxl10</a>	<a href="#">Fbxl10.aSep08</a>	<a href="#">304495</a>	55141	694		218	F-box and leucine-rich repeat protein 10 (Fbxl10) alternative variant aSep08, mRNA.
<a href="#">Fbxl11</a>	<a href="#">Fbxl11.bSep08</a>	<a href="#">361700</a>	3137	571	2	94	F-box and leucine-rich repeat protein 11 (Fbxl11) alternative variant bSep08, mRNA.
<a href="#">Fbxl12</a>	<a href="#">Fbxl12.bSep08</a>	<a href="#">313782</a>	3579	751	2	179	F-box and leucine-rich repeat protein 12 (Fbxl12) alternative variant bSep08, mRNA.
<a href="#">Fbxl17</a>	<a href="#">Fbxl17.bSep08</a>	<a href="#">316663</a>	156019	529	5	159	F-box and leucine-rich repeat protein 17 (Fbxl17) alternative variant bSep08, mRNA.
<a href="#">Fbxl17</a>	<a href="#">Fbxl17.cSep08</a>	<a href="#">316663</a>	99794	644	3	57	F-box and leucine-rich repeat protein 17 (Fbxl17) alternative variant cSep08, mRNA.
<a href="#">Fbxl17</a>	<a href="#">Fbxl17.dSep08</a>	<a href="#">316663</a>	86107	1450	4	135	F-box and leucine-rich repeat protein 17 (15.4 kD) (Fbxl17) alternative variant dSep08, mRNA.
<a href="#">Fbxl19</a>	<a href="#">Fbxl19.bSep08</a>	<a href="#">308999</a>	15897	2445	3	298	F-box and leucine-rich repeat protein 19 (Fbxl19) alternative variant bSep08, mRNA.
<a href="#">Fbxl20</a>	<a href="#">Fbxl20.aSep08</a>	<a href="#">64039</a>	49202	1794	4	585	F-box and leucine-rich repeat protein 20 (Fbxl20) alternative variant aSep08, mRNA.
<a href="#">Fbxl20</a>	<a href="#">Fbxl20.bSep08</a>	<a href="#">64039</a>	57840	2450	15	438	F-box and leucine-rich repeat protein 20 (48.5 kD) (Fbxl20) alternative variant bSep08, mRNA.
<a href="#">Fbxl20</a>	<a href="#">Fbxl20.eSep08</a>	<a href="#">64039</a>	2478	228	2	76	F-box and leucine-rich repeat protein 20 (Fbxl20) alternative variant eSep08, mRNA.
<a href="#">Fbxo3</a>	<a href="#">Fbxo3.bSep08</a>	<a href="#">690634</a>	7646	772	3	97	F-box protein 3 (Fbxo3) alternative variant bSep08, mRNA.
<a href="#">Fbxo3</a>	<a href="#">Fbxo3.cSep08</a>	<a href="#">690634</a>	16485	753	4	94	F-box protein 3 (Fbxo3) alternative variant cSep08, mRNA.
<a href="#">Fbxo6</a>	<a href="#">Fbxo6.bSep08</a>	<a href="#">192351</a>	4980	785	5	204	F-box protein 6 (Fbxo6) alternative variant bSep08, mRNA.
<a href="#">Fbxo6</a>	<a href="#">Fbxo6.cSep08</a>	<a href="#">192351</a>	2282	380	2	126	F-box protein 6 (Fbxo6) alternative variant cSep08, mRNA.
<a href="#">Fbxo7</a>	<a href="#">Fbxo7.bSep08</a>	<a href="#">366854</a>	11136	625	5	208	F-box protein 7 (Fbxo7) alternative variant bSep08, mRNA.
<a href="#">Fbxo7</a>	<a href="#">Fbxo7.cSep08</a>	<a href="#">366854</a>	11497	1138	5	194	F-box protein 7 (Fbxo7) alternative variant cSep08, mRNA.
<a href="#">Fbxo7</a>	<a href="#">Fbxo7.dSep08</a>	<a href="#">366854</a>	11403	707	5	137	F-box protein 7 (Fbxo7) alternative variant dSep08, mRNA.
<a href="#">Fbxo7</a>	<a href="#">Fbxo7.eSep08</a>	<a href="#">366854</a>	2138	702	2	85	F-box protein 7 (Fbxo7) alternative variant eSep08, mRNA.
<a href="#">Fbxo8</a>	<a href="#">Fbxo8.bSep08</a>	<a href="#">306436</a>	21871	466	4	63	F-box protein 8 (7.5 kD) (Fbxo8) alternative variant bSep08, mRNA.
<a href="#">Fbxo8</a>	<a href="#">Fbxo8.cSep08</a>	<a href="#">306436</a>	1472	694	2	45	F-box protein 8 (5.6 kD) (Fbxo8) alternative variant cSep08, mRNA.

<a href="#">Fbxo9</a>	<a href="#">Fbxo9.bSep08</a>	<a href="#">300849</a>	13178	1710	8	167	f-box protein 9 (20.2 kD) (Fbxo9) alternative variant bSep08, mRNA.
<a href="#">Fbxo9</a>	<a href="#">Fbxo9.cSep08</a>	<a href="#">300849</a>	20072	807	8	147	f-box protein 9 (Fbxo9) alternative variant cSep08, mRNA.
<a href="#">Fbxo9</a>	<a href="#">Fbxo9.dSep08</a>	<a href="#">300849</a>	3832	864	2	136	f-box protein 9 (Fbxo9) alternative variant dSep08, mRNA.
<a href="#">Fbxo10</a>	<a href="#">Fbxo10.aSep08</a>	<a href="#">362511</a>	6070	1019	4	300	F-box protein 10 (Fbxo10) alternative variant aSep08, mRNA.
<a href="#">Fbxo10</a>	<a href="#">Fbxo10.bSep08</a>	<a href="#">362511</a>	6573	971	2	101	F-box protein 10 (Fbxo10) alternative variant bSep08, mRNA.
<a href="#">Fbxo11</a>	<a href="#">Fbxo11.bSep08</a>	<a href="#">301674</a>	554	343	2	95	F-box protein 11 (Fbxo11) alternative variant bSep08, mRNA.
<a href="#">Fbxo16</a>	<a href="#">Fbxo16.bSep08</a>	<a href="#">305970</a>	23101	537	1	92	F-box protein 16 (Fbxo16) alternative variant bSep08, mRNA.
<a href="#">Fbxo17</a>	<a href="#">Fbxo17.bSep08</a>	<a href="#">292757</a>	15475	1697	7	250	F-box protein 17 (28.2 kD) (Fbxo17) alternative variant bSep08, mRNA.
<a href="#">Fbxo17</a>	<a href="#">Fbxo17.cSep08</a>	<a href="#">292757</a>	2317	519	3	152	F-box protein 17 (Fbxo17) alternative variant cSep08, mRNA.
<a href="#">Fbxo18</a>	<a href="#">Fbxo18.bSep08</a>	<a href="#">291293</a>	6258	756	6	252	F-box protein 18 (Fbxo18) alternative variant bSep08, mRNA.
<a href="#">Fbxo18</a>	<a href="#">Fbxo18.cSep08</a>	<a href="#">291293</a>	2697	608	4	166	F-box protein 18 (Fbxo18) alternative variant cSep08, mRNA.
<a href="#">Fbxo18</a>	<a href="#">Fbxo18.dSep08</a>	<a href="#">291293</a>	4103	554	5	145	F-box protein 18 (Fbxo18) alternative variant dSep08, mRNA.
<a href="#">Fbxo18</a>	<a href="#">Fbxo18.eSep08</a>	<a href="#">291293</a>	1138	751	2	123	F-box protein 18 (14.7 kD) (Fbxo18) alternative variant eSep08, mRNA.
<a href="#">Fbxo18</a>	<a href="#">Fbxo18.gSep08</a>	<a href="#">291293</a>	13356	840	3	175	F-box protein 18 (Fbxo18) alternative variant gSep08, mRNA.
<a href="#">Fbxo21</a>	<a href="#">Fbxo21.bSep08</a>	<a href="#">360818</a>	19247	1492	1	486	F-box protein 21 (55.3 kD) (Fbxo21) alternative variant bSep08, mRNA.
<a href="#">Fbxo22</a>	<a href="#">Fbxo22.bSep08</a>	<a href="#">300724</a>	12611	796	5	251	F-box protein 22 (Fbxo22) alternative variant bSep08, mRNA.
<a href="#">Fbxo22</a>	<a href="#">Fbxo22.cSep08</a>	<a href="#">300724</a>	15149	744	5	226	F-box protein 22 (Fbxo22) alternative variant cSep08, mRNA.
<a href="#">Fbxo22</a>	<a href="#">Fbxo22.dSep08</a>	<a href="#">300724</a>	12695	806	5	155	F-box protein 22 (Fbxo22) alternative variant dSep08, mRNA.
<a href="#">Fbxo22</a>	<a href="#">Fbxo22.eSep08</a>	<a href="#">300724</a>	12271	706	5	126	F-box protein 22 (Fbxo22) alternative variant eSep08, mRNA.
<a href="#">Fbxo22</a>	<a href="#">Fbxo22.fSep08</a>	<a href="#">300724</a>	9858	781	3	100	F-box protein 22 (Fbxo22) alternative variant fSep08, mRNA.
<a href="#">Fbxo23</a>	<a href="#">Fbxo23.bSep08</a>	<a href="#">306771</a>	3383	583	4	138	F-box only protein 23 (Fbxo23) alternative variant bSep08, mRNA.
<a href="#">Fbxo23</a>	<a href="#">Fbxo23.cSep08</a>	<a href="#">306771</a>	1356	550	3	130	F-box only protein 23 (Fbxo23) alternative variant cSep08, mRNA.
<a href="#">Fbxo23</a>	<a href="#">Fbxo23.dSep08</a>	<a href="#">306771</a>	1776	507	4	118	F-box only protein 23 (Fbxo23) alternative variant dSep08, mRNA.
<a href="#">Fbxo23</a>	<a href="#">Fbxo23.eSep08</a>	<a href="#">306771</a>	3238	376	3	70	F-box only protein 23 (Fbxo23) alternative variant eSep08, mRNA.

<a href="#">Fbxo24</a>	<a href="#">Fbxo24.aSep08</a>	<a href="#">304374</a>	5915	1314	3	438	F-box protein 24 (Fbxo24) alternative variant aSep08, mRNA.
<a href="#">Fbxo24</a>	<a href="#">Fbxo24.cSep08</a>	<a href="#">304374</a>	2915	755	1	251	F-box protein 24 (Fbxo24) alternative variant cSep08, mRNA.
<a href="#">Fbxo25</a>	<a href="#">Fbxo25.aSep08</a>	<a href="#">364637</a>	11308	1385	4	147	F-box protein 25 (Fbxo25) alternative variant aSep08, mRNA.
<a href="#">Fbxo25</a>	<a href="#">Fbxo25.dSep08</a>	<a href="#">364637</a>	4162	755	2	60	F-box protein 25 (7.1 kD) (Fbxo25) alternative variant dSep08, mRNA.
<a href="#">Fbxo25</a>	<a href="#">Fbxo25.eSep08</a>	<a href="#">364637</a>	3980	600	3	48	F-box protein 25 (Fbxo25) alternative variant eSep08, mRNA.
<a href="#">Fbxo25</a>	<a href="#">Fbxo25.fSep08</a>	<a href="#">364637</a>	1323	746	2	40	F-box protein 25 (4.9 kD) (Fbxo25) alternative variant fSep08, mRNA.
<a href="#">Fbxo28</a>	<a href="#">Fbxo28.bSep08</a>	<a href="#">305105</a>	24718	699	5	176	F-box protein 28 (Fbxo28) alternative variant bSep08, mRNA.
<a href="#">Fbxo30</a>	<a href="#">Fbxo30.bSep08</a>	<a href="#">308283</a>	8290	762	2	200	F-box protein 30 (Fbxo30) alternative variant bSep08, mRNA.
<a href="#">Fbxo33</a>	<a href="#">Fbxo33.aSep08</a>	<a href="#">314157</a>	32088	3429	4	558	F-box protein 33 (Fbxo33) alternative variant aSep08, mRNA.
<a href="#">Fbxo34</a>	<a href="#">Fbxo34.bSep08</a>	<a href="#">305830</a>	72728	3102	2	681	F-box protein 34 (74.9 kD) (Fbxo34) alternative variant bSep08, mRNA.
<a href="#">Fbxo34</a>	<a href="#">Fbxo34.cSep08</a>	<a href="#">305830</a>	70206	810	3	66	F-box protein 34 (Fbxo34) alternative variant cSep08, mRNA.
<a href="#">Fbxo36</a>	<a href="#">Fbxo36.bSep08</a>	<a href="#">363268</a>	79257	707	1	138	F-box protein 36 (Fbxo36) alternative variant bSep08, mRNA.
<a href="#">Fbxo36</a>	<a href="#">Fbxo36.cSep08</a>	<a href="#">363268</a>	78917	303	1	62	F-box protein 36 (Fbxo36) alternative variant cSep08, mRNA.
<a href="#">Fbxo38</a>	<a href="#">Fbxo38.bSep08</a>	<a href="#">307390</a>	23325	1095	8	154	F-box protein 38 (17.9 kD) (Fbxo38) alternative variant bSep08, mRNA.
<a href="#">Fbxo38</a>	<a href="#">Fbxo38.cSep08</a>	<a href="#">307390</a>	7555	431	4	143	F-box protein 38 (Fbxo38) alternative variant cSep08, mRNA.
<a href="#">Fbxo39</a>	<a href="#">Fbxo39.bSep08</a>	<a href="#">303287</a>	2246	693	2	126	F-box protein 39 (3.8 kD) (Fbxo39) alternative variant bSep08, mRNA.
<a href="#">Fbxo39</a>	<a href="#">Fbxo39.cSep08</a>	<a href="#">303287</a>	1047	476	1	80	F-box protein 39 (10.0 kD) (Fbxo39) alternative variant cSep08, mRNA.
<a href="#">Fbxo40</a>	<a href="#">Fbxo40.aSep08</a>	<a href="#">363790</a>	4060	1741		101	F-box protein 40 (Fbxo40) mRNA.
<a href="#">Fbxo42</a>	<a href="#">Fbxo42.bSep08</a>	<a href="#">362646</a>	6700	2185	5	467	F-box protein 42 (Fbxo42) alternative variant bSep08, mRNA.
<a href="#">Fbxo43</a>	<a href="#">Fbxo43.aSep08</a>	<a href="#">315034</a>	2097	752	2	155	F-box protein 43 (Fbxo43) alternative variant aSep08, mRNA.
<a href="#">Fbxo44</a>	<a href="#">Fbxo44.aSep08</a>	<a href="#">500587</a>	9154	1682	6	255	F-box protein 44 (29.8 kD) (Fbxo44) alternative variant aSep08, mRNA.
<a href="#">Fbxo44</a>	<a href="#">Fbxo44.bSep08</a>	<a href="#">500587</a>	4076	760	4	150	F-box protein 44 (Fbxo44) alternative variant bSep08, mRNA.
<a href="#">Fbxo44</a>	<a href="#">Fbxo44.cSep08</a>	<a href="#">500587</a>	1286	424	2	141	F-box protein 44 (Fbxo44) alternative variant cSep08, mRNA.
<a href="#">Fbxo44</a>	<a href="#">Fbxo44.dSep08</a>	<a href="#">500587</a>	4039	451	4	139	F-box protein 44 (Fbxo44) alternative variant dSep08, mRNA.

<a href="#">Fbxo46</a>	<a href="#">Fbxo46.bSep08</a>	<a href="#">292686</a>	14420	1062	2	271	F-box protein 46 (Fbxo46) alternative variant bSep08, mRNA.
<a href="#">Fbxo46</a>	<a href="#">Fbxo46.cSep08</a>	<a href="#">292686</a>	3636	378	1	80	F-box protein 46 (Fbxo46) alternative variant cSep08, mRNA.
<a href="#">Fbxw2</a>	<a href="#">Fbxw2.bSep08</a>	<a href="#">311881</a>	21212	1991	6	354	F-box wd-40 domain protein 2 (40.4 kD) (Fbxw2) alternative variant bSep08, mRNA.
<a href="#">Fbxw2</a>	<a href="#">Fbxw2.cSep08</a>	<a href="#">311881</a>	20709	1423	4	193	repeat-containing protein 2 (22.1 kD) (Fbxw2) alternative variant cSep08, mRNA.
<a href="#">Fbxw2</a>	<a href="#">Fbxw2.fSep08</a>	<a href="#">311881</a>	15489	652	4	43	putative protein (Fbxw2) alternative variant fSep08, mRNA.
<a href="#">Fbxw2</a>	<a href="#">Fbxw2.hSep08</a>	<a href="#">311881</a>	798	697	2	91	putative protein (Fbxw2) alternative variant hSep08, mRNA.
<a href="#">Fbxw2</a>	<a href="#">Fbxw2.iSep08</a>	<a href="#">311881</a>	3230	472	3	33	putative protein (3.7 kD) (Fbxw2) alternative variant iSep08, mRNA.
<a href="#">Fbxw4</a>	<a href="#">Fbxw4.bSep08</a>	<a href="#">309444</a>	102643	1964	7	238	F-box and WD-40 domain protein 4 (25.9 kD) (Fbxw4) alternative variant bSep08, mRNA.
<a href="#">Fbxw4</a>	<a href="#">Fbxw4.cSep08</a>	<a href="#">309444</a>	102002	1149	7	122	F-box and WD-40 domain protein 4 (Fbxw4) alternative variant cSep08, mRNA.
<a href="#">Fbxw5</a>	<a href="#">Fbxw5.bSep08</a>	<a href="#">362081</a>	4194	1770	9	494	F-box WD-40 domain protein 5 (55.9 kD) (Fbxw5) alternative variant bSep08, mRNA.
<a href="#">Fbxw5</a>	<a href="#">Fbxw5.cSep08</a>	<a href="#">362081</a>	1359	788	4	222	WD-40 repeat (Fbxw5) alternative variant cSep08, mRNA.
<a href="#">Fbxw5</a>	<a href="#">Fbxw5.dSep08</a>	<a href="#">362081</a>	1701	1411	4	198	WD-40 repeat (Fbxw5) alternative variant dSep08, mRNA.
<a href="#">Fbxw5</a>	<a href="#">Fbxw5.eSep08</a>	<a href="#">362081</a>	1957	839	4	156	cyclin-like F-box and WD-40 repeat (18.1 kD) (Fbxw5) alternative variant eSep08, mRNA.
<a href="#">Fbxw5</a>	<a href="#">Fbxw5.fSep08</a>	<a href="#">362081</a>	1007	939	2	150	WD-40 repeat (Fbxw5) alternative variant fSep08, mRNA.
<a href="#">Fbxw8</a>	<a href="#">Fbxw8.aSep08</a>	<a href="#">304522</a>	30058	2441	5	211	F-box and WD-40 domain protein 8 (Fbxw8) alternative variant aSep08, mRNA.
<a href="#">Fbxw8</a>	<a href="#">Fbxw8.bSep08</a>	<a href="#">304522</a>	31818	583	3	151	F-box and WD-40 domain protein 8 (Fbxw8) alternative variant bSep08, mRNA.
<a href="#">Fbxw9</a>	<a href="#">Fbxw9.cSep08</a>	<a href="#">288921</a>	1061	901	3	121	F-box and WD-40 domain protein 9 (Fbxw9) alternative variant cSep08, mRNA.
<a href="#">Fbxw11</a>	<a href="#">Fbxw11.bSep08</a>	<a href="#">303024</a>	19448	3422	9	344	F-box and WD-40 domain protein 11 (Fbxw11) alternative variant bSep08, mRNA.
<a href="#">Fbxw11</a>	<a href="#">Fbxw11.cSep08</a>	<a href="#">303024</a>	76583	517	3	172	F-box and WD-40 domain protein 11 (Fbxw11) alternative variant cSep08, mRNA.
<a href="#">Fcer1a</a>	<a href="#">Fcer1a.aSep08</a>	<a href="#">25047</a>	4338	805		161	fc receptor, IgE, high affinity I, alpha polypeptide (Fcer1a) mRNA.
<a href="#">Fcer1g</a>	<a href="#">Fcer1g.cSep08</a>	<a href="#">25441</a>	3717	715	2	13	fc receptor, IgE, high affinity I, gamma polypeptide (1.5 kD) (Fcer1g) alternative variant cSep08, mRNA.
<a href="#">Fcer1g</a>	<a href="#">Fcer1g.dSep08</a>	<a href="#">25441</a>	4799	576	5	13	fc receptor, IgE, high affinity I, gamma polypeptide (1.5 kD) (Fcer1g) alternative variant dSep08, mRNA.
<a href="#">Fcer2a</a>	<a href="#">Fcer2a.cSep08</a>	<a href="#">171075</a>	1088	765	2	98	fc receptor, IgE, low affinity II, alpha polypeptide (10.7 kD) (Fcer2a) alternative variant cSep08, mRNA.
<a href="#">Fcer2a</a>	<a href="#">Fcer2a.dSep08</a>	<a href="#">171075</a>	936	829	2	37	fc receptor, IgE, low affinity II, alpha polypeptide (4.2 kD) (Fcer2a) alternative variant dSep08, mRNA.
<a href="#">Fcf2.0</a>	<a href="#">Fcf2.0.aSep08</a>		11327	2531	6	772	deoxynucleotidyltransferase terminal interacting protein 2 (Fcf2.0) alternative variant aSep08, mRNA.

<a href="#">Fcf2.0</a>	<a href="#">Fcf2.0.bSep08</a>		1819	1169	1	96	putative nuclear protein human specific (11.4 kD) (Fcf2.0) alternative variant bSep08, mRNA.
<a href="#">Fcgr2b</a>	<a href="#">Fcgr2b.aSep08</a>	<a href="#">289211</a>	14595	1535	8	342	fc receptor, IgG, low affinity IIb (38.0 kD) (Fcgr2b) alternative variant aSep08, complete mRNA.
<a href="#">Fcgr2b</a>	<a href="#">Fcgr2b.cSep08</a>	<a href="#">289211</a>	13821	713	6	237	fc receptor, IgG, low affinity IIb (Fcgr2b) alternative variant cSep08, mRNA.
<a href="#">Fcgr2b</a>	<a href="#">Fcgr2b.dSep08</a>	<a href="#">289211</a>	7497	791	4	196	fc receptor, IgG, low affinity IIb (Fcgr2b) alternative variant dSep08, mRNA.
<a href="#">Fcgr2b</a>	<a href="#">Fcgr2b.eSep08</a>	<a href="#">289211</a>	1445	831	2	109	fc receptor, IgG, low affinity IIb (Fcgr2b) alternative variant eSep08, mRNA.
<a href="#">Fcgr2b</a>	<a href="#">Fcgr2b.fSep08</a>	<a href="#">289211</a>	737	641	2	74	fc receptor, IgG, low affinity IIb (Fcgr2b) alternative variant fSep08, mRNA.
<a href="#">Fcgr3</a>	<a href="#">Fcgr3.aSep08</a>	<a href="#">116591</a>	7516	1067	3	186	fc receptor, IgG, low affinity III (Fcgr3) alternative variant aSep08, mRNA.
<a href="#">Fcgrt</a>	<a href="#">Fcgrt.bSep08</a>	<a href="#">29558</a>	9377	1267	5	274	fc receptor, IgG, alpha chain transporter (29.8 kD) (Fcgrt) alternative variant bSep08, complete mRNA.
<a href="#">Fcgrt</a>	<a href="#">Fcgrt.cSep08</a>	<a href="#">29558</a>	1777	668	3	171	fc receptor, IgG, alpha chain transporter (Fcgrt) alternative variant cSep08, mRNA.
<a href="#">Fcgrt</a>	<a href="#">Fcgrt.dSep08</a>	<a href="#">29558</a>	2390	1591	2	121	fc receptor, IgG, alpha chain transporter (13.7 kD) (Fcgrt) alternative variant dSep08, mRNA.
<a href="#">Fcgrt</a>	<a href="#">Fcgrt.eSep08</a>	<a href="#">29558</a>	7691	1200	3	75	fc receptor, IgG, alpha chain transporter (Fcgrt) alternative variant eSep08, mRNA.
<a href="#">FCH.0</a>	<a href="#">FCH.0.aSep08</a>		37498	418	6	139	proline-serine-threonine phosphatase interacting protein 2 (FCH.0) mRNA.
<a href="#">FCH.1</a>	<a href="#">FCH.1.aSep08</a>		271681	1282	3	148	tyrosine kinase (17.1 kD) (FCH.1) alternative variant aSep08, mRNA.
<a href="#">FCH.2</a>	<a href="#">FCH.2.aSep08</a>		59391	1794		517	FCH domain only 2 (FCH.2) alternative variant aSep08, mRNA.
<a href="#">FCH.2</a>	<a href="#">FCH.2.bSep08</a>		22588	652		217	FCH domain only 2 (FCH.2) alternative variant bSep08, mRNA.
<a href="#">Fcho1</a>	<a href="#">Fcho1.bSep08</a>	<a href="#">290639</a>	4827	2131	12	486	FCH domain only 1 (51.8 kD) (Fcho1) alternative variant bSep08, mRNA.
<a href="#">Fcho1</a>	<a href="#">Fcho1.cSep08</a>	<a href="#">290639</a>	6719	706	8	193	FCH domain only 1 (Fcho1) alternative variant cSep08, mRNA.
<a href="#">Fcho1</a>	<a href="#">Fcho1.dSep08</a>	<a href="#">290639</a>	1632	969	4	165	FCH domain only 1 (Fcho1) alternative variant dSep08, mRNA.
<a href="#">Fcho2</a>	<a href="#">Fcho2.aSep08</a>	<a href="#">309129</a>	30156	1782		347	FCH domain only 2 (Fcho2) alternative variant aSep08, mRNA.
<a href="#">Fcho2</a>	<a href="#">Fcho2.bSep08</a>	<a href="#">309129</a>	6810	524		174	FCH domain only 2 (Fcho2) alternative variant bSep08, mRNA.
<a href="#">Fcho2</a>	<a href="#">Fcho2.cSep08</a>	<a href="#">309129</a>	7519	2585	1	95	FCH domain only 2 (Fcho2) alternative variant cSep08, mRNA.
<a href="#">Fchsd1</a>	<a href="#">Fchsd1.bSep08</a>	<a href="#">307482</a>	3361	709	3	236	FCH and double SH3 domains 1 (Fchsd1) alternative variant bSep08, mRNA.
<a href="#">Fchsd2</a>	<a href="#">Fchsd2.bSep08</a>	<a href="#">308864</a>	142258	1910	7	482	FCH and double SH3 domains 2 (Fchsd2) alternative variant bSep08, mRNA.



<a href="#">Fchsd2</a>	<a href="#">Fchsd2.cSep08</a>	<a href="#">308864</a>	49083	920	9	306	FCH and double SH3 domains 2 (Fchsd2) alternative variant cSep08, mRNA.
<a href="#">Fchsd2</a>	<a href="#">Fchsd2.dSep08</a>	<a href="#">308864</a>	8942	344	2	79	FCH and double SH3 domains 2 (Fchsd2) alternative variant dSep08, mRNA.
<a href="#">Fchsd2</a>	<a href="#">Fchsd2.fSep08</a>	<a href="#">308864</a>	10049	2089	3	73	FCH and double SH3 domains 2 (Fchsd2) alternative variant fSep08, mRNA.
<a href="#">Fcna</a>	<a href="#">Fcna.bSep08</a>	<a href="#">83517</a>	2585	724	6	241	ficolin A (Fcna) alternative variant bSep08, mRNA.
<a href="#">Fcrl1</a>	<a href="#">Fcrl1.aSep08</a>	<a href="#">680665</a>	8510	1370	1	299	fc receptor-like 1 (Fcrl1) alternative variant aSep08, mRNA.
<a href="#">Fcrl1</a>	<a href="#">Fcrl1.bSep08</a>	<a href="#">680665</a>	6437	678	1	160	fc receptor-like 1 (Fcrl1) alternative variant bSep08, mRNA.
<a href="#">Fcrla</a>	<a href="#">Fcrla.aSep08</a>	<a href="#">304965</a>	10149	1285	6	363	fc receptor-like A (Fcrla) alternative variant aSep08, mRNA.
<a href="#">Fcrla</a>	<a href="#">Fcrla.bSep08</a>	<a href="#">304965</a>	4269	802	3	215	fc receptor-like A (Fcrla) alternative variant bSep08, mRNA.
<a href="#">Fcrla</a>	<a href="#">Fcrla.cSep08</a>	<a href="#">304965</a>	4906	446	2	104	fc receptor-like A (11.4 kD) (Fcrla) alternative variant cSep08, mRNA.
<a href="#">Fcrla</a>	<a href="#">Fcrla.dSep08</a>	<a href="#">304965</a>	2097	420	2	89	fc receptor-like A (Fcrla) alternative variant dSep08, mRNA.
<a href="#">Fcrla</a>	<a href="#">Fcrla.eSep08</a>	<a href="#">304965</a>	2613	458	3	87	fc receptor-like A (Fcrla) alternative variant eSep08, mRNA.
<a href="#">Fcrlb</a>	<a href="#">Fcrlb.aSep08</a>	<a href="#">498275</a>	966	814		226	fc receptor-like B (Fcrlb) mRNA.
<a href="#">Fdft1</a>	<a href="#">Fdft1.bSep08</a>	<a href="#">29580</a>	6409	659	3	219	farnesyl diphosphate farnesyl transferase 1 (Fdft1) alternative variant bSep08, mRNA.
<a href="#">Fdft1</a>	<a href="#">Fdft1.cSep08</a>	<a href="#">29580</a>	20913	696	5	168	farnesyl diphosphate farnesyl transferase 1 (Fdft1) alternative variant cSep08, mRNA.
<a href="#">Fdps</a>	<a href="#">Fdps.bSep08</a>	<a href="#">83791</a>	8776	2156	2	259	farnesyl diphosphate synthetase (30.0 kD) (Fdps) alternative variant bSep08, mRNA.
<a href="#">Fdps</a>	<a href="#">Fdps.cSep08</a>	<a href="#">83791</a>	8106	718	3	115	farnesyl diphosphate synthetase (13.0 kD) (Fdps) alternative variant cSep08, mRNA.
<a href="#">Fdx1l</a>	<a href="#">Fdx1l.bSep08</a>	<a href="#">313786</a>	5219	1143	4	153	ferredoxin 1-like (16.8 kD) (Fdx1l) alternative variant bSep08, mRNA.
<a href="#">Fdx1l</a>	<a href="#">Fdx1l.cSep08</a>	<a href="#">313786</a>	4834	633	4	114	ferredoxin 1-like (12.7 kD) (Fdx1l) alternative variant cSep08, mRNA.
<a href="#">Fdx1l</a>	<a href="#">Fdx1l.dSep08</a>	<a href="#">313786</a>	4827	770	4	114	ferredoxin 1-like (12.7 kD) (Fdx1l) alternative variant dSep08, mRNA.
<a href="#">Fdx1l</a>	<a href="#">Fdx1l.eSep08</a>	<a href="#">313786</a>	4870	1782	6	103	ferredoxin 1-like (Fdx1l) alternative variant eSep08, mRNA.
<a href="#">feeby</a>	<a href="#">feeby.aSep08</a>		1598	258		31	putative protein (feeby) mRNA.
<a href="#">feechy</a>	<a href="#">feechy.aSep08</a>		1933	612		28	putative protein (3.2 kD) (feechy) mRNA.
<a href="#">feefer</a>	<a href="#">feefer.aSep08</a>		31719	430		143	leucine rich repeat containing 16A (feefer) mRNA.
<a href="#">feeflo</a>	<a href="#">feeflo.aSep08</a>		5147	604		201	transient receptor potential cation channel subfamily M member 3 (feeflo) mRNA.
<a href="#">feeflu</a>	<a href="#">feeflu.bSep08</a>		1990	611		126	repeat-containing protein (feeflu) alternative variant bSep08, mRNA.
<a href="#">feekee</a>	<a href="#">feekee.aSep08</a>		15750	324		31	putative protein (feekee) mRNA.
<a href="#">feeloy</a>	<a href="#">feeloy.bSep08</a>		3480	419	3	74	CRA b like (8.1 kD) (feeloy) alternative variant bSep08, mRNA.
<a href="#">feemer</a>	<a href="#">feemer.aSep08</a>		793	463	1	121	d11lgp1 like precursor (13.5 kD) (feemer) alternative variant aSep08, mRNA.
<a href="#">feemer</a>	<a href="#">feemer.bSep08</a>		1032	421	2	110	d11lgp1 like (feemer) alternative variant bSep08, mRNA.
<a href="#">feenoy</a>	<a href="#">feenoy.aSep08</a>		11052	428		60	CRA a like (feenoy) mRNA.

<a href="#">feepor</a>	<a href="#">feepor.aSep08</a>		25043	658		190	tetratricopeptide repeat 12 (feepor) mRNA.
<a href="#">feesa</a>	<a href="#">feesa.aSep08</a>		1402	284		48	putative protein (5.9 kD) (feesa) mRNA.
<a href="#">feeshee</a>	<a href="#">feeshee.aSep08</a>		14715	262		87	protein tyrosine phosphatase non-receptor type substrate I like (feeshee) mRNA.
<a href="#">feeto</a>	<a href="#">feeto.aSep08</a>		548	390		109	putative protein (feeto) mRNA.
<a href="#">feevar</a>	<a href="#">feevar.aSep08</a>		1531	620		103	putative protein (feevar) mRNA.
<a href="#">feewey</a>	<a href="#">feewey.aSep08</a>		21024	629	4	55	CRA b like (feewey) alternative variant aSep08, mRNA.
<a href="#">feewey</a>	<a href="#">feewey.bSep08</a>		10493	757	4	43	CRA b like (4.9 kD) (feewey) alternative variant bSep08, complete mRNA.
<a href="#">feewey</a>	<a href="#">feewey.cSep08</a>		20980	1083	3	42	CRA c like (4.8 kD) (feewey) alternative variant cSep08, mRNA.
<a href="#">feewey</a>	<a href="#">feewey.dSep08</a>		17536	827	2	42	CRA c like (4.8 kD) (feewey) alternative variant dSep08, mRNA.
<a href="#">feewey</a>	<a href="#">feewey.eSep08</a>		22013	624	4	10	putative protein (1.2 kD) (feewey) alternative variant eSep08, mRNA.
<a href="#">Feld-I_B.0</a>	<a href="#">Feld-I_B.0.aSep08</a>		1929	494		69	putative protein of mammalian origin (Feld-I_B.0) mRNA.
<a href="#">Fen1</a>	<a href="#">Fen1.bSep08</a>	<a href="#">84490</a>	3058	850	1	205	flap structure-specific endonuclease 1 (Fen1) alternative variant bSep08, mRNA.
<a href="#">Fer1I3</a>	<a href="#">Fer1I3.aSep08</a>	<a href="#">309499</a>	28403	2449	16	616	myoferlin (71.4 kD) (Fer1I3) alternative variant aSep08, mRNA.
<a href="#">Fer1I3</a>	<a href="#">Fer1I3.bSep08</a>	<a href="#">309499</a>	7836	780	5	166	myoferlin (Fer1I3) alternative variant bSep08, mRNA.
<a href="#">Fer1I3</a>	<a href="#">Fer1I3.dSep08</a>	<a href="#">309499</a>	2245	234	3	47	putative protein (5.1 kD) (Fer1I3) alternative variant dSep08, mRNA.
<a href="#">Fer2.0</a>	<a href="#">Fer2.0.aSep08</a>		7816	620		67	aldehyde oxidase (7.4 kD) (Fer2.0) mRNA.
<a href="#">FerB.0</a>	<a href="#">FerB.0.aSep08</a>		9336	399		133	myoferlin (FerB.0) mRNA.
<a href="#">ferby</a>	<a href="#">ferby.aSep08</a>		2057	512		38	putative protein (ferby) mRNA.
<a href="#">ferchy</a>	<a href="#">ferchy.aSep08</a>		1206	822		44	putative protein (5.2 kD) (ferchy) mRNA.
<a href="#">ferfer</a>	<a href="#">ferfer.aSep08</a>		32208	720		238	leucine rich repeat containing 16A (ferfer) mRNA.
<a href="#">ferflo</a>	<a href="#">ferflo.aSep08</a>		6292	854		51	putative protein (6.1 kD) (ferflo) mRNA.
<a href="#">ferflu</a>	<a href="#">ferflu.aSep08</a>		13690	685		108	putative protein (ferflu) mRNA.
<a href="#">ferkee</a>	<a href="#">ferkee.aSep08</a>		3672	598		68	putative protein (ferkee) mRNA.
<a href="#">ferloy</a>	<a href="#">ferloy.aSep08</a>		7737	970		211	solute carrier family 35 member A5 (24.2 kD) (ferloy) mRNA.
<a href="#">fermer</a>	<a href="#">fermer.aSep08</a>		2131	670	1	223	alpha-N-acetylglucosaminidase (fermer) alternative variant aSep08, mRNA.
<a href="#">fermer</a>	<a href="#">fermer.bSep08</a>		1862	466	1	137	alpha-N-acetylglucosaminidase (fermer) alternative variant bSep08, mRNA.
<a href="#">Fermt2</a>	<a href="#">Fermt2.bSep08</a>	<a href="#">289992</a>	11424	1344	7	320	fermitin family homolog 2 (Fermt2) alternative variant bSep08, mRNA.
<a href="#">Fermt2</a>	<a href="#">Fermt2.cSep08</a>	<a href="#">289992</a>	5558	932	5	227	fermitin family homolog 2 (Fermt2) alternative variant cSep08, mRNA.
<a href="#">Fermt2</a>	<a href="#">Fermt2.dSep08</a>	<a href="#">289992</a>	44076	747	3	133	fermitin family homolog 2 (Fermt2) alternative variant dSep08, mRNA.
<a href="#">Fermt2</a>	<a href="#">Fermt2.eSep08</a>	<a href="#">289992</a>	2970	849	3	132	fermitin family homolog 2 (15.5 kD) (Fermt2) alternative variant eSep08, mRNA.

<a href="#">Fermt2</a>	<a href="#">Fermt2.fSep08</a>	<a href="#">289992</a>	568	259	2	82	fermitin family homolog 2 (Fermt2) alternative variant fSep08, mRNA.
<a href="#">Fermt3</a>	<a href="#">Fermt3.cSep08</a>	<a href="#">309186</a>	773	630	2	71	fermitin family homolog 3 (Drosophila) (Fermt3) alternative variant cSep08, mRNA.
<a href="#">FERM_C.0</a>	<a href="#">FERM_C.0.aSep08</a>		16720	1857		489	putative protein of eukaryotic origin (FERM_C.0) alternative variant aSep08, mRNA.
<a href="#">FERM_C.0</a>	<a href="#">FERM_C.0.bSep08</a>		7506	1801		164	putative protein of vertebrate origin (FERM_C.0) alternative variant bSep08, mRNA.
<a href="#">FERM_N.0</a>	<a href="#">FERM_N.0.aSep08</a>		11381	1006		335	protein Tyrosine phosphatase non-receptor type 13 (FERM_N.0) mRNA.
<a href="#">FERM_N.1</a>	<a href="#">FERM_N.1.aSep08</a>		55714	777	5	206	putative protein of bilateral origin (FERM_N.1) alternative variant aSep08, mRNA.
<a href="#">FERM_N.1</a>	<a href="#">FERM_N.1.bSep08</a>		33853	398	1	69	putative protein (FERM_N.1) alternative variant bSep08, mRNA.
<a href="#">fernoy</a>	<a href="#">fernoy.aSep08</a>		27330	4160	18	1145	transmembrane protein 131 (fernoy) alternative variant aSep08, mRNA.
<a href="#">fernoy</a>	<a href="#">fernoy.bSep08</a>		3850	1332	2	298	transmembrane protein 131 (31.1 kD) (fernoy) alternative variant bSep08, mRNA.
<a href="#">fernoy</a>	<a href="#">fernoy.cSep08</a>		4559	715	2	167	transmembrane protein 131 (fernoy) alternative variant cSep08, mRNA.
<a href="#">ferpor</a>	<a href="#">ferpor.aSep08</a>		1440	300		40	putative protein (ferpor) mRNA.
<a href="#">fersa</a>	<a href="#">fersa.aSep08</a>		6313	769		50	putative protein (6.1 kD) (fersa) mRNA.
<a href="#">fershee</a>	<a href="#">fershee.aSep08</a>		636	329		31	putative protein (fershee) mRNA.
<a href="#">Fert2</a>	<a href="#">Fert2.bSep08</a>	<a href="#">301737</a>	46902	398	4	115	fer (fms/fps related) protein kinase, testis specific 2 (Fert2) alternative variant bSep08, mRNA.
<a href="#">Fert2</a>	<a href="#">Fert2.cSep08</a>	<a href="#">301737</a>	66784	374	3	86	fer (fms/fps related) protein kinase, testis specific 2 (Fert2) alternative variant cSep08, mRNA.
<a href="#">ferto</a>	<a href="#">ferto.aSep08</a>		1940	771		133	putative protein of mammalian origin (ferto) mRNA.
<a href="#">fervar</a>	<a href="#">fervar.aSep08</a>		52070	329		92	carboxypeptidase 6 (fervar) mRNA.
<a href="#">ferwey</a>	<a href="#">ferwey.aSep08</a>		18596	543		180	kinase 1 (ferwey) mRNA.
<a href="#">Fes</a>	<a href="#">Fes.bSep08</a>	<a href="#">361597</a>	9160	3235	18	500	feline sarcoma oncogene CRA d (Fes) alternative variant bSep08, mRNA.
<a href="#">Fes</a>	<a href="#">Fes.cSep08</a>	<a href="#">361597</a>	923	556	3	182	feline sarcoma oncogene (Fes) alternative variant cSep08, mRNA.
<a href="#">Fes</a>	<a href="#">Fes.dSep08</a>	<a href="#">361597</a>	1183	919	2	123	feline sarcoma oncogene (13.9 kD) (Fes) alternative variant dSep08, mRNA.
<a href="#">Fes</a>	<a href="#">Fes.eSep08</a>	<a href="#">361597</a>	1471	469	4	110	feline sarcoma oncogene CRA d (Fes) alternative variant eSep08, mRNA.
<a href="#">Fes</a>	<a href="#">Fes.gSep08</a>	<a href="#">361597</a>	1330	401	3	82	feline sarcoma oncogene (Fes) alternative variant gSep08, mRNA.
<a href="#">Fetub</a>	<a href="#">Fetub.bSep08</a>	<a href="#">83928</a>	8003	1141	7	331	fetuin beta (Fetub) alternative variant bSep08, mRNA.
<a href="#">Fetub</a>	<a href="#">Fetub.cSep08</a>	<a href="#">83928</a>	9220	1229	7	281	fetuin beta (Fetub) alternative variant cSep08, mRNA.
<a href="#">Fetub</a>	<a href="#">Fetub.dSep08</a>	<a href="#">83928</a>	7794	976	7	278	fetuin beta (Fetub) alternative variant dSep08, mRNA.
<a href="#">Fetub</a>	<a href="#">Fetub.eSep08</a>	<a href="#">83928</a>	7008	764	5	222	fetuin beta (24.5 kD) (Fetub) alternative variant eSep08, mRNA.
<a href="#">Fetub</a>	<a href="#">Fetub.fSep08</a>	<a href="#">83928</a>	5069	713	4	138	fetuin beta (Fetub) alternative variant fSep08, mRNA.

<a href="#">Fetub</a>	<a href="#">Fetub.hSep08</a>	<a href="#">83928</a>	1481	641	2	53	fetuin beta (5.9 kD) (Fetub) alternative variant hSep08, mRNA.
<a href="#">feyby</a>	<a href="#">feyby.aSep08</a>		20165	580		103	bcl-6 corepressor (feyby) mRNA.
<a href="#">feychy</a>	<a href="#">feychy.aSep08</a>		773	673		34	putative protein (3.8 kD) (feychy) mRNA.
<a href="#">feyfer</a>	<a href="#">feyfer.aSep08</a>		6419	241		80	leucine rich repeat containing 16A (feyfer) mRNA.
<a href="#">feyflo</a>	<a href="#">feyflo.aSep08</a>		10141	342		113	putative protein of bilateral origin (feyflo) mRNA.
<a href="#">feyflu</a>	<a href="#">feyflu.aSep08</a>		16510	871	3	55	putative protein (6.4 kD) (feyflu) alternative variant aSep08, mRNA.
<a href="#">feyflu</a>	<a href="#">feyflu.bSep08</a>		12709	306	2	54	putative protein (feyflu) alternative variant bSep08, mRNA.
<a href="#">feykee</a>	<a href="#">feykee.aSep08</a>		7265	769		256	putative protein of ancient origin (feykee) mRNA.
<a href="#">feyloy</a>	<a href="#">feyloy.aSep08</a>		824	750		25	putative protein (2.9 kD) (feyloy) mRNA.
<a href="#">feymer</a>	<a href="#">feymer.aSep08</a>		1148	376		124	tubulin gamma 1 (feymer) mRNA.
<a href="#">feynoy</a>	<a href="#">feynoy.aSep08</a>		4159	627		149	transmembrane protein 131 (feynoy) mRNA.
<a href="#">feypor</a>	<a href="#">feypor.aSep08</a>		784	730		75	putative cytoplasmic protein (8.9 kD) (feypor) mRNA.
<a href="#">feysa</a>	<a href="#">feysa.aSep08</a>		950	425		124	putative protein (feysa) mRNA.
<a href="#">feyshee</a>	<a href="#">feyshee.aSep08</a>		4627	1326	4	104	putative protein, with a coiled coil domain, of vertebrate origin (feyshee) alternative variant aSep08, mRNA.
<a href="#">feyto</a>	<a href="#">feyto.aSep08</a>		3917	891		86	putative protein (feyto) mRNA.
<a href="#">feyvar</a>	<a href="#">feyvar.aSep08</a>		8504	565		98	putative protein, with a transmembrane domain, of mammalian origin (feyvar) mRNA.
<a href="#">feywey</a>	<a href="#">feywey.aSep08</a>		5524	1725	4	128	kinase 1 (feywey) alternative variant aSep08, mRNA.
<a href="#">Fez1</a>	<a href="#">Fez1.bSep08</a>	<a href="#">81730</a>	9608	610	1	73	fasciculation and elongation protein zeta 1 (zygin I) (Fez1) alternative variant bSep08, mRNA.
<a href="#">Fez2</a>	<a href="#">Fez2.bSep08</a>	<a href="#">94269</a>	23809	747	6	198	fasciculation and elongation protein zeta 2 (zygin II) (Fez2) alternative variant bSep08, mRNA.
<a href="#">Fez2</a>	<a href="#">Fez2.cSep08</a>	<a href="#">94269</a>	24151	776	3	176	fasciculation and elongation protein zeta 2 (zygin II) (Fez2) alternative variant cSep08, mRNA.
<a href="#">Fez2</a>	<a href="#">Fez2.dSep08</a>	<a href="#">94269</a>	24601	1368	5	172	fasciculation and elongation protein zeta 2 (zygin II) (Fez2) alternative variant dSep08, mRNA.
<a href="#">Fez2</a>	<a href="#">Fez2.eSep08</a>	<a href="#">94269</a>	5030	396	2	131	fasciculation and elongation protein zeta 2 (zygin II) (Fez2) alternative variant eSep08, mRNA.
<a href="#">FFD_TFG.0</a>	<a href="#">FFD_TFG.0.aSep08</a>		2214	734		87	lsm14 homolog b CRA e (FFD_TFG.0) mRNA.
<a href="#">FG-GAP.0</a>	<a href="#">FG-GAP.0.aSep08</a>		68113	1541		513	integrin alpha 8 CRA a (FG-GAP.0) mRNA.
<a href="#">Fga</a>	<a href="#">Fga.aSep08</a>	<a href="#">361969</a>	3387	2015	4	604	fibrinogen, alpha polypeptide (Fga) alternative variant aSep08, mRNA.
<a href="#">Fgb</a>	<a href="#">Fgb.bSep08</a>	<a href="#">24366</a>	5356	814	5	271	fibrinogen, B beta polypeptide (Fgb) alternative variant bSep08, mRNA.
<a href="#">Fgd1</a>	<a href="#">Fgd1.bSep08</a>	<a href="#">363460</a>	18530	1785	1	390	DH (Fgd1) alternative variant bSep08, mRNA.
<a href="#">Fgd2</a>	<a href="#">Fgd2.bSep08</a>	<a href="#">309653</a>	13426	2357	14	319	pleckstrin-like and zinc finger, FYVE-type (36.3 kD) (Fgd2) alternative variant bSep08, mRNA.
<a href="#">Fgd2</a>	<a href="#">Fgd2.cSep08</a>	<a href="#">309653</a>	7449	711	6	196	zinc finger, FYVE-type (Fgd2) alternative variant cSep08, mRNA.
<a href="#">Fgd2</a>	<a href="#">Fgd2.dSep08</a>	<a href="#">309653</a>	2186	691	3	96	putative protein (Fgd2) alternative variant dSep08, mRNA.
<a href="#">Fgd3</a>	<a href="#">Fgd3.bSep08</a>	<a href="#">361223</a>	10163	417	4	139	zinc finger, FYVE-type (Fgd3) alternative variant bSep08, mRNA.

<a href="#">Fgd4</a>	<a href="#">Fgd4.bSep08</a>	<a href="#">246174</a>	24108	508		71	putative protein of mammalian origin (Fgd4) alternative variant bSep08, mRNA.
<a href="#">Fgd5</a>	<a href="#">Fgd5.bSep08</a>	<a href="#">362402</a>	3771	369	5	122	putative protein of eukaryotic origin (Fgd5) alternative variant bSep08, mRNA.
<a href="#">Fgd5</a>	<a href="#">Fgd5.cSep08</a>	<a href="#">362402</a>	4269	801	3	72	putative protein of bilateral origin (Fgd5) alternative variant cSep08, mRNA.
<a href="#">Fgd6</a>	<a href="#">Fgd6.aSep08</a>	<a href="#">500824</a>	3253	533	3	104	pleckstrin-like (Fgd6) alternative variant aSep08, mRNA.
<a href="#">Fgf1</a>	<a href="#">Fgf1.aSep08</a>	<a href="#">25317</a>	73472	989	5	155	fibroblast growth factor 1 (17.4 kD) (Fgf1) alternative variant aSep08, mRNA.
<a href="#">Fgf1</a>	<a href="#">Fgf1.cSep08</a>	<a href="#">25317</a>	57080	385	2	68	fibroblast growth factor 1 (Fgf1) alternative variant cSep08, mRNA.
<a href="#">Fgf7</a>	<a href="#">Fgf7.bSep08</a>	<a href="#">29348</a>	2820	1827	1	25	fibroblast growth factor 7 (Fgf7) alternative variant bSep08, mRNA.
<a href="#">Fgf9</a>	<a href="#">Fgf9.bSep08</a>	<a href="#">25444</a>	38212	448	2	129	fibroblast growth factor 9 (Fgf9) alternative variant bSep08, mRNA.
<a href="#">Fgf9</a>	<a href="#">Fgf9.cSep08</a>	<a href="#">25444</a>	37549	624	2	126	fibroblast growth factor 9 (Fgf9) alternative variant cSep08, mRNA.
<a href="#">Fgf9</a>	<a href="#">Fgf9.dSep08</a>	<a href="#">25444</a>	37556	1545	2	91	fibroblast growth factor 9 (10.9 kD) (Fgf9) alternative variant dSep08, mRNA.
<a href="#">Fgf12</a>	<a href="#">Fgf12.bSep08</a>	<a href="#">170630</a>	577082	1299	6	181	fibroblast growth factor 12 (20.4 kD) (Fgf12) alternative variant bSep08, mRNA.
<a href="#">Fgf13</a>	<a href="#">Fgf13.bSep08</a>	<a href="#">84488</a>	294827	790	1	185	fibroblast growth factor 13 (Fgf13) alternative variant bSep08, mRNA.
<a href="#">Fgf21</a>	<a href="#">Fgf21.bSep08</a>	<a href="#">170580</a>	4879	720	3	106	fibroblast growth factor 21 (11.8 kD) (Fgf21) alternative variant bSep08, mRNA.
<a href="#">Fgfr1</a>	<a href="#">Fgfr1.aSep08</a>	<a href="#">79114</a>	55088	3802	10	822	fibroblast growth factor receptor 1 (91.9 kD) (Fgfr1) alternative variant aSep08, mRNA.
<a href="#">Fgfr1</a>	<a href="#">Fgfr1.bSep08</a>	<a href="#">79114</a>	9070	1783	3	434	fibroblast growth factor receptor 1 (Fgfr1) alternative variant bSep08, mRNA.
<a href="#">Fgfr1</a>	<a href="#">Fgfr1.cSep08</a>	<a href="#">79114</a>	4425	701	2	182	fibroblast growth factor receptor 1 (Fgfr1) alternative variant cSep08, mRNA.
<a href="#">Fgfr1</a>	<a href="#">Fgfr1.dSep08</a>	<a href="#">79114</a>	43932	1255	6	180	fibroblast growth factor receptor 1 (Fgfr1) alternative variant dSep08, mRNA.
<a href="#">Fgfr1op2</a>	<a href="#">Fgfr1op2.bSep08</a>	<a href="#">362463</a>	19320	1110	6	215	FGFR1 oncogene partner 2 (24.9 kD) (Fgfr1op2) alternative variant bSep08, mRNA.
<a href="#">Fgfr1op2</a>	<a href="#">Fgfr1op2.cSep08</a>	<a href="#">362463</a>	2874	330	2	75	FGFR1 oncogene partner 2 (Fgfr1op2) alternative variant cSep08, mRNA.
<a href="#">Fgfr1op2</a>	<a href="#">Fgfr1op2.dSep08</a>	<a href="#">362463</a>	1475	358	2	69	FGFR1 oncogene partner 2 (Fgfr1op2) alternative variant dSep08, mRNA.
<a href="#">Fgfr2</a>	<a href="#">Fgfr2.dSep08</a>	<a href="#">25022</a>	8977	607	5	157	fibroblast growth factor receptor 2 (Fgfr2) alternative variant dSep08, mRNA.
<a href="#">Fgfr2</a>	<a href="#">Fgfr2.eSep08</a>	<a href="#">25022</a>	10371	375	3	77	fibroblast growth factor receptor 2 (Fgfr2) alternative variant eSep08, mRNA.
<a href="#">Fgfr3</a>	<a href="#">Fgfr3.bSep08</a>	<a href="#">84489</a>	975	691	3	184	fibroblast growth factor receptor 3 (Fgfr3) alternative variant bSep08, mRNA.
<a href="#">Fgfr3</a>	<a href="#">Fgfr3.cSep08</a>	<a href="#">84489</a>	2468	2162	3	184	fibroblast growth factor receptor 3 (20.4 kD) (Fgfr3) alternative variant cSep08, mRNA.

<a href="#">Fgfr4</a>	<a href="#">Fgfr4.bSep08</a>	<a href="#">25114</a>	2765	1298	5	237	fibroblast growth factor receptor 4 (26.4 kD) (Fgfr4) alternative variant bSep08, mRNA.
<a href="#">Fgfr4</a>	<a href="#">Fgfr4.cSep08</a>	<a href="#">25114</a>	512	396	2	81	fibroblast growth factor receptor 4 (9.0 kD) (Fgfr4) alternative variant cSep08, mRNA.
<a href="#">Fgfr1</a>	<a href="#">Fgfr1.bSep08</a>	<a href="#">360903</a>	977	738	1	245	fibroblast growth factor receptor-like 1 (Fgfr1) alternative variant bSep08, mRNA.
<a href="#">Fgg</a>	<a href="#">Fgg.aSep08</a>	<a href="#">24367</a>	7281	2056	3	444	fibrinogen, gamma polypeptide (50.5 kD) (Fgg) alternative variant aSep08, complete mRNA.
<a href="#">Fgg</a>	<a href="#">Fgg.cSep08</a>	<a href="#">24367</a>	2596	350	2	116	fibrinogen, gamma polypeptide (Fgg) alternative variant cSep08, mRNA.
<a href="#">Fgr</a>	<a href="#">Fgr.bSep08</a>	<a href="#">79113</a>	12446	803	2	198	gardner-Rasheed feline sarcoma viral (Fgr) oncogene homolog (Fgr) alternative variant bSep08, mRNA.
<a href="#">Fgr</a>	<a href="#">Fgr.cSep08</a>	<a href="#">79113</a>	21648	650	1	165	gardner-Rasheed feline sarcoma viral (Fgr) oncogene homolog (Fgr) alternative variant cSep08, mRNA.
<a href="#">FH2.0</a>	<a href="#">FH2.0.aSep08</a>		2771	1598	6	446	actin-binding FH2 (FH2.0) alternative variant aSep08, mRNA.
<a href="#">FH2.0</a>	<a href="#">FH2.0.bSep08</a>		3170	1504	7	428	actin-binding FH2 (FH2.0) alternative variant bSep08, mRNA.
<a href="#">FH2.0</a>	<a href="#">FH2.0.cSep08</a>		1361	529	3	145	putative protein, with a coiled coil domain, of eukaryotic origin (FH2.0) alternative variant cSep08, mRNA.
<a href="#">FH2.1</a>	<a href="#">FH2.1.aSep08</a>		27206	952		317	actin-binding FH2 (FH2.1) mRNA.
<a href="#">FH2.2</a>	<a href="#">FH2.2.aSep08</a>		2175	636		211	diaphanous 1 (FH2.2) mRNA.
<a href="#">FH2.3</a>	<a href="#">FH2.3.aSep08</a>		2238	1068		355	formin-like 3 CRA a (FH2.3) mRNA.
<a href="#">FH2.4</a>	<a href="#">FH2.4.aSep08</a>		5842	1728		575	inverted (FH2.4) mRNA.
<a href="#">FHA.0</a>	<a href="#">FHA.0.aSep08</a>		3924	738	3	245	mediator of DNA damage checkpoint 1 (FHA.0) alternative variant aSep08, mRNA.
<a href="#">FHA.0</a>	<a href="#">FHA.0.bSep08</a>		2479	331	1	45	mediator of DNA damage checkpoint 1 (FHA.0) alternative variant bSep08, mRNA.
<a href="#">FHA.1</a>	<a href="#">FHA.1.aSep08</a>		50688	544		73	centrosomal protein 170kDa (FHA.1) mRNA.
<a href="#">Fhdc1</a>	<a href="#">Fhdc1.bSep08</a>	<a href="#">295161</a>	11853	2899		823	putative protein, with 2 coiled coil domains, of eukaryotic origin (Fhdc1) alternative variant bSep08, mRNA.
<a href="#">Fhit</a>	<a href="#">Fhit.aSep08</a>	<a href="#">60398</a>	30210	414		34	fragile histidine triad gene (Fhit) alternative variant aSep08, mRNA.
<a href="#">Fhit</a>	<a href="#">Fhit.bSep08</a>	<a href="#">60398</a>	846925	1803		6	fragile histidine triad gene (0.8 kD) (Fhit) alternative variant bSep08, mRNA.
<a href="#">Fhl1</a>	<a href="#">Fhl1.aSep08</a>	<a href="#">25177</a>	57705	1046	8	310	four and a half LIM domains 1 (Fhl1) alternative variant aSep08, mRNA.
<a href="#">Fhl1</a>	<a href="#">Fhl1.bSep08</a>	<a href="#">25177</a>	5258	926	6	305	four and a half LIM domains 1 (Fhl1) alternative variant bSep08, mRNA.
<a href="#">Fhl1</a>	<a href="#">Fhl1.eSep08</a>	<a href="#">25177</a>	55648	691	5	167	four and a half LIM domains 1 (Fhl1) alternative variant eSep08, mRNA.
<a href="#">Fhl2</a>	<a href="#">Fhl2.bSep08</a>	<a href="#">63839</a>	72683	1404	6	279	four and a half LIM domains 2 (32.1 kD) (Fhl2) alternative variant bSep08, complete mRNA.
<a href="#">Fhl2</a>	<a href="#">Fhl2.cSep08</a>	<a href="#">63839</a>	4700	879	2	129	four and a half LIM domains 2 (14.6 kD) (Fhl2) alternative variant cSep08, mRNA.

<a href="#">Fhod1</a>	<a href="#">Fhod1.aSep08</a>	<a href="#">291964</a>	11053	855	7	284	putative protein of metazoan origin (Fhod1) alternative variant aSep08, mRNA.
<a href="#">Fhod1</a>	<a href="#">Fhod1.bSep08</a>	<a href="#">291964</a>	1040	694	4	230	putative protein, with a coiled coil domain, of metazoan origin (Fhod1) alternative variant bSep08, mRNA.
<a href="#">Fibp</a>	<a href="#">Fibp.bSep08</a>	<a href="#">282837</a>	4191	1099	4	346	fibroblast growth factor (acidic) intracellular binding protein (Fibp) alternative variant bSep08, mRNA.
<a href="#">Fibp</a>	<a href="#">Fibp.cSep08</a>	<a href="#">282837</a>	1142	619	2	83	fibroblast growth factor (acidic) intracellular binding protein (9.7 kD) (Fibp) alternative variant cSep08, mRNA.
<a href="#">Fibp</a>	<a href="#">Fibp.dSep08</a>	<a href="#">282837</a>	1178	379	3	82	fibroblast growth factor (acidic) intracellular binding protein (Fibp) alternative variant dSep08, mRNA.
<a href="#">Fibrinogen_C.0</a>	<a href="#">Fibrinogen_C.0.aSep08</a>		1499	405		134	angiopoietin-like 7 (Fibrinogen_C.0) mRNA.
<a href="#">Ficd</a>	<a href="#">Ficd.aSep08</a>	<a href="#">288741</a>	4863	3188	3	458	filamentation induced by cAMP protein Fic precursor (51.8 kD) (Ficd) alternative variant aSep08, mRNA.
<a href="#">Fig4</a>	<a href="#">Fig4.bSep08</a>	<a href="#">309855</a>	58670	1784	6	463	FIG4 homolog (S. cerevisiae) (Fig4) alternative variant bSep08, mRNA.
<a href="#">Fig4</a>	<a href="#">Fig4.cSep08</a>	<a href="#">309855</a>	13760	684	5	125	FIG4 homolog (S. cerevisiae) (Fig4) alternative variant cSep08, mRNA.
<a href="#">Fig4</a>	<a href="#">Fig4.dSep08</a>	<a href="#">309855</a>	20734	433	5	73	FIG4 homolog (S. cerevisiae) (Fig4) alternative variant dSep08, mRNA.
<a href="#">Filamin.0</a>	<a href="#">Filamin.0.aSep08</a>		782	427		142	filamin Gamma (Filamin.0) mRNA.
<a href="#">Filip1</a>	<a href="#">Filip1.bSep08</a>	<a href="#">246776</a>	41315	934	2	82	filamin A interacting protein 1 (Filip1) alternative variant bSep08, mRNA.
<a href="#">Filip1</a>	<a href="#">Filip1.cSep08</a>	<a href="#">246776</a>	24674	230	2	24	filamin A interacting protein 1 (Filip1) alternative variant cSep08, mRNA.
<a href="#">Fip111</a>	<a href="#">Fip111.bSep08</a>	<a href="#">289582</a>	53464	1541	15	471	FIP1 like 1 (S. cerevisiae) (Fip111) alternative variant bSep08, mRNA.
<a href="#">Fip111</a>	<a href="#">Fip111.cSep08</a>	<a href="#">289582</a>	39090	761	9	253	FIP1 like 1 (S. cerevisiae) (Fip111) alternative variant cSep08, mRNA.
<a href="#">Fip111</a>	<a href="#">Fip111.dSep08</a>	<a href="#">289582</a>	37343	735	7	244	FIP1 like 1 (S. cerevisiae) (Fip111) alternative variant dSep08, mRNA.
<a href="#">Fip111</a>	<a href="#">Fip111.eSep08</a>	<a href="#">289582</a>	10990	650	7	177	FIP1 like 1 (S. cerevisiae) (Fip111) alternative variant eSep08, mRNA.
<a href="#">Fis1</a>	<a href="#">Fis1.aSep08</a>	<a href="#">288584</a>	4105	780	5	185	fission 1 (mitochondrial outer membrane) homolog (yeast) (Fis1) alternative variant aSep08, mRNA.
<a href="#">Fis1</a>	<a href="#">Fis1.bSep08</a>	<a href="#">288584</a>	14936	1188	4	179	fission 1 (mitochondrial outer membrane) homolog (yeast) (20.2 kD) (Fis1) alternative variant bSep08, mRNA.
<a href="#">Fiz1</a>	<a href="#">Fiz1.bSep08</a>	<a href="#">292584</a>	1982	625	2	176	FLT3-interacting zinc finger 1 (Fiz1) alternative variant bSep08, mRNA.
<a href="#">Fkbp1a</a>	<a href="#">Fkbp1a.bSep08</a>	<a href="#">25639</a>	3398	1218	2	75	FK506 binding protein 1a (Fkbp1a) alternative variant bSep08, mRNA.
<a href="#">Fkbp2</a>	<a href="#">Fkbp2.dSep08</a>	<a href="#">293702</a>	1346	402	4	55	FK506 binding protein 2 (6.2 kD) (Fkbp2) alternative variant dSep08, mRNA.
<a href="#">Fkbp2</a>	<a href="#">Fkbp2.eSep08</a>	<a href="#">293702</a>	731	359	3	47	FK506 binding protein 2 (5.2 kD) (Fkbp2) alternative variant eSep08, mRNA.
<a href="#">Fkbp3</a>	<a href="#">Fkbp3.aSep08</a>	<a href="#">299104</a>	11536	956	7	224	FK506 binding protein 3 (25.2 kD) (Fkbp3) alternative variant aSep08, complete mRNA.

<a href="#">Fkbp4</a>	<a href="#">Fkbp4.aSep08</a>	<a href="#">260321</a>	8400	2790	2	512	FK506 binding protein 4 (Fkbp4) alternative variant aSep08, mRNA.
<a href="#">Fkbp4</a>	<a href="#">Fkbp4.bSep08</a>	<a href="#">260321</a>	2234	713	1	148	FK506 binding protein 4 (16.6 kD) (Fkbp4) alternative variant bSep08, mRNA.
<a href="#">Fkbp7</a>	<a href="#">Fkbp7.aSep08</a>	<a href="#">295672</a>	11683	869	4	241	FK506 binding protein 7 (Fkbp7) alternative variant aSep08, mRNA.
<a href="#">Fkbp7</a>	<a href="#">Fkbp7.bSep08</a>	<a href="#">295672</a>	11572	761	4	206	FK506 binding protein 7 (Fkbp7) alternative variant bSep08, mRNA.
<a href="#">Fkbp7</a>	<a href="#">Fkbp7.fSep08</a>	<a href="#">295672</a>	1436	752	2	41	FK506 binding protein 7 (4.9 kD) (Fkbp7) alternative variant fSep08, mRNA.
<a href="#">Fkbp8</a>	<a href="#">Fkbp8.bSep08</a>	<a href="#">290652</a>	4438	991	7	330	FK506 binding protein 8 (Fkbp8) alternative variant bSep08, mRNA.
<a href="#">Fkbp8</a>	<a href="#">Fkbp8.cSep08</a>	<a href="#">290652</a>	4543	1706	5	292	FK506 binding protein 8 (31.1 kD) (Fkbp8) alternative variant cSep08, mRNA.
<a href="#">Fkbp8</a>	<a href="#">Fkbp8.dSep08</a>	<a href="#">290652</a>	2791	803	5	212	FK506 binding protein 8 (Fkbp8) alternative variant dSep08, mRNA.
<a href="#">Fkbp8</a>	<a href="#">Fkbp8.eSep08</a>	<a href="#">290652</a>	2784	1586	2	161	FK506 binding protein 8 (16.9 kD) (Fkbp8) alternative variant eSep08, mRNA.
<a href="#">Fkbp8</a>	<a href="#">Fkbp8.fSep08</a>	<a href="#">290652</a>	1021	763	2	91	FK506 binding protein 8 (Fkbp8) alternative variant fSep08, mRNA.
<a href="#">Fkbp9</a>	<a href="#">Fkbp9.bSep08</a>	<a href="#">297123</a>	46266	843	3	280	FK506 binding protein 9 (Fkbp9) alternative variant bSep08, mRNA.
<a href="#">Fkbp9</a>	<a href="#">Fkbp9.cSep08</a>	<a href="#">297123</a>	24165	855	2	202	FK506 binding protein 9 (Fkbp9) alternative variant cSep08, mRNA.
<a href="#">Fkbp10</a>	<a href="#">Fkbp10.bSep08</a>	<a href="#">360627</a>	11999	389	2	73	FK506 binding protein 10 (Fkbp10) alternative variant bSep08, mRNA.
<a href="#">Fkbp10</a>	<a href="#">Fkbp10.cSep08</a>	<a href="#">360627</a>	518	312	2	42	FK506 binding protein 10 (Fkbp10) alternative variant cSep08, mRNA.
<a href="#">Fkbp14</a>	<a href="#">Fkbp14.bSep08</a>	<a href="#">362366</a>	3514	747	2	101	FK506 binding protein 14 (Fkbp14) alternative variant bSep08, mRNA.
<a href="#">Fkbp15</a>	<a href="#">Fkbp15.aSep08</a>	<a href="#">362528</a>	32240	1051	7	350	fk506 binding protein 15 like (Fkbp15) alternative variant aSep08, mRNA.
<a href="#">Fkbp15</a>	<a href="#">Fkbp15.bSep08</a>	<a href="#">362528</a>	42931	1279	3	222	putative nuclear protein of mammalian origin (23.8 kD) (Fkbp15) alternative variant bSep08, mRNA.
<a href="#">Fkbp15</a>	<a href="#">Fkbp15.cSep08</a>	<a href="#">362528</a>	3794	1009	3	159	putative protein of mammalian origin (Fkbp15) alternative variant cSep08, mRNA.
<a href="#">Fkbp15</a>	<a href="#">Fkbp15.dSep08</a>	<a href="#">362528</a>	29034	815	6	128	fk506 binding protein 15 like (Fkbp15) alternative variant dSep08, mRNA.
<a href="#">Fktn</a>	<a href="#">Fktn.bSep08</a>	<a href="#">362520</a>	15281	536	1	178	fukutin (Fktn) alternative variant bSep08, mRNA.
<a href="#">flabor</a>	<a href="#">flabor.aSep08</a>		5017	1138		47	putative protein (flabor) mRNA.
<a href="#">flachy</a>	<a href="#">flachy.aSep08</a>		2369	727		72	putative nuclear protein (8.0 kD) (flachy) mRNA.
<a href="#">fladoy</a>	<a href="#">fladoy.aSep08</a>		1954	563		83	putative protein (fladoy) mRNA.
<a href="#">flafu</a>	<a href="#">flafu.aSep08</a>		2370	410		136	PI-3-kinase-related kinase smg-1 (flafu) mRNA.
<a href="#">flafly</a>	<a href="#">flafly.aSep08</a>		35041	394		34	putative protein (flafly) mRNA.
<a href="#">flagar</a>	<a href="#">flagar.aSep08</a>		793	230		30	putative protein (flagar) mRNA.
<a href="#">flaja</a>	<a href="#">flaja.aSep08</a>		1317	727		167	putative protein (18.8 kD) (flaja) mRNA.



<a href="#">flajey</a>	<a href="#">flajey.aSep08</a>		727	378		50	putative protein (flajey) mRNA.
<a href="#">flakee</a>	<a href="#">flakee.aSep08</a>		23340	1825		608	centromere protein F (flakee) alternative variant aSep08, mRNA.
<a href="#">flakee</a>	<a href="#">flakee.bSep08</a>		10685	586		195	centromere protein F (flakee) alternative variant bSep08, mRNA.
<a href="#">flalo</a>	<a href="#">flalo.aSep08</a>		1852	613	6	204	phosphatidylinositol transfer protein membrane-associated 2 CRA b (flalo) alternative variant aSep08, mRNA.
<a href="#">flamee</a>	<a href="#">flamee.aSep08</a>		1901	1063		42	putative protein (4.4 kD) (flamee) mRNA.
<a href="#">flanoy</a>	<a href="#">flanoy.aSep08</a>		5496	968		322	elastin microfibril interfacier 2 (flanoy) mRNA.
<a href="#">flapor</a>	<a href="#">flapor.bSep08</a>		1071	487	2	88	putative protein (flapor) alternative variant bSep08, mRNA.
<a href="#">flarbor</a>	<a href="#">flarbor.aSep08</a>		3544	302		21	putative protein (2.6 kD) (flarbor) mRNA.
<a href="#">flarchy</a>	<a href="#">flarchy.aSep08</a>		1550	523		40	putative protein (flarchy) mRNA.
<a href="#">flardoy</a>	<a href="#">flardoy.aSep08</a>		21618	392		130	dmx-like 1 (flardoy) mRNA.
<a href="#">flarflu</a>	<a href="#">flarflu.aSep08</a>		2342	635		96	PI-3-kinase-related kinase smg-1 (flarflu) mRNA.
<a href="#">flarfly</a>	<a href="#">flarfly.aSep08</a>		1192	622		70	neural cell expressed developmentally gene 4 (7.6 kD) (flarfly) mRNA.
<a href="#">flargar</a>	<a href="#">flargar.aSep08</a>		7268	377		36	putative protein (flargar) mRNA.
<a href="#">flarja</a>	<a href="#">flarja.aSep08</a>		3763	335		63	putative protein (6.7 kD) (flarja) mRNA.
<a href="#">flarjey</a>	<a href="#">flarjey.aSep08</a>		3512	755		71	putative nuclear protein (8.0 kD) (flarjey) mRNA.
<a href="#">flarkee</a>	<a href="#">flarkee.aSep08</a>		3542	405		134	angel homolog 2 CRA b (flarkee) mRNA.
<a href="#">flarlo</a>	<a href="#">flarlo.aSep08</a>		2146	310		102	vacuolar sorting protein 33A (flarlo) mRNA.
<a href="#">flarmee</a>	<a href="#">flarmee.aSep08</a>		10343	763		84	putative protein (flarmee) mRNA.
<a href="#">flaroy</a>	<a href="#">flaroy.aSep08</a>		732	411		67	putative protein (flaroy) mRNA.
<a href="#">flarpor</a>	<a href="#">flarpor.aSep08</a>		14869	404		124	protein tyrosine phosphatase non-receptor type 23 CRA a (flarpor) mRNA.
<a href="#">flarroy</a>	<a href="#">flarroy.aSep08</a>		6421	2100	14	650	ATP-binding cassette sub-family A member 7 like (flarroy) alternative variant aSep08, mRNA.
<a href="#">flarroy</a>	<a href="#">flarroy.bSep08</a>		1120	1017	1	168	member 7 (flarroy) alternative variant bSep08, mRNA.
<a href="#">flarsa</a>	<a href="#">flarsa.aSep08</a>		3992	1755	3	22	putative protein (flarsa) alternative variant aSep08, mRNA.
<a href="#">flartu</a>	<a href="#">flartu.aSep08</a>		2333	373		107	serologically defined colon cancer antigen 1 CRA f like (flartu) mRNA.
<a href="#">flarvo</a>	<a href="#">flarvo.bSep08</a>		1581	1167		71	putative mitochondrial protein (8.1 kD) (flarvo) alternative variant bSep08, mRNA.
<a href="#">flarwer</a>	<a href="#">flarwer.aSep08</a>		19951	794		264	CRA b (flarwer) mRNA.
<a href="#">flasa</a>	<a href="#">flasa.aSep08</a>		1398	855	1	67	putative protein (flasa) alternative variant aSep08, mRNA.
<a href="#">flasa</a>	<a href="#">flasa.bSep08</a>		9781	576	3	60	putative protein (6.9 kD) (flasa) alternative variant bSep08, mRNA.
<a href="#">flashee</a>	<a href="#">flashee.aSep08</a>		3593	1088	2	74	putative protein (8.5 kD) (flashee) alternative variant aSep08, mRNA.
<a href="#">flashee</a>	<a href="#">flashee.bSep08</a>		2002	738	1	58	putative protein (6.7 kD) (flashee) alternative variant bSep08, mRNA.
<a href="#">flatu</a>	<a href="#">flatu.aSep08</a>		2375	373		123	fam179b (flatu) mRNA.
<a href="#">flavo</a>	<a href="#">flavo.aSep08</a>		11900	556		89	putative protein (9.8 kD) (flavo) mRNA.
<a href="#">flawbor</a>	<a href="#">flawbor.aSep08</a>		65011	654		73	putative protein (flawbor) mRNA.

<a href="#">flawchy</a>	<a href="#">flawchy.bSep08</a>		3489	424	4	33	putative protein (3.8 kD) (flawchy) alternative variant bSep08, mRNA.
<a href="#">flawdoy</a>	<a href="#">flawdoy.aSep08</a>		5166	1897		537	dmx-like 1 (flawdoy) mRNA.
<a href="#">flower</a>	<a href="#">flower.aSep08</a>		818	529	1	87	putative protein (flower) alternative variant aSep08, mRNA.
<a href="#">flower</a>	<a href="#">flower.bSep08</a>		1560	510	2	49	putative protein (flower) alternative variant bSep08, mRNA.
<a href="#">flawflu</a>	<a href="#">flawflu.aSep08</a>		792	683		67	PI-3-kinase-related kinase smg-1 (flawflu) mRNA.
<a href="#">flawfly</a>	<a href="#">flawfly.aSep08</a>		11558	386		116	serine active site containing 1 (flawfly) mRNA.
<a href="#">flawgar</a>	<a href="#">flawgar.aSep08</a>		22241	728		156	putative protein (flawgar) mRNA.
<a href="#">flawja</a>	<a href="#">flawja.bSep08</a>		931	333	3	71	putative protein (flawja) alternative variant bSep08, mRNA.
<a href="#">flawjey</a>	<a href="#">flawjey.aSep08</a>		1049	783	2	64	putative protein (flawjey) alternative variant aSep08, mRNA.
<a href="#">flawjey</a>	<a href="#">flawjey.bSep08</a>		2990	605	4	61	putative protein (flawjey) alternative variant bSep08, mRNA.
<a href="#">flawkee</a>	<a href="#">flawkee.aSep08</a>		2281	576		38	putative protein (4.7 kD) (flawkee) mRNA.
<a href="#">flawlo</a>	<a href="#">flawlo.aSep08</a>		3787	493		150	leucine rich repeat containing 43 CRA b (flawlo) mRNA.
<a href="#">flawmee</a>	<a href="#">flawmee.aSep08</a>		16938	1542		50	putative protein (5.4 kD) (flawmee) alternative variant aSep08, mRNA.
<a href="#">flawpor</a>	<a href="#">flawpor.aSep08</a>		975	618		108	CRA c like (flawpor) mRNA.
<a href="#">flawroy</a>	<a href="#">flawroy.aSep08</a>		916	468	2	84	putative mitochondrial protein (9.1 kD) (flawroy) alternative variant aSep08, mRNA.
<a href="#">flawroy</a>	<a href="#">flawroy.bSep08</a>		1059	459	2	41	putative protein (flawroy) alternative variant bSep08, mRNA.
<a href="#">flawsa</a>	<a href="#">flawsa.aSep08</a>		456	410		39	putative protein (4.4 kD) (flawsa) mRNA.
<a href="#">flawshee</a>	<a href="#">flawshee.aSep08</a>		655	247		22	putative protein (flawshee) mRNA.
<a href="#">flawtu</a>	<a href="#">flawtu.aSep08</a>		24409	2727	8	451	son of sevenless 2 (flawtu) alternative variant aSep08, mRNA.
<a href="#">flawtu</a>	<a href="#">flawtu.bSep08</a>		3985	931	3	213	son of sevenless homolog 2 (flawtu) alternative variant bSep08, mRNA.
<a href="#">flawtu</a>	<a href="#">flawtu.cSep08</a>		1884	928	2	78	putative protein (flawtu) alternative variant cSep08, mRNA.
<a href="#">flawtu</a>	<a href="#">flawtu.dSep08</a>		5024	376	4	64	son of sevenless homolog 2 (flawtu) alternative variant dSep08, mRNA.
<a href="#">flawvo</a>	<a href="#">flawvo.aSep08</a>		43052	443		44	putative protein (4.9 kD) (flawvo) mRNA.
<a href="#">flawwer</a>	<a href="#">flawwer.aSep08</a>		5389	741		211	CRA b (flawwer) mRNA.
<a href="#">Flcn</a>	<a href="#">Flcn.bSep08</a>	<a href="#">303185</a>	2742	686	3	206	folliculin (Flcn) alternative variant bSep08, mRNA.
<a href="#">Flcn</a>	<a href="#">Flcn.cSep08</a>	<a href="#">303185</a>	9020	962	5	189	folliculin (Flcn) alternative variant cSep08, mRNA.
<a href="#">Flcn</a>	<a href="#">Flcn.dSep08</a>	<a href="#">303185</a>	8824	806	5	173	folliculin (Flcn) alternative variant dSep08, mRNA.
<a href="#">Flcn</a>	<a href="#">Flcn.eSep08</a>	<a href="#">303185</a>	3818	1260	4	167	folliculin (18.0 kD) (Flcn) alternative variant eSep08, mRNA.
<a href="#">Flcn</a>	<a href="#">Flcn.fSep08</a>	<a href="#">303185</a>	3437	422	2	14	folliculin (Flcn) alternative variant fSep08, mRNA.
<a href="#">fleebor</a>	<a href="#">fleebor.aSep08</a>		989	379		26	putative protein (3.0 kD) (fleebor) mRNA.
<a href="#">fleechy</a>	<a href="#">fleechy.aSep08</a>		44050	677		56	putative protein (6.5 kD) (fleechy) mRNA.
<a href="#">fleadoy</a>	<a href="#">fleadoy.aSep08</a>		22521	884		294	dmx-like 1 (fleadoy) mRNA.
<a href="#">fleeflu</a>	<a href="#">fleeflu.aSep08</a>		937	141		47	kinase (fleeflu) mRNA.

<a href="#">fleefly</a>	<a href="#">fleefly.aSep08</a>		26292	381		24	putative protein (2.3 kD) (fleefly) mRNA.
<a href="#">fleegar</a>	<a href="#">fleegar.aSep08</a>		2057	1050		105	putative protein (fleegar) mRNA.
<a href="#">fleeja</a>	<a href="#">fleeja.aSep08</a>		4116	534		54	putative protein (fleeja) mRNA.
<a href="#">fleejey</a>	<a href="#">fleejey.aSep08</a>		3613	499	3	141	putative protein of mammalian origin (fleejey) alternative variant aSep08, mRNA.
<a href="#">fleekee</a>	<a href="#">fleekee.aSep08</a>		5399	327		21	putative protein (fleekee) mRNA.
<a href="#">fleelo</a>	<a href="#">fleelo.aSep08</a>		2274	436		145	mlx interacting protein CRA a (fleelo) mRNA.
<a href="#">fleemee</a>	<a href="#">fleemee.aSep08</a>		6357	356		118	pitpnm family member 3 (fleemee) mRNA.
<a href="#">fleepor</a>	<a href="#">fleepor.aSep08</a>		936	540		180	CRA a (fleepor) mRNA.
<a href="#">fleeroy</a>	<a href="#">fleeroy.aSep08</a>		760	643		37	putative protein (fleeroy) mRNA.
<a href="#">fleesa</a>	<a href="#">fleesa.aSep08</a>		16156	445		34	putative protein (fleesa) mRNA.
<a href="#">fleeshee</a>	<a href="#">fleeshee.aSep08</a>		6705	845		42	putative protein (4.9 kD) (fleeshee) mRNA.
<a href="#">fleetu</a>	<a href="#">fleetu.aSep08</a>		4504	649		102	ninein (fleetu) mRNA.
<a href="#">fleevo</a>	<a href="#">fleevo.aSep08</a>		10862	164		40	putative protein (fleevo) mRNA.
<a href="#">fleewer</a>	<a href="#">fleewer.aSep08</a>		6173	340		59	CRA b like (fleewer) mRNA.
<a href="#">flerbor</a>	<a href="#">flerbor.aSep08</a>		13835	550		51	putative protein (5.8 kD) (flerbor) mRNA.
<a href="#">flerchy</a>	<a href="#">flerchy.aSep08</a>		6381	557		98	putative protein (flerchy) mRNA.
<a href="#">flerdoy</a>	<a href="#">flerdoy.aSep08</a>		1504	487		35	putative protein (4.0 kD) (flerdoy) mRNA.
<a href="#">flerflu</a>	<a href="#">flerflu.aSep08</a>		17632	451		97	transmembrane channel-like 5 (flerflu) mRNA.
<a href="#">flerfly</a>	<a href="#">flerfly.aSep08</a>		1602	1418		61	putative protein (6.8 kD) (flerfly) mRNA.
<a href="#">flergar</a>	<a href="#">flergar.aSep08</a>		5133	448		148	histone acetyltransferase myst3 (flergar) mRNA.
<a href="#">flerja</a>	<a href="#">flerja.aSep08</a>		599	381		111	putative protein (flerja) mRNA.
<a href="#">flerjey</a>	<a href="#">flerjey.aSep08</a>		6633	504		33	putative protein (flerjey) mRNA.
<a href="#">flerkee</a>	<a href="#">flerkee.aSep08</a>		11286	815		59	putative protein (6.8 kD) (flerkee) mRNA.
<a href="#">flerlo</a>	<a href="#">flerlo.aSep08</a>		25028	535	4	123	putative protein (13.7 kD) (flerlo) alternative variant aSep08, mRNA.
<a href="#">flerlo</a>	<a href="#">flerlo.bSep08</a>		25575	545	4	123	putative protein (13.7 kD) (flerlo) alternative variant bSep08, mRNA.
<a href="#">flermee</a>	<a href="#">flermee.aSep08</a>		5132	229		75	uncharacterized protein like (flermee) mRNA.
<a href="#">flerpor</a>	<a href="#">flerpor.aSep08</a>		699	403		134	CRA c like (flerpor) mRNA.
<a href="#">flerroy</a>	<a href="#">flerroy.aSep08</a>		12227	349		51	putative protein (flerroy) mRNA.
<a href="#">flersa</a>	<a href="#">flersa.aSep08</a>		3569	401		133	putative protein (flersa) mRNA.
<a href="#">flershee</a>	<a href="#">flershee.aSep08</a>		13834	677	7	225	sortilin (flershee) alternative variant aSep08, mRNA.
<a href="#">flershee</a>	<a href="#">flershee.bSep08</a>		49214	605	8	201	sortilin (flershee) alternative variant bSep08, mRNA.
<a href="#">flershee</a>	<a href="#">flershee.cSep08</a>		26487	528	3	52	sortilin (flershee) alternative variant cSep08, mRNA.
<a href="#">flertu</a>	<a href="#">flertu.aSep08</a>		6230	1266	4	46	putative protein (4.9 kD) (flertu) alternative variant aSep08, mRNA.
<a href="#">flervo</a>	<a href="#">flervo.aSep08</a>		362	253		83	putative protein (flervo) mRNA.
<a href="#">flerwer</a>	<a href="#">flerwer.aSep08</a>		5697	345		55	CRA b like (flerwer) mRNA.
<a href="#">fleybor</a>	<a href="#">fleybor.aSep08</a>		1044	392		53	putative protein (6.2 kD) (fleybor) mRNA.
<a href="#">fleychy</a>	<a href="#">fleychy.aSep08</a>		11486	728		28	putative protein (3.2 kD) (fleychy) mRNA.
<a href="#">fleydoy</a>	<a href="#">fleydoy.aSep08</a>		1706	355		51	putative protein (6.0 kD) (fleydoy) mRNA.

<a href="#">fleyflu</a>	<a href="#">fleyflu.aSep08</a>		16034	256		18	putative protein (fleyflu) mRNA.
<a href="#">fleyfly</a>	<a href="#">fleyfly.aSep08</a>		588	290		24	putative protein (fleyfly) mRNA.
<a href="#">fleygar</a>	<a href="#">fleygar.aSep08</a>		3116	954		193	myst histone acetyltransferase 3 (fleygar) alternative variant aSep08, mRNA.
<a href="#">fleygar</a>	<a href="#">fleygar.bSep08</a>		2516	441	1	51	putative protein (fleygar) alternative variant bSep08, mRNA.
<a href="#">fleyja</a>	<a href="#">fleyja.aSep08</a>		3066	621		66	putative protein (7.9 kD) (fleyja) mRNA.
<a href="#">fleyjey</a>	<a href="#">fleyjey.aSep08</a>		4408	227		75	adenylate cyclase 1 (fleyjey) mRNA.
<a href="#">fleykee</a>	<a href="#">fleykee.aSep08</a>		2921	1243		163	integrator complex (fleykee) mRNA.
<a href="#">fleylo</a>	<a href="#">fleylo.aSep08</a>		8199	737		245	WD repeat domain 66 (fleylo) mRNA.
<a href="#">fleymee</a>	<a href="#">fleymee.aSep08</a>		741	356		79	uncharacterized protein like (fleymee) mRNA.
<a href="#">fleypor</a>	<a href="#">fleypor.aSep08</a>		10508	420		139	putative protein (fleypor) mRNA.
<a href="#">fleyroy</a>	<a href="#">fleyroy.aSep08</a>		707	544		180	putative protein of mammalian origin (fleyroy) mRNA.
<a href="#">fleysa</a>	<a href="#">fleysa.aSep08</a>		5398	359		119	extra spindle poles like 1 (fleysa) mRNA.
<a href="#">fleyshee</a>	<a href="#">fleyshee.aSep08</a>		5158	642		214	cadherin EGF LAG seven-pass G-type receptor (fleyshee) mRNA.
<a href="#">fleytu</a>	<a href="#">fleytu.aSep08</a>		824	720		81	putative protein (fleytu) mRNA.
<a href="#">fleyvo</a>	<a href="#">fleyvo.aSep08</a>		15045	457	2	55	putative protein (fleyvo) alternative variant aSep08, mRNA.
<a href="#">fleyvo</a>	<a href="#">fleyvo.bSep08</a>		1368	334	1	39	putative protein (fleyvo) alternative variant bSep08, mRNA.
<a href="#">fleywer</a>	<a href="#">fleywer.aSep08</a>		4295	745		82	putative cytoplasmic protein (10.2 kD) (fleywer) mRNA.
<a href="#">Flii</a>	<a href="#">Flii.aSep08</a>	<a href="#">287375</a>	14647	4725	14	1270	flightless I homolog (Drosophila) (144.9 kD) (Flii) alternative variant aSep08, mRNA.
<a href="#">Flii</a>	<a href="#">Flii.bSep08</a>	<a href="#">287375</a>	1616	744	1	247	flightless I homolog (Drosophila) (Flii) alternative variant bSep08, mRNA.
<a href="#">Flii</a>	<a href="#">Flii.cSep08</a>	<a href="#">287375</a>	2235	1734	2	174	flightless I homolog (Drosophila) (Flii) alternative variant cSep08, mRNA.
<a href="#">Flna</a>	<a href="#">Flna.bSep08</a>	<a href="#">293860</a>	4922	1092	9	363	filamin A (Flna) alternative variant bSep08, mRNA.
<a href="#">Flna</a>	<a href="#">Flna.cSep08</a>	<a href="#">293860</a>	3426	1821	5	223	filamin alpha (23.4 kD) (Flna) alternative variant cSep08, mRNA.
<a href="#">Flna</a>	<a href="#">Flna.dSep08</a>	<a href="#">293860</a>	2013	657	4	195	filamin Alpha (Flna) alternative variant dSep08, mRNA.
<a href="#">Flna</a>	<a href="#">Flna.eSep08</a>	<a href="#">293860</a>	1166	707	4	191	filamin Alpha (Flna) alternative variant eSep08, mRNA.
<a href="#">Flna</a>	<a href="#">Flna.fSep08</a>	<a href="#">293860</a>	425	280	3	93	filamin alpha (Flna) alternative variant fSep08, mRNA.
<a href="#">Flnb</a>	<a href="#">Flnb.bSep08</a>	<a href="#">306204</a>	9556	720	7	240	filamin, beta (Flnb) alternative variant bSep08, mRNA.
<a href="#">Flnc</a>	<a href="#">Flnc.aSep08</a>	<a href="#">362332</a>	10962	4339	23	1219	filamin C, gamma (actin binding protein 280) (Flnc) alternative variant aSep08, mRNA.
<a href="#">Flnc</a>	<a href="#">Flnc.bSep08</a>	<a href="#">362332</a>	14670	654	4	159	filamin C, gamma (actin binding protein 280) (Flnc) alternative variant bSep08, mRNA.
<a href="#">flobor</a>	<a href="#">flobor.aSep08</a>		8205	735		34	putative protein (flobor) mRNA.
<a href="#">flochy</a>	<a href="#">flochy.aSep08</a>		6781	271		76	putative protein (flochy) mRNA.
<a href="#">flodoy</a>	<a href="#">flodoy.aSep08</a>		1481	523		174	dmx-like 1 (flodoy) mRNA.
<a href="#">floflu</a>	<a href="#">floflu.aSep08</a>		1346	432		143	PI-3-kinase-related kinase smg-1 (floflu) mRNA.
<a href="#">flofly</a>	<a href="#">flofly.aSep08</a>		10892	927	5	86	putative protein (9.5 kD) (flofly) alternative variant aSep08, mRNA.

<a href="#">flofly</a>	<a href="#">flofly.cSep08</a>		5811	329	2	54	putative protein (5.9 kD) (flofly) alternative variant cSep08, mRNA.
<a href="#">flogar</a>	<a href="#">flogar.aSep08</a>		27549	697		86	putative protein (flogar) mRNA.
<a href="#">floja</a>	<a href="#">floja.aSep08</a>		3630	305		101	exportin 7 (floja) mRNA.
<a href="#">flojey</a>	<a href="#">flojey.aSep08</a>		558	351		117	zinc finger MIZ-type containing 2 (flojey) mRNA.
<a href="#">flokee</a>	<a href="#">flokee.aSep08</a>		3767	708		75	putative protein (8.2 kD) (flokee) mRNA.
<a href="#">flogo</a>	<a href="#">flogo.aSep08</a>		13620	330		103	putative protein (flogo) mRNA.
<a href="#">flomee</a>	<a href="#">flomee.aSep08</a>		3651	355		45	putative protein (flomee) mRNA.
<a href="#">flonoy</a>	<a href="#">flonoy.aSep08</a>		1811	1481		67	putative protein (7.7 kD) (flonoy) mRNA.
<a href="#">flopore</a>	<a href="#">flopore.aSep08</a>		2336	608	4	71	putative protein (7.7 kD) (flopore) alternative variant aSep08, mRNA.
<a href="#">flopore</a>	<a href="#">flopore.cSep08</a>		2823	608	2	75	putative mitochondrial protein (8.2 kD) (flopore) alternative variant cSep08, mRNA.
<a href="#">florbor</a>	<a href="#">florbor.aSep08</a>		2944	965	1	37	putative protein (florbor) alternative variant aSep08, mRNA.
<a href="#">florbor</a>	<a href="#">florbor.bSep08</a>		2522	699	1	50	putative protein (5.4 kD) (florbor) alternative variant bSep08, mRNA.
<a href="#">florchy</a>	<a href="#">florchy.aSep08</a>		1196	389		73	putative protein (florchy) mRNA.
<a href="#">flordoy</a>	<a href="#">flordoy.aSep08</a>		6760	608		14	putative protein (flordoy) mRNA.
<a href="#">florflu</a>	<a href="#">florflu.aSep08</a>		29572	1777		35	putative protein (florflu) mRNA.
<a href="#">florfly</a>	<a href="#">florfly.aSep08</a>		1239	574		39	putative protein (4.6 kD) (florfly) mRNA.
<a href="#">florgar</a>	<a href="#">florgar.aSep08</a>		5940	426		92	putative protein (florgar) mRNA.
<a href="#">florja</a>	<a href="#">florja.aSep08</a>		49673	384		56	reverse transcriptase like (6.7 kD) (florja) mRNA.
<a href="#">florjey</a>	<a href="#">florjey.aSep08</a>		16473	561	1	156	tensin 3 (florjey) alternative variant aSep08, mRNA.
<a href="#">florjey</a>	<a href="#">florjey.bSep08</a>		14573	267	1	71	putative protein (florjey) alternative variant bSep08, mRNA.
<a href="#">florkee</a>	<a href="#">florkee.aSep08</a>		1456	522		124	putative protein (13.7 kD) (florkee) mRNA.
<a href="#">florlo</a>	<a href="#">florlo.aSep08</a>		1879	741	2	122	putative protein (florlo) alternative variant aSep08, mRNA.
<a href="#">flormee</a>	<a href="#">flormee.aSep08</a>		1311	322		106	uncharacterized protein (flormee) mRNA.
<a href="#">floroy</a>	<a href="#">floroy.aSep08</a>		536	256		84	ABC transporter (floroy) mRNA.
<a href="#">florpor</a>	<a href="#">florpor.aSep08</a>		1898	244		80	putative protein (florpor) mRNA.
<a href="#">florroy</a>	<a href="#">florroy.aSep08</a>		6854	3792	5	369	putative protein, with a coiled coil domain, of vertebrate origin (florroy) alternative variant aSep08, mRNA.
<a href="#">florroy</a>	<a href="#">florroy.bSep08</a>		643	415	1	138	putative protein of vertebrate origin (florroy) alternative variant bSep08, mRNA.
<a href="#">florsa</a>	<a href="#">florsa.aSep08</a>		12174	514		43	putative protein (florsa) mRNA.
<a href="#">florshee</a>	<a href="#">florshee.aSep08</a>		14300	422		39	modulator 2 (florshee) mRNA.
<a href="#">flortu</a>	<a href="#">flortu.aSep08</a>		12523	717	2	80	putative protein (flortu) alternative variant aSep08, mRNA.
<a href="#">flortu</a>	<a href="#">flortu.bSep08</a>		12616	491	3	33	putative protein (flortu) alternative variant bSep08, mRNA.
<a href="#">florvo</a>	<a href="#">florvo.aSep08</a>		2297	370		123	midasin (florvo) mRNA.
<a href="#">florwer</a>	<a href="#">florwer.aSep08</a>		516	410		41	putative protein (4.7 kD) (florwer) mRNA.
<a href="#">flosa</a>	<a href="#">flosa.aSep08</a>		5960	413		49	putative protein (flosa) mRNA.
<a href="#">floshee</a>	<a href="#">floshee.aSep08</a>		2352	383		39	putative protein (4.5 kD) (floshee) mRNA.
<a href="#">Flot1</a>	<a href="#">Flot1.bSep08</a>	<a href="#">64665</a>	7670	1242	10	352	flotillin (39.8 kD) (Flot1) alternative variant bSep08, mRNA.

<a href="#">Flot1</a>	<a href="#">Flot1.cSep08</a>	<a href="#">64665</a>	7242	817	8	176	flotillin 1 (Flot1) alternative variant cSep08, mRNA.
<a href="#">Flot1</a>	<a href="#">Flot1.dSep08</a>	<a href="#">64665</a>	2339	636	7	143	flotillin 1 CRA a (Flot1) alternative variant dSep08, mRNA.
<a href="#">Flot1</a>	<a href="#">Flot1.eSep08</a>	<a href="#">64665</a>	563	415	2	84	flotillin 1 CRA a (Flot1) alternative variant eSep08, mRNA.
<a href="#">Flot1</a>	<a href="#">Flot1.fSep08</a>	<a href="#">64665</a>	2136	467	2	75	flotillin 1 (Flot1) alternative variant fSep08, mRNA.
<a href="#">Flot1</a>	<a href="#">Flot1.hSep08</a>	<a href="#">64665</a>	560	315	2	32	putative protein (Flot1) alternative variant hSep08, mRNA.
<a href="#">Flot2</a>	<a href="#">Flot2.bSep08</a>	<a href="#">83764</a>	19870	769	7	256	flotillin 2 (Flot2) alternative variant bSep08, mRNA.
<a href="#">Flot2</a>	<a href="#">Flot2.cSep08</a>	<a href="#">83764</a>	20031	754	7	181	flotillin 2 (Flot2) alternative variant cSep08, mRNA.
<a href="#">Flot2</a>	<a href="#">Flot2.eSep08</a>	<a href="#">83764</a>	18022	397	4	81	flotillin (Flot2) alternative variant eSep08, mRNA.
<a href="#">flotu</a>	<a href="#">flotu.aSep08</a>		6260	2058	6	175	serologically defined colon cancer antigen 1 like (flotu) alternative variant aSep08, mRNA.
<a href="#">flovo</a>	<a href="#">flovo.aSep08</a>		4764	1361	4	93	putative protein (flovo) alternative variant aSep08, mRNA.
<a href="#">flower</a>	<a href="#">flower.aSep08</a>		7711	524		89	carrier family (flower) mRNA.
<a href="#">floybor</a>	<a href="#">floybor.aSep08</a>		1511	618		68	putative protein (floybor) mRNA.
<a href="#">floychy</a>	<a href="#">floychy.aSep08</a>		1345	370		35	putative protein (floychy) mRNA.
<a href="#">floydoy</a>	<a href="#">floydoy.aSep08</a>		899	330		39	putative protein of mammalian origin (floydoy) mRNA.
<a href="#">floyflu</a>	<a href="#">floyflu.aSep08</a>		1351	985		44	putative protein (5.1 kD) (floyflu) mRNA.
<a href="#">floyfly</a>	<a href="#">floyfly.aSep08</a>		7941	1681	4	98	ab2-076 like (floyfly) alternative variant aSep08, mRNA.
<a href="#">floygar</a>	<a href="#">floygar.bSep08</a>		14865	3307	4	225	putative protein of vertebrate origin (floygar) alternative variant bSep08, mRNA.
<a href="#">floyja</a>	<a href="#">floyja.aSep08</a>		2941	311		65	putative protein (floyja) mRNA.
<a href="#">floyjey</a>	<a href="#">floyjey.aSep08</a>		13684	406		135	tensin 3 (floyjey) mRNA.
<a href="#">floykee</a>	<a href="#">floykee.aSep08</a>		123300	386		128	hedgehog acyltransferase (floykee) mRNA.
<a href="#">floylo</a>	<a href="#">floylo.aSep08</a>		4083	603	2	33	putative protein (floylo) alternative variant aSep08, mRNA.
<a href="#">floylo</a>	<a href="#">floylo.cSep08</a>		4083	316	2	12	putative protein (floylo) alternative variant cSep08, mRNA.
<a href="#">floymee</a>	<a href="#">floymee.aSep08</a>		2478	792		70	putative protein (floymee) mRNA.
<a href="#">floypor</a>	<a href="#">floypor.aSep08</a>		2225	471		111	als2 C-terminal like (floypor) mRNA.
<a href="#">floyroy</a>	<a href="#">floyroy.aSep08</a>		1340	349		50	putative protein (floyroy) mRNA.
<a href="#">floyrsa</a>	<a href="#">floyrsa.aSep08</a>		991	558		183	putative protein (floyrsa) mRNA.
<a href="#">floyrshee</a>	<a href="#">floyrshee.aSep08</a>		4061	374		75	putative protein (floyrshee) mRNA.
<a href="#">floytu</a>	<a href="#">floytu.aSep08</a>		10143	840		280	CRA a (floytu) mRNA.
<a href="#">floyvo</a>	<a href="#">floyvo.aSep08</a>		14822	1781		573	midasin (floyvo) mRNA.
<a href="#">floywer</a>	<a href="#">floywer.aSep08</a>		1451	525		135	putative protein (15.0 kD) (floywer) mRNA.
<a href="#">Flt1</a>	<a href="#">Flt1.bSep08</a>	<a href="#">54251</a>	17386	2454	1	185	FMS-like tyrosine kinase 1 (Flt1) alternative variant bSep08, mRNA.
<a href="#">Flt3</a>	<a href="#">Flt3.aSep08</a>	<a href="#">140635</a>	13173	1255		274	FMS-like tyrosine kinase 3 (Flt3) mRNA.
<a href="#">Flt3_lig.0</a>	<a href="#">Flt3_lig.0.aSep08</a>		5187	1161	6	253	flt3 ligand (Flt3_lig.0) alternative variant aSep08, mRNA.
<a href="#">Flt3_lig.0</a>	<a href="#">Flt3_lig.0.bSep08</a>		2706	395	1	131	CRA d like (Flt3_lig.0) alternative variant bSep08, mRNA.
<a href="#">Flt3_lig.0</a>	<a href="#">Flt3_lig.0.cSep08</a>		2052	390	5	115	ligand (Flt3_lig.0) alternative variant cSep08, mRNA.
<a href="#">flubor</a>	<a href="#">flubor.aSep08</a>		905	759		60	putative protein (flubor) mRNA.
<a href="#">fluchy</a>	<a href="#">fluchy.aSep08</a>		77776	412		85	putative protein (fluchy) mRNA.
<a href="#">flufly</a>	<a href="#">flufly.aSep08</a>		1752	557		185	kinase smg1 (flufly) mRNA.

<a href="#">flufly</a>	<a href="#">flufly.aSep08</a>		28561	529		176	AT rich interactive domain 1B (flufly) mRNA.
<a href="#">flugar</a>	<a href="#">flugar.bSep08</a>		811	305	2	75	putative protein (flugar) alternative variant bSep08, mRNA.
<a href="#">fluja</a>	<a href="#">fluja.aSep08</a>		2476	447		60	putative protein (fluja) mRNA.
<a href="#">flujey</a>	<a href="#">flujey.aSep08</a>		6867	985	1	77	zinc finger MIZ-type containing 2 (8.0 kD) (flujey) alternative variant aSep08, mRNA.
<a href="#">flujey</a>	<a href="#">flujey.bSep08</a>		6284	399	1	67	putative protein (flujey) alternative variant bSep08, mRNA.
<a href="#">flukee</a>	<a href="#">flukee.aSep08</a>		88055	706		35	putative protein (flukee) mRNA.
<a href="#">flulo</a>	<a href="#">flulo.aSep08</a>		17908	419	4	78	putative protein (7.9 kD) (flulo) alternative variant aSep08, mRNA.
<a href="#">flulo</a>	<a href="#">flulo.bSep08</a>		944	469	2	51	putative protein (flulo) alternative variant bSep08, mRNA.
<a href="#">flumee</a>	<a href="#">flumee.aSep08</a>		945	480		111	containing 1 (flumee) mRNA.
<a href="#">flunoy</a>	<a href="#">flunoy.aSep08</a>		27450	1183		332	putative protein of eukaryotic origin (flunoy) mRNA.
<a href="#">flupor</a>	<a href="#">flupor.aSep08</a>		1429	437		41	type VII collagen (flupor) mRNA.
<a href="#">fluoy</a>	<a href="#">fluoy.aSep08</a>		919	255		54	solute carrier family 37 member 3 like (fluoy) mRNA.
<a href="#">flusa</a>	<a href="#">flusa.aSep08</a>		745	229		46	putative protein (flusa) mRNA.
<a href="#">flushee</a>	<a href="#">flushee.aSep08</a>		55432	1396		75	putative nuclear protein (8.6 kD) (flushee) mRNA.
<a href="#">flutu</a>	<a href="#">flutu.aSep08</a>		5754	370	2	47	putative protein (flutu) alternative variant aSep08, mRNA.
<a href="#">flutu</a>	<a href="#">flutu.bSep08</a>		6353	273	2	26	putative protein (flutu) alternative variant bSep08, mRNA.
<a href="#">fluvo</a>	<a href="#">fluvo.aSep08</a>		9611	471		24	putative protein (2.5 kD) (fluvo) mRNA.
<a href="#">fluwer</a>	<a href="#">fluwer.aSep08</a>		2284	306		102	domain 22 (fluwer) mRNA.
<a href="#">Flvcr2</a>	<a href="#">Flvcr2.bSep08</a>	<a href="#">314323</a>	9654	1690		73	feline leukemia virus subgroup C cellular receptor family, member 2 (Flvcr2) alternative variant bSep08, mRNA.
<a href="#">flybor</a>	<a href="#">flybor.aSep08</a>		3119	245		25	putative protein (flybor) mRNA.
<a href="#">flychy</a>	<a href="#">flychy.aSep08</a>		12240	501		166	akt substrate as250 (flychy) mRNA.
<a href="#">flydoy</a>	<a href="#">flydoy.aSep08</a>		1471	764		54	putative protein (5.9 kD) (flydoy) mRNA.
<a href="#">flyflu</a>	<a href="#">flyflu.aSep08</a>		4703	642		213	kinase (flyflu) mRNA.
<a href="#">flyfly</a>	<a href="#">flyfly.aSep08</a>		2933	699	3	67	putative protein (flyfly) alternative variant aSep08, mRNA.
<a href="#">flyfly</a>	<a href="#">flyfly.bSep08</a>		1895	343	1	32	CRA a like (flyfly) alternative variant bSep08, mRNA.
<a href="#">flygar</a>	<a href="#">flygar.aSep08</a>		5000	383		127	heparan-alpha-glucosaminide n-acetyltransferase (flygar) mRNA.
<a href="#">flyja</a>	<a href="#">flyja.aSep08</a>		11356	515	2	171	bone morphogenetic protein 1 (flyja) alternative variant aSep08, mRNA.
<a href="#">flyja</a>	<a href="#">flyja.bSep08</a>		8522	516	1	112	bone morphogenetic protein 1 CRA c (flyja) alternative variant bSep08, mRNA.
<a href="#">flyjey</a>	<a href="#">flyjey.aSep08</a>		22535	773		34	putative protein (3.7 kD) (flyjey) mRNA.
<a href="#">flykee</a>	<a href="#">flykee.aSep08</a>		4381	777		52	putative protein (6.0 kD) (flykee) mRNA.
<a href="#">flylo</a>	<a href="#">flylo.aSep08</a>		1425	692	2	161	putative protein (6.2 kD) (flylo) alternative variant aSep08, mRNA.
<a href="#">flylo</a>	<a href="#">flylo.cSep08</a>		469	335	2	34	putative protein (4.0 kD) (flylo) alternative variant cSep08, mRNA.
<a href="#">flymee</a>	<a href="#">flymee.aSep08</a>		1743	1154		151	putative protein (15.8 kD) (flymee) mRNA.
<a href="#">flynoy</a>	<a href="#">flynoy.aSep08</a>		10113	736		244	putative protein of metazoan origin (flynoy) mRNA.

<a href="#">flypor</a>	<a href="#">flypor.aSep08</a>		1453	727	4	209	solute carrier family 26 member 6 CRA a (flypor) alternative variant aSep08, mRNA.
<a href="#">flypor</a>	<a href="#">flypor.bSep08</a>		2018	393	1	90	solute carrier family 26 member 6 CRA a (flypor) alternative variant bSep08, mRNA.
<a href="#">flyroy</a>	<a href="#">flyroy.aSep08</a>		687	322	2	91	putative protein (flyroy) alternative variant aSep08, mRNA.
<a href="#">flysa</a>	<a href="#">flysa.aSep08</a>		2466	1543	2	116	putative protein (flysa) alternative variant aSep08, mRNA.
<a href="#">flyshee</a>	<a href="#">flyshee.aSep08</a>		2845	470		65	putative protein of mammalian origin (7.5 kD) (flyshee) mRNA.
<a href="#">flytu</a>	<a href="#">flytu.aSep08</a>		2160	325		23	putative protein (flytu) mRNA.
<a href="#">flyvo</a>	<a href="#">flyvo.aSep08</a>		12581	734		69	putative protein (7.8 kD) (flyvo) mRNA.
<a href="#">Flywch1</a>	<a href="#">Flywch1.aSep08</a>	<a href="#">360488</a>	16307	2264	7	662	zinc finger, FLYWCH-type (Flywch1) alternative variant aSep08, mRNA.
<a href="#">Flywch1</a>	<a href="#">Flywch1.bSep08</a>	<a href="#">360488</a>	8510	1807	5	464	zinc finger, FLYWCH-type (Flywch1) alternative variant bSep08, mRNA.
<a href="#">Flywch1</a>	<a href="#">Flywch1.cSep08</a>	<a href="#">360488</a>	8662	1833	4	263	zinc finger, FLYWCH-type (Flywch1) alternative variant cSep08, mRNA.
<a href="#">Flywch1</a>	<a href="#">Flywch1.dSep08</a>	<a href="#">360488</a>	5836	318	2	63	CRA a like (Flywch1) alternative variant dSep08, mRNA.
<a href="#">Flywch1</a>	<a href="#">Flywch1.eSep08</a>	<a href="#">360488</a>	8231	727	4	98	putative protein (10.8 kD) (Flywch1) alternative variant eSep08, mRNA.
<a href="#">flywer</a>	<a href="#">flywer.aSep08</a>		4789	576		103	putative protein (11.8 kD) (flywer) mRNA.
<a href="#">Fmnl1</a>	<a href="#">Fmnl1.bSep08</a>	<a href="#">287746</a>	7721	970	9	322	formin-like 1 (Fmnl1) alternative variant bSep08, mRNA.
<a href="#">Fmnl1</a>	<a href="#">Fmnl1.cSep08</a>	<a href="#">287746</a>	3136	1139	8	281	formin-like 1 (Fmnl1) alternative variant cSep08, mRNA.
<a href="#">Fmo1</a>	<a href="#">Fmo1.bSep08</a>	<a href="#">25256</a>	15889	693	4	181	flavin containing monooxygenase 1 (Fmo1) alternative variant bSep08, mRNA.
<a href="#">Fmo2</a>	<a href="#">Fmo2.bSep08</a>	<a href="#">246245</a>	11562	753	5	192	flavin containing monooxygenase 2 (Fmo2) alternative variant bSep08, mRNA.
<a href="#">Fmo2</a>	<a href="#">Fmo2.dSep08</a>	<a href="#">246245</a>	31125	957	3	68	flavin containing monooxygenase 2 (Fmo2) alternative variant dSep08, mRNA.
<a href="#">Fmo4</a>	<a href="#">Fmo4.cSep08</a>	<a href="#">246247</a>	13139	1779	4	464	flavin containing monooxygenase 4 (Fmo4) alternative variant cSep08, mRNA.
<a href="#">Fmo4</a>	<a href="#">Fmo4.dSep08</a>	<a href="#">246247</a>	3994	663	3	153	flavin containing monooxygenase 4 (Fmo4) alternative variant dSep08, mRNA.
<a href="#">Fmo4</a>	<a href="#">Fmo4.eSep08</a>	<a href="#">246247</a>	2627	377	2	107	flavin containing monooxygenase 4 (Fmo4) alternative variant eSep08, mRNA.
<a href="#">Fmo4</a>	<a href="#">Fmo4.fSep08</a>	<a href="#">246247</a>	1803	749	2	77	putative protein (8.6 kD) (Fmo4) alternative variant fSep08, mRNA.
<a href="#">Fmo5</a>	<a href="#">Fmo5.bSep08</a>	<a href="#">246248</a>	10073	747	4	238	flavin containing monooxygenase 5 (Fmo5) alternative variant bSep08, mRNA.
<a href="#">Fmo5</a>	<a href="#">Fmo5.cSep08</a>	<a href="#">246248</a>	8516	749	3	207	flavin containing monooxygenase 5 (Fmo5) alternative variant cSep08, mRNA.
<a href="#">Fmo5</a>	<a href="#">Fmo5.dSep08</a>	<a href="#">246248</a>	10053	683	5	119	flavin containing monooxygenase 5 (Fmo5) alternative variant dSep08, mRNA.
<a href="#">Fmo6</a>	<a href="#">Fmo6.aSep08</a>	<a href="#">304922</a>	11613	562		146	flavin containing monooxygenase 6 (Fmo6) mRNA.
<a href="#">Fmp27_GFWDK.0</a>	<a href="#">Fmp27_GFWDK.0.aSep08</a>		2644	1180		393	CRA a (Fmp27_GFWDK.0) mRNA.



<a href="#">Fmr1</a>	<a href="#">Fmr1.bSep08</a>	<a href="#">24948</a>	23117	3835	13	450	fragile X mental retardation syndrome 1 homolog (50.0 kD) (Fmr1) alternative variant bSep08, mRNA.
<a href="#">Fmr1</a>	<a href="#">Fmr1.cSep08</a>	<a href="#">24948</a>	26026	1204	12	243	fragile X mental retardation syndrome 1 homolog (Fmr1) alternative variant cSep08, mRNA.
<a href="#">Fmr1</a>	<a href="#">Fmr1.eSep08</a>	<a href="#">24948</a>	1409	770	2	65	fragile X mental retardation syndrome 1 homolog (Fmr1) alternative variant eSep08, mRNA.
<a href="#">Fmr1nb</a>	<a href="#">Fmr1nb.aSep08</a>	<a href="#">293918</a>	22651	849	2	242	fragile X mental retardation 1 neighbor (27.1 kD) (Fmr1nb) alternative variant aSep08, mRNA.
<a href="#">Fmr1nb</a>	<a href="#">Fmr1nb.bSep08</a>	<a href="#">293918</a>	20600	544		145	fragile X mental retardation 1 neighbor (Fmr1nb) alternative variant bSep08, mRNA.
<a href="#">Fn1</a>	<a href="#">Fn1.bSep08</a>	<a href="#">25661</a>	32960	3111	22	1036	fibronectin 1 (Fn1) alternative variant bSep08, mRNA.
<a href="#">Fn1</a>	<a href="#">Fn1.cSep08</a>	<a href="#">25661</a>	8082	1249	8	416	fibronectin 1 (Fn1) alternative variant cSep08, mRNA.
<a href="#">Fn1</a>	<a href="#">Fn1.dSep08</a>	<a href="#">25661</a>	1601	495	2	164	fibronectin 1 (Fn1) alternative variant dSep08, mRNA.
<a href="#">fn3.0</a>	<a href="#">fn3.0.aSep08</a>		5391	997	4	320	novel protein containing fibronectin type 3 FN3 domains (fn3.0) alternative variant aSep08, mRNA.
<a href="#">fn3.0</a>	<a href="#">fn3.0.bSep08</a>		2938	653	1	126	novel protein containing fibronectin type 3 FN3 domains like (fn3.0) alternative variant bSep08, mRNA.
<a href="#">fn3.1</a>	<a href="#">fn3.1.aSep08</a>		1964	1682	1	209	tenascin XB (fn3.1) alternative variant aSep08, mRNA.
<a href="#">fn3.1</a>	<a href="#">fn3.1.bSep08</a>		4808	406	2	135	tenascin XB (fn3.1) alternative variant bSep08, mRNA.
<a href="#">fn3.2</a>	<a href="#">fn3.2.aSep08</a>		13182	709		236	myomesin 2 (fn3.2) mRNA.
<a href="#">fn3.3</a>	<a href="#">fn3.3.aSep08</a>		4079	354		118	usherin (fn3.3) mRNA.
<a href="#">fn3.4</a>	<a href="#">fn3.4.aSep08</a>		33109	428		142	sidekick 1 (fn3.4) mRNA.
<a href="#">fn3.5</a>	<a href="#">fn3.5.aSep08</a>		9633	1027		342	2 -binding protein like (fn3.5) mRNA.
<a href="#">fn3.6</a>	<a href="#">fn3.6.aSep08</a>		71693	3900	5	522	roundabout 1 (fn3.6) alternative variant aSep08, mRNA.
<a href="#">fn3.6</a>	<a href="#">fn3.6.bSep08</a>		18708	775	2	258	roundabout homolog 1 (fn3.6) alternative variant bSep08, mRNA.
<a href="#">fn3.7</a>	<a href="#">fn3.7.aSep08</a>		37997	1672		449	roundabout homolog 2 (fn3.7) mRNA.
<a href="#">fn3.8</a>	<a href="#">fn3.8.aSep08</a>		32125	2466	11	437	ABI gene family member 3 binding protein like (fn3.8) alternative variant aSep08, mRNA.
<a href="#">fn3.8</a>	<a href="#">fn3.8.bSep08</a>		2350	742	1	98	ABI gene family member 3 binding protein like (fn3.8) alternative variant bSep08, mRNA.
<a href="#">fn3.9</a>	<a href="#">fn3.9.aSep08</a>		14675	612		203	sidekick homolog 2 (fn3.9) mRNA.
<a href="#">fn3.10</a>	<a href="#">fn3.10.aSep08</a>		17327	773		193	myomesin 1 (fn3.10) mRNA.
<a href="#">fn3.11</a>	<a href="#">fn3.11.aSep08</a>		11054	775	5	258	neogenin (fn3.11) alternative variant aSep08, mRNA.
<a href="#">fn3.11</a>	<a href="#">fn3.11.bSep08</a>		5778	340	1	80	neogenin (fn3.11) alternative variant bSep08, mRNA.
<a href="#">fn3.12</a>	<a href="#">fn3.12.aSep08</a>		14024	1786		506	neogenin (fn3.12) alternative variant aSep08, mRNA.
<a href="#">fn3.12</a>	<a href="#">fn3.12.bSep08</a>		16786	831		276	neogenin (fn3.12) alternative variant bSep08, mRNA.
<a href="#">fn3.13</a>	<a href="#">fn3.13.aSep08</a>		1761	357		118	type XII alpha 1 (fn3.13) mRNA.
<a href="#">fn3.14</a>	<a href="#">fn3.14.aSep08</a>		3422	501		167	type XII alpha 1 (fn3.14) mRNA.
<a href="#">fn3.15</a>	<a href="#">fn3.15.aSep08</a>		14267	758		252	type XII alpha 1 (fn3.15) mRNA.
<a href="#">fn3.16</a>	<a href="#">fn3.16.aSep08</a>		6798	2611	5	182	fibronectin, type III (fn3.16) mRNA.
<a href="#">fn3.17</a>	<a href="#">fn3.17.aSep08</a>		16353	413		137	receptor protein tyrosine phosphatase hPTP-J (fn3.17) mRNA.
<a href="#">fn3.19</a>	<a href="#">fn3.19.aSep08</a>		854	644		214	titin (fn3.19) mRNA.

<a href="#">fn3.20</a>	<a href="#">fn3.20.aSep08</a>		1607	464		154	titin CRA a (fn3.20) mRNA.
<a href="#">fn3.21</a>	<a href="#">fn3.21.aSep08</a>		3982	1175		391	titin CRA a (fn3.21) mRNA.
<a href="#">fn3.22</a>	<a href="#">fn3.22.aSep08</a>		1956	824		274	titin N2-B (fn3.22) mRNA.
<a href="#">fn3.24</a>	<a href="#">fn3.24.aSep08</a>		6423	971		148	fibronectin, type III (fn3.24) mRNA.
<a href="#">fn3.25</a>	<a href="#">fn3.25.aSep08</a>		83432	1069	9	189	fibronectin, type III (fn3.25) alternative variant aSep08, mRNA.
<a href="#">fn3.25</a>	<a href="#">fn3.25.bSep08</a>		17052	368	2	45	putative protein (5.0 kD) (fn3.25) alternative variant bSep08, mRNA.
<a href="#">Fnbp1</a>	<a href="#">Fnbp1.bSep08</a>	<a href="#">192348</a>	20866	1218	6	278	formin binding protein 1 (Fnbp1) alternative variant bSep08, mRNA.
<a href="#">Fnbp1</a>	<a href="#">Fnbp1.dSep08</a>	<a href="#">192348</a>	10057	2575	5	185	formin binding protein 1 (Fnbp1) alternative variant dSep08, mRNA.
<a href="#">Fnbp1</a>	<a href="#">Fnbp1.eSep08</a>	<a href="#">192348</a>	19510	515	5	171	formin binding protein 1 (Fnbp1) alternative variant eSep08, mRNA.
<a href="#">Fnbp1</a>	<a href="#">Fnbp1.fSep08</a>	<a href="#">192348</a>	3032	298	3	99	formin binding protein 1 (Fnbp1) alternative variant fSep08, mRNA.
<a href="#">Fnbp1l</a>	<a href="#">Fnbp1l.bSep08</a>	<a href="#">310839</a>	37645	612	2	90	formin binding protein 1-like (Fnbp1l) alternative variant bSep08, mRNA.
<a href="#">Fnbp1l</a>	<a href="#">Fnbp1l.cSep08</a>	<a href="#">310839</a>	2061	826	2	51	formin binding protein 1-like (5.8 kD) (Fnbp1l) alternative variant cSep08, mRNA.
<a href="#">Fnbp4</a>	<a href="#">Fnbp4.bSep08</a>	<a href="#">311183</a>	30343	3769	17	967	formin binding protein 4 (Fnbp4) alternative variant bSep08, mRNA.
<a href="#">Fnbp4</a>	<a href="#">Fnbp4.cSep08</a>	<a href="#">311183</a>	6229	485	5	147	formin binding protein 4 (Fnbp4) alternative variant cSep08, mRNA.
<a href="#">Fnbp4</a>	<a href="#">Fnbp4.eSep08</a>	<a href="#">311183</a>	1667	334	2	71	formin binding protein 4 (Fnbp4) alternative variant eSep08, mRNA.
<a href="#">Fndc3a</a>	<a href="#">Fndc3a.bSep08</a>	<a href="#">306022</a>	152666	1317	10	439	fibronectin, type III (Fndc3a) alternative variant bSep08, mRNA.
<a href="#">Fndc3a</a>	<a href="#">Fndc3a.cSep08</a>	<a href="#">306022</a>	4695	463	5	153	putative protein of ancient origin (Fndc3a) alternative variant cSep08, mRNA.
<a href="#">Fndc3a</a>	<a href="#">Fndc3a.dSep08</a>	<a href="#">306022</a>	130479	439	5	146	putative protein (Fndc3a) alternative variant dSep08, mRNA.
<a href="#">Fndc3a</a>	<a href="#">Fndc3a.eSep08</a>	<a href="#">306022</a>	12932	418	3	139	putative protein (Fndc3a) alternative variant eSep08, mRNA.
<a href="#">Fndc3b</a>	<a href="#">Fndc3b.aSep08</a>	<a href="#">294925</a>	7064	875	6	252	fibronectin, type III (Fndc3b) alternative variant aSep08, mRNA.
<a href="#">Fndc3b</a>	<a href="#">Fndc3b.bSep08</a>	<a href="#">294925</a>	22674	801	3	181	fibronectin, type III (Fndc3b) alternative variant bSep08, mRNA.
<a href="#">Fndc7</a>	<a href="#">Fndc7.bSep08</a>	<a href="#">310787</a>	7887	409	2	83	putative protein of vertebrate origin (Fndc7) alternative variant bSep08, mRNA.
<a href="#">Fntb</a>	<a href="#">Fntb.bSep08</a>	<a href="#">64511</a>	33937	756	7	251	farnesyltransferase, CAAX box, beta (Fntb) alternative variant bSep08, mRNA.
<a href="#">foby</a>	<a href="#">foby.aSep08</a>		3091	1389	2	117	putative protein (foby) alternative variant aSep08, mRNA.
<a href="#">foby</a>	<a href="#">foby.bSep08</a>		2952	547	3	71	putative protein (foby) alternative variant bSep08, mRNA.
<a href="#">foby</a>	<a href="#">foby.cSep08</a>		3085	505	4	79	putative protein (foby) alternative variant cSep08, mRNA.
<a href="#">foby</a>	<a href="#">foby.dSep08</a>		3120	456	3	61	putative protein (foby) alternative variant dSep08, mRNA.

<a href="#">fochy</a>	<a href="#">fochy.aSep08</a>		17057	768		83	putative cytoplasmic protein (9.5 kD) (fochy) mRNA.
<a href="#">fofer</a>	<a href="#">fofer.aSep08</a>		991	830		62	putative protein (fofer) mRNA.
<a href="#">foflo</a>	<a href="#">foflo.aSep08</a>		47944	415		31	putative protein (foflo) mRNA.
<a href="#">foflu</a>	<a href="#">foflu.aSep08</a>		4197	321		32	putative protein (3.4 kD) (foflu) mRNA.
<a href="#">fokee</a>	<a href="#">fokee.aSep08</a>		853	694		49	putative protein (5.3 kD) (fokee) mRNA.
<a href="#">Folh1</a>	<a href="#">Folh1.bSep08</a>	<a href="#">85309</a>	37471	691		183	folate hydrolase (Folh1) alternative variant bSep08, mRNA.
<a href="#">foloy</a>	<a href="#">foloy.aSep08</a>		1916	550		75	putative protein of vertebrate origin (foloy) mRNA.
<a href="#">Folr1</a>	<a href="#">Folr1.bSep08</a>	<a href="#">171049</a>	626	197	1	42	folate receptor 1 (adult) (Folr1) alternative variant bSep08, mRNA.
<a href="#">Folr2</a>	<a href="#">Folr2.bSep08</a>	<a href="#">293154</a>	5658	1785	2	6	folate receptor 2 (fetal) (0.6 kD) (Folr2) alternative variant bSep08, mRNA.
<a href="#">fomer</a>	<a href="#">fomer.aSep08</a>		702	333		35	putative protein (4.0 kD) (fomer) mRNA.
<a href="#">fonoy</a>	<a href="#">fonoy.aSep08</a>		499	291		96	sema domain immunoglobulin transmembrane short cytoplasmic 4C CRA a (fonoy) mRNA.
<a href="#">fopor</a>	<a href="#">fopor.aSep08</a>		815	501		88	putative protein (10.0 kD) (fopor) mRNA.
<a href="#">forby</a>	<a href="#">forby.aSep08</a>		22702	724		240	retinitis pigmentosa GTPase regulator (forby) mRNA.
<a href="#">forchy</a>	<a href="#">forchy.aSep08</a>		2135	395		131	lymphocyte antigen 75 like (forchy) mRNA.
<a href="#">forfer</a>	<a href="#">forfer.aSep08</a>		28792	831		276	leucine rich repeat containing 16A (forfer) mRNA.
<a href="#">forflo</a>	<a href="#">forflo.aSep08</a>		14150	846		35	putative protein (4.0 kD) (forflo) mRNA.
<a href="#">forflu</a>	<a href="#">forflu.aSep08</a>		1280	262		65	putative protein (forflu) mRNA.
<a href="#">forkee</a>	<a href="#">forkee.aSep08</a>		2953	1026		154	putative mitochondrial protein (16.7 kD) (forkee) mRNA.
<a href="#">forloy</a>	<a href="#">forloy.aSep08</a>		4309	421		110	sid1 transmembrane family member 1 (forloy) mRNA.
<a href="#">former</a>	<a href="#">former.aSep08</a>		724	607		51	putative protein (former) mRNA.
<a href="#">fornoy</a>	<a href="#">fornoy.aSep08</a>		2183	355		118	transmembrane protein 131 (fornoy) mRNA.
<a href="#">forpor</a>	<a href="#">forpor.aSep08</a>		6940	186		54	putative protein (forpor) mRNA.
<a href="#">forsa</a>	<a href="#">forsa.aSep08</a>		207762	469		76	oxidation resistance 1 (forsa) alternative variant aSep08, mRNA.
<a href="#">forsa</a>	<a href="#">forsa.bSep08</a>		207717	483	1	74	oxidation resistance 1 like (forsa) alternative variant bSep08, mRNA.
<a href="#">forsa</a>	<a href="#">forsa.cSep08</a>		207687	641	1	71	putative protein (forsa) alternative variant cSep08, mRNA.
<a href="#">forshee</a>	<a href="#">forshee.aSep08</a>		1102	718	1	161	gag-pro-pol polyprotein (17.7 kD) (forshee) alternative variant aSep08, mRNA.
<a href="#">forshee</a>	<a href="#">forshee.bSep08</a>		5075	1966	1	149	putative protein (16.5 kD) (forshee) alternative variant bSep08, complete mRNA.
<a href="#">forshee</a>	<a href="#">forshee.cSep08</a>		2622	705	2	74	putative nuclear protein (8.6 kD) (forshee) alternative variant cSep08, complete mRNA.
<a href="#">forto</a>	<a href="#">forto.aSep08</a>		571	446		73	putative nuclear protein (8.9 kD) (forto) mRNA.
<a href="#">forvar</a>	<a href="#">forvar.aSep08</a>		3495	701		81	putative protein (9.4 kD) (forvar) mRNA.
<a href="#">forwey</a>	<a href="#">forwey.aSep08</a>		14056	846	5	236	glutamine transaminase 1 CRA a (forwey) alternative variant aSep08, mRNA.
<a href="#">forwey</a>	<a href="#">forwey.bSep08</a>		1694	241	2	80	transaminase 1 CRA b (forwey) alternative variant bSep08, mRNA.
<a href="#">fosa</a>	<a href="#">fosa.aSep08</a>		12156	1768		66	putative protein (fosa) mRNA.

<a href="#">foshee</a>	<a href="#">foshee.aSep08</a>		363	140		46	membrane-associated ring finger 6 CRA a (foshee) mRNA.
<a href="#">foto</a>	<a href="#">foto.aSep08</a>		3222	1476	6	263	putative protein, with a coiled coil domain (foto) alternative variant aSep08, mRNA.
<a href="#">foto</a>	<a href="#">foto.cSep08</a>		810	716	2	106	putative protein of vertebrate origin (foto) alternative variant cSep08, mRNA.
<a href="#">fovar</a>	<a href="#">fovar.aSep08</a>		9272	490		49	putative protein (5.7 kD) (fovar) mRNA.
<a href="#">fowey</a>	<a href="#">fowey.aSep08</a>		8051	237		66	putative protein (fowey) mRNA.
<a href="#">Foxa2</a>	<a href="#">Foxa2.aSep08</a>	<a href="#">25099</a>	3019	2010		450	forkhead box A2 (Foxa2) mRNA.
<a href="#">Foxh1</a>	<a href="#">Foxh1.aSep08</a>	<a href="#">300054</a>	2031	1629	1	373	forkhead box H1 (Foxh1) alternative variant aSep08, mRNA.
<a href="#">Foxh1</a>	<a href="#">Foxh1.bSep08</a>	<a href="#">300054</a>	1116	756	1	252	forkhead box H1 (Foxh1) alternative variant bSep08, mRNA.
<a href="#">Foxk2</a>	<a href="#">Foxk2.aSep08</a>	<a href="#">303753</a>	23863	2154	8	514	forkhead box K2 (Foxk2) alternative variant aSep08, mRNA.
<a href="#">Foxk2</a>	<a href="#">Foxk2.cSep08</a>	<a href="#">303753</a>	17612	1782	6	397	forkhead box K2 (Foxk2) alternative variant cSep08, mRNA.
<a href="#">Foxk2</a>	<a href="#">Foxk2.dSep08</a>	<a href="#">303753</a>	30678	535	2	98	forkhead box K2 (Foxk2) alternative variant dSep08, mRNA.
<a href="#">Foxm1</a>	<a href="#">Foxm1.bSep08</a>	<a href="#">58921</a>	3457	485	5	161	forkhead box M1 CRA a (Foxm1) alternative variant bSep08, mRNA.
<a href="#">Foxm1</a>	<a href="#">Foxm1.cSep08</a>	<a href="#">58921</a>	9929	872	2	112	putative protein (12.2 kD) (Foxm1) alternative variant cSep08, mRNA.
<a href="#">Foxn3</a>	<a href="#">Foxn3.bSep08</a>	<a href="#">314374</a>	4106	545	2	133	forkhead box N3 (Foxn3) alternative variant bSep08, mRNA.
<a href="#">Foxn3</a>	<a href="#">Foxn3.cSep08</a>	<a href="#">314374</a>	4084	587	3	72	forkhead box N3 (Foxn3) alternative variant cSep08, mRNA.
<a href="#">Foxn3</a>	<a href="#">Foxn3.dSep08</a>	<a href="#">314374</a>	84990	774	4	62	forkhead box N3 (7.3 kD) (Foxn3) alternative variant dSep08, mRNA.
<a href="#">Foxo1</a>	<a href="#">Foxo1.aSep08</a>	<a href="#">84482</a>	73563	641		213	forkhead box O1 (Foxo1) mRNA.
<a href="#">Foxo3a</a>	<a href="#">Foxo3a.bSep08</a>	<a href="#">294515</a>	1188	386	3	94	forkhead box O3a (Foxo3a) alternative variant bSep08, mRNA.
<a href="#">Foxp1</a>	<a href="#">Foxp1.bSep08</a>	<a href="#">297480</a>	37746	2244	9	329	forkhead box P1 (Foxp1) alternative variant bSep08, mRNA.
<a href="#">Foxp1</a>	<a href="#">Foxp1.cSep08</a>	<a href="#">297480</a>	49070	727	6	242	forkhead box P1 (Foxp1) alternative variant cSep08, mRNA.
<a href="#">Foxp1</a>	<a href="#">Foxp1.dSep08</a>	<a href="#">297480</a>	264071	773	6	187	forkhead box P1 (Foxp1) alternative variant dSep08, mRNA.
<a href="#">Foxp2</a>	<a href="#">Foxp2.aSep08</a>	<a href="#">500037</a>	31479	1965		227	forkhead box P2 (Foxp2) alternative variant aSep08, mRNA.
<a href="#">Foxp2</a>	<a href="#">Foxp2.bSep08</a>	<a href="#">500037</a>	24935	723		85	forkhead box P2 (Foxp2) alternative variant bSep08, mRNA.
<a href="#">Foxp4</a>	<a href="#">Foxp4.bSep08</a>	<a href="#">363185</a>	2976	2217	3	120	forkhead box P4 CRA a (12.4 kD) (Foxp4) alternative variant bSep08, mRNA.
<a href="#">Foxp4</a>	<a href="#">Foxp4.cSep08</a>	<a href="#">363185</a>	2345	420	3	72	putative protein (8.2 kD) (Foxp4) alternative variant cSep08, mRNA.

<a href="#">Foxr1</a>	<a href="#">Foxr1.aSep08</a>	<a href="#">315601</a>	1999	397	3	119	forkhead box R1 (Foxr1) alternative variant aSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.aSep08</a>	<a href="#">315547</a>	5667	969	8	306	FAD dependent oxidoreductase (Foxred1) alternative variant aSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.bSep08</a>	<a href="#">315547</a>	7418	2177	11	299	FAD dependent oxidoreductase (33.4 kD) (Foxred1) alternative variant bSep08, complete mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.cSep08</a>	<a href="#">315547</a>	5442	846	7	271	FAD-dependent oxidoreductase domain-containing protein 1 (Foxred1) alternative variant cSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.dSep08</a>	<a href="#">315547</a>	5346	746	7	175	FAD-dependent oxidoreductase domain-containing protein 1 (Foxred1) alternative variant dSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.eSep08</a>	<a href="#">315547</a>	1145	393	2	116	putative protein of vertebrate origin (Foxred1) alternative variant eSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.fSep08</a>	<a href="#">315547</a>	3439	684	5	98	putative protein of ancient origin (Foxred1) alternative variant fSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.hSep08</a>	<a href="#">315547</a>	3963	1013	5	90	putative protein of ancient origin (Foxred1) alternative variant hSep08, mRNA.
<a href="#">Foxred1</a>	<a href="#">Foxred1.iSep08</a>	<a href="#">315547</a>	2328	772	4	55	putative protein of ancient origin (Foxred1) alternative variant iSep08, mRNA.
<a href="#">Foxred2</a>	<a href="#">Foxred2.aSep08</a>	<a href="#">315112</a>	2847	380		126	putative protein of metazoan origin (Foxred2) mRNA.
<a href="#">foyby</a>	<a href="#">foyby.aSep08</a>		4088	688	1	76	retinitis pigmentosa GTPase regulator (foyby) alternative variant aSep08, mRNA.
<a href="#">foyby</a>	<a href="#">foyby.bSep08</a>		6001	430	2	43	retinitis pigmentosa GTPase regulator (foyby) alternative variant bSep08, mRNA.
<a href="#">foychy</a>	<a href="#">foychy.aSep08</a>		12994	669		106	phospholipase A2 receptor 1 (foychy) mRNA.
<a href="#">foyer</a>	<a href="#">foyer.aSep08</a>		27619	616	5	205	leucine rich repeat containing 16A (foyer) alternative variant aSep08, mRNA.
<a href="#">foyflo</a>	<a href="#">foyflo.aSep08</a>		5348	740		84	putative nuclear protein (9.6 kD) (foyflo) mRNA.
<a href="#">foyflu</a>	<a href="#">foyflu.aSep08</a>		19872	707		32	putative protein (3.7 kD) (foyflu) mRNA.
<a href="#">foykee</a>	<a href="#">foykee.aSep08</a>		1528	353		117	lactase-phlorizin hydrolase (foykee) mRNA.
<a href="#">foyloy</a>	<a href="#">foyloy.aSep08</a>		10896	341		15	putative protein (1.7 kD) (foyloy) mRNA.
<a href="#">foymee</a>	<a href="#">foymee.aSep08</a>		13263	743		139	ATP-binding cassette sub-family C member 1 like (foymee) mRNA.
<a href="#">foymer</a>	<a href="#">foymer.aSep08</a>		4240	677		38	putative protein (foymer) mRNA.
<a href="#">foynoy</a>	<a href="#">foynoy.aSep08</a>		17950	710		86	CRA a (foynoy) mRNA.
<a href="#">foypor</a>	<a href="#">foypor.aSep08</a>		43220	534		39	putative protein (4.1 kD) (foypor) mRNA.
<a href="#">foysa</a>	<a href="#">foysa.aSep08</a>		1091	479		53	putative protein (foysa) mRNA.
<a href="#">foyshee</a>	<a href="#">foyshee.aSep08</a>		83856	432		94	putative protein (10.7 kD) (foyshee) mRNA.
<a href="#">foyto</a>	<a href="#">foyto.aSep08</a>		2686	759		123	putative protein of mammalian origin (foyto) mRNA.
<a href="#">foyvar</a>	<a href="#">foyvar.aSep08</a>		13029	801		29	putative protein (3.0 kD) (foyvar) mRNA.
<a href="#">foywey</a>	<a href="#">foywey.aSep08</a>		2201	720		89	putative protein (10.2 kD) (foywey) mRNA.
<a href="#">Fpgt</a>	<a href="#">Fpgt.bSep08</a>	<a href="#">310935</a>	20822	1721	2	137	fucose-1-phosphate guanylyltransferase (15.4 kD) (Fpgt) alternative variant bSep08, complete mRNA.
<a href="#">Frag1</a>	<a href="#">Frag1.bSep08</a>	<a href="#">116675</a>	7911	1949	6	281	FGF receptor activating protein 1 (Frag1) alternative variant bSep08, mRNA.

<a href="#">Frag1</a>	<a href="#">Frag1.cSep08</a>	<a href="#">116675</a>	12558	717	4	238	FGF receptor activating protein 1 (Frag1) alternative variant cSep08, mRNA.
<a href="#">Frag1</a>	<a href="#">Frag1.dSep08</a>	<a href="#">116675</a>	12806	688	5	228	FGF receptor activating protein 1 (Frag1) alternative variant dSep08, mRNA.
<a href="#">Frag1</a>	<a href="#">Frag1.eSep08</a>	<a href="#">116675</a>	1600	759	4	164	FGF receptor activating protein 1 (Frag1) alternative variant eSep08, mRNA.
<a href="#">Frag1</a>	<a href="#">Frag1.fSep08</a>	<a href="#">116675</a>	12836	717	4	69	FGF receptor activating protein 1 (Frag1) alternative variant fSep08, mRNA.
<a href="#">Frag1</a>	<a href="#">Frag1.gSep08</a>	<a href="#">116675</a>	11814	580	2	35	FGF receptor activating protein 1 (3.8 kD) (Frag1) alternative variant gSep08, mRNA.
<a href="#">Fras1</a>	<a href="#">Fras1.aSep08</a>	<a href="#">289486</a>	30484	1781		387	fraser syndrome 1 homolog (human) (Fras1) mRNA.
<a href="#">Frem1</a>	<a href="#">Frem1.aSep08</a>	<a href="#">298185</a>	14851	2242		309	fras1 related extracellular matrix 1 (Frem1) mRNA.
<a href="#">Frem2</a>	<a href="#">Frem2.aSep08</a>	<a href="#">310418</a>	23669	373		124	fras1 related extracellular matrix protein 2 (Frem2) mRNA.
<a href="#">Frem3</a>	<a href="#">Frem3.aSep08</a>	<a href="#">307767</a>	5145	577		118	fras1 related extracellular matrix protein 3 (Frem3) mRNA.
<a href="#">Frmd4a</a>	<a href="#">Frmd4a.aSep08</a>	<a href="#">307128</a>	9425	1102		366	putative protein of vertebrate origin (Frmd4a) mRNA.
<a href="#">Frmd4b</a>	<a href="#">Frmd4b.aSep08</a>	<a href="#">252858</a>	16986	408		135	putative protein of eukaryotic origin (Frmd4b) mRNA.
<a href="#">Frmd5</a>	<a href="#">Frmd5.aSep08</a>	<a href="#">311362</a>	231937	435		110	putative protein of metazoan origin (Frmd5) mRNA.
<a href="#">Frmd6</a>	<a href="#">Frmd6.aSep08</a>	<a href="#">257646</a>	50889	494		82	putative protein (Frmd6) mRNA.
<a href="#">Frmd8</a>	<a href="#">Frmd8.bSep08</a>	<a href="#">309172</a>	20583	1784	7	354	putative protein of eukaryotic origin (Frmd8) alternative variant bSep08, mRNA.
<a href="#">Frmd8</a>	<a href="#">Frmd8.cSep08</a>	<a href="#">309172</a>	846	302	2	81	putative protein of vertebrate origin (Frmd8) alternative variant cSep08, mRNA.
<a href="#">Frmpd1</a>	<a href="#">Frmpd1.aSep08</a>	<a href="#">313244</a>	6304	3185	2	1003	putative protein of vertebrate origin (Frmpd1) alternative variant aSep08, mRNA.
<a href="#">Frmpd1</a>	<a href="#">Frmpd1.cSep08</a>	<a href="#">313244</a>	4816	409	2	103	putative protein of metazoan origin (Frmpd1) alternative variant cSep08, mRNA.
<a href="#">Frrs1</a>	<a href="#">Frrs1.aSep08</a>	<a href="#">310810</a>	19076	414		79	ferric-chelate reductase 1 (Frrs1) mRNA.
<a href="#">Frs3</a>	<a href="#">Frs3.bSep08</a>	<a href="#">316213</a>	1324	336	3	111	fibroblast growth factor receptor substrate 3 (Frs3) alternative variant bSep08, mRNA.
<a href="#">Fscn2</a>	<a href="#">Fscn2.bSep08</a>	<a href="#">303741</a>	1667	1157	1	280	fascin homolog 2, actin-bundling protein, retinal (Strongylocentrotus purpuratus) (Fscn2) alternative variant bSep08, mRNA.
<a href="#">Fsd1</a>	<a href="#">Fsd1.aSep08</a>	<a href="#">301506</a>	6471	1259		336	fibronectin type 3 and SPRY domain-containing protein (Fsd1) mRNA.
<a href="#">Fsd2</a>	<a href="#">Fsd2.aSep08</a>	<a href="#">308779</a>	7946	1258		189	SPla/Ryanodine receptor SPRY (Fsd2) mRNA.
<a href="#">Fsip1</a>	<a href="#">Fsip1.bSep08</a>	<a href="#">296074</a>	19265	955	4	317	fibrous sheath interacting protein 1 (Fsip1) alternative variant bSep08, mRNA.
<a href="#">Fsip1</a>	<a href="#">Fsip1.cSep08</a>	<a href="#">296074</a>	13121	1062	4	204	fibrous sheath interacting protein 1 (22.9 kD) (Fsip1) alternative variant cSep08, mRNA.
<a href="#">Fsip1</a>	<a href="#">Fsip1.dSep08</a>	<a href="#">296074</a>	5125	586	1	195	fibrous sheath interacting protein 1 (Fsip1) alternative variant dSep08, mRNA.
<a href="#">Fstl4</a>	<a href="#">Fstl4.bSep08</a>	<a href="#">303130</a>	56248	877	2	292	folliculin-like 4 (Fstl4) alternative variant bSep08, mRNA.
<a href="#">Ftcd</a>	<a href="#">Ftcd.bSep08</a>	<a href="#">89833</a>	6184	910	5	169	formiminotransferase cyclodeaminase CRA b (18.6 kD) (Ftcd) alternative variant bSep08, mRNA.
<a href="#">Ftcd</a>	<a href="#">Ftcd.cSep08</a>	<a href="#">89833</a>	3257	764	3	151	formiminotransferase cyclodeaminase CRA b (Ftcd) alternative variant cSep08, mRNA.

<a href="#">Ftcd</a>	<a href="#">Ftcd.dSep08</a>	<a href="#">89833</a>	3520	720	4	122	formiminotransferase cyclodeaminase CRA a (Ftcd) alternative variant dSep08, mRNA.
<a href="#">Ftcd</a>	<a href="#">Ftcd.eSep08</a>	<a href="#">89833</a>	6835	1070	3	76	formiminotransferase cyclodeaminase CRA c (8.4 kD) (Ftcd) alternative variant eSep08, mRNA.
<a href="#">Ftl1</a>	<a href="#">Ftl1.bSep08</a>	<a href="#">29292</a>	1052	764	2	101	ferritin light (11.6 kD) (Ftl1) alternative variant bSep08, mRNA.
<a href="#">Ftl1</a>	<a href="#">Ftl1.cSep08</a>	<a href="#">29292</a>	1459	815	3	87	ferritin light (9.7 kD) (Ftl1) alternative variant cSep08, mRNA.
<a href="#">Fto</a>	<a href="#">Fto.bSep08</a>	<a href="#">291905</a>	182710	792	2	106	fat mass and obesity associated (12.4 kD) (Fto) alternative variant bSep08, mRNA.
<a href="#">Fts</a>	<a href="#">Fts.bSep08</a>	<a href="#">291906</a>	9141	971	10	272	fused toes (Fts) alternative variant bSep08, mRNA.
<a href="#">Fts</a>	<a href="#">Fts.cSep08</a>	<a href="#">291906</a>	6800	622	5	124	fused toes (Fts) alternative variant cSep08, mRNA.
<a href="#">Fts</a>	<a href="#">Fts.dSep08</a>	<a href="#">291906</a>	7726	740	7	107	fused toes (Fts) alternative variant dSep08, mRNA.
<a href="#">Ftsj1</a>	<a href="#">Ftsj1.aSep08</a>	<a href="#">363450</a>	3199	1005	6	193	ftsj homolog 1 (E. coli) and similar to ribosomal protein L31 (Ftsj1) alternative variant aSep08, mRNA.
<a href="#">Ftsj1</a>	<a href="#">Ftsj1.aSep08</a>	<a href="#">679519</a>	3199	1005	6	193	ftsj homolog 1 (E. coli) and similar to ribosomal protein L31 (Ftsj1) alternative variant aSep08, mRNA.
<a href="#">Ftsj1</a>	<a href="#">Ftsj1.bSep08</a>	<a href="#">363450</a>	1820	828	4	186	ftsj homolog 1 (E. coli) and similar to ribosomal protein L31 (20.8 kD) (Ftsj1) alternative variant bSep08, mRNA.
<a href="#">Ftsj1</a>	<a href="#">Ftsj1.bSep08</a>	<a href="#">679519</a>	1820	828	4	186	ftsj homolog 1 (E. coli) and similar to ribosomal protein L31 (20.8 kD) (Ftsj1) alternative variant bSep08, mRNA.
<a href="#">Ftsj2</a>	<a href="#">Ftsj2.bSep08</a>	<a href="#">304323</a>	543	304	1	46	FtsJ homolog 2 (E. coli) (Ftsj2) alternative variant bSep08, mRNA.
<a href="#">Ftsj3</a>	<a href="#">Ftsj3.bSep08</a>	<a href="#">303608</a>	1080	772	1	169	FtsJ homolog 3 (E. coli) (Ftsj3) alternative variant bSep08, mRNA.
<a href="#">Fubp1</a>	<a href="#">Fubp1.aSep08</a>	<a href="#">654496</a>	5587	1045	1	169	far upstream element (FUZE) binding protein 1 (Fubp1) alternative variant aSep08, mRNA.
<a href="#">Fubp1</a>	<a href="#">Fubp1.bSep08</a>	<a href="#">654496</a>	4890	357	1	119	far upstream element (FUZE) binding protein 1 (Fubp1) alternative variant bSep08, mRNA.
<a href="#">Fubp3</a>	<a href="#">Fubp3.bSep08</a>	<a href="#">362106</a>	28177	420	6	140	far upstream element (FUZE) binding protein 3 (Fubp3) alternative variant bSep08, mRNA.
<a href="#">Fubp3</a>	<a href="#">Fubp3.cSep08</a>	<a href="#">362106</a>	4659	954	7	115	far upstream element (FUZE) binding protein 3 (12.4 kD) (Fubp3) alternative variant cSep08, mRNA.
<a href="#">Fubp3</a>	<a href="#">Fubp3.dSep08</a>	<a href="#">362106</a>	3997	685	4	105	far upstream element (FUZE) binding protein 3 (Fubp3) alternative variant dSep08, mRNA.
<a href="#">fuby</a>	<a href="#">fuby.aSep08</a>		14931	396		132	ubiquitin specific peptidase 9 X-linked CRA a (fuby) mRNA.
<a href="#">Fuca1</a>	<a href="#">Fuca1.bSep08</a>	<a href="#">24375</a>	9862	1159	4	185	fucosidase, alpha-L- 1, tissue (Fuca1) alternative variant bSep08, mRNA.
<a href="#">Fuca2</a>	<a href="#">Fuca2.cSep08</a>	<a href="#">292485</a>	7260	410	2	44	fucosidase, alpha-L- 2, plasma (Fuca2) alternative variant cSep08, mRNA.
<a href="#">fuchy</a>	<a href="#">fuchy.aSep08</a>		30394	752		44	putative protein (fuchy) mRNA.
<a href="#">fufer</a>	<a href="#">fufer.aSep08</a>		3576	584		58	putative protein (fufer) mRNA.
<a href="#">fuflo</a>	<a href="#">fuflo.aSep08</a>		1829	346		48	putative protein of mammalian origin (fuflo) mRNA.
<a href="#">fuflu</a>	<a href="#">fuflu.aSep08</a>		1074	627	2	90	putative protein (fuflu) alternative variant aSep08, mRNA.
<a href="#">Fuk</a>	<a href="#">Fuk.bSep08</a>	<a href="#">307848</a>	4412	1361	6	281	fucokinase (Fuk) alternative variant bSep08, mRNA.
<a href="#">Fuk</a>	<a href="#">Fuk.cSep08</a>	<a href="#">307848</a>	3834	332	3	99	fucokinase (Fuk) alternative variant cSep08, mRNA.

<a href="#">fukee</a>	<a href="#">fukee.aSep08</a>		2793	376		53	putative protein (fukee) mRNA.
<a href="#">fuloy</a>	<a href="#">fuloy.aSep08</a>		17880	452		51	putative protein of mammalian origin (fuloy) mRNA.
<a href="#">fumer</a>	<a href="#">fumer.aSep08</a>		4163	418		38	putative protein (fumer) mRNA.
<a href="#">funoy</a>	<a href="#">funoy.aSep08</a>		415	266		64	sema domain immunoglobulin transmembrane short cytoplasmic 4C CRA a (funoy) mRNA.
<a href="#">fupor</a>	<a href="#">fupor.aSep08</a>		86681	821		56	putative protein (6.2 kD) (fupor) mRNA.
<a href="#">Furin</a>	<a href="#">Furin.bSep08</a>	<a href="#">54281</a>	7629	435	1	57	furin (paired basic amino acid cleaving enzyme) (Furin) alternative variant bSep08, mRNA.
<a href="#">Furin-like.0</a>	<a href="#">Furin-like.0.aSep08</a>		11994	757		193	epidermal growth factor receptor CRA b (Furin-like.0) mRNA.
<a href="#">Fus</a>	<a href="#">Fus.bSep08</a>	<a href="#">317385</a>	7996	870	8	269	fusion, derived from t(12;16) malignant liposarcoma (human) (Fus) alternative variant bSep08, mRNA.
<a href="#">Fus</a>	<a href="#">Fus.dSep08</a>	<a href="#">317385</a>	1982	807	3	110	fusion, derived from t(12;16) malignant liposarcoma (human) (Fus) alternative variant dSep08, mRNA.
<a href="#">Fus</a>	<a href="#">Fus.eSep08</a>	<a href="#">317385</a>	3006	1621	2	55	fusion, derived from t(12;16) malignant liposarcoma (human) (6.2 kD) (Fus) alternative variant eSep08, mRNA.
<a href="#">fusa</a>	<a href="#">fusa.aSep08</a>		4865	351		44	putative protein (4.7 kD) (fusa) mRNA.
<a href="#">fushee</a>	<a href="#">fushee.aSep08</a>		1403	796		50	putative protein (5.5 kD) (fushee) mRNA.
<a href="#">Fusip1</a>	<a href="#">Fusip1.aSep08</a>	<a href="#">362630</a>	8408	1068	6	262	FUS interacting protein (serine-arginine rich) 1 (31.3 kD) (Fusip1) alternative variant aSep08, mRNA.
<a href="#">Fusip1</a>	<a href="#">Fusip1.bSep08</a>	<a href="#">362630</a>	10005	2662	6	261	FUS interacting protein (serine-arginine rich) 1 (31.2 kD) (Fusip1) alternative variant bSep08, complete mRNA.
<a href="#">Fusip1</a>	<a href="#">Fusip1.cSep08</a>	<a href="#">362630</a>	13330	2606	6	183	FUS interacting protein (serine-arginine rich) 1 (22.2 kD) (Fusip1) alternative variant cSep08, complete mRNA.
<a href="#">Fusip1</a>	<a href="#">Fusip1.dSep08</a>	<a href="#">362630</a>	11414	687	6	182	FUS interacting protein (serine-arginine rich) 1 (22.1 kD) (Fusip1) alternative variant dSep08, mRNA.
<a href="#">Fusip1</a>	<a href="#">Fusip1.fSep08</a>	<a href="#">362630</a>	1790	428	2	88	FUS interacting protein (serine-arginine rich) 1 (Fusip1) alternative variant fSep08, mRNA.
<a href="#">Fut2</a>	<a href="#">Fut2.aSep08</a>	<a href="#">58924</a>	18910	1276	3	284	fucosyltransferase 2 (secretor status included) (Fut2) alternative variant aSep08, mRNA.
<a href="#">Fut2</a>	<a href="#">Fut2.cSep08</a>	<a href="#">58924</a>	16629	800	4	163	fucosyltransferase 2 (secretor status included) (Fut2) alternative variant cSep08, mRNA.
<a href="#">Fut2</a>	<a href="#">Fut2.dSep08</a>	<a href="#">58924</a>	16427	595	4	71	fucosyltransferase 2 (secretor status included) (Fut2) alternative variant dSep08, mRNA.
<a href="#">Fut2</a>	<a href="#">Fut2.eSep08</a>	<a href="#">58924</a>	16217	375	4	41	fucosyltransferase 2 (secretor status included) (Fut2) alternative variant eSep08, mRNA.
<a href="#">futo</a>	<a href="#">futo.aSep08</a>		2369	559		185	putative protein of vertebrate origin (futo) mRNA.
<a href="#">fuvar</a>	<a href="#">fuvar.aSep08</a>		1204	790		32	putative protein (3.5 kD) (fuvar) mRNA.
<a href="#">fuwey</a>	<a href="#">fuwey.aSep08</a>		639	291		30	putative protein (3.5 kD) (fuwey) mRNA.
<a href="#">Fuz</a>	<a href="#">Fuz.aSep08</a>	<a href="#">308577</a>	2121	1224	4	407	fuzzy homolog (Drosophila) (Fuz) alternative variant aSep08, mRNA.
<a href="#">Fuz</a>	<a href="#">Fuz.cSep08</a>	<a href="#">308577</a>	2917	812	6	223	fuzzy homolog (Drosophila) (Fuz) alternative variant cSep08, mRNA.
<a href="#">Fuz</a>	<a href="#">Fuz.dSep08</a>	<a href="#">308577</a>	757	402	3	123	fuzzy homolog (Drosophila) (Fuz) alternative variant dSep08, mRNA.



<a href="#">Fxc1</a>	<a href="#">Fxc1.aSep08</a>	<a href="#">84384</a>	2847	670	4	100	fractured callus expressed transcript 1 (11.4 kD) (Fxc1) alternative variant aSep08, complete mRNA.
<a href="#">Fxn</a>	<a href="#">Fxn.aSep08</a>	<a href="#">499335</a>	23885	668	5	145	frataxin (Fxn) alternative variant aSep08, mRNA.
<a href="#">Fxr1</a>	<a href="#">Fxr1.aSep08</a>	<a href="#">361927</a>	44972	1268	13	367	fragile X mental retardation, autosomal homolog 1 (Fxr1) alternative variant aSep08, mRNA.
<a href="#">Fxr1</a>	<a href="#">Fxr1.bSep08</a>	<a href="#">361927</a>	2495	422	2	131	fragile X mental retardation, autosomal homolog 1 (Fxr1) alternative variant bSep08, mRNA.
<a href="#">Fxr1</a>	<a href="#">Fxr1.cSep08</a>	<a href="#">361927</a>	1250	640	2	77	fragile X mental retardation, autosomal homolog 1 (8.6 kD) (Fxr1) alternative variant cSep08, mRNA.
<a href="#">Fxr2</a>	<a href="#">Fxr2.bSep08</a>	<a href="#">287433</a>	1790	1167	5	280	fragile X mental retardation 2 (30.4 kD) (Fxr2) alternative variant bSep08, mRNA.
<a href="#">Fxr2</a>	<a href="#">Fxr2.cSep08</a>	<a href="#">287433</a>	10628	1952	8	219	fragile X mental retardation autosomal homolog 2 (Fxr2) alternative variant cSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.aSep08</a>	<a href="#">58971</a>	3283	615	7	114	fxyd domain-containing ion transport regulator 1 CRA c (12.6 kD) (Fxyd1andFxyd7) alternative variant aSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.aSep08</a>	<a href="#">63848</a>	3283	615	7	114	fxyd domain-containing ion transport regulator 1 CRA c (12.6 kD) (Fxyd1andFxyd7) alternative variant aSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.bSep08</a>	<a href="#">58971</a>	1506	1243	2	108	FXYP domain-containing ion transport regulator 1 (11.7 kD) (Fxyd1andFxyd7) alternative variant bSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.bSep08</a>	<a href="#">63848</a>	1506	1243	2	108	FXYP domain-containing ion transport regulator 1 (11.7 kD) (Fxyd1andFxyd7) alternative variant bSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.dSep08</a>	<a href="#">58971</a>	4085	434	6	104	FXYP domain-containing ion transport regulator 1 (Fxyd1andFxyd7) alternative variant dSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.dSep08</a>	<a href="#">63848</a>	4085	434	6	104	FXYP domain-containing ion transport regulator 1 (Fxyd1andFxyd7) alternative variant dSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.eSep08</a>	<a href="#">58971</a>	4353	777	7	95	fxyd domain-containing ion transport regulator 1 CRA c (10.6 kD) (Fxyd1andFxyd7) alternative variant eSep08, complete mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.eSep08</a>	<a href="#">63848</a>	4353	777	7	95	fxyd domain-containing ion transport regulator 1 CRA c (10.6 kD) (Fxyd1andFxyd7) alternative variant eSep08, complete mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.fSep08</a>	<a href="#">58971</a>	3126	370	7	92	fxyd domain-containing ion transport regulator 1 CRA c (Fxyd1andFxyd7) alternative variant fSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.fSep08</a>	<a href="#">63848</a>	3126	370	7	92	fxyd domain-containing ion transport regulator 1 CRA c (Fxyd1andFxyd7) alternative variant fSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.gSep08</a>	<a href="#">58971</a>	2529	375	4	56	FXYP (Fxyd1andFxyd7) alternative variant gSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.gSep08</a>	<a href="#">63848</a>	2529	375	4	56	FXYP (Fxyd1andFxyd7) alternative variant gSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.hSep08</a>	<a href="#">58971</a>	3115	294	6	50	FXYP domain-containing ion transport regulator 1 (5.4 kD) (Fxyd1andFxyd7) alternative variant hSep08, mRNA.
<a href="#">Fxyd1andFxyd7</a>	<a href="#">Fxyd1andFxyd7.hSep08</a>	<a href="#">63848</a>	3115	294	6	50	FXYP domain-containing ion transport regulator 1 (5.4 kD) (Fxyd1andFxyd7) alternative variant hSep08, mRNA.
<a href="#">Fxyd2</a>	<a href="#">Fxyd2.aSep08</a>	<a href="#">29639</a>	2017	407	5	75	FXYP domain-containing ion transport regulator 2 (Fxyd2) alternative variant aSep08, mRNA.

<a href="#">Fxd2</a>	<a href="#">Fxd2.bSep08</a>	<a href="#">29639</a>	4259	595	6	74	FXD domain-containing ion transport regulator 2 (Fxd2) alternative variant bSep08, mRNA.
<a href="#">Fxd2</a>	<a href="#">Fxd2.dSep08</a>	<a href="#">29639</a>	4715	928	6	66	FXD domain-containing ion transport regulator 2 (7.3 kD) (Fxd2) alternative variant dSep08, mRNA.
<a href="#">Fxd3</a>	<a href="#">Fxd3.bSep08</a>	<a href="#">116831</a>	881	464	2	56	FXD domain-containing ion transport regulator 3 (Fxd3) alternative variant bSep08, mRNA.
<a href="#">Fxd4</a>	<a href="#">Fxd4.aSep08</a>	<a href="#">64190</a>	1556	370		46	FXD domain-containing ion transport regulator 4 (4.7 kD) (Fxd4) mRNA.
<a href="#">Fxd5</a>	<a href="#">Fxd5.aSep08</a>	<a href="#">60338</a>	9388	820	9	188	FXD domain-containing ion transport regulator 5 (Fxd5) alternative variant aSep08, mRNA.
<a href="#">Fxd6</a>	<a href="#">Fxd6.aSep08</a>	<a href="#">63847</a>	23218	531	1	169	FXD domain-containing ion transport regulator 6 (Fxd6) alternative variant aSep08, mRNA.
<a href="#">Fxd6</a>	<a href="#">Fxd6.cSep08</a>	<a href="#">63847</a>	23212	522	1	166	FXD domain-containing ion transport regulator 6 (Fxd6) alternative variant cSep08, mRNA.
<a href="#">Fyb</a>	<a href="#">Fyb.bSep08</a>	<a href="#">499537</a>	95065	1662	5	478	FYN binding protein (Fyb) alternative variant bSep08, mRNA.
<a href="#">Fyb</a>	<a href="#">Fyb.cSep08</a>	<a href="#">499537</a>	24426	1000	9	255	FYN binding protein (Fyb) alternative variant cSep08, mRNA.
<a href="#">Fyb</a>	<a href="#">Fyb.dSep08</a>	<a href="#">499537</a>	9903	405	5	134	FYN binding protein (Fyb) alternative variant dSep08, mRNA.
<a href="#">fyby</a>	<a href="#">fyby.aSep08</a>		4284	481		160	ubiquitin specific peptidase 9 X-linked CRA a (fyby) mRNA.
<a href="#">fyby</a>	<a href="#">fyby.aSep08</a>		46338	555		185	formin-like 2 CRA a (fyby) mRNA.
<a href="#">Fyco1</a>	<a href="#">Fyco1.bSep08</a>	<a href="#">301085</a>	26942	411	3	136	putative protein of vertebrate origin (Fyco1) alternative variant bSep08, mRNA.
<a href="#">Fyco1</a>	<a href="#">Fyco1.cSep08</a>	<a href="#">301085</a>	14240	442	4	67	putative protein, with a coiled coil domain (Fyco1) alternative variant cSep08, mRNA.
<a href="#">fyfer</a>	<a href="#">fyfer.aSep08</a>		25563	353		10	putative protein (1.1 kD) (fyfer) mRNA.
<a href="#">fyflo</a>	<a href="#">fyflo.aSep08</a>		35171	853	4	97	putative protein (fyflo) alternative variant aSep08, mRNA.
<a href="#">fyflo</a>	<a href="#">fyflo.bSep08</a>		25792	273	2	27	putative protein (3.0 kD) (fyflo) alternative variant bSep08, mRNA.
<a href="#">fyflu</a>	<a href="#">fyflu.aSep08</a>		2773	871		116	CRA a (fyflu) mRNA.
<a href="#">fykee</a>	<a href="#">fykee.aSep08</a>		3736	688		44	putative protein (fykee) mRNA.
<a href="#">fyloy</a>	<a href="#">fyloy.aSep08</a>		56651	377		61	heavy like (fyloy) mRNA.
<a href="#">fymer</a>	<a href="#">fymer.aSep08</a>		7391	554		67	putative protein (7.4 kD) (fymer) mRNA.
<a href="#">Fyn</a>	<a href="#">Fyn.bSep08</a>	<a href="#">25150</a>	54146	2690	11	537	fyn proto-oncogene (Fyn) alternative variant bSep08, mRNA.
<a href="#">Fyn</a>	<a href="#">Fyn.cSep08</a>	<a href="#">25150</a>	51171	2926	11	426	fyn proto-oncogene (48.7 kD) (Fyn) alternative variant cSep08, mRNA.
<a href="#">Fyn</a>	<a href="#">Fyn.dSep08</a>	<a href="#">25150</a>	19443	1028	4	160	fyn proto-oncogene (Fyn) alternative variant dSep08, mRNA.
<a href="#">fynoy</a>	<a href="#">fynoy.aSep08</a>		10687	767		84	v-crk sarcoma virus CT10 oncogene homolog like (9.6 kD) (fynoy) mRNA.
<a href="#">fypor</a>	<a href="#">fypor.aSep08</a>		2806	664		45	putative protein (5.3 kD) (fypor) mRNA.
<a href="#">FYRN.0</a>	<a href="#">FYRN.0.aSep08</a>		2975	1161		386	mixed-lineage leukemia 3 like (FYRN.0) mRNA.
<a href="#">FYRN.1</a>	<a href="#">FYRN.1.aSep08</a>		1934	467		155	myeloid lymphoid mixed-lineage leukemia like (FYRN.1) mRNA.

<a href="#">fysa</a>	<a href="#">fysa.aSep08</a>		7180	629		104	polyprotein (11.7 kD) (fysa) mRNA.
<a href="#">fyshee</a>	<a href="#">fyshee.aSep08</a>		633	453		27	putative protein (fyshee) mRNA.
<a href="#">fyto</a>	<a href="#">fyto.aSep08</a>		11127	300		100	inverted (fyto) mRNA.
<a href="#">Fytt1</a>	<a href="#">Fytt1.aSep08</a>	<a href="#">360726</a>	29402	3576	9	306	forty-two-three domain-containing protein 1 (Fytt1) alternative variant aSep08, mRNA.
<a href="#">Fytt1</a>	<a href="#">Fytt1.bSep08</a>	<a href="#">360726</a>	22126	682	5	207	putative protein of vertebrate origin (Fytt1) alternative variant bSep08, mRNA.
<a href="#">Fytt1</a>	<a href="#">Fytt1.dSep08</a>	<a href="#">360726</a>	6228	722	4	87	putative protein of vertebrate origin (Fytt1) alternative variant dSep08, mRNA.
<a href="#">Fytt1</a>	<a href="#">Fytt1.eSep08</a>	<a href="#">360726</a>	2225	614	2	31	putative protein (3.7 kD) (Fytt1) alternative variant eSep08, mRNA.
<a href="#">fyvar</a>	<a href="#">fyvar.aSep08</a>		1857	698		232	putative protein of metazoan origin (fyvar) mRNA.
<a href="#">fywey</a>	<a href="#">fywey.aSep08</a>		2180	629		58	putative protein (fywey) mRNA.
<a href="#">Fzr1</a>	<a href="#">Fzr1.aSep08</a>	<a href="#">314642</a>	12517	2386	14	493	fizzy/cell division cycle 20 related 1 (Drosophila) (54.6 kD) (Fzr1) alternative variant aSep08, mRNA.
<a href="#">Fzr1</a>	<a href="#">Fzr1.bSep08</a>	<a href="#">314642</a>	1266	757	3	240	fizzy/cell division cycle 20 related 1 (Drosophila) (Fzr1) alternative variant bSep08, mRNA.
<a href="#">Fzr1</a>	<a href="#">Fzr1.cSep08</a>	<a href="#">314642</a>	533	453	2	54	fizzy/cell division cycle 20 related 1 (Drosophila) (Fzr1) alternative variant cSep08, mRNA.
<a href="#">G-patch.0</a>	<a href="#">G-patch.0.aSep08</a>		122388	2943		182	d111/G-patch (G-patch.0) mRNA.
<a href="#">G2F.0</a>	<a href="#">G2F.0.aSep08</a>		9666	535	2	178	nidogen (G2F.0) alternative variant aSep08, mRNA.
<a href="#">G3bp1</a>	<a href="#">G3bp1.aSep08</a>	<a href="#">171092</a>	29685	2708	12	502	ras-GTPase-activating protein SH3-domain-binding like (G3bp1) alternative variant aSep08, mRNA.
<a href="#">G3bp1</a>	<a href="#">G3bp1.bSep08</a>	<a href="#">171092</a>	3732	982	5	209	ras-GTPase-activating protein SH3-domain-binding like (G3bp1) alternative variant bSep08, mRNA.
<a href="#">G3bp1</a>	<a href="#">G3bp1.cSep08</a>	<a href="#">171092</a>	5119	915	3	73	ras-GTPase-activating protein SH3-domain-binding like (G3bp1) alternative variant cSep08, mRNA.
<a href="#">G3bp2</a>	<a href="#">G3bp2.aSep08</a>	<a href="#">305240</a>	32703	3571	12	562	GTPase activating protein (SH3 domain) binding protein 2 (G3bp2) alternative variant aSep08, mRNA.
<a href="#">G3bp2</a>	<a href="#">G3bp2.cSep08</a>	<a href="#">305240</a>	98855	944	8	275	GTPase activating protein (SH3 domain) binding protein 2 (G3bp2) alternative variant cSep08, mRNA.
<a href="#">G6pc</a>	<a href="#">G6pc.bSep08</a>	<a href="#">25634</a>	7915	1641	4	286	glucose-6-phosphatase, catalytic (G6pc) alternative variant bSep08, mRNA.
<a href="#">G6pc</a>	<a href="#">G6pc.cSep08</a>	<a href="#">25634</a>	7636	1037	4	171	glucose-6-phosphatase, catalytic (19.1 kD) (G6pc) alternative variant cSep08, mRNA.
<a href="#">G6pc</a>	<a href="#">G6pc.dSep08</a>	<a href="#">25634</a>	2820	626	2	105	glucose-6-phosphatase, catalytic (G6pc) alternative variant dSep08, mRNA.
<a href="#">G6pc3</a>	<a href="#">G6pc3.bSep08</a>	<a href="#">303565</a>	5248	914	7	285	glucose 6 phosphatase, catalytic, 3 (G6pc3) alternative variant bSep08, mRNA.
<a href="#">G6pc3</a>	<a href="#">G6pc3.cSep08</a>	<a href="#">303565</a>	3451	721	5	209	glucose 6 phosphatase, catalytic, 3 (G6pc3) alternative variant cSep08, mRNA.
<a href="#">G6pc3</a>	<a href="#">G6pc3.dSep08</a>	<a href="#">303565</a>	2996	757	4	162	glucose 6 phosphatase, catalytic, 3 (G6pc3) alternative variant dSep08, mRNA.
<a href="#">G6pc3</a>	<a href="#">G6pc3.eSep08</a>	<a href="#">303565</a>	2896	1115	2	125	glucose 6 phosphatase, catalytic, 3 (14.5 kD) (G6pc3) alternative variant eSep08, mRNA.

<a href="#">G6pdx</a>	<a href="#">G6pdx.bSep08</a>	<a href="#">24377</a>	1559	789	4	193	glucose-6-phosphate dehydrogenase X-linked (G6pdx) alternative variant bSep08, mRNA.
<a href="#">G6pdx</a>	<a href="#">G6pdx.cSep08</a>	<a href="#">24377</a>	1797	1005	4	178	glucose-6-phosphate dehydrogenase X-linked (G6pdx) alternative variant cSep08, mRNA.
<a href="#">G6pdx</a>	<a href="#">G6pdx.dSep08</a>	<a href="#">24377</a>	832	729	1	43	glucose-6-phosphate dehydrogenase X-linked (G6pdx) alternative variant dSep08, mRNA.
<a href="#">G7c</a>	<a href="#">G7c.bSep08</a>	<a href="#">309611</a>	5106	1263	4	23	g7c protein (G7c) alternative variant bSep08, mRNA.
<a href="#">Gaa</a>	<a href="#">Gaa.bSep08</a>	<a href="#">367562</a>	6510	1054	4	312	glucosidase, alpha, acid (Gaa) alternative variant bSep08, mRNA.
<a href="#">Gaa</a>	<a href="#">Gaa.cSep08</a>	<a href="#">367562</a>	6080	1142	5	250	glucosidase, alpha, acid (Gaa) alternative variant cSep08, mRNA.
<a href="#">Gaa</a>	<a href="#">Gaa.dSep08</a>	<a href="#">367562</a>	1027	312	3	103	glucosidase, alpha, acid (Gaa) alternative variant dSep08, mRNA.
<a href="#">Gab1</a>	<a href="#">Gab1.bSep08</a>	<a href="#">361388</a>	67326	757	5	214	binding protein 1 like (Gab1) alternative variant bSep08, mRNA.
<a href="#">Gab1</a>	<a href="#">Gab1.cSep08</a>	<a href="#">361388</a>	10578	774	6	160	binding protein 1 like (Gab1) alternative variant cSep08, mRNA.
<a href="#">Gab1</a>	<a href="#">Gab1.dSep08</a>	<a href="#">361388</a>	10613	719	5	152	growth factor receptor bound protein 1 CRA b (Gab1) alternative variant dSep08, mRNA.
<a href="#">Gab1</a>	<a href="#">Gab1.eSep08</a>	<a href="#">361388</a>	9661	327	2	59	growth factor receptor bound protein 1 CRA a (Gab1) alternative variant eSep08, mRNA.
<a href="#">Gab1</a>	<a href="#">Gab1.gSep08</a>	<a href="#">361388</a>	836	592	2	27	putative protein (Gab1) alternative variant gSep08, mRNA.
<a href="#">Gabarapl2</a>	<a href="#">Gabarapl2.bSep08</a>	<a href="#">64670</a>	9658	618	2	40	GABA(A) receptor-associated protein like 2 (Gabarapl2) alternative variant bSep08, mRNA.
<a href="#">Gabbr1</a>	<a href="#">Gabbr1.bSep08</a>	<a href="#">81657</a>	3366	1552	4	298	gamma-aminobutyric acid (GABA) B receptor 1 (Gabbr1) alternative variant bSep08, mRNA.
<a href="#">Gabbr1</a>	<a href="#">Gabbr1.cSep08</a>	<a href="#">81657</a>	1148	603	2	121	gamma-aminobutyric acid (GABA) B receptor 1 (Gabbr1) alternative variant cSep08, mRNA.
<a href="#">Gabbr1</a>	<a href="#">Gabbr1.dSep08</a>	<a href="#">81657</a>	3237	303	2	100	gamma-aminobutyric acid (GABA) B receptor 1 (Gabbr1) alternative variant dSep08, mRNA.
<a href="#">Gabbr2</a>	<a href="#">Gabbr2.bSep08</a>	<a href="#">83633</a>	15596	786	5	226	gamma-aminobutyric acid (GABA) B receptor 2 (Gabbr2) alternative variant bSep08, mRNA.
<a href="#">Gabbr2</a>	<a href="#">Gabbr2.dSep08</a>	<a href="#">83633</a>	87119	754	5	95	gamma-aminobutyric acid (GABA) B receptor 2 (Gabbr2) alternative variant dSep08, mRNA.
<a href="#">Gabra1</a>	<a href="#">Gabra1.bSep08</a>	<a href="#">29705</a>	30036	861	1	150	gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 1 (Gabra1) alternative variant bSep08, mRNA.
<a href="#">Gabrd</a>	<a href="#">Gabrd.aSep08</a>	<a href="#">29689</a>	11893	1782		449	gamma-aminobutyric acid (GABA-A) receptor, subunit delta (50.6 kD) (Gabrd) mRNA.
<a href="#">Gabrg1</a>	<a href="#">Gabrg1.bSep08</a>	<a href="#">140674</a>	12523	391	3	95	gamma-aminobutyric acid (GABA) A receptor, gamma 1 (Gabrg1) alternative variant bSep08, mRNA.
<a href="#">Gabrg1</a>	<a href="#">Gabrg1.cSep08</a>	<a href="#">140674</a>	21631	499	3	37	gamma-aminobutyric acid (GABA) A receptor, gamma 1 (4.2 kD) (Gabrg1) alternative variant cSep08, mRNA.
<a href="#">Gabrg2</a>	<a href="#">Gabrg2.bSep08</a>	<a href="#">29709</a>	51675	2805	2	229	gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 2 (Gabrg2) alternative variant bSep08, mRNA.
<a href="#">Gabrp</a>	<a href="#">Gabrp.bSep08</a>	<a href="#">81658</a>	11628	537	5	135	gamma-aminobutyric acid (GABA-A) receptor, pi (Gabrp) alternative variant bSep08, mRNA.

<a href="#">gaby</a>	<a href="#">gaby.aSep08</a>		16853	311	2	22	putative protein (2.7 kD) (gaby) alternative variant aSep08, mRNA.
<a href="#">gaby</a>	<a href="#">gaby.bSep08</a>		1654	191	1	45	putative protein (gaby) alternative variant bSep08, mRNA.
<a href="#">gachy</a>	<a href="#">gachy.aSep08</a>		2111	392		21	putative protein (gachy) mRNA.
<a href="#">Gad1</a>	<a href="#">Gad1.bSep08</a>	<a href="#">24379</a>	33940	3105	1	444	glutamic acid decarboxylase 1 (Gad1) alternative variant bSep08, mRNA.
<a href="#">Gad2</a>	<a href="#">Gad2.bSep08</a>	<a href="#">24380</a>	4297	572	3	88	glutamic acid decarboxylase 2 (Gad2) alternative variant bSep08, mRNA.
<a href="#">Gadd45b</a>	<a href="#">Gadd45b.bSep08</a>	<a href="#">299626</a>	1409	743	2	128	growth arrest and DNA-damage-inducible 45 beta (Gadd45b) alternative variant bSep08, mRNA.
<a href="#">Gadd45gip1</a>	<a href="#">Gadd45gip1.aSep08</a>	<a href="#">288916</a>	2562	1224		228	growth arrest and DNA-damage-inducible, gamma interacting protein 1 (Gadd45gip1) complete mRNA.
<a href="#">gafer</a>	<a href="#">gafer.aSep08</a>		9772	486	2	64	putative protein (7.3 kD) (gafer) alternative variant aSep08, mRNA.
<a href="#">gafer</a>	<a href="#">gafer.bSep08</a>		9679	532	3	51	putative protein (5.8 kD) (gafer) alternative variant bSep08, mRNA.
<a href="#">gafer</a>	<a href="#">gafer.cSep08</a>		1311	405	1	28	putative protein (3.4 kD) (gafer) alternative variant cSep08, mRNA.
<a href="#">gaflo</a>	<a href="#">gaflo.aSep08</a>		2701	730	1	61	putative protein (gaflo) mRNA.
<a href="#">gaflu</a>	<a href="#">gaflu.aSep08</a>		2091	420		77	putative protein (gaflu) mRNA.
<a href="#">Gag_p30.0</a>	<a href="#">Gag_p30.0.bSep08</a>		4427	355	2	105	polyprotein (Gag_p30.0) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.1</a>	<a href="#">Gag_p30.1.aSep08</a>		4218	367		109	polyprotein (Gag_p30.1) mRNA.
<a href="#">Gag_p30.2</a>	<a href="#">Gag_p30.2.aSep08</a>		3355	358		106	polyprotein (Gag_p30.2) mRNA.
<a href="#">Gag_p30.6</a>	<a href="#">Gag_p30.6.aSep08</a>		3439	355		105	polyprotein (Gag_p30.6) mRNA.
<a href="#">Gag_p30.8</a>	<a href="#">Gag_p30.8.bSep08</a>		3272	367	3	109	polyprotein (Gag_p30.8) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.9</a>	<a href="#">Gag_p30.9.bSep08</a>		4641	1685	2	122	polyprotein (13.8 kD) (Gag_p30.9) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.9</a>	<a href="#">Gag_p30.9.cSep08</a>		7801	737	3	88	polyprotein (9.7 kD) (Gag_p30.9) alternative variant cSep08, mRNA.
<a href="#">Gag_p30.9</a>	<a href="#">Gag_p30.9.dSep08</a>		1865	759	2	59	polyprotein -pol (6.4 kD) (Gag_p30.9) alternative variant dSep08, mRNA.
<a href="#">Gag_p30.11</a>	<a href="#">Gag_p30.11.aSep08</a>		3333	356		85	polyprotein (Gag_p30.11) mRNA.
<a href="#">Gag_p30.12</a>	<a href="#">Gag_p30.12.aSep08</a>		3283	367	3	109	polyprotein (Gag_p30.12) alternative variant aSep08, mRNA.
<a href="#">Gag_p30.17</a>	<a href="#">Gag_p30.17.aSep08</a>		3395	355		105	polyprotein (Gag_p30.17) mRNA.
<a href="#">Gag_p30.18</a>	<a href="#">Gag_p30.18.bSep08</a>		2197	355	2	105	polyprotein (Gag_p30.18) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.19</a>	<a href="#">Gag_p30.19.bSep08</a>		3310	355	2	48	putative protein (Gag_p30.19) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.20</a>	<a href="#">Gag_p30.20.bSep08</a>		3397	355	2	105	polyprotein (Gag_p30.20) alternative variant bSep08, mRNA.
<a href="#">Gag_p30.23</a>	<a href="#">Gag_p30.23.aSep08</a>		3325	355		105	polyprotein (Gag_p30.23) mRNA.
<a href="#">Gak</a>	<a href="#">Gak.cSep08</a>	<a href="#">81659</a>	466	284	2	29	cyclin G associated kinase (3.2 kD) (Gak) alternative variant cSep08, mRNA.

<a href="#">gakee</a>	<a href="#">gakee.aSep08</a>		5308	712		237	lactase-phlorizin hydrolase (gakee) mRNA.
<a href="#">Gal3st4</a>	<a href="#">Gal3st4.bSep08</a>	<a href="#">498166</a>	1469	363	1	41	galactose-3-O-sulfotransferase 4 (Gal3st4) alternative variant bSep08, mRNA.
<a href="#">Galactosyl_T.0</a>	<a href="#">Galactosyl_T.0.aSep08</a>		16745	3017	8	315	2 beta 1 3 (Galactosyl_T.0) alternative variant aSep08, mRNA.
<a href="#">Galc</a>	<a href="#">Galc.bSep08</a>	<a href="#">314360</a>	25081	725	8	241	galactosylceramidase (Galc) alternative variant bSep08, mRNA.
<a href="#">Galc</a>	<a href="#">Galc.cSep08</a>	<a href="#">314360</a>	19555	808	8	216	galactosylceramidase (Galc) alternative variant cSep08, mRNA.
<a href="#">Gale</a>	<a href="#">Gale.bSep08</a>	<a href="#">114860</a>	1609	517	4	172	galactose-4-epimerase, UDP (Gale) alternative variant bSep08, mRNA.
<a href="#">Gale</a>	<a href="#">Gale.cSep08</a>	<a href="#">114860</a>	1908	774	7	171	galactose-4-epimerase, UDP (Gale) alternative variant cSep08, mRNA.
<a href="#">Gale</a>	<a href="#">Gale.dSep08</a>	<a href="#">114860</a>	961	719	3	131	galactose-4-epimerase, UDP (Gale) alternative variant dSep08, mRNA.
<a href="#">Gale</a>	<a href="#">Gale.eSep08</a>	<a href="#">114860</a>	1772	1349	5	69	galactose-4-epimerase, UDP (7.6 kD) (Gale) alternative variant eSep08, mRNA.
<a href="#">Galk1</a>	<a href="#">Galk1.bSep08</a>	<a href="#">287835</a>	9225	1501	4	217	galactokinase 1 (Galk1) alternative variant bSep08, mRNA.
<a href="#">Galk1</a>	<a href="#">Galk1.cSep08</a>	<a href="#">287835</a>	954	387	3	63	galactokinase 1 (Galk1) alternative variant cSep08, mRNA.
<a href="#">Galk2</a>	<a href="#">Galk2.bSep08</a>	<a href="#">296117</a>	251830	1784	10	402	galactokinase 2 (44.0 kD) (Galk2) alternative variant bSep08, mRNA.
<a href="#">Galk2</a>	<a href="#">Galk2.cSep08</a>	<a href="#">296117</a>	220468	934	7	256	galactokinase 2 (Galk2) alternative variant cSep08, mRNA.
<a href="#">Galk2</a>	<a href="#">Galk2.dSep08</a>	<a href="#">296117</a>	7716	726	2	95	galactokinase 2 (10.3 kD) (Galk2) alternative variant dSep08, mRNA.
<a href="#">Galk2</a>	<a href="#">Galk2.eSep08</a>	<a href="#">296117</a>	167385	367	3	74	galactokinase 2 (Galk2) alternative variant eSep08, mRNA.
<a href="#">Galnac4s-6st</a>	<a href="#">Galnac4s-6st.bSep08</a>	<a href="#">286974</a>	59906	2592	2	403	B cell RAG associated protein (46.6 kD) (Galnac4s-6st) alternative variant bSep08, mRNA.
<a href="#">Galns</a>	<a href="#">Galns.bSep08</a>	<a href="#">292073</a>	18523	1228	8	284	galactosamine (N-acetyl)-6-sulfate sulfatase (Galns) alternative variant bSep08, mRNA.
<a href="#">Galns</a>	<a href="#">Galns.cSep08</a>	<a href="#">292073</a>	2453	792	2	72	galactosamine (N-acetyl)-6-sulfate sulfatase (Galns) alternative variant cSep08, mRNA.
<a href="#">Galnt1</a>	<a href="#">Galnt1.bSep08</a>	<a href="#">79214</a>	18779	467	1	29	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (3.5 kD) (Galnt1) alternative variant bSep08, mRNA.
<a href="#">Galnt2</a>	<a href="#">Galnt2.aSep08</a>	<a href="#">292090</a>	48058	2116	2	529	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (Galnt2) alternative variant aSep08, mRNA.
<a href="#">Galnt3</a>	<a href="#">Galnt3.bSep08</a>	<a href="#">366061</a>	5930	345	2	77	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (Galnt3) alternative variant bSep08, mRNA.
<a href="#">Galnt7</a>	<a href="#">Galnt7.bSep08</a>	<a href="#">29750</a>	20641	3673	7	303	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (Galnt7) alternative variant bSep08, mRNA.
<a href="#">Galnt7</a>	<a href="#">Galnt7.cSep08</a>	<a href="#">29750</a>	27066	2345	6	300	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (Galnt7) alternative variant cSep08, mRNA.

<a href="#">Galnt7</a>	<a href="#">Galnt7.dSep08</a>	<a href="#">29750</a>	28984	582	3	99	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (Galnt7) alternative variant dSep08, mRNA.
<a href="#">Galnt10</a>	<a href="#">Galnt10.bSep08</a>	<a href="#">170501</a>	66617	440	2	134	putative protein (Galnt10) alternative variant bSep08, mRNA.
<a href="#">Galnt10</a>	<a href="#">Galnt10.cSep08</a>	<a href="#">170501</a>	77426	374	2	124	N-acetylgalactosaminyltransferase 10 (Galnt10) alternative variant cSep08, mRNA.
<a href="#">Galnt10</a>	<a href="#">Galnt10.dSep08</a>	<a href="#">170501</a>	2854	889	3	103	N-acetylgalactosaminyltransferase 10 (Galnt10) alternative variant dSep08, mRNA.
<a href="#">Galnt11</a>	<a href="#">Galnt11.bSep08</a>	<a href="#">311952</a>	16252	851	5	283	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (Galnt11) alternative variant bSep08, mRNA.
<a href="#">Galnt11</a>	<a href="#">Galnt11.cSep08</a>	<a href="#">311952</a>	6219	771	5	220	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (Galnt11) alternative variant cSep08, mRNA.
<a href="#">Galnt11</a>	<a href="#">Galnt11.dSep08</a>	<a href="#">311952</a>	6256	751	3	141	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (Galnt11) alternative variant dSep08, mRNA.
<a href="#">Galnt11</a>	<a href="#">Galnt11.eSep08</a>	<a href="#">311952</a>	3689	971	2	82	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (Galnt11) alternative variant eSep08, mRNA.
<a href="#">Galnt12</a>	<a href="#">Galnt12.aSep08</a>	<a href="#">313233</a>	2825	755		88	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 12 (Galnt12) mRNA.
<a href="#">Galnt13</a>	<a href="#">Galnt13.bSep08</a>	<a href="#">311039</a>	137433	876	3	100	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (Galnt13) alternative variant bSep08, mRNA.
<a href="#">Galnt13</a>	<a href="#">Galnt13.cSep08</a>	<a href="#">311039</a>	72434	1210	4	41	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (Galnt13) alternative variant cSep08, mRNA.
<a href="#">Galnt13</a>	<a href="#">Galnt13.dSep08</a>	<a href="#">311039</a>	8915	675	2	33	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (3.6 kD) (Galnt13) alternative variant dSep08, mRNA.
<a href="#">Galnt13</a>	<a href="#">Galnt13.eSep08</a>	<a href="#">311039</a>	71839	614	3	56	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (6.6 kD) (Galnt13) alternative variant eSep08, mRNA.
<a href="#">Galnt13</a>	<a href="#">Galnt13.fSep08</a>	<a href="#">311039</a>	70802	427	3	41	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (Galnt13) alternative variant fSep08, mRNA.
<a href="#">Galnt11</a>	<a href="#">Galnt11.aSep08</a>	<a href="#">362760</a>	26041	1269	13	220	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 1 (Galnt11) mRNA.
<a href="#">galoy</a>	<a href="#">galoy.aSep08</a>		5789	645		116	putative protein of vertebrate origin (galoy) mRNA.
<a href="#">Galt</a>	<a href="#">Galt.bSep08</a>	<a href="#">298003</a>	1453	709	5	175	uridyl transferase CRA b (19.5 kD) (Galt) alternative variant bSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.cSep08</a>	<a href="#">298003</a>	1291	612	4	170	uridyl transferase (Galt) alternative variant cSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.dSep08</a>	<a href="#">298003</a>	2002	766	8	129	uridyl transferase CRA b (Galt) alternative variant dSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.eSep08</a>	<a href="#">298003</a>	1205	893	2	111	uridyl transferase (Galt) alternative variant eSep08, mRNA.

<a href="#">Galt</a>	<a href="#">Galt.fSep08</a>	<a href="#">298003</a>	1345	723	4	88	uridyl transferase CRA b (Galt) alternative variant fSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.gSep08</a>	<a href="#">298003</a>	796	525	2	75	uridyl transferase (8.6 kD) (Galt) alternative variant gSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.hSep08</a>	<a href="#">298003</a>	1730	1629	2	97	uridyl transferase CRA b (11.0 kD) (Galt) alternative variant hSep08, mRNA.
<a href="#">Galt</a>	<a href="#">Galt.iSep08</a>	<a href="#">298003</a>	865	775	2	87	uridyl transferase CRA b (9.7 kD) (Galt) alternative variant iSep08, mRNA.
<a href="#">gamer</a>	<a href="#">gamer.aSep08</a>		3226	258		35	putative protein (gamer) mRNA.
<a href="#">Gamt</a>	<a href="#">Gamt.bSep08</a>	<a href="#">25257</a>	2006	507	4	169	guanidinoacetate methyltransferase (Gamt) alternative variant bSep08, mRNA.
<a href="#">Gan</a>	<a href="#">Gan.bSep08</a>	<a href="#">307893</a>	29112	399	2	132	giant axonal neuropathy (Gan) alternative variant bSep08, mRNA.
<a href="#">Ganab</a>	<a href="#">Ganab.bSep08</a>	<a href="#">293721</a>	15185	1372	11	457	alpha glucosidase 2 alpha neutral subunit (Ganab) alternative variant bSep08, mRNA.
<a href="#">Ganab</a>	<a href="#">Ganab.cSep08</a>	<a href="#">293721</a>	1040	376	3	67	alpha glucosidase 2 alpha neutral subunit (Ganab) alternative variant cSep08, mRNA.
<a href="#">ganoy</a>	<a href="#">ganoy.aSep08</a>		2019	779	1	115	putative protein (ganoy) alternative variant aSep08, mRNA.
<a href="#">ganoy</a>	<a href="#">ganoy.bSep08</a>		2061	612		83	putative mitochondrial protein (9.4 kD) (ganoy) alternative variant bSep08, complete mRNA.
<a href="#">Gap43</a>	<a href="#">Gap43.bSep08</a>	<a href="#">29423</a>	7209	554	1	54	growth associated protein 43 (Gap43) alternative variant bSep08, mRNA.
<a href="#">Gapdh</a>	<a href="#">Gapdh.bSep08</a>	<a href="#">24383</a>	4143	976	7	325	glyceraldehyde-3-phosphate dehydrogenase (Gapdh) alternative variant bSep08, mRNA.
<a href="#">Gapdh</a>	<a href="#">Gapdh.cSep08</a>	<a href="#">24383</a>	3150	736	6	229	glyceraldehyde-3-phosphate dehydrogenase (Gapdh) alternative variant cSep08, mRNA.
<a href="#">Gapdhs</a>	<a href="#">Gapdhs.bSep08</a>	<a href="#">66020</a>	10120	1496	9	432	glyceraldehyde-3-phosphate dehydrogenase, spermatogenic (46.7 kD) (Gapdhs) alternative variant bSep08, mRNA.
<a href="#">Gapdhs</a>	<a href="#">Gapdhs.cSep08</a>	<a href="#">66020</a>	8152	562	2	174	glyceraldehyde-3-phosphate dehydrogenase, spermatogenic (19.2 kD) (Gapdhs) alternative variant cSep08, mRNA.
<a href="#">Gapdhs</a>	<a href="#">Gapdhs.dSep08</a>	<a href="#">66020</a>	8146	621	2	142	glyceraldehyde-3-phosphate dehydrogenase, spermatogenic (15.4 kD) (Gapdhs) alternative variant dSep08, mRNA.
<a href="#">gapor</a>	<a href="#">gapor.aSep08</a>		2593	521		173	kinase 2 CRA b (gapor) mRNA.
<a href="#">Gapvd1</a>	<a href="#">Gapvd1.aSep08</a>	<a href="#">311880</a>	29000	4042	16	591	GTPase activating protein and VPS9 domains 1 (Gapvd1) alternative variant aSep08, mRNA.
<a href="#">Gapvd1</a>	<a href="#">Gapvd1.bSep08</a>	<a href="#">311880</a>	13106	719	6	239	GTPase activating protein and VPS9 domains 1 (Gapvd1) alternative variant bSep08, mRNA.
<a href="#">Gapvd1</a>	<a href="#">Gapvd1.dSep08</a>	<a href="#">311880</a>	7384	772	4	111	GTPase activating protein and VPS9 domains 1 (Gapvd1) alternative variant dSep08, mRNA.
<a href="#">garby</a>	<a href="#">garby.aSep08</a>		451	301		63	putative protein (7.3 kD) (garby) mRNA.
<a href="#">garchy</a>	<a href="#">garchy.aSep08</a>		4542	275		87	putative protein (garchy) mRNA.
<a href="#">garfer</a>	<a href="#">garfer.aSep08</a>		3793	1062	2	108	ab2-143 like (11.6 kD) (garfer) alternative variant aSep08, mRNA.



<a href="#">garflo</a>	<a href="#">garflo.aSep08</a>		18413	802	2	88	putative protein (garflo) alternative variant aSep08, mRNA.
<a href="#">garflo</a>	<a href="#">garflo.bSep08</a>		2737	642	3	59	putative protein (garflo) alternative variant bSep08, mRNA.
<a href="#">garflu</a>	<a href="#">garflu.aSep08</a>		14803	726		65	putative protein (7.1 kD) (garflu) mRNA.
<a href="#">garkee</a>	<a href="#">garkee.aSep08</a>		7204	735		54	putative protein (5.8 kD) (garkee) mRNA.
<a href="#">garloy</a>	<a href="#">garloy.aSep08</a>		1134	571		59	putative protein (6.1 kD) (garloy) mRNA.
<a href="#">garmee</a>	<a href="#">garmee.aSep08</a>		107995	739		50	putative protein (5.5 kD) (garmee) mRNA.
<a href="#">garmer</a>	<a href="#">garmer.aSep08</a>		12755	748		178	putative cytoplasmic protein of eukaryotic origin (20.8 kD) (garmer) mRNA.
<a href="#">Garnl1</a>	<a href="#">Garnl1.aSep08</a>	<a href="#">56785</a>	137638	3741	15	802	GTPase activating RANGAP domain-like 1 (Garnl1) alternative variant aSep08, mRNA.
<a href="#">Garnl1</a>	<a href="#">Garnl1.bSep08</a>	<a href="#">56785</a>	38418	370	2	105	GTPase activating RANGAP domain-like 1 (Garnl1) alternative variant bSep08, mRNA.
<a href="#">Garnl4</a>	<a href="#">Garnl4.bSep08</a>	<a href="#">303298</a>	8524	432	4	144	GTPase activating RANGAP domain-like 4 (Garnl4) alternative variant bSep08, mRNA.
<a href="#">Garnl4</a>	<a href="#">Garnl4.cSep08</a>	<a href="#">303298</a>	5780	395	5	93	GTPase activating RANGAP domain-like 4 (Garnl4) alternative variant cSep08, mRNA.
<a href="#">garnoy</a>	<a href="#">garnoy.aSep08</a>		517	400		133	family member 3 (garnoy) mRNA.
<a href="#">Garp</a>	<a href="#">Garp.aSep08</a>	<a href="#">293135</a>	9016	370		80	glycoprotein A repetitions predominant (Garp) mRNA.
<a href="#">garpor</a>	<a href="#">garpor.aSep08</a>		7592	684		100	putative secreted or extracellular protein precursor (11.0 kD) (garpor) mRNA.
<a href="#">Gars</a>	<a href="#">Gars.aSep08</a>	<a href="#">297113</a>	41014	2459		751	glycyl-tRNA synthetase (Gars) mRNA.
<a href="#">garsa</a>	<a href="#">garsa.aSep08</a>		9502	601		100	putative protein (garsa) mRNA.
<a href="#">garshee</a>	<a href="#">garshee.aSep08</a>		7690	631		90	putative protein (9.6 kD) (garshee) mRNA.
<a href="#">Gart</a>	<a href="#">Gart.aSep08</a>	<a href="#">288259</a>	15090	2463	14	732	phosphoribosylglycinamide formyltransferase (Gart) alternative variant aSep08, mRNA.
<a href="#">Gart</a>	<a href="#">Gart.bSep08</a>	<a href="#">288259</a>	12512	1850	11	433	phosphoribosylglycinamide formyltransferase (45.7 kD) (Gart) alternative variant bSep08, mRNA.
<a href="#">Gart</a>	<a href="#">Gart.cSep08</a>	<a href="#">288259</a>	7487	481	5	155	phosphoribosylglycinamide formyltransferase (Gart) alternative variant cSep08, mRNA.
<a href="#">Gart</a>	<a href="#">Gart.dSep08</a>	<a href="#">288259</a>	1782	278	2	92	phosphoribosylglycinamide formyltransferase (Gart) alternative variant dSep08, mRNA.
<a href="#">garto</a>	<a href="#">garto.aSep08</a>		36244	476		103	guanine nucleotide exchange factor (garto) mRNA.
<a href="#">garvar</a>	<a href="#">garvar.aSep08</a>		945	835		74	putative protein of mammalian origin (garvar) mRNA.
<a href="#">garwey</a>	<a href="#">garwey.aSep08</a>		3000	506		27	putative protein (garwey) mRNA.
<a href="#">Gas2</a>	<a href="#">Gas2.bSep08</a>	<a href="#">499156</a>	18709	1562	2	85	growth arrest-specific 2 (9.6 kD) (Gas2) alternative variant bSep08, mRNA.
<a href="#">Gas2</a>	<a href="#">Gas2.cSep08</a>	<a href="#">499156</a>	7806	300	3	42	growth arrest-specific 2 (Gas2) alternative variant cSep08, mRNA.
<a href="#">Gas2l1</a>	<a href="#">Gas2l1.aSep08</a>	<a href="#">360973</a>	4977	2555	6	712	growth arrest-specific 2 like 1 (Gas2l1) alternative variant aSep08, mRNA.
<a href="#">Gas2l1</a>	<a href="#">Gas2l1.dSep08</a>	<a href="#">360973</a>	3670	647	3	134	growth arrest-specific 2 like 1 (Gas2l1) alternative variant dSep08, mRNA.
<a href="#">Gas2l1</a>	<a href="#">Gas2l1.eSep08</a>	<a href="#">360973</a>	1147	601	4	56	growth arrest-specific 2 like 1 (Gas2l1) alternative variant eSep08, mRNA.
<a href="#">Gas2l3</a>	<a href="#">Gas2l3.aSep08</a>	<a href="#">680280</a>	10138	654		195	growth arrest-specific 2 like 3 (Gas2l3) mRNA.

<a href="#">Gas5</a>	<a href="#">Gas5.aSep08</a>	<a href="#">81714</a>	3326	2549	2	37	growth arrest specific 5 (4.1 kD) (Gas5) alternative variant aSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.bSep08</a>	<a href="#">81714</a>	3314	1144	6	38	growth arrest specific 5 (4.4 kD) (Gas5) alternative variant bSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.dSep08</a>	<a href="#">81714</a>	3319	974	6	37	growth arrest specific 5 (4.1 kD) (Gas5) alternative variant dSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.eSep08</a>	<a href="#">81714</a>	3315	871	8	38	growth arrest specific 5 (4.4 kD) (Gas5) alternative variant eSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.fSep08</a>	<a href="#">81714</a>	1675	714	4	23	growth arrest specific 5 (Gas5) alternative variant fSep08, mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.gSep08</a>	<a href="#">81714</a>	3303	694	8	38	growth arrest specific 5 (4.4 kD) (Gas5) alternative variant gSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.hSep08</a>	<a href="#">81714</a>	3327	613	8	37	growth arrest specific 5 (4.1 kD) (Gas5) alternative variant hSep08, complete mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.iSep08</a>	<a href="#">81714</a>	1984	515	4	41	growth arrest specific 5 (Gas5) alternative variant iSep08, mRNA.
<a href="#">Gas5</a>	<a href="#">Gas5.jSep08</a>	<a href="#">81714</a>	3322	429	9	38	growth arrest specific 5 (4.4 kD) (Gas5) alternative variant jSep08, complete mRNA.
<a href="#">Gas6</a>	<a href="#">Gas6.bSep08</a>	<a href="#">58935</a>	13506	405	4	82	growth arrest specific 6 (Gas6) alternative variant bSep08, mRNA.
<a href="#">Gas8</a>	<a href="#">Gas8.bSep08</a>	<a href="#">361438</a>	5598	341	1	113	growth arrest specific 8 (Gas8) alternative variant bSep08, mRNA.
<a href="#">gasa</a>	<a href="#">gasa.aSep08</a>		12742	686		71	putative protein (8.5 kD) (gasa) mRNA.
<a href="#">gashee</a>	<a href="#">gashee.aSep08</a>		994	510		87	putative protein (gashee) mRNA.
<a href="#">Gata2</a>	<a href="#">Gata2.bSep08</a>	<a href="#">25159</a>	7582	2168	5	362	GATA binding protein 2 (37.6 kD) (Gata2) alternative variant bSep08, mRNA.
<a href="#">Gata3</a>	<a href="#">Gata3.bSep08</a>	<a href="#">85471</a>	9210	710	1	236	GATA binding protein 3 (Gata3) alternative variant bSep08, mRNA.
<a href="#">Gata4</a>	<a href="#">Gata4.bSep08</a>	<a href="#">54254</a>	67555	1409	2	268	GATA binding protein 4 (Gata4) alternative variant bSep08, mRNA.
<a href="#">Gatad1</a>	<a href="#">Gatad1.aSep08</a>	<a href="#">500005</a>	11521	2483	4	370	putative protein of metazoan origin (Gatad1) alternative variant aSep08, mRNA.
<a href="#">Gatad1</a>	<a href="#">Gatad1.bSep08</a>	<a href="#">500005</a>	7771	811	4	197	putative protein of metazoan origin (Gatad1) alternative variant bSep08, mRNA.
<a href="#">Gatad1</a>	<a href="#">Gatad1.cSep08</a>	<a href="#">500005</a>	9283	738	3	185	putative protein of metazoan origin (21.1 kD) (Gatad1) alternative variant cSep08, mRNA.
<a href="#">Gatad1</a>	<a href="#">Gatad1.dSep08</a>	<a href="#">500005</a>	1119	626	2	171	putative protein of vertebrate origin (Gatad1) alternative variant dSep08, mRNA.
<a href="#">Gatad2a</a>	<a href="#">Gatad2a.aSep08</a>	<a href="#">290669</a>	30406	4551	9	580	putative protein, with a coiled coil domain, of metazoan origin (Gatad2a) alternative variant aSep08, mRNA.
<a href="#">Gatad2a</a>	<a href="#">Gatad2a.bSep08</a>	<a href="#">290669</a>	6049	2316	4	262	putative nuclear protein of metazoan origin (28.6 kD) (Gatad2a) alternative variant bSep08, mRNA.
<a href="#">Gatad2a</a>	<a href="#">Gatad2a.cSep08</a>	<a href="#">290669</a>	3686	757	6	252	putative protein, with a coiled coil domain, of vertebrate origin (Gatad2a) alternative variant cSep08, mRNA.
<a href="#">Gatad2a</a>	<a href="#">Gatad2a.dSep08</a>	<a href="#">290669</a>	21397	755	6	251	putative protein, with a coiled coil domain, of vertebrate origin (Gatad2a) alternative variant dSep08, mRNA.

<a href="#">Gatad2a</a>	<a href="#">Gatad2a.eSep08</a>	<a href="#">290669</a>	2057	844	1	82	putative protein of vertebrate origin (8.8 kD) (Gatad2a) alternative variant eSep08, mRNA.
<a href="#">Gatc</a>	<a href="#">Gatc.aSep08</a>	<a href="#">360821</a>	5464	561	1	164	glutamyl-tRNA(Gln) amidotransferase, subunit C homolog (bacterial) (Gatc) alternative variant aSep08, mRNA.
<a href="#">Gatc</a>	<a href="#">Gatc.cSep08</a>	<a href="#">360821</a>	1897	497	1	99	glutamyl-tRNA(Gln) amidotransferase, subunit C homolog (bacterial) (Gatc) alternative variant cSep08, mRNA.
<a href="#">gato</a>	<a href="#">gato.aSep08</a>		12338	346		114	WD repeat domain 60 like (gato) mRNA.
<a href="#">Gats</a>	<a href="#">Gats.aSep08</a>	<a href="#">304410</a>	3302	807		268	opposite strand transcription unit to Stag3 (Gats) mRNA.
<a href="#">gavar</a>	<a href="#">gavar.aSep08</a>		11504	587		36	putative protein (4.0 kD) (gavar) mRNA.
<a href="#">gawby</a>	<a href="#">gawby.aSep08</a>		1095	295		67	putative protein (gawby) mRNA.
<a href="#">gawchy</a>	<a href="#">gawchy.aSep08</a>		5621	448		149	cobl-like 1 (gawchy) mRNA.
<a href="#">gawdar</a>	<a href="#">gawdar.aSep08</a>		1625	509		169	carboxylesterase (gawdar) mRNA.
<a href="#">gawey</a>	<a href="#">gawey.bSep08</a>		768	653	2	75	putative protein (8.7 kD) (gawey) alternative variant bSep08, mRNA.
<a href="#">gawfer</a>	<a href="#">gawfer.aSep08</a>		1620	454		54	putative protein (gawfer) mRNA.
<a href="#">gawflo</a>	<a href="#">gawflo.aSep08</a>		441	299		48	putative protein (gawflo) mRNA.
<a href="#">gawflu</a>	<a href="#">gawflu.aSep08</a>		2547	526		92	putative protein (gawflu) mRNA.
<a href="#">gawkee</a>	<a href="#">gawkee.aSep08</a>		4576	1815	2	191	slit-robo Rho GTPase activating protein 2 (20.4 kD) (gawkee) alternative variant aSep08, mRNA.
<a href="#">gawloy</a>	<a href="#">gawloy.aSep08</a>		27114	251		29	putative protein (gawloy) alternative variant aSep08, mRNA.
<a href="#">gawloy</a>	<a href="#">gawloy.bSep08</a>		26965	186	1	18	putative protein (2.0 kD) (gawloy) alternative variant bSep08, mRNA.
<a href="#">gawmee</a>	<a href="#">gawmee.aSep08</a>		8881	259		41	putative protein (gawmee) mRNA.
<a href="#">gawmer</a>	<a href="#">gawmer.aSep08</a>		472	415		52	putative protein (gawmer) mRNA.
<a href="#">gawnoy</a>	<a href="#">gawnoy.aSep08</a>		4739	1446		312	tbc1 domain family member 8 (gawnoy) mRNA.
<a href="#">gawpor</a>	<a href="#">gawpor.aSep08</a>		48187	244		26	putative protein (2.9 kD) (gawpor) mRNA.
<a href="#">gawsa</a>	<a href="#">gawsa.aSep08</a>		8833	539	3	179	transcription initiation factor tfiid (gawsa) alternative variant aSep08, mRNA.
<a href="#">gawsa</a>	<a href="#">gawsa.bSep08</a>		5046	597	2	114	transcription initiation factor tfiid (gawsa) alternative variant bSep08, mRNA.
<a href="#">gawshee</a>	<a href="#">gawshee.aSep08</a>		8426	835		94	putative protein (gawshee) mRNA.
<a href="#">gawto</a>	<a href="#">gawto.aSep08</a>		1669	635	2	95	putative protein (gawto) alternative variant aSep08, mRNA.
<a href="#">gawto</a>	<a href="#">gawto.bSep08</a>		4989	489	4	97	putative protein (gawto) alternative variant bSep08, mRNA.
<a href="#">gawvar</a>	<a href="#">gawvar.aSep08</a>		3617	563		113	putative protein (gawvar) mRNA.
<a href="#">gawwey</a>	<a href="#">gawwey.aSep08</a>		1323	558		42	putative protein (gawwey) mRNA.
<a href="#">Gba2</a>	<a href="#">Gba2.aSep08</a>	<a href="#">298399</a>	12554	4096	16	500	glucosidase beta 2 CRA a (56.8 kD) (Gba2) alternative variant aSep08, mRNA.
<a href="#">Gba2</a>	<a href="#">Gba2.bSep08</a>	<a href="#">298399</a>	2695	1332	9	443	glucosidase beta 2 CRA a (Gba2) alternative variant bSep08, mRNA.
<a href="#">Gba2</a>	<a href="#">Gba2.cSep08</a>	<a href="#">298399</a>	5465	1619	3	195	glucosidase beta 2 CRA a (22.0 kD) (Gba2) alternative variant cSep08, mRNA.
<a href="#">Gba2</a>	<a href="#">Gba2.eSep08</a>	<a href="#">298399</a>	784	705	2	76	glucosidase beta 2 CRA a (Gba2) alternative variant eSep08, mRNA.

<a href="#">Gba3</a>	<a href="#">Gba3.aSep08</a>	<a href="#">289687</a>	141913	2022		459	glucosidase, beta, acid 3 (cytosolic) (Gba3) mRNA.
<a href="#">Gbas</a>	<a href="#">Gbas.bSep08</a>	<a href="#">498174</a>	13694	1738	1	185	glioblastoma amplified sequence (Gbas) alternative variant bSep08, mRNA.
<a href="#">Gbe1</a>	<a href="#">Gbe1.aSep08</a>	<a href="#">288333</a>	156881	1119		330	glucan (1,4-alpha-), branching enzyme 1 (Gbe1) mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.aSep08</a>	<a href="#">309451</a>	32902	5744	35	1722	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant aSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.bSep08</a>	<a href="#">309451</a>	3798	892	7	297	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant bSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.cSep08</a>	<a href="#">309451</a>	2080	719	5	185	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant cSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.dSep08</a>	<a href="#">309451</a>	1108	709	4	138	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant dSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.eSep08</a>	<a href="#">309451</a>	13615	368	3	92	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant eSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.gSep08</a>	<a href="#">309451</a>	5831	519	2	73	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant gSep08, mRNA.
<a href="#">Gbf1</a>	<a href="#">Gbf1.iSep08</a>	<a href="#">309451</a>	979	316	2	12	golgi-specific brefeldin A-resistance factor 1 (Gbf1) alternative variant iSep08, mRNA.
<a href="#">Gbp2</a>	<a href="#">Gbp2.bSep08</a>	<a href="#">171164</a>	6029	1282		277	guanylate nucleotide binding protein 2 (Gbp2) alternative variant bSep08, mRNA.
<a href="#">Gbp4</a>	<a href="#">Gbp4.aSep08</a>	<a href="#">310917</a>	1996	360		119	guanylate binding protein 4 (Gbp4) mRNA.
<a href="#">Gbp5</a>	<a href="#">Gbp5.bSep08</a>	<a href="#">362050</a>	4032	1442	4	209	guanylate nucleotide binding protein 5 (Gbp5) alternative variant bSep08, mRNA.
<a href="#">Gc</a>	<a href="#">Gc.bSep08</a>	<a href="#">24384</a>	9876	217	2	72	group specific component (Gc) alternative variant bSep08, mRNA.
<a href="#">Gc</a>	<a href="#">Gc.cSep08</a>	<a href="#">24384</a>	6528	207	3	59	group specific component (Gc) alternative variant cSep08, mRNA.
<a href="#">Gca</a>	<a href="#">Gca.aSep08</a>	<a href="#">295647</a>	24956	458	5	122	grancalcin (Gca) alternative variant aSep08, mRNA.
<a href="#">Gcap14</a>	<a href="#">Gcap14.aSep08</a>	<a href="#">306306</a>	67063	938	6	312	granule cell antiserum positive 14 (Gcap14) alternative variant aSep08, mRNA.
<a href="#">Gcap14</a>	<a href="#">Gcap14.bSep08</a>	<a href="#">306306</a>	55741	667	5	191	granule cell antiserum positive 14 (Gcap14) alternative variant bSep08, mRNA.
<a href="#">Gcap14</a>	<a href="#">Gcap14.cSep08</a>	<a href="#">306306</a>	26660	675	4	159	granule cell antiserum positive 14 (Gcap14) alternative variant cSep08, mRNA.
<a href="#">Gcap14</a>	<a href="#">Gcap14.dSep08</a>	<a href="#">306306</a>	2760	284	1	94	granule cell antiserum positive 14 (Gcap14) alternative variant dSep08, mRNA.
<a href="#">Gcdh</a>	<a href="#">Gcdh.bSep08</a>	<a href="#">364975</a>	3138	1385	5	213	glutaryl-Coenzyme A dehydrogenase (23.4 kD) (Gcdh) alternative variant bSep08, mRNA.
<a href="#">Gcdh</a>	<a href="#">Gcdh.dSep08</a>	<a href="#">364975</a>	1216	978	3	67	glutaryl-Coenzyme A dehydrogenase (Gcdh) alternative variant dSep08, mRNA.
<a href="#">Gcgr</a>	<a href="#">Gcgr.bSep08</a>	<a href="#">24953</a>	7986	1885	10	287	glucagon receptor (32.4 kD) (Gcgr) alternative variant bSep08, mRNA.
<a href="#">Gcgr</a>	<a href="#">Gcgr.cSep08</a>	<a href="#">24953</a>	619	349	2	115	glucagon receptor (Gcgr) alternative variant cSep08, mRNA.
<a href="#">Gch1</a>	<a href="#">Gch1.bSep08</a>	<a href="#">29244</a>	115314	531	1	147	GTP cyclohydrolase 1 (Gch1) alternative variant bSep08, mRNA.

<a href="#">Gch1</a>	<a href="#">Gch1.cSep08</a>	<a href="#">29244</a>	4579	2157	1	83	GTP cyclohydrolase 1 (Gch1) alternative variant cSep08, mRNA.
<a href="#">Gck</a>	<a href="#">Gck.bSep08</a>	<a href="#">24385</a>	1222	755	2	85	glucokinase (9.2 kD) (Gck) alternative variant bSep08, mRNA.
<a href="#">Gclc</a>	<a href="#">Gclc.bSep08</a>	<a href="#">25283</a>	6974	1814	5	198	glutamate-cysteine ligase, catalytic subunit (Gclc) alternative variant bSep08, mRNA.
<a href="#">Gclc</a>	<a href="#">Gclc.cSep08</a>	<a href="#">25283</a>	4640	727	3	126	glutamate-cysteine ligase, catalytic subunit (Gclc) alternative variant cSep08, mRNA.
<a href="#">Gclm</a>	<a href="#">Gclm.bSep08</a>	<a href="#">29739</a>	9833	907	6	177	glutamate cysteine ligase, modifier subunit (Gclm) alternative variant bSep08, mRNA.
<a href="#">Gcn111</a>	<a href="#">Gcn111.aSep08</a>	<a href="#">690632</a>	25016	4471	25	1315	gcn1 general control of amino-acid synthesis 1-like 1 (Gcn111) alternative variant aSep08, mRNA.
<a href="#">Gcn111</a>	<a href="#">Gcn111.bSep08</a>	<a href="#">690632</a>	3202	524	3	167	gcn1 general control of amino-acid synthesis 1-like 1 (Gcn111) alternative variant bSep08, mRNA.
<a href="#">Gcn111</a>	<a href="#">Gcn111.dSep08</a>	<a href="#">690632</a>	496	410	2	91	gcn1 general control of amino-acid synthesis 1-like 1 CRA b (Gcn111) alternative variant dSep08, mRNA.
<a href="#">Gcn111</a>	<a href="#">Gcn111.eSep08</a>	<a href="#">690632</a>	970	479	2	74	gcn1 1 (Gcn111) alternative variant eSep08, mRNA.
<a href="#">Gcn5l2</a>	<a href="#">Gcn5l2.bSep08</a>	<a href="#">303539</a>	2003	352	1	117	GCN5 general control of amino acid synthesis-like 2 (yeast) (Gcn5l2) alternative variant bSep08, mRNA.
<a href="#">Gcnt2</a>	<a href="#">Gcnt2.bSep08</a>	<a href="#">306860</a>	31368	424	1	89	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (10.3 kD) (Gcnt2) alternative variant bSep08, mRNA.
<a href="#">Gcom1</a>	<a href="#">Gcom1.bSep08</a>	<a href="#">363091</a>	35395	777	6	233	GRINL1A combined protein (Gcom1) alternative variant bSep08, mRNA.
<a href="#">Gcsh</a>	<a href="#">Gcsh.bSep08</a>	<a href="#">171133</a>	5933	891	1	80	glycine cleavage system protein H (aminomethyl carrier) (Gcsh) alternative variant bSep08, mRNA.
<a href="#">Gda</a>	<a href="#">Gda.bSep08</a>	<a href="#">83585</a>	79286	5152	13	380	guanine deaminase (42.3 kD) (Gda) alternative variant bSep08, mRNA.
<a href="#">Gda</a>	<a href="#">Gda.cSep08</a>	<a href="#">83585</a>	20974	951	8	236	guanine deaminase (26.1 kD) (Gda) alternative variant cSep08, mRNA.
<a href="#">Gda</a>	<a href="#">Gda.fSep08</a>	<a href="#">83585</a>	1306	568	2	33	guanine deaminase (3.6 kD) (Gda) alternative variant fSep08, mRNA.
<a href="#">Gdap111</a>	<a href="#">Gdap111.dSep08</a>	<a href="#">311616</a>	14748	283	2	78	ganglioside-induced differentiation-associated protein 1-like 1 (Gdap111) alternative variant dSep08, mRNA.
<a href="#">Gdap111</a>	<a href="#">Gdap111.eSep08</a>	<a href="#">311616</a>	1101	1004	3	67	ganglioside-induced differentiation-associated protein 1-like 1 (Gdap111) alternative variant eSep08, mRNA.
<a href="#">Gdap2</a>	<a href="#">Gdap2.bSep08</a>	<a href="#">362004</a>	13441	1823	2	120	ganglioside-induced differentiation-associated-protein 2 (14.1 kD) (Gdap2) alternative variant bSep08, mRNA.
<a href="#">Gdf10</a>	<a href="#">Gdf10.aSep08</a>	<a href="#">79216</a>	8861	780	2	260	growth differentiation factor 10 (Gdf10) alternative variant aSep08, mRNA.
<a href="#">Gdi1</a>	<a href="#">Gdi1.bSep08</a>	<a href="#">25183</a>	6665	2554	8	266	dissociation inhibitor (30.0 kD) (Gdi1) alternative variant bSep08, complete mRNA.
<a href="#">Gdi1</a>	<a href="#">Gdi1.cSep08</a>	<a href="#">25183</a>	6673	3369	7	241	dissociation inhibitor (Gdi1) alternative variant cSep08, mRNA.
<a href="#">Gdi1</a>	<a href="#">Gdi1.dSep08</a>	<a href="#">25183</a>	1744	1548	1	112	dissociation inhibitor (12.5 kD) (Gdi1) alternative variant dSep08, mRNA.
<a href="#">Gdi2</a>	<a href="#">Gdi2.bSep08</a>	<a href="#">29662</a>	979	750	3	137	GDP dissociation inhibitor 2 (15.9 kD) (Gdi2) alternative variant bSep08, mRNA.

<a href="#">Gdnf</a>	<a href="#">Gdnf.aSep08</a>	<a href="#">25453</a>	20206	573		185	glial cell line derived neurotrophic factor (20.7 kD) (Gdnf) complete mRNA.
<a href="#">Gdpd1</a>	<a href="#">Gdpd1.aSep08</a>	<a href="#">303407</a>	43457	1776	1	533	putative protein of ancient origin (Gdpd1) alternative variant aSep08, mRNA.
<a href="#">Gdpd1</a>	<a href="#">Gdpd1.cSep08</a>	<a href="#">303407</a>	8132	1604	1	117	putative protein of ancient origin (Gdpd1) alternative variant cSep08, mRNA.
<a href="#">Gdpd2</a>	<a href="#">Gdpd2.bSep08</a>	<a href="#">302421</a>	1990	919	5	134	putative protein (15.3 kD) (Gdpd2) alternative variant bSep08, mRNA.
<a href="#">Gdpd2</a>	<a href="#">Gdpd2.cSep08</a>	<a href="#">302421</a>	2899	605	5	123	putative endoplasmic reticulum protein, with a transmembrane domain, of vertebrate origin (14.4 kD) (Gdpd2) alternative variant cSep08, mRNA.
<a href="#">Gdpd2</a>	<a href="#">Gdpd2.dSep08</a>	<a href="#">302421</a>	2699	425	4	112	putative endoplasmic reticulum protein, with a transmembrane domain, of vertebrate origin (13.0 kD) (Gdpd2) alternative variant dSep08, mRNA.
<a href="#">Gdpd2</a>	<a href="#">Gdpd2.eSep08</a>	<a href="#">302421</a>	791	647	2	78	putative protein of vertebrate origin (Gdpd2) alternative variant eSep08, mRNA.
<a href="#">Gdpd3</a>	<a href="#">Gdpd3.aSep08</a>	<a href="#">293490</a>	9458	998	9	156	putative protein of ancient origin (18.4 kD) (Gdpd3) alternative variant aSep08, mRNA.
<a href="#">geeby</a>	<a href="#">geeby.aSep08</a>		5887	915	10	256	histone deacetylase 6 (geeby) alternative variant aSep08, mRNA.
<a href="#">geechy</a>	<a href="#">geechy.aSep08</a>		15964	1018		339	nuclear protein 3 (geechy) mRNA.
<a href="#">geedar</a>	<a href="#">geedar.aSep08</a>		218	149		44	putative protein (geedar) mRNA.
<a href="#">geefer</a>	<a href="#">geefer.aSep08</a>		12600	672		11	putative protein (1.3 kD) (geefer) mRNA.
<a href="#">geeflo</a>	<a href="#">geeflo.aSep08</a>		691	571		34	putative protein (geeflo) mRNA.
<a href="#">geeflu</a>	<a href="#">geeflu.aSep08</a>		14617	488		39	putative protein (geeflu) mRNA.
<a href="#">geekee</a>	<a href="#">geekee.aSep08</a>		9034	542		40	putative protein (geekee) mRNA.
<a href="#">geeloy</a>	<a href="#">geeloy.aSep08</a>		26180	725		34	putative protein (4.0 kD) (geeloy) mRNA.
<a href="#">geemee</a>	<a href="#">geemee.aSep08</a>		13709	734		55	putative protein (geemee) mRNA.
<a href="#">geemer</a>	<a href="#">geemer.aSep08</a>		8352	748	2	130	putative protein of mammalian origin (geemer) alternative variant aSep08, mRNA.
<a href="#">geemer</a>	<a href="#">geemer.bSep08</a>		8341	632	1	129	putative protein of mammalian origin (geemer) alternative variant bSep08, mRNA.
<a href="#">geenoy</a>	<a href="#">geenoy.aSep08</a>		5276	584	5	194	tbc1 domain family member 8 (geenoy) alternative variant aSep08, mRNA.
<a href="#">geepor</a>	<a href="#">geepor.aSep08</a>		2654	413	2	111	putative protein of mammalian origin (geepor) alternative variant aSep08, mRNA.
<a href="#">geesa</a>	<a href="#">geesa.aSep08</a>		2154	381		127	transcription initiation factor tfiid (geesa) mRNA.
<a href="#">geeshee</a>	<a href="#">geeshee.aSep08</a>		3962	383		51	putative protein (5.8 kD) (geeshee) mRNA.
<a href="#">geeto</a>	<a href="#">geeto.aSep08</a>		7219	500		118	putative protein (geeto) mRNA.
<a href="#">geevar</a>	<a href="#">geevar.aSep08</a>		26779	604		31	putative protein (3.7 kD) (geevar) mRNA.
<a href="#">geewey</a>	<a href="#">geewey.aSep08</a>		1391	710		44	putative protein (5.0 kD) (geewey) mRNA.
<a href="#">Geft</a>	<a href="#">Geft.bSep08</a>	<a href="#">314904</a>	6829	2260	15	531	RhoA/RAC/CDC42 exchange factor (59.0 kD) (Geft) alternative variant bSep08, mRNA.
<a href="#">Geft</a>	<a href="#">Geft.cSep08</a>	<a href="#">314904</a>	3076	658	5	172	RhoA/RAC/CDC42 exchange factor (Geft) alternative variant cSep08, mRNA.

<a href="#">Geft</a>	<a href="#">Geft.dSep08</a>	<a href="#">314904</a>	1058	801	3	99	RhoA/RAC/CDC42 exchange factor (Geft) alternative variant dSep08, mRNA.
<a href="#">Gelsolin.0</a>	<a href="#">Gelsolin.0.aSep08</a>		7999	429	5	142	villin-like (Gelsolin.0) alternative variant aSep08, mRNA.
<a href="#">Gemin6</a>	<a href="#">Gemin6.aSep08</a>	<a href="#">362688</a>	4435	1312	2	179	gem (nuclear organelle) associated protein 6 (19.9 kD) (Gemin6) alternative variant aSep08, complete mRNA.
<a href="#">Gemin6</a>	<a href="#">Gemin6.cSep08</a>	<a href="#">362688</a>	2051	519	1	51	gem (nuclear organelle) associated protein 6 (6.1 kD) (Gemin6) alternative variant cSep08, mRNA.
<a href="#">Gen1</a>	<a href="#">Gen1.bSep08</a>	<a href="#">298884</a>	4553	353	1	72	putative protein (Gen1) alternative variant bSep08, mRNA.
<a href="#">gerby</a>	<a href="#">gerby.aSep08</a>		3822	491	2	144	transcription factor E3 CRA b (gerby) alternative variant aSep08, mRNA.
<a href="#">gerby</a>	<a href="#">gerby.bSep08</a>		3902	590	2	140	transcription factor (gerby) alternative variant bSep08, mRNA.
<a href="#">gerby</a>	<a href="#">gerby.cSep08</a>		2722	328	2	109	transcription factor E3 CRA b (gerby) alternative variant cSep08, mRNA.
<a href="#">gerchy</a>	<a href="#">gerchy.aSep08</a>		1960	457		16	putative protein (2.0 kD) (gerchy) mRNA.
<a href="#">gerdar</a>	<a href="#">gerdar.aSep08</a>		1187	584		68	putative protein (gerdar) mRNA.
<a href="#">gerfer</a>	<a href="#">gerfer.aSep08</a>		610	424		59	putative protein (gerfer) mRNA.
<a href="#">gerflo</a>	<a href="#">gerflo.aSep08</a>		625	335		67	putative protein (gerflo) mRNA.
<a href="#">gerflu</a>	<a href="#">gerflu.aSep08</a>		3366	368		122	putative protein of eukaryotic origin (gerflu) mRNA.
<a href="#">gerkee</a>	<a href="#">gerkee.aSep08</a>		10092	725		66	putative protein (gerkee) mRNA.
<a href="#">gerloy</a>	<a href="#">gerloy.aSep08</a>		8906	382		85	CRA b like (gerloy) mRNA.
<a href="#">germee</a>	<a href="#">germee.aSep08</a>		8698	673	2	40	putative protein (4.8 kD) (germee) alternative variant aSep08, mRNA.
<a href="#">germer</a>	<a href="#">germer.aSep08</a>		15892	843		40	putative protein (4.3 kD) (germer) mRNA.
<a href="#">gernoy</a>	<a href="#">gernoy.bSep08</a>		1762	507	2	35	putative protein (gernoy) alternative variant bSep08, mRNA.
<a href="#">gerpor</a>	<a href="#">gerpor.aSep08</a>		3065	313		63	putative protein (gerpor) mRNA.
<a href="#">gersa</a>	<a href="#">gersa.aSep08</a>		7139	623		136	transcription initiation factor tfiid (gersa) mRNA.
<a href="#">gershee</a>	<a href="#">gershee.aSep08</a>		26550	1684		76	putative secreted or extracellular protein precursor (8.8 kD) (gershee) mRNA.
<a href="#">gerto</a>	<a href="#">gerto.aSep08</a>		1076	338		112	fam59b (gerto) mRNA.
<a href="#">gervar</a>	<a href="#">gervar.aSep08</a>		8590	224		69	putative protein (gervar) mRNA.
<a href="#">gerwey</a>	<a href="#">gerwey.aSep08</a>		37748	532		66	putative protein (7.3 kD) (gerwey) mRNA.
<a href="#">geyby</a>	<a href="#">geyby.aSep08</a>		5146	610	1	115	putative protein, with a coiled coil domain, of bilateral origin (geyby) alternative variant aSep08, mRNA.
<a href="#">geyby</a>	<a href="#">geyby.bSep08</a>		3483	349	1	79	JM11 protein CRA c like (geyby) alternative variant bSep08, mRNA.
<a href="#">geychy</a>	<a href="#">geychy.aSep08</a>		3266	1007		335	sodium channel voltage-gated type VII alpha (geychy) mRNA.
<a href="#">geydar</a>	<a href="#">geydar.aSep08</a>		34778	636		41	putative protein (4.7 kD) (geydar) mRNA.
<a href="#">geyfer</a>	<a href="#">geyfer.aSep08</a>		37337	311		36	putative protein (geyfer) mRNA.
<a href="#">geyflo</a>	<a href="#">geyflo.aSep08</a>		43886	791		54	putative protein (geyflo) mRNA.
<a href="#">geyflu</a>	<a href="#">geyflu.aSep08</a>		6479	366		121	somatostatin receptor-interacting protein splice like (geyflu) mRNA.

<a href="#">geykee</a>	<a href="#">geykee.aSep08</a>		15471	672		83	CRA a like (9.7 kD) (geykee) mRNA.
<a href="#">geyloy</a>	<a href="#">geyloy.aSep08</a>		31947	252		38	putative protein (geyloy) mRNA.
<a href="#">geymee</a>	<a href="#">geymee.aSep08</a>		16561	754		71	putative protein (8.3 kD) (geymee) mRNA.
<a href="#">geymer</a>	<a href="#">geymer.bSep08</a>		2594	1448	3	6	putative protein (0.8 kD) (geymer) alternative variant bSep08, mRNA.
<a href="#">geynoy</a>	<a href="#">geynoy.aSep08</a>		1413	410		71	putative nuclear protein (7.8 kD) (geynoy) mRNA.
<a href="#">geypor</a>	<a href="#">geypor.aSep08</a>		18586	1195	2	60	putative protein (geypor) alternative variant aSep08, mRNA.
<a href="#">geypor</a>	<a href="#">geypor.bSep08</a>		24623	695	3	57	putative protein (geypor) alternative variant bSep08, mRNA.
<a href="#">geysa</a>	<a href="#">geysa.aSep08</a>		1420	1308		14	putative protein (1.7 kD) (geysa) mRNA.
<a href="#">geyshee</a>	<a href="#">geyshee.aSep08</a>		21022	388		106	putative protein (geyshee) mRNA.
<a href="#">geyto</a>	<a href="#">geyto.aSep08</a>		1282	479		37	putative protein (geyto) mRNA.
<a href="#">geyvar</a>	<a href="#">geyvar.aSep08</a>		7063	696		93	CRA a like (geyvar) mRNA.
<a href="#">geywey</a>	<a href="#">geywey.aSep08</a>		2536	204		47	putative protein (geywey) mRNA.
<a href="#">Gfap</a>	<a href="#">Gfap.bSep08</a>	<a href="#">24387</a>	6453	1080	7	299	glial fibrillary acidic protein (Gfap) alternative variant bSep08, mRNA.
<a href="#">Gfap</a>	<a href="#">Gfap.cSep08</a>	<a href="#">24387</a>	2537	996	4	160	glial fibrillary acidic protein (Gfap) alternative variant cSep08, mRNA.
<a href="#">Gfi1</a>	<a href="#">Gfi1.bSep08</a>	<a href="#">24388</a>	4713	422	1	101	growth factor independent 1 (Gfi1) alternative variant bSep08, mRNA.
<a href="#">Gfm2</a>	<a href="#">Gfm2.aSep08</a>	<a href="#">294672</a>	11019	2024	2	352	G elongation factor, mitochondrial 2 (Gfm2) alternative variant aSep08, mRNA.
<a href="#">Gfm2</a>	<a href="#">Gfm2.cSep08</a>	<a href="#">294672</a>	722	560	2	58	G elongation factor, mitochondrial 2 (Gfm2) alternative variant cSep08, mRNA.
<a href="#">Gfod1</a>	<a href="#">Gfod1.aSep08</a>	<a href="#">306842</a>	101943	355		118	putative protein of ancient origin (Gfod1) mRNA.
<a href="#">Gfpt1</a>	<a href="#">Gfpt1.aSep08</a>	<a href="#">297417</a>	18647	1444		324	glutamine fructose-6-phosphate transaminase 1 (Gfpt1) mRNA.
<a href="#">Gfra1</a>	<a href="#">Gfra1.bSep08</a>	<a href="#">25454</a>	191527	728	3	211	glial cell line derived neurotrophic factor family receptor alpha 1 (Gfra1) alternative variant bSep08, mRNA.
<a href="#">Gfra1</a>	<a href="#">Gfra1.cSep08</a>	<a href="#">25454</a>	218035	785	3	131	glial cell line derived neurotrophic factor family receptor alpha 1 (13.7 kD) (Gfra1) alternative variant cSep08, mRNA.
<a href="#">Gga1</a>	<a href="#">Gga1.bSep08</a>	<a href="#">300066</a>	10877	1201	10	296	golgi associated, gamma adaptin ear containing, ARF binding protein 1 (Gga1) alternative variant bSep08, mRNA.
<a href="#">Gga1</a>	<a href="#">Gga1.cSep08</a>	<a href="#">300066</a>	3027	777	4	115	golgi associated, gamma adaptin ear containing, ARF binding protein 1 (Gga1) alternative variant cSep08, mRNA.
<a href="#">Gga2</a>	<a href="#">Gga2.aSep08</a>	<a href="#">293455</a>	33574	2770		604	golgi associated, gamma adaptin ear containing, ARF binding protein 2 (66.2 kD) (Gga2) complete mRNA.
<a href="#">Gga3</a>	<a href="#">Gga3.bSep08</a>	<a href="#">360658</a>	2741	1178	7	392	golgi associated, gamma adaptin ear containing, ARF binding protein 3 (Gga3) alternative variant bSep08, mRNA.



<a href="#">Gga3</a>	<a href="#">Gga3.cSep08</a>	<a href="#">360658</a>	619	503	2	167	golgi associated, gamma adaptin ear containing, ARF binding protein 3 (Gga3) alternative variant cSep08, mRNA.
<a href="#">Gga3</a>	<a href="#">Gga3.dSep08</a>	<a href="#">360658</a>	790	419	3	71	golgi associated, gamma adaptin ear containing, ARF binding protein 3 (Gga3) alternative variant dSep08, mRNA.
<a href="#">Gga3</a>	<a href="#">Gga3.eSep08</a>	<a href="#">360658</a>	789	693	2	34	golgi associated, gamma adaptin ear containing, ARF binding protein 3 (3.7 kD) (Gga3) alternative variant eSep08, mRNA.
<a href="#">Ggcx</a>	<a href="#">Ggcx.bSep08</a>	<a href="#">81716</a>	2803	1290	6	230	gamma-glutamyl carboxylase CRA c (26.5 kD) (Ggcx) alternative variant bSep08, mRNA.
<a href="#">Ggcx</a>	<a href="#">Ggcx.cSep08</a>	<a href="#">81716</a>	5682	2005	5	224	gamma-glutamyl carboxylase CRA b (26.0 kD) (Ggcx) alternative variant cSep08, complete mRNA.
<a href="#">Ggcx</a>	<a href="#">Ggcx.dSep08</a>	<a href="#">81716</a>	5429	701	5	201	gamma-glutamyl carboxylase CRA c (Ggcx) alternative variant dSep08, mRNA.
<a href="#">Ggcx</a>	<a href="#">Ggcx.eSep08</a>	<a href="#">81716</a>	2880	856	3	95	gamma-glutamyl carboxylase CRA c (10.7 kD) (Ggcx) alternative variant eSep08, mRNA.
<a href="#">Ggn</a>	<a href="#">Ggn.bSep08</a>	<a href="#">292765</a>	1976	1398	2	438	gametogenetin (Ggn) alternative variant bSep08, mRNA.
<a href="#">Ggnbp2</a>	<a href="#">Ggnbp2.bSep08</a>	<a href="#">360584</a>	15164	1148	7	382	gametogenetin binding protein 2 (Ggnbp2) alternative variant bSep08, mRNA.
<a href="#">Ggps1</a>	<a href="#">Ggps1.bSep08</a>	<a href="#">291211</a>	2198	714	2	226	geranylgeranyl diphosphate synthase 1 (Ggps1) alternative variant bSep08, mRNA.
<a href="#">Ggt7</a>	<a href="#">Ggt7.bSep08</a>	<a href="#">156275</a>	7371	730	7	242	gamma-glutamyltransferase 7 (Ggt7) alternative variant bSep08, mRNA.
<a href="#">Ggt7</a>	<a href="#">Ggt7.cSep08</a>	<a href="#">156275</a>	5360	571	4	167	gamma-glutamyltransferase 7 (Ggt7) alternative variant cSep08, mRNA.
<a href="#">Ggt7</a>	<a href="#">Ggt7.dSep08</a>	<a href="#">156275</a>	3558	423	3	140	gamma-glutamyltransferase 7 (Ggt7) alternative variant dSep08, mRNA.
<a href="#">Ggta1</a>	<a href="#">Ggta1.bSep08</a>	<a href="#">246766</a>	77632	3396	7	371	glycoprotein galactosyltransferase alpha 1 3 (43.5 kD) (Ggta1) alternative variant bSep08, mRNA.
<a href="#">Ggta1</a>	<a href="#">Ggta1.cSep08</a>	<a href="#">246766</a>	40759	735	2	88	putative mitochondrial protein (9.9 kD) (Ggta1) alternative variant cSep08, mRNA.
<a href="#">Gh1</a>	<a href="#">Gh1.bSep08</a>	<a href="#">24391</a>	1753	583	4	173	growth hormone 1 (Gh1) alternative variant bSep08, mRNA.
<a href="#">Gh1</a>	<a href="#">Gh1.cSep08</a>	<a href="#">24391</a>	1979	665	4	167	growth hormone 1 (19.1 kD) (Gh1) alternative variant cSep08, complete mRNA.
<a href="#">Gh1</a>	<a href="#">Gh1.dSep08</a>	<a href="#">24391</a>	1977	337	1	79	growth hormone 1 (Gh1) alternative variant dSep08, mRNA.
<a href="#">Gh1</a>	<a href="#">Gh1.eSep08</a>	<a href="#">24391</a>	1144	348	2	32	growth hormone 1 (Gh1) alternative variant eSep08, mRNA.
<a href="#">Ghdc</a>	<a href="#">Ghdc.aSep08</a>	<a href="#">303542</a>	2044	1140	5	215	ghdc mouse gh3 domain-containing protein like (Ghdc) alternative variant aSep08, mRNA.
<a href="#">Ghdc</a>	<a href="#">Ghdc.bSep08</a>	<a href="#">303542</a>	1662	487	3	85	ghdc mouse gh3 domain-containing protein like (Ghdc) alternative variant bSep08, mRNA.
<a href="#">Ghitm</a>	<a href="#">Ghitm.bSep08</a>	<a href="#">290596</a>	5190	643	3	151	growth hormone inducible transmembrane protein (15.7 kD) (Ghitm) alternative variant bSep08, mRNA.

<a href="#">Ghitm</a>	<a href="#">Ghitm.cSep08</a>	<a href="#">290596</a>	3342	624	4	106	growth hormone inducible transmembrane protein (Ghitm) alternative variant cSep08, mRNA.
<a href="#">Ghr</a>	<a href="#">Ghr.bSep08</a>	<a href="#">25235</a>	255541	1183	8	279	growth hormone receptor (32.0 kD) (Ghr) alternative variant bSep08, mRNA.
<a href="#">Ghr</a>	<a href="#">Ghr.cSep08</a>	<a href="#">25235</a>	248858	725	7	176	growth hormone receptor (Ghr) alternative variant cSep08, mRNA.
<a href="#">Gigyf1</a>	<a href="#">Gigyf1.bSep08</a>	<a href="#">304378</a>	802	486	4	143	GRB10 interacting GYF protein 1 (Gigyf1) alternative variant bSep08, mRNA.
<a href="#">Gigyf1</a>	<a href="#">Gigyf1.cSep08</a>	<a href="#">304378</a>	863	785	2	142	GRB10 interacting GYF protein 1 (Gigyf1) alternative variant cSep08, mRNA.
<a href="#">Gimap1</a>	<a href="#">Gimap1.cSep08</a>	<a href="#">312312</a>	859	365	2	60	putative protein (Gimap1) alternative variant cSep08, mRNA.
<a href="#">Gimap5</a>	<a href="#">Gimap5.dSep08</a>	<a href="#">246774</a>	57380	734	4	40	GTPase, IMAP family member 5 (Gimap5) alternative variant dSep08, mRNA.
<a href="#">Gimap8</a>	<a href="#">Gimap8.bSep08</a>	<a href="#">500112</a>	3260	705	1	161	GTPase, IMAP family member 8 (Gimap8) alternative variant bSep08, mRNA.
<a href="#">Gin1</a>	<a href="#">Gin1.bSep08</a>	<a href="#">316687</a>	7023	1435	3	312	gypsy retrotransposon integrase 1 (Gin1) alternative variant bSep08, mRNA.
<a href="#">Gin1</a>	<a href="#">Gin1.cSep08</a>	<a href="#">316687</a>	9581	1897	5	273	gypsy retrotransposon integrase 1 (Gin1) alternative variant cSep08, mRNA.
<a href="#">Gin1</a>	<a href="#">Gin1.eSep08</a>	<a href="#">316687</a>	2253	382	2	69	gypsy retrotransposon integrase 1 (Gin1) alternative variant eSep08, mRNA.
<a href="#">Gin1</a>	<a href="#">Gin1.fSep08</a>	<a href="#">316687</a>	9688	288	3	69	gypsy retrotransposon integrase 1 (Gin1) alternative variant fSep08, mRNA.
<a href="#">Gins1</a>	<a href="#">Gins1.aSep08</a>	<a href="#">499914</a>	22200	1044	3	196	GINS complex subunit 1 (Psf1 homolog) (22.9 kD) (Gins1) alternative variant aSep08, mRNA.
<a href="#">Gins3</a>	<a href="#">Gins3.bSep08</a>	<a href="#">307639</a>	3136	651	1	61	GINS complex subunit 3 (Psf3 homolog) and similar to 60S ribosomal protein L21 (Gins3) alternative variant bSep08, mRNA.
<a href="#">Gins3</a>	<a href="#">Gins3.bSep08</a>	<a href="#">688952</a>	3136	651	1	61	GINS complex subunit 3 (Psf3 homolog) and similar to 60S ribosomal protein L21 (Gins3) alternative variant bSep08, mRNA.
<a href="#">Gins4</a>	<a href="#">Gins4.bSep08</a>	<a href="#">290842</a>	11188	783	8	198	GINS complex subunit 4 (Sld5 homolog) (22.9 kD) (Gins4) alternative variant bSep08, complete mRNA.
<a href="#">Gip</a>	<a href="#">Gip.bSep08</a>	<a href="#">25040</a>	7669	608	1	164	gastric inhibitory polypeptide (Gip) alternative variant bSep08, mRNA.
<a href="#">Gipc1</a>	<a href="#">Gipc1.bSep08</a>	<a href="#">83823</a>	8404	680	1	141	putative protein (Gipc1) alternative variant bSep08, mRNA.
<a href="#">Git1</a>	<a href="#">Git1.bSep08</a>	<a href="#">83709</a>	4900	3029	12	481	G protein-coupled receptor kinase interactor 1 (53.2 kD) (Git1) alternative variant bSep08, mRNA.
<a href="#">Git1</a>	<a href="#">Git1.cSep08</a>	<a href="#">83709</a>	1046	730	4	164	G protein-coupled receptor kinase interactor 1 (Git1) alternative variant cSep08, mRNA.
<a href="#">Git1</a>	<a href="#">Git1.dSep08</a>	<a href="#">83709</a>	7436	581	7	153	G protein-coupled receptor kinase interactor 1 (Git1) alternative variant dSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.bSep08</a>	<a href="#">304546</a>	3673	1778	3	513	G protein-coupled receptor kinase 2 (Git2) alternative variant bSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.cSep08</a>	<a href="#">304546</a>	29021	1537	13	502	G protein-coupled receptor kinase 2 (Git2) alternative variant cSep08, mRNA.

<a href="#">Git2</a>	<a href="#">Git2.dSep08</a>	<a href="#">304546</a>	18849	1777	7	471	G protein-coupled receptor kinase-interactor 2 (Git2) alternative variant dSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.eSep08</a>	<a href="#">304546</a>	31442	1753	12	443	G protein-coupled receptor kinase 2 (Git2) alternative variant eSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.fSep08</a>	<a href="#">304546</a>	25403	1351	11	356	G protein-coupled receptor kinase interacting ArfGAP 2 (Git2) alternative variant fSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.gSep08</a>	<a href="#">304546</a>	14378	737	6	245	G protein-coupled receptor kinase 2 (Git2) alternative variant gSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.hSep08</a>	<a href="#">304546</a>	10293	403	4	133	G protein-coupled receptor kinase interacting ArfGAP 2 (Git2) alternative variant hSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.iSep08</a>	<a href="#">304546</a>	10446	637	6	127	G protein-coupled receptor kinase-interactor 2 (Git2) alternative variant iSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.jSep08</a>	<a href="#">304546</a>	10811	875	5	126	G protein-coupled receptor kinase-interactor 2 (Git2) alternative variant jSep08, mRNA.
<a href="#">Git2</a>	<a href="#">Git2.kSep08</a>	<a href="#">304546</a>	3078	533	3	100	G protein-coupled receptor kinase interactor 2 (Git2) alternative variant kSep08, mRNA.
<a href="#">Giyd2</a>	<a href="#">Giyd2.bSep08</a>	<a href="#">293489</a>	3830	1114	4	258	putative mitochondrial protein of eukaryotic origin (28.8 kD) (Giyd2) alternative variant bSep08, mRNA.
<a href="#">Giyd2</a>	<a href="#">Giyd2.cSep08</a>	<a href="#">293489</a>	1128	935	4	212	putative protein of eukaryotic origin (Giyd2) alternative variant cSep08, mRNA.
<a href="#">Giyd2</a>	<a href="#">Giyd2.dSep08</a>	<a href="#">293489</a>	1402	972	4	163	putative protein of eukaryotic origin (Giyd2) alternative variant dSep08, mRNA.
<a href="#">Giyd2</a>	<a href="#">Giyd2.eSep08</a>	<a href="#">293489</a>	1310	575	5	159	putative protein of mammalian origin (Giyd2) alternative variant eSep08, mRNA.
<a href="#">Giyd2</a>	<a href="#">Giyd2.fSep08</a>	<a href="#">293489</a>	512	393	2	73	putative mitochondrial protein of eukaryotic origin (8.2 kD) (Giyd2) alternative variant fSep08, mRNA.
<a href="#">Gja1</a>	<a href="#">Gja1.bSep08</a>	<a href="#">24392</a>	8097	542	2	149	gap junction protein, alpha 1 (Gja1) alternative variant bSep08, mRNA.
<a href="#">Gjb1</a>	<a href="#">Gjb1.bSep08</a>	<a href="#">29584</a>	890	553	1	174	gap junction protein, beta 1 (Gjb1) alternative variant bSep08, mRNA.
<a href="#">Gjb3</a>	<a href="#">Gjb3.bSep08</a>	<a href="#">29585</a>	2938	484	2	69	gap junction protein, beta 3 (Gjb3) alternative variant bSep08, mRNA.
<a href="#">Gjb6</a>	<a href="#">Gjb6.bSep08</a>	<a href="#">84403</a>	10336	2092	2	106	putative mitochondrial protein (11.7 kD) (Gjb6) alternative variant bSep08, mRNA.
<a href="#">Gjb6</a>	<a href="#">Gjb6.cSep08</a>	<a href="#">84403</a>	7487	429	2	42	gap junction protein beta (Gjb6) alternative variant cSep08, mRNA.
<a href="#">Gjc2</a>	<a href="#">Gjc2.aSep08</a>	<a href="#">497913</a>	6796	1051		350	gap junction protein, gamma 2 (Gjc2) mRNA.
<a href="#">Gk5</a>	<a href="#">Gk5.aSep08</a>	<a href="#">367146</a>	7943	663		148	glycerol kinase 5 (putative) (Gk5) mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.aSep08</a>	<a href="#">361202</a>	39749	1778	8	442	G kinase anchoring protein 1 (Gkap1) alternative variant aSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.cSep08</a>	<a href="#">361202</a>	30957	827	7	254	G kinase anchoring protein 1 (Gkap1) alternative variant cSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.dSep08</a>	<a href="#">361202</a>	21991	710	7	236	G kinase anchoring protein 1 (Gkap1) alternative variant dSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.eSep08</a>	<a href="#">361202</a>	28591	733	6	177	G kinase anchoring protein 1 (Gkap1) alternative variant eSep08, mRNA.

<a href="#">Gkap1</a>	<a href="#">Gkap1.fSep08</a>	<a href="#">361202</a>	34980	499	7	165	G kinase anchoring protein 1 (Gkap1) alternative variant fSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.gSep08</a>	<a href="#">361202</a>	16104	545	5	160	G kinase anchoring protein 1 (Gkap1) alternative variant gSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.hSep08</a>	<a href="#">361202</a>	30907	453	6	139	G kinase anchoring protein 1 (Gkap1) alternative variant hSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.iSep08</a>	<a href="#">361202</a>	15876	307	2	102	G kinase anchoring protein 1 (Gkap1) alternative variant iSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.jSep08</a>	<a href="#">361202</a>	8914	1336	5	71	G kinase anchoring protein 1 (8.0 kD) (Gkap1) alternative variant jSep08, mRNA.
<a href="#">Gkap1</a>	<a href="#">Gkap1.kSep08</a>	<a href="#">361202</a>	39791	791	6	71	G kinase anchoring protein 1 (8.0 kD) (Gkap1) alternative variant kSep08, complete mRNA.
<a href="#">glabor</a>	<a href="#">glabor.aSep08</a>		1977	397		108	putative protein (glabor) mRNA.
<a href="#">glachy</a>	<a href="#">glachy.aSep08</a>		23630	2250		58	putative protein (6.8 kD) (glachy) mRNA.
<a href="#">gladoy</a>	<a href="#">gladoy.aSep08</a>		8838	478	3	121	protein CRA a (gladoy) alternative variant aSep08, mRNA.
<a href="#">gladoy</a>	<a href="#">gladoy.bSep08</a>		5894	447	1	89	protein CRA a (gladoy) alternative variant bSep08, mRNA.
<a href="#">glafly</a>	<a href="#">glafly.aSep08</a>		6235	664		33	putative protein (3.8 kD) (glafly) mRNA.
<a href="#">glafly</a>	<a href="#">glafly.aSep08</a>		2690	665		58	acetyl-Coenzyme A acetyltransferase 3 like (6.2 kD) (glafly) mRNA.
<a href="#">glagar</a>	<a href="#">glagar.aSep08</a>		7307	365		121	nima -related kinase 5 (glagar) mRNA.
<a href="#">glaja</a>	<a href="#">glaja.aSep08</a>		4388	1014		255	CRA b (glaja) mRNA.
<a href="#">glajey</a>	<a href="#">glajey.aSep08</a>		23515	762		55	putative protein (6.1 kD) (glajey) mRNA.
<a href="#">glakee</a>	<a href="#">glakee.aSep08</a>		947	581		83	putative protein (9.0 kD) (glakee) mRNA.
<a href="#">glalo</a>	<a href="#">glalo.aSep08</a>		1682	379		63	putative protein (glalo) mRNA.
<a href="#">glamee</a>	<a href="#">glamee.aSep08</a>		21383	617		9	putative protein (1.1 kD) (glamee) mRNA.
<a href="#">glapor</a>	<a href="#">glapor.aSep08</a>		856	357		39	putative protein (glapor) mRNA.
<a href="#">glarbor</a>	<a href="#">glarbor.aSep08</a>		17854	594	2	36	putative protein (3.9 kD) (glarbor) alternative variant aSep08, mRNA.
<a href="#">glarbor</a>	<a href="#">glarbor.bSep08</a>		15021	434	2	44	putative protein (glarbor) alternative variant bSep08, mRNA.
<a href="#">glarchy</a>	<a href="#">glarchy.aSep08</a>		37453	809	1	105	putative protein (11.5 kD) (glarchy) alternative variant aSep08, mRNA.
<a href="#">glarchy</a>	<a href="#">glarchy.bSep08</a>		52943	751	1	69	putative protein (7.6 kD) (glarchy) alternative variant bSep08, mRNA.
<a href="#">glardoy</a>	<a href="#">glardoy.aSep08</a>		1670	314		42	putative protein (glardoy) mRNA.
<a href="#">glarfee</a>	<a href="#">glarfee.aSep08</a>		3661	394		62	putative protein (6.9 kD) (glarfee) mRNA.
<a href="#">glarflu</a>	<a href="#">glarflu.aSep08</a>		4483	886		165	putative protein (17.5 kD) (glarflu) mRNA.
<a href="#">glarfly</a>	<a href="#">glarfly.aSep08</a>		9891	255		51	putative protein (glarfly) mRNA.
<a href="#">glargar</a>	<a href="#">glargar.aSep08</a>		4915	628		209	microcephaly primary autosomal recessive 1 (glargar) mRNA.
<a href="#">glarja</a>	<a href="#">glarja.aSep08</a>		4387	584		119	putative protein (glarja) mRNA.
<a href="#">glarjey</a>	<a href="#">glarjey.aSep08</a>		12875	806		183	epidermal growth factor receptor CRA b (glarjey) mRNA.
<a href="#">glarkee</a>	<a href="#">glarkee.aSep08</a>		30571	499	1	88	putative protein (glarkee) alternative variant aSep08, mRNA.

<a href="#">glarkee</a>	<a href="#">glarkee.bSep08</a>		1487	758	1	82	putative protein (glarkee) alternative variant bSep08, mRNA.
<a href="#">glarlo</a>	<a href="#">glarlo.aSep08</a>		5111	507		168	tectonic (glarlo) mRNA.
<a href="#">glarmee</a>	<a href="#">glarmee.aSep08</a>		1107	292		97	zinc finger zz-type ef-hand domain-containing protein 1 (glarmee) mRNA.
<a href="#">glaroy</a>	<a href="#">glaroy.aSep08</a>		8391	925		171	CRA b precursor (19.4 kD) (glaroy) mRNA.
<a href="#">glarpor</a>	<a href="#">glarpor.aSep08</a>		32263	651		217	cyclic AMP-regulated phosphoprotein (glarpor) mRNA.
<a href="#">glarroy</a>	<a href="#">glarroy.aSep08</a>		60651	778	2	79	putative protein (8.7 kD) (glarroy) alternative variant aSep08, mRNA.
<a href="#">glarroy</a>	<a href="#">glarroy.bSep08</a>		3835	631	1	60	putative protein (glarroy) alternative variant bSep08, mRNA.
<a href="#">glarsa</a>	<a href="#">glarsa.aSep08</a>		3073	492		35	putative protein (4.1 kD) (glarsa) mRNA.
<a href="#">glarshee</a>	<a href="#">glarshee.aSep08</a>		225855	592		197	dihydropyrimidine dehydrogenase (glarshee) mRNA.
<a href="#">glartu</a>	<a href="#">glartu.aSep08</a>		3769	334		110	uncharacterized protein homolog (glartu) mRNA.
<a href="#">glarvo</a>	<a href="#">glarvo.aSep08</a>		7390	806		50	putative protein (5.3 kD) (glarvo) mRNA.
<a href="#">glarwer</a>	<a href="#">glarwer.aSep08</a>		3602	1430		363	a kinase anchor protein 9 CRA b (glarwer) mRNA.
<a href="#">glasa</a>	<a href="#">glasa.bSep08</a>		49941	400	2	47	putative protein (glasa) alternative variant bSep08, mRNA.
<a href="#">glashee</a>	<a href="#">glashee.aSep08</a>		1018	264		87	pancreatic alpha-amylase (glashee) mRNA.
<a href="#">glatu</a>	<a href="#">glatu.aSep08</a>		888	722		44	putative protein (5.0 kD) (glatu) mRNA.
<a href="#">glavo</a>	<a href="#">glavo.aSep08</a>		1404	387		129	midasin (glavo) mRNA.
<a href="#">glawbor</a>	<a href="#">glawbor.aSep08</a>		2531	691		35	putative protein (glawbor) mRNA.
<a href="#">glawchy</a>	<a href="#">glawchy.aSep08</a>		364	304		29	putative protein (3.2 kD) (glawchy) mRNA.
<a href="#">glawdoy</a>	<a href="#">glawdoy.aSep08</a>		1936	518		48	putative protein (glawdoy) mRNA.
<a href="#">glawer</a>	<a href="#">glawer.aSep08</a>		7877	692		83	putative protein (glawer) mRNA.
<a href="#">glawfee</a>	<a href="#">glawfee.aSep08</a>		1927	890		51	putative protein (5.8 kD) (glawfee) mRNA.
<a href="#">glawflu</a>	<a href="#">glawflu.aSep08</a>		1396	750		49	CRA b like (glawflu) mRNA.
<a href="#">glawfly</a>	<a href="#">glawfly.aSep08</a>		105940	283		60	putative protein (glawfly) mRNA.
<a href="#">glawgar</a>	<a href="#">glawgar.aSep08</a>		2049	245		75	putative protein (glawgar) mRNA.
<a href="#">glawja</a>	<a href="#">glawja.aSep08</a>		2175	266	2	28	putative protein (glawja) alternative variant aSep08, mRNA.
<a href="#">glawjey</a>	<a href="#">glawjey.aSep08</a>		460	318		105	putative protein (glawjey) mRNA.
<a href="#">glawkee</a>	<a href="#">glawkee.aSep08</a>		11560	412		70	putative protein (glawkee) mRNA.
<a href="#">glawlo</a>	<a href="#">glawlo.aSep08</a>		1223	944		49	CRA a (glawlo) mRNA.
<a href="#">glawmee</a>	<a href="#">glawmee.aSep08</a>		11000	380		126	zinc finger zz-type ef-hand domain-containing protein 1 (glawmee) mRNA.
<a href="#">glawpor</a>	<a href="#">glawpor.aSep08</a>		12271	691	4	117	cyclic AMP-regulated phosphoprotein 21 CRA c (glawpor) alternative variant aSep08, mRNA.
<a href="#">glawpor</a>	<a href="#">glawpor.bSep08</a>		4008	470	3	88	cyclic AMP-regulated phosphoprotein 21 CRA e (9.6 kD) (glawpor) alternative variant bSep08, mRNA.
<a href="#">glawpor</a>	<a href="#">glawpor.cSep08</a>		4772	634	3	88	cyclic AMP-regulated phosphoprotein 21 CRA e (9.6 kD) (glawpor) alternative variant cSep08, mRNA.
<a href="#">glawpor</a>	<a href="#">glawpor.dSep08</a>		2444	652	2	87	cyclic AMP-regulated phosphoprotein 21 CRA e (9.5 kD) (glawpor) alternative variant dSep08, mRNA.

<a href="#">glawroy</a>	<a href="#">glawroy.aSep08</a>		13973	362		111	heterogeneous nuclear ribonucleoprotein M (glawroy) mRNA.
<a href="#">glawsa</a>	<a href="#">glawsa.aSep08</a>		2167	349		116	p450 2c2 (glawsa) mRNA.
<a href="#">glawshee</a>	<a href="#">glawshee.aSep08</a>		100398	454		151	dihydropyrimidine dehydrogenase (glawshee) mRNA.
<a href="#">glawtu</a>	<a href="#">glawtu.aSep08</a>		1419	139		45	leucine rich repeat containing 9 like (glawtu) mRNA.
<a href="#">glawvo</a>	<a href="#">glawvo.aSep08</a>		1508	415		25	putative protein (glawvo) mRNA.
<a href="#">glawwer</a>	<a href="#">glawwer.aSep08</a>		684	537		83	putative protein (glawwer) mRNA.
<a href="#">Glb1</a>	<a href="#">Glb1.bSep08</a>	<a href="#">316033</a>	50453	1514	12	337	galactosidase beta 1 (37.3 kD) (Glb1) alternative variant bSep08, mRNA.
<a href="#">Glb1</a>	<a href="#">Glb1.cSep08</a>	<a href="#">316033</a>	15904	727	6	242	galactosidase beta 1 CRA b (Glb1) alternative variant cSep08, mRNA.
<a href="#">Glb1</a>	<a href="#">Glb1.dSep08</a>	<a href="#">316033</a>	10797	738	5	163	galactosidase beta 1 CRA b (Glb1) alternative variant dSep08, mRNA.
<a href="#">Glb112</a>	<a href="#">Glb112.aSep08</a>	<a href="#">503194</a>	72133	2866		637	galactosidase, beta 1-like 2 (Glb112) mRNA.
<a href="#">Glce</a>	<a href="#">Glce.aSep08</a>	<a href="#">363073</a>	54646	394		62	glucuronic acid epimerase (Glce) mRNA.
<a href="#">Gldc</a>	<a href="#">Gldc.bSep08</a>	<a href="#">309312</a>	47923	2061	2	620	glycine dehydrogenase (decarboxylating) (Gldc) alternative variant bSep08, mRNA.
<a href="#">Gldc</a>	<a href="#">Gldc.cSep08</a>	<a href="#">309312</a>	25243	2428	9	361	glycine dehydrogenase (decarboxylating) (40.1 kD) (Gldc) alternative variant cSep08, mRNA.
<a href="#">Gldc</a>	<a href="#">Gldc.dSep08</a>	<a href="#">309312</a>	52835	697	1	169	glycine dehydrogenase (decarboxylating) (Gldc) alternative variant dSep08, complete mRNA.
<a href="#">Gldc</a>	<a href="#">Gldc.eSep08</a>	<a href="#">309312</a>	1174	514	1	74	glycine dehydrogenase (decarboxylating) (8.5 kD) (Gldc) alternative variant eSep08, mRNA.
<a href="#">Gle1</a>	<a href="#">Gle1.bSep08</a>	<a href="#">362098</a>	9804	634	6	187	GLE1 RNA export mediator (Gle1) alternative variant bSep08, mRNA.
<a href="#">Gle1</a>	<a href="#">Gle1.cSep08</a>	<a href="#">362098</a>	15782	1778	7	151	RNA export mediator (Gle1) alternative variant cSep08, mRNA.
<a href="#">Gle1</a>	<a href="#">Gle1.eSep08</a>	<a href="#">362098</a>	4642	508	3	65	GLE1 RNA export mediator (7.9 kD) (Gle1) alternative variant eSep08, mRNA.
<a href="#">Gle1</a>	<a href="#">Gle1.fSep08</a>	<a href="#">362098</a>	1524	892	2	60	GLE1 RNA export mediator (6.8 kD) (Gle1) alternative variant fSep08, mRNA.
<a href="#">Gle1</a>	<a href="#">Gle1.gSep08</a>	<a href="#">362098</a>	2739	768	2	44	GLE1 RNA export mediator (4.9 kD) (Gle1) alternative variant gSep08, mRNA.
<a href="#">gleebor</a>	<a href="#">gleebor.aSep08</a>		845	652		60	putative protein (6.8 kD) (gleebor) mRNA.
<a href="#">gleechy</a>	<a href="#">gleechy.aSep08</a>		4090	345		49	putative protein (gleechy) mRNA.
<a href="#">gleedoy</a>	<a href="#">gleedoy.aSep08</a>		18375	619		66	putative protein (7.9 kD) (gleedoy) mRNA.
<a href="#">gleefee</a>	<a href="#">gleefee.aSep08</a>		2505	254		65	putative protein (7.3 kD) (gleefee) mRNA.
<a href="#">gleeflu</a>	<a href="#">gleeflu.aSep08</a>		6614	366		119	CRA b (gleeflu) mRNA.
<a href="#">gleefly</a>	<a href="#">gleefly.aSep08</a>		3281	213		23	putative protein (2.7 kD) (gleefly) mRNA.
<a href="#">gleegar</a>	<a href="#">gleegar.bSep08</a>		3241	199		63	gag protein like (gleegar) alternative variant bSep08, mRNA.
<a href="#">gleeja</a>	<a href="#">gleeja.aSep08</a>		870	262		43	putative protein (gleeja) mRNA.
<a href="#">gleejey</a>	<a href="#">gleejey.aSep08</a>		7563	609		50	putative protein (5.8 kD) (gleejey) mRNA.
<a href="#">gleekee</a>	<a href="#">gleekee.aSep08</a>		552	457		63	putative protein (gleekee) mRNA.
<a href="#">gleelo</a>	<a href="#">gleelo.bSep08</a>		2139	676		60	tectonic (gleelo) alternative variant bSep08, mRNA.

<a href="#">gleemee</a>	<a href="#">gleemee.aSep08</a>		2734	567		129	zinc finger ZZ-type with EF hand domain 1 (gleemee) mRNA.
<a href="#">gleepor</a>	<a href="#">gleepor.aSep08</a>		23148	791		78	putative protein (8.7 kD) (gleepor) mRNA.
<a href="#">gleeroy</a>	<a href="#">gleeroy.aSep08</a>		3215	199		63	gag protein like (gleeroy) mRNA.
<a href="#">gleesa</a>	<a href="#">gleesa.aSep08</a>		1368	1057		200	CRA a like (gleesa) mRNA.
<a href="#">gleeshee</a>	<a href="#">gleeshee.aSep08</a>		10191	375		45	putative protein (4.9 kD) (gleeshee) mRNA.
<a href="#">gleetu</a>	<a href="#">gleetu.aSep08</a>		9307	511		161	repeat containing 9 (gleetu) mRNA.
<a href="#">gleevo</a>	<a href="#">gleevo.aSep08</a>		6804	800		97	putative protein (gleevo) mRNA.
<a href="#">gleewer</a>	<a href="#">gleewer.aSep08</a>		3998	657		33	putative protein (3.8 kD) (gleewer) mRNA.
<a href="#">glerbor</a>	<a href="#">glerbor.aSep08</a>		1041	516		89	putative protein (glerbor) mRNA.
<a href="#">glerchy</a>	<a href="#">glerchy.aSep08</a>		628	311		66	putative protein (glerchy) mRNA.
<a href="#">glerdoy</a>	<a href="#">glerdoy.aSep08</a>		2516	410		77	putative protein (glerdoy) mRNA.
<a href="#">glerfee</a>	<a href="#">glerfee.aSep08</a>		2844	659		96	putative protein (glerfee) mRNA.
<a href="#">glerflu</a>	<a href="#">glerflu.aSep08</a>		7979	588		140	otoancorin (glerflu) mRNA.
<a href="#">glerfly</a>	<a href="#">glerfly.aSep08</a>		7309	547	2	122	putative mitochondrial protein (13.7 kD) (glerfly) alternative variant aSep08, mRNA.
<a href="#">glerfly</a>	<a href="#">glerfly.bSep08</a>		1465	879	2	87	putative mitochondrial protein (9.8 kD) (glerfly) alternative variant bSep08, mRNA.
<a href="#">glergar</a>	<a href="#">glergar.aSep08</a>		65896	377		70	putative protein (glergar) mRNA.
<a href="#">glerja</a>	<a href="#">glerja.aSep08</a>		23889	393		59	putative protein (glerja) mRNA.
<a href="#">glerjey</a>	<a href="#">glerjey.aSep08</a>		5932	774		56	putative protein (6.4 kD) (glerjey) mRNA.
<a href="#">glerkee</a>	<a href="#">glerkee.aSep08</a>		1077	532		109	putative protein (glerkee) mRNA.
<a href="#">glerlo</a>	<a href="#">glerlo.aSep08</a>		20310	454		150	CRA a (glerlo) mRNA.
<a href="#">glermee</a>	<a href="#">glermee.aSep08</a>		4684	702		229	CRA b (glermee) mRNA.
<a href="#">glerpor</a>	<a href="#">glerpor.aSep08</a>		13652	777		26	putative protein (glerpor) mRNA.
<a href="#">glerroy</a>	<a href="#">glerroy.bSep08</a>		1852	517	2	34	putative protein (glerroy) alternative variant bSep08, mRNA.
<a href="#">glerroy</a>	<a href="#">glerroy.cSep08</a>		19159	413	3	35	putative protein (glerroy) alternative variant cSep08, mRNA.
<a href="#">glersa</a>	<a href="#">glersa.aSep08</a>	<a href="#">503178</a>	6772	680		97	tubulin tyrosine ligase-like 1 (glersa) mRNA.
<a href="#">glershee</a>	<a href="#">glershee.aSep08</a>		1609	731		84	putative protein (glershee) mRNA.
<a href="#">glertu</a>	<a href="#">glertu.aSep08</a>		618	407		135	CRA b (glertu) mRNA.
<a href="#">glervo</a>	<a href="#">glervo.aSep08</a>		12112	407		74	putative protein (glervo) mRNA.
<a href="#">glerwer</a>	<a href="#">glerwer.aSep08</a>		2488	442		76	putative nuclear protein (8.8 kD) (glerwer) mRNA.
<a href="#">gleybor</a>	<a href="#">gleybor.aSep08</a>		1288	579		39	putative protein (gleybor) mRNA.
<a href="#">gleychy</a>	<a href="#">gleychy.aSep08</a>		3906	673		50	putative protein (5.3 kD) (gleychy) mRNA.
<a href="#">gleydoy</a>	<a href="#">gleydoy.aSep08</a>		12956	806		256	treacher Collins Franceschetti syndrome 1 homolog like (gleydoy) mRNA.
<a href="#">gleyfee</a>	<a href="#">gleyfee.aSep08</a>		941	423		95	putative secreted or extracellular protein precursor (10.7 kD) (gleyfee) mRNA.
<a href="#">gleyflu</a>	<a href="#">gleyflu.aSep08</a>		1361	531		42	putative protein (gleyflu) mRNA.
<a href="#">gleyfly</a>	<a href="#">gleyfly.aSep08</a>		10278	542		66	putative protein (6.9 kD) (gleyfly) mRNA.
<a href="#">gleygar</a>	<a href="#">gleygar.aSep08</a>		8064	284		56	putative protein (gleygar) mRNA.

<a href="#">gleyja</a>	<a href="#">gleyja.aSep08</a>		1587	741	2	59	putative protein (6.7 kD) (gleyja) alternative variant aSep08, mRNA.
<a href="#">gleyjey</a>	<a href="#">gleyjey.aSep08</a>		8566	875		241	centrosomal protein 68 (gleyjey) mRNA.
<a href="#">gleykee</a>	<a href="#">gleykee.aSep08</a>		1276	535	1	38	putative protein (gleykee) alternative variant aSep08, mRNA.
<a href="#">gleykee</a>	<a href="#">gleykee.bSep08</a>		1294	471	1	43	putative protein (4.9 kD) (gleykee) alternative variant bSep08, mRNA.
<a href="#">gleylo</a>	<a href="#">gleylo.aSep08</a>		11681	505	2	138	putative protein, with a coiled coil domain, of vertebrate origin (gleylo) alternative variant aSep08, mRNA.
<a href="#">gleylo</a>	<a href="#">gleylo.bSep08</a>		5924	376	2	125	putative protein, with a coiled coil domain, of vertebrate origin (gleylo) alternative variant bSep08, mRNA.
<a href="#">gleylo</a>	<a href="#">gleylo.cSep08</a>		5924	477	3	120	putative protein (gleylo) alternative variant cSep08, mRNA.
<a href="#">gleylo</a>	<a href="#">gleylo.dSep08</a>		11651	576	3	120	putative protein, with a coiled coil domain, of vertebrate origin (gleylo) alternative variant dSep08, mRNA.
<a href="#">gleylo</a>	<a href="#">gleylo.eSep08</a>		11674	666	3	119	putative protein, with a coiled coil domain, of vertebrate origin (gleylo) alternative variant eSep08, mRNA.
<a href="#">gleymee</a>	<a href="#">gleymee.aSep08</a>		2717	677		100	putative protein (gleymee) mRNA.
<a href="#">gleypor</a>	<a href="#">gleypor.aSep08</a>		1086	313		84	putative protein (gleypor) mRNA.
<a href="#">gleyroy</a>	<a href="#">gleyroy.aSep08</a>		2707	799		66	putative protein (gleyroy) mRNA.
<a href="#">gleyshee</a>	<a href="#">gleyshee.aSep08</a>		15421	816		72	putative protein (gleyshee) mRNA.
<a href="#">gleytu</a>	<a href="#">gleytu.aSep08</a>		3351	571		111	pol protein (gleytu) mRNA.
<a href="#">gleyvo</a>	<a href="#">gleyvo.aSep08</a>		80310	376		60	putative protein (gleyvo) mRNA.
<a href="#">gleywer</a>	<a href="#">gleywer.aSep08</a>		466	364		18	putative protein (gleywer) mRNA.
<a href="#">Glg1</a>	<a href="#">Glg1.bSep08</a>	<a href="#">29476</a>	3684	602	4	160	golgi apparatus protein 1 (Glg1) alternative variant bSep08, mRNA.
<a href="#">Glg1</a>	<a href="#">Glg1.cSep08</a>	<a href="#">29476</a>	4162	2524	2	107	golgi apparatus protein 1 (Glg1) alternative variant cSep08, mRNA.
<a href="#">Glg1</a>	<a href="#">Glg1.dSep08</a>	<a href="#">29476</a>	1131	426	2	88	golgi apparatus protein 1 (Glg1) alternative variant dSep08, mRNA.
<a href="#">Glg1</a>	<a href="#">Glg1.eSep08</a>	<a href="#">29476</a>	2176	277	3	80	golgi apparatus protein 1 (Glg1) alternative variant eSep08, mRNA.
<a href="#">Gli3</a>	<a href="#">Gli3.aSep08</a>	<a href="#">140588</a>	75556	404		122	GLI-Kruppel family member GLI3 (Gli3) mRNA.
<a href="#">Glipr1</a>	<a href="#">Glipr1.bSep08</a>	<a href="#">299783</a>	10973	792	4	177	GLI pathogenesis-related 1 (glioma) (20.2 kD) (Glipr1) alternative variant bSep08, mRNA.
<a href="#">Glipr1</a>	<a href="#">Glipr1.cSep08</a>	<a href="#">299783</a>	10882	787	5	91	GLI pathogenesis-related 1 (glioma) (Glipr1) alternative variant cSep08, mRNA.
<a href="#">Glipr2</a>	<a href="#">Glipr2.aSep08</a>	<a href="#">679819</a>	20177	686		169	GLI pathogenesis-related 2 (Glipr2) mRNA.
<a href="#">Glis1</a>	<a href="#">Glis1.aSep08</a>	<a href="#">298732</a>	7839	914	4	197	GLIS family zinc finger 1 (Glis1) alternative variant aSep08, mRNA.
<a href="#">Glis1</a>	<a href="#">Glis1.cSep08</a>	<a href="#">298732</a>	3113	542	2	74	GLIS family zinc finger 1 (Glis1) alternative variant cSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.bSep08</a>	<a href="#">289437</a>	44369	1965	19	458	glomulin (51.6 kD) (Glmn) alternative variant bSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.cSep08</a>	<a href="#">289437</a>	17144	817	7	242	glomulin (Glmn) alternative variant cSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.dSep08</a>	<a href="#">289437</a>	13026	588	5	177	glomulin (Glmn) alternative variant dSep08, mRNA.



<a href="#">Glmn</a>	<a href="#">Glmn.eSep08</a>	<a href="#">289437</a>	12426	495	4	129	glomulin (Glmn) alternative variant eSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.fSep08</a>	<a href="#">289437</a>	12415	523	4	127	glomulin (14.1 kD) (Glmn) alternative variant fSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.gSep08</a>	<a href="#">289437</a>	9229	389	5	125	glomulin (Glmn) alternative variant gSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.hSep08</a>	<a href="#">289437</a>	15438	762	6	97	glomulin (11.4 kD) (Glmn) alternative variant hSep08, mRNA.
<a href="#">Glmn</a>	<a href="#">Glmn.iSep08</a>	<a href="#">289437</a>	3164	620	4	82	glomulin (9.6 kD) (Glmn) alternative variant iSep08, complete mRNA.
<a href="#">Globin.0</a>	<a href="#">Globin.0.aSep08</a>		1248	1043		201	hemoglobin theta (Globin.0) mRNA.
<a href="#">Globin.1</a>	<a href="#">Globin.1.aSep08</a>		1485	595	2	142	hemoglobin zeta (16.1 kD) (Globin.1) alternative variant aSep08, complete mRNA.
<a href="#">Globin.1</a>	<a href="#">Globin.1.bSep08</a>		623	518	1	102	hemoglobin zeta (11.5 kD) (Globin.1) alternative variant bSep08, mRNA.
<a href="#">globor</a>	<a href="#">globor.aSep08</a>		5409	369		119	putative protein (globor) mRNA.
<a href="#">glochy</a>	<a href="#">glochy.aSep08</a>		11648	1801	6	458	ninein-like (glochy) alternative variant aSep08, mRNA.
<a href="#">Glod4</a>	<a href="#">Glod4.aSep08</a>	<a href="#">363644</a>	6963	649		88	putative protein (Glod4) mRNA.
<a href="#">glodoy</a>	<a href="#">glodoy.aSep08</a>		2584	400		132	multiple EGF-like-domains 10 (glodoy) mRNA.
<a href="#">glofee</a>	<a href="#">glofee.aSep08</a>		3083	723		46	putative protein (5.1 kD) (glofee) mRNA.
<a href="#">gloflu</a>	<a href="#">gloflu.aSep08</a>		67005	1210		174	CRA b (gloflu) mRNA.
<a href="#">glofly</a>	<a href="#">glofly.aSep08</a>		10640	682		81	putative cytoplasmic protein (9.0 kD) (glofly) mRNA.
<a href="#">glogar</a>	<a href="#">glogar.aSep08</a>		10718	356		27	putative protein (glogar) mRNA.
<a href="#">gloja</a>	<a href="#">gloja.aSep08</a>		8315	1388		225	zinc finger ccch domain-containing protein 13 (gloja) mRNA.
<a href="#">glojey</a>	<a href="#">glojey.aSep08</a>		9509	293		41	putative protein (glojey) mRNA.
<a href="#">glokee</a>	<a href="#">glokee.aSep08</a>		13771	449		58	putative protein (glokee) mRNA.
<a href="#">glolo</a>	<a href="#">glolo.aSep08</a>		684	410		99	putative protein (glolo) mRNA.
<a href="#">glomee</a>	<a href="#">glomee.aSep08</a>		31337	390		130	spinster (glomee) mRNA.
<a href="#">glopor</a>	<a href="#">glopor.aSep08</a>		16432	361		120	cysteine rich domain (glopor) mRNA.
<a href="#">glorbor</a>	<a href="#">glorbor.aSep08</a>		2130	452		56	putative protein (6.5 kD) (glorbor) mRNA.
<a href="#">glorchy</a>	<a href="#">glorchy.aSep08</a>		1231	400		42	putative protein (glorchy) mRNA.
<a href="#">glordoy</a>	<a href="#">glordoy.aSep08</a>		10206	748		78	putative protein (8.7 kD) (glordoy) mRNA.
<a href="#">glorfee</a>	<a href="#">glorfee.aSep08</a>		5760	556		79	putative protein (glorfee) mRNA.
<a href="#">glorflu</a>	<a href="#">glorflu.aSep08</a>		7840	1786		167	glutamyl-tRNA synthetase 2 (glorflu) alternative variant aSep08, mRNA.
<a href="#">glorfly</a>	<a href="#">glorfly.aSep08</a>		139487	610		82	putative protein of eukaryotic origin (glorfly) mRNA.
<a href="#">glorgar</a>	<a href="#">glorgar.aSep08</a>		5356	277		50	myomesin 2 (glorgar) mRNA.
<a href="#">glorja</a>	<a href="#">glorja.aSep08</a>		27522	798		42	putative protein (4.8 kD) (glorja) mRNA.
<a href="#">glorjey</a>	<a href="#">glorjey.aSep08</a>		4102	733	4	80	putative mitochondrial protein (8.6 kD) (glorjey) alternative variant aSep08, mRNA.
<a href="#">glorjey</a>	<a href="#">glorjey.bSep08</a>		5103	773	4	65	putative protein (7.2 kD) (glorjey) alternative variant bSep08, mRNA.
<a href="#">glorjey</a>	<a href="#">glorjey.cSep08</a>		5047	734	4	67	putative protein (7.4 kD) (glorjey) alternative variant cSep08, mRNA.

<a href="#">glorlo</a>	<a href="#">glorlo.aSep08</a>		2623	326		31	putative protein (3.1 kD) (glorlo) mRNA.
<a href="#">glormee</a>	<a href="#">glormee.aSep08</a>		12148	407		58	putative protein (glormee) mRNA.
<a href="#">gloroy</a>	<a href="#">gloroy.aSep08</a>		58248	404		50	putative protein (5.5 kD) (gloroy) mRNA.
<a href="#">glorpor</a>	<a href="#">glorpor.aSep08</a>		18046	1789	4	471	putative protein (glorpor) alternative variant aSep08, mRNA.
<a href="#">glorpor</a>	<a href="#">glorpor.bSep08</a>		24249	731	5	89	putative protein (9.9 kD) (glorpor) alternative variant bSep08, mRNA.
<a href="#">glorroy</a>	<a href="#">glorroy.aSep08</a>		782	644		36	putative protein (4.0 kD) (glorroy) mRNA.
<a href="#">glorshee</a>	<a href="#">glorshee.aSep08</a>		2492	790		69	putative protein (7.6 kD) (glorshee) mRNA.
<a href="#">glortu</a>	<a href="#">glortu.aSep08</a>		4764	534		85	putative mitochondrial protein (9.3 kD) (glortu) mRNA.
<a href="#">glorvo</a>	<a href="#">glorvo.aSep08</a>		5807	310		21	putative protein (2.3 kD) (glorvo) mRNA.
<a href="#">glorwer</a>	<a href="#">glorwer.aSep08</a>		1110	596		73	putative protein (glorwer) mRNA.
<a href="#">glosa</a>	<a href="#">glosa.aSep08</a>		122020	442		85	putative protein (glosa) mRNA.
<a href="#">gloshee</a>	<a href="#">gloshee.aSep08</a>		5178	701		85	putative protein (gloshee) mRNA.
<a href="#">glotu</a>	<a href="#">glotu.aSep08</a>		5246	616	3	63	uncharacterized protein homolog like (glotu) alternative variant aSep08, mRNA.
<a href="#">glovo</a>	<a href="#">glovo.aSep08</a>		12346	2842		524	midasin (glovo) mRNA.
<a href="#">glower</a>	<a href="#">glower.aSep08</a>		1467	542		102	a kinase anchor protein 9 (glower) mRNA.
<a href="#">gloybor</a>	<a href="#">gloybor.aSep08</a>		27497	437	1	114	putative protein of eukaryotic origin (gloybor) alternative variant aSep08, mRNA.
<a href="#">gloybor</a>	<a href="#">gloybor.bSep08</a>		35246	1247	2	114	putative protein of eukaryotic origin (13.2 kD) (gloybor) alternative variant bSep08, mRNA.
<a href="#">gloychy</a>	<a href="#">gloychy.aSep08</a>		597	521		39	putative protein (4.5 kD) (gloychy) mRNA.
<a href="#">gloydoy</a>	<a href="#">gloydoy.aSep08</a>		4772	298		57	putative protein (gloydoy) mRNA.
<a href="#">gloyfee</a>	<a href="#">gloyfee.aSep08</a>		6950	399		62	uncharacterized protein like (gloyfee) mRNA.
<a href="#">gloyflu</a>	<a href="#">gloyflu.aSep08</a>		1962	474		157	glutamyl-tRNA synthetase 2 (gloyflu) mRNA.
<a href="#">gloyfly</a>	<a href="#">gloyfly.aSep08</a>		10342	348		115	ribosomal S6 kinase (gloyfly) mRNA.
<a href="#">gloygar</a>	<a href="#">gloygar.aSep08</a>		15154	303		63	putative protein (gloygar) mRNA.
<a href="#">gloyja</a>	<a href="#">gloyja.aSep08</a>		759	407		36	CRA a like (gloyja) mRNA.
<a href="#">gloyjey</a>	<a href="#">gloyjey.bSep08</a>		5478	810	4	51	putative protein (5.6 kD) (gloyjey) alternative variant bSep08, complete mRNA.
<a href="#">gloyjey</a>	<a href="#">gloyjey.cSep08</a>		5674	1026	3	50	putative protein (5.7 kD) (gloyjey) alternative variant cSep08, complete mRNA.
<a href="#">gloylo</a>	<a href="#">gloylo.aSep08</a>		2808	1069		356	putative protein of vertebrate origin (gloylo) mRNA.
<a href="#">gloymee</a>	<a href="#">gloymee.aSep08</a>		1901	1762		233	putative protein of vertebrate origin (26.1 kD) (gloymee) mRNA.
<a href="#">gloypor</a>	<a href="#">gloypor.aSep08</a>		7259	1166		125	F-box leucine-rich repeat protein 2 (gloypor) mRNA.
<a href="#">gloyroy</a>	<a href="#">gloyroy.aSep08</a>		66869	836		123	putative protein (14.5 kD) (gloyroy) mRNA.
<a href="#">gloyshee</a>	<a href="#">gloyshee.aSep08</a>		2487	471		78	putative protein (8.9 kD) (gloyshee) mRNA.
<a href="#">gloytu</a>	<a href="#">gloytu.aSep08</a>		9276	385		122	putative protein (gloytu) mRNA.
<a href="#">gloyvo</a>	<a href="#">gloyvo.aSep08</a>		651	371		109	putative protein of mammalian origin (gloyvo) mRNA.
<a href="#">gloywer</a>	<a href="#">gloywer.aSep08</a>		5022	477	3	18	putative protein (1.8 kD) (gloywer) alternative variant aSep08, mRNA.

<a href="#">Glx2</a>	<a href="#">Glx2.aSep08</a>	<a href="#">114022</a>	8704	1193	4	218	glutaredoxin 2 (thioltransferase) (23.5 kD) (Glx2) alternative variant aSep08, mRNA.
<a href="#">Glx2</a>	<a href="#">Glx2.bSep08</a>	<a href="#">114022</a>	13010	823	6	124	glutaredoxin 2 (thioltransferase) (14.0 kD) (Glx2) alternative variant bSep08, mRNA.
<a href="#">Glx2</a>	<a href="#">Glx2.cSep08</a>	<a href="#">114022</a>	9147	599	5	124	glutaredoxin 2 (thioltransferase) (14.0 kD) (Glx2) alternative variant cSep08, mRNA.
<a href="#">Glx2</a>	<a href="#">Glx2.dSep08</a>	<a href="#">114022</a>	10757	1984	5	124	glutaredoxin 2 (thioltransferase) (14.0 kD) (Glx2) alternative variant dSep08, mRNA.
<a href="#">Glx2</a>	<a href="#">Glx2.eSep08</a>	<a href="#">114022</a>	6771	331	3	109	glutaredoxin 2 (thioltransferase) (Glx2) alternative variant eSep08, mRNA.
<a href="#">Glx3</a>	<a href="#">Glx3.cSep08</a>	<a href="#">58815</a>	15882	679	3	43	glutaredoxin 3 (Glx3) alternative variant cSep08, mRNA.
<a href="#">Glx3</a>	<a href="#">Glx3.dSep08</a>	<a href="#">58815</a>	3017	456	3	24	glutaredoxin 3 (2.9 kD) (Glx3) alternative variant dSep08, mRNA.
<a href="#">Gls</a>	<a href="#">Gls.cSep08</a>	<a href="#">24398</a>	6362	392	5	130	glutaminase and similar to High mobility group protein 2 (HMG-2) (Gls) alternative variant cSep08, mRNA.
<a href="#">Gls</a>	<a href="#">Gls.cSep08</a>	<a href="#">685445</a>	6362	392	5	130	glutaminase and similar to High mobility group protein 2 (HMG-2) (Gls) alternative variant cSep08, mRNA.
<a href="#">Gls</a>	<a href="#">Gls.dSep08</a>	<a href="#">24398</a>	9178	390	3	45	glutaminase and similar to High mobility group protein 2 (HMG-2) (Gls) alternative variant dSep08, mRNA.
<a href="#">Gls</a>	<a href="#">Gls.dSep08</a>	<a href="#">685445</a>	9178	390	3	45	glutaminase and similar to High mobility group protein 2 (HMG-2) (Gls) alternative variant dSep08, mRNA.
<a href="#">Gls2</a>	<a href="#">Gls2.bSep08</a>	<a href="#">192268</a>	8014	992	6	259	glutaminase 2 (liver, mitochondrial) (Gls2) alternative variant bSep08, mRNA.
<a href="#">Gls2</a>	<a href="#">Gls2.cSep08</a>	<a href="#">192268</a>	3660	757	6	145	glutaminase 2 (liver, mitochondrial) (Gls2) alternative variant cSep08, mRNA.
<a href="#">Gls2</a>	<a href="#">Gls2.dSep08</a>	<a href="#">192268</a>	2407	289	3	96	glutaminase 2 (liver, mitochondrial) (Gls2) alternative variant dSep08, mRNA.
<a href="#">Glt1d1</a>	<a href="#">Glt1d1.aSep08</a>	<a href="#">304445</a>	54887	752		161	putative protein of ancient origin (Glt1d1) mRNA.
<a href="#">Glt6d1_predicted</a>	<a href="#">Glt6d1_predicted.aSep08</a>	<a href="#">296577</a>	8432	1083	3	297	glycosyl transferase, family 6 (Glt6d1_predicted) alternative variant aSep08, mRNA.
<a href="#">Glt8d1</a>	<a href="#">Glt8d1.bSep08</a>	<a href="#">306253</a>	13853	751	7	250	glycosyl transferase, family 8 (Glt8d1) alternative variant bSep08, mRNA.
<a href="#">Glt8d1</a>	<a href="#">Glt8d1.cSep08</a>	<a href="#">306253</a>	3497	475	4	158	glycosyl transferase, family 8 (Glt8d1) alternative variant cSep08, mRNA.
<a href="#">Glt8d1</a>	<a href="#">Glt8d1.eSep08</a>	<a href="#">306253</a>	1077	830	2	70	putative protein (8.2 kD) (Glt8d1) alternative variant eSep08, mRNA.
<a href="#">Glt8d1</a>	<a href="#">Glt8d1.gSep08</a>	<a href="#">306253</a>	819	533	2	25	putative protein (Glt8d1) alternative variant gSep08, mRNA.
<a href="#">Glt8d2</a>	<a href="#">Glt8d2.aSep08</a>	<a href="#">366859</a>	9004	1167		277	glycosyl transferase, family 8 (Glt8d2) mRNA.
<a href="#">Glt8d3</a>	<a href="#">Glt8d3.aSep08</a>	<a href="#">300173</a>	16240	1592	4	359	glycosyl transferase, family 8 (42.4 kD) (Glt8d3) alternative variant aSep08, mRNA.
<a href="#">Glt8d3</a>	<a href="#">Glt8d3.bSep08</a>	<a href="#">300173</a>	25274	1022	3	340	putative protein of ancient origin (Glt8d3) alternative variant bSep08, mRNA.
<a href="#">Glt8d3</a>	<a href="#">Glt8d3.cSep08</a>	<a href="#">300173</a>	23743	1039	2	248	putative protein of eukaryotic origin (Glt8d3) alternative variant cSep08, mRNA.
<a href="#">Glt8d4</a>	<a href="#">Glt8d4.aSep08</a>	<a href="#">688618</a>	15262	604		126	putative protein of metazoan origin (15.0 kD) (Glt8d4) mRNA.

<a href="#">Glt25d2</a>	<a href="#">Glt25d2.aSep08</a>	<a href="#">289081</a>	69192	681		226	putative protein of ancient origin (Glt25d2) mRNA.
<a href="#">Gltp</a>	<a href="#">Gltp.bSep08</a>	<a href="#">288707</a>	2334	776	2	86	glycolipid transfer protein (Gltp) alternative variant bSep08, mRNA.
<a href="#">Gltpd1</a>	<a href="#">Gltpd1.bSep08</a>	<a href="#">313771</a>	2934	911	2	153	putative protein of eukaryotic origin (Gltpd1) alternative variant bSep08, mRNA.
<a href="#">Gltpd1</a>	<a href="#">Gltpd1.cSep08</a>	<a href="#">313771</a>	2983	735	1	137	putative protein of eukaryotic origin (Gltpd1) alternative variant cSep08, mRNA.
<a href="#">Gltpd2</a>	<a href="#">Gltpd2.bSep08</a>	<a href="#">497943</a>	2567	1078	1	197	putative protein of mammalian origin (21.6 kD) (Gltpd2) alternative variant bSep08, complete mRNA.
<a href="#">Gltscr1</a>	<a href="#">Gltscr1.aSep08</a>	<a href="#">292622</a>	3943	2032		414	glioma tumor suppressor candidate region gene 1 (Gltscr1) mRNA.
<a href="#">Gltscr2</a>	<a href="#">Gltscr2.bSep08</a>	<a href="#">292624</a>	1116	506	1	63	glioma tumor suppressor candidate region gene 2 (Gltscr2) alternative variant bSep08, mRNA.
<a href="#">glubor</a>	<a href="#">glubor.aSep08</a>		15136	2092		242	family with sequence similarity 13 member C1 (glubor) mRNA.
<a href="#">gluchy</a>	<a href="#">gluchy.aSep08</a>		8236	784		232	CRA c (gluchy) mRNA.
<a href="#">gludoy</a>	<a href="#">gludoy.aSep08</a>		2501	340		77	putative protein (gludoy) mRNA.
<a href="#">gluflu</a>	<a href="#">gluflu.aSep08</a>		11229	305		59	putative protein (gluflu) mRNA.
<a href="#">gluflly</a>	<a href="#">gluflly.aSep08</a>		5503	436		32	putative protein (gluflly) mRNA.
<a href="#">glugar</a>	<a href="#">glugar.aSep08</a>		1659	301		55	putative protein (glugar) mRNA.
<a href="#">gluja</a>	<a href="#">gluja.aSep08</a>		813	381		126	CRA b like (gluja) mRNA.
<a href="#">glujey</a>	<a href="#">glujey.aSep08</a>		6205	241		57	putative protein (glujey) mRNA.
<a href="#">glukee</a>	<a href="#">glukee.aSep08</a>		798	393		71	putative mitochondrial protein (8.1 kD) (glukee) mRNA.
<a href="#">Glul</a>	<a href="#">Glul.bSep08</a>	<a href="#">24957</a>	27174	2458	2	314	glutamate-ammonia ligase (glutamine synthetase) (35.1 kD) (Glul) alternative variant bSep08, mRNA.
<a href="#">Glul</a>	<a href="#">Glul.cSep08</a>	<a href="#">24957</a>	25104	740	3	114	glutamate-ammonia ligase (glutamine synthetase) (Glul) alternative variant cSep08, mRNA.
<a href="#">glulo</a>	<a href="#">glulo.aSep08</a>		2402	504		36	putative protein (glulo) mRNA.
<a href="#">glumee</a>	<a href="#">glumee.aSep08</a>		3828	990	2	301	spinster homolog 2 (glumee) alternative variant aSep08, mRNA.
<a href="#">glumee</a>	<a href="#">glumee.bSep08</a>		2564	2129	2	197	CRA a (21.6 kD) (glumee) alternative variant bSep08, mRNA.
<a href="#">glumee</a>	<a href="#">glumee.cSep08</a>		3734	719	3	148	spinster homolog 2 (glumee) alternative variant cSep08, mRNA.
<a href="#">glupor</a>	<a href="#">glupor.aSep08</a>		7004	1305		133	doublecortin-like kinase 3 (glupor) mRNA.
<a href="#">gluroy</a>	<a href="#">gluroy.aSep08</a>		10415	1026		99	putative protein (gluroy) mRNA.
<a href="#">glusa</a>	<a href="#">glusa.aSep08</a>		2091	452	2	55	putative protein (6.3 kD) (glusa) mRNA.
<a href="#">glushee</a>	<a href="#">glushee.aSep08</a>		1190	845	1	131	polyprotein (glushee) alternative variant aSep08, mRNA.
<a href="#">glushee</a>	<a href="#">glushee.bSep08</a>		3797	822	1	62	putative protein (7.0 kD) (glushee) alternative variant bSep08, mRNA.
<a href="#">glutu</a>	<a href="#">glutu.aSep08</a>		3236	1286		79	putative protein (8.7 kD) (glutu) mRNA.
<a href="#">gluvo</a>	<a href="#">gluvo.aSep08</a>		1220	273		90	midasin (gluvo) mRNA.
<a href="#">gluwer</a>	<a href="#">gluwer.aSep08</a>		8556	773		257	a kinase anchor protein 9 (gluwer) mRNA.
<a href="#">Glyat</a>	<a href="#">Glyat.aSep08</a>	<a href="#">293779</a>	20912	1877	4	296	glycine-N-acyltransferase (33.9 kD) (Glyat) alternative variant aSep08, complete mRNA.

<a href="#">Glyat</a>	<a href="#">Glyat.cSep08</a>	<a href="#">293779</a>	18756	745	4	118	glycine-N-acyltransferase (Glyat) alternative variant cSep08, mRNA.
<a href="#">Glyat</a>	<a href="#">Glyat.dSep08</a>	<a href="#">293779</a>	13495	476	1	102	glycine-N-acyltransferase (Glyat) alternative variant dSep08, mRNA.
<a href="#">glybor</a>	<a href="#">glybor.aSep08</a>		616	286		25	putative protein (glybor) mRNA.
<a href="#">glychy</a>	<a href="#">glychy.aSep08</a>		596	246		36	putative protein (glychy) mRNA.
<a href="#">Glyco_hydro_1.0</a>	<a href="#">Glyco_hydro_1.0.aSep08</a>		5119	594		197	lactase (Glyco_hydro_1.0) mRNA.
<a href="#">Glyco_hydro_20b.0</a>	<a href="#">Glyco_hydro_20b.0.aSep08</a>		9201	586		166	hexosaminidase B (Glyco_hydro_20b.0) mRNA.
<a href="#">Glyco_hydro_31.0</a>	<a href="#">Glyco_hydro_31.0.aSep08</a>		24320	1513	13	503	glucosidase alpha (Glyco_hydro_31.0) alternative variant aSep08, mRNA.
<a href="#">Glyco_hydro_38.0</a>	<a href="#">Glyco_hydro_38.0.aSep08</a>		25307	3252	18	1083	mannosidase 2 alpha (Glyco_hydro_38.0) alternative variant aSep08, mRNA.
<a href="#">Glyco_hydro_38.0</a>	<a href="#">Glyco_hydro_38.0.bSep08</a>		6256	778	5	258	mannosidase 2 alpha B2 (Glyco_hydro_38.0) alternative variant bSep08, mRNA.
<a href="#">Glyco_hydro_38.0</a>	<a href="#">Glyco_hydro_38.0.cSep08</a>		1763	586	1	110	CRA f (Glyco_hydro_38.0) alternative variant cSep08, mRNA.
<a href="#">Glyco_hydro_38.0</a>	<a href="#">Glyco_hydro_38.0.dSep08</a>		5506	551	4	86	mannosidase 2 alpha (Glyco_hydro_38.0) alternative variant dSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.aSep08</a>		3026	1473	10	490	alpha-L-iduronidase (Glyco_hydro_39.0) alternative variant aSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.bSep08</a>		11619	840	7	247	iduronidase alpha-L- CRA b (Glyco_hydro_39.0) alternative variant bSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.cSep08</a>		2706	1911	4	160	alpha-L-iduronidase (18.1 kD) (Glyco_hydro_39.0) alternative variant cSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.dSep08</a>		1978	798	4	139	iduronidase alpha-L- CRA e (Glyco_hydro_39.0) alternative variant dSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.eSep08</a>		1837	734	5	131	alpha-L-iduronidase (Glyco_hydro_39.0) alternative variant eSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.fSep08</a>		1138	949	3	109	iduronidase alpha-L- CRA b (Glyco_hydro_39.0) alternative variant fSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.gSep08</a>		1541	750	4	87	alpha-L-iduronidase (9.4 kD) (Glyco_hydro_39.0) alternative variant gSep08, mRNA.
<a href="#">Glyco_hydro_39.0</a>	<a href="#">Glyco_hydro_39.0.iSep08</a>		1520	1095	2	65	putative protein (Glyco_hydro_39.0) alternative variant iSep08, mRNA.
<a href="#">Glyco_transf_6.0</a>	<a href="#">Glyco_transf_6.0.aSep08</a>		20636	1517	2	340	histo-blood group ABO transferase like (Glyco_transf_6.0) alternative variant aSep08, mRNA.
<a href="#">Glyco_transf_6.0</a>	<a href="#">Glyco_transf_6.0.bSep08</a>		5358	500	1	166	ABO blood group like (Glyco_transf_6.0) alternative variant bSep08, mRNA.
<a href="#">Glyco_transf_25.0</a>	<a href="#">Glyco_transf_25.0.aSep08</a>		4566	406		135	glycosyl transferase, family 25 (Glyco_transf_25.0) mRNA.
<a href="#">glydoy</a>	<a href="#">glydoy.aSep08</a>		5860	672		95	putative protein (glydoy) mRNA.
<a href="#">glyflu</a>	<a href="#">glyflu.aSep08</a>		5559	885		62	putative protein (7.1 kD) (glyflu) mRNA.
<a href="#">glyfly</a>	<a href="#">glyfly.aSep08</a>		3617	397		86	putative protein (glyfly) mRNA.
<a href="#">glygar</a>	<a href="#">glygar.aSep08</a>		9077	623		115	serine threonine-protein kinase nek5 like (glygar) mRNA.

<a href="#">glyja</a>	<a href="#">glyja.aSep08</a>		2595	784		261	CRA b like (glyja) mRNA.
<a href="#">glyjey</a>	<a href="#">glyjey.aSep08</a>		4346	679	2	39	putative protein (4.6 kD) (glyjey) alternative variant aSep08, mRNA.
<a href="#">glykee</a>	<a href="#">glykee.bSep08</a>		919	665	2	54	putative protein (glykee) alternative variant bSep08, mRNA.
<a href="#">glylo</a>	<a href="#">glylo.aSep08</a>		3022	745		89	putative protein (9.6 kD) (glylo) mRNA.
<a href="#">glymee</a>	<a href="#">glymee.aSep08</a>		970	711		167	putative mitochondrial protein (18.3 kD) (glymee) mRNA.
<a href="#">glypor</a>	<a href="#">glypor.aSep08</a>		1074	319		28	putative protein (3.1 kD) (glypor) mRNA.
<a href="#">glyroy</a>	<a href="#">glyroy.aSep08</a>		4724	244	1	73	putative protein (glyroy) alternative variant aSep08, mRNA.
<a href="#">glyroy</a>	<a href="#">glyroy.bSep08</a>		5353	749	1	119	putative protein (13.1 kD) (glyroy) alternative variant bSep08, mRNA.
<a href="#">glysa</a>	<a href="#">glysa.aSep08</a>		1805	951		61	putative protein (7.0 kD) (glysa) mRNA.
<a href="#">glyshee</a>	<a href="#">glyshee.aSep08</a>		9546	270		70	CRA b like (7.4 kD) (glyshee) alternative variant aSep08, mRNA.
<a href="#">glyshee</a>	<a href="#">glyshee.bSep08</a>		9528	399	1	61	CRA a like (glyshee) alternative variant bSep08, mRNA.
<a href="#">glytu</a>	<a href="#">glytu.aSep08</a>		1387	279		26	putative protein (glytu) mRNA.
<a href="#">glyvo</a>	<a href="#">glyvo.aSep08</a>		2187	330		110	midasin (glyvo) mRNA.
<a href="#">glywer</a>	<a href="#">glywer.aSep08</a>		8230	719		83	putative protein (glywer) mRNA.
<a href="#">Gmcl1</a>	<a href="#">Gmcl1.cSep08</a>	<a href="#">312516</a>	14316	832	5	200	germ cell-less homolog 1 (Drosophila) (22.3 kD) (Gmcl1) alternative variant cSep08, mRNA.
<a href="#">Gmcl1</a>	<a href="#">Gmcl1.eSep08</a>	<a href="#">312516</a>	6988	646	3	71	germ cell-less homolog 1 (Drosophila) (Gmcl1) alternative variant eSep08, mRNA.
<a href="#">Gmnds</a>	<a href="#">Gmnds.bSep08</a>	<a href="#">291095</a>	279052	825	5	114	GDP-mannose 4, 6-dehydratase (13.3 kD) (Gmnds) alternative variant bSep08, mRNA.
<a href="#">Gmeb2</a>	<a href="#">Gmeb2.bSep08</a>	<a href="#">83635</a>	8480	1781	1	403	glucocorticoid modulatory element binding protein 2 (Gmeb2) alternative variant bSep08, mRNA.
<a href="#">Gmfb</a>	<a href="#">Gmfb.bSep08</a>	<a href="#">81661</a>	12991	4047	7	105	glia maturation factor, beta (12.4 kD) (Gmfb) alternative variant bSep08, complete mRNA.
<a href="#">Gmfb</a>	<a href="#">Gmfb.cSep08</a>	<a href="#">81661</a>	9565	592	7	102	glia maturation factor, beta (11.4 kD) (Gmfb) alternative variant cSep08, complete mRNA.
<a href="#">Gmfb</a>	<a href="#">Gmfb.dSep08</a>	<a href="#">81661</a>	9505	723	8	69	glia maturation factor, beta (7.6 kD) (Gmfb) alternative variant dSep08, mRNA.
<a href="#">Gmfg</a>	<a href="#">Gmfg.bSep08</a>	<a href="#">113940</a>	4817	479	1	101	glia maturation factor, gamma (11.9 kD) (Gmfg) alternative variant bSep08, mRNA.
<a href="#">Gmfg</a>	<a href="#">Gmfg.cSep08</a>	<a href="#">113940</a>	10373	586	2	101	glia maturation factor, gamma (11.9 kD) (Gmfg) alternative variant cSep08, mRNA.
<a href="#">Gmfg</a>	<a href="#">Gmfg.dSep08</a>	<a href="#">113940</a>	5439	468	2	101	glia maturation factor, gamma (11.9 kD) (Gmfg) alternative variant dSep08, mRNA.
<a href="#">Gmip</a>	<a href="#">Gmip.bSep08</a>	<a href="#">306357</a>	1521	1406	2	154	gem-interacting protein (11.3 kD) (Gmip) alternative variant bSep08, mRNA.
<a href="#">Gmnn</a>	<a href="#">Gmnn.aSep08</a>	<a href="#">291137</a>	8212	918	6	206	geminin (23.1 kD) (Gmnn) alternative variant aSep08, mRNA.
<a href="#">Gmnn</a>	<a href="#">Gmnn.cSep08</a>	<a href="#">291137</a>	557	362	1	91	geminin (Gmnn) alternative variant cSep08, mRNA.
<a href="#">Gmppa</a>	<a href="#">Gmppa.bSep08</a>	<a href="#">501167</a>	5899	815	8	246	GDP-mannose pyrophosphorylase A (Gmppa) alternative variant bSep08, mRNA.

<a href="#">Gmppa</a>	<a href="#">Gmppa.cSep08</a>	<a href="#">501167</a>	3402	692	6	163	GDP-mannose pyrophosphorylase A (Gmppa) alternative variant cSep08, mRNA.
<a href="#">Gmppa</a>	<a href="#">Gmppa.dSep08</a>	<a href="#">501167</a>	2106	383	3	127	GDP-mannose pyrophosphorylase A (Gmppa) alternative variant dSep08, mRNA.
<a href="#">Gmppa</a>	<a href="#">Gmppa.eSep08</a>	<a href="#">501167</a>	1609	1302	3	88	GDP-mannose pyrophosphorylase A (9.7 kD) (Gmppa) alternative variant eSep08, mRNA.
<a href="#">Gmppa</a>	<a href="#">Gmppa.fSep08</a>	<a href="#">501167</a>	7460	610	3	17	GDP-mannose pyrophosphorylase A (1.8 kD) (Gmppa) alternative variant fSep08, complete mRNA.
<a href="#">Gmppb</a>	<a href="#">Gmppb.aSep08</a>	<a href="#">363145</a>	2608	1763		360	GDP-mannose pyrophosphorylase B (39.9 kD) (Gmppb) mRNA.
<a href="#">Gmpr</a>	<a href="#">Gmpr.bSep08</a>	<a href="#">117533</a>	10436	540	3	112	guanosine monophosphate reductase (Gmpr) alternative variant bSep08, mRNA.
<a href="#">Gmpr</a>	<a href="#">Gmpr.cSep08</a>	<a href="#">117533</a>	7327	1138	3	106	guanosine monophosphate reductase (11.5 kD) (Gmpr) alternative variant cSep08, mRNA.
<a href="#">Gmpr2</a>	<a href="#">Gmpr2.aSep08</a>	<a href="#">192357</a>	6228	1069	7	330	guanosine monophosphate reductase 2 (Gmpr2) alternative variant aSep08, mRNA.
<a href="#">Gmpr2</a>	<a href="#">Gmpr2.cSep08</a>	<a href="#">192357</a>	5653	736	7	157	guanosine monophosphate reductase 2 (Gmpr2) alternative variant cSep08, mRNA.
<a href="#">Gmpr2</a>	<a href="#">Gmpr2.dSep08</a>	<a href="#">192357</a>	5593	754	8	152	guanosine monophosphate reductase 2 (Gmpr2) alternative variant dSep08, mRNA.
<a href="#">Gmpr2</a>	<a href="#">Gmpr2.eSep08</a>	<a href="#">192357</a>	1925	1007	3	130	guanosine monophosphate reductase 2 (Gmpr2) alternative variant eSep08, mRNA.
<a href="#">Gmps</a>	<a href="#">Gmps.bSep08</a>	<a href="#">295088</a>	10101	663	5	214	guanine monphosphate synthetase (Gmps) alternative variant bSep08, mRNA.
<a href="#">Gmps</a>	<a href="#">Gmps.cSep08</a>	<a href="#">295088</a>	2589	476	3	151	guanine monphosphate synthetase (Gmps) alternative variant cSep08, mRNA.
<a href="#">Gmps</a>	<a href="#">Gmps.dSep08</a>	<a href="#">295088</a>	8728	709	4	112	guanine monphosphate synthetase (Gmps) alternative variant dSep08, mRNA.
<a href="#">Gmps</a>	<a href="#">Gmps.eSep08</a>	<a href="#">295088</a>	9537	457	4	85	guanine monphosphate synthetase (Gmps) alternative variant eSep08, mRNA.
<a href="#">Gna11</a>	<a href="#">Gna11.bSep08</a>	<a href="#">81662</a>	711	525	2	102	guanine nucleotide binding protein, alpha 11 (Gna11) alternative variant bSep08, mRNA.
<a href="#">Gna12</a>	<a href="#">Gna12.cSep08</a>	<a href="#">81663</a>	44164	659	1	213	guanine nucleotide binding protein, alpha 12 (Gna12) alternative variant cSep08, mRNA.
<a href="#">Gnai1</a>	<a href="#">Gnai1.aSep08</a>	<a href="#">25686</a>	30631	2749	6	287	guanine nucleotide binding protein (G protein), alpha inhibiting 1 (Gnai1) alternative variant aSep08, mRNA.
<a href="#">Gnai2</a>	<a href="#">Gnai2.bSep08</a>	<a href="#">81664</a>	1869	551	4	127	guanine nucleotide binding protein (G protein), alpha inhibiting 2 (Gnai2) alternative variant bSep08, mRNA.
<a href="#">Gnai2</a>	<a href="#">Gnai2.cSep08</a>	<a href="#">81664</a>	1275	390	3	91	guanine nucleotide binding protein (G protein), alpha inhibiting 2 (Gnai2) alternative variant cSep08, mRNA.
<a href="#">Gnal</a>	<a href="#">Gnal.aSep08</a>	<a href="#">24611</a>	47836	716		182	guanine nucleotide binding protein, alpha stimulating, olfactory type (Gnal) mRNA.
<a href="#">Gnao1</a>	<a href="#">Gnao1.cSep08</a>	<a href="#">50664</a>	127928	449	1	78	guanine nucleotide binding protein, alpha O (Gnao1) alternative variant cSep08, mRNA.
<a href="#">Gnaq</a>	<a href="#">Gnaq.aSep08</a>	<a href="#">81666</a>	244572	1800	4	558	guanine nucleotide binding protein, alpha q polypeptide (Gnaq) alternative variant aSep08, mRNA.

<a href="#">Gnaq</a>	<a href="#">Gnaq.cSep08</a>	<a href="#">81666</a>	1387	422	1	61	guanine nucleotide binding protein, alpha q polypeptide (6.6 kD) (Gnaq) alternative variant cSep08, mRNA.
<a href="#">Gnas</a>	<a href="#">Gnas.aSep08</a>	<a href="#">24896</a>	1154	973	2	264	neuroendocrine secretory protein 55 (29.8 kD) (Gnas) alternative variant aSep08, complete mRNA.
<a href="#">Gnas</a>	<a href="#">Gnas.bSep08</a>	<a href="#">24896</a>	656	564	1	105	putative protein (Gnas) alternative variant bSep08, mRNA.
<a href="#">Gnas</a>	<a href="#">Gnas.cSep08</a>	<a href="#">24896</a>	488	393	1	71	putative protein (Gnas) alternative variant cSep08, mRNA.
<a href="#">Gnat2</a>	<a href="#">Gnat2.aSep08</a>	<a href="#">365901</a>	2300	349		116	guanine nucleotide binding protein, alpha transducing 2 (Gnat2) mRNA.
<a href="#">Gnb1.1</a>	<a href="#">Gnb1.1.aSep08</a>	<a href="#">24400</a>	45922	1797	3	340	guanine nucleotide binding protein (G protein), beta 1 (37.4 kD) (Gnb1.1) alternative variant aSep08, mRNA.
<a href="#">Gnb1.1</a>	<a href="#">Gnb1.1.bSep08</a>	<a href="#">24400</a>	68956	3097	3	340	guanine nucleotide binding protein (G protein), beta 1 (37.4 kD) (Gnb1.1) alternative variant bSep08, mRNA.
<a href="#">Gnb1.1</a>	<a href="#">Gnb1.1.cSep08</a>	<a href="#">24400</a>	61090	697	2	148	guanine nucleotide binding protein (G protein), beta 1 (Gnb1.1) alternative variant cSep08, mRNA.
<a href="#">Gnb1.1</a>	<a href="#">Gnb1.1.dSep08</a>	<a href="#">24400</a>	53253	695	1	141	guanine nucleotide binding protein (G protein), beta 1 (Gnb1.1) alternative variant dSep08, mRNA.
<a href="#">Gnb1l</a>	<a href="#">Gnb1l.aSep08</a>	<a href="#">680266</a>	73816	1331	5	223	guanine nucleotide binding protein (G protein), beta polypeptide 1-like (24.4 kD) (Gnb1l) alternative variant aSep08, mRNA.
<a href="#">Gnb1l</a>	<a href="#">Gnb1l.bSep08</a>	<a href="#">680266</a>	4096	2197	1	75	guanine nucleotide binding protein (G protein), beta polypeptide 1-like (8.2 kD) (Gnb1l) alternative variant bSep08, mRNA.
<a href="#">Gnb2</a>	<a href="#">Gnb2.bSep08</a>	<a href="#">81667</a>	1824	869	6	289	guanine nucleotide binding protein (G protein), beta 2 (Gnb2) alternative variant bSep08, mRNA.
<a href="#">Gnb2</a>	<a href="#">Gnb2.cSep08</a>	<a href="#">81667</a>	2016	724	7	241	guanine nucleotide binding protein (G protein), beta 2 (Gnb2) alternative variant cSep08, mRNA.
<a href="#">Gnb2</a>	<a href="#">Gnb2.dSep08</a>	<a href="#">81667</a>	980	817	3	179	guanine nucleotide binding protein (G protein), beta 2 (Gnb2) alternative variant dSep08, mRNA.
<a href="#">Gnb2</a>	<a href="#">Gnb2.eSep08</a>	<a href="#">81667</a>	3681	763	5	174	guanine nucleotide binding protein (G protein), beta 2 (Gnb2) alternative variant eSep08, mRNA.
<a href="#">Gnb2</a>	<a href="#">Gnb2.fSep08</a>	<a href="#">81667</a>	1316	837	4	122	guanine nucleotide binding protein (G protein), beta 2 (13.6 kD) (Gnb2) alternative variant fSep08, mRNA.
<a href="#">Gnb2l1</a>	<a href="#">Gnb2l1.bSep08</a>	<a href="#">83427</a>	5543	1747	7	238	guanine nucleotide binding protein (G protein), beta polypeptide 2 like 1 (26.2 kD) (Gnb2l1) alternative variant bSep08, complete mRNA.
<a href="#">Gnb2l1</a>	<a href="#">Gnb2l1.dSep08</a>	<a href="#">83427</a>	1219	742	2	56	guanine nucleotide binding protein (G protein), beta polypeptide 2 like 1 (Gnb2l1) alternative variant dSep08, mRNA.
<a href="#">Gnb3</a>	<a href="#">Gnb3.bSep08</a>	<a href="#">60449</a>	2868	875	7	148	guanine nucleotide binding protein (G protein), beta 3 (Gnb3) alternative variant bSep08, mRNA.
<a href="#">Gnb3</a>	<a href="#">Gnb3.cSep08</a>	<a href="#">60449</a>	2303	420	5	96	guanine nucleotide binding protein (G protein), beta 3 (Gnb3) alternative variant cSep08, mRNA.
<a href="#">Gnb3</a>	<a href="#">Gnb3.dSep08</a>	<a href="#">60449</a>	2444	762	4	68	guanine nucleotide binding protein (G protein), beta 3 (7.6 kD) (Gnb3) alternative variant dSep08, mRNA.
<a href="#">Gnb3</a>	<a href="#">Gnb3.eSep08</a>	<a href="#">60449</a>	2319	713	3	47	guanine nucleotide binding protein (G protein), beta 3 (5.5 kD) (Gnb3) alternative variant eSep08, mRNA.



<a href="#">Gnb4</a>	<a href="#">Gnb4.aSep08</a>	<a href="#">294962</a>	9075	910	4	249	guanine nucleotide binding protein (G protein), beta 4 (Gnb4) alternative variant aSep08, mRNA.
<a href="#">Gnb4</a>	<a href="#">Gnb4.bSep08</a>	<a href="#">294962</a>	28493	715	7	73	guanine nucleotide binding protein (G protein), beta 4 (8.5 kD) (Gnb4) alternative variant bSep08, mRNA.
<a href="#">Gnb5</a>	<a href="#">Gnb5.aSep08</a>	<a href="#">83579</a>	41846	2067	10	298	guanine nucleotide binding protein (G protein), beta 5 (Gnb5) alternative variant aSep08, mRNA.
<a href="#">Gnb5</a>	<a href="#">Gnb5.cSep08</a>	<a href="#">83579</a>	5933	1277	4	103	guanine nucleotide binding protein (G protein), beta 5 (Gnb5) alternative variant cSep08, mRNA.
<a href="#">Gne</a>	<a href="#">Gne.bSep08</a>	<a href="#">114711</a>	16951	772	2	226	glucosamine (Gne) alternative variant bSep08, mRNA.
<a href="#">Gne</a>	<a href="#">Gne.cSep08</a>	<a href="#">114711</a>	19315	396	1	109	glucosamine (Gne) alternative variant cSep08, mRNA.
<a href="#">Gng3</a>	<a href="#">Gng3.aSep08</a>	<a href="#">114117</a>	1431	817		75	guanine nucleotide binding protein (G protein), gamma 3 (8.3 kD) (Gng3) mRNA.
<a href="#">Gng7</a>	<a href="#">Gng7.bSep08</a>	<a href="#">58979</a>	88013	398	4	54	putative protein (Gng7) alternative variant bSep08, mRNA.
<a href="#">Gng7</a>	<a href="#">Gng7.cSep08</a>	<a href="#">58979</a>	66083	473	4	48	putative protein (Gng7) alternative variant cSep08, mRNA.
<a href="#">Gng8</a>	<a href="#">Gng8.bSep08</a>	<a href="#">245986</a>	3345	577		43	guanine nucleotide binding protein (G protein), gamma 8 (Gng8) alternative variant bSep08, mRNA.
<a href="#">Gng12</a>	<a href="#">Gng12.aSep08</a>	<a href="#">114120</a>	135295	342	1	57	guanine nucleotide binding protein (G protein), gamma 12 (Gng12) alternative variant aSep08, mRNA.
<a href="#">Gng12</a>	<a href="#">Gng12.bSep08</a>	<a href="#">114120</a>	138129	395	1	45	guanine nucleotide binding protein (G protein), gamma 12 (Gng12) alternative variant bSep08, mRNA.
<a href="#">Gng12</a>	<a href="#">Gng12.cSep08</a>	<a href="#">114120</a>	140081	801	2	72	guanine nucleotide binding protein (G protein), gamma 12 (8.0 kD) (Gng12) alternative variant cSep08, complete mRNA.
<a href="#">Gng13</a>	<a href="#">Gng13.aSep08</a>	<a href="#">685451</a>	1905	354		92	guanine nucleotide binding protein (G protein), gamma 13 (Gng13) mRNA.
<a href="#">Gnl1</a>	<a href="#">Gnl1.bSep08</a>	<a href="#">309593</a>	2726	677	6	225	guanine nucleotide binding protein-like 1 (Gnl1) alternative variant bSep08, mRNA.
<a href="#">Gnl1</a>	<a href="#">Gnl1.cSep08</a>	<a href="#">309593</a>	1385	902	3	180	guanine nucleotide binding protein-like 1 (19.8 kD) (Gnl1) alternative variant cSep08, mRNA.
<a href="#">Gnl1</a>	<a href="#">Gnl1.dSep08</a>	<a href="#">309593</a>	1359	741	3	155	guanine nucleotide binding protein-like 1 (Gnl1) alternative variant dSep08, mRNA.
<a href="#">Gnl2</a>	<a href="#">Gnl2.bSep08</a>	<a href="#">362593</a>	1649	666	2	101	guanine nucleotide binding protein-like 2 (nucleolar) (11.9 kD) (Gnl2) alternative variant bSep08, mRNA.
<a href="#">Gnl2</a>	<a href="#">Gnl2.cSep08</a>	<a href="#">362593</a>	14778	605	5	65	guanine nucleotide binding protein-like 2 (nucleolar) (Gnl2) alternative variant cSep08, mRNA.
<a href="#">Gnl3</a>	<a href="#">Gnl3.bSep08</a>	<a href="#">290556</a>	1691	1377	4	353	guanine nucleotide binding protein-like 3 (Gnl3) alternative variant bSep08, mRNA.
<a href="#">Gnl3</a>	<a href="#">Gnl3.cSep08</a>	<a href="#">290556</a>	4386	1491	12	265	guanine nucleotide binding protein-like 3 (29.0 kD) (Gnl3) alternative variant cSep08, mRNA.
<a href="#">Gnl3</a>	<a href="#">Gnl3.dSep08</a>	<a href="#">290556</a>	5945	2850	10	115	guanine nucleotide binding protein-like 3 (13.9 kD) (Gnl3) alternative variant dSep08, complete mRNA.
<a href="#">Gnl3</a>	<a href="#">Gnl3.eSep08</a>	<a href="#">290556</a>	483	389	2	35	guanine nucleotide binding protein-like 3 (3.9 kD) (Gnl3) alternative variant eSep08, mRNA.
<a href="#">Gnpat</a>	<a href="#">Gnpat.cSep08</a>	<a href="#">84470</a>	1757	507	2	53	glyceronephosphate O-acyltransferase (6.0 kD) (Gnpat) alternative variant cSep08, mRNA.
<a href="#">Gnpda2</a>	<a href="#">Gnpda2.bSep08</a>	<a href="#">289608</a>	13434	645	5	214	glucosamine-6-phosphate deaminase 2 (Gnpda2) alternative variant bSep08, mRNA.

<a href="#">Gnpda2</a>	<a href="#">Gnpda2.cSep08</a>	<a href="#">289608</a>	15915	708	5	166	glucosamine-6-phosphate deaminase 2 (Gnpda2) alternative variant cSep08, mRNA.
<a href="#">Gnpda2</a>	<a href="#">Gnpda2.dSep08</a>	<a href="#">289608</a>	8018	553	4	152	glucosamine-6-phosphate deaminase 2 (Gnpda2) alternative variant dSep08, mRNA.
<a href="#">Gnpda2</a>	<a href="#">Gnpda2.fSep08</a>	<a href="#">289608</a>	15874	725	6	69	glucosamine-6-phosphate deaminase 2 (7.8 kD) (Gnpda2) alternative variant fSep08, mRNA.
<a href="#">Gnptab</a>	<a href="#">Gnptab.aSep08</a>	<a href="#">362865</a>	5462	1098	1	365	N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits (Gnptab) alternative variant aSep08, mRNA.
<a href="#">Gnptab</a>	<a href="#">Gnptab.bSep08</a>	<a href="#">362865</a>	6008	861	3	286	N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits (Gnptab) alternative variant bSep08, mRNA.
<a href="#">Gnptab</a>	<a href="#">Gnptab.cSep08</a>	<a href="#">362865</a>	13134	1478	6	221	N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits (Gnptab) alternative variant cSep08, mRNA.
<a href="#">Gnptg</a>	<a href="#">Gnptg.aSep08</a>	<a href="#">287134</a>	6852	3183	5	331	N-acetylglucosamine-1-phosphotransferase, gamma subunit (36.8 kD) (Gnptg) alternative variant aSep08, complete mRNA.
<a href="#">Gnptg</a>	<a href="#">Gnptg.cSep08</a>	<a href="#">287134</a>	4881	1296	4	281	N-acetylglucosamine-1-phosphotransferase, gamma subunit (31.1 kD) (Gnptg) alternative variant cSep08, complete mRNA.
<a href="#">Gnptg</a>	<a href="#">Gnptg.dSep08</a>	<a href="#">287134</a>	4632	910	4	224	N-acetylglucosamine-1-phosphotransferase, gamma subunit (Gnptg) alternative variant dSep08, mRNA.
<a href="#">Gnptg</a>	<a href="#">Gnptg.eSep08</a>	<a href="#">287134</a>	4137	717	2	180	N-acetylglucosamine-1-phosphotransferase, gamma subunit (Gnptg) alternative variant eSep08, mRNA.
<a href="#">Gns</a>	<a href="#">Gns.bSep08</a>	<a href="#">299825</a>	21702	3263	3	352	glucosamine (N-acetyl)-6-sulfatase (40.0 kD) (Gns) alternative variant bSep08, mRNA.
<a href="#">goby</a>	<a href="#">goby.aSep08</a>		1485	394		63	ftsj homolog 1 (goby) mRNA.
<a href="#">gochy</a>	<a href="#">gochy.aSep08</a>		36387	680		88	putative protein (9.3 kD) (gochy) mRNA.
<a href="#">gofer</a>	<a href="#">gofer.aSep08</a>		3952	1118			
<a href="#">goflo</a>	<a href="#">goflo.aSep08</a>		3287	420		33	putative protein (3.8 kD) (goflo) mRNA.
<a href="#">goflu</a>	<a href="#">goflu.aSep08</a>		3376	775		166	putative protein of mammalian origin (goflu) mRNA.
<a href="#">gokee</a>	<a href="#">gokee.aSep08</a>		2070	981		326	specifically androgen-regulated protein (gokee) mRNA.
<a href="#">Golga1</a>	<a href="#">Golga1.bSep08</a>	<a href="#">311919</a>	11039	601	5	87	golgi autoantigen, golgin subfamily a, 1 (Golga1) alternative variant bSep08, mRNA.
<a href="#">Golga1</a>	<a href="#">Golga1.cSep08</a>	<a href="#">311919</a>	822	382	2	73	golgi autoantigen, golgin subfamily a, 1 (Golga1) alternative variant cSep08, mRNA.
<a href="#">Golga2</a>	<a href="#">Golga2.aSep08</a>	<a href="#">64528</a>	3835	2570	8	419	golgi autoantigen, golgin subfamily a, 2 (Golga2) alternative variant aSep08, mRNA.
<a href="#">Golga2</a>	<a href="#">Golga2.bSep08</a>	<a href="#">64528</a>	7775	895	5	286	golgi autoantigen, golgin subfamily a, 2 (Golga2) alternative variant bSep08, mRNA.
<a href="#">Golga2</a>	<a href="#">Golga2.cSep08</a>	<a href="#">64528</a>	1778	806	5	268	golgi autoantigen, golgin subfamily a, 2 (Golga2) alternative variant cSep08, mRNA.
<a href="#">Golga3</a>	<a href="#">Golga3.bSep08</a>	<a href="#">312077</a>	30378	1784	10	569	golgi autoantigen, golgin subfamily a, 3 (Golga3) alternative variant bSep08, mRNA.
<a href="#">Golga3</a>	<a href="#">Golga3.cSep08</a>	<a href="#">312077</a>	2506	718	3	144	golgi autoantigen, golgin subfamily a, 3 (15.8 kD) (Golga3) alternative variant cSep08, mRNA.
<a href="#">Golga3</a>	<a href="#">Golga3.dSep08</a>	<a href="#">312077</a>	7736	425	4	141	golgi autoantigen, golgin subfamily a, 3 (Golga3) alternative variant dSep08, mRNA.

<a href="#">Golga3</a>	<a href="#">Golga3.eSep08</a>	<a href="#">312077</a>	1288	581	2	117	golgi autoantigen, golgin subfamily a, 3 (Golga3) alternative variant eSep08, mRNA.
<a href="#">Golga4</a>	<a href="#">Golga4.aSep08</a>	<a href="#">501069</a>	23087	3001	10	791	golgi autoantigen, golgin subfamily a, 4 (Golga4) alternative variant aSep08, mRNA.
<a href="#">Golga4</a>	<a href="#">Golga4.bSep08</a>	<a href="#">501069</a>	12139	549	5	88	golgi autoantigen, golgin subfamily a, 4 (Golga4) alternative variant bSep08, mRNA.
<a href="#">Golga5</a>	<a href="#">Golga5.bSep08</a>	<a href="#">299258</a>	2358	826	2	28	golgi autoantigen, golgin subfamily a, 5 (Golga5) alternative variant bSep08, mRNA.
<a href="#">Golga7</a>	<a href="#">Golga7.aSep08</a>	<a href="#">361171</a>	14712	1335	2	137	golgi autoantigen, golgin subfamily a, 7 (15.8 kD) (Golga7) alternative variant aSep08, mRNA.
<a href="#">Golgb1</a>	<a href="#">Golgb1.bSep08</a>	<a href="#">192243</a>	4850	595	5	148	golgi autoantigen, golgin subfamily b, macrogolgin 1 (Golgb1) alternative variant bSep08, mRNA.
<a href="#">Golgb1</a>	<a href="#">Golgb1.cSep08</a>	<a href="#">192243</a>	1738	1533	2	88	golgi autoantigen, golgin subfamily b, macrogolgin 1 (10.0 kD) (Golgb1) alternative variant cSep08, mRNA.
<a href="#">Golim4</a>	<a href="#">Golim4.aSep08</a>	<a href="#">310526</a>	20891	3035	9	392	golgi 4 (45.7 kD) (Golim4) alternative variant aSep08, mRNA.
<a href="#">Golim4</a>	<a href="#">Golim4.bSep08</a>	<a href="#">310526</a>	66634	1013	7	310	golgi 4 (Golim4) alternative variant bSep08, mRNA.
<a href="#">Golim4</a>	<a href="#">Golim4.cSep08</a>	<a href="#">310526</a>	91991	1595	4	249	golgi 4 (Golim4) alternative variant cSep08, mRNA.
<a href="#">Golim4</a>	<a href="#">Golim4.dSep08</a>	<a href="#">310526</a>	9770	738	5	245	golgi 4 (Golim4) alternative variant dSep08, mRNA.
<a href="#">GoLoco.0</a>	<a href="#">GoLoco.0.aSep08</a>		2028	734		84	LGN protein (GoLoco.0) mRNA.
<a href="#">goloy</a>	<a href="#">goloy.aSep08</a>		13396	938		58	limbic system-associated membrane protein (6.5 kD) (goloy) mRNA.
<a href="#">Golph2</a>	<a href="#">Golph2.aSep08</a>	<a href="#">306725</a>	17128	809		105	golgi phosphoprotein 2 and hypothetical protein LOC681199 (Golph2) mRNA.
<a href="#">Golph2</a>	<a href="#">Golph2.aSep08</a>	<a href="#">681199</a>	17128	809		105	golgi phosphoprotein 2 and hypothetical protein LOC681199 (Golph2) mRNA.
<a href="#">Golph3</a>	<a href="#">Golph3.bSep08</a>	<a href="#">78961</a>	20796	739	1	162	golgi phosphoprotein 3 (18.0 kD) (Golph3) alternative variant bSep08, mRNA.
<a href="#">Golsyn</a>	<a href="#">Golsyn.aSep08</a>	<a href="#">500865</a>	83984	1813	8	546	golgi-localized protein (Golsyn) alternative variant aSep08, mRNA.
<a href="#">Golsyn</a>	<a href="#">Golsyn.bSep08</a>	<a href="#">500865</a>	36042	410	2	92	golgi-localized protein (Golsyn) alternative variant bSep08, mRNA.
<a href="#">Golsyn</a>	<a href="#">Golsyn.cSep08</a>	<a href="#">500865</a>	27066	383	3	59	golgi-localized protein (Golsyn) alternative variant cSep08, mRNA.
<a href="#">Golt1b</a>	<a href="#">Golt1b.cSep08</a>	<a href="#">362460</a>	6571	460	3	99	golgi transport 1 homolog B (S. cerevisiae) (11.2 kD) (Golt1b) alternative variant cSep08, mRNA.
<a href="#">gomee</a>	<a href="#">gomee.aSep08</a>		15307	371		123	dna repair endonuclease (gomee) mRNA.
<a href="#">gomer</a>	<a href="#">gomer.aSep08</a>		876	780		64	putative protein (7.7 kD) (gomer) mRNA.
<a href="#">gonoy</a>	<a href="#">gonoy.aSep08</a>		1916	291		95	family member 3 (gonoy) mRNA.
<a href="#">Gopc</a>	<a href="#">Gopc.bSep08</a>	<a href="#">309774</a>	1319	334	1	27	golgi associated PDZ and coiled-coil motif containing (3.2 kD) (Gopc) alternative variant bSep08, mRNA.
<a href="#">gopor</a>	<a href="#">gopor.aSep08</a>		2289	662	4	82	putative protein (9.2 kD) (gopor) alternative variant aSep08, mRNA.
<a href="#">gopor</a>	<a href="#">gopor.bSep08</a>		905	710	2	109	putative protein (12.2 kD) (gopor) alternative variant bSep08, mRNA.

<a href="#">Gorasp1</a>	<a href="#">Gorasp1.bSep08</a>	<a href="#">56082</a>	6117	1926	3	194	golgi reassembly stacking protein 1 (Gorasp1) alternative variant bSep08, mRNA.
<a href="#">Gorasp1</a>	<a href="#">Gorasp1.cSep08</a>	<a href="#">56082</a>	4490	768	4	183	golgi reassembly stacking protein 1 (Gorasp1) alternative variant cSep08, mRNA.
<a href="#">Gorasp2</a>	<a href="#">Gorasp2.bSep08</a>	<a href="#">113961</a>	1604	252	2	71	golgi reassembly stacking protein 2 (Gorasp2) alternative variant bSep08, mRNA.
<a href="#">gorby</a>	<a href="#">gorby.aSep08</a>		1206	243		57	putative protein of eukaryotic origin (gorby) mRNA.
<a href="#">gorchy</a>	<a href="#">gorchy.aSep08</a>		4559	491		115	putative protein of eukaryotic origin (gorchy) mRNA.
<a href="#">gordar</a>	<a href="#">gordar.aSep08</a>		2377	234		37	putative protein (gordar) mRNA.
<a href="#">gorfer</a>	<a href="#">gorfer.aSep08</a>		1954	449		83	putative protein (9.3 kD) (gorfer) mRNA.
<a href="#">gorflo</a>	<a href="#">gorflo.aSep08</a>		1328	620		131	putative protein of mammalian origin (14.1 kD) (gorflo) mRNA.
<a href="#">gorflu</a>	<a href="#">gorflu.aSep08</a>		7572	524		87	putative protein (gorflu) mRNA.
<a href="#">gorkee</a>	<a href="#">gorkee.aSep08</a>		2606	909		88	putative protein (gorkee) mRNA.
<a href="#">gorloy</a>	<a href="#">gorloy.aSep08</a>		4137	410		136	poly polymerase 14 (gorloy) mRNA.
<a href="#">gormee</a>	<a href="#">gormee.aSep08</a>		9051	404		134	C-type lectin domain family 16 member A (gormee) mRNA.
<a href="#">gormer</a>	<a href="#">gormer.aSep08</a>		916	838		67	putative protein (gormer) mRNA.
<a href="#">gornoy</a>	<a href="#">gornoy.aSep08</a>		18573	697	4	56	putative protein of mammalian origin (6.0 kD) (gornoy) mRNA.
<a href="#">gorpor</a>	<a href="#">gorpor.aSep08</a>		2278	507		89	putative protein (gorpor) mRNA.
<a href="#">gorsa</a>	<a href="#">gorsa.aSep08</a>		1890	784		81	putative protein (9.2 kD) (gorsa) mRNA.
<a href="#">gorshee</a>	<a href="#">gorshee.aSep08</a>		6710	456		60	putative protein (6.8 kD) (gorshee) mRNA.
<a href="#">gorto</a>	<a href="#">gorto.aSep08</a>		1712	907		43	putative protein (gorto) mRNA.
<a href="#">gorvar</a>	<a href="#">gorvar.bSep08</a>		406	364	2	30	putative protein (gorvar) alternative variant bSep08, mRNA.
<a href="#">gorwey</a>	<a href="#">gorwey.aSep08</a>		2639	740		95	putative nuclear protein (10.4 kD) (gorwey) mRNA.
<a href="#">gosa</a>	<a href="#">gosa.aSep08</a>		13307	548		182	finger transcription factor trps1 (gosa) mRNA.
<a href="#">goshee</a>	<a href="#">goshee.aSep08</a>		1916	354		15	putative protein (goshee) mRNA.
<a href="#">Gosr2</a>	<a href="#">Gosr2.bSep08</a>	<a href="#">64154</a>	10022	611	4	155	golgi SNAP receptor complex member 2 (Gosr2) alternative variant bSep08, mRNA.
<a href="#">Got1</a>	<a href="#">Got1.bSep08</a>	<a href="#">24401</a>	16046	1975	10	327	glutamate oxaloacetate transaminase 1, soluble (36.8 kD) (Got1) alternative variant bSep08, mRNA.
<a href="#">Got1</a>	<a href="#">Got1.cSep08</a>	<a href="#">24401</a>	14554	592	3	118	glutamate oxaloacetate transaminase 1, soluble (Got1) alternative variant cSep08, mRNA.
<a href="#">Got111</a>	<a href="#">Got111.bSep08</a>	<a href="#">306540</a>	1427	649	1	154	glutamic-oxaloacetic transaminase 1-like 1 (17.5 kD) (Got111) alternative variant bSep08, mRNA.
<a href="#">Got111</a>	<a href="#">Got111.cSep08</a>	<a href="#">306540</a>	789	422	1	83	glutamic-oxaloacetic transaminase 1-like 1 (9.6 kD) (Got111) alternative variant cSep08, mRNA.
<a href="#">Got2</a>	<a href="#">Got2.bSep08</a>	<a href="#">25721</a>	13866	618	4	191	aspartate aminotransferase (Got2) alternative variant bSep08, mRNA.
<a href="#">Got2</a>	<a href="#">Got2.dSep08</a>	<a href="#">25721</a>	1359	656	2	63	aspartate aminotransferase (7.0 kD) (Got2) alternative variant dSep08, mRNA.
<a href="#">goto</a>	<a href="#">goto.aSep08</a>		43323	235		58	putative protein (goto) mRNA.
<a href="#">govar</a>	<a href="#">govar.aSep08</a>		5490	713	1	60	putative protein (govar) alternative variant aSep08, mRNA.

<a href="#">govar</a>	<a href="#">govar.bSep08</a>		5370	288	3	31	putative protein (3.8 kD) (govar) alternative variant bSep08, mRNA.
<a href="#">gowey</a>	<a href="#">gowey.aSep08</a>		26343	394	1	109	putative protein (gowey) alternative variant aSep08, mRNA.
<a href="#">gowey</a>	<a href="#">gowey.bSep08</a>		26454	417	1	105	putative protein (gowey) alternative variant bSep08, mRNA.
<a href="#">goyby</a>	<a href="#">goyby.aSep08</a>		8895	531		93	putative protein (goyby) mRNA.
<a href="#">goychy</a>	<a href="#">goychy.aSep08</a>		5265	377		125	sodium channel voltage-gated type VII alpha (goychy) mRNA.
<a href="#">goydar</a>	<a href="#">goydar.aSep08</a>		75241	303	1	32	putative protein (3.7 kD) (goydar) alternative variant aSep08, mRNA.
<a href="#">goydar</a>	<a href="#">goydar.bSep08</a>		17863	302	2	48	putative protein (goydar) alternative variant bSep08, mRNA.
<a href="#">goyfer</a>	<a href="#">goyfer.aSep08</a>		5267	344		42	putative protein (4.7 kD) (goyfer) mRNA.
<a href="#">goyflo</a>	<a href="#">goyflo.aSep08</a>		6821	292		65	putative protein (goyflo) mRNA.
<a href="#">goyflu</a>	<a href="#">goyflu.aSep08</a>		706	598		94	putative protein of mammalian origin (10.1 kD) (goyflu) mRNA.
<a href="#">goykee</a>	<a href="#">goykee.aSep08</a>		1847	466		33	putative protein (goykee) mRNA.
<a href="#">goyloy</a>	<a href="#">goyloy.aSep08</a>		6255	745		99	putative nuclear protein (10.8 kD) (goyloy) mRNA.
<a href="#">goymee</a>	<a href="#">goymee.aSep08</a>		97138	804	2	63	CRA a like (6.8 kD) (goymee) alternative variant aSep08, mRNA.
<a href="#">goymee</a>	<a href="#">goymee.bSep08</a>		40156	591	1	60	putative protein (goymee) alternative variant bSep08, mRNA.
<a href="#">goymer</a>	<a href="#">goymer.bSep08</a>		883	367		24	putative protein (goymer) alternative variant bSep08, mRNA.
<a href="#">goynoy</a>	<a href="#">goynoy.aSep08</a>		1747	777		101	putative protein (goynoy) mRNA.
<a href="#">goypor</a>	<a href="#">goypor.aSep08</a>		8750	1579	2	101	putative protein of eukaryotic origin (goypor) alternative variant aSep08, mRNA.
<a href="#">goysa</a>	<a href="#">goysa.aSep08</a>		3905	585		118	putative protein (13.2 kD) (goysa) alternative variant aSep08, mRNA.
<a href="#">goysa</a>	<a href="#">goysa.bSep08</a>		3795	614	1	59	putative protein (6.8 kD) (goysa) alternative variant bSep08, mRNA.
<a href="#">goyshee</a>	<a href="#">goyshee.aSep08</a>		7256	622		162	epithelial cell transforming 2 (goyshee) mRNA.
<a href="#">goyto</a>	<a href="#">goyto.aSep08</a>		1275	472	2	37	putative protein (goyto) alternative variant aSep08, mRNA.
<a href="#">goyvar</a>	<a href="#">goyvar.aSep08</a>		4389	367		50	putative protein (goyvar) mRNA.
<a href="#">goywey</a>	<a href="#">goywey.aSep08</a>		2108	693		230	plexin A1 CRA c (goywey) mRNA.
<a href="#">Gp2</a>	<a href="#">Gp2.aSep08</a>	<a href="#">171459</a>	7613	814		190	glycoprotein 2 (zymogen granule membrane) (Gp2) mRNA.
<a href="#">Gpa33</a>	<a href="#">Gpa33.aSep08</a>	<a href="#">360873</a>	20605	538	2	179	glycoprotein A33 (transmembrane) (Gpa33) alternative variant aSep08, mRNA.
<a href="#">Gpa33</a>	<a href="#">Gpa33.bSep08</a>	<a href="#">360873</a>	25480	515	3	69	glycoprotein A33 (transmembrane) (8.0 kD) (Gpa33) alternative variant bSep08, mRNA.
<a href="#">Gpaa1</a>	<a href="#">Gpaa1.bSep08</a>	<a href="#">300046</a>	2387	1542	2	262	GPI anchor attachment protein 1 (Gpaa1) alternative variant bSep08, mRNA.
<a href="#">Gpaa1</a>	<a href="#">Gpaa1.cSep08</a>	<a href="#">300046</a>	1125	716	2	180	GPI anchor attachment protein 1 (Gpaa1) alternative variant cSep08, mRNA.

<a href="#">Gpam</a>	<a href="#">Gpam.bSep08</a>	<a href="#">29653</a>	14205	561	6	161	glycerol-3-phosphate acyltransferase, mitochondrial (Gpam) alternative variant bSep08, mRNA.
<a href="#">Gpatc2</a>	<a href="#">Gpatc2.bSep08</a>	<a href="#">289362</a>	9590	738	1	158	putative protein of vertebrate origin (Gpatc2) alternative variant bSep08, mRNA.
<a href="#">Gpatch1</a>	<a href="#">Gpatch1.bSep08</a>	<a href="#">292810</a>	8537	436	2	56	putative protein (Gpatch1) alternative variant bSep08, mRNA.
<a href="#">Gpatch3</a>	<a href="#">Gpatch3.aSep08</a>	<a href="#">362615</a>	3049	655		218	putative protein of eukaryotic origin (Gpatch3) mRNA.
<a href="#">Gpatch4</a>	<a href="#">Gpatch4.bSep08</a>	<a href="#">295228</a>	7025	614	6	142	d111/G-patch (15.9 kD) (Gpatch4) alternative variant bSep08, complete mRNA.
<a href="#">Gpatch4</a>	<a href="#">Gpatch4.cSep08</a>	<a href="#">295228</a>	11530	2114	3	104	putative protein (11.7 kD) (Gpatch4) alternative variant cSep08, mRNA.
<a href="#">Gbbp1</a>	<a href="#">Gbbp1.bSep08</a>	<a href="#">294734</a>	48217	2038	1	538	GC-rich promoter binding protein 1 like (Gbbp1) alternative variant bSep08, mRNA.
<a href="#">Gbbp1</a>	<a href="#">Gbbp1.cSep08</a>	<a href="#">294734</a>	30749	2223	2	465	GC-rich promoter binding protein 1 like (Gbbp1) alternative variant cSep08, mRNA.
<a href="#">Gbbp1</a>	<a href="#">Gbbp1.dSep08</a>	<a href="#">294734</a>	29941	1478	1	368	GC-rich promoter binding protein 1 like (Gbbp1) alternative variant dSep08, mRNA.
<a href="#">Gbbp1</a>	<a href="#">Gbbp1.eSep08</a>	<a href="#">294734</a>	2627	1448		127	putative protein of mammalian origin (13.3 kD) (Gbbp1) alternative variant eSep08, mRNA.
<a href="#">Gpc4</a>	<a href="#">Gpc4.bSep08</a>	<a href="#">317322</a>	71938	380	2	87	glypican 4 (Gpc4) alternative variant bSep08, mRNA.
<a href="#">Gpc5</a>	<a href="#">Gpc5.bSep08</a>	<a href="#">306157</a>	95212	342	1	103	glypican 5 (Gpc5) alternative variant bSep08, mRNA.
<a href="#">Gpd1l</a>	<a href="#">Gpd1l.aSep08</a>	<a href="#">363159</a>	13104	565		120	glycerol-3-phosphate dehydrogenase 1-like (Gpd1l) mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.bSep08</a>	<a href="#">25062</a>	42178	710	5	220	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant bSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.bSep08</a>	<a href="#">680324</a>	42178	710	5	220	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant bSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.cSep08</a>	<a href="#">25062</a>	42167	719	5	220	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant cSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.cSep08</a>	<a href="#">680324</a>	42167	719	5	220	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant cSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.dSep08</a>	<a href="#">25062</a>	9923	514	5	161	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant dSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.dSep08</a>	<a href="#">680324</a>	9923	514	5	161	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (Gpd2) alternative variant dSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.fSep08</a>	<a href="#">25062</a>	3280	460	2	69	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (7.7 kD) (Gpd2) alternative variant fSep08, mRNA.
<a href="#">Gpd2</a>	<a href="#">Gpd2.fSep08</a>	<a href="#">680324</a>	3280	460	2	69	glycerol-3-phosphate dehydrogenase 2, mitochondrial and hypothetical protein LOC680324 (7.7 kD) (Gpd2) alternative variant fSep08, mRNA.

<a href="#">Gpha2</a>	<a href="#">Gpha2.aSep08</a>	<a href="#">171158</a>	725	423	2	141	glycoprotein hormone alpha 2 (Gpha2) alternative variant aSep08, mRNA.
<a href="#">Gpha2</a>	<a href="#">Gpha2.cSep08</a>	<a href="#">171158</a>	1080	995	2	40	glycoprotein hormone alpha 2 (Gpha2) alternative variant cSep08, mRNA.
<a href="#">Gphn</a>	<a href="#">Gphn.bSep08</a>	<a href="#">64845</a>	69288	1543	7	224	gephyrin (Gphn) alternative variant bSep08, mRNA.
<a href="#">Gphn</a>	<a href="#">Gphn.cSep08</a>	<a href="#">64845</a>	123630	420	4	139	gephyrin (Gphn) alternative variant cSep08, mRNA.
<a href="#">Gphn</a>	<a href="#">Gphn.dSep08</a>	<a href="#">64845</a>	23414	372	4	53	gephyrin (Gphn) alternative variant dSep08, mRNA.
<a href="#">Gphn</a>	<a href="#">Gphn.eSep08</a>	<a href="#">64845</a>	1883	722	2	48	gephyrin (Gphn) alternative variant eSep08, mRNA.
<a href="#">Gpi</a>	<a href="#">Gpi.aSep08</a>	<a href="#">292804</a>	27927	2064	18	558	glucose phosphate isomerase (62.8 kD) (Gpi) alternative variant aSep08, mRNA.
<a href="#">Gpi</a>	<a href="#">Gpi.bSep08</a>	<a href="#">292804</a>	10957	690	7	192	glucose phosphate isomerase (Gpi) alternative variant bSep08, mRNA.
<a href="#">Gpi</a>	<a href="#">Gpi.cSep08</a>	<a href="#">292804</a>	4034	547	5	148	glucose phosphate isomerase (Gpi) alternative variant cSep08, mRNA.
<a href="#">Gpi</a>	<a href="#">Gpi.dSep08</a>	<a href="#">292804</a>	24423	815	3	62	glucose phosphate isomerase (6.7 kD) (Gpi) alternative variant dSep08, mRNA.
<a href="#">Gpi</a>	<a href="#">Gpi.eSep08</a>	<a href="#">292804</a>	976	576	3	64	glucose phosphate isomerase (7.4 kD) (Gpi) alternative variant eSep08, mRNA.
<a href="#">Gpkow</a>	<a href="#">Gpkow.bSep08</a>	<a href="#">679890</a>	2499	486	5	129	gpkow protein (Gpkow) alternative variant bSep08, mRNA.
<a href="#">Gpkow</a>	<a href="#">Gpkow.cSep08</a>	<a href="#">679890</a>	8468	770	5	113	G patch domain KOW motifs (Gpkow) alternative variant cSep08, mRNA.
<a href="#">Gpld1</a>	<a href="#">Gpld1.aSep08</a>	<a href="#">291132</a>	19919	2325	15	559	glycosylphosphatidylinositol specific phospholipase D1 (Gpld1) alternative variant aSep08, mRNA.
<a href="#">Gpld1</a>	<a href="#">Gpld1.bSep08</a>	<a href="#">291132</a>	11685	625	7	208	glycosylphosphatidylinositol specific phospholipase D1 (Gpld1) alternative variant bSep08, mRNA.
<a href="#">Gpld1</a>	<a href="#">Gpld1.cSep08</a>	<a href="#">291132</a>	3416	791	2	48	glycosylphosphatidylinositol specific phospholipase D1 (Gpld1) alternative variant cSep08, mRNA.
<a href="#">Gpm6a</a>	<a href="#">Gpm6a.bSep08</a>	<a href="#">306439</a>	334934	654	4	217	glycoprotein m6a (Gpm6a) alternative variant bSep08, mRNA.
<a href="#">Gpm6a</a>	<a href="#">Gpm6a.cSep08</a>	<a href="#">306439</a>	88095	396	2	17	glycoprotein m6a (1.9 kD) (Gpm6a) alternative variant cSep08, mRNA.
<a href="#">Gpm6b</a>	<a href="#">Gpm6b.aSep08</a>	<a href="#">192179</a>	147903	2885	7	321	glycoprotein m6b (Gpm6b) alternative variant aSep08, mRNA.
<a href="#">Gpm6b</a>	<a href="#">Gpm6b.eSep08</a>	<a href="#">192179</a>	8851	470	2	61	glycoprotein m6b (Gpm6b) alternative variant eSep08, mRNA.
<a href="#">Gpm6b</a>	<a href="#">Gpm6b.fSep08</a>	<a href="#">192179</a>	1818	399	2	36	glycoprotein m6b (Gpm6b) alternative variant fSep08, mRNA.
<a href="#">Gpn2</a>	<a href="#">Gpn2.aSep08</a>	<a href="#">362614</a>	7123	887	4	270	GNP-loop GTPase 2 (Gpn2) alternative variant aSep08, mRNA.
<a href="#">Gpn2</a>	<a href="#">Gpn2.bSep08</a>	<a href="#">362614</a>	6483	1144	4	145	GNP-loop GTPase 2 (16.5 kD) (Gpn2) alternative variant bSep08, mRNA.
<a href="#">Gpn3</a>	<a href="#">Gpn3.bSep08</a>	<a href="#">360810</a>	8234	1405	6	230	GNP-loop GTPase 3 (Gpn3) alternative variant bSep08, mRNA.
<a href="#">Gpn3</a>	<a href="#">Gpn3.cSep08</a>	<a href="#">360810</a>	4795	406	3	113	GNP-loop GTPase 3 (12.6 kD) (Gpn3) alternative variant cSep08, mRNA.

<a href="#">Gpr19</a>	<a href="#">Gpr19.bSep08</a>	<a href="#">312787</a>	26536	882	3	264	G protein-coupled receptor 19 (Gpr19) alternative variant bSep08, mRNA.
<a href="#">Gpr19</a>	<a href="#">Gpr19.bSep08</a>	<a href="#">690642</a>	26536	882	3	264	G protein-coupled receptor 19 (Gpr19) alternative variant bSep08, mRNA.
<a href="#">Gpr19</a>	<a href="#">Gpr19.cSep08</a>	<a href="#">312787</a>	4529	411	3	73	putative protein (Gpr19) alternative variant cSep08, mRNA.
<a href="#">Gpr19</a>	<a href="#">Gpr19.cSep08</a>	<a href="#">690642</a>	4529	411	3	73	putative protein (Gpr19) alternative variant cSep08, mRNA.
<a href="#">Gpr3711</a>	<a href="#">Gpr3711.aSep08</a>	<a href="#">252939</a>	6851	2281	2	495	G protein-coupled receptor 37-like 1 (Gpr3711) alternative variant aSep08, mRNA.
<a href="#">Gpr3711</a>	<a href="#">Gpr3711.bSep08</a>	<a href="#">252939</a>	10218	685	2	144	G protein-coupled receptor 37-like 1 (Gpr3711) alternative variant bSep08, mRNA.
<a href="#">Gpr56</a>	<a href="#">Gpr56.aSep08</a>	<a href="#">260326</a>	19781	3428	11	687	G protein-coupled receptor 56 (77.3 kD) (Gpr56) alternative variant aSep08, mRNA.
<a href="#">Gpr56</a>	<a href="#">Gpr56.bSep08</a>	<a href="#">260326</a>	29302	1433	2	265	G protein-coupled receptor 56 (30.2 kD) (Gpr56) alternative variant bSep08, mRNA.
<a href="#">Gpr56</a>	<a href="#">Gpr56.cSep08</a>	<a href="#">260326</a>	3585	764	2	254	G protein-coupled receptor 56 (Gpr56) alternative variant cSep08, mRNA.
<a href="#">Gpr56</a>	<a href="#">Gpr56.dSep08</a>	<a href="#">260326</a>	10438	692	1	181	G protein-coupled receptor 56 (Gpr56) alternative variant dSep08, mRNA.
<a href="#">Gpr68</a>	<a href="#">Gpr68.cSep08</a>	<a href="#">314386</a>	56258	673	6	129	putative protein (Gpr68) alternative variant cSep08, mRNA.
<a href="#">Gpr85</a>	<a href="#">Gpr85.bSep08</a>	<a href="#">64020</a>	2149	663	1	155	G protein-coupled receptor 85 (Gpr85) alternative variant bSep08, mRNA.
<a href="#">Gpr88</a>	<a href="#">Gpr88.bSep08</a>	<a href="#">64443</a>	6296	709	3	85	G-protein coupled receptor 88 (9.0 kD) (Gpr88) alternative variant bSep08, mRNA.
<a href="#">Gpr89</a>	<a href="#">Gpr89.aSep08</a>	<a href="#">362003</a>	23831	1488	11	335	G protein-coupled receptor 89 (38.4 kD) (Gpr89) alternative variant aSep08, mRNA.
<a href="#">Gpr89</a>	<a href="#">Gpr89.bSep08</a>	<a href="#">362003</a>	24219	1556	6	330	G protein-coupled receptor 89 (Gpr89) alternative variant bSep08, mRNA.
<a href="#">Gpr89</a>	<a href="#">Gpr89.cSep08</a>	<a href="#">362003</a>	15843	711	7	237	G protein-coupled receptor 89 (Gpr89) alternative variant cSep08, mRNA.
<a href="#">Gpr89</a>	<a href="#">Gpr89.dSep08</a>	<a href="#">362003</a>	12540	403	4	134	G protein-coupled receptor 89 (Gpr89) alternative variant dSep08, mRNA.
<a href="#">Gpr97</a>	<a href="#">Gpr97.aSep08</a>	<a href="#">291854</a>	20924	637	6	189	G protein-coupled receptor 97 (Gpr97) alternative variant aSep08, mRNA.
<a href="#">Gpr97</a>	<a href="#">Gpr97.bSep08</a>	<a href="#">291854</a>	19247	517	4	155	G protein-coupled receptor 97 (Gpr97) alternative variant bSep08, mRNA.
<a href="#">Gpr97</a>	<a href="#">Gpr97.cSep08</a>	<a href="#">291854</a>	3012	733	3	108	G protein-coupled receptor 97 (Gpr97) alternative variant cSep08, mRNA.
<a href="#">Gpr98</a>	<a href="#">Gpr98.aSep08</a>	<a href="#">685383</a>	57485	6658		869	G protein-coupled receptor 98 (94.3 kD) (Gpr98) mRNA.
<a href="#">Gpr107</a>	<a href="#">Gpr107.bSep08</a>	<a href="#">311857</a>	4451	803	2	35	G protein-coupled receptor 107 (3.7 kD) (Gpr107) alternative variant bSep08, mRNA.
<a href="#">Gpr108</a>	<a href="#">Gpr108.bSep08</a>	<a href="#">316136</a>	8952	1587	9	357	G protein-coupled receptor 108 (39.3 kD) (Gpr108) alternative variant bSep08, mRNA.
<a href="#">Gpr108</a>	<a href="#">Gpr108.cSep08</a>	<a href="#">316136</a>	9823	2015	13	281	G protein-coupled receptor 108 (30.9 kD) (Gpr108) alternative variant cSep08, mRNA.
<a href="#">Gpr108</a>	<a href="#">Gpr108.dSep08</a>	<a href="#">316136</a>	7218	742	7	207	G protein-coupled receptor 108 (Gpr108) alternative variant dSep08, mRNA.



<a href="#">Gpr108</a>	<a href="#">Gpr108.eSep08</a>	<a href="#">316136</a>	1475	751	3	124	G protein-coupled receptor 108 (Gpr108) alternative variant eSep08, mRNA.
<a href="#">Gpr110</a>	<a href="#">Gpr110.aSep08</a>	<a href="#">301266</a>	12571	617		196	G protein-coupled receptor 110 (Gpr110) mRNA.
<a href="#">Gpr114</a>	<a href="#">Gpr114.bSep08</a>	<a href="#">307645</a>	4724	821	1	273	G protein-coupled receptor 114 (Gpr114) alternative variant bSep08, mRNA.
<a href="#">Gpr115</a>	<a href="#">Gpr115.aSep08</a>	<a href="#">501106</a>	8010	803		53	G protein-coupled receptor 115 (Gpr115) mRNA.
<a href="#">Gpr116</a>	<a href="#">Gpr116.bSep08</a>	<a href="#">245977</a>	11344	2318	8	666	G protein-coupled receptor 116 (Gpr116) alternative variant bSep08, mRNA.
<a href="#">Gpr116</a>	<a href="#">Gpr116.cSep08</a>	<a href="#">245977</a>	11138	814	5	271	G protein-coupled receptor 116 (Gpr116) alternative variant cSep08, mRNA.
<a href="#">Gpr116</a>	<a href="#">Gpr116.dSep08</a>	<a href="#">245977</a>	4848	722	3	184	G protein-coupled receptor 116 (Gpr116) alternative variant dSep08, mRNA.
<a href="#">Gpr116</a>	<a href="#">Gpr116.eSep08</a>	<a href="#">245977</a>	43723	679	5	150	G protein-coupled receptor 116 (Gpr116) alternative variant eSep08, mRNA.
<a href="#">Gpr120</a>	<a href="#">Gpr120.bSep08</a>	<a href="#">294075</a>	12989	863	1	201	G protein-coupled receptor 120 (Gpr120) alternative variant bSep08, mRNA.
<a href="#">Gpr125</a>	<a href="#">Gpr125.aSep08</a>	<a href="#">305408</a>	44679	2947		875	G protein-coupled receptor 125 (Gpr125) mRNA.
<a href="#">Gpr126</a>	<a href="#">Gpr126.aSep08</a>	<a href="#">308376</a>	19766	1771	7	349	G protein-coupled receptor 126 (39.5 kD) (Gpr126) alternative variant aSep08, complete mRNA.
<a href="#">Gpr137</a>	<a href="#">Gpr137.bSep08</a>	<a href="#">689984</a>	2193	882	1	141	G protein-coupled receptor 137 (Gpr137) alternative variant bSep08, mRNA.
<a href="#">Gpr137b</a>	<a href="#">Gpr137b.bSep08</a>	<a href="#">289287</a>	25036	571	3	160	G protein-coupled receptor 137B (Gpr137b) alternative variant bSep08, mRNA.
<a href="#">Gpr137b</a>	<a href="#">Gpr137b.cSep08</a>	<a href="#">289287</a>	6587	2240	4	145	G protein-coupled receptor 137B (Gpr137b) alternative variant cSep08, mRNA.
<a href="#">Gpr137b</a>	<a href="#">Gpr137b.dSep08</a>	<a href="#">289287</a>	833	383	2	88	G protein-coupled receptor 137B (Gpr137b) alternative variant dSep08, mRNA.
<a href="#">Gpr155</a>	<a href="#">Gpr155.bSep08</a>	<a href="#">311730</a>	3559	644	4	129	G protein-coupled receptor 155 (15.5 kD) (Gpr155) alternative variant bSep08, mRNA.
<a href="#">Gpr155</a>	<a href="#">Gpr155.cSep08</a>	<a href="#">311730</a>	3279	587	3	89	G protein-coupled receptor 155 (Gpr155) alternative variant cSep08, mRNA.
<a href="#">Gpr155</a>	<a href="#">Gpr155.dSep08</a>	<a href="#">311730</a>	4986	642	3	125	G protein-coupled receptor 155 (Gpr155) alternative variant dSep08, mRNA.
<a href="#">Gpr158</a>	<a href="#">Gpr158.aSep08</a>	<a href="#">291352</a>	346925	617		205	G protein-coupled receptor 158 (Gpr158) mRNA.
<a href="#">Gpr160</a>	<a href="#">Gpr160.cSep08</a>	<a href="#">499588</a>	56876	735	3	172	G protein-coupled receptor 160 (Gpr160) alternative variant cSep08, mRNA.
<a href="#">Gpr162</a>	<a href="#">Gpr162.bSep08</a>	<a href="#">362436</a>	1510	1363	2	41	G protein-coupled receptor 162 (4.8 kD) (Gpr162) alternative variant bSep08, mRNA.
<a href="#">Gpr172b</a>	<a href="#">Gpr172b.aSep08</a>	<a href="#">362942</a>	5459	5044	2	532	G protein-coupled receptor 172B (56.0 kD) (Gpr172b) alternative variant aSep08, mRNA.
<a href="#">Gpr175</a>	<a href="#">Gpr175.bSep08</a>	<a href="#">85494</a>	9367	1465	8	265	G protein-coupled receptor 175 (29.2 kD) (Gpr175) alternative variant bSep08, mRNA.
<a href="#">Gpr175</a>	<a href="#">Gpr175.cSep08</a>	<a href="#">85494</a>	8465	732	8	199	G protein-coupled receptor 175 (Gpr175) alternative variant cSep08, mRNA.
<a href="#">Gpr175</a>	<a href="#">Gpr175.dSep08</a>	<a href="#">85494</a>	5055	671	8	121	G protein-coupled receptor 175 (Gpr175) alternative variant dSep08, mRNA.

<a href="#">Gpr175</a>	<a href="#">Gpr175.eSep08</a>	<a href="#">85494</a>	5280	765	7	112	G protein-coupled receptor 175 (12.2 kD) (Gpr175) alternative variant eSep08, mRNA.
<a href="#">Gpr175</a>	<a href="#">Gpr175.fSep08</a>	<a href="#">85494</a>	8628	755	7	112	G protein-coupled receptor 175 (12.2 kD) (Gpr175) alternative variant fSep08, mRNA.
<a href="#">Gpr175</a>	<a href="#">Gpr175.hSep08</a>	<a href="#">85494</a>	3165	709	2	50	G protein-coupled receptor 175 (Gpr175) alternative variant hSep08, mRNA.
<a href="#">Gpr176</a>	<a href="#">Gpr176.aSep08</a>	<a href="#">117257</a>	6983	3441		447	G protein-coupled receptor 176 (Gpr176) mRNA.
<a href="#">Gpr177</a>	<a href="#">Gpr177.bSep08</a>	<a href="#">362065</a>	113588	1785	5	306	G protein-coupled receptor 177 (Gpr177) alternative variant bSep08, mRNA.
<a href="#">Gpr177</a>	<a href="#">Gpr177.cSep08</a>	<a href="#">362065</a>	29911	2904	4	163	G protein-coupled receptor 177 (18.5 kD) (Gpr177) alternative variant cSep08, mRNA.
<a href="#">Gpr177</a>	<a href="#">Gpr177.eSep08</a>	<a href="#">362065</a>	3543	641	3	74	G protein-coupled receptor 177 (Gpr177) alternative variant eSep08, mRNA.
<a href="#">Gpr180</a>	<a href="#">Gpr180.bSep08</a>	<a href="#">306165</a>	16772	877	6	292	G protein-coupled receptor 180 (Gpr180) alternative variant bSep08, mRNA.
<a href="#">Gpr180</a>	<a href="#">Gpr180.cSep08</a>	<a href="#">306165</a>	8176	444	3	78	G protein-coupled receptor 180 (Gpr180) alternative variant cSep08, mRNA.
<a href="#">Gpr180</a>	<a href="#">Gpr180.eSep08</a>	<a href="#">306165</a>	1907	539	2	50	G protein-coupled receptor 180 (Gpr180) alternative variant eSep08, mRNA.
<a href="#">Gpr182</a>	<a href="#">Gpr182.aSep08</a>	<a href="#">29307</a>	2330	1824	1	398	G protein-coupled receptor 182 (45.2 kD) (Gpr182) alternative variant aSep08, complete mRNA.
<a href="#">Gpr182</a>	<a href="#">Gpr182.cSep08</a>	<a href="#">29307</a>	1794	409	2	136	G protein-coupled receptor 182 (Gpr182) alternative variant cSep08, mRNA.
<a href="#">Gprc5b</a>	<a href="#">Gprc5b.aSep08</a>	<a href="#">293546</a>	20930	1419	4	458	G protein-coupled receptor, family C, group 5, member B (Gprc5b) alternative variant aSep08, mRNA.
<a href="#">Gprc5b</a>	<a href="#">Gprc5b.cSep08</a>	<a href="#">293546</a>	4708	3443	2	99	G protein-coupled receptor, family C, group 5, member B (Gprc5b) alternative variant cSep08, mRNA.
<a href="#">Gprc5c</a>	<a href="#">Gprc5c.aSep08</a>	<a href="#">287805</a>	18244	1827	2	467	G protein-coupled receptor, family C, group 5, member C (Gprc5c) alternative variant aSep08, mRNA.
<a href="#">Gprc5c</a>	<a href="#">Gprc5c.bSep08</a>	<a href="#">287805</a>	21766	3605	3	464	G protein-coupled receptor, family C, group 5, member C (50.8 kD) (Gprc5c) alternative variant bSep08, mRNA.
<a href="#">Gprc5c</a>	<a href="#">Gprc5c.cSep08</a>	<a href="#">287805</a>	12093	863	1	178	G protein-coupled receptor, family C, group 5, member C (Gprc5c) alternative variant cSep08, mRNA.
<a href="#">Gprc5c</a>	<a href="#">Gprc5c.dSep08</a>	<a href="#">287805</a>	12448	642	1	176	G protein-coupled receptor, family C, group 5, member C (Gprc5c) alternative variant dSep08, mRNA.
<a href="#">Gprc5d</a>	<a href="#">Gprc5d.bSep08</a>	<a href="#">500349</a>	10026	407		40	G protein-coupled receptor, family C, group 5, member D (Gprc5d) alternative variant bSep08, mRNA.
<a href="#">GPS.0</a>	<a href="#">GPS.0.aSep08</a>		5773	1777	7	395	g-protein coupled receptor 124 (GPS.0) alternative variant aSep08, mRNA.
<a href="#">GPS.0</a>	<a href="#">GPS.0.bSep08</a>		2232	787	6	262	G-protein coupled receptor 124 (GPS.0) alternative variant bSep08, mRNA.
<a href="#">Gps2</a>	<a href="#">Gps2.bSep08</a>	<a href="#">497941</a>	1885	785	8	261	G protein pathway suppressor 2 (Gps2) alternative variant bSep08, mRNA.
<a href="#">Gps2</a>	<a href="#">Gps2.cSep08</a>	<a href="#">497941</a>	1883	1186	7	208	G protein pathway suppressor 2 (Gps2) alternative variant cSep08, mRNA.
<a href="#">Gps2</a>	<a href="#">Gps2.dSep08</a>	<a href="#">497941</a>	1364	678	5	173	G protein pathway suppressor 2 (Gps2) alternative variant dSep08, mRNA.

<a href="#">Gps2</a>	<a href="#">Gps2.eSep08</a>	<a href="#">497941</a>	661	399	4	116	G protein pathway suppressor 2 (Gps2) alternative variant eSep08, mRNA.
<a href="#">Gps2</a>	<a href="#">Gps2.fSep08</a>	<a href="#">497941</a>	942	745	2	97	G protein pathway suppressor 2 (10.7 kD) (Gps2) alternative variant fSep08, mRNA.
<a href="#">Gps2l</a>	<a href="#">Gps2l.bSep08</a>	<a href="#">303248</a>	1537	673	6	224	G protein pathway suppressor 2-like (Gps2l) alternative variant bSep08, mRNA.
<a href="#">Gps2l</a>	<a href="#">Gps2l.cSep08</a>	<a href="#">303248</a>	1254	789	5	140	G protein pathway suppressor 2-like (Gps2l) alternative variant cSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.bSep08</a>	<a href="#">246254</a>	10919	2239	6	227	GoLoco (25.1 kD) (Gpsm1) alternative variant bSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.cSep08</a>	<a href="#">246254</a>	8831	643	4	213	tetratricopeptide TPR 1 (Gpsm1) alternative variant cSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.dSep08</a>	<a href="#">246254</a>	4585	774	4	166	GoLoco (18.1 kD) (Gpsm1) alternative variant dSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.eSep08</a>	<a href="#">246254</a>	2748	942	3	166	GoLoco (18.1 kD) (Gpsm1) alternative variant eSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.fSep08</a>	<a href="#">246254</a>	5147	294	3	97	putative protein of vertebrate origin (Gpsm1) alternative variant fSep08, mRNA.
<a href="#">Gpsm1</a>	<a href="#">Gpsm1.gSep08</a>	<a href="#">246254</a>	1100	265	2	88	putative protein (Gpsm1) alternative variant gSep08, mRNA.
<a href="#">Gpsm2</a>	<a href="#">Gpsm2.aSep08</a>	<a href="#">362021</a>	10213	909	4	302	LGN protein (Gpsm2) alternative variant aSep08, mRNA.
<a href="#">Gpsm2</a>	<a href="#">Gpsm2.bSep08</a>	<a href="#">362021</a>	15375	709	3	236	modulator 2 (Gpsm2) alternative variant bSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.bSep08</a>	<a href="#">191576</a>	26142	820	10	272	glycoprotein, synaptic 2 (Gpsn2) alternative variant bSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.cSep08</a>	<a href="#">191576</a>	25913	737	10	227	glycoprotein, synaptic 2 (Gpsn2) alternative variant cSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.dSep08</a>	<a href="#">191576</a>	1980	716	6	189	glycoprotein, synaptic 2 (Gpsn2) alternative variant dSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.eSep08</a>	<a href="#">191576</a>	40590	709	8	182	glycoprotein, synaptic 2 (Gpsn2) alternative variant eSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.fSep08</a>	<a href="#">191576</a>	1247	695	2	131	glycoprotein, synaptic 2 (Gpsn2) alternative variant fSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.gSep08</a>	<a href="#">191576</a>	25286	449	5	113	glycoprotein, synaptic 2 (13.2 kD) (Gpsn2) alternative variant gSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.hSep08</a>	<a href="#">191576</a>	25083	777	4	106	glycoprotein, synaptic 2 (Gpsn2) alternative variant hSep08, mRNA.
<a href="#">Gpsn2</a>	<a href="#">Gpsn2.iSep08</a>	<a href="#">191576</a>	26647	201	3	60	glycoprotein, synaptic 2 (Gpsn2) alternative variant iSep08, mRNA.
<a href="#">Gpt</a>	<a href="#">Gpt.bSep08</a>	<a href="#">81670</a>	1587	704	6	216	glutamic pyruvic transaminase soluble (Gpt) alternative variant bSep08, mRNA.
<a href="#">Gpt</a>	<a href="#">Gpt.cSep08</a>	<a href="#">81670</a>	1034	792	3	88	glutamic-pyruvate transaminase (10.0 kD) (Gpt) alternative variant cSep08, mRNA.
<a href="#">Gpt</a>	<a href="#">Gpt.dSep08</a>	<a href="#">81670</a>	1148	987	3	85	glutamic transaminase (9.7 kD) (Gpt) alternative variant dSep08, mRNA.
<a href="#">Gpt2</a>	<a href="#">Gpt2.bSep08</a>	<a href="#">307759</a>	10833	411	3	136	glutamic pyruvate transaminase (alanine aminotransferase) 2 (Gpt2) alternative variant bSep08, mRNA.

<a href="#">Gpx3</a>	<a href="#">Gpx3.bSep08</a>	<a href="#">64317</a>	7427	921	4	85	glutathione peroxidase 3 (9.3 kD) (Gpx3) alternative variant bSep08, mRNA.
<a href="#">Gpx4</a>	<a href="#">Gpx4.cSep08</a>	<a href="#">29328</a>	1619	726	4	69	glutathione peroxidase 4 (8.0 kD) (Gpx4) alternative variant cSep08, mRNA.
<a href="#">Gpx4</a>	<a href="#">Gpx4.eSep08</a>	<a href="#">29328</a>	844	678	3	52	glutathione peroxidase 4 (Gpx4) alternative variant eSep08, mRNA.
<a href="#">Gpx6</a>	<a href="#">Gpx6.bSep08</a>	<a href="#">259233</a>	2049	927	1	74	glutathione peroxidase 6 (Gpx6) alternative variant bSep08, mRNA.
<a href="#">GRAM.0</a>	<a href="#">GRAM.0.aSep08</a>		49413	783	6	260	GRAM (GRAM.0) alternative variant aSep08, mRNA.
<a href="#">GRAM.0</a>	<a href="#">GRAM.0.bSep08</a>		6757	438	1	52	putative protein of mammalian origin (GRAM.0) alternative variant bSep08, mRNA.
<a href="#">Gramd1a</a>	<a href="#">Gramd1a.bSep08</a>	<a href="#">361550</a>	21531	1601	8	302	putative protein of eukaryotic origin (Gramd1a) alternative variant bSep08, mRNA.
<a href="#">Gramd1a</a>	<a href="#">Gramd1a.cSep08</a>	<a href="#">361550</a>	5923	2004	6	239	gram 1A (27.0 kD) (Gramd1a) alternative variant cSep08, mRNA.
<a href="#">Gramd1a</a>	<a href="#">Gramd1a.dSep08</a>	<a href="#">361550</a>	4448	626	6	174	gram 1A (Gramd1a) alternative variant dSep08, mRNA.
<a href="#">Gramd1a</a>	<a href="#">Gramd1a.eSep08</a>	<a href="#">361550</a>	3326	715	3	48	putative protein (Gramd1a) alternative variant eSep08, mRNA.
<a href="#">Gramd1b</a>	<a href="#">Gramd1b.aSep08</a>	<a href="#">300644</a>	180076	2603	20	758	gram 1B (87.8 kD) (Gramd1b) alternative variant aSep08, mRNA.
<a href="#">Gramd1b</a>	<a href="#">Gramd1b.bSep08</a>	<a href="#">300644</a>	151906	750	8	249	gram 1B (Gramd1b) alternative variant bSep08, mRNA.
<a href="#">Gramd1b</a>	<a href="#">Gramd1b.cSep08</a>	<a href="#">300644</a>	5863	399	4	132	gram 1B (Gramd1b) alternative variant cSep08, mRNA.
<a href="#">Gramd1b</a>	<a href="#">Gramd1b.dSep08</a>	<a href="#">300644</a>	77948	392	4	83	putative protein (9.0 kD) (Gramd1b) alternative variant dSep08, mRNA.
<a href="#">Gramd1b</a>	<a href="#">Gramd1b.fSep08</a>	<a href="#">300644</a>	60673	404	3	43	putative protein (Gramd1b) alternative variant fSep08, mRNA.
<a href="#">Gramd1c</a>	<a href="#">Gramd1c.aSep08</a>	<a href="#">360717</a>	29923	939	7	313	putative protein of eukaryotic origin (Gramd1c) alternative variant aSep08, mRNA.
<a href="#">Gramd1c</a>	<a href="#">Gramd1c.bSep08</a>	<a href="#">360717</a>	3976	703	2	98	putative nuclear protein of vertebrate origin (11.2 kD) (Gramd1c) alternative variant bSep08, mRNA.
<a href="#">Gramd2</a>	<a href="#">Gramd2.aSep08</a>	<a href="#">300761</a>	6092	994	6	260	GRAM (Gramd2) alternative variant aSep08, mRNA.
<a href="#">Gramd2</a>	<a href="#">Gramd2.bSep08</a>	<a href="#">300761</a>	2947	842	1	110	putative nuclear protein of vertebrate origin (12.4 kD) (Gramd2) alternative variant bSep08, mRNA.
<a href="#">Gramd3</a>	<a href="#">Gramd3.bSep08</a>	<a href="#">307288</a>	50851	770	7	256	GRAM (Gramd3) alternative variant bSep08, mRNA.
<a href="#">Gramd3</a>	<a href="#">Gramd3.cSep08</a>	<a href="#">307288</a>	24021	842	9	246	GRAM (Gramd3) alternative variant cSep08, mRNA.
<a href="#">Gramd4</a>	<a href="#">Gramd4.aSep08</a>	<a href="#">315203</a>	64306	1018		337	death-inducing-protein (Gramd4) mRNA.
<a href="#">Grap</a>	<a href="#">Grap.bSep08</a>	<a href="#">363616</a>	2625	803	2	152	GRB2-related adaptor protein (17.2 kD) (Grap) alternative variant bSep08, mRNA.
<a href="#">Grap2</a>	<a href="#">Grap2.bSep08</a>	<a href="#">366962</a>	2245	470	3	103	GRB2-related adaptor protein 2 (Grap2) alternative variant bSep08, mRNA.
<a href="#">Grap2</a>	<a href="#">Grap2.cSep08</a>	<a href="#">366962</a>	5885	442	2	35	GRB2-related adaptor protein 2 (3.9 kD) (Grap2) alternative variant cSep08, mRNA.
<a href="#">Grb2</a>	<a href="#">Grb2.bSep08</a>	<a href="#">81504</a>	85503	1108	5	175	growth factor receptor bound protein 2 (20.3 kD) (Grb2) alternative variant bSep08, mRNA.
<a href="#">Grb2</a>	<a href="#">Grb2.dSep08</a>	<a href="#">81504</a>	79287	308	2	36	growth factor receptor bound protein 2 (Grb2) alternative variant dSep08, mRNA.

<a href="#">Grb2</a>	<a href="#">Grb2.eSep08</a>	<a href="#">81504</a>	84613	396	3	78	growth factor receptor bound protein 2 (Grb2) alternative variant eSep08, mRNA.
<a href="#">Grb7</a>	<a href="#">Grb7.bSep08</a>	<a href="#">84427</a>	971	761	3	119	growth factor receptor bound protein 7 (Grb7) alternative variant bSep08, mRNA.
<a href="#">Grb7</a>	<a href="#">Grb7.cSep08</a>	<a href="#">84427</a>	4530	867	3	103	growth factor receptor bound protein 7 (Grb7) alternative variant cSep08, mRNA.
<a href="#">Grb10</a>	<a href="#">Grb10.bSep08</a>	<a href="#">498416</a>	3734	565	2	130	growth factor receptor bound protein 10 (Grb10) alternative variant bSep08, mRNA.
<a href="#">Grb10</a>	<a href="#">Grb10.cSep08</a>	<a href="#">498416</a>	7319	3107	4	88	factor receptor-bound protein 10 (Grb10) alternative variant cSep08, mRNA.
<a href="#">Grhl1</a>	<a href="#">Grhl1.aSep08</a>	<a href="#">313993</a>	9026	645		214	grainyhead-like 1 (Drosophila) (Grhl1) mRNA.
<a href="#">Grhl2</a>	<a href="#">Grhl2.bSep08</a>	<a href="#">299979</a>	33538	887	2	184	grainyhead-like 2 (Drosophila) (Grhl2) alternative variant bSep08, mRNA.
<a href="#">Grhpr</a>	<a href="#">Grhpr.aSep08</a>	<a href="#">680021</a>	9230	1784	5	528	glyoxylate reductase/hydroxypyruvate reductase (Grhpr) alternative variant aSep08, complete mRNA.
<a href="#">Grhpr</a>	<a href="#">Grhpr.bSep08</a>	<a href="#">680021</a>	4349	634	4	148	glyoxylate reductase/hydroxypyruvate reductase (Grhpr) alternative variant bSep08, mRNA.
<a href="#">Grhpr</a>	<a href="#">Grhpr.cSep08</a>	<a href="#">680021</a>	3173	422	2	128	glyoxylate reductase/hydroxypyruvate reductase (Grhpr) alternative variant cSep08, mRNA.
<a href="#">Grhpr</a>	<a href="#">Grhpr.dSep08</a>	<a href="#">680021</a>	3179	1799	1	93	glyoxylate reductase/hydroxypyruvate reductase (10.0 kD) (Grhpr) alternative variant dSep08, complete mRNA.
<a href="#">Gria2</a>	<a href="#">Gria2.dSep08</a>	<a href="#">29627</a>	2369	1977	2	56	glutamate receptor, ionotropic, AMPA 2 (6.4 kD) (Gria2) alternative variant dSep08, mRNA.
<a href="#">Grid1</a>	<a href="#">Grid1.bSep08</a>	<a href="#">79219</a>	36550	330	2	110	glutamate receptor, ionotropic, delta 1 (Grid1) alternative variant bSep08, mRNA.
<a href="#">Grid2ip</a>	<a href="#">Grid2ip.bSep08</a>	<a href="#">288484</a>	690	404	1	121	glutamate receptor, ionotropic, delta 2 (Grid2) interacting protein 1 (Grid2ip) alternative variant bSep08, mRNA.
<a href="#">Grifin</a>	<a href="#">Grifin.bSep08</a>	<a href="#">117130</a>	1492	426	1	141	galectin-related inter-fiber protein (Grifin) alternative variant bSep08, mRNA.
<a href="#">Grik1</a>	<a href="#">Grik1.dSep08</a>	<a href="#">29559</a>	1761	478	2	128	glutamate receptor, ionotropic, kainate 1 (Grik1) alternative variant dSep08, mRNA.
<a href="#">Grik1</a>	<a href="#">Grik1.eSep08</a>	<a href="#">29559</a>	19399	772	2	113	glutamate receptor, ionotropic, kainate 1 (12.7 kD) (Grik1) alternative variant eSep08, mRNA.
<a href="#">Grik4</a>	<a href="#">Grik4.bSep08</a>	<a href="#">24406</a>	22142	749	1	179	glutamate receptor, ionotropic, kainate 4 (Grik4) alternative variant bSep08, mRNA.
<a href="#">Grik5</a>	<a href="#">Grik5.bSep08</a>	<a href="#">24407</a>	70192	1788	4	595	glutamate receptor, ionotropic, kainate 5 (Grik5) alternative variant bSep08, mRNA.
<a href="#">Grik5</a>	<a href="#">Grik5.cSep08</a>	<a href="#">24407</a>	4890	1021	7	340	glutamate receptor, ionotropic, kainate 5 (Grik5) alternative variant cSep08, mRNA.
<a href="#">Grik5</a>	<a href="#">Grik5.dSep08</a>	<a href="#">24407</a>	3238	394	3	131	glutamate receptor, ionotropic, kainate 5 (Grik5) alternative variant dSep08, mRNA.
<a href="#">Grik5</a>	<a href="#">Grik5.eSep08</a>	<a href="#">24407</a>	1005	740	2	90	glutamate receptor, ionotropic, kainate 5 (Grik5) alternative variant eSep08, mRNA.
<a href="#">Grin1</a>	<a href="#">Grin1.bSep08</a>	<a href="#">24408</a>	5266	867	6	221	glutamate receptor, ionotropic, N-methyl D-aspartate 1 (Grin1) alternative variant bSep08, mRNA.

<a href="#">Grina</a>	<a href="#">Grina.bSep08</a>	<a href="#">266668</a>	1451	511	1	170	glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1 (glutamate binding) (Grina) alternative variant bSep08, mRNA.
<a href="#">Grin1a</a>	<a href="#">Grin1a.bSep08</a>	<a href="#">192147</a>	1611	780	1	28	glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A (Grin1a) alternative variant bSep08, mRNA.
<a href="#">Grip1</a>	<a href="#">Grip1.bSep08</a>	<a href="#">84016</a>	28376	971	6	273	glutamate receptor interacting protein 1 (Grip1) alternative variant bSep08, mRNA.
<a href="#">Grip1</a>	<a href="#">Grip1.cSep08</a>	<a href="#">84016</a>	14098	585	3	111	glutamate receptor interacting protein 1 (Grip1) alternative variant cSep08, mRNA.
<a href="#">Grip2</a>	<a href="#">Grip2.bSep08</a>	<a href="#">171571</a>	3075	726	1	179	glutamate receptor interacting protein 2 (Grip2) alternative variant bSep08, mRNA.
<a href="#">Gripap1</a>	<a href="#">Gripap1.bSep08</a>	<a href="#">116493</a>	8513	562	8	187	GRIP1 associated protein 1 (Gripap1) alternative variant bSep08, mRNA.
<a href="#">Gripap1</a>	<a href="#">Gripap1.dSep08</a>	<a href="#">116493</a>	611	472	2	52	GRIP1 associated protein 1 (Gripap1) alternative variant dSep08, mRNA.
<a href="#">Gripap1</a>	<a href="#">Gripap1.eSep08</a>	<a href="#">116493</a>	9681	390	3	42	GRIP1 associated protein 1 (Gripap1) alternative variant eSep08, mRNA.
<a href="#">Grit</a>	<a href="#">Grit.aSep08</a>	<a href="#">315530</a>	41311	536	4	178	rho GTPase-activating protein (Grit) alternative variant aSep08, mRNA.
<a href="#">Grit</a>	<a href="#">Grit.bSep08</a>	<a href="#">315530</a>	22203	380	3	91	rho GTPase-activating protein (Grit) alternative variant bSep08, mRNA.
<a href="#">Grit</a>	<a href="#">Grit.cSep08</a>	<a href="#">315530</a>	15575	410	1	20	rho GTPase-activating protein (Grit) alternative variant cSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.bSep08</a>	<a href="#">59077</a>	37084	1102	7	317	G protein-coupled receptor kinase 4 (Grk4) alternative variant bSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.cSep08</a>	<a href="#">59077</a>	12897	731	5	205	G protein-coupled receptor kinase 4 (Grk4) alternative variant cSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.dSep08</a>	<a href="#">59077</a>	8165	610	3	196	G protein-coupled receptor kinase 4 (Grk4) alternative variant dSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.eSep08</a>	<a href="#">59077</a>	7433	560	3	76	G protein-coupled receptor kinase 4 (Grk4) alternative variant eSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.fSep08</a>	<a href="#">59077</a>	4894	457	4	53	G protein-coupled receptor kinase 4 (Grk4) alternative variant fSep08, mRNA.
<a href="#">Grk4</a>	<a href="#">Grk4.hSep08</a>	<a href="#">59077</a>	12125	445	5	40	G protein-coupled receptor kinase 4 (Grk4) alternative variant hSep08, mRNA.
<a href="#">Grk6</a>	<a href="#">Grk6.cSep08</a>	<a href="#">59076</a>	8160	1455	2	300	G protein-coupled receptor kinase 6 (33.9 kD) (Grk6) alternative variant cSep08, mRNA.
<a href="#">Grif1</a>	<a href="#">Grif1.aSep08</a>	<a href="#">306400</a>	64111	1606		408	glucocorticoid receptor DNA binding factor 1 (Grif1) mRNA.
<a href="#">Grm5</a>	<a href="#">Grm5.aSep08</a>	<a href="#">24418</a>	613778	3612	1	1203	glutamate receptor, metabotropic 5 (Grm5) alternative variant aSep08, mRNA.
<a href="#">Grm7</a>	<a href="#">Grm7.bSep08</a>	<a href="#">81672</a>	182695	1513	1	416	glutamate receptor, metabotropic 7 (Grm7) alternative variant bSep08, mRNA.
<a href="#">Grn</a>	<a href="#">Grn.bSep08</a>	<a href="#">29143</a>	689	591	2	158	granulin (Grn) alternative variant bSep08, mRNA.
<a href="#">Grn</a>	<a href="#">Grn.cSep08</a>	<a href="#">29143</a>	701	174	2	58	granulin (Grn) alternative variant cSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.bSep08</a>	<a href="#">192266</a>	68999	650	4	216	glutamine glutamic acid-rich protein (Grpca) alternative variant bSep08, mRNA.

<a href="#">Grpca</a>	<a href="#">Grpca.bSep08</a>	<a href="#">360395</a>	68999	650	4	216	glutamine glutamic acid-rich protein (Grpca) alternative variant bSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.cSep08</a>	<a href="#">192266</a>	5146	732	4	212	glutamine glutamic acid-rich protein (Grpca) alternative variant cSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.cSep08</a>	<a href="#">360395</a>	5146	732	4	212	glutamine glutamic acid-rich protein (Grpca) alternative variant cSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.dSep08</a>	<a href="#">192266</a>	68615	691	3	198	glutamine glutamic acid-rich protein (Grpca) alternative variant dSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.dSep08</a>	<a href="#">360395</a>	68615	691	3	198	glutamine glutamic acid-rich protein (Grpca) alternative variant dSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.eSep08</a>	<a href="#">192266</a>	5047	657	5	187	submandibular gland secretory protein like (Grpca) alternative variant eSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.eSep08</a>	<a href="#">360395</a>	5047	657	5	187	submandibular gland secretory protein like (Grpca) alternative variant eSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.fSep08</a>	<a href="#">192266</a>	3619	621	4	175	submandibular gland secretory protein (Grpca) alternative variant fSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.fSep08</a>	<a href="#">360395</a>	3619	621	4	175	submandibular gland secretory protein (Grpca) alternative variant fSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.gSep08</a>	<a href="#">192266</a>	1804	571	2		
<a href="#">Grpca</a>	<a href="#">Grpca.gSep08</a>	<a href="#">360395</a>	1804	571	2		
<a href="#">Grpca</a>	<a href="#">Grpca.hSep08</a>	<a href="#">192266</a>	1403	180	2	15	putative protein (1.7 kD) (Grpca) alternative variant hSep08, mRNA.
<a href="#">Grpca</a>	<a href="#">Grpca.hSep08</a>	<a href="#">360395</a>	1403	180	2	15	putative protein (1.7 kD) (Grpca) alternative variant hSep08, mRNA.
<a href="#">Grpel1</a>	<a href="#">Grpel1.aSep08</a>	<a href="#">79563</a>	6517	1857	2	230	GrpE-like 1, mitochondrial (25.8 kD) (Grpel1) alternative variant aSep08, mRNA.
<a href="#">Grpel2</a>	<a href="#">Grpel2.bSep08</a>	<a href="#">688777</a>	10821	759	5	207	GrpE-like 2, mitochondrial (Grpel2) alternative variant bSep08, mRNA.
<a href="#">Grpel2</a>	<a href="#">Grpel2.cSep08</a>	<a href="#">688777</a>	5514	418	2	41	GrpE-like 2, mitochondrial (Grpel2) alternative variant cSep08, mRNA.
<a href="#">Grsf1</a>	<a href="#">Grsf1.aSep08</a>	<a href="#">305256</a>	15937	2420	9	393	G-rich RNA sequence binding factor 1 (Grsf1) alternative variant aSep08, mRNA.
<a href="#">Grsf1</a>	<a href="#">Grsf1.bSep08</a>	<a href="#">305256</a>	9191	768	6	256	G-rich RNA sequence binding factor 1 (Grsf1) alternative variant bSep08, mRNA.
<a href="#">Grsf1</a>	<a href="#">Grsf1.cSep08</a>	<a href="#">305256</a>	4990	613	5	204	G-rich RNA sequence binding factor 1 (Grsf1) alternative variant cSep08, mRNA.
<a href="#">Grsf1</a>	<a href="#">Grsf1.dSep08</a>	<a href="#">305256</a>	1447	709	2	81	G-rich RNA sequence binding factor 1 (Grsf1) alternative variant dSep08, mRNA.
<a href="#">Grsf1</a>	<a href="#">Grsf1.eSep08</a>	<a href="#">305256</a>	3156	730	1	49	G-rich RNA sequence binding factor 1 (5.7 kD) (Grsf1) alternative variant eSep08, mRNA.
<a href="#">Grwd1</a>	<a href="#">Grwd1.bSep08</a>	<a href="#">308592</a>	3490	2190		313	glutamate-rich WD repeat containing 1 (34.5 kD) (Grwd1) alternative variant bSep08, mRNA.
<a href="#">Gsbs</a>	<a href="#">Gsbs.bSep08</a>	<a href="#">266705</a>	15880	731		149	G substrate (Gsbs) alternative variant bSep08, mRNA.
<a href="#">Gsdma1</a>	<a href="#">Gsdma1.bSep08</a>	<a href="#">360619</a>	11240	2545	8	235	gasdermin A1 (26.2 kD) (Gsdma1) alternative variant bSep08, complete mRNA.
<a href="#">Gsdma1</a>	<a href="#">Gsdma1.cSep08</a>	<a href="#">360619</a>	3607	651	2	141	gasdermin A1 (Gsdma1) alternative variant cSep08, mRNA.

<a href="#">Gsdmd</a>	<a href="#">Gsdmd.bSep08</a>	<a href="#">315084</a>	1179	395	2	131	gasdermin D (Gsdmd) alternative variant bSep08, mRNA.
<a href="#">Gsdmd</a>	<a href="#">Gsdmd.dSep08</a>	<a href="#">315084</a>	1473	576	3	91	gasdermin D (Gsdmd) alternative variant dSep08, mRNA.
<a href="#">Gsg1</a>	<a href="#">Gsg1.bSep08</a>	<a href="#">312793</a>	4643	705	1	157	germ cell associated 1 (Gsg1) alternative variant bSep08, mRNA.
<a href="#">Gsg1l</a>	<a href="#">Gsg1l.aSep08</a>	<a href="#">499263</a>	155264	1372	2	230	GSg1-like (26.2 kD) (Gsg1l) alternative variant aSep08, mRNA.
<a href="#">Gsg1l</a>	<a href="#">Gsg1l.bSep08</a>	<a href="#">499263</a>	129452	472	1	97	GSg1-like (Gsg1l) alternative variant bSep08, mRNA.
<a href="#">Gsk3b</a>	<a href="#">Gsk3b.bSep08</a>	<a href="#">84027</a>	75347	1349	9	366	glycogen synthase kinase 3 beta (41.6 kD) (Gsk3b) alternative variant bSep08, mRNA.
<a href="#">Gsk3b</a>	<a href="#">Gsk3b.cSep08</a>	<a href="#">84027</a>	20294	494	4	128	glycogen synthase kinase 3 beta (Gsk3b) alternative variant cSep08, mRNA.
<a href="#">Gsk3b</a>	<a href="#">Gsk3b.dSep08</a>	<a href="#">84027</a>	8356	281	3	80	glycogen synthase kinase 3 beta (Gsk3b) alternative variant dSep08, mRNA.
<a href="#">Gsn</a>	<a href="#">Gsn.aSep08</a>	<a href="#">296654</a>	26858	3372	1	491	gelsolin (54.6 kD) (Gsn) alternative variant aSep08, mRNA.
<a href="#">Gsn</a>	<a href="#">Gsn.bSep08</a>	<a href="#">296654</a>	5026	1023		260	gelsolin (Gsn) alternative variant bSep08, mRNA.
<a href="#">Gsn</a>	<a href="#">Gsn.cSep08</a>	<a href="#">296654</a>	11135	744	2	245	gelsolin (Gsn) alternative variant cSep08, mRNA.
<a href="#">Gsn</a>	<a href="#">Gsn.dSep08</a>	<a href="#">296654</a>	37523	827	2	220	gelsolin (Gsn) alternative variant dSep08, mRNA.
<a href="#">Gsn</a>	<a href="#">Gsn.eSep08</a>	<a href="#">296654</a>	26284	726	1	206	gelsolin (Gsn) alternative variant eSep08, mRNA.
<a href="#">Gspt1</a>	<a href="#">Gspt1.bSep08</a>	<a href="#">24420</a>	5748	771	4	180	g1 to S phase transition 1 (Gspt1) alternative variant bSep08, mRNA.
<a href="#">Gspt1</a>	<a href="#">Gspt1.cSep08</a>	<a href="#">24420</a>	12993	517	6	120	g1 to S phase transition 1 (Gspt1) alternative variant cSep08, mRNA.
<a href="#">Gspt1</a>	<a href="#">Gspt1.dSep08</a>	<a href="#">24420</a>	5087	361	4	120	g1 to S phase transition 1 (Gspt1) alternative variant dSep08, mRNA.
<a href="#">Gsr</a>	<a href="#">Gsr.bSep08</a>	<a href="#">116686</a>	17335	319	5	106	glutathione reductase (Gsr) alternative variant bSep08, mRNA.
<a href="#">Gstcd</a>	<a href="#">Gstcd.bSep08</a>	<a href="#">310855</a>	78420	859	7	286	putative protein, with a transmembrane domain, of ancient origin (Gstcd) alternative variant bSep08, mRNA.
<a href="#">Gstcd</a>	<a href="#">Gstcd.cSep08</a>	<a href="#">310855</a>	68369	669	6	223	putative protein of ancient origin (Gstcd) alternative variant cSep08, mRNA.
<a href="#">Gstcd</a>	<a href="#">Gstcd.eSep08</a>	<a href="#">310855</a>	5766	390	2	76	putative protein (Gstcd) alternative variant eSep08, mRNA.
<a href="#">Gstcd</a>	<a href="#">Gstcd.fSep08</a>	<a href="#">310855</a>	1616	579	2	66	putative protein of bilateral origin (7.0 kD) (Gstcd) alternative variant fSep08, mRNA.
<a href="#">Gstk1</a>	<a href="#">Gstk1.bSep08</a>	<a href="#">297029</a>	969	785	1	63	glutathione S-transferase kappa 1 (7.3 kD) (Gstk1) alternative variant bSep08, mRNA.
<a href="#">Gstm1</a>	<a href="#">Gstm1.bSep08</a>	<a href="#">24423</a>	3363	879	8	206	glutathione S-transferase mu 1 (24.3 kD) (Gstm1) alternative variant bSep08, mRNA.
<a href="#">Gstm1</a>	<a href="#">Gstm1.cSep08</a>	<a href="#">24423</a>	1212	287	4	93	A First-Sphere Second-Sphere Electrostatic Effects In The Active Site Of Class mu glutathione transferase (Gstm1) alternative variant cSep08, mRNA.
<a href="#">Gstm1</a>	<a href="#">Gstm1.eSep08</a>	<a href="#">24423</a>	1408	593	2	43	a First-Sphere Second-Sphere Electrostatic Effects In The Active Site Of Class mu glutathione transferase (5.0 kD) (Gstm1) alternative variant eSep08, mRNA.
<a href="#">Gstm2</a>	<a href="#">Gstm2.bSep08</a>	<a href="#">24424</a>	2307	402	6	133	glutathione S-transferase, mu 2 (Gstm2) alternative variant bSep08, mRNA.



<a href="#">Gstm2</a>	<a href="#">Gstm2.dSep08</a>	<a href="#">24424</a>	4323	1807	6	160	glutathione S-transferase, mu 2 (17.8 kD) (Gstm2) alternative variant dSep08, mRNA.
<a href="#">Gstm3</a>	<a href="#">Gstm3.bSep08</a>	<a href="#">81869</a>	5048	745	7	182	glutathione S-transferase M2 (21.4 kD) (Gstm3) alternative variant bSep08, mRNA.
<a href="#">Gstm3</a>	<a href="#">Gstm3.cSep08</a>	<a href="#">81869</a>	2640	585	6	170	glutathione S-transferase mu (Gstm3) alternative variant cSep08, mRNA.
<a href="#">Gstm3</a>	<a href="#">Gstm3.dSep08</a>	<a href="#">81869</a>	832	747	2	46	glutathione S-transferase M2 (Gstm3) alternative variant dSep08, mRNA.
<a href="#">Gstm3</a>	<a href="#">Gstm3.eSep08</a>	<a href="#">81869</a>	1156	889	2	42	glutathione S-transferase M1 (4.9 kD) (Gstm3) alternative variant eSep08, mRNA.
<a href="#">Gstm4</a>	<a href="#">Gstm4.bSep08</a>	<a href="#">57298</a>	5205	765	4	112	glutathione S-transferase M4 (Gstm4) alternative variant bSep08, mRNA.
<a href="#">Gstm4</a>	<a href="#">Gstm4.cSep08</a>	<a href="#">57298</a>	2351	652	2	75	glutathione S-transferase M4 (Gstm4) alternative variant cSep08, mRNA.
<a href="#">Gstm5</a>	<a href="#">Gstm5.bSep08</a>	<a href="#">64352</a>	2566	793	1	158	glutathione S-transferase, mu 5 (18.7 kD) (Gstm5) alternative variant bSep08, mRNA.
<a href="#">Gstm6l</a>	<a href="#">Gstm6l.aSep08</a>	<a href="#">295362</a>	6125	1755	3	244	glutathione S-transferase, mu 6-like (Gstm6l) alternative variant aSep08, mRNA.
<a href="#">Gstm6l</a>	<a href="#">Gstm6l.cSep08</a>	<a href="#">295362</a>	2390	599	2	47	glutathione S-transferase, mu 6-like (Gstm6l) alternative variant cSep08, mRNA.
<a href="#">Gsto1</a>	<a href="#">Gsto1.bSep08</a>	<a href="#">114846</a>	10372	1357	5	213	glutathione S-transferase omega 1 (24.9 kD) (Gsto1) alternative variant bSep08, complete mRNA.
<a href="#">Gsto1</a>	<a href="#">Gsto1.cSep08</a>	<a href="#">114846</a>	2651	687	2	97	glutathione S-transferase omega 1 (Gsto1) alternative variant cSep08, mRNA.
<a href="#">Gsto1</a>	<a href="#">Gsto1.dSep08</a>	<a href="#">114846</a>	1611	814	2	85	glutathione S-transferase omega 1 (9.9 kD) (Gsto1) alternative variant dSep08, mRNA.
<a href="#">Gsto1</a>	<a href="#">Gsto1.eSep08</a>	<a href="#">114846</a>	1467	285	3	60	glutathione S-transferase omega (Gsto1) alternative variant eSep08, mRNA.
<a href="#">Gsto2</a>	<a href="#">Gsto2.bSep08</a>	<a href="#">309465</a>	35394	1187	8	134	glutathione S-transferase omega 2 (15.4 kD) (Gsto2) alternative variant bSep08, mRNA.
<a href="#">Gsto2</a>	<a href="#">Gsto2.fSep08</a>	<a href="#">309465</a>	4893	148	2	48	glutathione S-transferase omega 2 (Gsto2) alternative variant fSep08, mRNA.
<a href="#">Gstp1</a>	<a href="#">Gstp1.bSep08</a>	<a href="#">24426</a>	617	461	2	132	glutathione S-transferase (Gstp1) alternative variant bSep08, mRNA.
<a href="#">Gstp1</a>	<a href="#">Gstp1.bSep08</a>	<a href="#">29438</a>	617	461	2	132	glutathione S-transferase (Gstp1) alternative variant bSep08, mRNA.
<a href="#">Gstp1</a>	<a href="#">Gstp1.cSep08</a>	<a href="#">24426</a>	1250	425	5	123	glutathione (Gstp1) alternative variant cSep08, mRNA.
<a href="#">Gstp1</a>	<a href="#">Gstp1.cSep08</a>	<a href="#">29438</a>	1250	425	5	123	glutathione (Gstp1) alternative variant cSep08, mRNA.
<a href="#">Gstt1</a>	<a href="#">Gstt1.bSep08</a>	<a href="#">25260</a>	15918	559	1	131	glutathione S-transferase theta 1 (Gstt1) alternative variant bSep08, mRNA.
<a href="#">Gstt2</a>	<a href="#">Gstt2.cSep08</a>	<a href="#">29487</a>	735	441	2	139	glutathione S-transferase, theta 2 (Gstt2) alternative variant cSep08, mRNA.
<a href="#">Gstt3</a>	<a href="#">Gstt3.aSep08</a>	<a href="#">499422</a>	6851	1661	4	300	glutathione S-transferase, theta 3 (Gstt3) alternative variant aSep08, mRNA.
<a href="#">Gstt3</a>	<a href="#">Gstt3.bSep08</a>	<a href="#">499422</a>	4701	1084	4	223	glutathione S-transferase, theta 3 (Gstt3) alternative variant bSep08, mRNA.

<a href="#">Gstt3</a>	<a href="#">Gstt3.dSep08</a>	<a href="#">499422</a>	4323	656	3	85	glutathione S-transferase, theta 3 (Gstt3) alternative variant dSep08, mRNA.
<a href="#">Gtdc1</a>	<a href="#">Gtdc1.bSep08</a>	<a href="#">362129</a>	41535	798	1	137	putative protein of ancient origin (Gtdc1) alternative variant bSep08, mRNA.
<a href="#">Gtf2a2</a>	<a href="#">Gtf2a2.cSep08</a>	<a href="#">83828</a>	13128	795	6	61	general transcription factor IIa 2 (7.1 kD) (Gtf2a2) alternative variant cSep08, complete mRNA.
<a href="#">Gtf2e1</a>	<a href="#">Gtf2e1.aSep08</a>	<a href="#">303918</a>	33821	2882		438	general transcription factor II E, polypeptide 1 (alpha subunit) (49.2 kD) (Gtf2e1) mRNA.
<a href="#">Gtf2e2</a>	<a href="#">Gtf2e2.bSep08</a>	<a href="#">306516</a>	35270	726	6	241	general transcription factor II E, polypeptide 2 (beta subunit) (Gtf2e2) alternative variant bSep08, mRNA.
<a href="#">Gtf2e2</a>	<a href="#">Gtf2e2.cSep08</a>	<a href="#">306516</a>	1478	1083	2	93	general transcription factor II E, polypeptide 2 (beta subunit) (10.9 kD) (Gtf2e2) alternative variant cSep08, mRNA.
<a href="#">Gtf2f1</a>	<a href="#">Gtf2f1.bSep08</a>	<a href="#">316123</a>	5562	746	6	194	general transcription factor IIF polypeptide 1 CRA a (Gtf2f1) alternative variant bSep08, mRNA.
<a href="#">Gtf2f1</a>	<a href="#">Gtf2f1.cSep08</a>	<a href="#">316123</a>	5527	729	6	184	general transcription factor IIF polypeptide 1 74kDa CRA b (Gtf2f1) alternative variant cSep08, mRNA.
<a href="#">Gtf2f1</a>	<a href="#">Gtf2f1.dSep08</a>	<a href="#">316123</a>	3705	701	5	109	general transcription factor IIF polypeptide 1 CRA a (12.6 kD) (Gtf2f1) alternative variant dSep08, mRNA.
<a href="#">Gtf2f1</a>	<a href="#">Gtf2f1.eSep08</a>	<a href="#">316123</a>	748	558	3	84	general transcription factor iif (9.5 kD) (Gtf2f1) alternative variant eSep08, mRNA.
<a href="#">Gtf2f2</a>	<a href="#">Gtf2f2.bSep08</a>	<a href="#">81674</a>	55825	483	1	102	general transcription factor IIF, polypeptide 2 (Gtf2f2) alternative variant bSep08, mRNA.
<a href="#">Gtf2h1</a>	<a href="#">Gtf2h1.bSep08</a>	<a href="#">361580</a>	10192	1229	3	279	general transcription factor II H, polypeptide 1 (Gtf2h1) alternative variant bSep08, mRNA.
<a href="#">Gtf2h2</a>	<a href="#">Gtf2h2.bSep08</a>	<a href="#">294693</a>	28144	2174	15	317	general transcription factor II H polypeptide 2 CRA b (35.5 kD) (Gtf2h2) alternative variant bSep08, complete mRNA.
<a href="#">Gtf2h2</a>	<a href="#">Gtf2h2.cSep08</a>	<a href="#">294693</a>	16984	1027	11	285	general transcription factor II H polypeptide 2 CRA b (32.3 kD) (Gtf2h2) alternative variant cSep08, mRNA.
<a href="#">Gtf2h2</a>	<a href="#">Gtf2h2.dSep08</a>	<a href="#">294693</a>	16804	881	11	191	general transcription factor II H polypeptide 2 CRA b (Gtf2h2) alternative variant dSep08, mRNA.
<a href="#">Gtf2h2</a>	<a href="#">Gtf2h2.eSep08</a>	<a href="#">294693</a>	8070	815	4	110	transcription factor 2 (12.4 kD) (Gtf2h2) alternative variant eSep08, mRNA.
<a href="#">Gtf2h2</a>	<a href="#">Gtf2h2.fSep08</a>	<a href="#">294693</a>	8448	515	5	76	general transcription factor IIH polypeptide 2B (Gtf2h2) alternative variant fSep08, mRNA.
<a href="#">Gtf2h3</a>	<a href="#">Gtf2h3.bSep08</a>	<a href="#">288651</a>	12364	616	9	139	general transcription factor IIH, polypeptide 3 (15.5 kD) (Gtf2h3) alternative variant bSep08, mRNA.
<a href="#">Gtf2h3</a>	<a href="#">Gtf2h3.cSep08</a>	<a href="#">288651</a>	1146	399	2	74	general transcription factor IIH, polypeptide 3 (Gtf2h3) alternative variant cSep08, mRNA.
<a href="#">Gtf2h4</a>	<a href="#">Gtf2h4.bSep08</a>	<a href="#">294236</a>	2971	790	4	221	general transcription factor II H, polypeptide 4 (Gtf2h4) alternative variant bSep08, mRNA.
<a href="#">Gtf2h4</a>	<a href="#">Gtf2h4.cSep08</a>	<a href="#">294236</a>	2695	764	2	210	general transcription factor II H, polypeptide 4 (Gtf2h4) alternative variant cSep08, mRNA.
<a href="#">Gtf2h5</a>	<a href="#">Gtf2h5.bSep08</a>	<a href="#">502227</a>	5857	780	3	63	general transcription factor IIH, polypeptide 5 (Gtf2h5) alternative variant bSep08, mRNA.
<a href="#">Gtf2h5</a>	<a href="#">Gtf2h5.cSep08</a>	<a href="#">502227</a>	5807	818	4	48	general transcription factor IIH, polypeptide 5 (Gtf2h5) alternative variant cSep08, mRNA.

<a href="#">Gtf2i</a>	<a href="#">Gtf2i.bSep08</a>	<a href="#">353256</a>	45453	1974	11	461	general transcription factor II I (Gtf2i) alternative variant bSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.cSep08</a>	<a href="#">353256</a>	52713	1783	12	400	general transcription factor II I (Gtf2i) alternative variant cSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.dSep08</a>	<a href="#">353256</a>	29540	1061	11	347	general transcription factor II I (Gtf2i) alternative variant dSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.eSep08</a>	<a href="#">353256</a>	22439	873	9	290	general transcription factor II I (Gtf2i) alternative variant eSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.fSep08</a>	<a href="#">353256</a>	3791	737	7	245	general transcription factor II I (Gtf2i) alternative variant fSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.gSep08</a>	<a href="#">353256</a>	4629	566	5	142	general transcription factor II I (Gtf2i) alternative variant gSep08, mRNA.
<a href="#">Gtf2i</a>	<a href="#">Gtf2i.iSep08</a>	<a href="#">353256</a>	1166	590	2	69	general transcription factor II I (Gtf2i) alternative variant iSep08, mRNA.
<a href="#">Gtf2ird1</a>	<a href="#">Gtf2ird1.bSep08</a>	<a href="#">246770</a>	24360	1283	11	371	general transcription factor II I repeat domain-containing 1 (Gtf2ird1) alternative variant bSep08, mRNA.
<a href="#">Gtf2ird1</a>	<a href="#">Gtf2ird1.cSep08</a>	<a href="#">246770</a>	22137	762	9	253	general transcription factor II I repeat domain-containing 1 (Gtf2ird1) alternative variant cSep08, mRNA.
<a href="#">Gtf2ird1</a>	<a href="#">Gtf2ird1.dSep08</a>	<a href="#">246770</a>	56501	540	5	180	general transcription factor II I repeat domain-containing 1 (Gtf2ird1) alternative variant dSep08, mRNA.
<a href="#">Gtf2ird1</a>	<a href="#">Gtf2ird1.eSep08</a>	<a href="#">246770</a>	7599	629	5	154	general transcription factor II I repeat domain-containing 1 (Gtf2ird1) alternative variant eSep08, mRNA.
<a href="#">Gtf3c1</a>	<a href="#">Gtf3c1.bSep08</a>	<a href="#">171063</a>	12947	741	6	195	general transcription factor III C 1 CRA b (22.6 kD) (Gtf3c1) alternative variant bSep08, mRNA.
<a href="#">Gtf3c1</a>	<a href="#">Gtf3c1.cSep08</a>	<a href="#">171063</a>	1479	688	2	133	general transcription factor III C 1 CRA c (Gtf3c1) alternative variant cSep08, mRNA.
<a href="#">Gtf3c1</a>	<a href="#">Gtf3c1.dSep08</a>	<a href="#">171063</a>	1304	388	3	119	general transcription factor III C 1 CRA b (Gtf3c1) alternative variant dSep08, mRNA.
<a href="#">Gtf3c1</a>	<a href="#">Gtf3c1.eSep08</a>	<a href="#">171063</a>	2802	473	3	103	general transcription factor III C 1 CRA b (Gtf3c1) alternative variant eSep08, mRNA.
<a href="#">Gtf3c1</a>	<a href="#">Gtf3c1.gSep08</a>	<a href="#">171063</a>	20913	770	3	74	general transcription factor 1 (Gtf3c1) alternative variant gSep08, mRNA.
<a href="#">Gtf3c2</a>	<a href="#">Gtf3c2.aSep08</a>	<a href="#">313914</a>	10934	2746	4	388	general transcription factor IIIC, polypeptide 2, beta (Gtf3c2) alternative variant aSep08, mRNA.
<a href="#">Gtf3c2</a>	<a href="#">Gtf3c2.bSep08</a>	<a href="#">313914</a>	1610	737	1	124	general transcription factor IIIC, polypeptide 2, beta (Gtf3c2) alternative variant bSep08, mRNA.
<a href="#">Gtf3c3</a>	<a href="#">Gtf3c3.bSep08</a>	<a href="#">316810</a>	1957	1314	2	74	general transcription factor IIIC, polypeptide 3 (Gtf3c3) alternative variant bSep08, mRNA.
<a href="#">Gtf3c5</a>	<a href="#">Gtf3c5.bSep08</a>	<a href="#">362095</a>	3947	1030	5	145	general transcription factor IIIC, polypeptide 5 (Gtf3c5) alternative variant bSep08, mRNA.
<a href="#">Gtf3c6andBxdc1</a>	<a href="#">Gtf3c6andBxdc1.cSep08</a>	<a href="#">294436</a>	9617	840	6	194	general transcription factor IIIC polypeptide 6 alpha (21.6 kD) (Gtf3c6andBxdc1) alternative variant cSep08, mRNA.
<a href="#">Gtf3c6andBxdc1</a>	<a href="#">Gtf3c6andBxdc1.cSep08</a>	<a href="#">361858</a>	9617	840	6	194	general transcription factor IIIC polypeptide 6 alpha (21.6 kD) (Gtf3c6andBxdc1) alternative variant cSep08, mRNA.
<a href="#">Gtf3c6andBxdc1</a>	<a href="#">Gtf3c6andBxdc1.dSep08</a>	<a href="#">294436</a>	8099	367	5	122	brix (Gtf3c6andBxdc1) alternative variant dSep08, mRNA.

<a href="#">Gtf3c6andBxdc1</a>	<a href="#">Gtf3c6andBxdc1.dSep08</a>	<a href="#">361858</a>	8099	367	5	122	brix (Gtf3c6andBxdc1) alternative variant dSep08, mRNA.
<a href="#">Gtl3</a>	<a href="#">Gtl3.aSep08</a>	<a href="#">307642</a>	13988	1194	6	193	gene trap locus 3 (22.7 kD) (Gtl3) alternative variant aSep08, complete mRNA.
<a href="#">Gtl3</a>	<a href="#">Gtl3.cSep08</a>	<a href="#">307642</a>	1576	1083	2	91	gene trap locus 3 (11.1 kD) (Gtl3) alternative variant cSep08, mRNA.
<a href="#">Gtlf3b</a>	<a href="#">Gtlf3b.aSep08</a>	<a href="#">363614</a>	7548	765	3	118	gene trap locus F3b (Gtlf3b) alternative variant aSep08, mRNA.
<a href="#">Gtlf3b</a>	<a href="#">Gtlf3b.cSep08</a>	<a href="#">363614</a>	10581	1374	3	65	gene trap locus F3b (7.1 kD) (Gtlf3b) alternative variant cSep08, mRNA.
<a href="#">Gtlf3b</a>	<a href="#">Gtlf3b.dSep08</a>	<a href="#">363614</a>	24773	549	3	48	gene trap locus F3b (5.5 kD) (Gtlf3b) alternative variant dSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.aSep08</a>	<a href="#">300077</a>	11457	3314	9	395	GTP binding protein 1 (Gtpbp1) alternative variant aSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.bSep08</a>	<a href="#">300077</a>	17803	1049	5	338	GTP binding protein 1 (Gtpbp1) alternative variant bSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.cSep08</a>	<a href="#">300077</a>	12737	1243	8	336	GTP binding protein 1 (Gtpbp1) alternative variant cSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.eSep08</a>	<a href="#">300077</a>	3204	776	3	258	GTP binding protein 1 (Gtpbp1) alternative variant eSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.fSep08</a>	<a href="#">300077</a>	5923	659	3	217	GTP binding protein 1 (Gtpbp1) alternative variant fSep08, mRNA.
<a href="#">Gtpbp1</a>	<a href="#">Gtpbp1.gSep08</a>	<a href="#">300077</a>	2634	1060	3	70	GTP binding protein 1 (7.6 kD) (Gtpbp1) alternative variant gSep08, mRNA.
<a href="#">Gtpbp2</a>	<a href="#">Gtpbp2.aSep08</a>	<a href="#">363195</a>	8350	3267	11	596	GTP binding protein 2 (Gtpbp2) alternative variant aSep08, mRNA.
<a href="#">Gtpbp2</a>	<a href="#">Gtpbp2.bSep08</a>	<a href="#">363195</a>	4813	748	6	249	GTP binding protein 2 (Gtpbp2) alternative variant bSep08, mRNA.
<a href="#">Gtpbp2</a>	<a href="#">Gtpbp2.cSep08</a>	<a href="#">363195</a>	2537	744	5	156	GTP binding protein 2 (Gtpbp2) alternative variant cSep08, mRNA.
<a href="#">Gtpbp2</a>	<a href="#">Gtpbp2.dSep08</a>	<a href="#">363195</a>	2000	1689	2	106	GTP binding protein 2 (Gtpbp2) alternative variant dSep08, mRNA.
<a href="#">Gtpbp2</a>	<a href="#">Gtpbp2.eSep08</a>	<a href="#">363195</a>	1491	1348	1	56	GTP binding protein 2 (Gtpbp2) alternative variant eSep08, mRNA.
<a href="#">Gtpbp3</a>	<a href="#">Gtpbp3.bSep08</a>	<a href="#">290633</a>	3104	759	7	202	GTP binding protein 3 (mitochondrial) (Gtpbp3) alternative variant bSep08, mRNA.
<a href="#">Gtpbp3</a>	<a href="#">Gtpbp3.cSep08</a>	<a href="#">290633</a>	850	744	2	40	GTP binding protein 3 (mitochondrial) (Gtpbp3) alternative variant cSep08, mRNA.
<a href="#">Gtpbp3</a>	<a href="#">Gtpbp3.dSep08</a>	<a href="#">290633</a>	742	367	2	31	GTP binding protein 3 (mitochondrial) (Gtpbp3) alternative variant dSep08, mRNA.
<a href="#">Gtpbp4</a>	<a href="#">Gtpbp4.aSep08</a>	<a href="#">114300</a>	7644	793	6	264	GTP binding protein 4 (Gtpbp4) alternative variant aSep08, mRNA.
<a href="#">Gtpbp4</a>	<a href="#">Gtpbp4.bSep08</a>	<a href="#">114300</a>	3309	404	1	40	GTP binding protein 4 (4.9 kD) (Gtpbp4) alternative variant bSep08, mRNA.
<a href="#">Gtpbp5</a>	<a href="#">Gtpbp5.bSep08</a>	<a href="#">296462</a>	12381	870	6	289	GTP binding protein 5 (Gtpbp5) alternative variant bSep08, mRNA.

<a href="#">Gtpbp5</a>	<a href="#">Gtpbp5.cSep08</a>	<a href="#">296462</a>	13986	797	6	219	GTP binding protein 5 (Gtpbp5) alternative variant cSep08, mRNA.
<a href="#">Gtpbp5</a>	<a href="#">Gtpbp5.dSep08</a>	<a href="#">296462</a>	13894	724	6	200	GTP binding protein 5 (Gtpbp5) alternative variant dSep08, mRNA.
<a href="#">Gtpbp5</a>	<a href="#">Gtpbp5.fSep08</a>	<a href="#">296462</a>	14527	1498	6	113	GTP binding protein 5 (12.3 kD) (Gtpbp5) alternative variant fSep08, mRNA.
<a href="#">Gtpbp5</a>	<a href="#">Gtpbp5.gSep08</a>	<a href="#">296462</a>	1426	488	2	62	GTP binding protein 5 (Gtpbp5) alternative variant gSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.aSep08</a>	<a href="#">363931</a>	3808	1510	10	492	GTP binding protein 6 like (Gtpbp6) alternative variant aSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.bSep08</a>	<a href="#">363931</a>	2198	665	4	205	GTP-binding protein 6 like (Gtpbp6) alternative variant bSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.cSep08</a>	<a href="#">363931</a>	2401	786	7	167	GTP binding protein 6 like (Gtpbp6) alternative variant cSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.dSep08</a>	<a href="#">363931</a>	844	579	3	154	putative protein (Gtpbp6) alternative variant dSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.eSep08</a>	<a href="#">363931</a>	2303	743	7	136	GTP binding protein 6 like (Gtpbp6) alternative variant eSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.fSep08</a>	<a href="#">363931</a>	2389	1262	5	131	GTP binding protein 6 like (Gtpbp6) alternative variant fSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.gSep08</a>	<a href="#">363931</a>	2294	764	6	107	GTP binding protein 6 like (Gtpbp6) alternative variant gSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.hSep08</a>	<a href="#">363931</a>	970	555	2	102	putative protein (Gtpbp6) alternative variant hSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.iSep08</a>	<a href="#">363931</a>	382	279	2	93	GTP binding protein 6 like (Gtpbp6) alternative variant iSep08, mRNA.
<a href="#">Gtpbp6</a>	<a href="#">Gtpbp6.jSep08</a>	<a href="#">363931</a>	1862	319	2	31	putative protein (3.3 kD) (Gtpbp6) alternative variant jSep08, mRNA.
<a href="#">Gtpbp8</a>	<a href="#">Gtpbp8.bSep08</a>	<a href="#">360714</a>	9791	1775	5	146	GTP-binding protein 8 (putative) (Gtpbp8) alternative variant bSep08, mRNA.
<a href="#">GTP_EFTU.1</a>	<a href="#">GTP_EFTU.1.aSep08</a>		9169	1270	1	423	eukaryotic translation initiation factor 5B (GTP_EFTU.1) alternative variant aSep08, mRNA.
<a href="#">GTP_EFTU.1</a>	<a href="#">GTP_EFTU.1.bSep08</a>		3895	670	1	222	eukaryotic translation initiation factor 5B (GTP_EFTU.1) alternative variant bSep08, mRNA.
<a href="#">GTP_EFTU.2</a>	<a href="#">GTP_EFTU.2.aSep08</a>		11803	542		180	mitochondrial elongation factor G2 (GTP_EFTU.2) mRNA.
<a href="#">Gtse1</a>	<a href="#">Gtse1.bSep08</a>	<a href="#">300126</a>	1723	1001	3	219	G two S phase expressed protein 1 (Gtse1) alternative variant bSep08, mRNA.
<a href="#">Gtsf1</a>	<a href="#">Gtsf1.aSep08</a>	<a href="#">315347</a>	18092	676	2	185	gametocyte specific factor 1 (Gtsf1) alternative variant aSep08, mRNA.
<a href="#">Guanylate_kin.0</a>	<a href="#">Guanylate_kin.0.aSep08</a>		48190	1112		235	membrane protein palmitoylated 7 CRA b (Guanylate_kin.0) mRNA.
<a href="#">Guanylate_kin.1</a>	<a href="#">Guanylate_kin.1.aSep08</a>		3421	1198		273	membrane protein palmitoylated 2 (Guanylate_kin.1) alternative variant aSep08, mRNA.
<a href="#">Guanylate_kin.1</a>	<a href="#">Guanylate_kin.1.bSep08</a>		3908	2464		105	membrane protein palmitoylated 2 (Guanylate_kin.1) alternative variant bSep08, mRNA.
<a href="#">guby</a>	<a href="#">guby.aSep08</a>		4851	779	2	48	putative protein (guby) alternative variant aSep08, mRNA.
<a href="#">guby</a>	<a href="#">guby.bSep08</a>		50583	885	5	75	putative protein (guby) alternative variant bSep08, mRNA.

<a href="#">Guca1a</a>	<a href="#">Guca1a.aSep08</a>	<a href="#">301233</a>	1596	471		122	guanylate cyclase activator 1a (retina) (Guca1a) mRNA.
<a href="#">guchy</a>	<a href="#">guchy.aSep08</a>		4521	299		37	putative protein (4.3 kD) (guchy) mRNA.
<a href="#">Gucy1b2</a>	<a href="#">Gucy1b2.bSep08</a>	<a href="#">25206</a>	37000	727	1	193	guanylate cyclase 1, soluble, beta 2 (Gucy1b2) alternative variant bSep08, mRNA.
<a href="#">Gucy1b3</a>	<a href="#">Gucy1b3.bSep08</a>	<a href="#">25202</a>	2081	450	3	135	guanylate cyclase 1, soluble, beta 3 (Gucy1b3) alternative variant bSep08, mRNA.
<a href="#">Gucy1b3</a>	<a href="#">Gucy1b3.cSep08</a>	<a href="#">25202</a>	16294	627	4	105	guanylate cyclase 1, soluble, beta 3 (12.1 kD) (Gucy1b3) alternative variant cSep08, mRNA.
<a href="#">Gucy1b3</a>	<a href="#">Gucy1b3.dSep08</a>	<a href="#">25202</a>	13535	633	3	96	guanylate cyclase 1, soluble, beta 3 (10.9 kD) (Gucy1b3) alternative variant dSep08, mRNA.
<a href="#">Gucy2c</a>	<a href="#">Gucy2c.aSep08</a>	<a href="#">25711</a>	15975	635		211	guanylate cyclase 2C (Gucy2c) mRNA.
<a href="#">Gucy2d</a>	<a href="#">Gucy2d.bSep08</a>	<a href="#">113911</a>	5811	679	1	167	guanylate cyclase 2d (Gucy2d) alternative variant bSep08, mRNA.
<a href="#">gufer</a>	<a href="#">gufer.aSep08</a>		7912	496		77	solute carrier family 17 member 1 (gufer) mRNA.
<a href="#">guflo</a>	<a href="#">guflo.aSep08</a>		2071	653		89	putative protein (10.6 kD) (guflo) mRNA.
<a href="#">guflu</a>	<a href="#">guflu.bSep08</a>		384	261	2	87	CRA a like (guflu) alternative variant bSep08, mRNA.
<a href="#">Guk1</a>	<a href="#">Guk1.aSep08</a>	<a href="#">303179</a>	2724	1108	5	226	guanylate kinase 1 (Guk1) alternative variant aSep08, mRNA.
<a href="#">Guk1</a>	<a href="#">Guk1.bSep08</a>	<a href="#">303179</a>	7715	670	7	208	guanylate kinase 1 (Guk1) alternative variant bSep08, mRNA.
<a href="#">Guk1</a>	<a href="#">Guk1.cSep08</a>	<a href="#">303179</a>	8046	423	4		
<a href="#">gukee</a>	<a href="#">gukee.aSep08</a>		1042	374		42	putative protein (gukee) mRNA.
<a href="#">Gulo</a>	<a href="#">Gulo.bSep08</a>	<a href="#">60671</a>	8975	696	5	190	gulonolactone (L-) oxidase (Gulo) alternative variant bSep08, mRNA.
<a href="#">Gulo</a>	<a href="#">Gulo.cSep08</a>	<a href="#">60671</a>	3842	1363	4	180	gulonolactone (L-) oxidase (Gulo) alternative variant cSep08, mRNA.
<a href="#">Gulo</a>	<a href="#">Gulo.dSep08</a>	<a href="#">60671</a>	1244	759	2	54	gulonolactone (L-) oxidase (6.6 kD) (Gulo) alternative variant dSep08, mRNA.
<a href="#">guloy</a>	<a href="#">guloy.aSep08</a>		11478	585		34	putative protein (3.8 kD) (guloy) mRNA.
<a href="#">gumee</a>	<a href="#">gumee.aSep08</a>		16654	762		56	putative protein (6.1 kD) (gumee) mRNA.
<a href="#">gumer</a>	<a href="#">gumer.aSep08</a>		3596	790		88	putative secreted or extracellular protein precursor (9.5 kD) (gumer) mRNA.
<a href="#">gunoy</a>	<a href="#">gunoy.aSep08</a>		6536	2260		289	eukaryotic translation initiation factor 5B CRA b (gunoy) mRNA.
<a href="#">gupor</a>	<a href="#">gupor.aSep08</a>		654	216		45	putative protein (gupor) mRNA.
<a href="#">gusa</a>	<a href="#">gusa.aSep08</a>		12417	321		11	putative protein (gusa) mRNA.
<a href="#">Gusb</a>	<a href="#">Gusb.bSep08</a>	<a href="#">24434</a>	3389	973	3	122	glucuronidase beta CRA a (14.4 kD) (Gusb) alternative variant bSep08, mRNA.
<a href="#">Gusb</a>	<a href="#">Gusb.cSep08</a>	<a href="#">24434</a>	5412	871	1	105	glucuronidase beta CRA a (Gusb) alternative variant cSep08, mRNA.
<a href="#">gushee</a>	<a href="#">gushee.aSep08</a>		9357	870		87	CRA b like (gushee) mRNA.
<a href="#">guto</a>	<a href="#">guto.aSep08</a>		3546	350		85	putative protein (guto) mRNA.
<a href="#">guvar</a>	<a href="#">guvar.aSep08</a>		935	361		88	putative protein of mammalian origin (guvar) mRNA.
<a href="#">guwey</a>	<a href="#">guwey.aSep08</a>		1582	414		29	putative protein (3.4 kD) (guwey) mRNA.
<a href="#">gyby</a>	<a href="#">gyby.aSep08</a>		4966	486		88	putative protein (9.6 kD) (gyby) mRNA.

<a href="#">gychy</a>	<a href="#">gychy.aSep08</a>		5167	608		202	grancalcin (gychy) mRNA.
<a href="#">GYF.0</a>	<a href="#">GYF.0.aSep08</a>		13483	738		245	trinucleotide repeat containing 15 CRA c (GYF.0) mRNA.
<a href="#">gyfer</a>	<a href="#">gyfer.aSep08</a>		557	437		37	putative protein (4.6 kD) (gyfer) mRNA.
<a href="#">gyflo</a>	<a href="#">gyflo.aSep08</a>		947	511		86	putative protein, with a transmembrane domain (gyflo) mRNA.
<a href="#">gyflu</a>	<a href="#">gyflu.aSep08</a>		2539	469		56	putative protein (6.4 kD) (gyflu) mRNA.
<a href="#">Gyg1</a>	<a href="#">Gyg1.bSep08</a>	<a href="#">81675</a>	41493	1155	8	270	glycogenin 1 (Gyg1) alternative variant bSep08, mRNA.
<a href="#">Gyg1</a>	<a href="#">Gyg1.cSep08</a>	<a href="#">81675</a>	42004	728	4	182	glycogenin 1 (Gyg1) alternative variant cSep08, mRNA.
<a href="#">Gyg1</a>	<a href="#">Gyg1.dSep08</a>	<a href="#">81675</a>	5893	606	3	140	glycogenin 1 (Gyg1) alternative variant dSep08, mRNA.
<a href="#">Gyk</a>	<a href="#">Gyk.bSep08</a>	<a href="#">79223</a>	11965	755	5	138	glycerol kinase (Gyk) alternative variant bSep08, mRNA.
<a href="#">Gyk</a>	<a href="#">Gyk.cSep08</a>	<a href="#">79223</a>	8208	293	4	84	glycerol kinase (Gyk) alternative variant cSep08, mRNA.
<a href="#">gykee</a>	<a href="#">gykee.aSep08</a>		5407	711		152	putative cytoplasmic protein of mammalian origin (16.5 kD) (gykee) mRNA.
<a href="#">gyloy</a>	<a href="#">gyloy.aSep08</a>		302	195		64	putative protein (gyloy) mRNA.
<a href="#">Gyltl1b</a>	<a href="#">Gyltl1b.aSep08</a>	<a href="#">311202</a>	6111	2381	15	691	glycosyltransferase-like 1B (79.4 kD) (Gyltl1b) alternative variant aSep08, mRNA.
<a href="#">Gyltl1b</a>	<a href="#">Gyltl1b.cSep08</a>	<a href="#">311202</a>	7902	926	7	229	glycosyltransferase-like 1B (Gyltl1b) alternative variant cSep08, mRNA.
<a href="#">gymee</a>	<a href="#">gymee.aSep08</a>		17094	762		56	putative protein (6.1 kD) (gymee) mRNA.
<a href="#">gymer</a>	<a href="#">gymer.aSep08</a>		18989	367		52	putative protein of vertebrate origin (gymer) mRNA.
<a href="#">gynoy</a>	<a href="#">gynoy.aSep08</a>		43188	590		74	putative protein (gynoy) mRNA.
<a href="#">gypor</a>	<a href="#">gypor.aSep08</a>		922	359		90	kinase 2 (gypor) mRNA.
<a href="#">Gys2</a>	<a href="#">Gys2.bSep08</a>	<a href="#">25623</a>	3742	750	1	123	glycogen synthase 2 (Gys2) alternative variant bSep08, mRNA.
<a href="#">Gys2</a>	<a href="#">Gys2.cSep08</a>	<a href="#">25623</a>	993	456	2	56	glycogen synthase 2 (6.3 kD) (Gys2) alternative variant cSep08, mRNA.
<a href="#">gysa</a>	<a href="#">gysa.aSep08</a>		140651	463		84	putative protein (9.3 kD) (gysa) mRNA.
<a href="#">gyshee</a>	<a href="#">gyshee.aSep08</a>		59851	733		54	putative protein (6.1 kD) (gyshee) mRNA.
<a href="#">gyto</a>	<a href="#">gyto.aSep08</a>		36879	636		212	CRA a 2 (gyto) mRNA.
<a href="#">gyvar</a>	<a href="#">gyvar.aSep08</a>		3126	803		69	putative protein (7.6 kD) (gyvar) mRNA.
<a href="#">gywey</a>	<a href="#">gywey.aSep08</a>		3222	199	2	63	gag protein like (gywey) alternative variant aSep08, mRNA.
<a href="#">Gzf1</a>	<a href="#">Gzf1.bSep08</a>	<a href="#">311508</a>	8302	2459		217	GDNF-inducible zinc finger protein 1 (24.4 kD) (Gzf1) alternative variant bSep08, mRNA.
<a href="#">Gzmb</a>	<a href="#">Gzmb.bSep08</a>	<a href="#">171528</a>	2099	740	4	246	granzyme B (Gzmb) alternative variant bSep08, mRNA.
<a href="#">Gzmn</a>	<a href="#">Gzmn.aSep08</a>	<a href="#">691668</a>	1709	401	2	97	granzyme N (Gzmn) alternative variant aSep08, mRNA.
<a href="#">Gzmn</a>	<a href="#">Gzmn.bSep08</a>	<a href="#">691668</a>	661	271	1	53	granzyme N (Gzmn) alternative variant bSep08, mRNA.
<a href="#">H2-T18</a>	<a href="#">H2-T18.bSep08</a>	<a href="#">406194</a>	8046	2465	7	268	histocompatibility 2, T region locus 18 (H2-T18) alternative variant bSep08, mRNA.
<a href="#">H2-T18</a>	<a href="#">H2-T18.cSep08</a>	<a href="#">406194</a>	3968	1437	5	231	histocompatibility 2, T region locus 18 (H2-T18) alternative variant cSep08, mRNA.
<a href="#">H2-T18</a>	<a href="#">H2-T18.eSep08</a>	<a href="#">406194</a>	610	475	2	121	histocompatibility 2, T region locus 18 (H2-T18) alternative variant eSep08, mRNA.
<a href="#">H2-T18</a>	<a href="#">H2-T18.fSep08</a>	<a href="#">406194</a>	3695	740	3	89	histocompatibility 2, T region locus 18 (H2-T18) alternative variant fSep08, mRNA.

<a href="#">H2-T18</a>	<a href="#">H2-T18.gSep08</a>	<a href="#">406194</a>	701	551	2	59	histocompatibility 2, T region locus 18 (H2-T18) alternative variant gSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.aSep08</a>	<a href="#">29384</a>	63678	1865	9	372	histone (39.6 kD) (H2afy) alternative variant aSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.cSep08</a>	<a href="#">29384</a>	56818	1356	8	341	histone H2A.1 (H2afy) alternative variant cSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.dSep08</a>	<a href="#">29384</a>	42322	742	6	228	histone (H2afy) alternative variant dSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.eSep08</a>	<a href="#">29384</a>	8822	508	4	141	histone (H2afy) alternative variant eSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.fSep08</a>	<a href="#">29384</a>	12147	1203	3	117	histone H2A.1 (H2afy) alternative variant fSep08, mRNA.
<a href="#">H2afy</a>	<a href="#">H2afy.gSep08</a>	<a href="#">29384</a>	567	290	2	49	putative protein (H2afy) alternative variant gSep08, mRNA.
<a href="#">H2afy2</a>	<a href="#">H2afy2.aSep08</a>	<a href="#">361844</a>	50489	2063	9	372	H2A histone family, member Y2 (40.1 kD) (H2afy2) alternative variant aSep08, mRNA.
<a href="#">H2afz.1</a>	<a href="#">H2afz.1.aSep08</a>	<a href="#">58940</a>	1017	386	3	128	H2A histone family, member Z (H2afz.1) alternative variant aSep08, mRNA.
<a href="#">H2afz.1</a>	<a href="#">H2afz.1.bSep08</a>	<a href="#">58940</a>	2299	997	5	128	H2A histone family, member Z (13.6 kD) (H2afz.1) alternative variant bSep08, mRNA.
<a href="#">H3f3b</a>	<a href="#">H3f3b.aSep08</a>	<a href="#">117056</a>	11588	1071	4	136	h3 histone, family 3B (15.3 kD) (H3f3b) alternative variant aSep08, mRNA.
<a href="#">H3f3b</a>	<a href="#">H3f3b.bSep08</a>	<a href="#">117056</a>	11084	416	3	116	h3 histone, family 3B (H3f3b) alternative variant bSep08, mRNA.
<a href="#">H3f3b.1</a>	<a href="#">H3f3b.1.bSep08</a>	<a href="#">117056</a>	2223	1739	3	125	h3 histone, family 3B (14.0 kD) (H3f3b.1) alternative variant bSep08, complete mRNA.
<a href="#">H13</a>	<a href="#">H13.aSep08</a>	<a href="#">311545</a>	35521	1973	13	419	histocompatibility 13 (H13) alternative variant aSep08, mRNA.
<a href="#">H13</a>	<a href="#">H13.bSep08</a>	<a href="#">311545</a>	16053	885	7	182	histocompatibility 13 CRA d (H13) alternative variant bSep08, mRNA.
<a href="#">H13</a>	<a href="#">H13.cSep08</a>	<a href="#">311545</a>	22116	443	5	114	histocompatibility 13 CRA e (H13) alternative variant cSep08, mRNA.
<a href="#">H13</a>	<a href="#">H13.dSep08</a>	<a href="#">311545</a>	3206	304	2	88	histocompatibility 13 CRA f (H13) alternative variant dSep08, mRNA.
<a href="#">H13</a>	<a href="#">H13.eSep08</a>	<a href="#">311545</a>	5944	382	4	60	histocompatibility 13 CRA c (H13) alternative variant eSep08, mRNA.
<a href="#">H28</a>	<a href="#">H28.aSep08</a>	<a href="#">310968</a>	31898	689	5	229	histocompatibility 28 (H28) alternative variant aSep08, mRNA.
<a href="#">H28</a>	<a href="#">H28.bSep08</a>	<a href="#">310968</a>	1766	596	2	198	histocompatibility 28 (H28) alternative variant bSep08, mRNA.
<a href="#">H28</a>	<a href="#">H28.cSep08</a>	<a href="#">310968</a>	29765	704	3	163	histocompatibility 28 (H28) alternative variant cSep08, mRNA.
<a href="#">HA2.0</a>	<a href="#">HA2.0.aSep08</a>		17589	1764		450	DEAH box polypeptide 29 (50.2 kD) (HA2.0) mRNA.
<a href="#">Habp2</a>	<a href="#">Habp2.bSep08</a>	<a href="#">292126</a>	35053	2049		521	hyaluronic acid binding protein 2 (57.7 kD) (Habp2) alternative variant bSep08, complete mRNA.
<a href="#">Hace1</a>	<a href="#">Hace1.bSep08</a>	<a href="#">361866</a>	12581	1776	1	513	putative protein (Hace1) alternative variant bSep08, mRNA.
<a href="#">Hac1andColq</a>	<a href="#">Hac1andColq.cSep08</a>	<a href="#">29755</a>	6305	534	5	123	2-hydroxyphytanoyl-CoA lyase (Hac1andColq) alternative variant cSep08, mRNA.
<a href="#">Hac1andColq</a>	<a href="#">Hac1andColq.cSep08</a>	<a href="#">85255</a>	6305	534	5	123	2-hydroxyphytanoyl-CoA lyase (Hac1andColq) alternative variant cSep08, mRNA.



<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.dSep08</a>	<a href="#">29755</a>	13275	693	5	118	2-hydroxyacyl-CoA lyase 1 (Hacl1andColq) alternative variant dSep08, mRNA.
<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.dSep08</a>	<a href="#">85255</a>	13275	693	5	118	2-hydroxyacyl-CoA lyase 1 (Hacl1andColq) alternative variant dSep08, mRNA.
<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.eSep08</a>	<a href="#">29755</a>	9873	697	5	96	2-hydroxyphytanoyl-CoA lyase (Hacl1andColq) alternative variant eSep08, mRNA.
<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.eSep08</a>	<a href="#">85255</a>	9873	697	5	96	2-hydroxyphytanoyl-CoA lyase (Hacl1andColq) alternative variant eSep08, mRNA.
<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.fSep08</a>	<a href="#">29755</a>	68106	1024	6	76	acetylcholinesterase collagen-like tail VIII (Hacl1andColq) alternative variant fSep08, mRNA.
<a href="#">Hacl1andColq</a>	<a href="#">Hacl1andColq.fSep08</a>	<a href="#">85255</a>	68106	1024	6	76	acetylcholinesterase collagen-like tail VIII (Hacl1andColq) alternative variant fSep08, mRNA.
<a href="#">Hadha</a>	<a href="#">Hadha.bSep08</a>	<a href="#">170670</a>	9960	1417	8	328	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (35.3 kD) (Hadha) alternative variant bSep08, mRNA.
<a href="#">Hadha</a>	<a href="#">Hadha.cSep08</a>	<a href="#">170670</a>	3512	783	4	223	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (Hadha) alternative variant cSep08, mRNA.
<a href="#">Hadha</a>	<a href="#">Hadha.dSep08</a>	<a href="#">170670</a>	5016	1006	3	83	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (Hadha) alternative variant dSep08, mRNA.
<a href="#">Hadha</a>	<a href="#">Hadha.eSep08</a>	<a href="#">170670</a>	1395	961	3	72	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (7.6 kD) (Hadha) alternative variant eSep08, mRNA.
<a href="#">Hadha</a>	<a href="#">Hadha.iSep08</a>	<a href="#">170670</a>	2826	408	3	46	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (Hadha) alternative variant iSep08, mRNA.
<a href="#">Hadhb</a>	<a href="#">Hadhb.bSep08</a>	<a href="#">171155</a>	20004	709	6	211	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (Hadhb) alternative variant bSep08, mRNA.
<a href="#">Hadhb</a>	<a href="#">Hadhb.cSep08</a>	<a href="#">171155</a>	22759	598	6	198	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (Hadhb) alternative variant cSep08, mRNA.
<a href="#">Hadhb</a>	<a href="#">Hadhb.dSep08</a>	<a href="#">171155</a>	17287	683	5	112	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (Hadhb) alternative variant dSep08, mRNA.

<a href="#">Hadhb</a>	<a href="#">Hadhb.eSep08</a>	<a href="#">171155</a>	11050	274	2	71	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (7.7 kD) (Hadhb) alternative variant eSep08, complete mRNA.
<a href="#">Hagh</a>	<a href="#">Hagh.bSep08</a>	<a href="#">24439</a>	4095	1245	3	102	hydroxyacyl glutathione hydrolase (11.8 kD) (Hagh) alternative variant bSep08, mRNA.
<a href="#">Haghl</a>	<a href="#">Haghl.bSep08</a>	<a href="#">302995</a>	2707	2015	5	132	hydroxyacylglutathione hydrolase-like (14.3 kD) (Haghl) alternative variant bSep08, complete mRNA.
<a href="#">Haghl</a>	<a href="#">Haghl.cSep08</a>	<a href="#">302995</a>	734	415	2	80	hydroxyacylglutathione hydrolase-like (Haghl) alternative variant cSep08, mRNA.
<a href="#">Haghl</a>	<a href="#">Haghl.dSep08</a>	<a href="#">302995</a>	1384	699	6	73	hydroxyacylglutathione hydrolase-like (Haghl) alternative variant dSep08, mRNA.
<a href="#">Hand2</a>	<a href="#">Hand2.aSep08</a>	<a href="#">64637</a>	2509	1113		112	heart and neural crest derivatives expressed transcript 2 (Hand2) mRNA.
<a href="#">Hao2</a>	<a href="#">Hao2.aSep08</a>	<a href="#">84029</a>	22586	1088	7	362	hydroxyacid oxidase 2 (long chain) (Hao2) alternative variant aSep08, mRNA.
<a href="#">Hao2</a>	<a href="#">Hao2.cSep08</a>	<a href="#">84029</a>	4678	838	5	251	hydroxyacid oxidase 2 (long chain) (Hao2) alternative variant cSep08, mRNA.
<a href="#">Hao2</a>	<a href="#">Hao2.dSep08</a>	<a href="#">84029</a>	4550	719	5	229	hydroxyacid oxidase 2 (long chain) (Hao2) alternative variant dSep08, mRNA.
<a href="#">Hao2</a>	<a href="#">Hao2.eSep08</a>	<a href="#">84029</a>	12657	715	5	216	hydroxyacid oxidase 2 (long chain) (Hao2) alternative variant eSep08, mRNA.
<a href="#">Hap1</a>	<a href="#">Hap1.cSep08</a>	<a href="#">29430</a>	880	594	2	108	huntingtin-associated protein 1 (12.2 kD) (Hap1) alternative variant cSep08, mRNA.
<a href="#">Hap1</a>	<a href="#">Hap1.dSep08</a>	<a href="#">29430</a>	814	437	2	86	huntingtin-associated protein 1 (Hap1) alternative variant dSep08, mRNA.
<a href="#">Hapln2</a>	<a href="#">Hapln2.aSep08</a>	<a href="#">64057</a>	5175	1430	4	453	hyaluronan and proteoglycan link protein 2 (Hapln2) alternative variant aSep08, mRNA.
<a href="#">Hapln2</a>	<a href="#">Hapln2.cSep08</a>	<a href="#">64057</a>	3783	590	1	101	hyaluronan and proteoglycan link protein 2 (10.6 kD) (Hapln2) alternative variant cSep08, mRNA.
<a href="#">Hapln2</a>	<a href="#">Hapln2.dSep08</a>	<a href="#">64057</a>	3660	434	1	90	hyaluronan and proteoglycan link protein 2 (9.6 kD) (Hapln2) alternative variant dSep08, mRNA.
<a href="#">Hapln4</a>	<a href="#">Hapln4.bSep08</a>	<a href="#">361129</a>	808	485	1	142	hyaluronan and proteoglycan link protein 4 (Hapln4) alternative variant bSep08, mRNA.
<a href="#">Harbi1l</a>	<a href="#">Harbi1l.aSep08</a>	<a href="#">362164</a>	35211	3742	16	479	harbinger transposase derived 1-like (52.6 kD) (Harbi1l) alternative variant aSep08, mRNA.
<a href="#">Harbi1l</a>	<a href="#">Harbi1l.bSep08</a>	<a href="#">362164</a>	6531	721	8	240	harbinger transposase derived 1-like (Harbi1l) alternative variant bSep08, mRNA.
<a href="#">Harbi1l</a>	<a href="#">Harbi1l.cSep08</a>	<a href="#">362164</a>	8472	502	6	133	harbinger transposase derived 1-like (Harbi1l) alternative variant cSep08, mRNA.
<a href="#">Harbi1l</a>	<a href="#">Harbi1l.dSep08</a>	<a href="#">362164</a>	2453	516	3	133	harbinger transposase derived 1-like (Harbi1l) alternative variant dSep08, mRNA.
<a href="#">HAT.0</a>	<a href="#">HAT.0.aSep08</a>		3989	718		209	crooked neck-like 1 (HAT.0) mRNA.
<a href="#">Hat1</a>	<a href="#">Hat1.bSep08</a>	<a href="#">296501</a>	40062	884	4	287	histone aminotransferase 1 (Hat1) alternative variant bSep08, mRNA.
<a href="#">Hat1</a>	<a href="#">Hat1.cSep08</a>	<a href="#">296501</a>	29121	721	2	204	histone aminotransferase 1 (23.8 kD) (Hat1) alternative variant cSep08, mRNA.

<a href="#">Havcr1</a>	<a href="#">Havcr1.bSep08</a>	<a href="#">286934</a>	32924	1997	9	307	hepatitis A virus cellular receptor 1 (34.0 kD) (Havcr1) alternative variant bSep08, mRNA.
<a href="#">Hba-a2</a>	<a href="#">Hba-a2.bSep08</a>	<a href="#">25632</a>	601	416	1	138	hemoglobin alpha, adult chain 2 (Hba-a2) alternative variant bSep08, mRNA.
<a href="#">Hbb</a>	<a href="#">Hbb.bSep08</a>	<a href="#">24440</a>	909	800	2	133	hemoglobin, beta (14.5 kD) (Hbb) alternative variant bSep08, mRNA.
<a href="#">Hbegf</a>	<a href="#">Hbegf.aSep08</a>	<a href="#">25433</a>	10929	2100	6	208	heparin-binding EGF-like growth factor (Hbegf) alternative variant aSep08, complete mRNA.
<a href="#">Hbg1</a>	<a href="#">Hbg1.bSep08</a>	<a href="#">94164</a>	1475	447	2	107	hemoglobin, gamma A (Hbg1) alternative variant bSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.bSep08</a>	<a href="#">293408</a>	35767	781	5	260	CRA d (Hbs1l) alternative variant bSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.cSep08</a>	<a href="#">293408</a>	35822	641	4	213	CRA d (Hbs1l) alternative variant cSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.dSep08</a>	<a href="#">293408</a>	35172	1043	10	162	CRA c (17.8 kD) (Hbs1l) alternative variant dSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.eSep08</a>	<a href="#">293408</a>	5020	662	3	156	CRA d (Hbs1l) alternative variant eSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.fSep08</a>	<a href="#">293408</a>	25567	427	4	108	CRA b (Hbs1l) alternative variant fSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.gSep08</a>	<a href="#">293408</a>	28387	507	3	92	CRA b like (Hbs1l) alternative variant gSep08, mRNA.
<a href="#">Hbs1l</a>	<a href="#">Hbs1l.iSep08</a>	<a href="#">293408</a>	2725	757	2	47	CRA c like (Hbs1l) alternative variant iSep08, mRNA.
<a href="#">Hbxip</a>	<a href="#">Hbxip.bSep08</a>	<a href="#">295357</a>	2766	1135	2	70	hepatitis B virus x interacting protein (7.5 kD) (Hbxip) alternative variant bSep08, mRNA.
<a href="#">Hcca2</a>	<a href="#">Hcca2.bSep08</a>	<a href="#">499288</a>	16875	688	4	229	HCCA2 protein (Hcca2) alternative variant bSep08, mRNA.
<a href="#">Hcca2</a>	<a href="#">Hcca2.cSep08</a>	<a href="#">499288</a>	27832	704	4	218	HCCA2 protein (Hcca2) alternative variant cSep08, mRNA.
<a href="#">Hcca2</a>	<a href="#">Hcca2.dSep08</a>	<a href="#">499288</a>	17188	1011	4	150	HCCA2 protein (17.0 kD) (Hcca2) alternative variant dSep08, mRNA.
<a href="#">Hcca2</a>	<a href="#">Hcca2.eSep08</a>	<a href="#">499288</a>	56899	1446	4	142	HCCA2 protein (16.5 kD) (Hcca2) alternative variant eSep08, complete mRNA.
<a href="#">Hcca2</a>	<a href="#">Hcca2.fSep08</a>	<a href="#">499288</a>	6215	387	1	67	HCCA2 protein (Hcca2) alternative variant fSep08, mRNA.
<a href="#">Hccs</a>	<a href="#">Hccs.aSep08</a>	<a href="#">317444</a>	9294	2145	4	184	holocytochrome c synthetase (22.0 kD) (Hccs) alternative variant aSep08, complete mRNA.
<a href="#">Hccs</a>	<a href="#">Hccs.bSep08</a>	<a href="#">317444</a>	3137	392	1	81	holocytochrome c synthetase (Hccs) alternative variant bSep08, mRNA.
<a href="#">Hcfc1</a>	<a href="#">Hcfc1.aSep08</a>	<a href="#">363519</a>	4160	2425	5	240	host cell factor C1 (Hcfc1) mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.aSep08</a>	<a href="#">287097</a>	2219	1269	4	157	host cell factor C1 regulator 1 (XPO1-dependent) (Hcfc1r1) alternative variant aSep08, mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.bSep08</a>	<a href="#">287097</a>	1668	979	3	119	host cell factor C1 regulator 1 (XPO1-dependent) (13.5 kD) (Hcfc1r1) alternative variant bSep08, mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.cSep08</a>	<a href="#">287097</a>	1377	560	4	119	host cell factor C1 regulator 1 (XPO1-dependent) (13.5 kD) (Hcfc1r1) alternative variant cSep08, mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.dSep08</a>	<a href="#">287097</a>	855	333	3	111	host cell factor C1 regulator 1 (XPO1-dependent) (Hcfc1r1) alternative variant dSep08, mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.eSep08</a>	<a href="#">287097</a>	1244	376	3	93	host cell factor C1 regulator 1 (XPO1-dependent) (Hcfc1r1) alternative variant eSep08, mRNA.
<a href="#">Hcfc1r1</a>	<a href="#">Hcfc1r1.fSep08</a>	<a href="#">287097</a>	1305	487	2	76	host cell factor C1 regulator 1 (XPO1-dependent) (Hcfc1r1) alternative variant fSep08, complete mRNA.
<a href="#">Hcls1</a>	<a href="#">Hcls1.bSep08</a>	<a href="#">288077</a>	19184	696	1	171	hematopoietic cell specific Lyn substrate 1 (Hcls1) alternative variant bSep08, mRNA.

<a href="#">Hcn3</a>	<a href="#">Hcn3.bSep08</a>	<a href="#">114245</a>	8008	744	4	248	hyperpolarization-activated cyclic nucleotide-gated potassium channel 3 (Hcn3) alternative variant bSep08, mRNA.
<a href="#">Hcrt</a>	<a href="#">Hcrt.aSep08</a>	<a href="#">25723</a>	1833	506	2	143	hypocretin (Hcrt) alternative variant aSep08, mRNA.
<a href="#">Hcst</a>	<a href="#">Hcst.cSep08</a>	<a href="#">474146</a>	2045	431	4	78	hematopoietic cell signal transducer (8.1 kD) (Hcst) alternative variant cSep08, mRNA.
<a href="#">HD.0</a>	<a href="#">HD.0.aSep08</a>		10058	579		193	SAM domain HD 1 (HD.0) mRNA.
<a href="#">Hdac1_predicted</a>	<a href="#">Hdac1_predicted.bSep08</a>	<a href="#">297893</a>	1413	746	3	118	histone deacetylase 1 (predicted) (Hdac1_predicted) alternative variant bSep08, mRNA.
<a href="#">Hdac2</a>	<a href="#">Hdac2.aSep08</a>	<a href="#">84577</a>	24400	2178	14	554	histone deacetylase 2 (Hdac2) alternative variant aSep08, mRNA.
<a href="#">Hdac2</a>	<a href="#">Hdac2.bSep08</a>	<a href="#">84577</a>	8659	698	3	58	histone deacetylase 2 (Hdac2) alternative variant bSep08, mRNA.
<a href="#">Hdac3</a>	<a href="#">Hdac3.aSep08</a>	<a href="#">84578</a>	9420	1618	12	334	histone deacetylase 3 (Hdac3) alternative variant aSep08, mRNA.
<a href="#">Hdac3</a>	<a href="#">Hdac3.bSep08</a>	<a href="#">84578</a>	5781	1813	10	294	histone deacetylase 3 (Hdac3) alternative variant bSep08, mRNA.
<a href="#">Hdac3</a>	<a href="#">Hdac3.cSep08</a>	<a href="#">84578</a>	11677	875	9	272	histone deacetylase 3 (Hdac3) alternative variant cSep08, mRNA.
<a href="#">Hdac3</a>	<a href="#">Hdac3.dSep08</a>	<a href="#">84578</a>	7134	871	7	198	histone deacetylase 3 (Hdac3) alternative variant dSep08, mRNA.
<a href="#">Hdac4</a>	<a href="#">Hdac4.aSep08</a>	<a href="#">363287</a>	46325	1905	5	634	histone deacetylase 4 (Hdac4) alternative variant aSep08, mRNA.
<a href="#">Hdac6</a>	<a href="#">Hdac6.aSep08</a>	<a href="#">84581</a>	5945	1849		500	histone deacetylase 6 (Hdac6) alternative variant aSep08, mRNA.
<a href="#">Hdac6</a>	<a href="#">Hdac6.bSep08</a>	<a href="#">84581</a>	9205	1789		486	histone deacetylase 6 (Hdac6) alternative variant bSep08, mRNA.
<a href="#">Hdac6</a>	<a href="#">Hdac6.cSep08</a>	<a href="#">84581</a>	8683	1122		374	histone deacetylase 6 (Hdac6) alternative variant cSep08, mRNA.
<a href="#">Hdac7a</a>	<a href="#">Hdac7a.aSep08</a>	<a href="#">84582</a>	15801	2919	17	640	histone deacetylase 7A (Hdac7a) alternative variant aSep08, mRNA.
<a href="#">Hdac7a</a>	<a href="#">Hdac7a.bSep08</a>	<a href="#">84582</a>	3356	879	4	246	histone deacetylase 7A (Hdac7a) alternative variant bSep08, mRNA.
<a href="#">Hdac7a</a>	<a href="#">Hdac7a.cSep08</a>	<a href="#">84582</a>	1556	425	3	141	histone deacetylase 7A (Hdac7a) alternative variant cSep08, mRNA.
<a href="#">Hdac7a</a>	<a href="#">Hdac7a.dSep08</a>	<a href="#">84582</a>	1001	740	2	107	histone deacetylase 7A (11.2 kD) (Hdac7a) alternative variant dSep08, mRNA.
<a href="#">Hdac8</a>	<a href="#">Hdac8.aSep08</a>	<a href="#">363481</a>	208503	1964	1	377	histone deacetylase 8 (41.8 kD) (Hdac8) alternative variant aSep08, mRNA.
<a href="#">Hdac10</a>	<a href="#">Hdac10.aSep08</a>	<a href="#">362981</a>	3677	1658	12	430	histone deacetylase 10 (Hdac10) alternative variant aSep08, mRNA.
<a href="#">Hdac10</a>	<a href="#">Hdac10.bSep08</a>	<a href="#">362981</a>	1845	747	7	248	histone deacetylase 10 (Hdac10) alternative variant bSep08, mRNA.
<a href="#">Hdac10</a>	<a href="#">Hdac10.cSep08</a>	<a href="#">362981</a>	1001	719	2	171	histone deacetylase 10 (Hdac10) alternative variant cSep08, mRNA.
<a href="#">Hdac10</a>	<a href="#">Hdac10.dSep08</a>	<a href="#">362981</a>	1014	648	3	166	histone deacetylase 10 (Hdac10) alternative variant dSep08, mRNA.

<a href="#">Hdac10</a>	<a href="#">Hdac10.eSep08</a>	<a href="#">362981</a>	1167	587	5	132	histone deacetylase 10 (Hdac10) alternative variant eSep08, mRNA.
<a href="#">Hdac11</a>	<a href="#">Hdac11.bSep08</a>	<a href="#">297453</a>	3954	470	3	138	histone deacetylase 11 CRA a (Hdac11) alternative variant bSep08, mRNA.
<a href="#">Hdac11</a>	<a href="#">Hdac11.cSep08</a>	<a href="#">297453</a>	7069	1407	3	94	putative protein (10.0 kD) (Hdac11) alternative variant cSep08, mRNA.
<a href="#">Hdac11</a>	<a href="#">Hdac11.dSep08</a>	<a href="#">297453</a>	2919	481	3	57	putative protein (Hdac11) alternative variant dSep08, mRNA.
<a href="#">Hdac11</a>	<a href="#">Hdac11.eSep08</a>	<a href="#">297453</a>	2323	729	2	31	histone deacetylase 11 (Hdac11) alternative variant eSep08, mRNA.
<a href="#">Hdc</a>	<a href="#">Hdc.bSep08</a>	<a href="#">24443</a>	8470	1252	5	326	histidine decarboxylase (Hdc) alternative variant bSep08, mRNA.
<a href="#">Hdc</a>	<a href="#">Hdc.cSep08</a>	<a href="#">24443</a>	3275	904	1	63	histidine decarboxylase (Hdc) alternative variant cSep08, mRNA.
<a href="#">Hddc2</a>	<a href="#">Hddc2.bSep08</a>	<a href="#">361462</a>	19446	910	6	155	metal-dependent phosphohydrolase, HD region, subdomain containing protein (18.0 kD) (Hddc2) alternative variant bSep08, mRNA.
<a href="#">Hddc3</a>	<a href="#">Hddc3.bSep08</a>	<a href="#">308758</a>	1391	825	4	144	metal-dependent phosphohydrolase, HD region, subdomain containing protein (16.4 kD) (Hddc3) alternative variant bSep08, mRNA.
<a href="#">Hddc3</a>	<a href="#">Hddc3.cSep08</a>	<a href="#">308758</a>	1261	810	3	129	putative protein of ancient origin (14.8 kD) (Hddc3) alternative variant cSep08, mRNA.
<a href="#">Hdgf</a>	<a href="#">Hdgf.bSep08</a>	<a href="#">114499</a>	10344	967	5	229	hepatoma-derived growth factor (Hdgf) alternative variant bSep08, mRNA.
<a href="#">Hdgf</a>	<a href="#">Hdgf.cSep08</a>	<a href="#">114499</a>	8030	800	5	213	hepatoma-derived growth factor (Hdgf) alternative variant cSep08, mRNA.
<a href="#">Hdgfrp2</a>	<a href="#">Hdgfrp2.bSep08</a>	<a href="#">171073</a>	16872	1502	9	450	hepatoma-derived growth factor, related protein 2 (Hdgfrp2) alternative variant bSep08, mRNA.
<a href="#">Hdgfrp2</a>	<a href="#">Hdgfrp2.cSep08</a>	<a href="#">171073</a>	1430	695	4	181	hepatoma-derived growth factor, related protein 2 (Hdgfrp2) alternative variant cSep08, mRNA.
<a href="#">Hdgfrp2</a>	<a href="#">Hdgfrp2.dSep08</a>	<a href="#">171073</a>	1600	1357	2	103	hepatoma-derived growth factor, related protein 2 (11.0 kD) (Hdgfrp2) alternative variant dSep08, mRNA.
<a href="#">Hdgfrp2</a>	<a href="#">Hdgfrp2.eSep08</a>	<a href="#">171073</a>	2357	708	2	63	hepatoma-derived growth factor, related protein 2 (Hdgfrp2) alternative variant eSep08, mRNA.
<a href="#">Hdhd2</a>	<a href="#">Hdhd2.bSep08</a>	<a href="#">361351</a>	23086	889	7	259	haloacid dehalogenase-like hydrolase (28.8 kD) (Hdhd2) alternative variant bSep08, mRNA.
<a href="#">Hdhd2</a>	<a href="#">Hdhd2.cSep08</a>	<a href="#">361351</a>	39868	1532	7	259	haloacid dehalogenase-like hydrolase (28.8 kD) (Hdhd2) alternative variant cSep08, complete mRNA.
<a href="#">Hdhd2</a>	<a href="#">Hdhd2.dSep08</a>	<a href="#">361351</a>	11119	1405	3	82	immediate early response protein 1 (9.0 kD) (Hdhd2) alternative variant dSep08, complete mRNA.
<a href="#">Hdhd2</a>	<a href="#">Hdhd2.eSep08</a>	<a href="#">361351</a>	53147	1799	4	40	immediate early response protein 1 (4.6 kD) (Hdhd2) alternative variant eSep08, mRNA.
<a href="#">Hdlbp</a>	<a href="#">Hdlbp.bSep08</a>	<a href="#">64474</a>	35892	2176	10	407	high density lipoprotein binding protein CRA f like (Hdlbp) alternative variant bSep08, mRNA.
<a href="#">Hdlbp</a>	<a href="#">Hdlbp.cSep08</a>	<a href="#">64474</a>	16251	1882	8	373	high density lipoprotein binding protein CRA b like (41.3 kD) (Hdlbp) alternative variant cSep08, mRNA.

<a href="#">Hdlbp</a>	<a href="#">Hdlbp.dSep08</a>	<a href="#">64474</a>	5885	3080	5	214	high density lipoprotein binding protein CRA e like (23.8 kD) (Hdlbp) alternative variant dSep08, mRNA.
<a href="#">Hdlbp</a>	<a href="#">Hdlbp.eSep08</a>	<a href="#">64474</a>	1112	733	3	112	high density lipoprotein binding protein CRA e like (Hdlbp) alternative variant eSep08, mRNA.
<a href="#">Hdlbp</a>	<a href="#">Hdlbp.fSep08</a>	<a href="#">64474</a>	1115	978	2	58	high density lipoprotein binding protein CRA e like (6.3 kD) (Hdlbp) alternative variant fSep08, mRNA.
<a href="#">Hdmcp</a>	<a href="#">Hdmcp.bSep08</a>	<a href="#">299316</a>	2448	1593	1	131	mitochondrial hepatocellular carcinoma-downregulated carrier protein (Hdmcp) alternative variant bSep08, mRNA.
<a href="#">HEAT.0</a>	<a href="#">HEAT.0.aSep08</a>		30892	3561		369	CLIP-associating protein 1 (HEAT.0) mRNA.
<a href="#">HEAT.1</a>	<a href="#">HEAT.1.aSep08</a>		2573	1267	8	330	CRA b (36.5 kD) (HEAT.1) alternative variant aSep08, mRNA.
<a href="#">HEAT.1</a>	<a href="#">HEAT.1.bSep08</a>		1112	600	4	118	CRA b (HEAT.1) alternative variant bSep08, mRNA.
<a href="#">HEAT.1</a>	<a href="#">HEAT.1.cSep08</a>		1251	738	4	84	CRA a like (9.5 kD) (HEAT.1) alternative variant cSep08, mRNA.
<a href="#">HEAT.1</a>	<a href="#">HEAT.1.dSep08</a>		714	641	2	34	CRA b like (HEAT.1) alternative variant dSep08, mRNA.
<a href="#">Heatr1</a>	<a href="#">Heatr1.bSep08</a>	<a href="#">361262</a>	3509	781	2	215	HEAT repeat containing 1 (Heatr1) alternative variant bSep08, mRNA.
<a href="#">Heatr1</a>	<a href="#">Heatr1.cSep08</a>	<a href="#">361262</a>	3742	714	2	165	HEAT repeat containing 1 (Heatr1) alternative variant cSep08, mRNA.
<a href="#">Heatr2</a>	<a href="#">Heatr2.bSep08</a>	<a href="#">304332</a>	32653	1814	2	413	HEAT repeat containing 2 (Heatr2) alternative variant bSep08, mRNA.
<a href="#">Heatr3</a>	<a href="#">Heatr3.aSep08</a>	<a href="#">361375</a>	15612	1933		244	HEAT repeat containing 3 (Heatr3) mRNA.
<a href="#">Heatr5a</a>	<a href="#">Heatr5a.aSep08</a>	<a href="#">362737</a>	16717	3037		539	HEAT repeat containing 5A (Heatr5a) mRNA.
<a href="#">Heatr5b</a>	<a href="#">Heatr5b.aSep08</a>	<a href="#">362683</a>	49310	3946	16	1127	HEAT repeat containing 5B (Heatr5b) alternative variant aSep08, mRNA.
<a href="#">Heatr5b</a>	<a href="#">Heatr5b.bSep08</a>	<a href="#">362683</a>	10222	1895	6	443	HEAT repeat containing 5B (Heatr5b) alternative variant bSep08, mRNA.
<a href="#">Heatr5b</a>	<a href="#">Heatr5b.cSep08</a>	<a href="#">362683</a>	7196	723	5	240	HEAT repeat containing 5B (Heatr5b) alternative variant cSep08, mRNA.
<a href="#">Heatr5b</a>	<a href="#">Heatr5b.dSep08</a>	<a href="#">362683</a>	2809	389	3	129	HEAT repeat containing 5B (Heatr5b) alternative variant dSep08, mRNA.
<a href="#">Heatr6</a>	<a href="#">Heatr6.bSep08</a>	<a href="#">497972</a>	25053	1784	7	493	HEAT repeat containing 6 (Heatr6) alternative variant bSep08, mRNA.
<a href="#">Heatr6</a>	<a href="#">Heatr6.cSep08</a>	<a href="#">497972</a>	3561	1323	3	272	HEAT repeat containing 6 (29.7 kD) (Heatr6) alternative variant cSep08, mRNA.
<a href="#">Hebp1</a>	<a href="#">Hebp1.aSep08</a>	<a href="#">362454</a>	36605	1050	2	225	heme binding protein 1 (Hebp1) alternative variant aSep08, mRNA.
<a href="#">HECT.0</a>	<a href="#">HECT.0.aSep08</a>		10563	1934	2	264	HECT (HECT.0) alternative variant aSep08, mRNA.
<a href="#">HECT.0</a>	<a href="#">HECT.0.bSep08</a>		3655	777	1	164	HECT (HECT.0) alternative variant bSep08, mRNA.
<a href="#">Hectd1</a>	<a href="#">Hectd1.aSep08</a>	<a href="#">362736</a>	46195	6390	30	1864	sad1/UNC-like, C-terminal and mib herc2 and HECT (Hectd1) alternative variant aSep08, mRNA.
<a href="#">Hectd1</a>	<a href="#">Hectd1.bSep08</a>	<a href="#">362736</a>	32137	1819	8	606	ankyrin (Hectd1) alternative variant bSep08, mRNA.
<a href="#">Hectd1</a>	<a href="#">Hectd1.cSep08</a>	<a href="#">362736</a>	62160	1788	4	469	putative protein of eukaryotic origin (Hectd1) alternative variant cSep08, mRNA.
<a href="#">Hectd1</a>	<a href="#">Hectd1.dSep08</a>	<a href="#">362736</a>	5232	1178	6	260	ankyrin (Hectd1) alternative variant dSep08, mRNA.
<a href="#">Hectd1</a>	<a href="#">Hectd1.eSep08</a>	<a href="#">362736</a>	6164	572	4	150	ankyrin (Hectd1) alternative variant eSep08, mRNA.

<a href="#">Hectd2</a>	<a href="#">Hectd2.bSep08</a>	<a href="#">309514</a>	45221	753	3	240	putative protein of metazoan origin (Hectd2) alternative variant bSep08, mRNA.
<a href="#">Hectd2</a>	<a href="#">Hectd2.cSep08</a>	<a href="#">309514</a>	45884	743	3	182	putative protein of metazoan origin (Hectd2) alternative variant cSep08, mRNA.
<a href="#">Hel308</a>	<a href="#">Hel308.aSep08</a>	<a href="#">360912</a>	17539	1038		346	DNA helicase HEL308 (Hel308) mRNA.
<a href="#">Helb</a>	<a href="#">Helb.aSep08</a>	<a href="#">500837</a>	5067	382		126	helicase (DNA) B (Helb) mRNA.
<a href="#">Helicase_C.0</a>	<a href="#">Helicase_C.0.aSep08</a>		6421	401		133	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 1 (Helicase_C.0) mRNA.
<a href="#">Helicase_C.1</a>	<a href="#">Helicase_C.1.aSep08</a>		2474	368		122	chromodomain helicase binding protein 9 like (Helicase_C.1) mRNA.
<a href="#">Helicase_C.2</a>	<a href="#">Helicase_C.2.aSep08</a>		3552	748		151	responsive protein zinc zd10 (Helicase_C.2) alternative variant aSep08, mRNA.
<a href="#">Helicase_C.3</a>	<a href="#">Helicase_C.3.aSep08</a>		19398	2042		498	dead box polypeptide 46 (Helicase_C.3) mRNA.
<a href="#">Helicase_C.4</a>	<a href="#">Helicase_C.4.aSep08</a>		23611	713		237	atp-dependent rna helicase tdrd9 (Helicase_C.4) mRNA.
<a href="#">Helicase_C.5</a>	<a href="#">Helicase_C.5.aSep08</a>		2996	522		173	CBP activator (Helicase_C.5) mRNA.
<a href="#">HELP.0</a>	<a href="#">HELP.0.aSep08</a>		12185	3924	22	896	echinoderm microtubule associated protein like 3 (95.6 kD) (HELP.0) alternative variant aSep08, mRNA.
<a href="#">HELP.0</a>	<a href="#">HELP.0.bSep08</a>		3797	1244	6	337	echinoderm microtubule associated protein like 3 (HELP.0) alternative variant bSep08, mRNA.
<a href="#">HELP.0</a>	<a href="#">HELP.0.cSep08</a>		3742	1244	7	179	echinoderm microtubule associated protein like 3 CRA e (19.7 kD) (HELP.0) alternative variant cSep08, mRNA.
<a href="#">HELP.0</a>	<a href="#">HELP.0.dSep08</a>		867	707	3	179	echinoderm microtubule associated protein like 3 (HELP.0) alternative variant dSep08, mRNA.
<a href="#">HELP.0</a>	<a href="#">HELP.0.eSep08</a>		1587	414	4	66	echinoderm microtubule associated protein like 3 CRA f (HELP.0) alternative variant eSep08, mRNA.
<a href="#">Helz</a>	<a href="#">Helz.bSep08</a>	<a href="#">287773</a>	11562	730	3	243	helicase with zinc finger domain (Helz) alternative variant bSep08, mRNA.
<a href="#">Helz</a>	<a href="#">Helz.cSep08</a>	<a href="#">287773</a>	13358	1247	3	107	helicase with zinc finger domain (Helz) alternative variant cSep08, mRNA.
<a href="#">Helz</a>	<a href="#">Helz.eSep08</a>	<a href="#">287773</a>	29057	406	5	80	helicase with zinc finger domain (Helz) alternative variant eSep08, mRNA.
<a href="#">Heme_oxygenase_e.0</a>	<a href="#">Heme_oxygenase.0.aSep08</a>		2169	1403		232	heme (Heme_oxygenase.0) mRNA.
<a href="#">Hemgn</a>	<a href="#">Hemgn.bSep08</a>	<a href="#">113882</a>	15985	1341	2	385	hemogen (Hemgn) alternative variant bSep08, mRNA.
<a href="#">Hemk1</a>	<a href="#">Hemk1.bSep08</a>	<a href="#">300989</a>	19362	797	1	265	HemK methyltransferase family member 1 (Hemk1) alternative variant bSep08, mRNA.
<a href="#">Heph</a>	<a href="#">Heph.bSep08</a>	<a href="#">117240</a>	12029	756	1	74	hephaestin (Heph) alternative variant bSep08, mRNA.
<a href="#">Herc1</a>	<a href="#">Herc1.aSep08</a>	<a href="#">315771</a>	19424	2391	11	645	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 (Herc1) alternative variant aSep08, mRNA.
<a href="#">Herc1</a>	<a href="#">Herc1.bSep08</a>	<a href="#">315771</a>	1031	541	1	153	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 (Herc1) alternative variant bSep08, mRNA.
<a href="#">Herc1</a>	<a href="#">Herc1.cSep08</a>	<a href="#">315771</a>	6721	684	2		

<a href="#">Herc2</a>	<a href="#">Herc2.bSep08</a>	<a href="#">308669</a>	697	412	2	136	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 2 (Herc2) alternative variant bSep08, mRNA.
<a href="#">Herc2</a>	<a href="#">Herc2.dSep08</a>	<a href="#">308669</a>	2540	401	4	67	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 2 (Herc2) alternative variant dSep08, mRNA.
<a href="#">Herc3</a>	<a href="#">Herc3.cSep08</a>	<a href="#">362377</a>	12345	716	4	97	hect domain and RLD 3 (Herc3) alternative variant cSep08, mRNA.
<a href="#">Herc3</a>	<a href="#">Herc3.eSep08</a>	<a href="#">362377</a>	4812	376	3	39	hect domain and RLD 3 (Herc3) alternative variant eSep08, mRNA.
<a href="#">Herc4</a>	<a href="#">Herc4.bSep08</a>	<a href="#">309758</a>	32029	2067	12	511	hect domain and RLD 4 (Herc4) alternative variant bSep08, mRNA.
<a href="#">Herc4</a>	<a href="#">Herc4.cSep08</a>	<a href="#">309758</a>	8952	505	4	164	hect domain and RLD 4 (Herc4) alternative variant cSep08, mRNA.
<a href="#">Herc6</a>	<a href="#">Herc6.aSep08</a>	<a href="#">362376</a>	3883	1792	5	211	potential ubiquitin ligase (Herc6) alternative variant aSep08, mRNA.
<a href="#">Herpud1</a>	<a href="#">Herpud1.bSep08</a>	<a href="#">85430</a>	6706	749	6	222	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1 (Herpud1) alternative variant bSep08, mRNA.
<a href="#">Herpud1</a>	<a href="#">Herpud1.cSep08</a>	<a href="#">85430</a>	5460	751	6	177	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1 (Herpud1) alternative variant cSep08, mRNA.
<a href="#">Herpud1</a>	<a href="#">Herpud1.eSep08</a>	<a href="#">85430</a>	1492	409	2	32	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1 (3.7 kD) (Herpud1) alternative variant eSep08, mRNA.
<a href="#">Herpud2</a>	<a href="#">Herpud2.aSep08</a>	<a href="#">300463</a>	23369	1234	4	202	HERPUD family member 2 (Herpud2) alternative variant aSep08, mRNA.
<a href="#">Herpud2</a>	<a href="#">Herpud2.bSep08</a>	<a href="#">300463</a>	2175	1037	3	195	HERPUD family member 2 (Herpud2) alternative variant bSep08, mRNA.
<a href="#">Herpud2</a>	<a href="#">Herpud2.dSep08</a>	<a href="#">300463</a>	2256	1089	3	111	HERPUD family member 2 (Herpud2) alternative variant dSep08, mRNA.
<a href="#">Herpud2</a>	<a href="#">Herpud2.eSep08</a>	<a href="#">300463</a>	2431	1973	2	54	member 2 (5.8 kD) (Herpud2) alternative variant eSep08, mRNA.
<a href="#">Hes6</a>	<a href="#">Hes6.bSep08</a>	<a href="#">316626</a>	1719	1325	4	224	hairy and enhancer of split 6 (Drosophila) (24.6 kD) (Hes6) alternative variant bSep08, complete mRNA.
<a href="#">Hes6</a>	<a href="#">Hes6.cSep08</a>	<a href="#">316626</a>	1214	995	3	221	hairy and enhancer of split 6 (Drosophila) (Hes6) alternative variant cSep08, mRNA.
<a href="#">Hes6</a>	<a href="#">Hes6.dSep08</a>	<a href="#">316626</a>	1206	1050	3	126	hairy and enhancer of split 6 (Drosophila) (14.0 kD) (Hes6) alternative variant dSep08, mRNA.
<a href="#">Hexa</a>	<a href="#">Hexa.bSep08</a>	<a href="#">300757</a>	5849	548	4	125	hexosaminidase A (Hexa) alternative variant bSep08, mRNA.
<a href="#">Hexb</a>	<a href="#">Hexb.cSep08</a>	<a href="#">294673</a>	4246	750	6	78	hexosaminidase B (Hexb) alternative variant cSep08, mRNA.
<a href="#">Hexb</a>	<a href="#">Hexb.dSep08</a>	<a href="#">294673</a>	1023	617	2	53	hexosaminidase B (Hexb) alternative variant dSep08, mRNA.



<a href="#">Hexim2</a>	<a href="#">Hexim2.aSep08</a>	<a href="#">303580</a>	4426	1289	1	313	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (35.5 kD) (Hexim2) alternative variant aSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.aSep08</a>	<a href="#">685659</a>	4426	1289	1	313	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (35.5 kD) (Hexim2) alternative variant aSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.cSep08</a>	<a href="#">303580</a>	4689	862	2	262	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant cSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.cSep08</a>	<a href="#">685659</a>	4689	862	2	262	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant cSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.dSep08</a>	<a href="#">303580</a>	4670	934	2	214	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant dSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.dSep08</a>	<a href="#">685659</a>	4670	934	2	214	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant dSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.eSep08</a>	<a href="#">303580</a>	4492	721	2	196	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant eSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.eSep08</a>	<a href="#">685659</a>	4492	721	2	196	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant eSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.fSep08</a>	<a href="#">303580</a>	5162	709	3	131	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant fSep08, mRNA.
<a href="#">Hexim2</a>	<a href="#">Hexim2.fSep08</a>	<a href="#">685659</a>	5162	709	3	131	hexamethylene bis-acetamide inducible 2 and hypothetical protein LOC685659 (Hexim2) alternative variant fSep08, mRNA.
<a href="#">Hey1</a>	<a href="#">Hey1.aSep08</a>	<a href="#">155437</a>	2024	562		165	hairy/enhancer-of-split related with YRPW motif 1 (Hey1) mRNA.
<a href="#">Hey2</a>	<a href="#">Hey2.aSep08</a>	<a href="#">155430</a>	6212	567		188	hairy/enhancer-of-split related with YRPW motif 2 (Hey2) mRNA.
<a href="#">Hfm1</a>	<a href="#">Hfm1.aSep08</a>	<a href="#">690161</a>	8031	896	3	24	HFM1, ATP-dependent DNA helicase homolog (S. cerevisiae) (2.7 kD) (Hfm1) alternative variant aSep08, mRNA.
<a href="#">Hgd</a>	<a href="#">Hgd.bSep08</a>	<a href="#">360719</a>	37679	758	10	237	homogentisate 1, 2-dioxygenase (Hgd) alternative variant bSep08, mRNA.
<a href="#">Hgd</a>	<a href="#">Hgd.cSep08</a>	<a href="#">360719</a>	30287	668	9	222	homogentisate 1, 2-dioxygenase (Hgd) alternative variant cSep08, mRNA.
<a href="#">Hgd</a>	<a href="#">Hgd.dSep08</a>	<a href="#">360719</a>	51862	1558	9	145	homogentisate 1, 2-dioxygenase (16.2 kD) (Hgd) alternative variant dSep08, complete mRNA.
<a href="#">Hgd</a>	<a href="#">Hgd.eSep08</a>	<a href="#">360719</a>	31399	720	6	145	homogentisate 1, 2-dioxygenase (16.2 kD) (Hgd) alternative variant eSep08, mRNA.
<a href="#">Hgd</a>	<a href="#">Hgd.fSep08</a>	<a href="#">360719</a>	4229	422	2	118	homogentisate 1, 2-dioxygenase (Hgd) alternative variant fSep08, mRNA.

<a href="#">Hgfac</a>	<a href="#">Hgfac.bSep08</a>	<a href="#">58947</a>	2830	787	3	229	hepatocyte growth factor activator (Hgfac) alternative variant bSep08, mRNA.
<a href="#">Hgfac</a>	<a href="#">Hgfac.cSep08</a>	<a href="#">58947</a>	1768	698	2	209	hepatocyte growth factor activator (Hgfac) alternative variant cSep08, mRNA.
<a href="#">Hgs</a>	<a href="#">Hgs.aSep08</a>	<a href="#">56084</a>	1450	941		150	HGF-regulated tyrosine kinase substrate (Hgs) alternative variant aSep08, mRNA.
<a href="#">HGTP_anticodon_0</a>	<a href="#">HGTP_anticodon.0.aSep08</a>		8890	1296		207	threonyl-tRNA synthetase (HGTP_anticodon.0) mRNA.
<a href="#">Hhat</a>	<a href="#">Hhat.aSep08</a>	<a href="#">289344</a>	41219	689		45	hedgehog acyltransferase (Hhat) mRNA.
<a href="#">Hhatl</a>	<a href="#">Hhatl.bSep08</a>	<a href="#">301073</a>	3415	766	5	221	hedgehog acyltransferase-like (Hhatl) alternative variant bSep08, mRNA.
<a href="#">Hhatl</a>	<a href="#">Hhatl.cSep08</a>	<a href="#">301073</a>	1225	1056	2	82	hedgehog acyltransferase-like (9.5 kD) (Hhatl) alternative variant cSep08, mRNA.
<a href="#">Hhatl</a>	<a href="#">Hhatl.eSep08</a>	<a href="#">301073</a>	3523	729	4	82	hedgehog acyltransferase-like (9.5 kD) (Hhatl) alternative variant eSep08, mRNA.
<a href="#">Hhex</a>	<a href="#">Hhex.bSep08</a>	<a href="#">79237</a>	2610	765	2	12	hematopoietically expressed homeobox (1.3 kD) (Hhex) alternative variant bSep08, mRNA.
<a href="#">Hhex</a>	<a href="#">Hhex.cSep08</a>	<a href="#">79237</a>	649	327	2	53	hematopoietically expressed homeobox (6.0 kD) (Hhex) alternative variant cSep08, mRNA.
<a href="#">Hhip</a>	<a href="#">Hhip.aSep08</a>	<a href="#">291936</a>	3209	727		94	hedgehog-interacting protein (Hhip) mRNA.
<a href="#">Hhipl1</a>	<a href="#">Hhipl1.aSep08</a>	<a href="#">362781</a>	5606	548		176	hedgehog interacting protein-like 1 (Hhipl1) mRNA.
<a href="#">Hiat1</a>	<a href="#">Hiat1.bSep08</a>	<a href="#">100134827</a>	3461	1762	3	155	hippocampus abundant gene transcript 1 (16.5 kD) (Hiat1) alternative variant bSep08, mRNA.
<a href="#">Hiat1</a>	<a href="#">Hiat1.cSep08</a>	<a href="#">100134827</a>	5018	837	2	58	hippocampus abundant gene transcript 1 (Hiat1) alternative variant cSep08, mRNA.
<a href="#">Hibch</a>	<a href="#">Hibch.aSep08</a>	<a href="#">301384</a>	58494	1673	2	355	3-hydroxyisobutyryl-Coenzyme A hydrolase (Hibch) alternative variant aSep08, mRNA.
<a href="#">Hic2</a>	<a href="#">Hic2.aSep08</a>	<a href="#">287940</a>	25985	728		174	hypermethylated in cancer 2 (Hic2) mRNA.
<a href="#">Hif1a</a>	<a href="#">Hif1a.aSep08</a>	<a href="#">29560</a>	45782	4620	15	828	hypoxia inducible factor 1, alpha subunit (92.7 kD) (Hif1a) alternative variant aSep08, mRNA.
<a href="#">Hif1a</a>	<a href="#">Hif1a.bSep08</a>	<a href="#">29560</a>	5240	2120	5	307	hypoxia inducible factor 1, alpha subunit (Hif1a) alternative variant bSep08, mRNA.
<a href="#">Hif1a</a>	<a href="#">Hif1a.cSep08</a>	<a href="#">29560</a>	24627	501	5	134	hypoxia inducible factor 1, alpha subunit (Hif1a) alternative variant cSep08, mRNA.
<a href="#">Hif1an</a>	<a href="#">Hif1an.bSep08</a>	<a href="#">309434</a>	9847	982		197	hypoxia-inducible factor 1, alpha subunit inhibitor (22.8 kD) (Hif1an) alternative variant bSep08, mRNA.
<a href="#">Higd1a</a>	<a href="#">Higd1a.bSep08</a>	<a href="#">140937</a>	8056	519	4	136	HIG1 domain family, member 1A (Higd1a) alternative variant bSep08, mRNA.
<a href="#">Higd1b</a>	<a href="#">Higd1b.aSep08</a>	<a href="#">287738</a>	2204	500	2	98	HIG1 domain family, member 1B (10.9 kD) (Higd1b) alternative variant aSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.bSep08</a>	<a href="#">290999</a>	30942	867	8	279	putative protein of eukaryotic origin (Higd2aandFaf2) alternative variant bSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.bSep08</a>	<a href="#">291000</a>	30942	867	8	279	putative protein of eukaryotic origin (Higd2aandFaf2) alternative variant bSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.cSep08</a>	<a href="#">290999</a>	34519	817	7	262	putative protein, with a coiled coil domain, of eukaryotic origin (Higd2aandFaf2) alternative variant cSep08, mRNA.

<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.cSep08</a>	<a href="#">291000</a>	34519	817	7	262	putative protein, with a coiled coil domain, of eukaryotic origin (Higd2aandFaf2) alternative variant cSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.dSep08</a>	<a href="#">290999</a>	29005	623	7	201	putative protein of eukaryotic origin (Higd2aandFaf2) alternative variant dSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.dSep08</a>	<a href="#">291000</a>	29005	623	7	201	putative protein of eukaryotic origin (Higd2aandFaf2) alternative variant dSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.eSep08</a>	<a href="#">290999</a>	34431	806	8	154	putative cytoplasmic protein of metazoan origin (17.9 kD) (Higd2aandFaf2) alternative variant eSep08, complete mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.eSep08</a>	<a href="#">291000</a>	34431	806	8	154	putative cytoplasmic protein of metazoan origin (17.9 kD) (Higd2aandFaf2) alternative variant eSep08, complete mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.hSep08</a>	<a href="#">290999</a>	57601	378	4	68	putative protein of bilateral origin (Higd2aandFaf2) alternative variant hSep08, mRNA.
<a href="#">Higd2aandFaf2</a>	<a href="#">Higd2aandFaf2.hSep08</a>	<a href="#">291000</a>	57601	378	4	68	putative protein of bilateral origin (Higd2aandFaf2) alternative variant hSep08, mRNA.
<a href="#">HIN.0</a>	<a href="#">HIN.0.aSep08</a>		3496	509		121	activated gene Interferon 20 (HIN.0) mRNA.
<a href="#">Hint2</a>	<a href="#">Hint2.bSep08</a>	<a href="#">313491</a>	2205	641	3	110	histidine triad nucleotide binding protein 2 CRA b like (11.6 kD) (Hint2) alternative variant bSep08, complete mRNA.
<a href="#">Hint2</a>	<a href="#">Hint2.cSep08</a>	<a href="#">313491</a>	880	778	1	67	putative protein (Hint2) alternative variant cSep08, mRNA.
<a href="#">Hint3</a>	<a href="#">Hint3.aSep08</a>	<a href="#">246769</a>	9862	958	5	175	histidine triad nucleotide binding protein 3 (19.7 kD) (Hint3) alternative variant aSep08, mRNA.
<a href="#">Hint3</a>	<a href="#">Hint3.bSep08</a>	<a href="#">246769</a>	9588	557	4	120	histidine triad nucleotide binding protein 3 (Hint3) alternative variant bSep08, mRNA.
<a href="#">Hint3</a>	<a href="#">Hint3.eSep08</a>	<a href="#">246769</a>	5644	256	2	43	histidine triad nucleotide binding protein 3 (Hint3) alternative variant eSep08, mRNA.
<a href="#">Hip1</a>	<a href="#">Hip1.aSep08</a>	<a href="#">192154</a>	61715	2608	25	808	huntingtin interacting protein 1 (Hip1) alternative variant aSep08, mRNA.
<a href="#">Hipk1</a>	<a href="#">Hipk1.bSep08</a>	<a href="#">365895</a>	10280	743	6	155	homeodomain interacting protein kinase 1 (Hipk1) alternative variant bSep08, mRNA.
<a href="#">Hipk1</a>	<a href="#">Hipk1.cSep08</a>	<a href="#">365895</a>	3767	372	3	87	homeodomain interacting protein kinase 1 (Hipk1) alternative variant cSep08, mRNA.
<a href="#">Hipk1</a>	<a href="#">Hipk1.dSep08</a>	<a href="#">365895</a>	763	232	2	69	homeodomain interacting protein kinase 1 (Hipk1) alternative variant dSep08, mRNA.
<a href="#">Hipk2</a>	<a href="#">Hipk2.bSep08</a>	<a href="#">362342</a>	33066	1216	8	405	putative protein of metazoan origin (Hipk2) alternative variant bSep08, mRNA.
<a href="#">Hipk2</a>	<a href="#">Hipk2.cSep08</a>	<a href="#">362342</a>	13737	412	4	137	putative protein of fungal and metazoan origin (Hipk2) alternative variant cSep08, mRNA.
<a href="#">Hipk2</a>	<a href="#">Hipk2.dSep08</a>	<a href="#">362342</a>	3345	2673	3	116	putative protein (12.9 kD) (Hipk2) alternative variant dSep08, mRNA.
<a href="#">Hipk3</a>	<a href="#">Hipk3.bSep08</a>	<a href="#">83617</a>	2921	758	3	252	homeodomain interacting protein kinase 3 (Hipk3) alternative variant bSep08, mRNA.
<a href="#">Hipk3</a>	<a href="#">Hipk3.cSep08</a>	<a href="#">83617</a>	2884	724	3	241	homeodomain interacting protein kinase 3 (Hipk3) alternative variant cSep08, mRNA.
<a href="#">Hipk3</a>	<a href="#">Hipk3.dSep08</a>	<a href="#">83617</a>	2577	835	3	237	homeodomain interacting protein kinase 3 (Hipk3) alternative variant dSep08, mRNA.

<a href="#">Hipk3</a>	<a href="#">Hipk3.eSep08</a>	<a href="#">83617</a>	2544	697	2	54	homeodomain interacting protein kinase 3 (Hipk3) alternative variant eSep08, mRNA.
<a href="#">Hirip3</a>	<a href="#">Hirip3.bSep08</a>	<a href="#">361650</a>	2551	2041	5	432	HIRA interacting protein 3 (Hirip3) alternative variant bSep08, mRNA.
<a href="#">Hirip3</a>	<a href="#">Hirip3.cSep08</a>	<a href="#">361650</a>	766	686	2	91	HIRA interacting protein 3 (Hirip3) alternative variant cSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.cSep08</a>	<a href="#">311355</a>	12720	759	7	253	putative protein of bilateral origin (Hisppd2a) alternative variant cSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.dSep08</a>	<a href="#">311355</a>	14327	718	6	239	putative protein of metazoan origin (Hisppd2a) alternative variant dSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.eSep08</a>	<a href="#">311355</a>	12444	595	5	197	putative protein (Hisppd2a) alternative variant eSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.fSep08</a>	<a href="#">311355</a>	6150	640	5	181	putative protein of eukaryotic origin (Hisppd2a) alternative variant fSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.gSep08</a>	<a href="#">311355</a>	1007	353	4	117	putative protein of eukaryotic origin (Hisppd2a) alternative variant gSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.hSep08</a>	<a href="#">311355</a>	790	350	3	116	putative protein of eukaryotic origin (Hisppd2a) alternative variant hSep08, mRNA.
<a href="#">Hisppd2a</a>	<a href="#">Hisppd2a.iSep08</a>	<a href="#">311355</a>	1046	458	3	107	putative protein of eukaryotic origin (Hisppd2a) alternative variant iSep08, mRNA.
<a href="#">Hist1h2bn</a>	<a href="#">Hist1h2bn.bSep08</a>	<a href="#">291157</a>	2300	425	2	124	histone cluster 1, H2bn (Hist1h2bn) alternative variant bSep08, mRNA.
<a href="#">Hist2h2bb</a>	<a href="#">Hist2h2bb.aSep08</a>	<a href="#">295278</a>	6022	629	2	131	histone cluster 2, H2bb (14.4 kD) (Hist2h2bb) alternative variant aSep08, mRNA.
<a href="#">Histone.7</a>	<a href="#">Histone.7.bSep08</a>		724	392	2	125	histone H3.2 (Histone.7) alternative variant bSep08, mRNA.
<a href="#">Histone.8</a>	<a href="#">Histone.8.aSep08</a>		19549	639		212	son of sevenless (Histone.8) mRNA.
<a href="#">Hist_deacetyl.0</a>	<a href="#">Hist_deacetyl.0.aSep08</a>		11346	2985	7	913	histone deacetylase 5 (Hist_deacetyl.0) alternative variant aSep08, mRNA.
<a href="#">Hist_deacetyl.0</a>	<a href="#">Hist_deacetyl.0.bSep08</a>		3309	337	2	97	histone deacetylase 5 (Hist_deacetyl.0) alternative variant bSep08, mRNA.
<a href="#">Hivep1</a>	<a href="#">Hivep1.bSep08</a>	<a href="#">117140</a>	3637	2052	2	512	human immunodeficiency virus type I enhancer binding protein 1 (Hivep1) alternative variant bSep08, mRNA.
<a href="#">Hivep1</a>	<a href="#">Hivep1.eSep08</a>	<a href="#">117140</a>	45848	374	3	124	human immunodeficiency virus type I enhancer binding protein 1 (Hivep1) alternative variant eSep08, mRNA.
<a href="#">Hivep1</a>	<a href="#">Hivep1.gSep08</a>	<a href="#">117140</a>	942	230	2	76	human immunodeficiency virus type I enhancer binding protein 1 (Hivep1) alternative variant gSep08, mRNA.
<a href="#">Hivep2</a>	<a href="#">Hivep2.bSep08</a>	<a href="#">29721</a>	6154	5393		1652	human immunodeficiency virus type I enhancer binding protein 2 (Hivep2) alternative variant bSep08, mRNA.
<a href="#">Hivep2</a>	<a href="#">Hivep2.cSep08</a>	<a href="#">29721</a>	8357	3319		555	human immunodeficiency virus type I enhancer binding protein 2 (Hivep2) alternative variant cSep08, mRNA.
<a href="#">Hjurp</a>	<a href="#">Hjurp.aSep08</a>	<a href="#">316602</a>	2136	775		258	holliday junction recognition protein (Hjurp) mRNA.
<a href="#">Hk2</a>	<a href="#">Hk2.bSep08</a>	<a href="#">25059</a>	6039	2912	2	231	hexokinase 2 (25.6 kD) (Hk2) alternative variant bSep08, mRNA.
<a href="#">Hkr3</a>	<a href="#">Hkr3.bSep08</a>	<a href="#">362668</a>	2607	1411	5	292	GLI-Kruppel family member HKR3 (32.4 kD) (Hkr3) alternative variant bSep08, mRNA.

<a href="#">Hkr3</a>	<a href="#">Hkr3.cSep08</a>	<a href="#">362668</a>	1206	703	2	223	GLI-Kruppel family member HKR3 (Hkr3) alternative variant cSep08, mRNA.
<a href="#">Hla-dma</a>	<a href="#">Hla-dma.bSep08</a>	<a href="#">294274</a>	3356	1913	2	195	major histocompatibility complex, class II, DM alpha (21.7 kD) (Hla-dma) alternative variant bSep08, mRNA.
<a href="#">Hla-dmb</a>	<a href="#">Hla-dmb.bSep08</a>	<a href="#">294273</a>	775	511	2	40	major histocompatibility complex, class II, DM beta (4.6 kD) (Hla-dmb) alternative variant bSep08, mRNA.
<a href="#">Hlcs</a>	<a href="#">Hlcs.aSep08</a>	<a href="#">288240</a>	811	572		190	holocarboxylase synthetase (biotin- [propionyl-Coenzyme A-carboxylase (ATP-hydrolysing)] ligase) (Hlcs) mRNA.
<a href="#">HLH.1</a>	<a href="#">HLH.1.aSep08</a>		1945	732		131	basic helix-loop-helix dimerisation region bHLH (HLH.1) mRNA.
<a href="#">HLH.2</a>	<a href="#">HLH.2.bSep08</a>		27516	756	5	154	max 1 (17.6 kD) (HLH.2) alternative variant bSep08, mRNA.
<a href="#">HLH.3</a>	<a href="#">HLH.3.aSep08</a>		3759	1995		149	max 1 (HLH.3) mRNA.
<a href="#">Hmbox1</a>	<a href="#">Hmbox1.aSep08</a>	<a href="#">305968</a>	27236	1830	4	159	homeobox containing 1 (Hmbox1) alternative variant aSep08, mRNA.
<a href="#">Hmbox1</a>	<a href="#">Hmbox1.bSep08</a>	<a href="#">305968</a>	2452	872	1	49	homeobox containing 1 (Hmbox1) alternative variant bSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.bSep08</a>	<a href="#">25709</a>	4447	781	11	260	hydroxymethylbilane synthase (Hmbs) alternative variant bSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.cSep08</a>	<a href="#">25709</a>	4495	751	10	250	hydroxymethylbilane synthase (Hmbs) alternative variant cSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.dSep08</a>	<a href="#">25709</a>	3925	731	9	243	hydroxymethylbilane synthase (Hmbs) alternative variant dSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.eSep08</a>	<a href="#">25709</a>	2574	682	7	149	hydroxymethylbilane synthase (Hmbs) alternative variant eSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.fSep08</a>	<a href="#">25709</a>	2633	567	4	105	hydroxymethylbilane synthase CRA a (Hmbs) alternative variant fSep08, mRNA.
<a href="#">Hmbs</a>	<a href="#">Hmbs.gSep08</a>	<a href="#">25709</a>	2308	361	2	97	putative protein of mammalian origin (Hmbs) alternative variant gSep08, mRNA.
<a href="#">Hmcn1</a>	<a href="#">Hmcn1.aSep08</a>	<a href="#">289094</a>	21441	2004	6	328	hemicentin 1 (Hmcn1) alternative variant aSep08, mRNA.
<a href="#">Hmcn2</a>	<a href="#">Hmcn2.bSep08</a>	<a href="#">686132</a>	818	381	2	112	hemicentin 2 (Hmcn2) alternative variant bSep08, mRNA.
<a href="#">Hmg20b</a>	<a href="#">Hmg20b.bSep08</a>	<a href="#">362825</a>	3733	828	8	275	high mobility group 20 B (Hmg20b) alternative variant bSep08, mRNA.
<a href="#">Hmg20b</a>	<a href="#">Hmg20b.cSep08</a>	<a href="#">362825</a>	831	661	3	186	high mobility group 20 B (Hmg20b) alternative variant cSep08, mRNA.
<a href="#">Hmg20b</a>	<a href="#">Hmg20b.dSep08</a>	<a href="#">362825</a>	2680	1005	5	71	high mobility group 20 B (Hmg20b) alternative variant dSep08, mRNA.
<a href="#">Hmg20b</a>	<a href="#">Hmg20b.eSep08</a>	<a href="#">362825</a>	669	586	2	160	high mobility group 20 B (Hmg20b) alternative variant eSep08, mRNA.
<a href="#">Hmg20b</a>	<a href="#">Hmg20b.gSep08</a>	<a href="#">362825</a>	1437	765	2	66	high mobility group 20 B (Hmg20b) alternative variant gSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.aSep08</a>	<a href="#">117062</a>	6499	927	3	120	high mobility group AT-hook 1 and hypothetical protein LOC689053 (Hmga1) alternative variant aSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.aSep08</a>	<a href="#">689053</a>	6499	927	3	120	high mobility group AT-hook 1 and hypothetical protein LOC689053 (Hmga1) alternative variant aSep08, mRNA.

<a href="#">Hmga1</a>	<a href="#">Hmga1.bSep08</a>	<a href="#">117062</a>	6527	988	3	107	high mobility group AT-hook 1 and hypothetical protein LOC689053 (11.7 kD) (Hmga1) alternative variant bSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.bSep08</a>	<a href="#">689053</a>	6527	988	3	107	high mobility group AT-hook 1 and hypothetical protein LOC689053 (11.7 kD) (Hmga1) alternative variant bSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.cSep08</a>	<a href="#">117062</a>	6094	667	4	107	high mobility group AT-hook 1 and hypothetical protein LOC689053 (11.7 kD) (Hmga1) alternative variant cSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.cSep08</a>	<a href="#">689053</a>	6094	667	4	107	high mobility group AT-hook 1 and hypothetical protein LOC689053 (11.7 kD) (Hmga1) alternative variant cSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.eSep08</a>	<a href="#">117062</a>	6956	868	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant eSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.eSep08</a>	<a href="#">689053</a>	6956	868	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant eSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.fSep08</a>	<a href="#">117062</a>	6494	922	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant fSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.fSep08</a>	<a href="#">689053</a>	6494	922	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant fSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.gSep08</a>	<a href="#">117062</a>	6221	761	4	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant gSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.gSep08</a>	<a href="#">689053</a>	6221	761	4	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant gSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.hSep08</a>	<a href="#">117062</a>	5124	691	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant hSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.hSep08</a>	<a href="#">689053</a>	5124	691	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant hSep08, mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.iSep08</a>	<a href="#">117062</a>	7132	1558	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant iSep08, complete mRNA.
<a href="#">Hmga1</a>	<a href="#">Hmga1.iSep08</a>	<a href="#">689053</a>	7132	1558	3	96	high mobility group AT-hook 1 and hypothetical protein LOC689053 (10.7 kD) (Hmga1) alternative variant iSep08, complete mRNA.
<a href="#">Hmgb1</a>	<a href="#">Hmgb1.aSep08</a>	<a href="#">25459</a>	23295	759		205	high mobility group box 1 (Hmgb1) mRNA.
<a href="#">Hmgb2</a>	<a href="#">Hmgb2.bSep08</a>	<a href="#">29395</a>	1898	1078	4	164	high mobility group box 2 (19.0 kD) (Hmgb2) alternative variant bSep08, mRNA.
<a href="#">Hmgb2</a>	<a href="#">Hmgb2.cSep08</a>	<a href="#">29395</a>	1297	783	2	82	high mobility group box 2 (9.1 kD) (Hmgb2) alternative variant cSep08, mRNA.

<a href="#">Hmgb2</a>	<a href="#">Hmgb2.dSep08</a>	<a href="#">29395</a>	520	172	2	46	high mobility group box 2 (Hmgb2) alternative variant dSep08, mRNA.
<a href="#">Hmgb2l1</a>	<a href="#">Hmgb2l1.bSep08</a>	<a href="#">307667</a>	11373	2466	6	184	high mobility group box 2-like 1 (Hmgb2l1) alternative variant bSep08, mRNA.
<a href="#">Hmgb2l1</a>	<a href="#">Hmgb2l1.dSep08</a>	<a href="#">307667</a>	1504	738	2	91	high mobility group box 2-like 1 (10.3 kD) (Hmgb2l1) alternative variant dSep08, mRNA.
<a href="#">Hmgb3</a>	<a href="#">Hmgb3.aSep08</a>	<a href="#">305373</a>	5028	1467	1	241	high mobility group box 3 (27.6 kD) (Hmgb3) alternative variant aSep08, mRNA.
<a href="#">Hmgb3</a>	<a href="#">Hmgb3.bSep08</a>	<a href="#">305373</a>	5241	1527	1	200	high mobility group box 3 (23.0 kD) (Hmgb3) alternative variant bSep08, mRNA.
<a href="#">Hmgb3</a>	<a href="#">Hmgb3.cSep08</a>	<a href="#">305373</a>	1584	651	1	114	high mobility group box 3 (Hmgb3) alternative variant cSep08, mRNA.
<a href="#">Hmgcl</a>	<a href="#">Hmgcl.bSep08</a>	<a href="#">79238</a>	1580	673	2	144	3-hydroxy-3-methylglutaryl-Coenzyme A lyase (Hmgcl) alternative variant bSep08, mRNA.
<a href="#">Hmgcr</a>	<a href="#">Hmgcr.aSep08</a>	<a href="#">25675</a>	10327	3626		673	3-hydroxy-3-methylglutaryl-Coenzyme A reductase (Hmgcr) mRNA.
<a href="#">Hmgcs1</a>	<a href="#">Hmgcs1.bSep08</a>	<a href="#">29637</a>	10767	940	4	225	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (Hmgcs1) alternative variant bSep08, mRNA.
<a href="#">Hmgcs1</a>	<a href="#">Hmgcs1.cSep08</a>	<a href="#">29637</a>	8713	624	3	37	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (Hmgcs1) alternative variant cSep08, mRNA.
<a href="#">Hmgcs2</a>	<a href="#">Hmgcs2.bSep08</a>	<a href="#">24450</a>	7670	1590	1	146	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2 (16.5 kD) (Hmgcs2) alternative variant bSep08, mRNA.
<a href="#">Hmgn1</a>	<a href="#">Hmgn1.aSep08</a>	<a href="#">360704</a>	2902	477	4	102	high mobility group nucleosomal binding domain 1 (Hmgn1) alternative variant aSep08, mRNA.
<a href="#">Hmgn1</a>	<a href="#">Hmgn1.cSep08</a>	<a href="#">360704</a>	5996	1255	6	95	high mobility group nucleosomal binding domain 1 (10.0 kD) (Hmgn1) alternative variant cSep08, complete mRNA.
<a href="#">Hmgn1</a>	<a href="#">Hmgn1.dSep08</a>	<a href="#">360704</a>	3064	878	2	46	high mobility group nucleosomal binding domain 1 (5.4 kD) (Hmgn1) alternative variant dSep08, mRNA.
<a href="#">Hmgn2</a>	<a href="#">Hmgn2.bSep08</a>	<a href="#">114637</a>	2722	870	3	72	high mobility group nucleosomal binding domain 2 (7.5 kD) (Hmgn2) alternative variant bSep08, mRNA.
<a href="#">Hmgn2</a>	<a href="#">Hmgn2.cSep08</a>	<a href="#">114637</a>	1326	730	2	42	high mobility group nucleosomal binding domain 2 (5.2 kD) (Hmgn2) alternative variant cSep08, mRNA.
<a href="#">Hmgn3</a>	<a href="#">Hmgn3.bSep08</a>	<a href="#">113990</a>	36634	828	6	126	high mobility group nucleosomal binding domain 3 (Hmgn3) alternative variant bSep08, mRNA.
<a href="#">Hmgn3</a>	<a href="#">Hmgn3.cSep08</a>	<a href="#">113990</a>	36669	822	6	119	high mobility group nucleosomal binding domain 3 (Hmgn3) alternative variant cSep08, mRNA.
<a href="#">Hmgn3</a>	<a href="#">Hmgn3.dSep08</a>	<a href="#">113990</a>	2419	657	3	43	high mobility group nucleosomal binding domain 3 (5.0 kD) (Hmgn3) alternative variant dSep08, mRNA.
<a href="#">HMG_box.2</a>	<a href="#">HMG_box.2.aSep08</a>		12365	723		241	SRY -box 30 (HMG_box.2) mRNA.
<a href="#">HMG_box.4</a>	<a href="#">HMG_box.4.aSep08</a>		1224	574		191	myeloid lymphoid mixed-lineage leukemia 2 CRA a like (HMG_box.4) mRNA.
<a href="#">Hmha1</a>	<a href="#">Hmha1.bSep08</a>	<a href="#">314618</a>	4654	1213	6	208	histocompatibility ha-1 (Hmha1) alternative variant bSep08, mRNA.
<a href="#">Hmha1</a>	<a href="#">Hmha1.cSep08</a>	<a href="#">314618</a>	1496	642	2	190	histocompatibility ha-1 (Hmha1) alternative variant cSep08, mRNA.
<a href="#">Hmha1</a>	<a href="#">Hmha1.dSep08</a>	<a href="#">314618</a>	1549	762	3	151	histocompatibility ha-1 (Hmha1) alternative variant dSep08, mRNA.

<a href="#">Hmha1</a>	<a href="#">Hmha1.eSep08</a>	<a href="#">314618</a>	1356	662	2	92	histocompatibility ha-1 (10.5 kD) (Hmha1) alternative variant eSep08, mRNA.
<a href="#">Hmha1</a>	<a href="#">Hmha1.fSep08</a>	<a href="#">314618</a>	761	467	3	45	putative protein (Hmha1) alternative variant fSep08, mRNA.
<a href="#">Hmnr</a>	<a href="#">Hmnr.aSep08</a>	<a href="#">25460</a>	21236	2514	12	396	hyaluronan mediated motility receptor (RHAMM) (Hmnr) alternative variant aSep08, mRNA.
<a href="#">Hmnr</a>	<a href="#">Hmnr.cSep08</a>	<a href="#">25460</a>	1198	493	2	63	hyaluronan mediated motility receptor (RHAMM) (Hmnr) alternative variant cSep08, mRNA.
<a href="#">Hmox1</a>	<a href="#">Hmox1.bSep08</a>	<a href="#">24451</a>	3584	982	2	139	heme oxygenase (decycling) 1 (Hmox1) alternative variant bSep08, mRNA.
<a href="#">Hmox2</a>	<a href="#">Hmox2.bSep08</a>	<a href="#">79239</a>	45256	900	3	262	heme oxygenase (decycling) 2 (Hmox2) alternative variant bSep08, mRNA.
<a href="#">Hmox2</a>	<a href="#">Hmox2.cSep08</a>	<a href="#">79239</a>	44535	865	2	238	heme oxygenase (decycling) 2 (Hmox2) alternative variant cSep08, mRNA.
<a href="#">Hmox2</a>	<a href="#">Hmox2.dSep08</a>	<a href="#">79239</a>	44237	777	2	171	heme oxygenase (decycling) 2 (Hmox2) alternative variant dSep08, mRNA.
<a href="#">Hmx1</a>	<a href="#">Hmx1.bSep08</a>	<a href="#">360960</a>	1367	1277	1	177	h6 homeo box 1 (Hmx1) alternative variant bSep08, mRNA.
<a href="#">Hmx2</a>	<a href="#">Hmx2.bSep08</a>	<a href="#">293538</a>	7640	1206		81	h6 homeo box 2 (8.9 kD) (Hmx2) alternative variant bSep08, mRNA.
<a href="#">Hn1</a>	<a href="#">Hn1.bSep08</a>	<a href="#">287828</a>	16274	852	5	132	hematological and neurological expressed sequence 1 (13.8 kD) (Hn1) alternative variant bSep08, mRNA.
<a href="#">Hn1l</a>	<a href="#">Hn1l.aSep08</a>	<a href="#">360492</a>	19484	1092	6	205	hematological and neurological expressed 1-like (Hn1l) alternative variant aSep08, mRNA.
<a href="#">Hn1l</a>	<a href="#">Hn1l.cSep08</a>	<a href="#">360492</a>	13447	416	3	112	hematological and neurological expressed 1-like (Hn1l) alternative variant cSep08, mRNA.
<a href="#">Hnf1b</a>	<a href="#">Hnf1b.bSep08</a>	<a href="#">25640</a>	5315	695		181	HNF1 homeobox B (Hnf1b) alternative variant bSep08, mRNA.
<a href="#">Hnf4g</a>	<a href="#">Hnf4g.bSep08</a>	<a href="#">365744</a>	17434	1302	7	414	hepatocyte nuclear factor 4, gamma (Hnf4g) alternative variant bSep08, mRNA.
<a href="#">Hnf4g</a>	<a href="#">Hnf4g.cSep08</a>	<a href="#">365744</a>	18391	1318	8	364	hepatocyte nuclear factor 4, gamma (Hnf4g) alternative variant cSep08, mRNA.
<a href="#">Hnrnpa1</a>	<a href="#">Hnrnpa1.aSep08</a>	<a href="#">29578</a>	3350	1464	10	328	heterogeneous nuclear ribonucleoprotein A1 (33.7 kD) (Hnrnpa1) alternative variant aSep08, mRNA.
<a href="#">Hnrnpa1</a>	<a href="#">Hnrnpa1.cSep08</a>	<a href="#">29578</a>	3126	442	4	43	heterogeneous nuclear ribonucleoprotein A1 (4.4 kD) (Hnrnpa1) alternative variant cSep08, mRNA.
<a href="#">Hnrnpa1</a>	<a href="#">Hnrnpa1.dSep08</a>	<a href="#">29578</a>	1125	1024	2	69	heterogeneous nuclear ribonucleoprotein A1 (8.0 kD) (Hnrnpa1) alternative variant dSep08, mRNA.
<a href="#">Hnrnpa1</a>	<a href="#">Hnrnpa1.eSep08</a>	<a href="#">29578</a>	1599	528	3	39	heterogeneous nuclear ribonucleoprotein A1 (4.2 kD) (Hnrnpa1) alternative variant eSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.bSep08</a>	<a href="#">362361</a>	6278	1544	8	261	nuclear ribonucleoprotein A2 B1 (27.6 kD) (Hnrnpa2b1) alternative variant bSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.bSep08</a>	<a href="#">685548</a>	6278	1544	8	261	nuclear ribonucleoprotein A2 B1 (27.6 kD) (Hnrnpa2b1) alternative variant bSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.cSep08</a>	<a href="#">362361</a>	5333	2867	7	232	nuclear ribonucleoprotein A2 B1 (Hnrnpa2b1) alternative variant cSep08, mRNA.



<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.cSep08</a>	<a href="#">685548</a>	5333	2867	7	232	nuclear ribonucleoprotein A2 B1 (Hnrnpa2b1) alternative variant cSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.dSep08</a>	<a href="#">362361</a>	3639	1053	6	178	nuclear ribonucleoprotein A2 b1 (Hnrnpa2b1) alternative variant dSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.dSep08</a>	<a href="#">685548</a>	3639	1053	6	178	nuclear ribonucleoprotein A2 b1 (Hnrnpa2b1) alternative variant dSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.eSep08</a>	<a href="#">362361</a>	3166	658	3	96	CRA d like (10.4 kD) (Hnrnpa2b1) alternative variant eSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.eSep08</a>	<a href="#">685548</a>	3166	658	3	96	CRA d like (10.4 kD) (Hnrnpa2b1) alternative variant eSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.fSep08</a>	<a href="#">362361</a>	1274	497	3	86	CRA d like (9.4 kD) (Hnrnpa2b1) alternative variant fSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.fSep08</a>	<a href="#">685548</a>	1274	497	3	86	CRA d like (9.4 kD) (Hnrnpa2b1) alternative variant fSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.gSep08</a>	<a href="#">362361</a>	3469	697	4	75	CRA c like (8.3 kD) (Hnrnpa2b1) alternative variant gSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.gSep08</a>	<a href="#">685548</a>	3469	697	4	75	CRA c like (8.3 kD) (Hnrnpa2b1) alternative variant gSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.hSep08</a>	<a href="#">362361</a>	1302	811	2	63	CRA d like (6.8 kD) (Hnrnpa2b1) alternative variant hSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.hSep08</a>	<a href="#">685548</a>	1302	811	2	63	CRA d like (6.8 kD) (Hnrnpa2b1) alternative variant hSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.iSep08</a>	<a href="#">362361</a>	1329	775	2	42	CRA c like (4.7 kD) (Hnrnpa2b1) alternative variant iSep08, mRNA.
<a href="#">Hnrnpa2b1</a>	<a href="#">Hnrnpa2b1.iSep08</a>	<a href="#">685548</a>	1329	775	2	42	CRA c like (4.7 kD) (Hnrnpa2b1) alternative variant iSep08, mRNA.
<a href="#">Hnrnpa3</a>	<a href="#">Hnrnpa3.cSep08</a>	<a href="#">362152</a>	4404	4317	1	79	heterogeneous nuclear ribonucleoprotein A3 (8.9 kD) (Hnrnpa3) alternative variant cSep08, mRNA.
<a href="#">Hnrnpa3</a>	<a href="#">Hnrnpa3.dSep08</a>	<a href="#">362152</a>	4394	1602	2	55	heterogeneous nuclear ribonucleoprotein A3 (6.2 kD) (Hnrnpa3) alternative variant dSep08, mRNA.
<a href="#">Hnrnpa3</a>	<a href="#">Hnrnpa3.eSep08</a>	<a href="#">362152</a>	1109	929	2	48	heterogeneous nuclear ribonucleoprotein A3 (5.3 kD) (Hnrnpa3) alternative variant eSep08, mRNA.
<a href="#">Hnrnpa3</a>	<a href="#">Hnrnpa3.fSep08</a>	<a href="#">362152</a>	1908	660	2	58	heterogeneous nuclear ribonucleoprotein A3 (Hnrnpa3) alternative variant fSep08, mRNA.
<a href="#">Hnrnpab</a>	<a href="#">Hnrnpab.bSep08</a>	<a href="#">83498</a>	4545	1214	6	224	heterogeneous nuclear ribonucleoprotein A/B (Hnrnpab) alternative variant bSep08, mRNA.
<a href="#">Hnrnpab</a>	<a href="#">Hnrnpab.cSep08</a>	<a href="#">83498</a>	653	570	2	75	heterogeneous nuclear ribonucleoprotein A/B (7.1 kD) (Hnrnpab) alternative variant cSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.aSep08</a>	<a href="#">290046</a>	30366	1951	9	313	heterogeneous nuclear ribonucleoprotein C (34.4 kD) (Hnrnpc) alternative variant aSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.bSep08</a>	<a href="#">290046</a>	29660	1198	8	306	heterogeneous nuclear ribonucleoprotein C (33.7 kD) (Hnrnpc) alternative variant bSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.cSep08</a>	<a href="#">290046</a>	30161	2914	9	305	heterogeneous nuclear ribonucleoprotein C (33.6 kD) (Hnrnpc) alternative variant cSep08, complete mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.dSep08</a>	<a href="#">290046</a>	29684	1230	9	300	heterogeneous nuclear ribonucleoprotein C (33.1 kD) (Hnrnpc) alternative variant dSep08, mRNA.

<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.eSep08</a>	<a href="#">290046</a>	29820	1345	9	293	heterogeneous nuclear ribonucleoprotein C (32.3 kD) (Hnrnpc) alternative variant eSep08, complete mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.fSep08</a>	<a href="#">290046</a>	30171	1667	8	292	heterogeneous nuclear ribonucleoprotein C (32.2 kD) (Hnrnpc) alternative variant fSep08, complete mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.gSep08</a>	<a href="#">290046</a>	29491	1013	9	290	heterogeneous nuclear ribonucleoprotein C (Hnrnpc) alternative variant gSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.hSep08</a>	<a href="#">290046</a>	22592	1257	6	213	heterogeneous nuclear ribonucleoprotein C (Hnrnpc) alternative variant hSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.iSep08</a>	<a href="#">290046</a>	28847	754	8	190	heterogeneous nuclear ribonucleoprotein C (Hnrnpc) alternative variant iSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.jSep08</a>	<a href="#">290046</a>	27467	825	7	167	heterogeneous nuclear ribonucleoprotein C (Hnrnpc) alternative variant jSep08, mRNA.
<a href="#">Hnrnpc</a>	<a href="#">Hnrnpc.mSep08</a>	<a href="#">290046</a>	7736	1594	2	72	heterogeneous nuclear ribonucleoprotein C (7.8 kD) (Hnrnpc) alternative variant mSep08, mRNA.
<a href="#">Hnrmpf</a>	<a href="#">Hnrmpf.dSep08</a>	<a href="#">64200</a>	18408	424	4	113	heterogeneous nuclear ribonucleoprotein F (Hnrmpf) alternative variant dSep08, mRNA.
<a href="#">Hnrmpf</a>	<a href="#">Hnrmpf.fSep08</a>	<a href="#">64200</a>	21754	3068	4	94	heterogeneous nuclear ribonucleoprotein F (10.8 kD) (Hnrmpf) alternative variant fSep08, mRNA.
<a href="#">Hnrmpf</a>	<a href="#">Hnrmpf.gSep08</a>	<a href="#">64200</a>	15158	900	3	102	heterogeneous nuclear ribonucleoprotein F (Hnrmpf) alternative variant gSep08, mRNA.
<a href="#">Hnrnph2</a>	<a href="#">Hnrnph2.bSep08</a>	<a href="#">308650</a>	5889	2380	2	66	putative protein (Hnrnph2) alternative variant bSep08, mRNA.
<a href="#">Hnrnph3</a>	<a href="#">Hnrnph3.bSep08</a>	<a href="#">361838</a>	3029	1494	6	177	heterogeneous nuclear ribonucleoprotein H3 (18.0 kD) (Hnrnph3) alternative variant bSep08, mRNA.
<a href="#">Hnrnph3</a>	<a href="#">Hnrnph3.cSep08</a>	<a href="#">361838</a>	1768	1088	2	88	heterogeneous nuclear ribonucleoprotein H3 (Hnrnph3) alternative variant cSep08, mRNA.
<a href="#">Hnrnpk</a>	<a href="#">Hnrnpk.aSep08</a>	<a href="#">117282</a>	11280	2994	16	464	heterogeneous nuclear ribonucleoprotein K (51.0 kD) (Hnrnpk) alternative variant aSep08, mRNA.
<a href="#">Hnrnpk</a>	<a href="#">Hnrnpk.cSep08</a>	<a href="#">117282</a>	7846	1006	9	289	heterogeneous nuclear ribonucleoprotein K (Hnrnpk) alternative variant cSep08, mRNA.
<a href="#">Hnrnpk</a>	<a href="#">Hnrnpk.dSep08</a>	<a href="#">117282</a>	1441	867	4	118	heterogeneous nuclear ribonucleoprotein K (Hnrnpk) alternative variant dSep08, mRNA.
<a href="#">Hnrnpk</a>	<a href="#">Hnrnpk.eSep08</a>	<a href="#">117282</a>	3355	972	4	97	heterogeneous nuclear ribonucleoprotein K (Hnrnpk) alternative variant eSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.cSep08</a>	<a href="#">80846</a>	3478	1208	4	295	heterogeneous nuclear ribonucleoprotein L (32.6 kD) (Hnrnpl) alternative variant cSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.dSep08</a>	<a href="#">80846</a>	9372	820	8	272	heterogeneous nuclear ribonucleoprotein L CRA b (Hnrnpl) alternative variant dSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.eSep08</a>	<a href="#">80846</a>	4214	826	7	270	heterogeneous nuclear ribonucleoprotein L (Hnrnpl) alternative variant eSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.gSep08</a>	<a href="#">80846</a>	1331	1220	2	135	putative protein of mammalian origin (Hnrnpl) alternative variant gSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.hSep08</a>	<a href="#">80846</a>	961	847	2	120	heterogeneous nuclear ribonucleoprotein hnRNP-L (Hnrnpl) alternative variant hSep08, mRNA.
<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.iSep08</a>	<a href="#">80846</a>	1092	810	2	90	heterogeneous nuclear ribonucleoprotein L (Hnrnpl) alternative variant iSep08, mRNA.

<a href="#">Hnrnpl</a>	<a href="#">Hnrnpl.kSep08</a>	<a href="#">80846</a>	5252	791	7	141	heterogeneous nuclear ribonucleoprotein L (Hnrnpl) alternative variant kSep08, mRNA.
<a href="#">Hnrnpr</a>	<a href="#">Hnrnpr.cSep08</a>	<a href="#">319110</a>	17107	648	6	185	heterogeneous nuclear ribonucleoprotein R (Hnrnpr) alternative variant cSep08, mRNA.
<a href="#">Hnrnpr</a>	<a href="#">Hnrnpr.dSep08</a>	<a href="#">319110</a>	32971	2582	8	177	heterogeneous nuclear ribonucleoprotein R (19.6 kD) (Hnrnpr) alternative variant dSep08, complete mRNA.
<a href="#">Hnrnpr</a>	<a href="#">Hnrnpr.fSep08</a>	<a href="#">319110</a>	5819	413	2	95	heterogeneous nuclear ribonucleoprotein R (10.5 kD) (Hnrnpr) alternative variant fSep08, mRNA.
<a href="#">Hnrnpu</a>	<a href="#">Hnrnpu.bSep08</a>	<a href="#">117280</a>	2083	993	4	103	heterogeneous nuclear ribonucleoprotein U (Hnrnpu) alternative variant bSep08, mRNA.
<a href="#">Hnrnpu</a>	<a href="#">Hnrnpu.cSep08</a>	<a href="#">117280</a>	922	834	2	63	heterogeneous nuclear ribonucleoprotein U (Hnrnpu) alternative variant cSep08, mRNA.
<a href="#">Hnrnpul1</a>	<a href="#">Hnrnpul1.bSep08</a>	<a href="#">361522</a>	5721	908	4	274	heterogeneous nuclear ribonucleoprotein U-like 1 (Hnrnpul1) alternative variant bSep08, mRNA.
<a href="#">Hnrnpul1</a>	<a href="#">Hnrnpul1.cSep08</a>	<a href="#">361522</a>	6088	361	3	120	heterogeneous nuclear ribonucleoprotein U-like 1 (Hnrnpul1) alternative variant cSep08, mRNA.
<a href="#">Hnrnpul1</a>	<a href="#">Hnrnpul1.dSep08</a>	<a href="#">361522</a>	3829	392	2	93	heterogeneous nuclear ribonucleoprotein U-like 1 (10.7 kD) (Hnrnpul1) alternative variant dSep08, mRNA.
<a href="#">Hnrnpul1</a>	<a href="#">Hnrnpul1.eSep08</a>	<a href="#">361522</a>	1515	1416	2	100	heterogeneous nuclear ribonucleoprotein U-like 1 (10.5 kD) (Hnrnpul1) alternative variant eSep08, mRNA.
<a href="#">Hnrnpul2</a>	<a href="#">Hnrnpul2.aSep08</a>	<a href="#">309197</a>	6213	899	7	294	heterogeneous nuclear ribonucleoprotein U-like 2 (Hnrnpul2) alternative variant aSep08, mRNA.
<a href="#">Hnrnpul2</a>	<a href="#">Hnrnpul2.bSep08</a>	<a href="#">309197</a>	2207	1028	3	140	heterogeneous nuclear ribonucleoprotein U-like 2 (Hnrnpul2) alternative variant bSep08, mRNA.
<a href="#">Hnrpd</a>	<a href="#">Hnrpd.eSep08</a>	<a href="#">79256</a>	3148	639	4	166	heterogeneous nuclear ribonucleoprotein D (Hnrpd) alternative variant eSep08, mRNA.
<a href="#">Hnrpd</a>	<a href="#">Hnrpd.fSep08</a>	<a href="#">79256</a>	2284	675	3	120	heterogeneous nuclear ribonucleoprotein D (Hnrpd) alternative variant fSep08, mRNA.
<a href="#">Hnrpd</a>	<a href="#">Hnrpd.gSep08</a>	<a href="#">79256</a>	3135	2044	2	54	heterogeneous nuclear ribonucleoprotein D (Hnrpd) alternative variant gSep08, mRNA.
<a href="#">Hnrpd</a>	<a href="#">Hnrpd.hSep08</a>	<a href="#">79256</a>	707	269	2	35	heterogeneous nuclear ribonucleoprotein D (Hnrpd) alternative variant hSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.aSep08</a>	<a href="#">305178</a>	4140	1397	9	333	heterogeneous nuclear ribonucleoprotein D-like (Hnrpdl) alternative variant aSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.cSep08</a>	<a href="#">305178</a>	3763	2747	6	270	heterogeneous nuclear ribonucleoprotein D-like (30.1 kD) (Hnrpdl) alternative variant cSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.dSep08</a>	<a href="#">305178</a>	4707	2482	8	270	heterogeneous nuclear ribonucleoprotein D-like (30.1 kD) (Hnrpdl) alternative variant dSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.eSep08</a>	<a href="#">305178</a>	476	388	2	98	heterogeneous nuclear ribonucleoprotein D-like (Hnrpdl) alternative variant eSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.fSep08</a>	<a href="#">305178</a>	917	812	2	66	heterogeneous nuclear ribonucleoprotein D-like (Hnrpdl) alternative variant fSep08, mRNA.
<a href="#">Hnrpdl</a>	<a href="#">Hnrpdl.gSep08</a>	<a href="#">305178</a>	988	642	2	61	heterogeneous nuclear ribonucleoprotein D-like (Hnrpdl) alternative variant gSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.aSep08</a>	<a href="#">140931</a>	9246	2130	13	496	heterogeneous nuclear ribonucleoprotein (Hnrph1) alternative variant aSep08, mRNA.

<a href="#">Hnrph1</a>	<a href="#">Hnrph1.cSep08</a>	<a href="#">140931</a>	6381	1156	10	306	heterogeneous nuclear ribonucleoprotein (Hnrph1) alternative variant cSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.dSep08</a>	<a href="#">140931</a>	4796	1772	8	248	heterogeneous nuclear ribonucleoprotein (Hnrph1) alternative variant dSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.eSep08</a>	<a href="#">140931</a>	4082	1289	8	226	heterogeneous nuclear ribonucleoprotein (24.1 kD) (Hnrph1) alternative variant eSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.fSep08</a>	<a href="#">140931</a>	8723	2187	11	203	heterogeneous nuclear ribonucleoprotein H1 CRA b (Hnrph1) alternative variant fSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.gSep08</a>	<a href="#">140931</a>	3047	1146	4	161	heterogeneous nuclear ribonucleoprotein H1 CRA b (Hnrph1) alternative variant gSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.hSep08</a>	<a href="#">140931</a>	2087	382	3	115	heterogeneous nuclear ribonucleoprotein H1 CRA a (Hnrph1) alternative variant hSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.iSep08</a>	<a href="#">140931</a>	1365	562	3	93	heterogeneous nuclear ribonucleoprotein (Hnrph1) alternative variant iSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.jSep08</a>	<a href="#">140931</a>	2425	679	3	77	putative protein (7.4 kD) (Hnrph1) alternative variant jSep08, mRNA.
<a href="#">Hnrph1</a>	<a href="#">Hnrph1.kSep08</a>	<a href="#">140931</a>	2102	1696	2	39	putative protein (Hnrph1) alternative variant kSep08, mRNA.
<a href="#">Hnrpll</a>	<a href="#">Hnrpll.aSep08</a>	<a href="#">313842</a>	31027	2967	13	661	heterogeneous nuclear ribonucleoprotein L-like (Hnrpll) alternative variant aSep08, mRNA.
<a href="#">Hnrpll</a>	<a href="#">Hnrpll.bSep08</a>	<a href="#">313842</a>	10428	734	3	244	heterogeneous nuclear ribonucleoprotein L-like (Hnrpll) alternative variant bSep08, mRNA.
<a href="#">Hnrpll</a>	<a href="#">Hnrpll.cSep08</a>	<a href="#">313842</a>	3576	721	4	135	heterogeneous nuclear ribonucleoprotein L-like (Hnrpll) alternative variant cSep08, mRNA.
<a href="#">Hnrpm</a>	<a href="#">Hnrpm.aSep08</a>	<a href="#">116655</a>	19207	1683	9	472	heterogeneous nuclear ribonucleoprotein M (Hnrpm) alternative variant aSep08, mRNA.
<a href="#">Hnrpm</a>	<a href="#">Hnrpm.bSep08</a>	<a href="#">116655</a>	2888	701	1	63	heterogeneous nuclear ribonucleoprotein M (7.1 kD) (Hnrpm) alternative variant bSep08, mRNA.
<a href="#">Hnt</a>	<a href="#">Hnt.bSep08</a>	<a href="#">50864</a>	2240	333	2	42	neurotrimin (Hnt) alternative variant bSep08, mRNA.
<a href="#">Homeobox.1</a>	<a href="#">Homeobox.1.aSep08</a>		6029	614	5	82	putative mitochondrial protein (8.8 kD) (Homeobox.1) alternative variant aSep08, mRNA.
<a href="#">Homeobox.1</a>	<a href="#">Homeobox.1.cSep08</a>		6067	432	4	72	otx2 (Homeobox.1) alternative variant cSep08, mRNA.
<a href="#">Homeobox.1</a>	<a href="#">Homeobox.1.dSep08</a>		5815	388	3	50	otx2 (Homeobox.1) alternative variant dSep08, mRNA.
<a href="#">Homeobox.7</a>	<a href="#">Homeobox.7.aSep08</a>		8062	580		193	zinc finger 4 (Homeobox.7) mRNA.
<a href="#">Homeobox.9</a>	<a href="#">Homeobox.9.aSep08</a>		3585	625		199	iroquois protein (Homeobox.9) mRNA.
<a href="#">Homer2</a>	<a href="#">Homer2.bSep08</a>	<a href="#">29547</a>	30933	837	7	278	homer homolog 2 (Drosophila) (Homer2) alternative variant bSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.aSep08</a>	<a href="#">29548</a>	2273	1796	5	474	putative protein (Homer3) alternative variant aSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.bSep08</a>	<a href="#">29548</a>	8169	1210	9	369	homer homolog 3 CRA a (Homer3) alternative variant bSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.dSep08</a>	<a href="#">29548</a>	4129	965	6	145	homer homolog 3 CRA b (16.4 kD) (Homer3) alternative variant dSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.fSep08</a>	<a href="#">29548</a>	3323	949	4	119	homer homolog 3 (13.3 kD) (Homer3) alternative variant fSep08, mRNA.

<a href="#">Homer3</a>	<a href="#">Homer3.gSep08</a>	<a href="#">29548</a>	2750	614	4	104	homer homolog 3 (Homer3) alternative variant gSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.hSep08</a>	<a href="#">29548</a>	2162	669	3	94	homer homolog 3 CRA e (Homer3) alternative variant hSep08, mRNA.
<a href="#">Homer3</a>	<a href="#">Homer3.iSep08</a>	<a href="#">29548</a>	1073	463	3	88	homer homolog 3 CRA b (Homer3) alternative variant iSep08, mRNA.
<a href="#">Hook1</a>	<a href="#">Hook1.bSep08</a>	<a href="#">313370</a>	11060	1756	1	488	hook homolog 1 (Drosophila) (Hook1) alternative variant bSep08, mRNA.
<a href="#">Hook2</a>	<a href="#">Hook2.aSep08</a>	<a href="#">304669</a>	4522	1042	8	345	hook homolog 2 (Drosophila) (Hook2) alternative variant aSep08, mRNA.
<a href="#">Hook2</a>	<a href="#">Hook2.bSep08</a>	<a href="#">304669</a>	4246	866	7	288	hook homolog 2 (Drosophila) (Hook2) alternative variant bSep08, mRNA.
<a href="#">Hook2</a>	<a href="#">Hook2.cSep08</a>	<a href="#">304669</a>	2163	564	6	177	hook homolog 2 (Drosophila) (20.7 kD) (Hook2) alternative variant cSep08, mRNA.
<a href="#">Hook3</a>	<a href="#">Hook3.aSep08</a>	<a href="#">306548</a>	85818	2494		701	hook homolog 3 (Drosophila) (Hook3) alternative variant aSep08, mRNA.
<a href="#">Hopx</a>	<a href="#">Hopx.aSep08</a>	<a href="#">171160</a>	7000	418	2	110	HOP homeobox (Hopx) alternative variant aSep08, mRNA.
<a href="#">Hopx</a>	<a href="#">Hopx.bSep08</a>	<a href="#">171160</a>	26582	870	3	99	HOP homeobox (11.2 kD) (Hopx) alternative variant bSep08, mRNA.
<a href="#">Hopx</a>	<a href="#">Hopx.cSep08</a>	<a href="#">171160</a>	27830	965	4	87	HOP homeobox (Hopx) alternative variant cSep08, mRNA.
<a href="#">Hopx</a>	<a href="#">Hopx.eSep08</a>	<a href="#">171160</a>	25999	426	4	70	HOP homeobox (7.4 kD) (Hopx) alternative variant eSep08, mRNA.
<a href="#">Hox9_act.0</a>	<a href="#">Hox9_act.0.aSep08</a>		2804	1046		246	homeobox protein (Hox9_act.0) mRNA.
<a href="#">Hoxa1</a>	<a href="#">Hoxa1.bSep08</a>	<a href="#">25607</a>	1519	830	3	128	homeo box A1 (14.4 kD) (Hoxa1) alternative variant bSep08, mRNA.
<a href="#">Hoxa5</a>	<a href="#">Hoxa5.aSep08</a>	<a href="#">79241</a>	2904	1947	2	375	homeo box A5 (Hoxa5) alternative variant aSep08, mRNA.
<a href="#">Hoxa5</a>	<a href="#">Hoxa5.bSep08</a>	<a href="#">79241</a>	1523	1032	2	126	homeo box A5 (Hoxa5) alternative variant bSep08, mRNA.
<a href="#">Hoxa5</a>	<a href="#">Hoxa5.cSep08</a>	<a href="#">79241</a>	21172	727	3	72	homeo box A5 (Hoxa5) alternative variant cSep08, mRNA.
<a href="#">Hoxa5</a>	<a href="#">Hoxa5.dSep08</a>	<a href="#">79241</a>	8929	591	2	23	homeo box A5 (2.6 kD) (Hoxa5) alternative variant dSep08, mRNA.
<a href="#">Hoxa9</a>	<a href="#">Hoxa9.aSep08</a>	<a href="#">500126</a>	1734	585	1	181	homeobox A9 (Hoxa9) alternative variant aSep08, mRNA.
<a href="#">Hoxa9</a>	<a href="#">Hoxa9.bSep08</a>	<a href="#">500126</a>	2237	915	2	138	homeobox A9 (16.6 kD) (Hoxa9) alternative variant bSep08, mRNA.
<a href="#">Hoxa10</a>	<a href="#">Hoxa10.aSep08</a>	<a href="#">368057</a>	3015	1844	2	175	homeo box A10 (Hoxa10) alternative variant aSep08, mRNA.
<a href="#">Hoxa10</a>	<a href="#">Hoxa10.bSep08</a>	<a href="#">368057</a>	1196	961	2	136	homeo box A10 (Hoxa10) alternative variant bSep08, mRNA.
<a href="#">Hoxa10</a>	<a href="#">Hoxa10.dSep08</a>	<a href="#">368057</a>	8589	1061	2	94	homeo box A10 (11.5 kD) (Hoxa10) alternative variant dSep08, mRNA.
<a href="#">Hoxa10</a>	<a href="#">Hoxa10.eSep08</a>	<a href="#">368057</a>	8414	717	2	79	homeo box A10 (8.8 kD) (Hoxa10) alternative variant eSep08, mRNA.
<a href="#">Hoxa13</a>	<a href="#">Hoxa13.aSep08</a>	<a href="#">500129</a>	1378	654		141	homeo box A13 (Hoxa13) mRNA.
<a href="#">Hoxb1</a>	<a href="#">Hoxb1.aSep08</a>	<a href="#">303491</a>	1224	788		239	homeo box B1 (Hoxb1) mRNA.
<a href="#">Hoxb2</a>	<a href="#">Hoxb2.aSep08</a>	<a href="#">303489</a>	1551	810	2	270	homeo box B2 (Hoxb2) alternative variant aSep08, mRNA.
<a href="#">Hoxb3</a>	<a href="#">Hoxb3.bSep08</a>	<a href="#">303488</a>	14383	2405	4	428	homeo box B3 (44.1 kD) (Hoxb3) alternative variant bSep08, mRNA.

<a href="#">Hoxb3</a>	<a href="#">Hoxb3.cSep08</a>	<a href="#">303488</a>	52740	599	4	129	homeo box B3 (Hoxb3) alternative variant cSep08, mRNA.
<a href="#">Hoxb6</a>	<a href="#">Hoxb6.bSep08</a>	<a href="#">497986</a>	2001	911	2	219	homeo box B6 (Hoxb6) alternative variant bSep08, mRNA.
<a href="#">Hoxc6</a>	<a href="#">Hoxc6.aSep08</a>	<a href="#">252885</a>	2359	1635	3	421	homeo box C6 (Hoxc6) alternative variant aSep08, mRNA.
<a href="#">Hoxc10</a>	<a href="#">Hoxc10.aSep08</a>	<a href="#">315338</a>	4098	739	2	181	homeo box C10 (Hoxc10) alternative variant aSep08, mRNA.
<a href="#">Hoxc10</a>	<a href="#">Hoxc10.bSep08</a>	<a href="#">315338</a>	4465	251	3	68	homeo box C10 (Hoxc10) alternative variant bSep08, mRNA.
<a href="#">Hoxd9</a>	<a href="#">Hoxd9.aSep08</a>	<a href="#">688999</a>	2387	1833		366	homeo box D9 (Hoxd9) alternative variant aSep08, mRNA.
<a href="#">Hoxd10</a>	<a href="#">Hoxd10.aSep08</a>	<a href="#">303991</a>	11662	758	2	96	homeo box D10 (Hoxd10) alternative variant aSep08, mRNA.
<a href="#">Hoxd10</a>	<a href="#">Hoxd10.bSep08</a>	<a href="#">303991</a>	4139	1229	1	70	homeo box D10 (8.1 kD) (Hoxd10) alternative variant bSep08, mRNA.
<a href="#">Hp</a>	<a href="#">Hp.bSep08</a>	<a href="#">24464</a>	4165	871	5	260	haptoglobin (Hp) alternative variant bSep08, mRNA.
<a href="#">Hp</a>	<a href="#">Hp.cSep08</a>	<a href="#">24464</a>	3707	332	5	110	haptoglobin (Hp) alternative variant cSep08, mRNA.
<a href="#">Hp</a>	<a href="#">Hp.eSep08</a>	<a href="#">24464</a>	1127	259	3	86	haptoglobin (Hp) alternative variant eSep08, mRNA.
<a href="#">Hp</a>	<a href="#">Hp.fSep08</a>	<a href="#">24464</a>	2331	358	3	67	haptoglobin (7.5 kD) (Hp) alternative variant fSep08, complete mRNA.
<a href="#">Hp1bp3</a>	<a href="#">Hp1bp3.bSep08</a>	<a href="#">313647</a>	16270	785	7	261	heterochromatin protein 1, binding protein 3 (Hp1bp3) alternative variant bSep08, mRNA.
<a href="#">Hp1bp3</a>	<a href="#">Hp1bp3.cSep08</a>	<a href="#">313647</a>	13622	1337	6	238	heterochromatin protein 1, binding protein 3 (26.3 kD) (Hp1bp3) alternative variant cSep08, mRNA.
<a href="#">Hp1bp3</a>	<a href="#">Hp1bp3.dSep08</a>	<a href="#">313647</a>	5336	2074	4	229	heterochromatin protein 1, binding protein 3 (Hp1bp3) alternative variant dSep08, mRNA.
<a href="#">Hp1bp3</a>	<a href="#">Hp1bp3.fSep08</a>	<a href="#">313647</a>	4612	502	3	71	heterochromatin protein 1, binding protein 3 (Hp1bp3) alternative variant fSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.aSep08</a>	<a href="#">50871</a>	116414	1721	5	193	hippocalcin-like 1 CRA b (22.3 kD) (Hpcal1) alternative variant aSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.cSep08</a>	<a href="#">50871</a>	111934	544	2	97	CRA a like (Hpcal1) alternative variant cSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.dSep08</a>	<a href="#">50871</a>	78828	768	4	93	CRA a like (10.0 kD) (Hpcal1) alternative variant dSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.eSep08</a>	<a href="#">50871</a>	38274	368	4	76	putative protein (Hpcal1) alternative variant eSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.fSep08</a>	<a href="#">50871</a>	109232	783	7	81	putative protein (Hpcal1) alternative variant fSep08, mRNA.
<a href="#">Hpcal1</a>	<a href="#">Hpcal1.gSep08</a>	<a href="#">50871</a>	72477	387	4	52	putative protein (Hpcal1) alternative variant gSep08, mRNA.
<a href="#">Hpd</a>	<a href="#">Hpd.aSep08</a>	<a href="#">29531</a>	10223	1362	10	401	4-hydroxyphenylpyruvic acid dioxygenase (Hpd) alternative variant aSep08, mRNA.
<a href="#">Hpd</a>	<a href="#">Hpd.bSep08</a>	<a href="#">29531</a>	7925	868	9	289	4-hydroxyphenylpyruvic acid dioxygenase (Hpd) alternative variant bSep08, mRNA.
<a href="#">Hpd</a>	<a href="#">Hpd.cSep08</a>	<a href="#">29531</a>	6263	726	8	241	4-hydroxyphenylpyruvic acid dioxygenase (Hpd) alternative variant cSep08, mRNA.
<a href="#">Hpd</a>	<a href="#">Hpd.eSep08</a>	<a href="#">29531</a>	4715	777	5	104	4-hydroxyphenylpyruvic acid dioxygenase (Hpd) alternative variant eSep08, mRNA.
<a href="#">Hpn</a>	<a href="#">Hpn.bSep08</a>	<a href="#">29135</a>	11578	1054	1	351	hepsin (Hpn) alternative variant bSep08, mRNA.
<a href="#">Hpn</a>	<a href="#">Hpn.cSep08</a>	<a href="#">29135</a>	4634	1147	4	279	hepsin (Hpn) alternative variant cSep08, mRNA.

<a href="#">Hprt1</a>	<a href="#">Hprt1.aSep08</a>	<a href="#">24465</a>	43060	4356	11	364	PHD finger protein 6 (41.1 kD) (Hprt1) alternative variant aSep08, mRNA.
<a href="#">Hprt1</a>	<a href="#">Hprt1.bSep08</a>	<a href="#">24465</a>	43070	1802	7	351	PHD finger protein 6 (Hprt1) alternative variant bSep08, mRNA.
<a href="#">Hprt1</a>	<a href="#">Hprt1.dSep08</a>	<a href="#">24465</a>	30551	720	8	197	hypoxanthine phosphoribosyltransferase (Hprt1) alternative variant dSep08, mRNA.
<a href="#">Hprt1</a>	<a href="#">Hprt1.eSep08</a>	<a href="#">24465</a>	26666	689	6	161	hypoxanthine phosphoribosyltransferase (Hprt1) alternative variant eSep08, mRNA.
<a href="#">Hprt1</a>	<a href="#">Hprt1.fSep08</a>	<a href="#">24465</a>	5292	496	4	92	PHD finger protein 6 (10.3 kD) (Hprt1) alternative variant fSep08, mRNA.
<a href="#">Hprt1</a>	<a href="#">Hprt1.gSep08</a>	<a href="#">24465</a>	91856	300	3	39	putative protein (Hprt1) alternative variant gSep08, mRNA.
<a href="#">Hps1</a>	<a href="#">Hps1.aSep08</a>	<a href="#">114638</a>	20383	1806	16	531	hermansky-Pudlak syndrome 1 homolog (human) (Hps1) alternative variant aSep08, mRNA.
<a href="#">Hps1</a>	<a href="#">Hps1.bSep08</a>	<a href="#">114638</a>	12809	737	7	191	hermansky-Pudlak syndrome 1 homolog (human) (Hps1) alternative variant bSep08, mRNA.
<a href="#">Hps1</a>	<a href="#">Hps1.cSep08</a>	<a href="#">114638</a>	2854	839	2	81	hermansky-Pudlak syndrome 1 homolog (human) (Hps1) alternative variant cSep08, mRNA.
<a href="#">Hps1</a>	<a href="#">Hps1.eSep08</a>	<a href="#">114638</a>	1580	593	2	50	hermansky-Pudlak syndrome 1 homolog (human) (Hps1) alternative variant eSep08, mRNA.
<a href="#">Hps3</a>	<a href="#">Hps3.bSep08</a>	<a href="#">310288</a>	15264	1345	5	79	hermansky-Pudlak syndrome 3 homolog (human) (9.0 kD) (Hps3) alternative variant bSep08, mRNA.
<a href="#">Hps4</a>	<a href="#">Hps4.bSep08</a>	<a href="#">304555</a>	6250	1569	4	351	hermansky-Pudlak syndrome 4 homolog (human) (37.6 kD) (Hps4) alternative variant bSep08, mRNA.
<a href="#">Hps4</a>	<a href="#">Hps4.cSep08</a>	<a href="#">304555</a>	8048	1539	6	323	hermansky-Pudlak syndrome 4 homolog (human) (34.3 kD) (Hps4) alternative variant cSep08, mRNA.
<a href="#">Hpse</a>	<a href="#">Hpse.bSep08</a>	<a href="#">64537</a>	7949	1888	3	141	heparanase (Hpse) alternative variant bSep08, mRNA.
<a href="#">Hpx</a>	<a href="#">Hpx.bSep08</a>	<a href="#">58917</a>	5825	1501	7	445	hemopexin (Hpx) alternative variant bSep08, mRNA.
<a href="#">Hpx</a>	<a href="#">Hpx.cSep08</a>	<a href="#">58917</a>	6585	912	7	299	hemopexin (Hpx) alternative variant cSep08, mRNA.
<a href="#">Hpx</a>	<a href="#">Hpx.dSep08</a>	<a href="#">58917</a>	958	877	2	119	hemopexin (13.0 kD) (Hpx) alternative variant dSep08, mRNA.
<a href="#">Hr</a>	<a href="#">Hr.bSep08</a>	<a href="#">60563</a>	754	377	3	125	hairless (Hr) alternative variant bSep08, mRNA.
<a href="#">Hr</a>	<a href="#">Hr.cSep08</a>	<a href="#">60563</a>	1873	379	3	114	hairless (Hr) alternative variant cSep08, mRNA.
<a href="#">Hr</a>	<a href="#">Hr.dSep08</a>	<a href="#">60563</a>	849	532	3	107	hairless (Hr) alternative variant dSep08, mRNA.
<a href="#">Hr</a>	<a href="#">Hr.eSep08</a>	<a href="#">60563</a>	915	283	2	48	putative protein (Hr) alternative variant eSep08, mRNA.
<a href="#">Hras</a>	<a href="#">Hras.cSep08</a>	<a href="#">293621</a>	6921	695	3	63	harvey rat sarcoma virus oncogene (Hras) alternative variant cSep08, mRNA.
<a href="#">Hrasls</a>	<a href="#">Hrasls.bSep08</a>	<a href="#">288025</a>	16136	482	1	61	HRAS-like suppressor (7.0 kD) (Hrasls) alternative variant bSep08, mRNA.
<a href="#">Hrbl</a>	<a href="#">Hrbl.aSep08</a>	<a href="#">304375</a>	14621	1927	7	510	HIV-1 Rev binding protein-like (Hrbl) alternative variant aSep08, mRNA.
<a href="#">Hrbl</a>	<a href="#">Hrbl.cSep08</a>	<a href="#">304375</a>	2085	451	3	95	HIV-1 Rev binding protein-like (Hrbl) alternative variant cSep08, mRNA.
<a href="#">Hrg</a>	<a href="#">Hrg.bSep08</a>	<a href="#">171016</a>	33424	1501	7	497	histidine-rich glycoprotein (Hrg) alternative variant bSep08, mRNA.

<a href="#">Hrh3</a>	<a href="#">Hrh3.aSep08</a>	<a href="#">85268</a>	5416	2237	3	379	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant aSep08, mRNA.
<a href="#">Hrh3</a>	<a href="#">Hrh3.aSep08</a>	<a href="#">690396</a>	5416	2237	3	379	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant aSep08, mRNA.
<a href="#">Hrh3</a>	<a href="#">Hrh3.bSep08</a>	<a href="#">85268</a>	1163	992	2	245	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant bSep08, mRNA.
<a href="#">Hrh3</a>	<a href="#">Hrh3.bSep08</a>	<a href="#">690396</a>	1163	992	2	245	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant bSep08, mRNA.
<a href="#">Hrh3</a>	<a href="#">Hrh3.cSep08</a>	<a href="#">85268</a>	3202	888	3	184	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant cSep08, mRNA.
<a href="#">Hrh3</a>	<a href="#">Hrh3.cSep08</a>	<a href="#">690396</a>	3202	888	3	184	histamine receptor H3 and similar to SR protein related family member (rsr-1) (Hrh3) alternative variant cSep08, mRNA.
<a href="#">HRM.0</a>	<a href="#">HRM.0.aSep08</a>		3514	634		210	brain-specific angiogenesis inhibitor 1 (HRM.0) mRNA.
<a href="#">HRM.1</a>	<a href="#">HRM.1.aSep08</a>		18057	388		129	vasoactive intestinal peptide receptor 2 (HRM.1) mRNA.
<a href="#">HRM.2</a>	<a href="#">HRM.2.aSep08</a>		1970	506		168	cadherin EGF LAG seven-pass G-type receptor (HRM.2) mRNA.
<a href="#">Hrpap20</a>	<a href="#">Hrpap20.bSep08</a>	<a href="#">362495</a>	5268	2410	2	136	hormone-regulated proliferation associated protein 20 (15.7 kD) (Hrpap20) alternative variant bSep08, complete mRNA.
<a href="#">Hrpap20</a>	<a href="#">Hrpap20.cSep08</a>	<a href="#">362495</a>	3922	952	3	115	hormone-regulated proliferation associated protein 20 (13.2 kD) (Hrpap20) alternative variant cSep08, mRNA.
<a href="#">Hs1bp3</a>	<a href="#">Hs1bp3.aSep08</a>	<a href="#">313950</a>	20100	2414		192	HCLS1 binding protein 3 (Hs1bp3) mRNA.
<a href="#">Hs2st1</a>	<a href="#">Hs2st1.aSep08</a>	<a href="#">292155</a>	32060	2693	4	259	heparan sulfate 2-O-sulfotransferase 1 (Hs2st1) alternative variant aSep08, mRNA.
<a href="#">Hs2st1</a>	<a href="#">Hs2st1.bSep08</a>	<a href="#">292155</a>	3523	658	1	134	heparan sulfate 2-O-sulfotransferase 1 (Hs2st1) alternative variant bSep08, mRNA.
<a href="#">Hs3st5</a>	<a href="#">Hs3st5.bSep08</a>	<a href="#">294449</a>	2338	560	2	43	heparan sulfate (glucosamine) 3-O-sulfotransferase 5 (Hs3st5) alternative variant bSep08, mRNA.
<a href="#">Hs3st6</a>	<a href="#">Hs3st6.bSep08</a>	<a href="#">684979</a>	921	406	1	59	heparan sulfate (glucosamine) 3-O-sulfotransferase 6 (Hs3st6) alternative variant bSep08, mRNA.
<a href="#">HS6ST.0</a>	<a href="#">HS6ST.0.aSep08</a>		1363	568		189	heparan sulfate 6-O-sulfotransferase 2 (HS6ST.0) mRNA.
<a href="#">HSA.0</a>	<a href="#">HSA.0.aSep08</a>		6195	589		196	CBP activator (HSA.0) mRNA.
<a href="#">HSA.1</a>	<a href="#">HSA.1.aSep08</a>		13715	584	5	194	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 2 (HSA.1) alternative variant aSep08, mRNA.
<a href="#">Hsbp1</a>	<a href="#">Hsbp1.bSep08</a>	<a href="#">286899</a>	1042	744	2	75	heat shock factor binding protein 1 (9.0 kD) (Hsbp1) alternative variant bSep08, mRNA.
<a href="#">Hsbp1</a>	<a href="#">Hsbp1.cSep08</a>	<a href="#">286899</a>	5084	1227	5	76	heat shock factor binding protein 1 (8.6 kD) (Hsbp1) alternative variant cSep08, complete mRNA.
<a href="#">HSBP1.1</a>	<a href="#">HSBP1.1.aSep08</a>		5801	440		72	CRA a (8.1 kD) (HSBP1.1) mRNA.



<a href="#">Hscb</a>	<a href="#">Hscb.bSep08</a>	<a href="#">360826</a>	10339	781	6	152	HscB iron-sulfur cluster co-chaperone homolog (E. coli) (Hscb) alternative variant bSep08, mRNA.
<a href="#">Hscb</a>	<a href="#">Hscb.cSep08</a>	<a href="#">360826</a>	10301	593	5	146	HscB iron-sulfur cluster co-chaperone homolog (E. coli) (Hscb) alternative variant cSep08, mRNA.
<a href="#">Hsd3b</a>	<a href="#">Hsd3b.aSep08</a>	<a href="#">24470</a>	18753	687	1	228	steroid delta-isomerase, 3 beta (Hsd3b) alternative variant aSep08, mRNA.
<a href="#">Hsd3b</a>	<a href="#">Hsd3b.bSep08</a>	<a href="#">24470</a>	18840	847	1	142	steroid delta-isomerase, 3 beta (Hsd3b) alternative variant bSep08, mRNA.
<a href="#">Hsd3b7</a>	<a href="#">Hsd3b7.bSep08</a>	<a href="#">246211</a>	2375	992	3	330	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7 (Hsd3b7) alternative variant bSep08, mRNA.
<a href="#">Hsd3b7</a>	<a href="#">Hsd3b7.cSep08</a>	<a href="#">246211</a>	2375	989	3	329	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7 (Hsd3b7) alternative variant cSep08, mRNA.
<a href="#">Hsd11b1</a>	<a href="#">Hsd11b1.aSep08</a>	<a href="#">25116</a>	50064	1216	4	288	hydroxysteroid 11-beta dehydrogenase 1 (31.9 kD) (Hsd11b1) alternative variant aSep08, mRNA.
<a href="#">Hsd11b1</a>	<a href="#">Hsd11b1.cSep08</a>	<a href="#">25116</a>	24578	835	2	278	hydroxysteroid 11-beta dehydrogenase 1 (Hsd11b1) alternative variant cSep08, mRNA.
<a href="#">Hsd11b1</a>	<a href="#">Hsd11b1.dSep08</a>	<a href="#">25116</a>	28495	874	3	201	hydroxysteroid 11-beta dehydrogenase 1 (Hsd11b1) alternative variant dSep08, mRNA.
<a href="#">Hsd11b2</a>	<a href="#">Hsd11b2.cSep08</a>	<a href="#">25117</a>	1421	1306	2	95	hydroxysteroid 11-beta dehydrogenase 2 (Hsd11b2) alternative variant cSep08, mRNA.
<a href="#">Hsd17b2</a>	<a href="#">Hsd17b2.bSep08</a>	<a href="#">79243</a>	21437	1019	3	144	hydroxysteroid (17-beta) dehydrogenase 2 (15.9 kD) (Hsd17b2) alternative variant bSep08, mRNA.
<a href="#">Hsd17b2</a>	<a href="#">Hsd17b2.cSep08</a>	<a href="#">79243</a>	43032	687	3	112	hydroxysteroid (17-beta) dehydrogenase 2 (Hsd17b2) alternative variant cSep08, mRNA.
<a href="#">Hsd17b4</a>	<a href="#">Hsd17b4.bSep08</a>	<a href="#">79244</a>	13408	524	7	174	hydroxysteroid (17-beta) dehydrogenase 4 (Hsd17b4) alternative variant bSep08, mRNA.
<a href="#">Hsd17b4</a>	<a href="#">Hsd17b4.cSep08</a>	<a href="#">79244</a>	8383	1615	2	41	hydroxysteroid (17-beta) dehydrogenase 4 (4.7 kD) (Hsd17b4) alternative variant cSep08, mRNA.
<a href="#">Hsd17b6</a>	<a href="#">Hsd17b6.bSep08</a>	<a href="#">286964</a>	3345	606		96	hydroxysteroid (17-beta) dehydrogenase 6 (Hsd17b6) alternative variant bSep08, mRNA.
<a href="#">Hsd17b7</a>	<a href="#">Hsd17b7.bSep08</a>	<a href="#">29540</a>	4578	655	3	66	hydroxysteroid (17-beta) dehydrogenase 7 (Hsd17b7) alternative variant bSep08, mRNA.
<a href="#">Hsd17b7</a>	<a href="#">Hsd17b7.cSep08</a>	<a href="#">29540</a>	802	597	2	38	hydroxysteroid (17-beta) dehydrogenase 7 (Hsd17b7) alternative variant cSep08, mRNA.
<a href="#">Hsd17b10</a>	<a href="#">Hsd17b10.cSep08</a>	<a href="#">63864</a>	1041	684	2	92	hydroxysteroid (17-beta) dehydrogenase 10 (9.2 kD) (Hsd17b10) alternative variant cSep08, mRNA.
<a href="#">Hsd17b11</a>	<a href="#">Hsd17b11.bSep08</a>	<a href="#">289456</a>	31284	791	5	185	hydroxysteroid (17-beta) dehydrogenase 11 (Hsd17b11) alternative variant bSep08, mRNA.
<a href="#">Hsd17b11</a>	<a href="#">Hsd17b11.cSep08</a>	<a href="#">289456</a>	27476	749	5	173	hydroxysteroid (17-beta) dehydrogenase 11 (Hsd17b11) alternative variant cSep08, mRNA.
<a href="#">Hsd17b12</a>	<a href="#">Hsd17b12.bSep08</a>	<a href="#">84013</a>	89900	366	4	115	hydroxysteroid (17-beta) dehydrogenase 12 (Hsd17b12) alternative variant bSep08, mRNA.
<a href="#">Hsd17b12</a>	<a href="#">Hsd17b12.cSep08</a>	<a href="#">84013</a>	1011	740	2	35	hydroxysteroid (17-beta) dehydrogenase 12 (4.2 kD) (Hsd17b12) alternative variant cSep08, mRNA.

<a href="#">Hsd17b13</a>	<a href="#">Hsd17b13.bSep08</a>	<a href="#">305150</a>	2227	730	1	45	hydroxysteroid (17-beta) dehydrogenase 13 (5.4 kD) (Hsd17b13) alternative variant bSep08, mRNA.
<a href="#">Hsd17b14</a>	<a href="#">Hsd17b14.aSep08</a>	<a href="#">691018</a>	9875	673		167	hydroxysteroid (17-beta) dehydrogenase 14 (Hsd17b14) mRNA.
<a href="#">Hsd11</a>	<a href="#">Hsd11.bSep08</a>	<a href="#">361418</a>	9186	518	3	172	hydroxysteroid dehydrogenase like 1 (Hsd11) alternative variant bSep08, mRNA.
<a href="#">Hsf1</a>	<a href="#">Hsf1.aSep08</a>	<a href="#">79245</a>	26698	2157	5	582	heat shock transcription factor 1 (Hsf1) alternative variant aSep08, mRNA.
<a href="#">Hsf1</a>	<a href="#">Hsf1.bSep08</a>	<a href="#">79245</a>	25857	1417	1	218	heat shock transcription factor 1 (Hsf1) alternative variant bSep08, mRNA.
<a href="#">Hsf1</a>	<a href="#">Hsf1.cSep08</a>	<a href="#">79245</a>	986	694	3	109	heat shock transcription factor 1 (Hsf1) alternative variant cSep08, mRNA.
<a href="#">Hsf2</a>	<a href="#">Hsf2.bSep08</a>	<a href="#">64441</a>	1814	552	3	121	heat shock factor 2 (Hsf2) alternative variant bSep08, mRNA.
<a href="#">Hsf2</a>	<a href="#">Hsf2.cSep08</a>	<a href="#">64441</a>	4956	389	4	92	heat shock factor 2 CRA b (Hsf2) alternative variant cSep08, mRNA.
<a href="#">Hsf2bp</a>	<a href="#">Hsf2bp.bSep08</a>	<a href="#">499413</a>	84091	1531	2	192	heat shock transcription factor 2 binding protein (Hsf2bp) alternative variant bSep08, mRNA.
<a href="#">Hsf4</a>	<a href="#">Hsf4.bSep08</a>	<a href="#">291960</a>	3059	883	6	236	heat shock transcription factor 4 (Hsf4) alternative variant bSep08, mRNA.
<a href="#">Hsf4</a>	<a href="#">Hsf4.cSep08</a>	<a href="#">291960</a>	1025	352	3	117	heat shock transcription factor 4 (Hsf4) alternative variant cSep08, mRNA.
<a href="#">Hsf4</a>	<a href="#">Hsf4.dSep08</a>	<a href="#">291960</a>	1312	370	5	111	heat shock transcription factor 4 (Hsf4) alternative variant dSep08, mRNA.
<a href="#">HSP70.0</a>	<a href="#">HSP70.0.aSep08</a>		3815	2135	8	627	heat shock protein (HSP70.0) alternative variant aSep08, mRNA.
<a href="#">HSP70.0</a>	<a href="#">HSP70.0.bSep08</a>		698	613	2	107	shock protein (HSP70.0) alternative variant bSep08, mRNA.
<a href="#">Hsp90aa1</a>	<a href="#">Hsp90aa1.bSep08</a>	<a href="#">299331</a>	690	592	2	157	heat shock protein 90, alpha (cytosolic), class A member 1 (Hsp90aa1) alternative variant bSep08, mRNA.
<a href="#">Hsp90aa1</a>	<a href="#">Hsp90aa1.cSep08</a>	<a href="#">299331</a>	1834	1337	2	123	heat shock protein 90, alpha (cytosolic), class A member 1 (13.8 kD) (Hsp90aa1) alternative variant cSep08, mRNA.
<a href="#">Hsp90ab1</a>	<a href="#">Hsp90ab1.bSep08</a>	<a href="#">301252</a>	4030	1649	8	518	heat shock protein 90kDa alpha (cytosolic), class B member 1 (59.6 kD) (Hsp90ab1) alternative variant bSep08, mRNA.
<a href="#">Hsp90ab1</a>	<a href="#">Hsp90ab1.cSep08</a>	<a href="#">301252</a>	1747	1219	3	293	heat shock protein 90kDa alpha (cytosolic), class B member 1 (Hsp90ab1) alternative variant cSep08, mRNA.
<a href="#">Hsp90ab1</a>	<a href="#">Hsp90ab1.dSep08</a>	<a href="#">301252</a>	1704	940	4	282	heat shock protein 90kDa alpha (cytosolic), class B member 1 (Hsp90ab1) alternative variant dSep08, mRNA.
<a href="#">Hsp90ab1</a>	<a href="#">Hsp90ab1.eSep08</a>	<a href="#">301252</a>	2501	890	4	199	heat shock protein 90kDa alpha (cytosolic), class B member 1 (21.8 kD) (Hsp90ab1) alternative variant eSep08, mRNA.
<a href="#">Hspa2</a>	<a href="#">Hspa2.aSep08</a>	<a href="#">60460</a>	2717	2476	2	662	heat shock protein 2 (Hspa2) alternative variant aSep08, mRNA.
<a href="#">Hspa4</a>	<a href="#">Hspa4.bSep08</a>	<a href="#">266759</a>	5487	507	5	168	heat shock protein 4 CRA c (Hspa4) alternative variant bSep08, mRNA.

<a href="#">Hspa4</a>	<a href="#">Hspa4.cSep08</a>	<a href="#">266759</a>	1371	567	3	131	heat shock protein 4 (14.4 kD) (Hspa4) alternative variant cSep08, mRNA.
<a href="#">Hspa4</a>	<a href="#">Hspa4.dSep08</a>	<a href="#">266759</a>	831	479	2	110	heat shock protein 4 CRA d (12.0 kD) (Hspa4) alternative variant dSep08, mRNA.
<a href="#">Hspa4</a>	<a href="#">Hspa4.eSep08</a>	<a href="#">266759</a>	3172	838	2	95	heat shock protein 4 (Hspa4) alternative variant eSep08, mRNA.
<a href="#">Hspa4</a>	<a href="#">Hspa4.fSep08</a>	<a href="#">266759</a>	1103	401	2	47	heat shock protein 4 CRA d (Hspa4) alternative variant fSep08, mRNA.
<a href="#">Hspa4l</a>	<a href="#">Hspa4l.bSep08</a>	<a href="#">294993</a>	21138	637	1	212	heat shock protein 4 like (Hspa4l) alternative variant bSep08, mRNA.
<a href="#">Hspa5</a>	<a href="#">Hspa5.bSep08</a>	<a href="#">25617</a>	1710	830	2	202	heat shock protein 5 (22.1 kD) (Hspa5) alternative variant bSep08, mRNA.
<a href="#">Hspa5</a>	<a href="#">Hspa5.cSep08</a>	<a href="#">25617</a>	1335	771	1	174	heat shock protein 5 (19.0 kD) (Hspa5) alternative variant cSep08, mRNA.
<a href="#">Hspa9</a>	<a href="#">Hspa9.bSep08</a>	<a href="#">291671</a>	5296	2059	7	281	heat shock 70kDa protein 9 (30.6 kD) (Hspa9) alternative variant bSep08, mRNA.
<a href="#">Hspa9</a>	<a href="#">Hspa9.cSep08</a>	<a href="#">291671</a>	5127	925	2	105	heat shock 70kDa protein 9A (11.6 kD) (Hspa9) alternative variant cSep08, mRNA.
<a href="#">Hspa12b</a>	<a href="#">Hspa12b.aSep08</a>	<a href="#">311427</a>	3845	1196	6	398	heat shock protein 12B (Hspa12b) alternative variant aSep08, mRNA.
<a href="#">Hspa12b</a>	<a href="#">Hspa12b.cSep08</a>	<a href="#">311427</a>	11687	841	8	193	heat shock protein 12B (Hspa12b) alternative variant cSep08, mRNA.
<a href="#">Hspa12b</a>	<a href="#">Hspa12b.dSep08</a>	<a href="#">311427</a>	6070	639	5	95	heat shock protein 12B (Hspa12b) alternative variant dSep08, mRNA.
<a href="#">Hspa12b</a>	<a href="#">Hspa12b.eSep08</a>	<a href="#">311427</a>	4265	605	3	65	heat shock protein 12B (6.9 kD) (Hspa12b) alternative variant eSep08, mRNA.
<a href="#">Hspa14</a>	<a href="#">Hspa14.bSep08</a>	<a href="#">307133</a>	5942	581	6	189	heat shock protein 14 (Hspa14) alternative variant bSep08, mRNA.
<a href="#">Hspa14</a>	<a href="#">Hspa14.cSep08</a>	<a href="#">307133</a>	2046	1414	2	44	heat shock protein 14 (4.8 kD) (Hspa14) alternative variant cSep08, mRNA.
<a href="#">Hspb1</a>	<a href="#">Hspb1.cSep08</a>	<a href="#">24471</a>	696	585	2	84	heat shock protein 1 (9.5 kD) (Hspb1) alternative variant cSep08, mRNA.
<a href="#">Hspb2</a>	<a href="#">Hspb2.bSep08</a>	<a href="#">161476</a>	1208	503	2	100	heat shock protein 2 (Hspb2) alternative variant bSep08, mRNA.
<a href="#">Hspb2</a>	<a href="#">Hspb2.cSep08</a>	<a href="#">161476</a>	1232	628	2	61	heat shock protein 2 (6.6 kD) (Hspb2) alternative variant cSep08, mRNA.
<a href="#">Hspb6</a>	<a href="#">Hspb6.bSep08</a>	<a href="#">192245</a>	2915	1289	1	127	heat shock protein, alpha-crystallin-related, B6 (13.8 kD) (Hspb6) alternative variant bSep08, complete mRNA.
<a href="#">Hspb7</a>	<a href="#">Hspb7.bSep08</a>	<a href="#">50565</a>	1949	711	1	95	heat shock protein family, member 7 (cardiovascular) (Hspb7) alternative variant bSep08, mRNA.
<a href="#">Hspb8</a>	<a href="#">Hspb8.bSep08</a>	<a href="#">113906</a>	26854	588	2	128	heat shock protein 8 (Hspb8) alternative variant bSep08, mRNA.
<a href="#">Hspb8</a>	<a href="#">Hspb8.cSep08</a>	<a href="#">113906</a>	7432	529	2	27	heat shock protein 8 (Hspb8) alternative variant cSep08, mRNA.
<a href="#">Hspb11</a>	<a href="#">Hspb11.bSep08</a>	<a href="#">685284</a>	29374	707	6	141	heat shock protein family B (small), member 11 (Hspb11) alternative variant bSep08, mRNA.

<a href="#">Hspb11</a>	<a href="#">Hspb11.cSep08</a>	<a href="#">685284</a>	19312	329	4	109	heat shock protein family B (small), member 11 (Hspb11) alternative variant cSep08, mRNA.
<a href="#">Hspb11</a>	<a href="#">Hspb11.dSep08</a>	<a href="#">685284</a>	20831	306	3	67	heat shock protein family B (small), member 11 (Hspb11) alternative variant dSep08, mRNA.
<a href="#">Hsppb1</a>	<a href="#">Hsppb1.aSep08</a>	<a href="#">246146</a>	23489	1431	6	357	hsp70-interacting protein (39.2 kD) (Hsppb1) alternative variant aSep08, mRNA.
<a href="#">Hsppb1</a>	<a href="#">Hsppb1.cSep08</a>	<a href="#">246146</a>	11360	766	3	211	hsp70-interacting protein (Hsppb1) alternative variant cSep08, mRNA.
<a href="#">Hsppb1</a>	<a href="#">Hsppb1.dSep08</a>	<a href="#">246146</a>	6385	963	2	204	hsp70-interacting protein (Hsppb1) alternative variant dSep08, mRNA.
<a href="#">Hsppb1</a>	<a href="#">Hsppb1.eSep08</a>	<a href="#">246146</a>	5884	788	1	194	hsp70-interacting protein (Hsppb1) alternative variant eSep08, mRNA.
<a href="#">Hspc105</a>	<a href="#">Hspc105.aSep08</a>	<a href="#">307897</a>	11168	383		127	NAD(P) dependent steroid dehydrogenase-like (Hspc105) mRNA.
<a href="#">Hspc159</a>	<a href="#">Hspc159.bSep08</a>	<a href="#">360983</a>	3174	544	2	102	galectin-related protein (11.6 kD) (Hspc159) alternative variant bSep08, mRNA.
<a href="#">Hspc159</a>	<a href="#">Hspc159.cSep08</a>	<a href="#">360983</a>	1458	716	3	58	galectin-related protein (Hspc159) alternative variant cSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.bSep08</a>	<a href="#">63868</a>	8063	1367	8	341	heat shock protein 1 (chaperonin) (Hspd1) alternative variant bSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.cSep08</a>	<a href="#">63868</a>	3735	896	6	298	heat shock protein 1 (chaperonin) (Hspd1) alternative variant cSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.dSep08</a>	<a href="#">63868</a>	6597	883	7	272	heat shock protein 1 (chaperonin) (Hspd1) alternative variant dSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.eSep08</a>	<a href="#">63868</a>	1984	597	3	199	heat shock protein 1 (chaperonin) (Hspd1) alternative variant eSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.fSep08</a>	<a href="#">63868</a>	4497	532	4	164	heat shock protein 1 (chaperonin) (Hspd1) alternative variant fSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.gSep08</a>	<a href="#">63868</a>	3566	630	5	160	heat shock protein 1 (chaperonin) (Hspd1) alternative variant gSep08, mRNA.
<a href="#">Hspd1</a>	<a href="#">Hspd1.hSep08</a>	<a href="#">63868</a>	1577	656	2	45	heat shock protein 1 (chaperonin) (Hspd1) alternative variant hSep08, mRNA.
<a href="#">Hspe1</a>	<a href="#">Hspe1.aSep08</a>	<a href="#">25462</a>	2928	502	2	114	heat shock 10 kDa protein 1 (chaperonin 10) (12.5 kD) (Hspe1) alternative variant aSep08, mRNA.
<a href="#">Hspe1</a>	<a href="#">Hspe1.cSep08</a>	<a href="#">25462</a>	2458	535	2	71	heat shock 10 kDa protein 1 (chaperonin 10) (7.5 kD) (Hspe1) alternative variant cSep08, mRNA.
<a href="#">Hsph1</a>	<a href="#">Hsph1.bSep08</a>	<a href="#">288444</a>	1779	371	4	123	heat shock 105kDa/110kDa protein 1 (Hsph1) alternative variant bSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.bSep08</a>	<a href="#">192218</a>	4483	901	9	300	HIV-1 tat interactive protein, homolog (human) (Htatif) alternative variant bSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.cSep08</a>	<a href="#">192218</a>	2021	718	7	238	HIV-1 tat interactive protein, homolog (human) (Htatif) alternative variant cSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.dSep08</a>	<a href="#">192218</a>	2424	726	7	229	HIV-1 tat interactive protein, homolog (human) (26.0 kD) (Htatif) alternative variant dSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.eSep08</a>	<a href="#">192218</a>	1753	687	6	227	HIV-1 tat interactive protein, homolog (human) (Htatif) alternative variant eSep08, mRNA.

<a href="#">Htatif</a>	<a href="#">Htatif.fSep08</a>	<a href="#">192218</a>	1150	384	5	126	HIV-1 tat interactive protein, homolog (human) (Htatif) alternative variant fSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.gSep08</a>	<a href="#">192218</a>	908	401	4	119	HIV-1 tat interactive protein, homolog (human) (13.6 kD) (Htatif) alternative variant gSep08, mRNA.
<a href="#">Htatif</a>	<a href="#">Htatif.iSep08</a>	<a href="#">192218</a>	459	371	2	26	HIV-1 tat interactive protein, homolog (human) (Htatif) alternative variant iSep08, mRNA.
<a href="#">Htatif2</a>	<a href="#">Htatif2.aSep08</a>	<a href="#">292935</a>	14777	939	4	242	HIV-1 tat interactive protein 2, homolog (human) (26.9 kD) (Htatif2) alternative variant aSep08, mRNA.
<a href="#">Htatif2</a>	<a href="#">Htatif2.cSep08</a>	<a href="#">292935</a>	11793	486	2	161	HIV-1 tat interactive protein 2, homolog (human) (Htatif2) alternative variant cSep08, mRNA.
<a href="#">Htatsf1</a>	<a href="#">Htatsf1.bSep08</a>	<a href="#">317612</a>	4111	761	4	253	HIV TAT specific factor 1 (Htatsf1) alternative variant bSep08, mRNA.
<a href="#">Htatsf1</a>	<a href="#">Htatsf1.cSep08</a>	<a href="#">317612</a>	4175	738	5	222	HIV TAT specific factor 1 (Htatsf1) alternative variant cSep08, mRNA.
<a href="#">Htatsf1</a>	<a href="#">Htatsf1.dSep08</a>	<a href="#">317612</a>	2231	967	1	54	HIV TAT specific factor 1 (Htatsf1) alternative variant dSep08, mRNA.
<a href="#">Htf9c</a>	<a href="#">Htf9c.bSep08</a>	<a href="#">287953</a>	2437	1133	4	341	HpaII tiny fragments locus 9c (38.0 kD) (Htf9c) alternative variant bSep08, mRNA.
<a href="#">Htf9c</a>	<a href="#">Htf9c.cSep08</a>	<a href="#">287953</a>	2074	992	7	221	HpaII tiny fragments locus 9c (24.1 kD) (Htf9c) alternative variant cSep08, mRNA.
<a href="#">Htf9c</a>	<a href="#">Htf9c.dSep08</a>	<a href="#">287953</a>	1090	723	4	153	HpaII tiny fragments locus 9c (16.9 kD) (Htf9c) alternative variant dSep08, mRNA.
<a href="#">Htf9c</a>	<a href="#">Htf9c.eSep08</a>	<a href="#">287953</a>	1044	553	2	112	HpaII tiny fragments locus 9c (Htf9c) alternative variant eSep08, mRNA.
<a href="#">Htr3a</a>	<a href="#">Htr3a.bSep08</a>	<a href="#">79246</a>	1187	661	1	150	5-hydroxytryptamine (serotonin) receptor 3a (Htr3a) alternative variant bSep08, mRNA.
<a href="#">Htr3b</a>	<a href="#">Htr3b.bSep08</a>	<a href="#">58963</a>	18064	568	2	97	5-hydroxytryptamine (serotonin) receptor 3b (Htr3b) alternative variant bSep08, mRNA.
<a href="#">Htr3b</a>	<a href="#">Htr3b.cSep08</a>	<a href="#">58963</a>	18743	388	1	57	5-hydroxytryptamine (serotonin) receptor 3b (3.4 kD) (Htr3b) alternative variant cSep08, mRNA.
<a href="#">Htra1</a>	<a href="#">Htra1.bSep08</a>	<a href="#">65164</a>	31645	633	1	153	HtrA serine peptidase 1 (Htra1) alternative variant bSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.bSep08</a>	<a href="#">297376</a>	2161	698	6	171	protease serine (Htra2) alternative variant bSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.cSep08</a>	<a href="#">297376</a>	2577	1219	4	158	serine protease (16.4 kD) (Htra2) alternative variant cSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.dSep08</a>	<a href="#">297376</a>	2189	826	6	136	protease serine precursor (15.2 kD) (Htra2) alternative variant dSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.eSep08</a>	<a href="#">297376</a>	2998	1368	7	136	protease serine precursor (15.2 kD) (Htra2) alternative variant eSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.fSep08</a>	<a href="#">297376</a>	1013	924	2	132	serine protease (Htra2) alternative variant fSep08, mRNA.
<a href="#">Htra2</a>	<a href="#">Htra2.hSep08</a>	<a href="#">297376</a>	1856	364	5	65	serine protease (Htra2) alternative variant hSep08, mRNA.
<a href="#">Htra3</a>	<a href="#">Htra3.aSep08</a>	<a href="#">360959</a>	28359	2070		391	HtrA serine peptidase 3 (Htra3) mRNA.
<a href="#">Hunk</a>	<a href="#">Hunk.aSep08</a>	<a href="#">288275</a>	6069	1785		356	hormonally upregulated Neu-associated kinase (Hunk) alternative variant aSep08, mRNA.
<a href="#">Hus1andRGD1566034</a>	<a href="#">Hus1andRGD1566034.cSep08</a>	<a href="#">364193</a>	3561	599	4	116	checkpoint protein Hus1 (Hus1andRGD1566034) alternative variant cSep08, mRNA.

<a href="#">Hus1andRGD1566034</a>	<a href="#">Hus1andRGD1566034.cSep08</a>	<a href="#">498411</a>	3561	599	4	116	checkpoint protein Hus1 (Hus1andRGD1566034) alternative variant cSep08, mRNA.
<a href="#">Hyal2</a>	<a href="#">Hyal2.bSep08</a>	<a href="#">64468</a>	2948	827	1	226	hyaluronoglucosaminidase 2 (Hyal2) alternative variant bSep08, mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.bSep08</a>	<a href="#">300993</a>	3287	1425	2	334	N-acetyltransferase 6 (36.6 kD) (Hyal3) alternative variant bSep08, complete mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.cSep08</a>	<a href="#">300993</a>	3329	1502	2	312	N-acetyltransferase 6 (34.2 kD) (Hyal3) alternative variant cSep08, mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.dSep08</a>	<a href="#">300993</a>	6390	1243	4	259	hyaluronoglucosaminidase 3 CRA b (Hyal3) alternative variant dSep08, mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.fSep08</a>	<a href="#">300993</a>	2730	743	2	129	N-acetyltransferase 6 (Hyal3) alternative variant fSep08, mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.gSep08</a>	<a href="#">300993</a>	5461	1221	2	118	hyaluronoglucosaminidase 3 CRA b (Hyal3) alternative variant gSep08, mRNA.
<a href="#">Hyal3</a>	<a href="#">Hyal3.hSep08</a>	<a href="#">300993</a>	6162	924	2	98	hyaluronoglucosaminidase 3 CRA b (11.3 kD) (Hyal3) alternative variant hSep08, mRNA.
<a href="#">Hyal5</a>	<a href="#">Hyal5.bSep08</a>	<a href="#">500052</a>	13921	757	2	45	hyaluronoglucosaminidase 5 (5.3 kD) (Hyal5) alternative variant bSep08, mRNA.
<a href="#">Hydin</a>	<a href="#">Hydin.aSep08</a>	<a href="#">292017</a>	18146	2781		665	hydrocephalus inducing (74.8 kD) (Hydin) mRNA.
<a href="#">Hyd_WA.0</a>	<a href="#">Hyd_WA.0.aSep08</a>		3473	476		158	CRA a (Hyd_WA.0) mRNA.
<a href="#">Hyou1</a>	<a href="#">Hyou1.cSep08</a>	<a href="#">192235</a>	2395	627	3	208	hypoxia up-regulated 1 (Hyou1) alternative variant cSep08, mRNA.
<a href="#">Hyou1</a>	<a href="#">Hyou1.dSep08</a>	<a href="#">192235</a>	1831	402	1	133	hypoxia up-regulated 1 (Hyou1) alternative variant dSep08, mRNA.
<a href="#">Hypk</a>	<a href="#">Hypk.aSep08</a>	<a href="#">311359</a>	1377	613	3	130	huntingtin interacting protein K (Hypk) alternative variant aSep08, mRNA.
<a href="#">Hypk</a>	<a href="#">Hypk.bSep08</a>	<a href="#">311359</a>	771	413	1	75	huntingtin interacting protein K (Hypk) alternative variant bSep08, mRNA.
<a href="#">I-set.0</a>	<a href="#">I-set.0.aSep08</a>		78246	2118	5	670	palladin (I-set.0) alternative variant aSep08, mRNA.
<a href="#">I-set.0</a>	<a href="#">I-set.0.bSep08</a>		13196	2084	6	294	palladin (I-set.0) alternative variant bSep08, mRNA.
<a href="#">I-set.0</a>	<a href="#">I-set.0.cSep08</a>		140593	788	2	262	palladin (I-set.0) alternative variant cSep08, mRNA.
<a href="#">I-set.2</a>	<a href="#">I-set.2.aSep08</a>		731103	652		152	roundabout homolog 2 (I-set.2) mRNA.
<a href="#">I-set.3</a>	<a href="#">I-set.3.aSep08</a>		6309	420		139	down syndrome cell adhesion molecule CHD2-42 like (I-set.3) mRNA.
<a href="#">I-set.4</a>	<a href="#">I-set.4.aSep08</a>		46630	410		114	down syndrome cell adhesion molecule CHD2-42 like (I-set.4) mRNA.
<a href="#">I-set.5</a>	<a href="#">I-set.5.aSep08</a>		442	358		119	obscurin (I-set.5) mRNA.
<a href="#">I-set.6</a>	<a href="#">I-set.6.aSep08</a>		2155	713	5	199	obscurin-like 1 (I-set.6) alternative variant aSep08, mRNA.
<a href="#">I-set.6</a>	<a href="#">I-set.6.bSep08</a>		1618	624	3	88	obscurin-like 1 (9.8 kD) (I-set.6) alternative variant bSep08, mRNA.
<a href="#">I-set.7</a>	<a href="#">I-set.7.aSep08</a>		4170	1549	6	516	CRA a (I-set.7) alternative variant aSep08, mRNA.
<a href="#">I-set.7</a>	<a href="#">I-set.7.bSep08</a>		1279	1164	2	246	obscurin-like 1 (27.3 kD) (I-set.7) alternative variant bSep08, mRNA.
<a href="#">I-set.8</a>	<a href="#">I-set.8.aSep08</a>		25696	1163		387	neogenin (I-set.8) mRNA.
<a href="#">I-set.9</a>	<a href="#">I-set.9.aSep08</a>		5109	667		111	peroxidasin (I-set.9) mRNA.

<a href="#">I-set.10</a>	<a href="#">I-set.10.aSep08</a>		3726	1265		421	heparan sulfate proteoglycan (I-set.10) mRNA.
<a href="#">I-set.11</a>	<a href="#">I-set.11.aSep08</a>		5519	1653		535	leucine-rich repeats immunoglobulin-like domains 1 (I-set.11) alternative variant aSep08, mRNA.
<a href="#">I-set.11</a>	<a href="#">I-set.11.bSep08</a>		4424	1777		403	leucine-rich repeats immunoglobulin-like domains 1 (I-set.11) alternative variant bSep08, mRNA.
<a href="#">I-set.12</a>	<a href="#">I-set.12.aSep08</a>		868	593		183	titin CRA a (I-set.12) mRNA.
<a href="#">I-set.13</a>	<a href="#">I-set.13.aSep08</a>		485	378		125	titin (I-set.13) mRNA.
<a href="#">I-set.14</a>	<a href="#">I-set.14.aSep08</a>		924	467		155	titin (I-set.14) mRNA.
<a href="#">I-set.15</a>	<a href="#">I-set.15.aSep08</a>		1063	705		235	titin (I-set.15) mRNA.
<a href="#">I-set.16</a>	<a href="#">I-set.16.aSep08</a>		1594	1171		390	titin (I-set.16) mRNA.
<a href="#">I-set.17</a>	<a href="#">I-set.17.aSep08</a>		4366	952		317	titin (I-set.17) mRNA.
<a href="#">lapp</a>	<a href="#">lapp.bSep08</a>	<a href="#">24476</a>	2430	1089	2	80	islet amyloid polypeptide (lapp) alternative variant bSep08, mRNA.
<a href="#">lars2</a>	<a href="#">lars2.aSep08</a>	<a href="#">364070</a>	39435	1783	8	443	isoleucine-tRNA synthetase 2, mitochondrial (lars2) alternative variant aSep08, mRNA.
<a href="#">lars2</a>	<a href="#">lars2.bSep08</a>	<a href="#">364070</a>	7856	921	6	251	isoleucine-tRNA synthetase 2, mitochondrial (lars2) alternative variant bSep08, mRNA.
<a href="#">lars2</a>	<a href="#">lars2.cSep08</a>	<a href="#">364070</a>	3043	673	6	224	isoleucine-tRNA synthetase 2, mitochondrial (lars2) alternative variant cSep08, mRNA.
<a href="#">lars2</a>	<a href="#">lars2.dSep08</a>	<a href="#">364070</a>	3343	797	4	181	isoleucine-tRNA synthetase 2, mitochondrial (lars2) alternative variant dSep08, mRNA.
<a href="#">lars_predicted</a>	<a href="#">lars_predicted.aSep08</a>	<a href="#">306804</a>	31000	3293		931	isoleucine-tRNA synthetase (predicted) (lars_predicted) mRNA.
<a href="#">IBB.0</a>	<a href="#">IBB.0.aSep08</a>		21386	365		110	karyopherin alpha 6 (IBB.0) mRNA.
<a href="#">IBN_N.0</a>	<a href="#">IBN_N.0.aSep08</a>		37826	406		135	transportin 1 (IBN_N.0) mRNA.
<a href="#">lbtck</a>	<a href="#">lbtck.aSep08</a>	<a href="#">315858</a>	27104	2563	6	255	CRA c (lbtck) alternative variant aSep08, mRNA.
<a href="#">lbtck</a>	<a href="#">lbtck.bSep08</a>	<a href="#">315858</a>	24909	1461	4	193	inhibitor of Bruton's tyrosine kinase (20.9 kD) (lbtck) alternative variant bSep08, mRNA.
<a href="#">lbtck</a>	<a href="#">lbtck.cSep08</a>	<a href="#">315858</a>	16498	1087	9	159	inhibitor of Bruton's tyrosine kinase (17.9 kD) (lbtck) alternative variant cSep08, mRNA.
<a href="#">lbtck</a>	<a href="#">lbtck.dSep08</a>	<a href="#">315858</a>	13011	459	3	152	inhibitor of Bruton's tyrosine kinase (lbtck) alternative variant dSep08, mRNA.
<a href="#">lbtck</a>	<a href="#">lbtck.eSep08</a>	<a href="#">315858</a>	3414	737	1	31	putative protein (3.6 kD) (lbtck) alternative variant eSep08, mRNA.
<a href="#">Ica1</a>	<a href="#">Ica1.bSep08</a>	<a href="#">81024</a>	149262	1832	12	459	islet cell autoantigen 1 (52.7 kD) (Ica1) alternative variant bSep08, mRNA.
<a href="#">Ica1</a>	<a href="#">Ica1.cSep08</a>	<a href="#">81024</a>	110118	877	7	234	islet cell autoantigen 1 (Ica1) alternative variant cSep08, mRNA.
<a href="#">Ica1</a>	<a href="#">Ica1.dSep08</a>	<a href="#">81024</a>	39402	717	6	192	islet cell autoantigen 1 (Ica1) alternative variant dSep08, mRNA.
<a href="#">Ica1</a>	<a href="#">Ica1.eSep08</a>	<a href="#">81024</a>	12745	518	1	143	islet cell autoantigen 1 (Ica1) alternative variant eSep08, mRNA.
<a href="#">Ica1l</a>	<a href="#">Ica1l.bSep08</a>	<a href="#">316432</a>	7397	1576	2	112	islet cell autoantigen 1-like (Ica1l) alternative variant bSep08, mRNA.
<a href="#">Icam2</a>	<a href="#">Icam2.bSep08</a>	<a href="#">360647</a>	4324	1340	4	127	intercellular adhesion molecule 2 (14.1 kD) (Icam2) alternative variant bSep08, mRNA.

<a href="#">Icam2</a>	<a href="#">Icam2.cSep08</a>	<a href="#">360647</a>	5330	679	5	116	intercellular adhesion molecule 2 (Icam2) alternative variant cSep08, mRNA.
<a href="#">Icam2</a>	<a href="#">Icam2.eSep08</a>	<a href="#">360647</a>	4047	589	4	72	intercellular adhesion molecule 2 (Icam2) alternative variant eSep08, mRNA.
<a href="#">Icam5</a>	<a href="#">Icam5.aSep08</a>	<a href="#">313785</a>	1364	752		245	intercellular adhesion molecule 5, telencephalin (Icam5) mRNA.
<a href="#">Ick</a>	<a href="#">Ick.bSep08</a>	<a href="#">84411</a>	1597	644	2	159	intestinal cell kinase (Ick) alternative variant bSep08, mRNA.
<a href="#">Ick</a>	<a href="#">Ick.dSep08</a>	<a href="#">84411</a>	22381	517	2	39	intestinal cell kinase (4.6 kD) (Ick) alternative variant dSep08, mRNA.
<a href="#">Ick</a>	<a href="#">Ick.eSep08</a>	<a href="#">84411</a>	3232	385	2	54	intestinal cell kinase (Ick) alternative variant eSep08, mRNA.
<a href="#">Icmt</a>	<a href="#">Icmt.bSep08</a>	<a href="#">170818</a>	2033	532	1	155	isoprenylcysteine carboxyl methyltransferase (Icmt) alternative variant bSep08, mRNA.
<a href="#">Icosl</a>	<a href="#">Icosl.aSep08</a>	<a href="#">499415</a>	5824	2073	1	354	icos ligand (Icosl) alternative variant aSep08, mRNA.
<a href="#">Icosl</a>	<a href="#">Icosl.bSep08</a>	<a href="#">499415</a>	4625	871	1	290	icos ligand (Icosl) alternative variant bSep08, mRNA.
<a href="#">Ict1</a>	<a href="#">Ict1.aSep08</a>	<a href="#">303673</a>	6495	826	3	205	immature colon carcinoma transcript 1 (Ict1) alternative variant aSep08, mRNA.
<a href="#">Ict1</a>	<a href="#">Ict1.bSep08</a>	<a href="#">303673</a>	6347	1282	1	101	immature colon carcinoma transcript 1 (Ict1) alternative variant bSep08, mRNA.
<a href="#">Id1</a>	<a href="#">Id1.bSep08</a>	<a href="#">25261</a>	1129	913	2	154	inhibitor of DNA binding 1 (16.3 kD) (Id1) alternative variant bSep08, complete mRNA.
<a href="#">Id3</a>	<a href="#">Id3.aSep08</a>	<a href="#">25585</a>	948	833	1	119	inhibitor of DNA binding 3 (13.1 kD) (Id3) alternative variant aSep08, mRNA.
<a href="#">Ide</a>	<a href="#">Ide.bSep08</a>	<a href="#">25700</a>	30419	1781	8	431	insulin degrading enzyme (Ide) alternative variant bSep08, mRNA.
<a href="#">Idh1</a>	<a href="#">Idh1.bSep08</a>	<a href="#">24479</a>	15787	397	3	102	isocitrate dehydrogenase 1 (NADP+), soluble (Idh1) alternative variant bSep08, mRNA.
<a href="#">Idh1</a>	<a href="#">Idh1.cSep08</a>	<a href="#">24479</a>	4002	338	2	41	isocitrate dehydrogenase 1 (NADP+), soluble (4.9 kD) (Idh1) alternative variant cSep08, complete mRNA.
<a href="#">Idh2</a>	<a href="#">Idh2.bSep08</a>	<a href="#">361596</a>	16360	925	6	259	isocitrate dehydrogenase 2 (NADP+), mitochondrial (Idh2) alternative variant bSep08, mRNA.
<a href="#">Idh3a</a>	<a href="#">Idh3a.bSep08</a>	<a href="#">114096</a>	14606	787	8	259	isocitrate dehydrogenase 3 (NAD+) alpha (Idh3a) alternative variant bSep08, mRNA.
<a href="#">Idh3a</a>	<a href="#">Idh3a.cSep08</a>	<a href="#">114096</a>	9561	663	6	221	isocitrate dehydrogenase 3 (NAD+) alpha (Idh3a) alternative variant cSep08, mRNA.
<a href="#">Idh3a</a>	<a href="#">Idh3a.dSep08</a>	<a href="#">114096</a>	17210	617	8	195	isocitrate dehydrogenase 3 (NAD+) alpha (Idh3a) alternative variant dSep08, mRNA.
<a href="#">Idh3B</a>	<a href="#">Idh3B.bSep08</a>	<a href="#">94173</a>	2271	670	7	207	isocitrate dehydrogenase 3 (NAD+) beta (Idh3B) alternative variant bSep08, mRNA.
<a href="#">Idh3g</a>	<a href="#">Idh3g.bSep08</a>	<a href="#">25179</a>	8882	1668	10	339	isocitrate dehydrogenase 3 gamma CRA c (37.0 kD) (Idh3g) alternative variant bSep08, mRNA.
<a href="#">Idh3g</a>	<a href="#">Idh3g.cSep08</a>	<a href="#">25179</a>	7567	731	9	243	isocitrate dehydrogenase 3 gamma (Idh3g) alternative variant cSep08, mRNA.
<a href="#">Idh3g</a>	<a href="#">Idh3g.dSep08</a>	<a href="#">25179</a>	7430	1552	5	187	isocitrate dehydrogenase 3 gamma (20.7 kD) (Idh3g) alternative variant dSep08, mRNA.



<a href="#">ldh3g</a>	<a href="#">ldh3g.eSep08</a>	<a href="#">25179</a>	7110	746	7	180	isocitrate dehydrogenase 3 gamma (ldh3g) alternative variant eSep08, mRNA.
<a href="#">ldh3g</a>	<a href="#">ldh3g.fSep08</a>	<a href="#">25179</a>	7223	672	7	166	isocitrate dehydrogenase 3 gamma CRA a (ldh3g) alternative variant fSep08, mRNA.
<a href="#">ldh3g</a>	<a href="#">ldh3g.gSep08</a>	<a href="#">25179</a>	6860	511	7	95	isocitrate dehydrogenase 3 gamma (ldh3g) alternative variant gSep08, mRNA.
<a href="#">ldh3g</a>	<a href="#">ldh3g.hSep08</a>	<a href="#">25179</a>	805	428	3	60	isocitrate dehydrogenase 3 gamma (6.6 kD) (ldh3g) alternative variant hSep08, mRNA.
<a href="#">ldh3g</a>	<a href="#">ldh3g.iSep08</a>	<a href="#">25179</a>	5060	696	2	35	putative protein (ldh3g) alternative variant iSep08, mRNA.
<a href="#">lds</a>	<a href="#">lds.aSep08</a>	<a href="#">363513</a>	16253	1255	2	417	iduronate 2-sulfatase (lds) alternative variant aSep08, mRNA.
<a href="#">lds</a>	<a href="#">lds.bSep08</a>	<a href="#">363513</a>	6487	846	1	281	iduronate 2-sulfatase (lds) alternative variant bSep08, mRNA.
<a href="#">lds</a>	<a href="#">lds.cSep08</a>	<a href="#">363513</a>	4299	1200	1	191	iduronate 2-sulfatase (21.7 kD) (lds) alternative variant cSep08, mRNA.
<a href="#">lds</a>	<a href="#">lds.dSep08</a>	<a href="#">363513</a>	3776	673	1	63	iduronate 2-sulfatase (lds) alternative variant dSep08, mRNA.
<a href="#">ler3</a>	<a href="#">ler3.aSep08</a>	<a href="#">294235</a>	1208	1101	2	160	immediate early response 3 (17.6 kD) (ler3) alternative variant aSep08, complete mRNA.
<a href="#">ler3</a>	<a href="#">ler3.bSep08</a>	<a href="#">294235</a>	1258	439	3	146	immediate early response 3 (ler3) alternative variant bSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.aSep08</a>	<a href="#">170512</a>	6529	857	6	226	CRA b like (lfi27l) alternative variant aSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.aSep08</a>	<a href="#">360415</a>	6529	857	6	226	CRA b like (lfi27l) alternative variant aSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.bSep08</a>	<a href="#">170512</a>	4668	405	4	134	CRA b like (lfi27l) alternative variant bSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.bSep08</a>	<a href="#">360415</a>	4668	405	4	134	CRA b like (lfi27l) alternative variant bSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.cSep08</a>	<a href="#">170512</a>	3281	717	3	119	CRA b like (lfi27l) alternative variant cSep08, mRNA.
<a href="#">lfi27l</a>	<a href="#">lfi27l.cSep08</a>	<a href="#">360415</a>	3281	717	3	119	CRA b like (lfi27l) alternative variant cSep08, mRNA.
<a href="#">lfi30</a>	<a href="#">lfi30.bSep08</a>	<a href="#">290644</a>	3171	1134	1	212	interferon gamma inducible protein 30 (lfi30) alternative variant bSep08, mRNA.
<a href="#">lfi35</a>	<a href="#">lfi35.bSep08</a>	<a href="#">287719</a>	7815	833	6	277	interferon-induced protein 35 (lfi35) alternative variant bSep08, mRNA.
<a href="#">lfi44</a>	<a href="#">lfi44.bSep08</a>	<a href="#">310969</a>	17749	1490	8	416	interferon-induced protein 44 (lfi44) alternative variant bSep08, mRNA.
<a href="#">lfi44</a>	<a href="#">lfi44.cSep08</a>	<a href="#">310969</a>	4039	765	5	187	interferon-induced protein 44 (lfi44) alternative variant cSep08, mRNA.
<a href="#">lfi44</a>	<a href="#">lfi44.dSep08</a>	<a href="#">310969</a>	8426	715	4	135	interferon-induced protein 44 (lfi44) alternative variant dSep08, mRNA.
<a href="#">lfi203</a>	<a href="#">lfi203.aSep08</a>	<a href="#">498288</a>	5378	545		101	interferon activated gene 203 (lfi203) mRNA.
<a href="#">lfi204</a>	<a href="#">lfi204.bSep08</a>	<a href="#">304988</a>	7560	916	1	192	interferon activated gene 204 (lfi204) alternative variant bSep08, mRNA.
<a href="#">lfih1</a>	<a href="#">lfih1.bSep08</a>	<a href="#">499801</a>	7920	693	3	85	interferon induced with helicase C domain 1 (lfih1) alternative variant bSep08, mRNA.
<a href="#">lfit3</a>	<a href="#">lfit3.aSep08</a>	<a href="#">309526</a>	4920	1816		411	interferon-induced protein with tetratricopeptide repeats 3 (48.1 kD) (lfit3) mRNA.
<a href="#">lfitm2</a>	<a href="#">lfitm2.bSep08</a>	<a href="#">114709</a>	624	382	2	98	interferon induced transmembrane protein 2 (lfitm2) alternative variant bSep08, mRNA.

<a href="#">Ifitm2</a>	<a href="#">Ifitm2.cSep08</a>	<a href="#">114709</a>	4384	536	2	85	interferon induced transmembrane protein 2 (9.4 kD) (Ifitm2) alternative variant cSep08, complete mRNA.
<a href="#">Ifitm3</a>	<a href="#">Ifitm3.aSep08</a>	<a href="#">361673</a>	1158	652	2	169	interferon induced transmembrane protein 3 (Ifitm3) alternative variant aSep08, mRNA.
<a href="#">Ifitm6</a>	<a href="#">Ifitm6.aSep08</a>	<a href="#">309104</a>	1310	687		129	interferon induced transmembrane protein 6 (Ifitm6) mRNA.
<a href="#">Ifitd1</a>	<a href="#">Ifitd1.aSep08</a>	<a href="#">500362</a>	8015	759		180	putative protein of mammalian origin (Ifitd1) mRNA.
<a href="#">Ifnar1</a>	<a href="#">Ifnar1.bSep08</a>	<a href="#">288264</a>	5863	1761	4	216	interferon (alpha and beta) receptor 1 (Ifnar1) alternative variant bSep08, mRNA.
<a href="#">Ifngr1</a>	<a href="#">Ifngr1.bSep08</a>	<a href="#">116465</a>	16723	967	6	231	interferon gamma receptor 1 (Ifngr1) alternative variant bSep08, mRNA.
<a href="#">Ifngr1</a>	<a href="#">Ifngr1.cSep08</a>	<a href="#">116465</a>	1724	884	2	202	interferon gamma receptor 1 (Ifngr1) alternative variant cSep08, mRNA.
<a href="#">Ifrd2</a>	<a href="#">Ifrd2.bSep08</a>	<a href="#">300994</a>	1560	718	1	139	interferon-related developmental regulator 2 (Ifrd2) alternative variant bSep08, mRNA.
<a href="#">Ifit20</a>	<a href="#">Ifit20.bSep08</a>	<a href="#">287541</a>	5488	890	4	132	intraflagellar transport 20 homolog (Chlamydomonas) (15.3 kD) (Ifit20) alternative variant bSep08, mRNA.
<a href="#">Ifit20</a>	<a href="#">Ifit20.cSep08</a>	<a href="#">287541</a>	3236	967	3	131	intraflagellar transport 20 homolog (Chlamydomonas) (14.9 kD) (Ifit20) alternative variant cSep08, mRNA.
<a href="#">Ifit20</a>	<a href="#">Ifit20.dSep08</a>	<a href="#">287541</a>	5081	586	4	114	intraflagellar transport 20 homolog (Chlamydomonas) (13.2 kD) (Ifit20) alternative variant dSep08, mRNA.
<a href="#">Ifit52</a>	<a href="#">Ifit52.aSep08</a>	<a href="#">362265</a>	24393	2140	13	394	intraflagellar transport 52 homolog (Chlamydomonas) (Ifit52) alternative variant aSep08, mRNA.
<a href="#">Ifit52</a>	<a href="#">Ifit52.bSep08</a>	<a href="#">362265</a>	24468	2143	13	343	intraflagellar transport 52 homolog (Chlamydomonas) (Ifit52) alternative variant bSep08, complete mRNA.
<a href="#">Ifit52</a>	<a href="#">Ifit52.cSep08</a>	<a href="#">362265</a>	2218	704	4	120	intraflagellar transport 52 homolog (Chlamydomonas) (Ifit52) alternative variant cSep08, mRNA.
<a href="#">Ifit57</a>	<a href="#">Ifit57.bSep08</a>	<a href="#">303968</a>	8641	262		86	intraflagellar transport 57 homolog (Chlamydomonas) (Ifit57) alternative variant bSep08, mRNA.
<a href="#">Ifit74</a>	<a href="#">Ifit74.bSep08</a>	<a href="#">313365</a>	28347	742	5	149	intraflagellar transport 74 homolog (Chlamydomonas) (Ifit74) alternative variant bSep08, mRNA.
<a href="#">Ifit74</a>	<a href="#">Ifit74.cSep08</a>	<a href="#">313365</a>	16694	744	7	115	intraflagellar transport 74 homolog (Chlamydomonas) (Ifit74) alternative variant cSep08, mRNA.
<a href="#">Ifit74</a>	<a href="#">Ifit74.dSep08</a>	<a href="#">313365</a>	2144	979	2	42	intraflagellar transport 74 homolog (Chlamydomonas) (4.8 kD) (Ifit74) alternative variant dSep08, mRNA.
<a href="#">Ifit80</a>	<a href="#">Ifit80.bSep08</a>	<a href="#">295106</a>	8261	616	2	139	intraflagellar transport 80 homolog (Chlamydomonas) (15.6 kD) (Ifit80) alternative variant bSep08, mRNA.
<a href="#">Ifit81</a>	<a href="#">Ifit81.bSep08</a>	<a href="#">373066</a>	59149	1260	11	343	carnitine deficiency-associated expressed in ventricle 1 (Ifit81) alternative variant bSep08, mRNA.
<a href="#">Ifit81</a>	<a href="#">Ifit81.cSep08</a>	<a href="#">373066</a>	3798	428	2	81	putative protein (Ifit81) alternative variant cSep08, mRNA.
<a href="#">Ifit81</a>	<a href="#">Ifit81.dSep08</a>	<a href="#">373066</a>	4923	347	3	68	intraflagellar transport 81 homolog CRA a (Ifit81) alternative variant dSep08, mRNA.
<a href="#">Ifit81</a>	<a href="#">Ifit81.eSep08</a>	<a href="#">373066</a>	2049	296	2	48	intraflagellar transport 81 homolog CRA a (Ifit81) alternative variant eSep08, mRNA.
<a href="#">Ifit81</a>	<a href="#">Ifit81.fSep08</a>	<a href="#">373066</a>	646	297	2	39	intraflagellar transport 81 homolog CRA a (Ifit81) alternative variant fSep08, mRNA.

<a href="#">lft88</a>	<a href="#">lft88.bSep08</a>	<a href="#">305918</a>	6091	403	4	90	intraflagellar transport 88 homolog (Chlamydomonas) (lft88) alternative variant bSep08, mRNA.
<a href="#">lft122</a>	<a href="#">lft122.aSep08</a>	<a href="#">312651</a>	55588	2392	18	739	intraflagellar transport 122 homolog (Chlamydomonas) (83.6 kD) (lft122) alternative variant aSep08, mRNA.
<a href="#">lft122</a>	<a href="#">lft122.bSep08</a>	<a href="#">312651</a>	14343	646	6	215	intraflagellar transport 122 homolog (Chlamydomonas) (lft122) alternative variant bSep08, mRNA.
<a href="#">ig.0</a>	<a href="#">ig.0.aSep08</a>		15111	1365	2	273	gamma receptor iii (30.8 kD) (ig.0) alternative variant aSep08, mRNA.
<a href="#">ig.0</a>	<a href="#">ig.0.bSep08</a>		7389	703	1	148	receptor (ig.0) alternative variant bSep08, mRNA.
<a href="#">ig.1</a>	<a href="#">ig.1.aSep08</a>		23374	330		109	sidekick 1 (ig.1) mRNA.
<a href="#">ig.4</a>	<a href="#">ig.4.aSep08</a>		2539	856		285	heparan sulfate proteoglycan (ig.4) mRNA.
<a href="#">ig.5</a>	<a href="#">ig.5.aSep08</a>		1546	553		112	glycoprotein pregnancy specific 1 (ig.5) mRNA.
<a href="#">lgbp1</a>	<a href="#">lgbp1.bSep08</a>	<a href="#">58845</a>	11944	820	1	100	immunoglobulin (CD79A) binding protein 1 (11.7 kD) (lgbp1) alternative variant bSep08, mRNA.
<a href="#">lgf1</a>	<a href="#">lgf1.fSep08</a>	<a href="#">24482</a>	72949	756	5	121	insulin-like growth factor 1 (lgf1) alternative variant fSep08, mRNA.
<a href="#">lgf1</a>	<a href="#">lgf1.gSep08</a>	<a href="#">24482</a>	5524	304	3	73	insulin-like growth factor 1 (lgf1) alternative variant gSep08, mRNA.
<a href="#">lgf1r</a>	<a href="#">lgf1r.bSep08</a>	<a href="#">25718</a>	21998	2541	3	356	insulin-like growth factor I receptor (lgf1r) alternative variant bSep08, mRNA.
<a href="#">lgf2</a>	<a href="#">lgf2.bSep08</a>	<a href="#">24483</a>	3969	885	3	180	insulin-like growth factor 2 (20.1 kD) (lgf2) alternative variant bSep08, mRNA.
<a href="#">lgf2</a>	<a href="#">lgf2.cSep08</a>	<a href="#">24483</a>	6744	368	2	122	insulin-like growth factor 2 (lgf2) alternative variant cSep08, mRNA.
<a href="#">lgf2</a>	<a href="#">lgf2.dSep08</a>	<a href="#">24483</a>	1684	467	1	84	insulin-like growth factor 2 (lgf2) alternative variant dSep08, mRNA.
<a href="#">lgf2bp2</a>	<a href="#">lgf2bp2.aSep08</a>	<a href="#">303824</a>	30983	3309	10	513	insulin-like growth factor 2 mRNA binding protein 2 (lgf2bp2) mRNA.
<a href="#">lgf2bp3</a>	<a href="#">lgf2bp3.aSep08</a>	<a href="#">312320</a>	133540	1816	11	457	insulin-like growth factor 2 mRNA binding protein 3 (lgf2bp3) alternative variant aSep08, mRNA.
<a href="#">lgf2bp3</a>	<a href="#">lgf2bp3.bSep08</a>	<a href="#">312320</a>	21564	2629	6	188	insulin-like growth factor 2 mRNA binding protein 3 (lgf2bp3) alternative variant bSep08, mRNA.
<a href="#">lgf2bp3</a>	<a href="#">lgf2bp3.cSep08</a>	<a href="#">312320</a>	28290	667	5	186	insulin-like growth factor 2 mRNA binding protein 3 (lgf2bp3) alternative variant cSep08, mRNA.
<a href="#">lgf2bp3</a>	<a href="#">lgf2bp3.eSep08</a>	<a href="#">312320</a>	28180	701	5	145	insulin-like growth factor 2 mRNA binding protein 3 (lgf2bp3) alternative variant eSep08, mRNA.
<a href="#">lgf2r</a>	<a href="#">lgf2r.bSep08</a>	<a href="#">25151</a>	2083	416	3	138	insulin-like growth factor 2 receptor (lgf2r) alternative variant bSep08, mRNA.
<a href="#">lgf2r</a>	<a href="#">lgf2r.dSep08</a>	<a href="#">25151</a>	1605	747	2	34	insulin-like growth factor 2 receptor (lgf2r) alternative variant dSep08, mRNA.
<a href="#">lgfbp7</a>	<a href="#">lgfbp7.bSep08</a>	<a href="#">289560</a>	2750	512	3	94	insulin-like growth factor binding protein 7 (lgfbp7) alternative variant bSep08, mRNA.
<a href="#">lgn1</a>	<a href="#">lgn1.aSep08</a>	<a href="#">304823</a>	992	472		42	putative protein (lgn1) mRNA.
<a href="#">lghg</a>	<a href="#">lghg.bSep08</a>	<a href="#">299354</a>	1646	1087		326	immunoglobulin heavy chain (gamma polypeptide) (lghg) alternative variant bSep08, mRNA.

<a href="#">Ighmbp2</a>	<a href="#">Ighmbp2.bSep08</a>	<a href="#">29532</a>	3202	751	2	76	immunoglobulin mu binding protein 2 (Ighmbp2) alternative variant bSep08, mRNA.
<a href="#">Igj</a>	<a href="#">Igj.aSep08</a>	<a href="#">360922</a>	7058	1046	1	159	immunoglobulin joining chain (17.8 kD) (Igj) alternative variant aSep08, mRNA.
<a href="#">Igj</a>	<a href="#">Igj.bSep08</a>	<a href="#">360922</a>	6836	743		132	immunoglobulin joining chain (14.7 kD) (Igj) alternative variant bSep08, mRNA.
<a href="#">Igsf1</a>	<a href="#">Igsf1.bSep08</a>	<a href="#">302822</a>	2203	927	5	308	immunoglobulin superfamily, member 1 (Igsf1) alternative variant bSep08, mRNA.
<a href="#">Igsf1</a>	<a href="#">Igsf1.cSep08</a>	<a href="#">302822</a>	2107	1162	4	261	immunoglobulin superfamily, member 1 (Igsf1) alternative variant cSep08, mRNA.
<a href="#">Igsf1</a>	<a href="#">Igsf1.dSep08</a>	<a href="#">302822</a>	4870	1721	5	232	immunoglobulin superfamily, member 1 (26.4 kD) (Igsf1) alternative variant dSep08, mRNA.
<a href="#">Igsf1</a>	<a href="#">Igsf1.fSep08</a>	<a href="#">302822</a>	406	309	2	87	immunoglobulin superfamily, member 1 (Igsf1) alternative variant fSep08, mRNA.
<a href="#">Igsf2</a>	<a href="#">Igsf2.aSep08</a>	<a href="#">310727</a>	4678	727		242	immunoglobulin superfamily, member 2 (Igsf2) mRNA.
<a href="#">Igsf7</a>	<a href="#">Igsf7.aSep08</a>	<a href="#">287813</a>	4253	687	2	190	immunoglobulin superfamily, member 7 (Igsf7) alternative variant aSep08, mRNA.
<a href="#">Igsf7</a>	<a href="#">Igsf7.bSep08</a>	<a href="#">287813</a>	2538	757	1	103	immunoglobulin superfamily, member 7 (11.4 kD) (Igsf7) alternative variant bSep08, mRNA.
<a href="#">Igsf8</a>	<a href="#">Igsf8.bSep08</a>	<a href="#">304979</a>	3844	442	2	147	immunoglobulin superfamily, member 8 (Igsf8) alternative variant bSep08, mRNA.
<a href="#">Igsf11</a>	<a href="#">Igsf11.bSep08</a>	<a href="#">303926</a>	123977	664	2	120	immunoglobulin superfamily, member 11 (Igsf11) alternative variant bSep08, mRNA.
<a href="#">Igsf21</a>	<a href="#">Igsf21.aSep08</a>	<a href="#">298591</a>	1449	467		104	immunoglobulin superfamily, member 21 (Igsf21) mRNA.
<a href="#">Igtp</a>	<a href="#">Igtp.aSep08</a>	<a href="#">303163</a>	9302	2692	3	474	interferon gamma induced GTPase (Igtp) alternative variant aSep08, mRNA.
<a href="#">Igtp</a>	<a href="#">Igtp.bSep08</a>	<a href="#">303163</a>	837	434	2	125	interferon gamma induced GTPase (Igtp) alternative variant bSep08, mRNA.
<a href="#">Igtp</a>	<a href="#">Igtp.cSep08</a>	<a href="#">303163</a>	1256	1107	3	55	interferon gamma induced GTPase (Igtp) alternative variant cSep08, mRNA.
<a href="#">Ihpk2</a>	<a href="#">Ihpk2.bSep08</a>	<a href="#">59268</a>	9370	2207	4	345	inositol hexaphosphate kinase 2 (Ihpk2) alternative variant bSep08, mRNA.
<a href="#">Ihpk2</a>	<a href="#">Ihpk2.cSep08</a>	<a href="#">59268</a>	1298	521	3	173	inositol hexaphosphate kinase 2 CRA d (Ihpk2) alternative variant cSep08, mRNA.
<a href="#">Ihpk2</a>	<a href="#">Ihpk2.eSep08</a>	<a href="#">59268</a>	1490	604	2	111	inositol hexaphosphate kinase 2 (Ihpk2) alternative variant eSep08, mRNA.
<a href="#">Ihpk2</a>	<a href="#">Ihpk2.fSep08</a>	<a href="#">59268</a>	948	520	2	53	putative protein (Ihpk2) alternative variant fSep08, mRNA.
<a href="#">Iip45</a>	<a href="#">Iip45.bSep08</a>	<a href="#">298643</a>	4775	857	6	244	invasion inhibitory protein 45 (Iip45) alternative variant bSep08, mRNA.
<a href="#">Iip45</a>	<a href="#">Iip45.cSep08</a>	<a href="#">298643</a>	4840	784	4	196	invasion inhibitory protein 45 (Iip45) alternative variant cSep08, mRNA.
<a href="#">Iip45</a>	<a href="#">Iip45.dSep08</a>	<a href="#">298643</a>	2311	675	3	162	invasion inhibitory protein 45 (Iip45) alternative variant dSep08, mRNA.
<a href="#">Iip45</a>	<a href="#">Iip45.eSep08</a>	<a href="#">298643</a>	1818	697	4	155	invasion inhibitory protein 45 (Iip45) alternative variant eSep08, mRNA.
<a href="#">Iip45</a>	<a href="#">Iip45.fSep08</a>	<a href="#">298643</a>	1633	459	4	153	invasion inhibitory protein 45 (Iip45) alternative variant fSep08, mRNA.

<a href="#">lip45</a>	<a href="#">lip45.gSep08</a>	<a href="#">298643</a>	2192	560	3	123	invasion inhibitory protein 45 (lip45) alternative variant gSep08, mRNA.
<a href="#">lip45</a>	<a href="#">lip45.hSep08</a>	<a href="#">298643</a>	1903	427	2	99	invasion inhibitory protein 45 (10.9 kD) (lip45) alternative variant hSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.bSep08</a>	<a href="#">291659</a>	4590	605	6	177	IK cytokine down-regulator of HLA II (20.5 kD) (lk) alternative variant bSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.cSep08</a>	<a href="#">291659</a>	4632	790	5	135	ik cytokine (15.8 kD) (lk) alternative variant cSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.dSep08</a>	<a href="#">291659</a>	5133	630	4	115	IK cytokine down-regulator of HLA II (lk) alternative variant dSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.eSep08</a>	<a href="#">291659</a>	2152	747	5	105	IK cytokine (12.4 kD) (lk) alternative variant eSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.fSep08</a>	<a href="#">291659</a>	1330	781	3	91	IK cytokine (10.7 kD) (lk) alternative variant fSep08, mRNA.
<a href="#">lk</a>	<a href="#">lk.gSep08</a>	<a href="#">291659</a>	1061	234	2	48	putative protein (lk) alternative variant gSep08, mRNA.
<a href="#">lkbkap</a>	<a href="#">lkbkap.bSep08</a>	<a href="#">140934</a>	40714	1779	7	441	inhibitor of kappa light polypeptide enhancer in B-cells, kinase complex-associated protein (lkbkap) alternative variant bSep08, mRNA.
<a href="#">lkbkb</a>	<a href="#">lkbkb.bSep08</a>	<a href="#">84351</a>	3450	700	5	214	inhibitor of kappaB kinase beta (lkbkb) alternative variant bSep08, mRNA.
<a href="#">lkbkb</a>	<a href="#">lkbkb.dSep08</a>	<a href="#">84351</a>	1293	725	2	100	inhibitor of kappaB kinase beta (lkbkb) alternative variant dSep08, mRNA.
<a href="#">lkbkb</a>	<a href="#">lkbkb.gSep08</a>	<a href="#">84351</a>	8702	502	2	30	inhibitor of kappaB kinase beta (lkbkb) alternative variant gSep08, mRNA.
<a href="#">lkbke</a>	<a href="#">lkbke.bSep08</a>	<a href="#">363984</a>	11256	730	8	174	inhibitor of kappaB kinase epsilon (lkbke) alternative variant bSep08, mRNA.
<a href="#">lkbke</a>	<a href="#">lkbke.cSep08</a>	<a href="#">363984</a>	3110	736	4	166	inhibitor of kappaB kinase epsilon (lkbke) alternative variant cSep08, mRNA.
<a href="#">lkbke</a>	<a href="#">lkbke.dSep08</a>	<a href="#">363984</a>	3553	679	4	44	inhibitor of kappaB kinase epsilon (lkbke) alternative variant dSep08, mRNA.
<a href="#">lkzf2</a>	<a href="#">lkzf2.bSep08</a>	<a href="#">301476</a>	1367	530	1	63	IKAROS family zinc finger 2 (lkzf2) alternative variant bSep08, mRNA.
<a href="#">lkzf5</a>	<a href="#">lkzf5.bSep08</a>	<a href="#">309031</a>	14444	370	3	83	IKAROS family zinc finger 5 (lkzf5) alternative variant bSep08, mRNA.
<a href="#">lkzf5</a>	<a href="#">lkzf5.cSep08</a>	<a href="#">309031</a>	12555	339	2	57	IKAROS family zinc finger 5 (6.5 kD) (lkzf5) alternative variant cSep08, mRNA.
<a href="#">ll1a</a>	<a href="#">ll1a.bSep08</a>	<a href="#">24493</a>	6900	1685	3	186	interleukin 1 alpha (ll1a) alternative variant bSep08, mRNA.
<a href="#">ll1a</a>	<a href="#">ll1a.cSep08</a>	<a href="#">24493</a>	2580	701	1	74	interleukin 1 alpha (ll1a) alternative variant cSep08, mRNA.
<a href="#">ll1r1</a>	<a href="#">ll1r1.bSep08</a>	<a href="#">25663</a>	40566	2521	13	573	interleukin 1 receptor, type I (66.3 kD) (ll1r1) alternative variant bSep08, mRNA.
<a href="#">ll1r1</a>	<a href="#">ll1r1.cSep08</a>	<a href="#">25663</a>	38643	1763	12	525	interleukin 1 receptor, type I (60.8 kD) (ll1r1) alternative variant cSep08, mRNA.
<a href="#">ll1r2</a>	<a href="#">ll1r2.bSep08</a>	<a href="#">117022</a>	7310	614	1	185	interleukin 1 receptor, type II (20.6 kD) (ll1r2) alternative variant bSep08, mRNA.
<a href="#">ll1rap</a>	<a href="#">ll1rap.bSep08</a>	<a href="#">25466</a>	25162	1334	6	295	interleukin 1 receptor accessory protein (ll1rap) alternative variant bSep08, mRNA.

<a href="#">Il1rap</a>	<a href="#">Il1rap.cSep08</a>	<a href="#">25466</a>	10664	650	3	150	interleukin 1 receptor accessory protein (Il1rap) alternative variant cSep08, mRNA.
<a href="#">Il1rapl2</a>	<a href="#">Il1rapl2.aSep08</a>	<a href="#">300913</a>	52304	423		89	interleukin 1 receptor accessory protein-like 2 (Il1rapl2) mRNA.
<a href="#">Il1rl2</a>	<a href="#">Il1rl2.bSep08</a>	<a href="#">171106</a>	3351	499	1	121	interleukin 1 receptor-like 2 (Il1rl2) alternative variant bSep08, mRNA.
<a href="#">Il1rn</a>	<a href="#">Il1rn.bSep08</a>	<a href="#">60582</a>	4170	517	3	172	interleukin 1 receptor antagonist (Il1rn) alternative variant bSep08, mRNA.
<a href="#">Il2rg</a>	<a href="#">Il2rg.bSep08</a>	<a href="#">140924</a>	2799	854	6	206	interleukin 2 receptor (Il2rg) alternative variant bSep08, mRNA.
<a href="#">Il2rg</a>	<a href="#">Il2rg.cSep08</a>	<a href="#">140924</a>	1135	692	3	192	interleukin 2 receptor CRA a precursor (22.2 kD) (Il2rg) alternative variant cSep08, mRNA.
<a href="#">Il2rg</a>	<a href="#">Il2rg.dSep08</a>	<a href="#">140924</a>	1239	712	3	148	interleukin 2 receptor (16.3 kD) (Il2rg) alternative variant dSep08, mRNA.
<a href="#">Il2rg</a>	<a href="#">Il2rg.eSep08</a>	<a href="#">140924</a>	1077	740	2	61	interleukin 2 receptor (Il2rg) alternative variant eSep08, mRNA.
<a href="#">Il3ra</a>	<a href="#">Il3ra.aSep08</a>	<a href="#">246144</a>	6092	1655	9	275	interleukin 3 receptor, alpha chain (29.9 kD) (Il3ra) alternative variant aSep08, mRNA.
<a href="#">Il3ra</a>	<a href="#">Il3ra.cSep08</a>	<a href="#">246144</a>	4133	643	6	196	interleukin 3 receptor, alpha chain (Il3ra) alternative variant cSep08, mRNA.
<a href="#">Il3ra</a>	<a href="#">Il3ra.dSep08</a>	<a href="#">246144</a>	1266	646	5	155	interleukin 3 receptor, alpha chain (Il3ra) alternative variant dSep08, mRNA.
<a href="#">Il4ra</a>	<a href="#">Il4ra.bSep08</a>	<a href="#">25084</a>	4139	1241	2	306	interleukin 4 receptor, alpha (Il4ra) alternative variant bSep08, mRNA.
<a href="#">Il4ra</a>	<a href="#">Il4ra.cSep08</a>	<a href="#">25084</a>	1519	773	1	257	interleukin 4 receptor, alpha (Il4ra) alternative variant cSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.bSep08</a>	<a href="#">25205</a>	12150	5949	6	421	interleukin 6 signal transducer (Il6st) alternative variant bSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.cSep08</a>	<a href="#">25205</a>	15871	749	6	176	interleukin 6 signal transducer (Il6st) alternative variant cSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.dSep08</a>	<a href="#">25205</a>	5290	479	4	159	interleukin 6 signal transducer (Il6st) alternative variant dSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.eSep08</a>	<a href="#">25205</a>	14808	633	4	138	interleukin 6 signal transducer (Il6st) alternative variant eSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.fSep08</a>	<a href="#">25205</a>	3037	732	3	155	interleukin 6 signal transducer (Il6st) alternative variant fSep08, mRNA.
<a href="#">Il6st</a>	<a href="#">Il6st.gSep08</a>	<a href="#">25205</a>	14517	394	5	36	interleukin 6 signal transducer (Il6st) alternative variant gSep08, mRNA.
<a href="#">Il9</a>	<a href="#">Il9.bSep08</a>	<a href="#">116558</a>	46485	527	1	141	interleukin 9 (Il9) alternative variant bSep08, mRNA.
<a href="#">Il10rb</a>	<a href="#">Il10rb.bSep08</a>	<a href="#">304091</a>	13205	834	2	253	interleukin 10 receptor, beta (Il10rb) alternative variant bSep08, mRNA.
<a href="#">Il11ra1</a>	<a href="#">Il11ra1.bSep08</a>	<a href="#">245983</a>	6000	814	8	254	interleukin 11 receptor, alpha chain 1 (Il11ra1) alternative variant bSep08, mRNA.
<a href="#">Il11ra1</a>	<a href="#">Il11ra1.cSep08</a>	<a href="#">245983</a>	1799	601	5	199	interleukin 11 receptor, alpha chain 1 (Il11ra1) alternative variant cSep08, mRNA.
<a href="#">Il11ra1</a>	<a href="#">Il11ra1.eSep08</a>	<a href="#">245983</a>	1273	434	2	62	interleukin 11 receptor, alpha chain 1 (Il11ra1) alternative variant eSep08, mRNA.

<a href="#">Il12a</a>	<a href="#">Il12a.aSep08</a>	<a href="#">84405</a>	3535	761		101	interleukin 12a (11.6 kD) (Il12a) mRNA.
<a href="#">Il12rb1</a>	<a href="#">Il12rb1.aSep08</a>	<a href="#">171333</a>	4294	903		301	interleukin 12 receptor, beta 1 (Il12rb1) mRNA.
<a href="#">Il12rb2</a>	<a href="#">Il12rb2.aSep08</a>	<a href="#">171334</a>	40214	770		256	interleukin 12 receptor, beta 2 (Il12rb2) mRNA.
<a href="#">Il15</a>	<a href="#">Il15.bSep08</a>	<a href="#">25670</a>	6410	744	4	67	interleukin 15 (7.8 kD) (Il15) alternative variant bSep08, mRNA.
<a href="#">Il16</a>	<a href="#">Il16.bSep08</a>	<a href="#">116996</a>	22200	1662	7	242	interleukin 16 (Il16) alternative variant bSep08, mRNA.
<a href="#">Il16</a>	<a href="#">Il16.cSep08</a>	<a href="#">116996</a>	32058	569	2	78	interleukin 16 (Il16) alternative variant cSep08, mRNA.
<a href="#">Il17b</a>	<a href="#">Il17b.aSep08</a>	<a href="#">116472</a>	4382	694	2	180	interleukin 17B (20.4 kD) (Il17b) alternative variant aSep08, mRNA.
<a href="#">Il17b</a>	<a href="#">Il17b.bSep08</a>	<a href="#">116472</a>	2766	971	1	122	interleukin 17B (13.9 kD) (Il17b) alternative variant bSep08, mRNA.
<a href="#">Il17d</a>	<a href="#">Il17d.aSep08</a>	<a href="#">691799</a>	1996	1613	3	108	interleukin 17D (Il17d) alternative variant aSep08, mRNA.
<a href="#">Il17ra</a>	<a href="#">Il17ra.bSep08</a>	<a href="#">312679</a>	8948	3331	8	684	interleukin 17 receptor A (Il17ra) alternative variant bSep08, mRNA.
<a href="#">Il17rc</a>	<a href="#">Il17rc.aSep08</a>	<a href="#">297520</a>	6402	1316	10	379	interleukin 17 receptor C CRA c (42.2 kD) (Il17rc) alternative variant aSep08, mRNA.
<a href="#">Il17rc</a>	<a href="#">Il17rc.bSep08</a>	<a href="#">297520</a>	6692	858	11	285	interleukin 17 receptor C (Il17rc) alternative variant bSep08, mRNA.
<a href="#">Il17rc</a>	<a href="#">Il17rc.cSep08</a>	<a href="#">297520</a>	1584	827	5	175	interleukin 17 receptor C (19.0 kD) (Il17rc) alternative variant cSep08, mRNA.
<a href="#">Il17rc</a>	<a href="#">Il17rc.dSep08</a>	<a href="#">297520</a>	10504	976	12	166	interleukin 17 receptor C (18.4 kD) (Il17rc) alternative variant dSep08, mRNA.
<a href="#">Il18</a>	<a href="#">Il18.bSep08</a>	<a href="#">29197</a>	4180	715	3	149	interleukin 18 (17.2 kD) (Il18) alternative variant bSep08, mRNA.
<a href="#">Il18</a>	<a href="#">Il18.cSep08</a>	<a href="#">29197</a>	10286	1093	7	147	interleukin 18 (17.1 kD) (Il18) alternative variant cSep08, mRNA.
<a href="#">Il18</a>	<a href="#">Il18.dSep08</a>	<a href="#">29197</a>	9817	745	2	71	interleukin 18 (7.5 kD) (Il18) alternative variant dSep08, mRNA.
<a href="#">Il18</a>	<a href="#">Il18.eSep08</a>	<a href="#">29197</a>	3562	340	3	66	interleukin 18 (Il18) alternative variant eSep08, mRNA.
<a href="#">Il18bp</a>	<a href="#">Il18bp.bSep08</a>	<a href="#">84388</a>	736	438	2	39	interleukin 18 binding protein (Il18bp) alternative variant bSep08, mRNA.
<a href="#">Il20rb</a>	<a href="#">Il20rb.aSep08</a>	<a href="#">501043</a>	12037	1267		171	interleukin 20 receptor beta (18.9 kD) (Il20rb) mRNA.
<a href="#">Il21r</a>	<a href="#">Il21r.bSep08</a>	<a href="#">308977</a>	2307	812	1	270	interleukin 21 receptor (Il21r) alternative variant bSep08, mRNA.
<a href="#">Il22ra1</a>	<a href="#">Il22ra1.aSep08</a>	<a href="#">362629</a>	9780	622		159	interleukin 22 receptor, alpha 1 (Il22ra1) mRNA.
<a href="#">Il27ra</a>	<a href="#">Il27ra.bSep08</a>	<a href="#">288905</a>	3611	1214	1	257	interleukin 27 receptor, alpha (Il27ra) alternative variant bSep08, mRNA.
<a href="#">Il28ra</a>	<a href="#">Il28ra.aSep08</a>	<a href="#">362628</a>	14329	496		165	interleukin 28 receptor alpha (Il28ra) mRNA.
<a href="#">Il33</a>	<a href="#">Il33.aSep08</a>	<a href="#">361749</a>	35610	1144	9	264	interleukin 33 (29.5 kD) (Il33) alternative variant aSep08, mRNA.
<a href="#">Il33</a>	<a href="#">Il33.cSep08</a>	<a href="#">361749</a>	7114	637	6	165	interleukin 33 (Il33) alternative variant cSep08, mRNA.
<a href="#">Ilf2</a>	<a href="#">Ilf2.bSep08</a>	<a href="#">310612</a>	11839	1604	14	422	interleukin enhancer binding factor 2 (Ilf2) alternative variant bSep08, mRNA.
<a href="#">Ilf2</a>	<a href="#">Ilf2.cSep08</a>	<a href="#">310612</a>	4689	1114	8	334	interleukin enhancer binding factor 2 (Ilf2) alternative variant cSep08, mRNA.

<a href="#">Ilf2</a>	<a href="#">Ilf2.dSep08</a>	<a href="#">310612</a>	6057	768	5	119	interleukin enhancer binding factor 2 (Ilf2) alternative variant dSep08, mRNA.
<a href="#">Ilf2</a>	<a href="#">Ilf2.eSep08</a>	<a href="#">310612</a>	5252	276	4	80	interleukin enhancer binding factor 2 (Ilf2) alternative variant eSep08, mRNA.
<a href="#">Ilf2</a>	<a href="#">Ilf2.fSep08</a>	<a href="#">310612</a>	6171	675	5	70	interleukin enhancer binding factor 2 (Ilf2) alternative variant fSep08, mRNA.
<a href="#">Ilf3</a>	<a href="#">Ilf3.bSep08</a>	<a href="#">84472</a>	13895	2306	12	537	interleukin enhancer binding factor 3 (Ilf3) alternative variant bSep08, mRNA.
<a href="#">Ilf3</a>	<a href="#">Ilf3.cSep08</a>	<a href="#">84472</a>	4536	2416	7	296	interleukin enhancer binding factor 3 (Ilf3) alternative variant cSep08, mRNA.
<a href="#">Ilf3</a>	<a href="#">Ilf3.dSep08</a>	<a href="#">84472</a>	27851	1073	8	286	interleukin enhancer binding factor 3 (Ilf3) alternative variant dSep08, mRNA.
<a href="#">Ilk</a>	<a href="#">Ilk.bSep08</a>	<a href="#">170922</a>	6112	1806	11	287	integrin linked kinase (Ilk) alternative variant bSep08, mRNA.
<a href="#">Ilk</a>	<a href="#">Ilk.cSep08</a>	<a href="#">170922</a>	5230	1734	6	228	integrin linked kinase (25.7 kD) (Ilk) alternative variant cSep08, mRNA.
<a href="#">Ilk</a>	<a href="#">Ilk.dSep08</a>	<a href="#">170922</a>	1252	840	4	150	integrin linked kinase (Ilk) alternative variant dSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.bSep08</a>	<a href="#">64538</a>	15316	683	6	227	integrin-linked kinase-associated serine threonine phosphatase 2C (Ilkap) alternative variant bSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.cSep08</a>	<a href="#">64538</a>	6559	1039	5	193	integrin-linked kinase-associated serine threonine phosphatase 2C precursor (21.6 kD) (Ilkap) alternative variant cSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.dSep08</a>	<a href="#">64538</a>	9355	769	7	150	integrin-linked kinase-associated serine threonine phosphatase 2C (Ilkap) alternative variant dSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.eSep08</a>	<a href="#">64538</a>	5130	729	4	148	integrin-linked kinase-associated serine threonine phosphatase 2C (Ilkap) alternative variant eSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.fSep08</a>	<a href="#">64538</a>	15376	735	7	138	integrin-linked kinase-associated serine threonine phosphatase 2C (Ilkap) alternative variant fSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.gSep08</a>	<a href="#">64538</a>	2609	924	3	109	integrin-linked kinase-associated serine threonine phosphatase 2C (12.0 kD) (Ilkap) alternative variant gSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.hSep08</a>	<a href="#">64538</a>	14441	556	6	105	integrin-linked kinase-associated serine threonine phosphatase 2C CRA c (Ilkap) alternative variant hSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.iSep08</a>	<a href="#">64538</a>	11864	478	5	87	integrin-linked kinase-associated serine threonine phosphatase 2C CRA c (Ilkap) alternative variant iSep08, mRNA.
<a href="#">Ilkap</a>	<a href="#">Ilkap.kSep08</a>	<a href="#">64538</a>	601	367	2	71	integrin-linked kinase-associated serine threonine phosphatase 2C (7.8 kD) (Ilkap) alternative variant kSep08, mRNA.
<a href="#">Ilvbl</a>	<a href="#">Ilvbl.bSep08</a>	<a href="#">362843</a>	1080	714	1	156	ilvB (bacterial acetolactate synthase)-like (Ilvbl) alternative variant bSep08, mRNA.
<a href="#">Immp1l</a>	<a href="#">Immp1l.aSep08</a>	<a href="#">691145</a>	27746	474		157	IMP1 inner mitochondrial membrane peptidase-like (S. cerevisiae) (Immp1l) mRNA.
<a href="#">Immt</a>	<a href="#">Immt.bSep08</a>	<a href="#">312444</a>	36170	2490	13	717	inner membrane protein, mitochondrial (Immt) alternative variant bSep08, mRNA.



<a href="#">Immt</a>	<a href="#">Immt.cSep08</a>	<a href="#">312444</a>	23246	818	7	272	inner membrane protein, mitochondrial (Immt) alternative variant cSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.dSep08</a>	<a href="#">312444</a>	6462	387	4	128	inner membrane protein, mitochondrial (Immt) alternative variant dSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.eSep08</a>	<a href="#">312444</a>	4629	366	4	121	inner membrane protein, mitochondrial (Immt) alternative variant eSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.fSep08</a>	<a href="#">312444</a>	4418	417	3	77	inner membrane protein, mitochondrial (Immt) alternative variant fSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.gSep08</a>	<a href="#">312444</a>	15502	375	4	68	inner membrane protein, mitochondrial (Immt) alternative variant gSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.jSep08</a>	<a href="#">312444</a>	9802	364	4	13	inner membrane protein, mitochondrial (1.6 kD) (Immt) alternative variant jSep08, mRNA.
<a href="#">Immt</a>	<a href="#">Immt.kSep08</a>	<a href="#">312444</a>	1294	318	2	28	inner membrane protein, mitochondrial (3.1 kD) (Immt) alternative variant kSep08, mRNA.
<a href="#">Imp4</a>	<a href="#">Imp4.bSep08</a>	<a href="#">316317</a>	5777	2862	6	206	imp4 U3 small nucleolar ribonucleoprotein homolog (23.6 kD) (Imp4) alternative variant bSep08, complete mRNA.
<a href="#">Imp4</a>	<a href="#">Imp4.cSep08</a>	<a href="#">316317</a>	889	659	4	169	imp4 U3 small nucleolar ribonucleoprotein homolog CRA b (Imp4) alternative variant cSep08, mRNA.
<a href="#">Imp4</a>	<a href="#">Imp4.dSep08</a>	<a href="#">316317</a>	3672	718	2	75	imp4 U3 small nucleolar ribonucleoprotein (8.9 kD) (Imp4) alternative variant dSep08, mRNA.
<a href="#">Impa1</a>	<a href="#">Impa1.aSep08</a>	<a href="#">83523</a>	19399	1800	5	436	4-monophosphatase 1 (Impa1) alternative variant aSep08, mRNA.
<a href="#">Impa1</a>	<a href="#">Impa1.cSep08</a>	<a href="#">83523</a>	22380	1999	8	161	4-monophosphatase 1 (17.4 kD) (Impa1) alternative variant cSep08, complete mRNA.
<a href="#">Impa1</a>	<a href="#">Impa1.dSep08</a>	<a href="#">83523</a>	3598	820	3	101	putative protein (Impa1) alternative variant dSep08, mRNA.
<a href="#">Impa1</a>	<a href="#">Impa1.eSep08</a>	<a href="#">83523</a>	3291	1929	2	78	4-monophosphatase 1 (8.3 kD) (Impa1) alternative variant eSep08, mRNA.
<a href="#">Impa2</a>	<a href="#">Impa2.bSep08</a>	<a href="#">282636</a>	11103	722	4	154	inositol (myo)-1(or 4)-monophosphatase 2 (Impa2) alternative variant bSep08, mRNA.
<a href="#">Impa2</a>	<a href="#">Impa2.cSep08</a>	<a href="#">282636</a>	18507	392	3	130	inositol (myo)-1(or 4)-monophosphatase 2 (Impa2) alternative variant cSep08, mRNA.
<a href="#">Impad1</a>	<a href="#">Impad1.aSep08</a>	<a href="#">312952</a>	10860	2587		192	inositol monophosphatase (Impad1) mRNA.
<a href="#">Impdh1</a>	<a href="#">Impdh1.bSep08</a>	<a href="#">362329</a>	8681	1162	10	386	inosine 5'-phosphate dehydrogenase 1 (Impdh1) alternative variant bSep08, mRNA.
<a href="#">Impdh2</a>	<a href="#">Impdh2.bSep08</a>	<a href="#">301005</a>	4595	1800	14	366	dehydrogenase 2 (39.5 kD) (Impdh2) alternative variant bSep08, complete mRNA.
<a href="#">Impdh2</a>	<a href="#">Impdh2.cSep08</a>	<a href="#">301005</a>	746	669	2	115	inosine monophosphate dehydrogenase 2 (Impdh2) alternative variant cSep08, mRNA.
<a href="#">Impdh2</a>	<a href="#">Impdh2.dSep08</a>	<a href="#">301005</a>	2296	652	7	104	dehydrogenase 2 (11.0 kD) (Impdh2) alternative variant dSep08, mRNA.
<a href="#">Impg1</a>	<a href="#">Impg1.aSep08</a>	<a href="#">66014</a>	29423	401		89	interphotoreceptor matrix proteoglycan 1 (Impg1) mRNA.
<a href="#">Inca1</a>	<a href="#">Inca1.aSep08</a>	<a href="#">360555</a>	10345	700	2	232	inhibitor of CDK interacting with cyclin A1 (Inca1) alternative variant aSep08, mRNA.
<a href="#">Incenp</a>	<a href="#">Incenp.bSep08</a>	<a href="#">293733</a>	11885	1901	4	447	inner centromere protein (Incenp) alternative variant bSep08, mRNA.
<a href="#">Incenp</a>	<a href="#">Incenp.cSep08</a>	<a href="#">293733</a>	2700	570	2	50	inner centromere protein (Incenp) alternative variant cSep08, mRNA.

<a href="#">Ing3</a>	<a href="#">Ing3.bSep08</a>	<a href="#">312154</a>	16104	601	5	150	inhibitor of growth family, member 3 (Ing3) alternative variant bSep08, mRNA.
<a href="#">Ing3</a>	<a href="#">Ing3.cSep08</a>	<a href="#">312154</a>	17895	454	5	103	inhibitor of growth family, member 3 (Ing3) alternative variant cSep08, mRNA.
<a href="#">Ing3</a>	<a href="#">Ing3.dSep08</a>	<a href="#">312154</a>	2689	402	3	86	inhibitor of growth family, member 3 (10.4 kD) (Ing3) alternative variant dSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.bSep08</a>	<a href="#">297597</a>	7839	878	6	244	inhibitor of growth family, member 4 (28.1 kD) (Ing4) alternative variant bSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.cSep08</a>	<a href="#">297597</a>	7827	869	6	222	inhibitor of growth family, member 4 (25.3 kD) (Ing4) alternative variant cSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.dSep08</a>	<a href="#">297597</a>	7682	715	6	217	inhibitor of growth family, member 4 (Ing4) alternative variant dSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.eSep08</a>	<a href="#">297597</a>	6292	595	4	198	inhibitor of growth family, member 4 (Ing4) alternative variant eSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.fSep08</a>	<a href="#">297597</a>	8081	1002	6	164	inhibitor of growth family, member 4 (19.1 kD) (Ing4) alternative variant fSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.gSep08</a>	<a href="#">297597</a>	8117	1353	5	163	inhibitor of growth family, member 4 (18.8 kD) (Ing4) alternative variant gSep08, mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.hSep08</a>	<a href="#">297597</a>	7000	775	2	158	inhibitor of growth family, member 4 (18.4 kD) (Ing4) alternative variant hSep08, complete mRNA.
<a href="#">Ing4</a>	<a href="#">Ing4.iSep08</a>	<a href="#">297597</a>	1162	1039	2	72	inhibitor of growth family, member 4 (7.9 kD) (Ing4) alternative variant iSep08, mRNA.
<a href="#">Inoc1</a>	<a href="#">Inoc1.aSep08</a>	<a href="#">296084</a>	37003	2319	12	493	INO80 complex homolog 1 ( <i>S. cerevisiae</i> ) (Inoc1) alternative variant aSep08, mRNA.
<a href="#">Inpp1</a>	<a href="#">Inpp1.aSep08</a>	<a href="#">316376</a>	30008	2162	4	396	inositol polyphosphate-1-phosphatase (43.4 kD) (Inpp1) alternative variant aSep08, mRNA.
<a href="#">Inpp1</a>	<a href="#">Inpp1.cSep08</a>	<a href="#">316376</a>	29005	1045	4	230	inositol polyphosphate-1-phosphatase (Inpp1) alternative variant cSep08, mRNA.
<a href="#">Inpp4a</a>	<a href="#">Inpp4a.bSep08</a>	<a href="#">80849</a>	16650	1006	9	278	inositol polyphosphate-4-phosphatase, type 1 (Inpp4a) alternative variant bSep08, mRNA.
<a href="#">Inpp5a</a>	<a href="#">Inpp5a.bSep08</a>	<a href="#">365382</a>	156942	776	8	258	inositol polyphosphate-5-phosphatase A (Inpp5a) alternative variant bSep08, mRNA.
<a href="#">Inpp5a</a>	<a href="#">Inpp5a.cSep08</a>	<a href="#">365382</a>	121949	659	7	219	inositol polyphosphate-5-phosphatase A (Inpp5a) alternative variant cSep08, mRNA.
<a href="#">Inpp5a</a>	<a href="#">Inpp5a.dSep08</a>	<a href="#">365382</a>	20135	369	3	64	inositol polyphosphate-5-phosphatase A (Inpp5a) alternative variant dSep08, mRNA.
<a href="#">Inpp5b</a>	<a href="#">Inpp5b.aSep08</a>	<a href="#">362590</a>	59932	3211	20	841	inositol polyphosphate-5-phosphatase B (Inpp5b) alternative variant aSep08, mRNA.
<a href="#">Inpp5b</a>	<a href="#">Inpp5b.bSep08</a>	<a href="#">362590</a>	30457	545	5	181	inositol polyphosphate-5-phosphatase B (Inpp5b) alternative variant bSep08, mRNA.
<a href="#">Inpp5b</a>	<a href="#">Inpp5b.cSep08</a>	<a href="#">362590</a>	6681	507	5	90	inositol polyphosphate-5-phosphatase B (Inpp5b) alternative variant cSep08, mRNA.
<a href="#">Inpp5b</a>	<a href="#">Inpp5b.dSep08</a>	<a href="#">362590</a>	1962	995	1	59	inositol polyphosphate-5-phosphatase B (6.7 kD) (Inpp5b) alternative variant dSep08, mRNA.
<a href="#">Inpp5f</a>	<a href="#">Inpp5f.bSep08</a>	<a href="#">309008</a>	5196	859	5	156	inositol polyphosphate-5-phosphatase F (Inpp5f) alternative variant bSep08, mRNA.

<a href="#">Inpp5f</a>	<a href="#">Inpp5f.eSep08</a>	<a href="#">309008</a>	725	314	2	34	inositol polyphosphate-5-phosphatase F (Inpp5f) alternative variant eSep08, mRNA.
<a href="#">Inpp1</a>	<a href="#">Inpp1.bSep08</a>	<a href="#">65038</a>	3984	645	4	113	inositol polyphosphate phosphatase-like 1 (Inpp1) alternative variant bSep08, mRNA.
<a href="#">Ins2</a>	<a href="#">Ins2.aSep08</a>	<a href="#">24506</a>	1077	459	3	141	insulin 2 (Ins2) alternative variant aSep08, mRNA.
<a href="#">Ins145_P3_rec.0</a>	<a href="#">Ins145_P3_rec.0.aSep08</a>		229802	443		123	ryanodine receptor (Ins145_P3_rec.0) mRNA.
<a href="#">Ins145_P3_rec.1</a>	<a href="#">Ins145_P3_rec.1.aSep08</a>		17144	577		192	inositol 1 receptor CRA b (Ins145_P3_rec.1) mRNA.
<a href="#">Insc</a>	<a href="#">Insc.bSep08</a>	<a href="#">293166</a>	48031	513	1	170	inscuteable homolog (Drosophila) (Insc) alternative variant bSep08, mRNA.
<a href="#">Insig1</a>	<a href="#">Insig1.bSep08</a>	<a href="#">64194</a>	2313	579	2	182	insulin induced gene 1 (Insig1) alternative variant bSep08, mRNA.
<a href="#">Insig1</a>	<a href="#">Insig1.cSep08</a>	<a href="#">64194</a>	8137	2309	5	165	insulin induced gene 1 (Insig1) alternative variant cSep08, mRNA.
<a href="#">Insig2</a>	<a href="#">Insig2.bSep08</a>	<a href="#">288985</a>	27526	2514	2	225	insulin induced gene 2 (25.0 kD) (Insig2) alternative variant bSep08, mRNA.
<a href="#">Insl3</a>	<a href="#">Insl3.bSep08</a>	<a href="#">114215</a>	1576	1444	1	97	insulin-like 3 (10.5 kD) (Insl3) alternative variant bSep08, mRNA.
<a href="#">Insrr</a>	<a href="#">Insrr.aSep08</a>	<a href="#">60663</a>	5899	1410	1	469	insulin receptor-related receptor (Insrr) alternative variant aSep08, mRNA.
<a href="#">Insrr</a>	<a href="#">Insrr.bSep08</a>	<a href="#">60663</a>	5899	1233	2	410	insulin receptor-related receptor (44.8 kD) (Insrr) alternative variant bSep08, mRNA.
<a href="#">Integrase.2</a>	<a href="#">Integrase.2.aSep08</a>		1508	1365		115	integrase like (12.8 kD) (Integrase.2) mRNA.
<a href="#">Integrin_alpha.0</a>	<a href="#">Integrin_alpha.0.aSep08</a>		2854	619		132	integrin alpha 2b (14.9 kD) (Integrin_alpha.0) mRNA.
<a href="#">Integrin_alpha.1</a>	<a href="#">Integrin_alpha.1.aSep08</a>		11140	2954	5	194	integrin (Integrin_alpha.1) alternative variant aSep08, mRNA.
<a href="#">Integrin_alpha.1</a>	<a href="#">Integrin_alpha.1.bSep08</a>		6259	409	3	71	integrin alpha 6 CRA c (Integrin_alpha.1) alternative variant bSep08, mRNA.
<a href="#">Integrin_alpha2.0</a>	<a href="#">Integrin_alpha2.0.aSep08</a>		60602	758		252	integrin alpha 8 CRA a (Integrin_alpha2.0) mRNA.
<a href="#">Integrin_b_cyt.0</a>	<a href="#">Integrin_b_cyt.0.aSep08</a>		4414	2553		78	putative protein (11.6 kD) (Integrin_b_cyt.0) mRNA.
<a href="#">Integrin_b_cyt.1</a>	<a href="#">Integrin_b_cyt.1.aSep08</a>		1048	665		145	integrin (Integrin_b_cyt.1) mRNA.
<a href="#">Ints3</a>	<a href="#">Ints3.aSep08</a>	<a href="#">361988</a>	18030	2363	17	564	integrator complex subunit 3 (Ints3) alternative variant aSep08, mRNA.
<a href="#">Ints3</a>	<a href="#">Ints3.bSep08</a>	<a href="#">361988</a>	12443	842	7	247	integrator complex subunit 3 (Ints3) alternative variant bSep08, mRNA.
<a href="#">Ints4</a>	<a href="#">Ints4.aSep08</a>	<a href="#">308837</a>	63755	3152		583	integrator complex subunit 4 (65.9 kD) (Ints4) mRNA.
<a href="#">Ints6</a>	<a href="#">Ints6.aSep08</a>	<a href="#">361057</a>	52033	3261	15	768	integrator complex (Ints6) alternative variant aSep08, mRNA.
<a href="#">Ints6</a>	<a href="#">Ints6.cSep08</a>	<a href="#">361057</a>	9673	1568	5	288	integrator complex (32.6 kD) (Ints6) alternative variant cSep08, mRNA.
<a href="#">Ints6</a>	<a href="#">Ints6.dSep08</a>	<a href="#">361057</a>	1909	776	3	226	putative protein (Ints6) alternative variant dSep08, mRNA.

<a href="#">Ints6</a>	<a href="#">Ints6.eSep08</a>	<a href="#">361057</a>	26826	700	6	182	integrator complex (20.1 kD) (Ints6) alternative variant eSep08, mRNA.
<a href="#">Ints7</a>	<a href="#">Ints7.aSep08</a>	<a href="#">289382</a>	17641	795		247	integrator complex subunit 7 (Ints7) mRNA.
<a href="#">Ints9</a>	<a href="#">Ints9.aSep08</a>	<a href="#">290322</a>	83591	2612	17	658	integrator complex subunit 9 (74.1 kD) (Ints9) alternative variant aSep08, mRNA.
<a href="#">Ints10</a>	<a href="#">Ints10.bSep08</a>	<a href="#">290679</a>	7335	592	5	164	integrator complex subunit 10 (Ints10) alternative variant bSep08, mRNA.
<a href="#">Ints10</a>	<a href="#">Ints10.cSep08</a>	<a href="#">290679</a>	5757	730	2	80	integrator complex subunit 10 (9.3 kD) (Ints10) alternative variant cSep08, mRNA.
<a href="#">Ints12</a>	<a href="#">Ints12.bSep08</a>	<a href="#">295448</a>	14607	873	6	248	integrator complex subunit 12 (Ints12) alternative variant bSep08, mRNA.
<a href="#">Intu</a>	<a href="#">Intu.aSep08</a>	<a href="#">361938</a>	33644	922		281	inturned planar cell polarity effector homolog (Drosophila) (Intu) mRNA.
<a href="#">Invs</a>	<a href="#">Invs.cSep08</a>	<a href="#">313228</a>	9498	648	5	156	inversin (Invs) alternative variant cSep08, mRNA.
<a href="#">Invs</a>	<a href="#">Invs.dSep08</a>	<a href="#">313228</a>	7949	425	2	141	inversin (Invs) alternative variant dSep08, mRNA.
<a href="#">Invs</a>	<a href="#">Invs.eSep08</a>	<a href="#">313228</a>	5623	2642	3	125	inversin (Invs) alternative variant eSep08, mRNA.
<a href="#">Invs</a>	<a href="#">Invs.fSep08</a>	<a href="#">313228</a>	8965	373	3	124	inversin (Invs) alternative variant fSep08, mRNA.
<a href="#">Ion_trans.0</a>	<a href="#">Ion_trans.0.aSep08</a>		14266	583		193	sodium channel Nav1.5a (Ion_trans.0) mRNA.
<a href="#">Ipmk</a>	<a href="#">Ipmk.bSep08</a>	<a href="#">171458</a>	31578	1614	4	296	inositol polyphosphate multikinase (33.7 kD) (Ipmk) alternative variant bSep08, mRNA.
<a href="#">Ipmk</a>	<a href="#">Ipmk.cSep08</a>	<a href="#">171458</a>	26257	935	5	173	inositol polyphosphate multikinase (19.7 kD) (Ipmk) alternative variant cSep08, mRNA.
<a href="#">Ipo4</a>	<a href="#">Ipo4.bSep08</a>	<a href="#">290228</a>	3709	1340	9	382	importin 4 (Ipo4) alternative variant bSep08, mRNA.
<a href="#">Ipo4</a>	<a href="#">Ipo4.cSep08</a>	<a href="#">290228</a>	1992	728	8	201	importin 4 (Ipo4) alternative variant cSep08, mRNA.
<a href="#">Ipo5</a>	<a href="#">Ipo5.aSep08</a>	<a href="#">306182</a>	35724	4552	16	1099	importin 5 (Ipo5) alternative variant aSep08, mRNA.
<a href="#">Ipo5</a>	<a href="#">Ipo5.bSep08</a>	<a href="#">306182</a>	2026	422	1	90	importin 5 (Ipo5) alternative variant bSep08, mRNA.
<a href="#">Ipo7</a>	<a href="#">Ipo7.bSep08</a>	<a href="#">308939</a>	1995	266	2	76	importin 7 (Ipo7) alternative variant bSep08, mRNA.
<a href="#">Ipo9</a>	<a href="#">Ipo9.bSep08</a>	<a href="#">304817</a>	38511	1988	11	580	importin 9 (Ipo9) alternative variant bSep08, mRNA.
<a href="#">Ipo9</a>	<a href="#">Ipo9.cSep08</a>	<a href="#">304817</a>	12221	822	5	241	importin 9 (Ipo9) alternative variant cSep08, mRNA.
<a href="#">Ipo9</a>	<a href="#">Ipo9.dSep08</a>	<a href="#">304817</a>	5111	2543	6	235	importin 9 (27.1 kD) (Ipo9) alternative variant dSep08, mRNA.
<a href="#">Ipo9</a>	<a href="#">Ipo9.eSep08</a>	<a href="#">304817</a>	1110	519	2	109	importin 9 (Ipo9) alternative variant eSep08, mRNA.
<a href="#">Ipo11</a>	<a href="#">Ipo11.aSep08</a>	<a href="#">310056</a>	23147	630		169	importin 11 (Ipo11) mRNA.
<a href="#">Ipo13</a>	<a href="#">Ipo13.bSep08</a>	<a href="#">116458</a>	3629	759	7	252	importin 13 (Ipo13) alternative variant bSep08, mRNA.
<a href="#">Ipo13</a>	<a href="#">Ipo13.cSep08</a>	<a href="#">116458</a>	3049	759	6	182	importin 13 (Ipo13) alternative variant cSep08, mRNA.
<a href="#">Ipo13</a>	<a href="#">Ipo13.dSep08</a>	<a href="#">116458</a>	1046	833	3	67	importin 13 (Ipo13) alternative variant dSep08, mRNA.
<a href="#">Ipp</a>	<a href="#">Ipp.bSep08</a>	<a href="#">298439</a>	9538	559	4	122	IAP promoted placental gene (Ipp) alternative variant bSep08, mRNA.
<a href="#">IQ.0</a>	<a href="#">IQ.0.aSep08</a>		1047	379		126	myosin 5B CRA b (IQ.0) mRNA.
<a href="#">IQ.1</a>	<a href="#">IQ.1.aSep08</a>		1269	378		87	IQ motif containing e (IQ.1) mRNA.
<a href="#">IQ.2</a>	<a href="#">IQ.2.aSep08</a>		35262	4266	3	1047	myosin 18A (IQ.2) alternative variant aSep08, mRNA.
<a href="#">IQ.2</a>	<a href="#">IQ.2.bSep08</a>		15016	1269	2	261	CRA a (IQ.2) alternative variant bSep08, mRNA.
<a href="#">IQ.2</a>	<a href="#">IQ.2.cSep08</a>		886	574	2	73	putative protein (IQ.2) alternative variant cSep08, mRNA.

<a href="#">IQ.2</a>	<a href="#">IQ.2.dSep08</a>		1412	328	1	18	putative protein (2.4 kD) (IQ.2) alternative variant dSep08, mRNA.
<a href="#">IQ.3</a>	<a href="#">IQ.3.aSep08</a>		3476	703		124	myosin-xix (IQ.3) alternative variant aSep08, mRNA.
<a href="#">IQ.4</a>	<a href="#">IQ.4.aSep08</a>		17767	1066	9	354	myosin Ib (IQ.4) alternative variant aSep08, mRNA.
<a href="#">IQ.4</a>	<a href="#">IQ.4.bSep08</a>		4793	816	1	105	myosin ib (IQ.4) alternative variant bSep08, mRNA.
<a href="#">IQ.5</a>	<a href="#">IQ.5.aSep08</a>		1267	267		89	calmodulin binding transcription activator 1 like (IQ.5) mRNA.
<a href="#">IQ.6</a>	<a href="#">IQ.6.aSep08</a>		2774	618		206	iqcal iq aaa domain-containing protein homolog (IQ.6) mRNA.
<a href="#">IQ.7</a>	<a href="#">IQ.7.aSep08</a>		18702	856	7	284	IQ motif containing GTPase activating protein 2 (IQ.7) alternative variant aSep08, mRNA.
<a href="#">IQ.8</a>	<a href="#">IQ.8.aSep08</a>		8738	244		66	leucine-rich repeats IQ motif containing 4 like (IQ.8) mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.bSep08</a>	<a href="#">303915</a>	29903	1337	6	306	IQ calmodulin-binding motif containing 1 (lqcb1) alternative variant bSep08, mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.cSep08</a>	<a href="#">303915</a>	20869	798	7	235	IQ calmodulin-binding motif containing 1 (lqcb1) alternative variant cSep08, mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.dSep08</a>	<a href="#">303915</a>	20823	822	8	215	IQ calmodulin-binding motif containing 1 (lqcb1) alternative variant dSep08, mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.eSep08</a>	<a href="#">303915</a>	16346	699	7	182	IQ calmodulin-binding motif containing 1 (lqcb1) alternative variant eSep08, mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.fSep08</a>	<a href="#">303915</a>	3238	391	3	77	IQ calmodulin-binding motif containing 1 (lqcb1) alternative variant fSep08, mRNA.
<a href="#">lqcb1</a>	<a href="#">lqcb1.gSep08</a>	<a href="#">303915</a>	849	383	2	7	IQ calmodulin-binding motif containing 1 (0.9 kD) (lqcb1) alternative variant gSep08, mRNA.
<a href="#">lqcd</a>	<a href="#">lqcd.aSep08</a>	<a href="#">498184</a>	11489	2113	5	366	IQ motif containing D (41.6 kD) (lqcd) alternative variant aSep08, mRNA.
<a href="#">lqcd</a>	<a href="#">lqcd.bSep08</a>	<a href="#">498184</a>	10180	563	1	187	IQ motif containing D (lqcd) alternative variant bSep08, mRNA.
<a href="#">lqce</a>	<a href="#">lqce.aSep08</a>	<a href="#">304318</a>	16214	1090		363	IQ motif containing E (lqce) mRNA.
<a href="#">lqcg</a>	<a href="#">lqcg.bSep08</a>	<a href="#">363796</a>	15248	747	1	167	IQ motif containing G (lqcg) alternative variant bSep08, mRNA.
<a href="#">lqch</a>	<a href="#">lqch.bSep08</a>	<a href="#">300776</a>	11337	727	2	242	IQ motif containing H (lqch) alternative variant bSep08, mRNA.
<a href="#">lqgap1</a>	<a href="#">lqgap1.bSep08</a>	<a href="#">361598</a>	5137	610	4	197	IQ motif containing GTPase activating protein 1 (lqgap1) alternative variant bSep08, mRNA.
<a href="#">lqgap3</a>	<a href="#">lqgap3.aSep08</a>	<a href="#">310621</a>	10949	1758		576	IQ motif containing GTPase activating protein 3 (lqgap3) alternative variant aSep08, mRNA.
<a href="#">lqwd1</a>	<a href="#">lqwd1.aSep08</a>	<a href="#">289181</a>	96496	2468	17	721	IQ motif and WD repeats 1 (lqwd1) alternative variant aSep08, mRNA.
<a href="#">lqwd1</a>	<a href="#">lqwd1.bSep08</a>	<a href="#">289181</a>	36697	741	7	246	IQ motif and WD repeats 1 (lqwd1) alternative variant bSep08, mRNA.
<a href="#">lqwd1</a>	<a href="#">lqwd1.cSep08</a>	<a href="#">289181</a>	41744	720	6	240	IQ motif and WD repeats 1 (lqwd1) alternative variant cSep08, mRNA.
<a href="#">Irak1</a>	<a href="#">Irak1.bSep08</a>	<a href="#">363520</a>	6056	1207	7	348	receptor-associated kinase 1 (Irak1) alternative variant bSep08, mRNA.

<a href="#">Irak1</a>	<a href="#">Irak1.cSep08</a>	<a href="#">363520</a>	1605	1138	3	173	receptor-associated kinase 1 (19.3 kD) (Irak1) alternative variant cSep08, mRNA.
<a href="#">Irak1</a>	<a href="#">Irak1.dSep08</a>	<a href="#">363520</a>	2877	427	4	141	receptor-associated kinase 1 (Irak1) alternative variant dSep08, mRNA.
<a href="#">Irak1bp1</a>	<a href="#">Irak1bp1.aSep08</a>	<a href="#">300862</a>	16520	1067	4	270	interleukin-1 receptor-associated kinase 1 binding protein 1 (Irak1bp1) alternative variant aSep08, mRNA.
<a href="#">Irak1bp1</a>	<a href="#">Irak1bp1.cSep08</a>	<a href="#">300862</a>	7074	419	2	62	interleukin-1 receptor-associated kinase 1 binding protein 1 (Irak1bp1) alternative variant cSep08, mRNA.
<a href="#">Irak1bp1</a>	<a href="#">Irak1bp1.dSep08</a>	<a href="#">300862</a>	16327	1195	5	42	interleukin-1 receptor-associated kinase 1 binding protein 1 (Irak1bp1) alternative variant dSep08, mRNA.
<a href="#">Irak2</a>	<a href="#">Irak2.bSep08</a>	<a href="#">362418</a>	27432	468	3	151	interleukin-1 receptor-associated kinase 2 (Irak2) alternative variant bSep08, mRNA.
<a href="#">Ireb2</a>	<a href="#">Ireb2.bSep08</a>	<a href="#">64831</a>	3829	3184	2	96	iron responsive element binding protein 2 (10.5 kD) (Ireb2) alternative variant bSep08, mRNA.
<a href="#">Ireb2</a>	<a href="#">Ireb2.cSep08</a>	<a href="#">64831</a>	2595	375	2	44	iron responsive element binding protein 2 (Ireb2) alternative variant cSep08, mRNA.
<a href="#">Irf1</a>	<a href="#">Irf1.bSep08</a>	<a href="#">24508</a>	1295	295	1	98	interferon regulatory factor 1 (Irf1) alternative variant bSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.bSep08</a>	<a href="#">292892</a>	4928	2291	7	360	interferon regulatory factor 3 CRA a (40.4 kD) (Irf3) alternative variant bSep08, complete mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.cSep08</a>	<a href="#">292892</a>	16955	794	6	196	interferon regulatory factor 3 (21.4 kD) (Irf3) alternative variant cSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.dSep08</a>	<a href="#">292892</a>	2852	1314	4	172	interferon regulatory factor 3 (Irf3) alternative variant dSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.eSep08</a>	<a href="#">292892</a>	3455	1491	5	158	interferon regulatory factor 3 (18.2 kD) (Irf3) alternative variant eSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.fSep08</a>	<a href="#">292892</a>	3126	734	6	143	interferon regulatory factor 3 (16.3 kD) (Irf3) alternative variant fSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.gSep08</a>	<a href="#">292892</a>	3258	824	5	142	interferon regulatory factor 3 (16.0 kD) (Irf3) alternative variant gSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.hSep08</a>	<a href="#">292892</a>	2694	715	4	141	interferon regulatory factor 3 (16.1 kD) (Irf3) alternative variant hSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.iSep08</a>	<a href="#">292892</a>	2807	802	4	141	interferon regulatory factor 3 (16.1 kD) (Irf3) alternative variant iSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.jSep08</a>	<a href="#">292892</a>	1428	468	3	134	interferon regulatory factor 3 (14.9 kD) (Irf3) alternative variant jSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.kSep08</a>	<a href="#">292892</a>	3776	427	4	72	putative protein human specific (Irf3) alternative variant kSep08, mRNA.
<a href="#">Irf3</a>	<a href="#">Irf3.nSep08</a>	<a href="#">292892</a>	1062	748	2	58	putative protein (Irf3) alternative variant nSep08, mRNA.
<a href="#">Irf5</a>	<a href="#">Irf5.bSep08</a>	<a href="#">296953</a>	1799	848	4	282	interferon regulatory factor 5 (Irf5) alternative variant bSep08, mRNA.
<a href="#">Irf5</a>	<a href="#">Irf5.cSep08</a>	<a href="#">296953</a>	626	416	1	138	interferon regulatory factor 5 (Irf5) alternative variant cSep08, mRNA.
<a href="#">Irf7</a>	<a href="#">Irf7.bSep08</a>	<a href="#">293624</a>	1473	766	4	203	interferon regulatory factor 7 (Irf7) alternative variant bSep08, mRNA.
<a href="#">Irs2</a>	<a href="#">Irs2.aSep08</a>	<a href="#">29376</a>	19954	2021		176	insulin receptor substrate 2 (Irs2) mRNA.

<a href="#">Irx1</a>	<a href="#">Irx1.aSep08</a>	<a href="#">306659</a>	5144	1659		418	iroquois related homeobox 1 (Drosophila) (Irx1) complete mRNA.
<a href="#">Irx2</a>	<a href="#">Irx2.cSep08</a>	<a href="#">306657</a>	2006	663	4	81	iroquois related homeobox 2 (Drosophila) (Irx2) alternative variant cSep08, mRNA.
<a href="#">Irx3</a>	<a href="#">Irx3.aSep08</a>	<a href="#">307721</a>	1341	682		103	iroquois related homeobox 3 (Drosophila) (Irx3) mRNA.
<a href="#">Irx4</a>	<a href="#">Irx4.cSep08</a>	<a href="#">306655</a>	6277	615	4	60	putative protein (6.6 kD) (Irx4) alternative variant cSep08, mRNA.
<a href="#">Irx4</a>	<a href="#">Irx4.dSep08</a>	<a href="#">306655</a>	946	477	2	31	iroquois 4a protein like (Irx4) alternative variant dSep08, mRNA.
<a href="#">Irx4</a>	<a href="#">Irx4.eSep08</a>	<a href="#">306655</a>	1524	1054	3	31	iroquois 4a protein like (Irx4) alternative variant eSep08, mRNA.
<a href="#">Isca1</a>	<a href="#">Isca1.bSep08</a>	<a href="#">290985</a>	11404	735	4	126	iron-sulfur cluster assembly 1 homolog (S. cerevisiae) (Isca1) alternative variant bSep08, mRNA.
<a href="#">Isca1</a>	<a href="#">Isca1.cSep08</a>	<a href="#">290985</a>	9761	878	3	86	iron-sulfur cluster assembly 1 homolog (S. cerevisiae) (Isca1) alternative variant cSep08, mRNA.
<a href="#">Iscu</a>	<a href="#">Iscu.bSep08</a>	<a href="#">288740</a>	1769	383	3	101	IscU iron-sulfur cluster scaffold homolog (E. coli) (Iscu) alternative variant bSep08, mRNA.
<a href="#">Iscu</a>	<a href="#">Iscu.cSep08</a>	<a href="#">288740</a>	1599	345	2	75	IscU iron-sulfur cluster scaffold homolog (E. coli) (Iscu) alternative variant cSep08, mRNA.
<a href="#">isg12(b)</a>	<a href="#">isg12(b).aSep08</a>	<a href="#">299269</a>	1467	540	1	136	putative ISG12(b) protein (15.3 kD) (isg12(b)) alternative variant aSep08, mRNA.
<a href="#">Isg2011</a>	<a href="#">Isg2011.bSep08</a>	<a href="#">361594</a>	6828	767	1	255	interferon stimulated exonuclease gene 20-like 1 (Isg2011) alternative variant bSep08, mRNA.
<a href="#">Isl1</a>	<a href="#">Isl1.bSep08</a>	<a href="#">64444</a>	4864	1716	3	209	ISL1 transcription factor, LIM/homeodomain 1 (23.5 kD) (Isl1) alternative variant bSep08, mRNA.
<a href="#">Isl1</a>	<a href="#">Isl1.cSep08</a>	<a href="#">64444</a>	2013	436	3	145	ISL1 transcription factor, LIM/homeodomain 1 (Isl1) alternative variant cSep08, mRNA.
<a href="#">Isl2</a>	<a href="#">Isl2.aSep08</a>	<a href="#">57233</a>	4011	1337	3	153	insulin related protein 2 (islet 2) (Isl2) alternative variant aSep08, mRNA.
<a href="#">Isl2</a>	<a href="#">Isl2.bSep08</a>	<a href="#">57233</a>	2093	595	2	126	insulin related protein 2 (islet 2) (Isl2) alternative variant bSep08, mRNA.
<a href="#">Isoc2b</a>	<a href="#">Isoc2b.bSep08</a>	<a href="#">361501</a>	13897	769	4	141	isochorismatase hydrolase (Isoc2b) alternative variant bSep08, mRNA.
<a href="#">Isoc2b</a>	<a href="#">Isoc2b.cSep08</a>	<a href="#">361501</a>	7240	699	2	128	putative protein of ancient origin (Isoc2b) alternative variant cSep08, mRNA.
<a href="#">Isoc2b</a>	<a href="#">Isoc2b.dSep08</a>	<a href="#">361501</a>	13843	530	2	121	putative cytoplasmic protein of ancient origin (13.4 kD) (Isoc2b) alternative variant dSep08, mRNA.
<a href="#">Isoc2b</a>	<a href="#">Isoc2b.eSep08</a>	<a href="#">361501</a>	21066	1237	2	69	putative protein (Isoc2b) alternative variant eSep08, mRNA.
<a href="#">Isy1</a>	<a href="#">Isy1.bSep08</a>	<a href="#">362394</a>	8512	790	6	165	ISY1 splicing factor homolog (S. cerevisiae) (Isy1) alternative variant bSep08, mRNA.
<a href="#">Isyna1</a>	<a href="#">Isyna1.bSep08</a>	<a href="#">290651</a>	2803	1920	10	441	myo-inositol 1-phosphate synthase A1 (48.0 kD) (Isyna1) alternative variant bSep08, complete mRNA.
<a href="#">Isyna1</a>	<a href="#">Isyna1.cSep08</a>	<a href="#">290651</a>	1545	1174	5	158	myo-inositol 1-phosphate synthase A1 (Isyna1) alternative variant cSep08, mRNA.
<a href="#">Isyna1</a>	<a href="#">Isyna1.dSep08</a>	<a href="#">290651</a>	941	705	3	148	myo-inositol 1-phosphate synthase A1 (16.0 kD) (Isyna1) alternative variant dSep08, mRNA.

<a href="#">Itfg1</a>	<a href="#">Itfg1.bSep08</a>	<a href="#">171083</a>	6232	877	4	133	integrin alpha fg-gap repeat containing 1 (Itfg1) alternative variant bSep08, mRNA.
<a href="#">Itfg1</a>	<a href="#">Itfg1.cSep08</a>	<a href="#">171083</a>	6637	772	3	99	integrin alpha fg-gap repeat containing 1 (11.4 kD) (Itfg1) alternative variant cSep08, mRNA.
<a href="#">Itfg2</a>	<a href="#">Itfg2.aSep08</a>	<a href="#">362441</a>	12501	1397	7	431	integrin alpha FG-GAP repeat containing 2 (Itfg2) alternative variant aSep08, mRNA.
<a href="#">Itfg2</a>	<a href="#">Itfg2.bSep08</a>	<a href="#">362441</a>	8642	619	1	205	integrin alpha FG-GAP repeat containing 2 (Itfg2) alternative variant bSep08, mRNA.
<a href="#">Itfg3</a>	<a href="#">Itfg3.bSep08</a>	<a href="#">360502</a>	25803	763	3	192	integrin alpha FG-GAP repeat containing 3 (Itfg3) alternative variant bSep08, mRNA.
<a href="#">Itfg3</a>	<a href="#">Itfg3.cSep08</a>	<a href="#">360502</a>	25262	837	5	109	oculocerebrorenal syndrome of Lowe like (Itfg3) alternative variant cSep08, mRNA.
<a href="#">Itfg3</a>	<a href="#">Itfg3.dSep08</a>	<a href="#">360502</a>	23103	415	2	73	putative protein (Itfg3) alternative variant dSep08, mRNA.
<a href="#">Itga1</a>	<a href="#">Itga1.bSep08</a>	<a href="#">25118</a>	12615	760	5	248	integrin alpha 1 (Itga1) alternative variant bSep08, mRNA.
<a href="#">Itga1</a>	<a href="#">Itga1.cSep08</a>	<a href="#">25118</a>	15502	352	2	82	integrin alpha 1 (Itga1) alternative variant cSep08, mRNA.
<a href="#">Itga2</a>	<a href="#">Itga2.aSep08</a>	<a href="#">170921</a>	4435	508		168	integrin, alpha 2 (Itga2) mRNA.
<a href="#">Itga3</a>	<a href="#">Itga3.bSep08</a>	<a href="#">360606</a>	10961	1769	1	209	integrin alpha 3 (Itga3) alternative variant bSep08, mRNA.
<a href="#">Itga5</a>	<a href="#">Itga5.bSep08</a>	<a href="#">315346</a>	2270	374	4	124	integrin alpha 5 (fibronectin receptor alpha) (Itga5) alternative variant bSep08, mRNA.
<a href="#">Itga6</a>	<a href="#">Itga6.aSep08</a>	<a href="#">114517</a>	19337	1871	12	623	integrin, alpha 6 (Itga6) alternative variant aSep08, mRNA.
<a href="#">Itga6</a>	<a href="#">Itga6.bSep08</a>	<a href="#">114517</a>	1532	381	1	111	integrin, alpha 6 (Itga6) alternative variant bSep08, mRNA.
<a href="#">Itga7</a>	<a href="#">Itga7.bSep08</a>	<a href="#">81008</a>	741	232	2	76	integrin alpha 7 (Itga7) alternative variant bSep08, mRNA.
<a href="#">Itga9</a>	<a href="#">Itga9.aSep08</a>	<a href="#">685004</a>	54056	2230		250	integrin alpha 9 (Itga9) mRNA.
<a href="#">Itga11</a>	<a href="#">Itga11.bSep08</a>	<a href="#">315744</a>	13211	994	9	290	integrin, alpha 11 (Itga11) alternative variant bSep08, mRNA.
<a href="#">Itga11</a>	<a href="#">Itga11.cSep08</a>	<a href="#">315744</a>	14446	1305	8	278	integrin, alpha 11 (Itga11) alternative variant cSep08, mRNA.
<a href="#">Itgal</a>	<a href="#">Itgal.bSep08</a>	<a href="#">308995</a>	9306	2190	2	293	integrin alpha L (32.6 kD) (Itgal) alternative variant bSep08, mRNA.
<a href="#">Itgam</a>	<a href="#">Itgam.aSep08</a>	<a href="#">25021</a>	10927	287		95	integrin alpha M (Itgam) mRNA.
<a href="#">Itgav</a>	<a href="#">Itgav.bSep08</a>	<a href="#">296456</a>	9117	1452	6	199	integrin alpha V (Itgav) alternative variant bSep08, mRNA.
<a href="#">Itgax</a>	<a href="#">Itgax.aSep08</a>	<a href="#">499271</a>	4094	648		215	integrin alpha X (Itgax) mRNA.
<a href="#">Itgb1bp1</a>	<a href="#">Itgb1bp1.bSep08</a>	<a href="#">298914</a>	13441	800	6	177	integrin 1 protein (Itgb1bp1) alternative variant bSep08, mRNA.
<a href="#">Itgb1bp1</a>	<a href="#">Itgb1bp1.cSep08</a>	<a href="#">298914</a>	10216	708	6	159	integrin 1 protein (Itgb1bp1) alternative variant cSep08, mRNA.
<a href="#">Itgb1bp1</a>	<a href="#">Itgb1bp1.dSep08</a>	<a href="#">298914</a>	13048	558	5	127	integrin 1 protein (Itgb1bp1) alternative variant dSep08, mRNA.
<a href="#">Itgb1bp1</a>	<a href="#">Itgb1bp1.eSep08</a>	<a href="#">298914</a>	5407	1388	3	106	integrin 1 protein (11.2 kD) (Itgb1bp1) alternative variant eSep08, mRNA.
<a href="#">Itgb1bp1</a>	<a href="#">Itgb1bp1.gSep08</a>	<a href="#">298914</a>	2095	1631	2	60	integrin 1 protein (6.5 kD) (Itgb1bp1) alternative variant gSep08, mRNA.
<a href="#">Itgb1bp2</a>	<a href="#">Itgb1bp2.aSep08</a>	<a href="#">317258</a>	4624	1270		349	integrin beta 1 binding protein 2 (38.6 kD) (Itgb1bp2) mRNA.



<a href="#">Itgb3bp</a>	<a href="#">Itgb3bp.aSep08</a>	<a href="#">362548</a>	66032	1296	8	225	integrin beta 3 binding protein (beta3-endonexin) (Itgb3bp) alternative variant aSep08, mRNA.
<a href="#">Itgb3bp</a>	<a href="#">Itgb3bp.bSep08</a>	<a href="#">362548</a>	65611	851	7	222	integrin beta 3 binding protein (beta3-endonexin) (Itgb3bp) alternative variant bSep08, mRNA.
<a href="#">Itgb3bp</a>	<a href="#">Itgb3bp.cSep08</a>	<a href="#">362548</a>	65653	744	5	180	integrin beta 3 binding protein (beta3-endonexin) (Itgb3bp) alternative variant cSep08, mRNA.
<a href="#">Itgb3bp</a>	<a href="#">Itgb3bp.eSep08</a>	<a href="#">362548</a>	34046	620	3	163	integrin beta 3 binding protein (beta3-endonexin) (Itgb3bp) alternative variant eSep08, mRNA.
<a href="#">Itgb3bp</a>	<a href="#">Itgb3bp.fSep08</a>	<a href="#">362548</a>	24295	390	3	37	integrin beta 3 binding protein (beta3-endonexin) (Itgb3bp) alternative variant fSep08, mRNA.
<a href="#">Itgb4</a>	<a href="#">Itgb4.bSep08</a>	<a href="#">25724</a>	4145	1112	6	370	integrin beta 4 (Itgb4) alternative variant bSep08, mRNA.
<a href="#">Itgb4</a>	<a href="#">Itgb4.cSep08</a>	<a href="#">25724</a>	377	297	2	98	integrin beta 4 (Itgb4) alternative variant cSep08, mRNA.
<a href="#">Itgb4</a>	<a href="#">Itgb4.dSep08</a>	<a href="#">25724</a>	1200	481	2	98	integrin beta 4 (Itgb4) alternative variant dSep08, mRNA.
<a href="#">Itgb5</a>	<a href="#">Itgb5.bSep08</a>	<a href="#">257645</a>	74476	3089	12	618	integrin, beta 5 (Itgb5) alternative variant bSep08, mRNA.
<a href="#">Itgb5</a>	<a href="#">Itgb5.cSep08</a>	<a href="#">257645</a>	33130	600	3	144	integrin, beta 5 (Itgb5) alternative variant cSep08, mRNA.
<a href="#">Itgb5</a>	<a href="#">Itgb5.dSep08</a>	<a href="#">257645</a>	10308	389	4	129	integrin, beta 5 (Itgb5) alternative variant dSep08, mRNA.
<a href="#">Itgb5</a>	<a href="#">Itgb5.eSep08</a>	<a href="#">257645</a>	9735	921	2	95	integrin, beta 5 (Itgb5) alternative variant eSep08, mRNA.
<a href="#">Itgb7</a>	<a href="#">Itgb7.aSep08</a>	<a href="#">25713</a>	11644	1899		633	integrin, beta 7 (Itgb7) mRNA.
<a href="#">Itgb8</a>	<a href="#">Itgb8.bSep08</a>	<a href="#">362800</a>	38931	605	3	118	integrin beta 8 (Itgb8) alternative variant bSep08, mRNA.
<a href="#">Itgb8</a>	<a href="#">Itgb8.dSep08</a>	<a href="#">362800</a>	49054	396	3	70	integrin beta 8 (Itgb8) alternative variant dSep08, mRNA.
<a href="#">Itih1</a>	<a href="#">Itih1.aSep08</a>	<a href="#">306251</a>	14348	3078	19	904	inter-alpha trypsin inhibitor, heavy chain 1 (100.6 kD) (Itih1) alternative variant aSep08, complete mRNA.
<a href="#">Itih1</a>	<a href="#">Itih1.bSep08</a>	<a href="#">306251</a>	2443	812	1	170	inter-alpha trypsin inhibitor, heavy chain 1 (Itih1) alternative variant bSep08, mRNA.
<a href="#">Itih3</a>	<a href="#">Itih3.aSep08</a>	<a href="#">50693</a>	8660	2046	7	607	inter-alpha trypsin inhibitor, heavy chain 3 (Itih3) alternative variant aSep08, mRNA.
<a href="#">Itih3</a>	<a href="#">Itih3.bSep08</a>	<a href="#">50693</a>	1433	383	1	88	inter-alpha trypsin inhibitor, heavy chain 3 (Itih3) alternative variant bSep08, mRNA.
<a href="#">ITI_HC_C.0</a>	<a href="#">ITI_HC_C.0.aSep08</a>		7498	1777	3	558	inter-alpha inhibitor (ITI_HC_C.0) alternative variant aSep08, mRNA.
<a href="#">ITI_HC_C.0</a>	<a href="#">ITI_HC_C.0.bSep08</a>		7447	1239	1	293	inter-alpha-trypsin inhibitor heavy chain H3 (32.2 kD) (ITI_HC_C.0) alternative variant bSep08, mRNA.
<a href="#">Itpa</a>	<a href="#">Itpa.bSep08</a>	<a href="#">311422</a>	11741	1240	7	130	inosine triphosphatase (nucleoside triphosphate pyrophosphatase) (14.5 kD) (Itpa) alternative variant bSep08, complete mRNA.
<a href="#">Itpk1</a>	<a href="#">Itpk1.aSep08</a>	<a href="#">500709</a>	121799	740	2	246	inositol 1,3,4-triphosphate 5/6 kinase (Itpk1) alternative variant aSep08, mRNA.
<a href="#">Itpk1</a>	<a href="#">Itpk1.bSep08</a>	<a href="#">500709</a>	42576	476	1	100	inositol 1,3,4-triphosphate 5/6 kinase (Itpk1) alternative variant bSep08, mRNA.
<a href="#">Itpk1</a>	<a href="#">Itpk1.cSep08</a>	<a href="#">500709</a>	25562	364	1	100	inositol 1,3,4-triphosphate 5/6 kinase (Itpk1) alternative variant cSep08, mRNA.
<a href="#">Itpka</a>	<a href="#">Itpka.bSep08</a>	<a href="#">81677</a>	717	501	1	97	inositol 1,4,5-trisphosphate 3-kinase A (Itpka) alternative variant bSep08, mRNA.
<a href="#">Itpkb</a>	<a href="#">Itpkb.bSep08</a>	<a href="#">54260</a>	4013	620	4	206	inositol 1,4,5-trisphosphate 3-kinase B (Itpkb) alternative variant bSep08, mRNA.

<a href="#">Itpkb</a>	<a href="#">Itpkb.cSep08</a>	<a href="#">54260</a>	4608	361	2	120	inositol 1,4,5-trisphosphate 3-kinase B (Itpkb) alternative variant cSep08, mRNA.
<a href="#">Itpr1</a>	<a href="#">Itpr1.aSep08</a>	<a href="#">25262</a>	115496	395		62	inositol 1,4,5-triphosphate receptor 1 (Itpr1) mRNA.
<a href="#">Itpr2</a>	<a href="#">Itpr2.aSep08</a>	<a href="#">81678</a>	64478	1788	5	290	inositol 1,4,5-triphosphate receptor 2 (32.9 kD) (Itpr2) alternative variant aSep08, mRNA.
<a href="#">Itpr2</a>	<a href="#">Itpr2.bSep08</a>	<a href="#">81678</a>	26549	738	4	246	inositol 1,4,5-triphosphate receptor 2 (Itpr2) alternative variant bSep08, mRNA.
<a href="#">Itsn2</a>	<a href="#">Itsn2.aSep08</a>	<a href="#">313934</a>	18864	2351	10	495	intersectin 2 (57.3 kD) (Itsn2) alternative variant aSep08, mRNA.
<a href="#">Itsn2</a>	<a href="#">Itsn2.bSep08</a>	<a href="#">313934</a>	1849	997	1	126	intersectin 2 (14.7 kD) (Itsn2) alternative variant bSep08, mRNA.
<a href="#">Ivd</a>	<a href="#">Ivd.bSep08</a>	<a href="#">24513</a>	5948	761	6	253	isovaleryl coenzyme A dehydrogenase (Ivd) alternative variant bSep08, mRNA.
<a href="#">Ivd</a>	<a href="#">Ivd.cSep08</a>	<a href="#">24513</a>	8653	2119	7	193	isovaleryl coenzyme A dehydrogenase (21.2 kD) (Ivd) alternative variant cSep08, mRNA.
<a href="#">Ivns1abp</a>	<a href="#">Ivns1abp.bSep08</a>	<a href="#">289089</a>	8129	824	6	274	influenza virus NS1A binding protein (Ivns1abp) alternative variant bSep08, mRNA.
<a href="#">Ivns1abp</a>	<a href="#">Ivns1abp.cSep08</a>	<a href="#">289089</a>	12467	490	4	135	influenza virus NS1A binding protein (15.2 kD) (Ivns1abp) alternative variant cSep08, mRNA.
<a href="#">Ivns1abp</a>	<a href="#">Ivns1abp.dSep08</a>	<a href="#">289089</a>	1048	843	2	131	influenza virus NS1A binding protein (Ivns1abp) alternative variant dSep08, mRNA.
<a href="#">Iws1</a>	<a href="#">Iws1.bSep08</a>	<a href="#">291705</a>	3397	733	2	244	IWS1 homolog (S. cerevisiae) (Iws1) alternative variant bSep08, mRNA.
<a href="#">Iws1</a>	<a href="#">Iws1.cSep08</a>	<a href="#">291705</a>	14382	799	6	188	IWS1 homolog (S. cerevisiae) (Iws1) alternative variant cSep08, mRNA.
<a href="#">Iws1</a>	<a href="#">Iws1.dSep08</a>	<a href="#">291705</a>	9331	261	2	75	IWS1 homolog (S. cerevisiae) (Iws1) alternative variant dSep08, mRNA.
<a href="#">Iws1</a>	<a href="#">Iws1.eSep08</a>	<a href="#">291705</a>	3082	482	2	53	IWS1 homolog (S. cerevisiae) (Iws1) alternative variant eSep08, mRNA.
<a href="#">I_LWEQ.0</a>	<a href="#">I_LWEQ.0.aSep08</a>		11379	2749	1	184	huntingtin interacting protein 1 (I_LWEQ.0) alternative variant aSep08, mRNA.
<a href="#">I_LWEQ.0</a>	<a href="#">I_LWEQ.0.bSep08</a>		5819	558	2	82	huntingtin interacting protein 1 (I_LWEQ.0) alternative variant bSep08, mRNA.
<a href="#">jaby</a>	<a href="#">jaby.aSep08</a>		2459	256		29	putative protein (jaby) mRNA.
<a href="#">jachy</a>	<a href="#">jachy.aSep08</a>		6846	417		35	putative protein (jachy) mRNA.
<a href="#">jadar</a>	<a href="#">jadar.aSep08</a>		17430	586	3	43	putative protein (jadar) alternative variant aSep08, mRNA.
<a href="#">jadar</a>	<a href="#">jadar.bSep08</a>		17882	512	3	54	CRA a like (6.4 kD) (jadar) alternative variant bSep08, mRNA.
<a href="#">jadar</a>	<a href="#">jadar.cSep08</a>		1547	479	3	54	CRA a like (6.4 kD) (jadar) alternative variant cSep08, mRNA.
<a href="#">jafer</a>	<a href="#">jafer.aSep08</a>		4554	389		72	putative protein of ancient origin (jafer) mRNA.
<a href="#">jaflo</a>	<a href="#">jaflo.aSep08</a>		3190	199		63	gag protein like (jaflo) mRNA.
<a href="#">jaflu</a>	<a href="#">jaflu.aSep08</a>		907	367	2	122	putative protein of mammalian origin (jaflu) alternative variant aSep08, mRNA.
<a href="#">jaflu</a>	<a href="#">jaflu.bSep08</a>		786	304	2	90	putative protein (jaflu) alternative variant bSep08, mRNA.
<a href="#">Jag2</a>	<a href="#">Jag2.aSep08</a>	<a href="#">29147</a>	6336	2819	9	561	jagged 2 (Jag2) alternative variant aSep08, mRNA.

<a href="#">Jag2</a>	<a href="#">Jag2.bSep08</a>	<a href="#">29147</a>	713	525	1	163	jagged 2 (Jag2) alternative variant bSep08, mRNA.
<a href="#">Jak1</a>	<a href="#">Jak1.cSep08</a>	<a href="#">84598</a>	1828	633	3	88	janus kinase 1 (Jak1) alternative variant cSep08, mRNA.
<a href="#">Jak3</a>	<a href="#">Jak3.bSep08</a>	<a href="#">25326</a>	1345	764	2	74	janus kinase 3 (Jak3) alternative variant bSep08, mRNA.
<a href="#">jakee</a>	<a href="#">jakee.aSep08</a>		19028	1444	1	481	putative protein, with 2 coiled coil domains, of vertebrate origin (jakee) alternative variant aSep08, mRNA.
<a href="#">jakee</a>	<a href="#">jakee.bSep08</a>		12855	748		248	putative protein, with 2 coiled coil domains, of vertebrate origin (jakee) alternative variant bSep08, mRNA.
<a href="#">Jakmip1</a>	<a href="#">Jakmip1.bSep08</a>	<a href="#">305434</a>	65262	436	1	69	janus kinase and microtubule interacting protein 1 (Jakmip1) alternative variant bSep08, mRNA.
<a href="#">Jakmip2</a>	<a href="#">Jakmip2.bSep08</a>	<a href="#">307479</a>	10372	666	2	37	janus kinase and microtubule interacting protein 2 (4.2 kD) (Jakmip2) alternative variant bSep08, mRNA.
<a href="#">jaloy</a>	<a href="#">jaloy.aSep08</a>		2038	686		123	putative nuclear protein (13.6 kD) (jaloy) mRNA.
<a href="#">jamee</a>	<a href="#">jamee.aSep08</a>		2295	278		62	putative protein (jamee) mRNA.
<a href="#">jamer</a>	<a href="#">jamer.aSep08</a>		1062	462		52	putative protein (jamer) mRNA.
<a href="#">janoy</a>	<a href="#">janoy.aSep08</a>		999	556		56	putative protein (janoy) mRNA.
<a href="#">japor</a>	<a href="#">japor.aSep08</a>		12529	724		38	putative protein (4.1 kD) (japor) mRNA.
<a href="#">jarby</a>	<a href="#">jarby.aSep08</a>		6625	1742		527	putative protein (jarby) mRNA.
<a href="#">jarchy</a>	<a href="#">jarchy.aSep08</a>		1485	812		31	putative protein (3.6 kD) (jarchy) mRNA.
<a href="#">jardar</a>	<a href="#">jardar.aSep08</a>		7407	420		139	cyclic nucleotide-gated channel beta like (jardar) mRNA.
<a href="#">jarfer</a>	<a href="#">jarfer.aSep08</a>		19149	465		154	gli3 (jarfer) mRNA.
<a href="#">jarflo</a>	<a href="#">jarflo.aSep08</a>		2389	302		28	putative protein (jarflo) mRNA.
<a href="#">jarflu</a>	<a href="#">jarflu.aSep08</a>		1844	830	1	187	transient receptor potential cation channel subfamily M member 4 (jarflu) alternative variant aSep08, mRNA.
<a href="#">jarflu</a>	<a href="#">jarflu.bSep08</a>		2000	708	2	128	transient receptor potential cation channel subfamily M member 4 (jarflu) alternative variant bSep08, mRNA.
<a href="#">Jarid1b</a>	<a href="#">Jarid1b.bSep08</a>	<a href="#">304809</a>	7255	678	4	216	jumonji, AT rich interactive domain 1B (Rbp2 like) (Jarid1b) alternative variant bSep08, mRNA.
<a href="#">Jarid1b</a>	<a href="#">Jarid1b.cSep08</a>	<a href="#">304809</a>	3180	754	3	205	jumonji, AT rich interactive domain 1B (Rbp2 like) (Jarid1b) alternative variant cSep08, mRNA.
<a href="#">jarkee</a>	<a href="#">jarkee.aSep08</a>		1778	590		57	putative protein (6.3 kD) (jarkee) mRNA.
<a href="#">jarloy</a>	<a href="#">jarloy.aSep08</a>		4809	781		123	putative protein (jarloy) mRNA.
<a href="#">jarmee</a>	<a href="#">jarmee.aSep08</a>		10871	403		134	putative protein of metazoan origin (jarmee) mRNA.
<a href="#">jarmer</a>	<a href="#">jarmer.aSep08</a>		679	403		75	putative nuclear protein (9.0 kD) (jarmer) mRNA.
<a href="#">jarnoy</a>	<a href="#">jarnoy.aSep08</a>		3824	357		114	ankyrin armadillo repeat containing (jarnoy) mRNA.
<a href="#">jarpor</a>	<a href="#">jarpor.aSep08</a>		948	327		108	dmx-like 2 (jarpor) mRNA.
<a href="#">jarsa</a>	<a href="#">jarsa.aSep08</a>		6442	1028		125	gag-pol protein like (jarsa) mRNA.
<a href="#">jarshee</a>	<a href="#">jarshee.aSep08</a>		4984	440		76	putative nuclear protein (8.5 kD) (jarshee) mRNA.
<a href="#">jarvar</a>	<a href="#">jarvar.aSep08</a>		2880	227		75	WD repeat domain 65 like (jarvar) mRNA.
<a href="#">jarwey</a>	<a href="#">jarwey.aSep08</a>		1090	838		169	putative protein (jarwey) mRNA.
<a href="#">jasa</a>	<a href="#">jasa.aSep08</a>		1378	432		38	putative protein (4.2 kD) (jasa) mRNA.
<a href="#">jashee</a>	<a href="#">jashee.aSep08</a>		709	488		89	putative protein (jashee) mRNA.
<a href="#">jato</a>	<a href="#">jato.aSep08</a>		526	416		119	immunoglobulin heavy chain (jato) mRNA.
<a href="#">javar</a>	<a href="#">javar.aSep08</a>		4623	746		248	putative protein of bilateral origin (javar) mRNA.

<a href="#">jawby</a>	<a href="#">jawby.aSep08</a>		4070	674		29	putative protein (jawby) mRNA.
<a href="#">jawchy</a>	<a href="#">jawchy.aSep08</a>		5257	751		85	putative protein of mammalian origin (jawchy) mRNA.
<a href="#">jawdar</a>	<a href="#">jawdar.aSep08</a>		9710	365	3	95	putative protein (jawdar) alternative variant aSep08, mRNA.
<a href="#">jawey</a>	<a href="#">jawey.aSep08</a>		1547	498		70	putative protein (jawey) mRNA.
<a href="#">jawfer</a>	<a href="#">jawfer.aSep08</a>		541	440		83	putative mitochondrial protein (9.3 kD) (jawfer) mRNA.
<a href="#">jawflo</a>	<a href="#">jawflo.aSep08</a>		1844	1048	2	57	putative protein (jawflo) alternative variant aSep08, mRNA.
<a href="#">jawflu</a>	<a href="#">jawflu.aSep08</a>		2979	344		114	transient receptor potential cation channel subfamily M member 4 (jawflu) mRNA.
<a href="#">jawkee</a>	<a href="#">jawkee.aSep08</a>		1451	652		89	putative protein (9.6 kD) (jawkee) mRNA.
<a href="#">jawloy</a>	<a href="#">jawloy.aSep08</a>		4581	602		36	putative protein (4.4 kD) (jawloy) mRNA.
<a href="#">jawmee</a>	<a href="#">jawmee.aSep08</a>		2884	513	2	76	putative protein (jawmee) alternative variant aSep08, mRNA.
<a href="#">jawmer</a>	<a href="#">jawmer.aSep08</a>		9989	670		54	putative protein (jawmer) mRNA.
<a href="#">jawnoy</a>	<a href="#">jawnoy.aSep08</a>		3437	885		27	putative protein (3.3 kD) (jawnoy) alternative variant aSep08, mRNA.
<a href="#">jawnoy</a>	<a href="#">jawnoy.bSep08</a>		3168	677	1	20	putative protein (2.2 kD) (jawnoy) alternative variant bSep08, mRNA.
<a href="#">jawpor</a>	<a href="#">jawpor.aSep08</a>		5003	457		59	putative protein (jawpor) mRNA.
<a href="#">jawsa</a>	<a href="#">jawsa.aSep08</a>		2235	348		39	putative protein (4.5 kD) (jawsa) mRNA.
<a href="#">jawshee</a>	<a href="#">jawshee.aSep08</a>		1271	395	2	83	ski-like (jawshee) alternative variant aSep08, mRNA.
<a href="#">jawvar</a>	<a href="#">jawvar.aSep08</a>		5657	568		26	putative protein (3.1 kD) (jawvar) mRNA.
<a href="#">jawwey</a>	<a href="#">jawwey.aSep08</a>		8127	712		16	putative protein (jawwey) mRNA.
<a href="#">Jazf1</a>	<a href="#">Jazf1.aSep08</a>	<a href="#">685879</a>	200215	771	5	251	JAZF zinc finger 1 (Jazf1) alternative variant aSep08, mRNA.
<a href="#">Jazf1</a>	<a href="#">Jazf1.bSep08</a>	<a href="#">685879</a>	146476	448	4	102	JAZF zinc finger 1 (Jazf1) alternative variant bSep08, mRNA.
<a href="#">jeeby</a>	<a href="#">jeeby.aSep08</a>		7717	2097		85	putative nuclear protein (9.6 kD) (jeeby) mRNA.
<a href="#">jeechy</a>	<a href="#">jeechy.aSep08</a>		24631	873	3	46	putative protein (jeechy) alternative variant aSep08, mRNA.
<a href="#">jeechy</a>	<a href="#">jeechy.bSep08</a>		22840	396	1	43	putative protein (jeechy) alternative variant bSep08, mRNA.
<a href="#">jeedar</a>	<a href="#">jeedar.aSep08</a>		627	556	2	61	putative protein (jeedar) alternative variant aSep08, mRNA.
<a href="#">jeefer</a>	<a href="#">jeefer.aSep08</a>		1395	786		125	putative protein of mammalian origin (jeefer) mRNA.
<a href="#">jeeflo</a>	<a href="#">jeeflo.aSep08</a>		25773	401		45	putative protein (jeeflo) mRNA.
<a href="#">jeeflu</a>	<a href="#">jeeflu.aSep08</a>		6536	646		214	transient receptor potential cation channel subfamily M member 4 (jeeflu) mRNA.
<a href="#">jeekee</a>	<a href="#">jeekee.aSep08</a>		3733	387		82	protein tyrosine phosphatase receptor type f polypeptide interacting alpha 4 (jeekee) mRNA.
<a href="#">jeeloy</a>	<a href="#">jeeloy.aSep08</a>		35868	484		61	putative protein (6.0 kD) (jeeloy) mRNA.
<a href="#">jeemee</a>	<a href="#">jeemee.aSep08</a>		8401	427		142	cyclase 9 (jeemee) mRNA.
<a href="#">jeemer</a>	<a href="#">jeemer.aSep08</a>		1263	1006	1	191	putative protein (jeemer) alternative variant aSep08, mRNA.

<a href="#">jeemer</a>	<a href="#">jeemer.bSep08</a>		1451	720	2	142	putative protein (jeemer) alternative variant bSep08, mRNA.
<a href="#">jeenoy</a>	<a href="#">jeenoy.aSep08</a>		543	456		35	putative protein (jeenoy) alternative variant aSep08, mRNA.
<a href="#">jeepor</a>	<a href="#">jeepor.aSep08</a>		3929	220		73	putative protein (jeepor) mRNA.
<a href="#">jeesa</a>	<a href="#">jeesa.aSep08</a>		22297	655		66	putative protein (7.3 kD) (jeesa) mRNA.
<a href="#">jeeshee</a>	<a href="#">jeeshee.aSep08</a>		4490	434		62	putative protein (jeeshee) mRNA.
<a href="#">jeevar</a>	<a href="#">jeevar.aSep08</a>		3428	763	1	41	putative protein (jeevar) alternative variant aSep08, mRNA.
<a href="#">jeevar</a>	<a href="#">jeevar.bSep08</a>		4518	316	2	36	peptidyl prolyl isomerase H (jeevar) alternative variant bSep08, mRNA.
<a href="#">jeevar</a>	<a href="#">jeevar.cSep08</a>		9679	303	2	36	peptidyl prolyl isomerase H (jeevar) alternative variant cSep08, mRNA.
<a href="#">jeewey</a>	<a href="#">jeewey.aSep08</a>		1162	342	1	67	putative protein (jeewey) alternative variant aSep08, mRNA.
<a href="#">jeewey</a>	<a href="#">jeewey.bSep08</a>		34545	1324	7	70	putative protein (8.0 kD) (jeewey) alternative variant bSep08, mRNA.
<a href="#">jerby</a>	<a href="#">jerby.aSep08</a>	<a href="#">367754</a>	4398	499		165	putative protein of vertebrate origin (jerby) mRNA.
<a href="#">jerchy</a>	<a href="#">jerchy.aSep08</a>		16446	1368		94	guanine nucleotide exchange factor (jerchy) mRNA.
<a href="#">jerdar</a>	<a href="#">jerdar.aSep08</a>		10212	656		74	putative protein (jerdar) mRNA.
<a href="#">jerfer</a>	<a href="#">jerfer.aSep08</a>		18139	407	2	96	2 beta 1 3 (jerfer) alternative variant aSep08, mRNA.
<a href="#">jerfer</a>	<a href="#">jerfer.bSep08</a>		12215	730	1	88	2 beta 1 3 precursor (10.1 kD) (jerfer) alternative variant bSep08, mRNA.
<a href="#">jerflo</a>	<a href="#">jerflo.aSep08</a>		2969	1595		61	pantothenate kinase (jerflo) mRNA.
<a href="#">jerflu</a>	<a href="#">jerflu.aSep08</a>		2094	353		117	alpha 3 (jerflu) mRNA.
<a href="#">jerkee</a>	<a href="#">jerkee.aSep08</a>		533	325		68	protein tyrosine phosphatase receptor type f polypeptide interacting alpha (jerkee) mRNA.
<a href="#">jerloy</a>	<a href="#">jerloy.aSep08</a>		1041	588		63	putative protein (jerloy) mRNA.
<a href="#">jermee</a>	<a href="#">jermee.aSep08</a>		1568	872		290	putative protein of vertebrate origin (jermee) mRNA.
<a href="#">jermer</a>	<a href="#">jermer.aSep08</a>		33940	676		38	putative protein (4.3 kD) (jermer) mRNA.
<a href="#">jernoy</a>	<a href="#">jernoy.aSep08</a>		410	273		41	putative protein (jernoy) mRNA.
<a href="#">jerpor</a>	<a href="#">jerpor.aSep08</a>		146771	701		233	zinc finger protein 291 (jerpor) mRNA.
<a href="#">jersa</a>	<a href="#">jersa.aSep08</a>		57603	698		44	putative protein (5.1 kD) (jersa) mRNA.
<a href="#">jershee</a>	<a href="#">jershee.aSep08</a>		9173	619	2	38	putative protein (jershee) alternative variant aSep08, mRNA.
<a href="#">jershee</a>	<a href="#">jershee.bSep08</a>		8597	585	1	14	putative protein (jershee) alternative variant bSep08, mRNA.
<a href="#">jervar</a>	<a href="#">jervar.aSep08</a>		2462	362		120	CRA b (jervar) mRNA.
<a href="#">jerwey</a>	<a href="#">jerwey.aSep08</a>		5554	682		43	putative protein (jerwey) mRNA.
<a href="#">jeyby</a>	<a href="#">jeyby.aSep08</a>		6129	458		124	interleukin 13 receptor alpha 2 (jeyby) mRNA.
<a href="#">jeychy</a>	<a href="#">jeychy.aSep08</a>		548	379		107	putative protein (jeychy) mRNA.
<a href="#">jeydar</a>	<a href="#">jeydar.aSep08</a>		2125	716	2	62	putative protein (6.6 kD) (jeydar) alternative variant aSep08, mRNA.
<a href="#">jeyfer</a>	<a href="#">jeyfer.aSep08</a>		29364	459		75	putative protein (jeyfer) mRNA.
<a href="#">jeyflo</a>	<a href="#">jeyflo.aSep08</a>		1664	724		66	putative protein (7.3 kD) (jeyflo) mRNA.

<a href="#">jeyflu</a>	<a href="#">jeyflu.aSep08</a>		1902	746		248	putative protein of metazoan origin (jeyflu) mRNA.
<a href="#">jeykee</a>	<a href="#">jeykee.aSep08</a>		1323	623		47	putative protein of mammalian origin (jeykee) mRNA.
<a href="#">jeyloy</a>	<a href="#">jeyloy.aSep08</a>		1990	525	3	108	putative protein (jeyloy) alternative variant aSep08, mRNA.
<a href="#">jeymee</a>	<a href="#">jeymee.aSep08</a>		2142	313		84	putative protein (jeymee) mRNA.
<a href="#">jeymer</a>	<a href="#">jeymer.aSep08</a>		2286	713		237	peptidase M2, peptidyl-dipeptidase A (jeymer) mRNA.
<a href="#">jeynoy</a>	<a href="#">jeynoy.aSep08</a>		1762	390		90	myosin Ib CRA b (jeynoy) mRNA.
<a href="#">jeypor</a>	<a href="#">jeypor.aSep08</a>		12969	470		53	putative protein (5.9 kD) (jeypor) mRNA.
<a href="#">jeysa</a>	<a href="#">jeysa.aSep08</a>		1162	398		132	putative protein of mammalian origin (jeysa) mRNA.
<a href="#">jeyshee</a>	<a href="#">jeyshee.aSep08</a>		9530	576	2	176	putative protein (jeyshee) alternative variant aSep08, mRNA.
<a href="#">jeyshee</a>	<a href="#">jeyshee.cSep08</a>		1744	759	2	57	putative protein (6.3 kD) (jeyshee) alternative variant cSep08, mRNA.
<a href="#">jeyvar</a>	<a href="#">jeyvar.aSep08</a>		21171	593		197	family with sequence similarity 80 member (jeyvar) mRNA.
<a href="#">jeywey</a>	<a href="#">jeywey.bSep08</a>		2797	478		59	putative protein (6.5 kD) (jeywey) alternative variant bSep08, mRNA.
<a href="#">JmjC.0</a>	<a href="#">JmjC.0.aSep08</a>		15133	1759		586	CRA a (JmjC.0) alternative variant aSep08, mRNA.
<a href="#">JmjC.0</a>	<a href="#">JmjC.0.bSep08</a>		33744	1790		447	CRA b (JmjC.0) alternative variant bSep08, mRNA.
<a href="#">JmjC.0</a>	<a href="#">JmjC.0.cSep08</a>		10509	1230	7	394	CRA a (JmjC.0) alternative variant cSep08, mRNA.
<a href="#">JmjC.0</a>	<a href="#">JmjC.0.dSep08</a>		15528	2672	7	340	CRA b (38.7 kD) (JmjC.0) alternative variant dSep08, mRNA.
<a href="#">JmjC.1</a>	<a href="#">JmjC.1.aSep08</a>		95069	1130		261	transcription factor jumonji, JmjN (30.3 kD) (JmjC.1) mRNA.
<a href="#">Jmjd1a</a>	<a href="#">Jmjd1a.bSep08</a>	<a href="#">312440</a>	23076	3470	17	955	transcription factor jumonji (Jmjd1a) alternative variant bSep08, mRNA.
<a href="#">Jmjd1a</a>	<a href="#">Jmjd1a.cSep08</a>	<a href="#">312440</a>	11102	711	5	183	putative protein (Jmjd1a) alternative variant cSep08, mRNA.
<a href="#">Jmjd1c</a>	<a href="#">Jmjd1c.aSep08</a>	<a href="#">171120</a>	120674	807	4	268	CRA b (Jmjd1c) alternative variant aSep08, mRNA.
<a href="#">Jmjd1c</a>	<a href="#">Jmjd1c.bSep08</a>	<a href="#">171120</a>	3412	349	2	83	putative protein (Jmjd1c) alternative variant bSep08, mRNA.
<a href="#">Jmjd2a</a>	<a href="#">Jmjd2a.bSep08</a>	<a href="#">313539</a>	7942	1057	5	229	putative protein of eukaryotic origin (Jmjd2a) alternative variant bSep08, mRNA.
<a href="#">Jmjd2a</a>	<a href="#">Jmjd2a.cSep08</a>	<a href="#">313539</a>	11148	521	5	173	transcription factor jumonji, JmjN (Jmjd2a) alternative variant cSep08, mRNA.
<a href="#">Jmjd2a</a>	<a href="#">Jmjd2a.dSep08</a>	<a href="#">313539</a>	583	246	2	62	putative protein of metazoan origin (Jmjd2a) alternative variant dSep08, mRNA.
<a href="#">Jmjd2a</a>	<a href="#">Jmjd2a.eSep08</a>	<a href="#">313539</a>	1627	339	2	41	putative protein (4.4 kD) (Jmjd2a) alternative variant eSep08, mRNA.
<a href="#">Jmjd2c</a>	<a href="#">Jmjd2c.aSep08</a>	<a href="#">298144</a>	12634	950	5	310	putative protein (Jmjd2c) alternative variant aSep08, mRNA.
<a href="#">Jmjd2c</a>	<a href="#">Jmjd2c.bSep08</a>	<a href="#">298144</a>	52122	1756	7	293	putative protein of eukaryotic origin (Jmjd2c) alternative variant bSep08, mRNA.
<a href="#">Jmjd2c</a>	<a href="#">Jmjd2c.cSep08</a>	<a href="#">298144</a>	46795	1111	8	250	putative protein (Jmjd2c) alternative variant cSep08, mRNA.
<a href="#">Jmjd2c</a>	<a href="#">Jmjd2c.dSep08</a>	<a href="#">298144</a>	20287	504	4	105	putative protein of eukaryotic origin (Jmjd2c) alternative variant dSep08, mRNA.

<a href="#">Jmjd3</a>	<a href="#">Jmjd3.bSep08</a>	<a href="#">363630</a>	1356	650	5	155	putative protein of metazoan origin (Jmjd3) alternative variant bSep08, mRNA.
<a href="#">Jmjd3</a>	<a href="#">Jmjd3.cSep08</a>	<a href="#">363630</a>	742	636	2	116	putative protein of metazoan origin (Jmjd3) alternative variant cSep08, mRNA.
<a href="#">Jmjd5</a>	<a href="#">Jmjd5.bSep08</a>	<a href="#">308976</a>	14905	1059	5	291	putative protein of ancient origin (33.1 kD) (Jmjd5) alternative variant bSep08, complete mRNA.
<a href="#">Jmjd5</a>	<a href="#">Jmjd5.cSep08</a>	<a href="#">308976</a>	3979	737	3	130	putative cytoplasmic protein of ancient origin (14.9 kD) (Jmjd5) alternative variant cSep08, mRNA.
<a href="#">Jmjd5</a>	<a href="#">Jmjd5.eSep08</a>	<a href="#">308976</a>	3291	664	3	43	putative protein of eukaryotic origin (Jmjd5) alternative variant eSep08, mRNA.
<a href="#">Jmjd6</a>	<a href="#">Jmjd6.bSep08</a>	<a href="#">360665</a>	5731	1684	6	454	phosphatidylserine receptor CRA b (Jmjd6) alternative variant bSep08, mRNA.
<a href="#">Jmjd6</a>	<a href="#">Jmjd6.cSep08</a>	<a href="#">360665</a>	3013	611	3	135	putative protein of eukaryotic origin (Jmjd6) alternative variant cSep08, mRNA.
<a href="#">Jmjd6</a>	<a href="#">Jmjd6.eSep08</a>	<a href="#">360665</a>	2536	693	3	98	putative nuclear protein of vertebrate origin (10.4 kD) (Jmjd6) alternative variant eSep08, mRNA.
<a href="#">Jmjd6</a>	<a href="#">Jmjd6.fSep08</a>	<a href="#">360665</a>	1807	670	2	88	putative protein of vertebrate origin (9.2 kD) (Jmjd6) alternative variant fSep08, mRNA.
<a href="#">Jmjd6</a>	<a href="#">Jmjd6.gSep08</a>	<a href="#">360665</a>	2939	631	4	60	phosphatidylserine receptor transcript like (Jmjd6) alternative variant gSep08, mRNA.
<a href="#">Jnk-SapK_ap_N.0</a>	<a href="#">Jnk-SapK_ap_N.0.aSep08</a>		23739	1194	7	365	mitogen-activated protein kinase 8 interacting 3 (Jnk-SapK_ap_N.0) alternative variant aSep08, mRNA.
<a href="#">joby</a>	<a href="#">joby.aSep08</a>		5015	543		180	shroom family member 4 (joby) mRNA.
<a href="#">jochy</a>	<a href="#">jochy.aSep08</a>		10026	536	3	68	putative protein (jochy) mRNA.
<a href="#">jodar</a>	<a href="#">jodar.aSep08</a>		2263	532	2	126	putative protein (jodar) alternative variant aSep08, mRNA.
<a href="#">jofer</a>	<a href="#">jofer.aSep08</a>		34414	937		312	cell division cycle 2-like 5 (jofer) mRNA.
<a href="#">joflo</a>	<a href="#">joflo.aSep08</a>		1174	970		15	putative protein (1.8 kD) (joflo) mRNA.
<a href="#">joflu</a>	<a href="#">joflu.aSep08</a>		1277	712		67	putative protein (joflu) mRNA.
<a href="#">jokee</a>	<a href="#">jokee.aSep08</a>		1876	785		56	putative protein (6.2 kD) (jokee) mRNA.
<a href="#">joloy</a>	<a href="#">joloy.aSep08</a>		1027	682		46	putative protein (joloy) mRNA.
<a href="#">jomee</a>	<a href="#">jomee.aSep08</a>		3742	743		81	putative protein (jomee) mRNA.
<a href="#">jomer</a>	<a href="#">jomer.aSep08</a>		1716	728		76	putative mitochondrial protein (8.9 kD) (jomer) mRNA.
<a href="#">jonoy</a>	<a href="#">jonoy.aSep08</a>		15753	1885		182	basic immunoglobulin-like variable motif containing CRA b (jonoy) mRNA.
<a href="#">jopor</a>	<a href="#">jopor.aSep08</a>		47192	462		154	dmx-like 2 (jopor) mRNA.
<a href="#">jorby</a>	<a href="#">jorby.aSep08</a>		744	416		76	angiominin (jorby) mRNA.
<a href="#">jorchy</a>	<a href="#">jorchy.aSep08</a>		9676	1651		327	putative protein of vertebrate origin (jorchy) mRNA.
<a href="#">jordar</a>	<a href="#">jordar.aSep08</a>		7030	361		27	putative protein (jordar) mRNA.
<a href="#">jorfer</a>	<a href="#">jorfer.aSep08</a>		24665	743		54	putative protein (6.2 kD) (jorfer) mRNA.
<a href="#">jorflo</a>	<a href="#">jorflo.aSep08</a>		8484	489		52	putative protein (5.7 kD) (jorflo) mRNA.
<a href="#">jorflu</a>	<a href="#">jorflu.aSep08</a>		7272	487		24	putative protein (jorflu) mRNA.
<a href="#">jorkee</a>	<a href="#">jorkee.aSep08</a>		4918	967	1	50	putative protein (jorkee) alternative variant aSep08, mRNA.
<a href="#">jorkee</a>	<a href="#">jorkee.bSep08</a>		4892	765		44	putative protein (jorkee) alternative variant bSep08, mRNA.
<a href="#">jorloy</a>	<a href="#">jorloy.aSep08</a>		1115	438	1	102	mucin 4 (jorloy) alternative variant aSep08, mRNA.

<a href="#">jorloy</a>	<a href="#">jorloy.bSep08</a>		2153	603	2	94	mucin 4 (jorloy) alternative variant bSep08, mRNA.
<a href="#">jormee</a>	<a href="#">jormee.aSep08</a>		2643	753		121	putative protein of mammalian origin (jormee) mRNA.
<a href="#">jormer</a>	<a href="#">jormer.aSep08</a>		7550	693	2	42	putative protein (4.9 kD) (jormer) alternative variant aSep08, mRNA.
<a href="#">jornoy</a>	<a href="#">jornoy.aSep08</a>		5107	1603		60	myosin lb (jornoy) mRNA.
<a href="#">jorpor</a>	<a href="#">jorpor.aSep08</a>		6016	483		66	putative protein (7.3 kD) (jorpor) mRNA.
<a href="#">jorsa</a>	<a href="#">jorsa.aSep08</a>		96217	764	3	52	putative protein (jorsa) alternative variant aSep08, mRNA.
<a href="#">jorsa</a>	<a href="#">jorsa.bSep08</a>		76999	578	2	56	putative protein (jorsa) alternative variant bSep08, mRNA.
<a href="#">jorsa</a>	<a href="#">jorsa.cSep08</a>		75019	463	1	44	putative protein (jorsa) alternative variant cSep08, mRNA.
<a href="#">jorsa</a>	<a href="#">jorsa.dSep08</a>		75496	384	1	32	putative protein (jorsa) alternative variant dSep08, mRNA.
<a href="#">jorshee</a>	<a href="#">jorshee.aSep08</a>		1413	354		46	putative protein (jorshee) mRNA.
<a href="#">jorvar</a>	<a href="#">jorvar.aSep08</a>		82274	302		25	putative protein (jorvar) mRNA.
<a href="#">jorwey</a>	<a href="#">jorwey.aSep08</a>		7531	807		27	putative protein (jorwey) mRNA.
<a href="#">josa</a>	<a href="#">josa.aSep08</a>		21495	741		63	putative protein (josa) mRNA.
<a href="#">Josd2</a>	<a href="#">Josd2.aSep08</a>	<a href="#">292876</a>	3431	1339	5	362	putative protein of eukaryotic origin (Josd2) alternative variant aSep08, mRNA.
<a href="#">Josd2</a>	<a href="#">Josd2.cSep08</a>	<a href="#">292876</a>	3792	928	5	188	machado-Joseph disease protein MJD (20.8 kD) (Josd2) alternative variant cSep08, mRNA.
<a href="#">Josd2</a>	<a href="#">Josd2.dSep08</a>	<a href="#">292876</a>	2899	443	3	98	putative protein of eukaryotic origin (Josd2) alternative variant dSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.bSep08</a>	<a href="#">363017</a>	6400	1447	7	285	putative nuclear protein of vertebrate origin (32.7 kD) (Josd3) alternative variant bSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.cSep08</a>	<a href="#">363017</a>	7642	1255	11	185	putative protein of vertebrate origin (Josd3) alternative variant cSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.dSep08</a>	<a href="#">363017</a>	6377	759	11	113	putative protein of vertebrate origin (Josd3) alternative variant dSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.eSep08</a>	<a href="#">363017</a>	1297	1093	2	49	putative protein (5.9 kD) (Josd3) alternative variant eSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.gSep08</a>	<a href="#">363017</a>	2921	688	7	42	putative protein (4.8 kD) (Josd3) alternative variant gSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.iSep08</a>	<a href="#">363017</a>	1515	518	5	35	putative protein (4.1 kD) (Josd3) alternative variant iSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.jSep08</a>	<a href="#">363017</a>	1277	484	4	8	putative protein (Josd3) alternative variant jSep08, mRNA.
<a href="#">Josd3</a>	<a href="#">Josd3.kSep08</a>	<a href="#">363017</a>	4560	352	7	38	putative protein (4.5 kD) (Josd3) alternative variant kSep08, mRNA.
<a href="#">joshee</a>	<a href="#">joshee.aSep08</a>		8194	267		89	putative protein (joshee) mRNA.
<a href="#">jovar</a>	<a href="#">jovar.aSep08</a>		1019	791		183	transmembrane protein 125 (jovar) mRNA.
<a href="#">jowey</a>	<a href="#">jowey.aSep08</a>		8780	436		94	putative mitochondrial protein (9.9 kD) (jowey) mRNA.
<a href="#">joyby</a>	<a href="#">joyby.aSep08</a>		54481	396		60	putative protein (6.5 kD) (joyby) mRNA.
<a href="#">joychy</a>	<a href="#">joychy.aSep08</a>		1847	267		72	putative protein (joychy) mRNA.
<a href="#">joydar</a>	<a href="#">joydar.aSep08</a>		9593	448		38	putative protein (joydar) mRNA.
<a href="#">joyfer</a>	<a href="#">joyfer.aSep08</a>		2937	198		66	putative protein (joyfer) mRNA.
<a href="#">joyflo</a>	<a href="#">joyflo.bSep08</a>	<a href="#">687730</a>	19577	995	1	122	putative protein, with a transmembrane domain (13.3 kD) (joyflo) alternative variant bSep08, mRNA.



<a href="#">joyflu</a>	<a href="#">joyflu.aSep08</a>		4039	772		119	CRA b like (12.4 kD) (joyflu) mRNA.
<a href="#">joykee</a>	<a href="#">joykee.aSep08</a>		16315	564	4	81	protein phosphatase 1 regulatory (joykee) alternative variant aSep08, mRNA.
<a href="#">joyloy</a>	<a href="#">joyloy.aSep08</a>		6497	749		149	putative mitochondrial protein (16.5 kD) (joyloy) mRNA.
<a href="#">joymee</a>	<a href="#">joymee.aSep08</a>		1302	736		127	putative protein (14.1 kD) (joymee) mRNA.
<a href="#">joymer</a>	<a href="#">joymer.aSep08</a>		2048	674		80	putative secreted or extracellular protein precursor (9.0 kD) (joymer) mRNA.
<a href="#">joynoy</a>	<a href="#">joynoy.aSep08</a>		42476	302		33	putative protein (joynoy) mRNA.
<a href="#">joypor</a>	<a href="#">joypor.aSep08</a>		11662	304		40	putative protein (joypor) mRNA.
<a href="#">joysa</a>	<a href="#">joysa.aSep08</a>		5129	1591	2	56	putative protein (6.2 kD) (joysa) alternative variant aSep08, mRNA.
<a href="#">joysa</a>	<a href="#">joysa.bSep08</a>		42742	921	4	81	putative protein (9.2 kD) (joysa) alternative variant bSep08, mRNA.
<a href="#">joysa</a>	<a href="#">joysa.cSep08</a>		4043	505	1	26	putative protein (joysa) alternative variant cSep08, mRNA.
<a href="#">joyshee</a>	<a href="#">joyshee.aSep08</a>		51709	527		102	putative protein (joyshee) mRNA.
<a href="#">joyvar</a>	<a href="#">joyvar.aSep08</a>		494	257		85	putative protein (joyvar) mRNA.
<a href="#">joywey</a>	<a href="#">joywey.aSep08</a>		9600	678		83	putative protein (joywey) mRNA.
<a href="#">Jph2</a>	<a href="#">Jph2.bSep08</a>	<a href="#">296345</a>	3133	1759	3	46	junctional protein 2 (Jph2) alternative variant bSep08, mRNA.
<a href="#">Jph2</a>	<a href="#">Jph2.cSep08</a>	<a href="#">296345</a>	67582	839	2	65	junctional protein 2 (7.1 kD) (Jph2) alternative variant cSep08, mRNA.
<a href="#">Jph4</a>	<a href="#">Jph4.bSep08</a>	<a href="#">445271</a>	3765	2690	3	393	junctional protein 4 (42.0 kD) (Jph4) alternative variant bSep08, mRNA.
<a href="#">Jph4</a>	<a href="#">Jph4.cSep08</a>	<a href="#">445271</a>	1451	729	2	242	junctional protein 4 (Jph4) alternative variant cSep08, mRNA.
<a href="#">Jph4</a>	<a href="#">Jph4.dSep08</a>	<a href="#">445271</a>	3576	2411	3	221	junctional protein 4 (23.0 kD) (Jph4) alternative variant dSep08, mRNA.
<a href="#">Jsrp1</a>	<a href="#">Jsrp1.bSep08</a>	<a href="#">690423</a>	1693	687	2	66	junctional sarcoplasmic reticulum protein 1 (Jsrp1) alternative variant bSep08, mRNA.
<a href="#">Jsrp1</a>	<a href="#">Jsrp1.cSep08</a>	<a href="#">690423</a>	1809	410	4	55	junctional sarcoplasmic reticulum protein 1 (5.9 kD) (Jsrp1) alternative variant cSep08, mRNA.
<a href="#">Jtb</a>	<a href="#">Jtb.bSep08</a>	<a href="#">29439</a>	5342	1041	5	32	jumping translocation breakpoint (3.5 kD) (Jtb) alternative variant bSep08, mRNA.
<a href="#">Jtb</a>	<a href="#">Jtb.dSep08</a>	<a href="#">29439</a>	5086	747	4	32	jumping translocation breakpoint (3.5 kD) (Jtb) alternative variant dSep08, mRNA.
<a href="#">Jtb</a>	<a href="#">Jtb.eSep08</a>	<a href="#">29439</a>	1678	736	3	56	jumping translocation breakpoint (Jtb) alternative variant eSep08, mRNA.
<a href="#">Jtv1</a>	<a href="#">Jtv1.bSep08</a>	<a href="#">288480</a>	9324	815	1	253	JTV1 gene (Jtv1) alternative variant bSep08, mRNA.
<a href="#">juby</a>	<a href="#">juby.aSep08</a>		30171	605		201	shroom family member 4 (juby) mRNA.
<a href="#">juchy</a>	<a href="#">juchy.aSep08</a>		556	294		27	putative protein (juchy) mRNA.
<a href="#">judar</a>	<a href="#">judar.aSep08</a>		12069	911		60	putative protein (6.7 kD) (judar) mRNA.
<a href="#">jufer</a>	<a href="#">jufer.aSep08</a>		596	409		32	putative protein (jufer) mRNA.
<a href="#">juflo</a>	<a href="#">juflo.aSep08</a>		1457	294	2	31	putative protein (juflo) alternative variant aSep08, mRNA.
<a href="#">juflu</a>	<a href="#">juflu.aSep08</a>		421	342		113	proline rich 12 (juflu) mRNA.
<a href="#">jukee</a>	<a href="#">jukee.aSep08</a>		10464	340		55	putative protein (jukee) mRNA.
<a href="#">juloy</a>	<a href="#">juloy.aSep08</a>		812	470		61	putative protein (juloy) mRNA.

<a href="#">jumee</a>	<a href="#">jumee.aSep08</a>		13225	359		53	putative protein (jumee) mRNA.
<a href="#">jumer</a>	<a href="#">jumer.aSep08</a>		2758	876		84	putative protein (jumer) mRNA.
<a href="#">junoy</a>	<a href="#">junoy.aSep08</a>		13650	814		189	basic immunoglobulin-like variable motif containing CRA b (junoy) mRNA.
<a href="#">Jup</a>	<a href="#">Jup.bSep08</a>	<a href="#">81679</a>	14713	858	2	285	junction plakoglobin (Jup) alternative variant bSep08, mRNA.
<a href="#">Jup</a>	<a href="#">Jup.cSep08</a>	<a href="#">81679</a>	14288	568	1	189	junction plakoglobin (Jup) alternative variant cSep08, mRNA.
<a href="#">jupor</a>	<a href="#">jupor.aSep08</a>		3553	481		160	CRA a (jupor) mRNA.
<a href="#">jusa</a>	<a href="#">jusa.aSep08</a>		867	781		99	putative protein (jusa) mRNA.
<a href="#">jushee</a>	<a href="#">jushee.aSep08</a>		112917	663	4	171	putative protein (18.8 kD) (jushee) alternative variant aSep08, mRNA.
<a href="#">juvar</a>	<a href="#">juvar.aSep08</a>		806	444		86	putative protein of bilateral origin (juvar) mRNA.
<a href="#">juwey</a>	<a href="#">juwey.aSep08</a>		16035	454		70	putative protein (juwey) mRNA.
<a href="#">jyby</a>	<a href="#">jyby.aSep08</a>		55211	373		96	putative protein (jyby) mRNA.
<a href="#">jychy</a>	<a href="#">jychy.aSep08</a>		5570	479		37	putative protein (jychy) mRNA.
<a href="#">jydar</a>	<a href="#">jydar.aSep08</a>		1390	421		108	putative protein (jydar) mRNA.
<a href="#">jyfer</a>	<a href="#">jyfer.aSep08</a>		2408	666		80	putative protein (jyfer) mRNA.
<a href="#">jyflo</a>	<a href="#">jyflo.aSep08</a>		5977	1190		129	putative protein (14.8 kD) (jyflo) mRNA.
<a href="#">jyflu</a>	<a href="#">jyflu.aSep08</a>		891	759		230	interleukin 4 induced 1 like (jyflu) mRNA.
<a href="#">jykee</a>	<a href="#">jykee.bSep08</a>		5753	506	4	131	putative protein of vertebrate origin (jykee) alternative variant bSep08, mRNA.
<a href="#">jykee</a>	<a href="#">jykee.cSep08</a>		8766	3869	2	65	putative protein of vertebrate origin (jykee) alternative variant cSep08, mRNA.
<a href="#">jykee</a>	<a href="#">jykee.eSep08</a>		1071	419	2	107	putative protein (jykee) alternative variant eSep08, mRNA.
<a href="#">jykee</a>	<a href="#">jykee.gSep08</a>		419	328	2	30	putative protein (3.1 kD) (jykee) alternative variant gSep08, mRNA.
<a href="#">jyloy</a>	<a href="#">jyloy.aSep08</a>		2352	434		119	putative protein (jyloy) mRNA.
<a href="#">jymee</a>	<a href="#">jymee.aSep08</a>		618	384		31	putative protein (jymee) mRNA.
<a href="#">jymer</a>	<a href="#">jymer.aSep08</a>		4013	544		103	putative nuclear protein (11.9 kD) (jymer) mRNA.
<a href="#">jynoy</a>	<a href="#">jynoy.aSep08</a>		9323	735		74	putative protein (jynoy) mRNA.
<a href="#">jypor</a>	<a href="#">jypor.aSep08</a>		2770	390		129	dmx-like 2 (jypor) mRNA.
<a href="#">jysa</a>	<a href="#">jysa.aSep08</a>		20723	922	2	57	putative protein (jysa) alternative variant aSep08, mRNA.
<a href="#">jysa</a>	<a href="#">jysa.bSep08</a>		14958	770	2	67	putative protein (jysa) alternative variant bSep08, mRNA.
<a href="#">jyshee</a>	<a href="#">jyshee.aSep08</a>		5250	354		118	putative protein of ancient origin (jyshee) mRNA.
<a href="#">jyvar</a>	<a href="#">jyvar.aSep08</a>		5689	1801		600	putative protein of bilateral origin (jyvar) alternative variant aSep08, mRNA.
<a href="#">jyvar</a>	<a href="#">jyvar.bSep08</a>		1356	417		138	putative protein of bilateral origin (jyvar) alternative variant bSep08, mRNA.
<a href="#">jywey</a>	<a href="#">jywey.aSep08</a>		3853	369		123	putative protein of ancient origin (jywey) mRNA.
<a href="#">Kab</a>	<a href="#">Kab.aSep08</a>	<a href="#">171457</a>	9218	2627		208	KARP-1 binding protein 1 (Kab) mRNA.
<a href="#">kaby</a>	<a href="#">kaby.aSep08</a>		2925	716		42	putative protein (4.9 kD) (kaby) mRNA.
<a href="#">kachy</a>	<a href="#">kachy.aSep08</a>		481	295		28	putative protein (kachy) mRNA.

<a href="#">kadar</a>	<a href="#">kadar.aSep08</a>		4148	540		96	putative protein (kadar) mRNA.
<a href="#">kafer</a>	<a href="#">kafer.aSep08</a>		4674	502	3	90	putative protein (10.2 kD) (kafer) alternative variant aSep08, mRNA.
<a href="#">kafer</a>	<a href="#">kafer.bSep08</a>		4297	436	3	35	putative protein (4.2 kD) (kafer) alternative variant bSep08, mRNA.
<a href="#">kaflo</a>	<a href="#">kaflo.aSep08</a>		405	281		62	putative protein (kaflo) mRNA.
<a href="#">kaflu</a>	<a href="#">kaflu.aSep08</a>		1356	663		111	putative protein (kaflu) mRNA.
<a href="#">kakee</a>	<a href="#">kakee.aSep08</a>		994	183		60	protein phosphatase 1 regulatory (kakee) mRNA.
<a href="#">kaloy</a>	<a href="#">kaloy.aSep08</a>		39492	390		105	putative protein (kaloy) mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.bSep08</a>	<a href="#">84009</a>	22707	803	5	250	kalirin RhoGEF kinase CRA c (28.6 kD) (Kalrn) alternative variant bSep08, mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.cSep08</a>	<a href="#">84009</a>	59065	737	4	245	kalirin RhoGEF kinase (Kalrn) alternative variant cSep08, mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.dSep08</a>	<a href="#">84009</a>	5688	950	4	200	kalirin RhoGEF kinase CRA c (Kalrn) alternative variant dSep08, mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.eSep08</a>	<a href="#">84009</a>	22854	387	4	124	kalirin RhoGEF kinase (Kalrn) alternative variant eSep08, mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.fSep08</a>	<a href="#">84009</a>	2935	584	2	84	kalirin RhoGEF kinase CRA f (Kalrn) alternative variant fSep08, mRNA.
<a href="#">Kalrn</a>	<a href="#">Kalrn.gSep08</a>	<a href="#">84009</a>	40057	417	2	67	putative protein (Kalrn) alternative variant gSep08, mRNA.
<a href="#">kamee</a>	<a href="#">kamee.aSep08</a>		1054	573		103	putative protein (kamee) mRNA.
<a href="#">kamer</a>	<a href="#">kamer.aSep08</a>		47320	277		65	testis expressed 2 (kamer) mRNA.
<a href="#">Kank3</a>	<a href="#">Kank3.bSep08</a>	<a href="#">366848</a>	2798	2002	5	615	putative protein, with 2 coiled coil domains (Kank3) alternative variant bSep08, mRNA.
<a href="#">Kank3</a>	<a href="#">Kank3.cSep08</a>	<a href="#">366848</a>	856	762	2	70	putative protein (7.6 kD) (Kank3) alternative variant cSep08, mRNA.
<a href="#">Kank4</a>	<a href="#">Kank4.bSep08</a>	<a href="#">313385</a>	19345	1764	5	422	ankyrin (Kank4) alternative variant bSep08, mRNA.
<a href="#">Kank4</a>	<a href="#">Kank4.cSep08</a>	<a href="#">313385</a>	5345	758	3	62	putative protein of mammalian origin (7.0 kD) (Kank4) alternative variant cSep08, mRNA.
<a href="#">kanoy</a>	<a href="#">kanoy.aSep08</a>		1458	411		47	putative protein (kanoy) mRNA.
<a href="#">kapor</a>	<a href="#">kapor.aSep08</a>		7481	208	2	68	putative protein (kapor) alternative variant aSep08, mRNA.
<a href="#">kapor</a>	<a href="#">kapor.bSep08</a>		8274	1026	2	51	putative protein (5.8 kD) (kapor) alternative variant bSep08, complete mRNA.
<a href="#">kapor</a>	<a href="#">kapor.dSep08</a>		9011	398	3	51	putative protein (5.8 kD) (kapor) alternative variant dSep08, mRNA.
<a href="#">karby</a>	<a href="#">karby.aSep08</a>		5346	1034		74	putative protein (karby) mRNA.
<a href="#">karchy</a>	<a href="#">karchy.aSep08</a>		5452	463	1	40	putative protein (karchy) alternative variant aSep08, mRNA.
<a href="#">karchy</a>	<a href="#">karchy.bSep08</a>		1452	99	1	32	putative protein (karchy) alternative variant bSep08, mRNA.
<a href="#">kardar</a>	<a href="#">kardar.aSep08</a>		11180	300		64	putative protein (kardar) mRNA.
<a href="#">karfer</a>	<a href="#">karfer.aSep08</a>		24214	714		237	supervillin (karfer) mRNA.
<a href="#">karflo</a>	<a href="#">karflo.aSep08</a>		1010	229	1	66	putative protein (karflo) alternative variant aSep08, mRNA.
<a href="#">karflo</a>	<a href="#">karflo.bSep08</a>		835	299	1	32	putative protein (3.5 kD) (karflo) alternative variant bSep08, mRNA.

<a href="#">karflu</a>	<a href="#">karflu.aSep08</a>		21785	674		224	guanine nucleotide exchange factor (karflu) mRNA.
<a href="#">karjey</a>	<a href="#">karjey.aSep08</a>		6812	681	3	182	putative protein of vertebrate origin (karjey) alternative variant aSep08, mRNA.
<a href="#">karkee</a>	<a href="#">karkee.aSep08</a>		2106	428		47	putative protein (karkee) mRNA.
<a href="#">karloy</a>	<a href="#">karloy.aSep08</a>		2328	1213		41	putative protein (karloy) mRNA.
<a href="#">karmee</a>	<a href="#">karmee.aSep08</a>		8145	816	3	178	putative protein (karmee) alternative variant aSep08, mRNA.
<a href="#">karmee</a>	<a href="#">karmee.bSep08</a>		9444	752	2	37	putative protein (4.4 kD) (karmee) alternative variant bSep08, mRNA.
<a href="#">karmer</a>	<a href="#">karmer.aSep08</a>		18068	629		115	CRA b like (karmer) mRNA.
<a href="#">karnoy</a>	<a href="#">karnoy.aSep08</a>		2490	412		42	putative protein (karnoy) mRNA.
<a href="#">karpor</a>	<a href="#">karpor.aSep08</a>		2465	493		39	putative protein (karpor) mRNA.
<a href="#">Kars</a>	<a href="#">Kars.bSep08</a>	<a href="#">292028</a>	14015	1564	10	499	lysyl-tRNA synthetase (Kars) alternative variant bSep08, mRNA.
<a href="#">Kars</a>	<a href="#">Kars.cSep08</a>	<a href="#">292028</a>	4276	875	2	107	lysyl-tRNA synthetase (Kars) alternative variant cSep08, mRNA.
<a href="#">karsa</a>	<a href="#">karsa.aSep08</a>		18521	724	6	241	PHD finger protein 20-like 1 (karsa) alternative variant aSep08, mRNA.
<a href="#">karsa</a>	<a href="#">karsa.bSep08</a>		649	417	2	99	tudor domain-containing protein Phf2011 (karsa) alternative variant bSep08, mRNA.
<a href="#">karshee</a>	<a href="#">karshee.aSep08</a>		3181	199	2	63	putative protein (karshee) alternative variant aSep08, mRNA.
<a href="#">kartu</a>	<a href="#">kartu.aSep08</a>		3521	330		25	putative protein (2.9 kD) (kartu) mRNA.
<a href="#">karvar</a>	<a href="#">karvar.aSep08</a>		8727	1010		336	rearranged L-myc fusion (karvar) mRNA.
<a href="#">karwey</a>	<a href="#">karwey.aSep08</a>		1438	634		51	putative protein (5.7 kD) (karwey) mRNA.
<a href="#">kasa</a>	<a href="#">kasa.aSep08</a>		19188	661	3	23	putative protein (2.5 kD) (kasa) alternative variant aSep08, mRNA.
<a href="#">kasa</a>	<a href="#">kasa.bSep08</a>		77452	615	4	46	putative protein (kasa) alternative variant bSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.aSep08</a>		38533	546	2	131	CRA b like (kashee) alternative variant aSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.bSep08</a>		39961	960	3	91	CRA b like (10.4 kD) (kashee) alternative variant bSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.cSep08</a>		25845	584	3	67	putative protein (8.0 kD) (kashee) alternative variant cSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.dSep08</a>		124315	1405	5	91	CRA b like (10.4 kD) (kashee) alternative variant dSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.fSep08</a>		24908	661	3	55	CRA c like (6.1 kD) (kashee) alternative variant fSep08, mRNA.
<a href="#">kashee</a>	<a href="#">kashee.gSep08</a>		25743	658	3	60	putative protein (7.3 kD) (kashee) alternative variant gSep08, mRNA.
<a href="#">Kat3</a>	<a href="#">Kat3.aSep08</a>	<a href="#">541589</a>	14327	636		167	kynurenine aminotransferase III (Kat3) alternative variant aSep08, mRNA.
<a href="#">Kat3</a>	<a href="#">Kat3.bSep08</a>	<a href="#">541589</a>	7950	431		74	kynurenine aminotransferase III (Kat3) alternative variant bSep08, mRNA.
<a href="#">Katnal1</a>	<a href="#">Katnal1.bSep08</a>	<a href="#">288449</a>	44133	798	6	240	katanin p60 subunit A-like 1 (Katnal1) alternative variant bSep08, mRNA.

<a href="#">Katnal1</a>	<a href="#">Katnal1.cSep08</a>	<a href="#">288449</a>	43268	720	5	196	katanin p60 subunit A-like 1 (Katnal1) alternative variant cSep08, mRNA.
<a href="#">Katnal1</a>	<a href="#">Katnal1.dSep08</a>	<a href="#">288449</a>	24034	772	4	135	katanin p60 subunit A-like 1 (Katnal1) alternative variant dSep08, mRNA.
<a href="#">Katnal1</a>	<a href="#">Katnal1.eSep08</a>	<a href="#">288449</a>	2579	426	2	112	katanin p60 subunit A-like 1 (Katnal1) alternative variant eSep08, mRNA.
<a href="#">Katnal1</a>	<a href="#">Katnal1.fSep08</a>	<a href="#">288449</a>	8095	686	2	71	katanin p60 subunit A-like 1 (8.1 kD) (Katnal1) alternative variant fSep08, mRNA.
<a href="#">Katnb1</a>	<a href="#">Katnb1.aSep08</a>	<a href="#">291852</a>	19722	2936	20	659	katanin p80 (WD40-containing) subunit B 1 (72.6 kD) (Katnb1) alternative variant aSep08, mRNA.
<a href="#">Katnb1</a>	<a href="#">Katnb1.bSep08</a>	<a href="#">291852</a>	1863	974	7	230	katanin p80 (WD40-containing) subunit B 1 (Katnb1) alternative variant bSep08, mRNA.
<a href="#">Katnb1</a>	<a href="#">Katnb1.cSep08</a>	<a href="#">291852</a>	1163	697	4	138	katanin p80 (WD40-containing) subunit B 1 (Katnb1) alternative variant cSep08, mRNA.
<a href="#">Katnb1</a>	<a href="#">Katnb1.eSep08</a>	<a href="#">291852</a>	1500	505	2	79	katanin p80 (WD40-containing) subunit B 1 (Katnb1) alternative variant eSep08, mRNA.
<a href="#">katu</a>	<a href="#">katu.aSep08</a>		39750	636		188	cysteine-rich motor neuron 1 (katu) mRNA.
<a href="#">kavar</a>	<a href="#">kavar.aSep08</a>		6639	747		90	putative cytoplasmic protein (9.9 kD) (kavar) mRNA.
<a href="#">kawby</a>	<a href="#">kawby.aSep08</a>		1765	684		163	inter-alpha inhibitor H5-like (kawby) mRNA.
<a href="#">kawchy</a>	<a href="#">kawchy.aSep08</a>		23229	568	3	131	putative protein (kawchy) alternative variant aSep08, mRNA.
<a href="#">kawchy</a>	<a href="#">kawchy.bSep08</a>		6390	411	3	40	putative protein (kawchy) alternative variant bSep08, mRNA.
<a href="#">kawdar</a>	<a href="#">kawdar.aSep08</a>		1957	817		31	putative protein (3.7 kD) (kawdar) mRNA.
<a href="#">kawey</a>	<a href="#">kawey.aSep08</a>		15116	298		32	putative protein (kawey) mRNA.
<a href="#">kawfer</a>	<a href="#">kawfer.aSep08</a>		2453	445		148	supervillin (kawfer) mRNA.
<a href="#">kawflo</a>	<a href="#">kawflo.aSep08</a>		8464	343		114	kinesin family member 11 CRA b (kawflo) mRNA.
<a href="#">kawflu</a>	<a href="#">kawflu.aSep08</a>		4193	818	3	94	ab2-143 like (10.9 kD) (kawflu) alternative variant aSep08, mRNA.
<a href="#">kawjey</a>	<a href="#">kawjey.aSep08</a>		7327	460		153	putative protein, with 2 coiled coil domains, of vertebrate origin (kawjey) mRNA.
<a href="#">kawkee</a>	<a href="#">kawkee.aSep08</a>		11826	1958	4	610	neuron navigator 1 (kawkee) alternative variant aSep08, mRNA.
<a href="#">kawkee</a>	<a href="#">kawkee.bSep08</a>		3827	761	3	253	neuron navigator 1 (kawkee) alternative variant bSep08, mRNA.
<a href="#">kawkee</a>	<a href="#">kawkee.cSep08</a>		7468	1488	5	213	neuron navigator 1 (kawkee) alternative variant cSep08, mRNA.
<a href="#">kawloy</a>	<a href="#">kawloy.aSep08</a>		813	742		82	putative cytoplasmic protein (9.2 kD) (kawloy) mRNA.
<a href="#">kawmee</a>	<a href="#">kawmee.aSep08</a>		1015	685		41	putative protein (kawmee) mRNA.
<a href="#">kawmer</a>	<a href="#">kawmer.aSep08</a>		1228	247		34	putative protein (kawmer) mRNA.
<a href="#">kawnoy</a>	<a href="#">kawnoy.bSep08</a>		902	380	3	26	putative protein (3.0 kD) (kawnoy) alternative variant bSep08, mRNA.
<a href="#">kawpor</a>	<a href="#">kawpor.aSep08</a>		2771	927		107	putative protein (kawpor) mRNA.
<a href="#">kawsa</a>	<a href="#">kawsa.aSep08</a>		5183	2332	3	118	PHD finger protein 20-like 1 (kawsa) alternative variant aSep08, mRNA.

<a href="#">kawsa</a>	<a href="#">kawsa.bSep08</a>		5032	1782	3	58	PHD finger protein 20 (kawsa) alternative variant bSep08, mRNA.
<a href="#">kawshee</a>	<a href="#">kawshee.aSep08</a>		30761	434		63	e1-E2 ATPase like (kawshee) mRNA.
<a href="#">kawtu</a>	<a href="#">kawtu.aSep08</a>		2547	226		15	putative protein (1.6 kD) (kawtu) mRNA.
<a href="#">kawvar</a>	<a href="#">kawvar.aSep08</a>		1193	245		39	putative protein (kawvar) mRNA.
<a href="#">kawwey</a>	<a href="#">kawwey.aSep08</a>		19755	3931	2	123	putative protein of vertebrate origin (13.9 kD) (kawwey) alternative variant aSep08, mRNA.
<a href="#">kawwey</a>	<a href="#">kawwey.bSep08</a>		16257	344	1	76	guanylate kinase 1 (kawwey) alternative variant bSep08, mRNA.
<a href="#">Kazald1</a>	<a href="#">Kazald1.bSep08</a>	<a href="#">293997</a>	3083	763	3	105	kazal-type serine peptidase inhibitor domain 1 (11.7 kD) (Kazald1) alternative variant bSep08, mRNA.
<a href="#">Kazald1</a>	<a href="#">Kazald1.cSep08</a>	<a href="#">293997</a>	2475	2208	2	105	kazal-type serine peptidase inhibitor domain 1 (11.7 kD) (Kazald1) alternative variant cSep08, mRNA.
<a href="#">Kb23</a>	<a href="#">Kb23.bSep08</a>	<a href="#">407759</a>	6516	1562		479	type II keratin Kb23 (52.7 kD) (Kb23) alternative variant bSep08, mRNA.
<a href="#">Kbtbd3</a>	<a href="#">Kbtbd3.bSep08</a>	<a href="#">315394</a>	24057	2624	3	607	BTB/POZ and BTB/Kelch-associated and Kelch repeat containing protein and kelch (69.5 kD) (Kbtbd3) alternative variant bSep08, mRNA.
<a href="#">Kbtbd4</a>	<a href="#">Kbtbd4.aSep08</a>	<a href="#">311185</a>	5993	2410	4	518	BTB/POZ and K <sup>+</sup> channel tetramerisation containing protein and BTB/Kelch-associated and kelch and Kelch repeat containing protein (58.1 kD) (Kbtbd4) alternative variant aSep08, mRNA.
<a href="#">Kbtbd8</a>	<a href="#">Kbtbd8.bSep08</a>	<a href="#">500262</a>	5377	420	1	83	kelch repeat (Kbtbd8) alternative variant bSep08, mRNA.
<a href="#">Kcnab2</a>	<a href="#">Kcnab2.aSep08</a>	<a href="#">29738</a>	27770	596	1	198	potassium voltage-gated channel, shaker-related subfamily, beta member 2 (Kcnab2) alternative variant aSep08, mRNA.
<a href="#">Kcnab2</a>	<a href="#">Kcnab2.bSep08</a>	<a href="#">29738</a>	78426	456	1	99	potassium voltage-gated channel, shaker-related subfamily, beta member 2 (Kcnab2) alternative variant bSep08, mRNA.
<a href="#">Kcnc1</a>	<a href="#">Kcnc1.bSep08</a>	<a href="#">25327</a>	29978	1249	2	416	potassium voltage gated channel, Shaw-related subfamily, member 1 (Kcnc1) alternative variant bSep08, mRNA.
<a href="#">Kcnc1</a>	<a href="#">Kcnc1.cSep08</a>	<a href="#">25327</a>	4038	2107	2	98	potassium voltage gated channel, Shaw-related subfamily, member 1 (10.6 kD) (Kcnc1) alternative variant cSep08, mRNA.
<a href="#">Kcnd2</a>	<a href="#">Kcnd2.bSep08</a>	<a href="#">65180</a>	2218	727	1	111	potassium voltage gated channel, Shal-related family, member 2 (Kcnd2) alternative variant bSep08, mRNA.
<a href="#">Kcnd3</a>	<a href="#">Kcnd3.bSep08</a>	<a href="#">65195</a>	3641	493	4	164	potassium voltage gated channel, Shal-related family, member 3 (Kcnd3) alternative variant bSep08, mRNA.
<a href="#">Kcne1</a>	<a href="#">Kcne1.bSep08</a>	<a href="#">25471</a>	10332	736	2	130	potassium voltage-gated channel, Isk-related subfamily, member 1 (14.7 kD) (Kcne1) alternative variant bSep08, mRNA.
<a href="#">Kcne3</a>	<a href="#">Kcne3.bSep08</a>	<a href="#">63883</a>	6621	956	2	107	potassium voltage-gated channel, Isk-related subfamily, gene 3 (12.0 kD) (Kcne3) alternative variant bSep08, complete mRNA.
<a href="#">Kcng2</a>	<a href="#">Kcng2.cSep08</a>	<a href="#">307234</a>	19718	520	3	57	potassium voltage-gated channel, subfamily G, member 2 (6.3 kD) (Kcng2) alternative variant cSep08, mRNA.

<a href="#">Kcng4</a>	<a href="#">Kcng4.bSep08</a>	<a href="#">307900</a>	776	414	2	107	potassium voltage-gated channel, subfamily G, member 4 (Kcng4) alternative variant bSep08, mRNA.
<a href="#">Kcnh2</a>	<a href="#">Kcnh2.bSep08</a>	<a href="#">117018</a>	597	374	2	113	potassium voltage-gated channel, subfamily H (eag-related), member 2 (Kcnh2) alternative variant bSep08, mRNA.
<a href="#">Kcnp1</a>	<a href="#">Kcnp1.aSep08</a>	<a href="#">65023</a>	15104	1260		151	kv channel-interacting protein 1 (Kcnp1) mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.cSep08</a>	<a href="#">56817</a>	18647	869	7	246	channel-interacting protein 2 (Kcnp2andKchip2) alternative variant cSep08, mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.cSep08</a>	<a href="#">619380</a>	18647	869	7	246	channel-interacting protein 2 (Kcnp2andKchip2) alternative variant cSep08, mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.fSep08</a>	<a href="#">56817</a>	1662	407	3	38	CRA e like (4.4 kD) (Kcnp2andKchip2) alternative variant fSep08, mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.fSep08</a>	<a href="#">619380</a>	1662	407	3	38	CRA e like (4.4 kD) (Kcnp2andKchip2) alternative variant fSep08, mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.gSep08</a>	<a href="#">56817</a>	1895	301	3	46	CRA e like (Kcnp2andKchip2) alternative variant gSep08, mRNA.
<a href="#">Kcnp2andKchip2</a>	<a href="#">Kcnp2andKchip2.gSep08</a>	<a href="#">619380</a>	1895	301	3	46	CRA e like (Kcnp2andKchip2) alternative variant gSep08, mRNA.
<a href="#">Kcnp3</a>	<a href="#">Kcnp3.bSep08</a>	<a href="#">65199</a>	18959	803	4	122	kv channel interacting protein 3, calsenilin (Kcnp3) alternative variant bSep08, mRNA.
<a href="#">Kcnp3</a>	<a href="#">Kcnp3.cSep08</a>	<a href="#">65199</a>	3832	2166	4	116	kv channel interacting protein 3, calsenilin (Kcnp3) alternative variant cSep08, mRNA.
<a href="#">Kcnp4</a>	<a href="#">Kcnp4.bSep08</a>	<a href="#">259243</a>	3049	744	2	35	kv channel interacting protein 4 (4.1 kD) (Kcnp4) alternative variant bSep08, mRNA.
<a href="#">Kcnj1</a>	<a href="#">Kcnj1.bSep08</a>	<a href="#">24521</a>	29861	2298	2	264	potassium inwardly-rectifying channel, subfamily J, member 1 (29.8 kD) (Kcnj1) alternative variant bSep08, mRNA.
<a href="#">Kcnj1</a>	<a href="#">Kcnj1.cSep08</a>	<a href="#">24521</a>	3654	755	2	63	potassium inwardly-rectifying channel, subfamily J, member 1 (7.0 kD) (Kcnj1) alternative variant cSep08, mRNA.
<a href="#">Kcnj3</a>	<a href="#">Kcnj3.bSep08</a>	<a href="#">50599</a>	10264	1357	1	323	potassium inwardly-rectifying channel, subfamily J, member 3 (Kcnj3) alternative variant bSep08, mRNA.
<a href="#">Kcnj6</a>	<a href="#">Kcnj6.aSep08</a>	<a href="#">25743</a>	12392	535		44	potassium inwardly-rectifying channel, subfamily J, member 6 (Kcnj6) mRNA.
<a href="#">Kcnj8</a>	<a href="#">Kcnj8.bSep08</a>	<a href="#">25472</a>	3488	1822	2	303	potassium inwardly-rectifying channel, subfamily J, member 8 (Kcnj8) alternative variant bSep08, mRNA.
<a href="#">Kcnj9</a>	<a href="#">Kcnj9.cSep08</a>	<a href="#">116560</a>	7268	1595	3	116	potassium inwardly-rectifying channel, subfamily J, member 9 (12.9 kD) (Kcnj9) alternative variant cSep08, mRNA.
<a href="#">Kcnj13</a>	<a href="#">Kcnj13.bSep08</a>	<a href="#">94341</a>	6495	646	1	171	potassium inwardly-rectifying channel, subfamily J, member 13 (19.6 kD) (Kcnj13) alternative variant bSep08, complete mRNA.
<a href="#">Kcnj15</a>	<a href="#">Kcnj15.bSep08</a>	<a href="#">170847</a>	41854	1659	3	375	potassium inwardly-rectifying channel, subfamily J, member 15 (42.6 kD) (Kcnj15) alternative variant bSep08, mRNA.
<a href="#">Kcnj16</a>	<a href="#">Kcnj16.bSep08</a>	<a href="#">29719</a>	27730	324	2	26	potassium inwardly-rectifying channel, subfamily J, member 16 (Kcnj16) alternative variant bSep08, mRNA.

<a href="#">Kcnk1</a>	<a href="#">Kcnk1.bSep08</a>	<a href="#">59324</a>	1288	968	2	59	potassium channel, subfamily K, member 1 (Kcnk1) alternative variant bSep08, mRNA.
<a href="#">Kcnk1</a>	<a href="#">Kcnk1.cSep08</a>	<a href="#">59324</a>	1055	735	2	62	potassium channel, subfamily K, member 1 (Kcnk1) alternative variant cSep08, mRNA.
<a href="#">Kcnk1</a>	<a href="#">Kcnk1.dSep08</a>	<a href="#">59324</a>	773	453	2	39	potassium channel, subfamily K, member 1 (Kcnk1) alternative variant dSep08, mRNA.
<a href="#">Kcnk4</a>	<a href="#">Kcnk4.aSep08</a>	<a href="#">116489</a>	6466	1782	1	520	potassium channel, subfamily K, member 4 (Kcnk4) alternative variant aSep08, mRNA.
<a href="#">Kcnk4</a>	<a href="#">Kcnk4.cSep08</a>	<a href="#">116489</a>	1367	811	2	129	potassium channel, subfamily K, member 4 (13.5 kD) (Kcnk4) alternative variant cSep08, mRNA.
<a href="#">Kcnk6</a>	<a href="#">Kcnk6.bSep08</a>	<a href="#">116491</a>	877	803	2	216	potassium inwardly-rectifying channel, subfamily K, member 6 (Kcnk6) alternative variant bSep08, mRNA.
<a href="#">Kcnk18</a>	<a href="#">Kcnk18.bSep08</a>	<a href="#">445371</a>	15450	1200		362	potassium channel, subfamily K, member 18 (40.3 kD) (Kcnk18) alternative variant bSep08, mRNA.
<a href="#">Kcnma1</a>	<a href="#">Kcnma1.aSep08</a>	<a href="#">83731</a>	356431	3271	26	997	potassium large conductance calcium-activated channel, subfamily M, alpha member 1 (Kcnma1) alternative variant aSep08, mRNA.
<a href="#">Kcnma1</a>	<a href="#">Kcnma1.bSep08</a>	<a href="#">83731</a>	14538	851	4	187	potassium large conductance calcium-activated channel, subfamily M, alpha member 1 (Kcnma1) alternative variant bSep08, mRNA.
<a href="#">Kcnmb1</a>	<a href="#">Kcnmb1.bSep08</a>	<a href="#">29747</a>	4913	1058	2	87	potassium large conductance calcium-activated channel, subfamily M, beta member 1 (Kcnmb1) alternative variant bSep08, mRNA.
<a href="#">Kcnmb2</a>	<a href="#">Kcnmb2.bSep08</a>	<a href="#">294961</a>	321839	1544	5	174	potassium large conductance calcium-activated channel, subfamily M, beta member 2 (20.0 kD) (Kcnmb2) alternative variant bSep08, complete mRNA.
<a href="#">Kcnmb2</a>	<a href="#">Kcnmb2.cSep08</a>	<a href="#">294961</a>	37674	341	1	110	potassium large conductance calcium-activated channel, subfamily M, beta member 2 (12.9 kD) (Kcnmb2) alternative variant cSep08, mRNA.
<a href="#">Kcnn1</a>	<a href="#">Kcnn1.bSep08</a>	<a href="#">54261</a>	4451	608	4	202	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1 (Kcnn1) alternative variant bSep08, mRNA.
<a href="#">Kcnn1</a>	<a href="#">Kcnn1.cSep08</a>	<a href="#">54261</a>	2108	700	2	96	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1 (Kcnn1) alternative variant cSep08, mRNA.
<a href="#">Kcnn3</a>	<a href="#">Kcnn3.bSep08</a>	<a href="#">54263</a>	6401	691	1	116	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3 (12.7 kD) (Kcnn3) alternative variant bSep08, mRNA.
<a href="#">Kcnn4</a>	<a href="#">Kcnn4.bSep08</a>	<a href="#">65206</a>	7556	935	6	310	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4 (Kcnn4) alternative variant bSep08, mRNA.
<a href="#">Kcnn4</a>	<a href="#">Kcnn4.cSep08</a>	<a href="#">65206</a>	5386	896	5	135	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4 (15.8 kD) (Kcnn4) alternative variant cSep08, mRNA.
<a href="#">Kcnn4</a>	<a href="#">Kcnn4.dSep08</a>	<a href="#">65206</a>	3114	787	3	77	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4 (Kcnn4) alternative variant dSep08, mRNA.



<a href="#">Kcnq1</a>	<a href="#">Kcnq1.bSep08</a>	<a href="#">84020</a>	22824	862	3	206	potassium voltage-gated channel, subfamily Q, member 1 (Kcnq1) alternative variant bSep08, mRNA.
<a href="#">Kcnq1</a>	<a href="#">Kcnq1.cSep08</a>	<a href="#">84020</a>	3215	406	1	129	potassium voltage-gated channel, subfamily Q, member 1 (Kcnq1) alternative variant cSep08, mRNA.
<a href="#">Kcnq1</a>	<a href="#">Kcnq1.dSep08</a>	<a href="#">84020</a>	66139	1252	3	103	potassium voltage-gated channel, subfamily Q, member 1 (Kcnq1) alternative variant dSep08, mRNA.
<a href="#">Kcnq5l</a>	<a href="#">Kcnq5l.aSep08</a>	<a href="#">259273</a>	499563	1792	4	597	potassium voltage-gated channel, subfamily Q, member 5-like (Kcnq5l) alternative variant aSep08, mRNA.
<a href="#">Kcnq5l</a>	<a href="#">Kcnq5l.bSep08</a>	<a href="#">259273</a>	448899	465	3	154	potassium voltage-gated channel, subfamily Q, member 5-like (Kcnq5l) alternative variant bSep08, mRNA.
<a href="#">KCNQ_channel.0</a>	<a href="#">KCNQ_channel.0.aSep08</a>	<a href="#">689123</a>	61242	421		139	voltage-gated potassium channel KCNQ5 (KCNQ_channel.0) mRNA.
<a href="#">Kcns3</a>	<a href="#">Kcns3.bSep08</a>	<a href="#">83588</a>	58971	2329	1	491	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 (55.9 kD) (Kcns3) alternative variant bSep08, mRNA.
<a href="#">Kcnt1</a>	<a href="#">Kcnt1.bSep08</a>	<a href="#">60444</a>	6429	1021	5	287	potassium channel subfamily T member 1 (Kcnt1) alternative variant bSep08, mRNA.
<a href="#">Kcnt1</a>	<a href="#">Kcnt1.cSep08</a>	<a href="#">60444</a>	6321	1877	3	193	potassium channel subfamily T member 1 (21.5 kD) (Kcnt1) alternative variant cSep08, mRNA.
<a href="#">Kcnt1</a>	<a href="#">Kcnt1.dSep08</a>	<a href="#">60444</a>	750	625	2	140	putative protein (15.3 kD) (Kcnt1) alternative variant dSep08, mRNA.
<a href="#">Kcnt2</a>	<a href="#">Kcnt2.aSep08</a>	<a href="#">304827</a>	88321	406		71	potassium channel, subfamily T, member 2 (Kcnt2) mRNA.
<a href="#">Kctd1</a>	<a href="#">Kctd1.aSep08</a>	<a href="#">291772</a>	100561	2071	5	257	hypothetical protein LOC680411 (29.4 kD) (Kctd1) alternative variant aSep08, mRNA.
<a href="#">Kctd1</a>	<a href="#">Kctd1.aSep08</a>	<a href="#">680411</a>	100561	2071	5	257	hypothetical protein LOC680411 (29.4 kD) (Kctd1) alternative variant aSep08, mRNA.
<a href="#">Kctd1</a>	<a href="#">Kctd1.bSep08</a>	<a href="#">291772</a>	21609	760	4	175	hypothetical protein LOC680411 (20.4 kD) (Kctd1) alternative variant bSep08, mRNA.
<a href="#">Kctd1</a>	<a href="#">Kctd1.bSep08</a>	<a href="#">680411</a>	21609	760	4	175	hypothetical protein LOC680411 (20.4 kD) (Kctd1) alternative variant bSep08, mRNA.
<a href="#">Kctd2</a>	<a href="#">Kctd2.aSep08</a>	<a href="#">498024</a>	11655	1463		171	BTB POZ domain-containing protein kctd2 (Kctd2) mRNA.
<a href="#">Kctd3</a>	<a href="#">Kctd3.bSep08</a>	<a href="#">305055</a>	3610	1783	1	216	putative protein of vertebrate origin (Kctd3) alternative variant bSep08, mRNA.
<a href="#">Kctd9</a>	<a href="#">Kctd9.bSep08</a>	<a href="#">364410</a>	3637	1202	3	110	pentapeptide repeat (11.7 kD) (Kctd9) alternative variant bSep08, mRNA.
<a href="#">Kctd9</a>	<a href="#">Kctd9.cSep08</a>	<a href="#">364410</a>	8863	596	6	80	putative protein of bilateral origin (Kctd9) alternative variant cSep08, mRNA.
<a href="#">Kctd11</a>	<a href="#">Kctd11.bSep08</a>	<a href="#">363634</a>	3569	686	2	33	putative protein (Kctd11) alternative variant bSep08, mRNA.
<a href="#">Kctd13</a>	<a href="#">Kctd13.bSep08</a>	<a href="#">293497</a>	3841	1336	2	159	containing potassium channel tetramerisation domain 1 (18.0 kD) (Kctd13) alternative variant bSep08, mRNA.
<a href="#">Kctd13</a>	<a href="#">Kctd13.cSep08</a>	<a href="#">293497</a>	6473	820	3	152	containing potassium channel tetramerisation domain 1 (16.1 kD) (Kctd13) alternative variant cSep08, mRNA.
<a href="#">Kctd14</a>	<a href="#">Kctd14.aSep08</a>	<a href="#">308836</a>	4359	688	2	229	K <sup>+</sup> channel tetramerisation containing protein (Kctd14) alternative variant aSep08, mRNA.
<a href="#">Kctd14</a>	<a href="#">Kctd14.bSep08</a>	<a href="#">308836</a>	2404	1191	2	202	K <sup>+</sup> channel tetramerisation containing protein (23.4 kD) (Kctd14) alternative variant bSep08, mRNA.

<a href="#">Kctd15</a>	<a href="#">Kctd15.bSep08</a>	<a href="#">499129</a>	14859	2403	2	283	K+ channel tetramerisation containing protein (31.9 kD) (Kctd15) alternative variant bSep08, mRNA.
<a href="#">Kctd18</a>	<a href="#">Kctd18.bSep08</a>	<a href="#">301436</a>	7331	1088	4	209	putative nuclear protein of vertebrate origin (22.3 kD) (Kctd18) alternative variant bSep08, mRNA.
<a href="#">Kctd18</a>	<a href="#">Kctd18.cSep08</a>	<a href="#">301436</a>	7348	820	4	117	K+ channel tetramerisation containing protein (13.4 kD) (Kctd18) alternative variant cSep08, mRNA.
<a href="#">Kctd20</a>	<a href="#">Kctd20.aSep08</a>	<a href="#">294307</a>	10051	408		135	putative protein of eukaryotic origin (Kctd20) mRNA.
<a href="#">Kdelc1</a>	<a href="#">Kdelc1.aSep08</a>	<a href="#">316370</a>	12263	2137	10	502	KDEL (Lys-Asp-Glu-Leu) containing 1 (58.1 kD) (Kdelc1) alternative variant aSep08, mRNA.
<a href="#">Kdelc1</a>	<a href="#">Kdelc1.cSep08</a>	<a href="#">316370</a>	4819	673	5	184	KDEL (Lys-Asp-Glu-Leu) containing 1 (Kdelc1) alternative variant cSep08, mRNA.
<a href="#">Kdelc1</a>	<a href="#">Kdelc1.dSep08</a>	<a href="#">316370</a>	3773	684	4	161	KDEL (Lys-Asp-Glu-Leu) containing 1 (Kdelc1) alternative variant dSep08, mRNA.
<a href="#">Kdelc1</a>	<a href="#">Kdelc1.eSep08</a>	<a href="#">316370</a>	45204	641	5	146	KDEL (Lys-Asp-Glu-Leu) containing 1 (Kdelc1) alternative variant eSep08, mRNA.
<a href="#">Kdelc2</a>	<a href="#">Kdelc2.bSep08</a>	<a href="#">315664</a>	4155	644	3	146	KDEL (Lys-Asp-Glu-Leu) containing 2 (Kdelc2) alternative variant bSep08, mRNA.
<a href="#">Kdelr1</a>	<a href="#">Kdelr1.bSep08</a>	<a href="#">361577</a>	3542	287	3	95	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1 (Kdelr1) alternative variant bSep08, mRNA.
<a href="#">Kdelr2</a>	<a href="#">Kdelr2.aSep08</a>	<a href="#">304290</a>	15124	719	3	203	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 (Kdelr2) alternative variant aSep08, mRNA.
<a href="#">Kdelr2</a>	<a href="#">Kdelr2.bSep08</a>	<a href="#">304290</a>	2064	726	1	122	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 (Kdelr2) alternative variant bSep08, mRNA.
<a href="#">Kdelr3</a>	<a href="#">Kdelr3.bSep08</a>	<a href="#">315131</a>	7958	580	5	159	kdel receptor (Kdelr3) alternative variant bSep08, mRNA.
<a href="#">Kdelr3</a>	<a href="#">Kdelr3.cSep08</a>	<a href="#">315131</a>	2782	872	3	106	ER lumen protein retaining receptor (Kdelr3) alternative variant cSep08, mRNA.
<a href="#">Kdelr3</a>	<a href="#">Kdelr3.dSep08</a>	<a href="#">315131</a>	2134	907	2	78	kdel receptor 3 (9.2 kD) (Kdelr3) alternative variant dSep08, mRNA.
<a href="#">Ke2</a>	<a href="#">Ke2.bSep08</a>	<a href="#">309629</a>	1474	587	5	127	prefoldin (14.5 kD) (Ke2) alternative variant bSep08, mRNA.
<a href="#">keeby</a>	<a href="#">keeby.aSep08</a>		34748	718		238	constitutive coactivator of ppar-gamma-like Protein 2 (keeby) mRNA.
<a href="#">keechy</a>	<a href="#">keechy.aSep08</a>		1447	729		35	putative protein (3.9 kD) (keechy) mRNA.
<a href="#">keedar</a>	<a href="#">keedar.aSep08</a>		2724	760		83	putative protein (keedar) mRNA.
<a href="#">keefer</a>	<a href="#">keefer.aSep08</a>		31489	580	1	157	CRA a (keefer) alternative variant aSep08, mRNA.
<a href="#">keefer</a>	<a href="#">keefer.bSep08</a>		22220	638	3	120	CRA b (13.2 kD) (keefer) alternative variant bSep08, mRNA.
<a href="#">keeflo</a>	<a href="#">keeflo.aSep08</a>		9830	376		125	kinesin family member 11 CRA b (keeflo) mRNA.
<a href="#">keeflu</a>	<a href="#">keeflu.aSep08</a>		694	364		43	putative protein (4.7 kD) (keeflu) mRNA.
<a href="#">keejey</a>	<a href="#">keejey.aSep08</a>		1105	329		67	putative protein (keejey) mRNA.
<a href="#">keekee</a>	<a href="#">keekee.aSep08</a>		11254	1292		430	neuron navigator 1 (keekee) mRNA.
<a href="#">keeloy</a>	<a href="#">keeloy.aSep08</a>		6115	538	3	35	putative protein (keeloy) alternative variant aSep08, mRNA.

<a href="#">keemee</a>	<a href="#">keemee.aSep08</a>		15001	1091		166	putative protein (keemee) mRNA.
<a href="#">keemer</a>	<a href="#">keemer.aSep08</a>		36928	678		40	putative protein (4.3 kD) (keemer) mRNA.
<a href="#">keenoy</a>	<a href="#">keenoy.aSep08</a>		3468	468		55	putative protein (keenoy) mRNA.
<a href="#">keepor</a>	<a href="#">keepor.aSep08</a>		7543	612		150	enhancer of mRNA decapping 3 (keepor) mRNA.
<a href="#">keesa</a>	<a href="#">keesa.aSep08</a>		5843	700		33	putative protein (3.7 kD) (keesa) mRNA.
<a href="#">keeshee</a>	<a href="#">keeshee.aSep08</a>		8078	434		91	probable phospholipid-transporting ATPase if like (keeshee) mRNA.
<a href="#">keetu</a>	<a href="#">keetu.aSep08</a>		1568	770		97	putative protein (10.5 kD) (keetu) mRNA.
<a href="#">keevar</a>	<a href="#">keevar.aSep08</a>		4927	738		108	putative protein (11.5 kD) (keevar) mRNA.
<a href="#">keewey</a>	<a href="#">keewey.aSep08</a>		6318	826		31	putative protein (keewey) mRNA.
<a href="#">Kel</a>	<a href="#">Kel.aSep08</a>	<a href="#">297025</a>	11220	2236	8	292	kell blood group like (Kel) alternative variant aSep08, mRNA.
<a href="#">Kel</a>	<a href="#">Kel.bSep08</a>	<a href="#">297025</a>	12568	1123	10	276	kell blood group like (Kel) alternative variant bSep08, mRNA.
<a href="#">Kel</a>	<a href="#">Kel.cSep08</a>	<a href="#">297025</a>	3681	642	4	213	kell blood group like (Kel) alternative variant cSep08, mRNA.
<a href="#">Kel</a>	<a href="#">Kel.dSep08</a>	<a href="#">297025</a>	9800	761	8	209	kell blood group like (Kel) alternative variant dSep08, mRNA.
<a href="#">Kel</a>	<a href="#">Kel.eSep08</a>	<a href="#">297025</a>	2750	474	4	120	kell blood group like (Kel) alternative variant eSep08, mRNA.
<a href="#">Kel</a>	<a href="#">Kel.fSep08</a>	<a href="#">297025</a>	2094	821	3	60	kell blood group like (Kel) alternative variant fSep08, mRNA.
<a href="#">Kelch_1.1</a>	<a href="#">Kelch_1.1.aSep08</a>		4166	409		136	kelch-like 18 CRA c (Kelch_1.1) mRNA.
<a href="#">Kelch_1.2</a>	<a href="#">Kelch_1.2.aSep08</a>		29907	1783		533	kelch-like 32 (Kelch_1.2) alternative variant aSep08, mRNA.
<a href="#">Kelch_1.2</a>	<a href="#">Kelch_1.2.bSep08</a>		12693	508		108	kelch-like 32 (Kelch_1.2) alternative variant bSep08, mRNA.
<a href="#">kerby</a>	<a href="#">kerby.aSep08</a>		1309	530		102	enolase 1 like (10.9 kD) (kerby) mRNA.
<a href="#">kerchy</a>	<a href="#">kerchy.aSep08</a>		675	523		34	putative protein (kerchy) mRNA.
<a href="#">kerdar</a>	<a href="#">kerdar.aSep08</a>		4848	319		89	putative protein (kerdar) mRNA.
<a href="#">kerfer</a>	<a href="#">kerfer.aSep08</a>		12384	516		50	putative protein (kerfer) mRNA.
<a href="#">kerflo</a>	<a href="#">kerflo.aSep08</a>		6792	545		40	putative protein (kerflo) mRNA.
<a href="#">kerflu</a>	<a href="#">kerflu.aSep08</a>		6698	678	5	139	hermansky-Pudlak syndrome 5 like (kerflu) alternative variant aSep08, mRNA.
<a href="#">kerflu</a>	<a href="#">kerflu.bSep08</a>		4438	1391	4	120	hermansky-Pudlak syndrome 5 like (kerflu) alternative variant bSep08, mRNA.
<a href="#">kerjey</a>	<a href="#">kerjey.aSep08</a>		11956	680		101	putative mitochondrial protein (11.3 kD) (kerjey) mRNA.
<a href="#">kerkee</a>	<a href="#">kerkee.aSep08</a>		1934	772		257	neuron navigator 1 (kerkee) mRNA.
<a href="#">kerloy</a>	<a href="#">kerloy.aSep08</a>		3571	779		93	putative protein (10.1 kD) (kerloy) mRNA.
<a href="#">kermee</a>	<a href="#">kermee.aSep08</a>		11101	542		180	ATP-binding cassette sub-family A member 3 like (kermee) alternative variant aSep08, mRNA.
<a href="#">kermer</a>	<a href="#">kermer.aSep08</a>		1493	459	1	56	putative protein (6.5 kD) (kermer) alternative variant aSep08, mRNA.
<a href="#">kermer</a>	<a href="#">kermer.bSep08</a>		2630	1589	2	51	putative protein (kermer) alternative variant bSep08, mRNA.

<a href="#">kernoy</a>	<a href="#">kernoy.aSep08</a>		7702	560		45	putative protein (5.0 kD) (kernoy) mRNA.
<a href="#">kerpor</a>	<a href="#">kerpor.aSep08</a>		32080	792	2	34	CRA a like (kerpor) alternative variant aSep08, mRNA.
<a href="#">kerpor</a>	<a href="#">kerpor.bSep08</a>		20872	652	3	37	CRA a like (4.2 kD) (kerpor) alternative variant bSep08, mRNA.
<a href="#">kerpor</a>	<a href="#">kerpor.cSep08</a>		20821	468	2	46	CRA b like (5.3 kD) (kerpor) alternative variant cSep08, mRNA.
<a href="#">kersa</a>	<a href="#">kersa.aSep08</a>		25718	692	3	64	putative protein (kersa) alternative variant aSep08, mRNA.
<a href="#">kersa</a>	<a href="#">kersa.bSep08</a>		10323	394	2	48	putative protein (5.6 kD) (kersa) alternative variant bSep08, mRNA.
<a href="#">kershee</a>	<a href="#">kershee.aSep08</a>		21907	1939	6	288	ATPase Class VI type 11B (kershee) alternative variant aSep08, mRNA.
<a href="#">kershee</a>	<a href="#">kershee.cSep08</a>		5935	466	2	26	putative protein (3.0 kD) (kershee) alternative variant cSep08, mRNA.
<a href="#">kertu</a>	<a href="#">kertu.aSep08</a>		6666	871	3	290	deah box polypeptide 57 (kertu) alternative variant aSep08, mRNA.
<a href="#">kertu</a>	<a href="#">kertu.bSep08</a>		8839	613	5	204	deah box polypeptide 57 (kertu) alternative variant bSep08, mRNA.
<a href="#">kervar</a>	<a href="#">kervar.aSep08</a>		3367	391		130	microtubule-actin crosslinking factor 1 (kervar) mRNA.
<a href="#">kerwey</a>	<a href="#">kerwey.aSep08</a>		1306	418		61	putative protein (kerwey) mRNA.
<a href="#">keyby</a>	<a href="#">keyby.aSep08</a>		1279	409		33	putative protein (3.9 kD) (keyby) mRNA.
<a href="#">keychy</a>	<a href="#">keychy.aSep08</a>		13296	641		63	putative protein (7.5 kD) (keychy) mRNA.
<a href="#">keydar</a>	<a href="#">keydar.aSep08</a>		962	359		63	putative protein (7.1 kD) (keydar) mRNA.
<a href="#">keyfer</a>	<a href="#">keyfer.aSep08</a>		8110	630	2	143	enhancer of polycomb 1 (keyfer) alternative variant aSep08, mRNA.
<a href="#">keyfer</a>	<a href="#">keyfer.bSep08</a>		1379	390	3	130	enhancer of polycomb 1 (keyfer) alternative variant bSep08, mRNA.
<a href="#">keyflo</a>	<a href="#">keyflo.aSep08</a>		1049	646		51	putative protein (5.7 kD) (keyflo) mRNA.
<a href="#">keyflu</a>	<a href="#">keyflu.aSep08</a>		4408	1220	2	370	hermansky-Pudlak syndrome 5 like (keyflu) alternative variant aSep08, mRNA.
<a href="#">keyflu</a>	<a href="#">keyflu.bSep08</a>		3764	706	3	122	CRA a like (keyflu) alternative variant bSep08, mRNA.
<a href="#">keyflu</a>	<a href="#">keyflu.cSep08</a>		2982	400	1	82	hermansky-Pudlak syndrome 5 like (keyflu) alternative variant cSep08, mRNA.
<a href="#">keyjey</a>	<a href="#">keyjey.aSep08</a>		46769	653	4	154	ecotropic viral integration site 5 (keyjey) alternative variant aSep08, mRNA.
<a href="#">keyjey</a>	<a href="#">keyjey.bSep08</a>		46304	761	3	127	ecotropic viral integration site 5 (14.0 kD) (keyjey) alternative variant bSep08, complete mRNA.
<a href="#">keyjey</a>	<a href="#">keyjey.cSep08</a>		18749	400	2	40	putative protein (keyjey) alternative variant cSep08, mRNA.
<a href="#">keyjey</a>	<a href="#">keyjey.dSep08</a>		24079	368	2	50	putative protein (keyjey) alternative variant dSep08, mRNA.
<a href="#">keykee</a>	<a href="#">keykee.aSep08</a>		2787	226		75	putative protein (keykee) mRNA.
<a href="#">keyloy</a>	<a href="#">keyloy.aSep08</a>		36794	771		50	putative protein (5.7 kD) (keyloy) mRNA.
<a href="#">keymee</a>	<a href="#">keymee.aSep08</a>		3777	618	4	206	ATP-binding cassette sub-family A member 3 like (keymee) alternative variant aSep08, mRNA.
<a href="#">keymer</a>	<a href="#">keymer.aSep08</a>		11831	390		58	putative protein (keymer) mRNA.
<a href="#">keynoy</a>	<a href="#">keynoy.aSep08</a>		16362	297		99	putative protein (keynoy) mRNA.
<a href="#">keypor</a>	<a href="#">keypor.aSep08</a>		2705	449		92	putative protein (9.2 kD) (keypor) mRNA.

<a href="#">keysa</a>	<a href="#">keysa.aSep08</a>		18850	244		37	putative protein (keysa) mRNA.
<a href="#">keyshee</a>	<a href="#">keyshee.aSep08</a>		641	453		38	putative protein (keyshee) mRNA.
<a href="#">keytu</a>	<a href="#">keytu.aSep08</a>		4149	405		129	putative protein (keytu) mRNA.
<a href="#">keyvar</a>	<a href="#">keyvar.aSep08</a>		5332	559	4	185	microtubule-actin crosslinking factor 1 (keyvar) alternative variant aSep08, mRNA.
<a href="#">keywey</a>	<a href="#">keywey.aSep08</a>		3420	327		54	putative protein (keywey) mRNA.
<a href="#">Khdrbs1</a>	<a href="#">Khdrbs1.aSep08</a>	<a href="#">117268</a>	19948	1036	5	273	putative protein of metazoan origin (Khdrbs1) alternative variant aSep08, mRNA.
<a href="#">Khdrbs1</a>	<a href="#">Khdrbs1.bSep08</a>	<a href="#">117268</a>	8943	846	3	212	putative protein of metazoan origin (Khdrbs1) alternative variant bSep08, mRNA.
<a href="#">Khdrbs1</a>	<a href="#">Khdrbs1.cSep08</a>	<a href="#">117268</a>	13150	1383	2	51	putative protein of vertebrate origin (Khdrbs1) alternative variant cSep08, mRNA.
<a href="#">Khdrbs1</a>	<a href="#">Khdrbs1.dSep08</a>	<a href="#">117268</a>	23009	446	4	45	putative protein of vertebrate origin (Khdrbs1) alternative variant dSep08, mRNA.
<a href="#">Khdrbs3</a>	<a href="#">Khdrbs3.bSep08</a>	<a href="#">64015</a>	67798	788	6	124	putative protein of vertebrate origin (Khdrbs3) alternative variant bSep08, mRNA.
<a href="#">Khdrbs3</a>	<a href="#">Khdrbs3.cSep08</a>	<a href="#">64015</a>	60175	598	4	103	putative protein of vertebrate origin (Khdrbs3) alternative variant cSep08, mRNA.
<a href="#">Khk</a>	<a href="#">Khk.bSep08</a>	<a href="#">25659</a>	8350	682	5	162	ketohehexokinase (Khk) alternative variant bSep08, mRNA.
<a href="#">Khk</a>	<a href="#">Khk.dSep08</a>	<a href="#">25659</a>	3162	770	5	150	ketohehexokinase (16.2 kD) (Khk) alternative variant dSep08, mRNA.
<a href="#">Khk</a>	<a href="#">Khk.fSep08</a>	<a href="#">25659</a>	4239	1288	2	70	ketohehexokinase (7.7 kD) (Khk) alternative variant fSep08, mRNA.
<a href="#">Khsrp</a>	<a href="#">Khsrp.bSep08</a>	<a href="#">171137</a>	2888	1782	6	405	KH-type splicing regulatory protein (Khsrp) alternative variant bSep08, mRNA.
<a href="#">Khsrp</a>	<a href="#">Khsrp.cSep08</a>	<a href="#">171137</a>	1189	593	3	32	KH-type splicing regulatory protein (Khsrp) alternative variant cSep08, mRNA.
<a href="#">KH_1.3</a>	<a href="#">KH_1.3.aSep08</a>		9265	455		151	neuro-oncological ventral antigen like (KH_1.3) mRNA.
<a href="#">Kiaa0415</a>	<a href="#">Kiaa0415.aSep08</a>	<a href="#">641386</a>	1662	1549		110	KIAA0415 protein (Kiaa0415) mRNA.
<a href="#">Kidins220</a>	<a href="#">Kidins220.bSep08</a>	<a href="#">116478</a>	18161	3636	5	592	kinase D-interacting (Kidins220) alternative variant bSep08, mRNA.
<a href="#">Kidins220</a>	<a href="#">Kidins220.cSep08</a>	<a href="#">116478</a>	20201	913	7	280	kinase D-interacting (Kidins220) alternative variant cSep08, mRNA.
<a href="#">Kidins220</a>	<a href="#">Kidins220.dSep08</a>	<a href="#">116478</a>	12098	1256	4	246	putative protein, with a coiled coil domain, of vertebrate origin (Kidins220) alternative variant dSep08, mRNA.
<a href="#">Kidins220</a>	<a href="#">Kidins220.eSep08</a>	<a href="#">116478</a>	23795	1093	6	208	kinase D-interacting (Kidins220) alternative variant eSep08, mRNA.
<a href="#">Kidins220</a>	<a href="#">Kidins220.fSep08</a>	<a href="#">116478</a>	5360	548	4	110	kinase D-interacting (Kidins220) alternative variant fSep08, mRNA.
<a href="#">Kif1a</a>	<a href="#">Kif1a.aSep08</a>	<a href="#">363288</a>	23855	2573	2	706	kinesin family member 1A (Kif1a) alternative variant aSep08, mRNA.
<a href="#">Kif1a</a>	<a href="#">Kif1a.bSep08</a>	<a href="#">363288</a>	1640	573	1	70	kinesin family member 1A (Kif1a) alternative variant bSep08, mRNA.
<a href="#">Kif1b</a>	<a href="#">Kif1b.bSep08</a>	<a href="#">117548</a>	10521	1028	7	221	kinesin family member 1B (Kif1b) alternative variant bSep08, mRNA.

<a href="#">Kif1b</a>	<a href="#">Kif1b.cSep08</a>	<a href="#">117548</a>	6765	599	3	199	kinesin family member 1B (Kif1b) alternative variant cSep08, mRNA.
<a href="#">Kif1b</a>	<a href="#">Kif1b.dSep08</a>	<a href="#">117548</a>	40953	506	5	109	kinesin family member 1B (Kif1b) alternative variant dSep08, mRNA.
<a href="#">Kif1b</a>	<a href="#">Kif1b.eSep08</a>	<a href="#">117548</a>	40864	444	4	85	kinesin family member 1B (Kif1b) alternative variant eSep08, mRNA.
<a href="#">Kif1b</a>	<a href="#">Kif1b.fSep08</a>	<a href="#">117548</a>	1853	410	2	36	kinesin family member 1B (Kif1b) alternative variant fSep08, mRNA.
<a href="#">Kif1c</a>	<a href="#">Kif1c.bSep08</a>	<a href="#">113886</a>	29436	1789	6	574	kinesin family member 1C (Kif1c) alternative variant bSep08, mRNA.
<a href="#">Kif2a</a>	<a href="#">Kif2a.aSep08</a>	<a href="#">84391</a>	17286	3283	11	400	kinesin family member 2A (Kif2a) alternative variant aSep08, mRNA.
<a href="#">Kif2a</a>	<a href="#">Kif2a.cSep08</a>	<a href="#">84391</a>	2859	456	2	34	kinesin family member 2A (3.7 kD) (Kif2a) alternative variant cSep08, mRNA.
<a href="#">Kif2c</a>	<a href="#">Kif2c.cSep08</a>	<a href="#">171529</a>	3959	757	4	115	kinesin family member 2C (Kif2c) alternative variant cSep08, mRNA.
<a href="#">Kif2c</a>	<a href="#">Kif2c.dSep08</a>	<a href="#">171529</a>	3868	831	2	110	kinesin family member 2C (Kif2c) alternative variant dSep08, mRNA.
<a href="#">Kif2c</a>	<a href="#">Kif2c.eSep08</a>	<a href="#">171529</a>	14829	763	4	45	kinesin family member 2C (Kif2c) alternative variant eSep08, mRNA.
<a href="#">Kif3a</a>	<a href="#">Kif3a.aSep08</a>	<a href="#">84392</a>	19571	1345	1	448	kinesin family member 3a (Kif3a) alternative variant aSep08, mRNA.
<a href="#">Kif3a</a>	<a href="#">Kif3a.bSep08</a>	<a href="#">84392</a>	12231	692	1	169	kinesin family member 3a (Kif3a) alternative variant bSep08, mRNA.
<a href="#">Kif3b</a>	<a href="#">Kif3b.bSep08</a>	<a href="#">296284</a>	22267	780	1	197	kinesin family member 3B (Kif3b) alternative variant bSep08, mRNA.
<a href="#">Kif3b</a>	<a href="#">Kif3b.cSep08</a>	<a href="#">296284</a>	4828	499	1	118	kinesin family member 3B (Kif3b) alternative variant cSep08, mRNA.
<a href="#">Kif3c</a>	<a href="#">Kif3c.bSep08</a>	<a href="#">85248</a>	15677	2005	6	239	kinesin family member 3C (28.0 kD) (Kif3c) alternative variant bSep08, mRNA.
<a href="#">Kif4</a>	<a href="#">Kif4.aSep08</a>	<a href="#">84393</a>	44425	2466	2	517	kinesin family member 4 (Kif4) alternative variant aSep08, mRNA.
<a href="#">Kif4</a>	<a href="#">Kif4.bSep08</a>	<a href="#">84393</a>	30723	762	2	231	kinesin family member 4 (Kif4) alternative variant bSep08, mRNA.
<a href="#">Kif5a</a>	<a href="#">Kif5a.bSep08</a>	<a href="#">314906</a>	767	388	3	95	kinesin family member 5A (Kif5a) alternative variant bSep08, mRNA.
<a href="#">Kif5b</a>	<a href="#">Kif5b.bSep08</a>	<a href="#">117550</a>	13763	1785	8	482	kinesin family member 5B (Kif5b) alternative variant bSep08, mRNA.
<a href="#">Kif5b</a>	<a href="#">Kif5b.cSep08</a>	<a href="#">117550</a>	11363	3521	7	195	kinesin family member 5B (Kif5b) alternative variant cSep08, mRNA.
<a href="#">Kif5b</a>	<a href="#">Kif5b.dSep08</a>	<a href="#">117550</a>	3150	797	6	96	kinesin family member 5B (Kif5b) alternative variant dSep08, mRNA.
<a href="#">Kif5c</a>	<a href="#">Kif5c.bSep08</a>	<a href="#">311024</a>	26422	4097	4	142	kinesin family member 5C (Kif5c) alternative variant bSep08, mRNA.
<a href="#">Kif5c</a>	<a href="#">Kif5c.cSep08</a>	<a href="#">311024</a>	2974	529	2	130	kinesin family member 5C (Kif5c) alternative variant cSep08, mRNA.

<a href="#">Kif5c</a>	<a href="#">Kif5c.dSep08</a>	<a href="#">311024</a>	24414	443	3	102	kinesin family member 5C (Kif5c) alternative variant dSep08, mRNA.
<a href="#">Kif6</a>	<a href="#">Kif6.aSep08</a>	<a href="#">171291</a>	36654	387		128	kinesin family member 6 (Kif6) mRNA.
<a href="#">Kif7</a>	<a href="#">Kif7.aSep08</a>	<a href="#">293047</a>	1417	1165		158	kinesin family member 7 (Kif7) mRNA.
<a href="#">Kif9</a>	<a href="#">Kif9.aSep08</a>	<a href="#">501059</a>	14329	527		175	kinesin family member 9 (Kif9) mRNA.
<a href="#">Kif11</a>	<a href="#">Kif11.aSep08</a>	<a href="#">171304</a>	11452	2535	7	388	kinesin family member 11 (Kif11) alternative variant aSep08, mRNA.
<a href="#">Kif12</a>	<a href="#">Kif12.bSep08</a>	<a href="#">313254</a>	1240	438	4	139	kinesin family member 12 (Kif12) alternative variant bSep08, mRNA.
<a href="#">Kif12</a>	<a href="#">Kif12.cSep08</a>	<a href="#">313254</a>	974	796	2	77	kinesin family member 12 (Kif12) alternative variant cSep08, mRNA.
<a href="#">Kif12</a>	<a href="#">Kif12.dSep08</a>	<a href="#">313254</a>	739	501	2	72	kinesin family member 12 (7.6 kD) (Kif12) alternative variant dSep08, mRNA.
<a href="#">Kif13a</a>	<a href="#">Kif13a.bSep08</a>	<a href="#">308173</a>	13399	859	7	286	kinesin family member 13A (Kif13a) alternative variant bSep08, mRNA.
<a href="#">Kif13a</a>	<a href="#">Kif13a.cSep08</a>	<a href="#">308173</a>	13459	2316	3	275	kinesin family member 13A (Kif13a) alternative variant cSep08, mRNA.
<a href="#">Kif13a</a>	<a href="#">Kif13a.dSep08</a>	<a href="#">308173</a>	3630	408	3	136	kinesin family member 13A (Kif13a) alternative variant dSep08, mRNA.
<a href="#">Kif14</a>	<a href="#">Kif14.bSep08</a>	<a href="#">360849</a>	4819	439	2	146	kinesin family member 14 (Kif14) alternative variant bSep08, mRNA.
<a href="#">Kif14</a>	<a href="#">Kif14.cSep08</a>	<a href="#">360849</a>	5451	401	4	133	kinesin family member 14 (Kif14) alternative variant cSep08, mRNA.
<a href="#">Kif16b</a>	<a href="#">Kif16b.bSep08</a>	<a href="#">311478</a>	49697	1793	8	579	kinesin family member 16B (Kif16b) alternative variant bSep08, mRNA.
<a href="#">Kif16b</a>	<a href="#">Kif16b.cSep08</a>	<a href="#">311478</a>	5590	378	4	125	kinesin family member 16B (Kif16b) alternative variant cSep08, mRNA.
<a href="#">Kif18a</a>	<a href="#">Kif18a.aSep08</a>	<a href="#">362186</a>	26298	537		159	kinesin family member 18A (Kif18a) mRNA.
<a href="#">Kif18b</a>	<a href="#">Kif18b.bSep08</a>	<a href="#">303575</a>	10120	771	4	256	kinesin family member 18B (Kif18b) alternative variant bSep08, mRNA.
<a href="#">Kif18b</a>	<a href="#">Kif18b.cSep08</a>	<a href="#">303575</a>	11001	1402	4	121	kinesin family member 18B (Kif18b) alternative variant cSep08, mRNA.
<a href="#">Kif19a</a>	<a href="#">Kif19a.aSep08</a>	<a href="#">303659</a>	1556	416	4	138	kinesin family member 19A (Kif19a) alternative variant aSep08, mRNA.
<a href="#">Kif19a</a>	<a href="#">Kif19a.bSep08</a>	<a href="#">303659</a>	1544	300	2	100	kinesin family member 19A (Kif19a) alternative variant bSep08, mRNA.
<a href="#">Kif21a</a>	<a href="#">Kif21a.bSep08</a>	<a href="#">300158</a>	15867	801	6	267	kinesin family member 21A (Kif21a) alternative variant bSep08, mRNA.
<a href="#">Kif21a</a>	<a href="#">Kif21a.cSep08</a>	<a href="#">300158</a>	10883	1233	7	246	kinesin family member 21A (Kif21a) alternative variant cSep08, mRNA.
<a href="#">Kif21a</a>	<a href="#">Kif21a.dSep08</a>	<a href="#">300158</a>	7589	727	5	154	kinesin family member 21A (Kif21a) alternative variant dSep08, mRNA.
<a href="#">Kif21a</a>	<a href="#">Kif21a.eSep08</a>	<a href="#">300158</a>	10848	525	4	147	kinesin family member 21A (Kif21a) alternative variant eSep08, mRNA.
<a href="#">Kif21b</a>	<a href="#">Kif21b.bSep08</a>	<a href="#">289397</a>	10312	2175	13	708	kinesin family member 21B (Kif21b) alternative variant bSep08, mRNA.

<a href="#">Kif21b</a>	<a href="#">Kif21b.cSep08</a>	<a href="#">289397</a>	8352	801	8	266	kinesin family member 21B (Kif21b) alternative variant cSep08, mRNA.
<a href="#">Kif22</a>	<a href="#">Kif22.bSep08</a>	<a href="#">293502</a>	628	449	3	97	kinesin family member 22 (11.0 kD) (Kif22) alternative variant bSep08, mRNA.
<a href="#">Kif22</a>	<a href="#">Kif22.cSep08</a>	<a href="#">293502</a>	841	724	2	67	kinesin family member 22 (Kif22) alternative variant cSep08, mRNA.
<a href="#">Kif23</a>	<a href="#">Kif23.bSep08</a>	<a href="#">315740</a>	4706	682	4	227	kinesin family member 23 (Kif23) alternative variant bSep08, mRNA.
<a href="#">Kif23</a>	<a href="#">Kif23.cSep08</a>	<a href="#">315740</a>	2577	1208	3	70	kinesin family member 23 (Kif23) alternative variant cSep08, mRNA.
<a href="#">Kifap3</a>	<a href="#">Kifap3.bSep08</a>	<a href="#">289168</a>	25802	693	6	230	kinesin-associated protein 3 (Kifap3) alternative variant bSep08, mRNA.
<a href="#">Kifap3</a>	<a href="#">Kifap3.cSep08</a>	<a href="#">289168</a>	55203	1004	6	203	kinesin-associated protein 3 (Kifap3) alternative variant cSep08, mRNA.
<a href="#">Kifap3</a>	<a href="#">Kifap3.dSep08</a>	<a href="#">289168</a>	15481	935	6	198	kinesin-associated protein 3 (Kifap3) alternative variant dSep08, mRNA.
<a href="#">Kifap3</a>	<a href="#">Kifap3.eSep08</a>	<a href="#">289168</a>	6188	379	3	126	kinesin-associated protein 3 (Kifap3) alternative variant eSep08, mRNA.
<a href="#">Kifap3</a>	<a href="#">Kifap3.fSep08</a>	<a href="#">289168</a>	26613	405	2	72	kinesin-associated protein 3 (8.0 kD) (Kifap3) alternative variant fSep08, mRNA.
<a href="#">Kifc1</a>	<a href="#">Kifc1.bSep08</a>	<a href="#">294286</a>	9545	1929	8	589	kinesin family member C1 (Kifc1) alternative variant bSep08, mRNA.
<a href="#">KIFC2</a>	<a href="#">KIFC2.bSep08</a>	<a href="#">300053</a>	3285	1331	3	328	kinesin family member C2 (36.0 kD) (KIFC2) alternative variant bSep08, mRNA.
<a href="#">KIFC2</a>	<a href="#">KIFC2.cSep08</a>	<a href="#">300053</a>	4507	1641	9	208	kinesin family member C2 (KIFC2) alternative variant cSep08, mRNA.
<a href="#">KIFC2</a>	<a href="#">KIFC2.dSep08</a>	<a href="#">300053</a>	471	385	2	128	kinesin family member C2 CRA c (KIFC2) alternative variant dSep08, mRNA.
<a href="#">KIFC2</a>	<a href="#">KIFC2.eSep08</a>	<a href="#">300053</a>	1909	850	2	86	kinesin family member C2 (8.9 kD) (KIFC2) alternative variant eSep08, mRNA.
<a href="#">Kifc3</a>	<a href="#">Kifc3.bSep08</a>	<a href="#">307644</a>	1664	704	3	184	kinesin family member C3 (Kifc3) alternative variant bSep08, mRNA.
<a href="#">Kifc3</a>	<a href="#">Kifc3.cSep08</a>	<a href="#">307644</a>	23081	609	5	170	kinesin family member C3 (Kifc3) alternative variant cSep08, mRNA.
<a href="#">Kifc3</a>	<a href="#">Kifc3.dSep08</a>	<a href="#">307644</a>	23726	385	4	84	kinesin family member C3 (Kifc3) alternative variant dSep08, mRNA.
<a href="#">Kinesin.0</a>	<a href="#">Kinesin.0.aSep08</a>		10735	1025		341	kinesin family member 1A CRA b (Kinesin.0) mRNA.
<a href="#">Kinesin.1</a>	<a href="#">Kinesin.1.aSep08</a>		644	513		170	kinesin family member 7 (Kinesin.1) mRNA.
<a href="#">Kirrel3</a>	<a href="#">Kirrel3.bSep08</a>	<a href="#">315546</a>	28112	2017	12	553	kin of irre like 3 (Kirrel3) alternative variant bSep08, mRNA.
<a href="#">Kirrel3</a>	<a href="#">Kirrel3.cSep08</a>	<a href="#">315546</a>	13244	794	6	171	kin of irre like 3 CRA b (Kirrel3) alternative variant cSep08, mRNA.
<a href="#">Kirrel3</a>	<a href="#">Kirrel3.dSep08</a>	<a href="#">315546</a>	42829	474	4	128	kin of irre like 3 (Kirrel3) alternative variant dSep08, mRNA.
<a href="#">Kitl</a>	<a href="#">Kitl.bSep08</a>	<a href="#">60427</a>	20101	5190	3	91	kit ligand (10.0 kD) (Kitl) alternative variant bSep08, mRNA.
<a href="#">klabor</a>	<a href="#">klabor.aSep08</a>		2194	1800		131	putative protein of mammalian origin (13.8 kD) (klabor) mRNA.



<a href="#">klachy</a>	<a href="#">klachy.aSep08</a>		1700	364		86	putative protein (klachy) mRNA.
<a href="#">kladoy</a>	<a href="#">kladoy.aSep08</a>		2142	1530		214	protein SMF (kladoy) mRNA.
<a href="#">klafee</a>	<a href="#">klafee.aSep08</a>		2077	382		37	putative protein (klafee) mRNA.
<a href="#">klaflu</a>	<a href="#">klaflu.aSep08</a>		4646	1589		529	partner localizer of BRCA2 (klaflu) mRNA.
<a href="#">klafly</a>	<a href="#">klafly.aSep08</a>		5545	276		24	putative protein (2.8 kD) (klafly) mRNA.
<a href="#">klagar</a>	<a href="#">klagar.aSep08</a>		9722	672		57	putative protein (klagar) mRNA.
<a href="#">klaja</a>	<a href="#">klaja.aSep08</a>		16524	478		27	putative protein (2.9 kD) (klaja) mRNA.
<a href="#">klajey</a>	<a href="#">klajey.aSep08</a>		5589	1167		143	putative protein (16.0 kD) (klajey) mRNA.
<a href="#">klalo</a>	<a href="#">klalo.aSep08</a>		5196	379		126	putative protein of vertebrate origin (klalo) mRNA.
<a href="#">klamee</a>	<a href="#">klamee.aSep08</a>		2076	389		65	putative protein (7.5 kD) (klamee) mRNA.
<a href="#">klapey</a>	<a href="#">klapey.aSep08</a>		30509	410		51	putative protein (klapey) mRNA.
<a href="#">klapor</a>	<a href="#">klapor.aSep08</a>		14255	714	3	238	transcription complex CRA b (klapor) alternative variant aSep08, mRNA.
<a href="#">klapor</a>	<a href="#">klapor.bSep08</a>		7702	416	1	95	transcription complex CRA b (klapor) alternative variant bSep08, mRNA.
<a href="#">klarbor</a>	<a href="#">klarbor.aSep08</a>		803	288		56	putative protein (klarbor) mRNA.
<a href="#">klarchy</a>	<a href="#">klarchy.aSep08</a>		45479	655		24	putative protein (2.8 kD) (klarchy) mRNA.
<a href="#">klardoy</a>	<a href="#">klardoy.aSep08</a>		4358	631		53	putative protein (5.6 kD) (klardoy) mRNA.
<a href="#">klarfee</a>	<a href="#">klarfee.aSep08</a>		19045	498		33	putative protein (3.5 kD) (klarfee) mRNA.
<a href="#">klarflu</a>	<a href="#">klarflu.aSep08</a>		2646	674		68	putative protein (7.4 kD) (klarflu) mRNA.
<a href="#">klarfly</a>	<a href="#">klarfly.aSep08</a>		2251	621	3	81	myeloid lymphoid mixed-lineage leukemia like (klarfly) mRNA.
<a href="#">klargar</a>	<a href="#">klargar.aSep08</a>		3699	272		70	putative protein (klargar) mRNA.
<a href="#">klarja</a>	<a href="#">klarja.aSep08</a>		1308	398		31	putative protein (3.7 kD) (klarja) mRNA.
<a href="#">klarjey</a>	<a href="#">klarjey.aSep08</a>		48238	963		72	putative protein (klarjey) mRNA.
<a href="#">klarlo</a>	<a href="#">klarlo.aSep08</a>		58333	419		43	putative protein (klarlo) mRNA.
<a href="#">klarmee</a>	<a href="#">klarmee.aSep08</a>		4704	1501	2	159	CRA b like (17.4 kD) (klarmee) alternative variant aSep08, mRNA.
<a href="#">klarmee</a>	<a href="#">klarmee.bSep08</a>		4541	1770	2	67	putative protein (klarmee) alternative variant bSep08, mRNA.
<a href="#">klarmee</a>	<a href="#">klarmee.cSep08</a>		3577	1655	1	60	putative protein (7.3 kD) (klarmee) alternative variant cSep08, mRNA.
<a href="#">klarmee</a>	<a href="#">klarmee.dSep08</a>		4064	548	2	47	putative protein (klarmee) alternative variant dSep08, mRNA.
<a href="#">klaroy</a>	<a href="#">klaroy.aSep08</a>		1063	189		62	putative protein (klaroy) mRNA.
<a href="#">klarpey</a>	<a href="#">klarpey.aSep08</a>		4687	639		213	dynein 2 heavy (klarpey) mRNA.
<a href="#">klarpor</a>	<a href="#">klarpor.aSep08</a>		1077	495		57	putative protein (6.8 kD) (klarpor) mRNA.
<a href="#">klarroy</a>	<a href="#">klarroy.aSep08</a>		1734	760		56	putative protein (6.1 kD) (klarroy) mRNA.
<a href="#">klarshee</a>	<a href="#">klarshee.aSep08</a>		2717	358		119	synaptopodin 2 (klarshee) mRNA.
<a href="#">klartu</a>	<a href="#">klartu.aSep08</a>		34515	337		46	synaptotagmin XVI like (klartu) mRNA.
<a href="#">klarvo</a>	<a href="#">klarvo.aSep08</a>		813	276		58	putative protein (klarvo) mRNA.
<a href="#">klarwer</a>	<a href="#">klarwer.aSep08</a>		7084	427		69	putative protein of mammalian origin (7.8 kD) (klarwer) mRNA.

<a href="#">klashee</a>	<a href="#">klashee.aSep08</a>		21245	1035		8	putative protein (9.8 kD) (klashee) mRNA.
<a href="#">klatu</a>	<a href="#">klatu.aSep08</a>		24165	414		49	putative protein (5.4 kD) (klatu) mRNA.
<a href="#">klavo</a>	<a href="#">klavo.aSep08</a>		14115	267		33	putative protein (klavo) mRNA.
<a href="#">klawbor</a>	<a href="#">klawbor.aSep08</a>		3048	282		43	putative protein (klawbor) mRNA.
<a href="#">klawchy</a>	<a href="#">klawchy.aSep08</a>		3005	553		184	nuclear receptor coactivator 6 (klawchy) mRNA.
<a href="#">klawdoy</a>	<a href="#">klawdoy.aSep08</a>		3154	2462		110	putative protein (12.5 kD) (klawdoy) mRNA.
<a href="#">klawer</a>	<a href="#">klawer.aSep08</a>		38494	2733		50	putative protein (klawer) mRNA.
<a href="#">klawfee</a>	<a href="#">klawfee.aSep08</a>		3858	1263		72	ab2-143 like (8.0 kD) (klawfee) mRNA.
<a href="#">klawflu</a>	<a href="#">klawflu.aSep08</a>		6765	479	3	106	putative protein (11.6 kD) (klawflu) alternative variant aSep08, mRNA.
<a href="#">klawfly</a>	<a href="#">klawfly.aSep08</a>		24077	748		69	putative protein (7.7 kD) (klawfly) mRNA.
<a href="#">klawgar</a>	<a href="#">klawgar.aSep08</a>		5024	891		72	putative protein (klawgar) mRNA.
<a href="#">klawja</a>	<a href="#">klawja.aSep08</a>		5625	275		52	diacylglycerol kinase eta like (klawja) mRNA.
<a href="#">klawjey</a>	<a href="#">klawjey.aSep08</a>		12361	583		103	putative protein (klawjey) mRNA.
<a href="#">klawlo</a>	<a href="#">klawlo.aSep08</a>		1561	208		62	putative protein (klawlo) mRNA.
<a href="#">klawmee</a>	<a href="#">klawmee.aSep08</a>		3271	747		32	putative protein (3.5 kD) (klawmee) mRNA.
<a href="#">klawpey</a>	<a href="#">klawpey.aSep08</a>		5951	593		197	dynein cytoplasmic 2 heavy (klawpey) mRNA.
<a href="#">klawpor</a>	<a href="#">klawpor.aSep08</a>		650	452		97	putative protein (klawpor) mRNA.
<a href="#">klawroy</a>	<a href="#">klawroy.aSep08</a>		17775	314		104	regulatory factor X 4 CRA a (klawroy) mRNA.
<a href="#">klawshee</a>	<a href="#">klawshee.bSep08</a>		680	523	2	35	CRA b like (4.1 kD) (klawshee) alternative variant bSep08, mRNA.
<a href="#">klawshee</a>	<a href="#">klawshee.cSep08</a>		767	358	3	45	CRA a like (klawshee) alternative variant cSep08, mRNA.
<a href="#">klawtu</a>	<a href="#">klawtu.aSep08</a>		18839	1313		83	putative protein (klawtu) alternative variant aSep08, mRNA.
<a href="#">klawvo</a>	<a href="#">klawvo.aSep08</a>		48017	750		106	putative nuclear protein (11.9 kD) (klawvo) mRNA.
<a href="#">klawwer</a>	<a href="#">klawwer.aSep08</a>		1434	722	3	53	putative protein (klawwer) alternative variant aSep08, mRNA.
<a href="#">Klc1</a>	<a href="#">Klc1.dSep08</a>	<a href="#">171041</a>	18783	788	5	130	kinesin light chain 1 (Klc1) alternative variant dSep08, mRNA.
<a href="#">Klc1</a>	<a href="#">Klc1.eSep08</a>	<a href="#">171041</a>	8742	913	5	115	kinesin light chain 1 (Klc1) alternative variant eSep08, mRNA.
<a href="#">Klc1</a>	<a href="#">Klc1.fSep08</a>	<a href="#">171041</a>	7197	724	4	52	kinesin light chain 1 (Klc1) alternative variant fSep08, mRNA.
<a href="#">Klc1</a>	<a href="#">Klc1.hSep08</a>	<a href="#">171041</a>	1461	876	2	48	kinesin light chain 1 (Klc1) alternative variant hSep08, mRNA.
<a href="#">Klc1</a>	<a href="#">Klc1.jSep08</a>	<a href="#">171041</a>	5589	619	2	59	kinesin light chain 1 (6.1 kD) (Klc1) alternative variant jSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.aSep08</a>	<a href="#">309159</a>	9947	2920	16	692	kinesin light CRA b (Klc2) alternative variant aSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.cSep08</a>	<a href="#">309159</a>	4814	723	5	240	kinesin light CRA b (Klc2) alternative variant cSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.dSep08</a>	<a href="#">309159</a>	4102	654	4	218	kinesin light CRA b (Klc2) alternative variant dSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.eSep08</a>	<a href="#">309159</a>	4025	653	3	153	kinesin light CRA b (Klc2) alternative variant eSep08, mRNA.

<a href="#">Klc2</a>	<a href="#">Klc2.fSep08</a>	<a href="#">309159</a>	1748	864	5	148	kinesin light CRA b (15.5 kD) (Klc2) alternative variant fSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.gSep08</a>	<a href="#">309159</a>	1089	426	2	87	putative protein (Klc2) alternative variant gSep08, mRNA.
<a href="#">Klc2</a>	<a href="#">Klc2.iSep08</a>	<a href="#">309159</a>	973	290	2	47	putative protein (Klc2) alternative variant iSep08, mRNA.
<a href="#">Klc3</a>	<a href="#">Klc3.bSep08</a>	<a href="#">171549</a>	426	265	2	40	kinesin light chain 3 (Klc3) alternative variant bSep08, mRNA.
<a href="#">Klc4</a>	<a href="#">Klc4.bSep08</a>	<a href="#">316226</a>	4980	706	5	190	kinesin light chain 4 (Klc4) alternative variant bSep08, mRNA.
<a href="#">Klc4</a>	<a href="#">Klc4.cSep08</a>	<a href="#">316226</a>	1391	818	4	146	kinesin light chain 4 (Klc4) alternative variant cSep08, mRNA.
<a href="#">Klc4</a>	<a href="#">Klc4.dSep08</a>	<a href="#">316226</a>	2542	617	3	119	kinesin light chain 4 (Klc4) alternative variant dSep08, mRNA.
<a href="#">Klc4</a>	<a href="#">Klc4.fSep08</a>	<a href="#">316226</a>	1278	591	4	58	kinesin light chain 4 (Klc4) alternative variant fSep08, mRNA.
<a href="#">kleebor</a>	<a href="#">kleebor.aSep08</a>		2183	399		70	putative protein (7.2 kD) (kleebor) mRNA.
<a href="#">kleechy</a>	<a href="#">kleechy.aSep08</a>		5991	429		143	nuclear receptor coactivator 6 (kleechy) mRNA.
<a href="#">kleedoy</a>	<a href="#">kleedoy.aSep08</a>		7403	400		57	putative protein (kleedoy) mRNA.
<a href="#">kleefee</a>	<a href="#">kleefee.aSep08</a>		9826	512	3	145	cdc14 cell division cycle 14 homolog B CRA b (kleefee) alternative variant aSep08, mRNA.
<a href="#">kleefee</a>	<a href="#">kleefee.bSep08</a>		3817	282	1	68	cdc14 cell division cycle 14 homolog B CRA b (kleefee) alternative variant bSep08, mRNA.
<a href="#">kleeflu</a>	<a href="#">kleeflu.aSep08</a>		2263	585		63	putative protein (kleeflu) mRNA.
<a href="#">kleefly</a>	<a href="#">kleefly.aSep08</a>		12617	419		66	putative protein (kleefly) mRNA.
<a href="#">kleegar</a>	<a href="#">kleegar.aSep08</a>		4664	764		66	tumor necrosis factor superfamily member 13b like (kleegar) mRNA.
<a href="#">kleeja</a>	<a href="#">kleeja.aSep08</a>		15575	751		250	putative protein of eukaryotic origin (kleeja) mRNA.
<a href="#">kleejey</a>	<a href="#">kleejey.aSep08</a>		4697	542		105	putative protein, with a coiled coil domain (kleejey) mRNA.
<a href="#">kleelo</a>	<a href="#">kleelo.aSep08</a>		6437	392		83	putative protein (kleelo) mRNA.
<a href="#">kleemee</a>	<a href="#">kleemee.aSep08</a>		770	474	2	54	putative protein (kleemee) alternative variant aSep08, mRNA.
<a href="#">kleemee</a>	<a href="#">kleemee.bSep08</a>		3631	425	3	39	putative protein (4.5 kD) (kleemee) alternative variant bSep08, mRNA.
<a href="#">kleepey</a>	<a href="#">kleepey.aSep08</a>		26173	478		101	putative protein (kleepey) mRNA.
<a href="#">kleepor</a>	<a href="#">kleepor.aSep08</a>		206352	473		94	RNA binding motif single stranded interacting protein like (kleepor) mRNA.
<a href="#">kleeroy</a>	<a href="#">kleeroy.aSep08</a>		19878	588		195	polymerase III polypeptide B CRA b (kleeroy) mRNA.
<a href="#">kleeshaw</a>	<a href="#">kleeshaw.aSep08</a>		11539	664		61	putative protein (kleeshaw) mRNA.
<a href="#">kleeshee</a>	<a href="#">kleeshee.aSep08</a>		3522	377	1	33	LYST-interacting protein 8 like (kleeshee) mRNA.
<a href="#">kleetu</a>	<a href="#">kleetu.aSep08</a>		2935	837		43	putative protein (4.9 kD) (kleetu) mRNA.
<a href="#">kleevo</a>	<a href="#">kleevo.aSep08</a>		3205	747		78	putative protein of mammalian origin (kleevo) mRNA.
<a href="#">kleewer</a>	<a href="#">kleewer.aSep08</a>		1425	977		60	putative protein (6.8 kD) (kleewer) mRNA.
<a href="#">klerbor</a>	<a href="#">klerbor.aSep08</a>		573	298		49	putative protein (klerbor) mRNA.
<a href="#">klerchy</a>	<a href="#">klerchy.aSep08</a>		25826	447		45	putative protein (klerchy) mRNA.
<a href="#">klerdoy</a>	<a href="#">klerdoy.aSep08</a>		18504	625		203	SH3 domain tetratricopeptide repeats 2 (klerdoy) mRNA.

<a href="#">klerflu</a>	<a href="#">klerflu.aSep08</a>		38069	807	2	102	putative cytoplasmic protein (11.1 kD) (klerflu) alternative variant aSep08, mRNA.
<a href="#">klerflu</a>	<a href="#">klerflu.bSep08</a>		28392	872	3	102	putative cytoplasmic protein (11.1 kD) (klerflu) alternative variant bSep08, mRNA.
<a href="#">klerfly</a>	<a href="#">klerfly.aSep08</a>		1038	926		50	putative protein (5.5 kD) (klerfly) mRNA.
<a href="#">klergar</a>	<a href="#">klergar.aSep08</a>		637	500		71	putative protein (klergar) mRNA.
<a href="#">klerja</a>	<a href="#">klerja.aSep08</a>		10854	673		223	putative protein of metazoan origin (klerja) mRNA.
<a href="#">klerjey</a>	<a href="#">klerjey.aSep08</a>		16960	705		51	putative protein (5.6 kD) (klerjey) mRNA.
<a href="#">klerlo</a>	<a href="#">klerlo.aSep08</a>		2597	337		104	putative protein (klerlo) mRNA.
<a href="#">klermee</a>	<a href="#">klermee.aSep08</a>		2573	1191		91	putative protein (klermee) mRNA.
<a href="#">klerpey</a>	<a href="#">klerpey.aSep08</a>		6122	170		56	putative protein of mammalian origin (klerpey) mRNA.
<a href="#">klerpor</a>	<a href="#">klerpor.aSep08</a>		9727	347	4	38	putative protein (4.3 kD) (klerpor) alternative variant aSep08, mRNA.
<a href="#">klerpor</a>	<a href="#">klerpor.bSep08</a>		20316	342	3	76	putative protein (klerpor) alternative variant bSep08, mRNA.
<a href="#">klerpor</a>	<a href="#">klerpor.cSep08</a>		7666	315	3	38	putative protein (4.3 kD) (klerpor) alternative variant cSep08, mRNA.
<a href="#">klerroy</a>	<a href="#">klerroy.aSep08</a>		4762	348		111	polymerase III polypeptide B CRA b (klerroy) mRNA.
<a href="#">klershaw</a>	<a href="#">klershaw.bSep08</a>		2694	640	3	61	putative protein (6.9 kD) (klershaw) alternative variant bSep08, mRNA.
<a href="#">klershee</a>	<a href="#">klershee.aSep08</a>		21838	439		44	putative protein (klershee) mRNA.
<a href="#">klertu</a>	<a href="#">klertu.bSep08</a>		13244	396	2	72	putative protein (klertu) alternative variant bSep08, mRNA.
<a href="#">klervo</a>	<a href="#">klervo.aSep08</a>		4548	778		28	putative protein (klervo) mRNA.
<a href="#">klerwer</a>	<a href="#">klerwer.aSep08</a>		1239	438		69	putative protein (klerwer) mRNA.
<a href="#">kleybor</a>	<a href="#">kleybor.aSep08</a>		5751	731		70	putative protein (kleybor) mRNA.
<a href="#">kleychy</a>	<a href="#">kleychy.aSep08</a>		6376	853		67	putative protein (kleychy) mRNA.
<a href="#">kleydoy</a>	<a href="#">kleydoy.aSep08</a>		1668	378		99	putative protein (kleydoy) mRNA.
<a href="#">kleyfee</a>	<a href="#">kleyfee.aSep08</a>		95937	421		139	aminopeptidase O (kleyfee) mRNA.
<a href="#">kleyflu</a>	<a href="#">kleyflu.aSep08</a>		2174	681		111	putative protein (11.5 kD) (kleyflu) mRNA.
<a href="#">kleyfly</a>	<a href="#">kleyfly.bSep08</a>		5825	971	2	72	ab2-143 like (8.0 kD) (kleyfly) alternative variant bSep08, mRNA.
<a href="#">kleygar</a>	<a href="#">kleygar.aSep08</a>		144902	682		64	putative protein (7.4 kD) (kleygar) mRNA.
<a href="#">kleyja</a>	<a href="#">kleyja.aSep08</a>		6026	393		48	putative protein (kleyja) mRNA.
<a href="#">kleyjey</a>	<a href="#">kleyjey.aSep08</a>		3048	657		44	putative protein (kleyjey) mRNA.
<a href="#">kleylo</a>	<a href="#">kleylo.aSep08</a>		687	304		60	putative protein (6.9 kD) (kleylo) mRNA.
<a href="#">kleymee</a>	<a href="#">kleymee.aSep08</a>		963	258		85	family with sequence similarity 57 member A like (kleymee) mRNA.
<a href="#">kleypey</a>	<a href="#">kleypey.aSep08</a>		1722	692		103	putative protein (kleypey) mRNA.
<a href="#">kleypor</a>	<a href="#">kleypor.aSep08</a>		13792	480	1	45	putative protein (4.7 kD) (kleypor) alternative variant aSep08, mRNA.
<a href="#">kleypor</a>	<a href="#">kleypor.bSep08</a>		2026	307	1	50	putative protein (kleypor) alternative variant bSep08, mRNA.
<a href="#">kleyroy</a>	<a href="#">kleyroy.aSep08</a>		3868	531		48	putative protein (5.3 kD) (kleyroy) mRNA.

<a href="#">kleyshaw</a>	<a href="#">kleyshaw.aSep08</a>		10911	440		51	putative protein (kleyshaw) mRNA.
<a href="#">kleyshee</a>	<a href="#">kleyshee.aSep08</a>		393	262		51	la ribonucleoprotein domain family member 7 (kleyshee) mRNA.
<a href="#">kleytu</a>	<a href="#">kleytu.aSep08</a>		2002	571		88	putative protein (kleytu) mRNA.
<a href="#">kleyvo</a>	<a href="#">kleyvo.aSep08</a>		1124	378		76	fanconi anemia complementation group G (kleyvo) mRNA.
<a href="#">kleywer</a>	<a href="#">kleywer.aSep08</a>		1222	837			
<a href="#">Klf3</a>	<a href="#">Klf3.bSep08</a>	<a href="#">114845</a>	10506	608	3	119	kruppel-like factor 3 (Klf3) alternative variant bSep08, mRNA.
<a href="#">Klf3</a>	<a href="#">Klf3.cSep08</a>	<a href="#">114845</a>	3545	2565	2	101	putative mitochondrial protein (11.8 kD) (Klf3) alternative variant cSep08, mRNA.
<a href="#">Klf5</a>	<a href="#">Klf5.bSep08</a>	<a href="#">84410</a>	7247	853	2	222	kruppel-like factor 5 (Klf5) alternative variant bSep08, mRNA.
<a href="#">Klf8</a>	<a href="#">Klf8.aSep08</a>	<a href="#">302582</a>	2980	379		125	kruppel-like factor 8 (Klf8) mRNA.
<a href="#">Klf12</a>	<a href="#">Klf12.bSep08</a>	<a href="#">306110</a>	58869	624	4	54	kruppel-like factor 12 (Klf12) alternative variant bSep08, mRNA.
<a href="#">Klf12</a>	<a href="#">Klf12.cSep08</a>	<a href="#">306110</a>	4539	557	3	65	kruppel-like factor 12 (Klf12) alternative variant cSep08, mRNA.
<a href="#">Klf13</a>	<a href="#">Klf13.bSep08</a>	<a href="#">499171</a>	42686	471	1	111	kruppel-like factor 13 (Klf13) alternative variant bSep08, mRNA.
<a href="#">Klf15</a>	<a href="#">Klf15.bSep08</a>	<a href="#">85497</a>	4729	983	2	249	kruppel-like factor 15 (Klf15) alternative variant bSep08, mRNA.
<a href="#">Klf15</a>	<a href="#">Klf15.dSep08</a>	<a href="#">85497</a>	12494	971	3	214	kruppel-like factor 15 (Klf15) alternative variant dSep08, mRNA.
<a href="#">Klf15</a>	<a href="#">Klf15.eSep08</a>	<a href="#">85497</a>	1602	668	2	160	kruppel-like factor 15 (Klf15) alternative variant eSep08, mRNA.
<a href="#">Klhdc1</a>	<a href="#">Klhdc1.bSep08</a>	<a href="#">314190</a>	54483	2460	14	311	kelch repeat containing protein and kelch (Klhdc1) alternative variant bSep08, mRNA.
<a href="#">Klhdc1</a>	<a href="#">Klhdc1.cSep08</a>	<a href="#">314190</a>	27939	767	5	208	kelch repeat containing protein (Klhdc1) alternative variant cSep08, mRNA.
<a href="#">Klhdc1</a>	<a href="#">Klhdc1.eSep08</a>	<a href="#">314190</a>	833	335	2	73	putative protein of vertebrate origin (Klhdc1) alternative variant eSep08, mRNA.
<a href="#">Klhdc2</a>	<a href="#">Klhdc2.bSep08</a>	<a href="#">299113</a>	3919	2056	2	70	putative protein (7.8 kD) (Klhdc2) alternative variant bSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.bSep08</a>	<a href="#">363192</a>	1653	869	6	266	kelch repeat containing protein and kelch (Klhdc3) alternative variant bSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.cSep08</a>	<a href="#">363192</a>	3464	773	6	209	kelch repeat containing protein and kelch (Klhdc3) alternative variant cSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.dSep08</a>	<a href="#">363192</a>	1848	746	6	197	kelch repeat containing protein and kelch (Klhdc3) alternative variant dSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.eSep08</a>	<a href="#">363192</a>	5501	2935	10	196	kelch repeat containing protein and kelch (22.3 kD) (Klhdc3) alternative variant eSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.fSep08</a>	<a href="#">363192</a>	5539	875	6	179	kelch repeat containing protein and kelch (20.0 kD) (Klhdc3) alternative variant fSep08, mRNA.
<a href="#">Klhdc3</a>	<a href="#">Klhdc3.hSep08</a>	<a href="#">363192</a>	3190	438	4	103	kelch repeat containing protein and kelch (Klhdc3) alternative variant hSep08, mRNA.

<a href="#">Klhdc3</a>	<a href="#">Klhdc3.jSep08</a>	<a href="#">363192</a>	4123	1625	7	115	kelch repeat containing protein and kelch (Klhdc3) alternative variant jSep08, mRNA.
<a href="#">Klhdc8a</a>	<a href="#">Klhdc8a.aSep08</a>	<a href="#">305096</a>	4631	1297	2	252	kelch repeat containing protein and kelch (Klhdc8a) alternative variant aSep08, mRNA.
<a href="#">Klhdc8a</a>	<a href="#">Klhdc8a.bSep08</a>	<a href="#">305096</a>	3717	1796	4	216	kelch repeat containing protein and kelch (24.2 kD) (Klhdc8a) alternative variant bSep08, mRNA.
<a href="#">Klhdc8a</a>	<a href="#">Klhdc8a.cSep08</a>	<a href="#">305096</a>	3575	528	3	95	kelch repeat containing protein (Klhdc8a) alternative variant cSep08, mRNA.
<a href="#">Klhdc9</a>	<a href="#">Klhdc9.bSep08</a>	<a href="#">360878</a>	2528	661	4	183	kelch containing (Klhdc9) alternative variant bSep08, mRNA.
<a href="#">Klh12</a>	<a href="#">Klh12.aSep08</a>	<a href="#">290692</a>	40087	2496	11	467	kelch-like 2, Mayven (Drosophila) (Klh12) alternative variant aSep08, mRNA.
<a href="#">Klh12</a>	<a href="#">Klh12.cSep08</a>	<a href="#">290692</a>	2432	507	2	53	kelch-like 2, Mayven (Drosophila) (Klh12) alternative variant cSep08, mRNA.
<a href="#">Klh15</a>	<a href="#">Klh15.bSep08</a>	<a href="#">305351</a>	9502	1322	4	135	kelch-like 5 (Drosophila) (Klh15) alternative variant bSep08, mRNA.
<a href="#">Klh16</a>	<a href="#">Klh16.bSep08</a>	<a href="#">287974</a>	6219	2127	2	206	kelch-like 6 (Drosophila) (22.5 kD) (Klh16) alternative variant bSep08, mRNA.
<a href="#">Klh18</a>	<a href="#">Klh18.bSep08</a>	<a href="#">289457</a>	2169	367	2	76	putative protein (Klh18) alternative variant bSep08, mRNA.
<a href="#">Klh18</a>	<a href="#">Klh18.dSep08</a>	<a href="#">289457</a>	20837	669	4	76	putative protein (Klh18) alternative variant dSep08, mRNA.
<a href="#">Klh12</a>	<a href="#">Klh12.bSep08</a>	<a href="#">266772</a>	11171	827	3	108	kelch-like 12 (Drosophila) (Klh12) alternative variant bSep08, mRNA.
<a href="#">Klh12</a>	<a href="#">Klh12.cSep08</a>	<a href="#">266772</a>	3390	481	2	83	kelch-like 12 (Drosophila) (Klh12) alternative variant cSep08, mRNA.
<a href="#">Klh13</a>	<a href="#">Klh13.aSep08</a>	<a href="#">313445</a>	136072	457		78	kelch-like 13 (Drosophila) (Klh13) mRNA.
<a href="#">Klh15</a>	<a href="#">Klh15.cSep08</a>	<a href="#">314111</a>	1433	588	2	195	kelch-like 15 (Drosophila) (Klh15) alternative variant cSep08, mRNA.
<a href="#">Klh18</a>	<a href="#">Klh18.aSep08</a>	<a href="#">316012</a>	41287	506	3	168	kelch-like 18 (Drosophila) (Klh18) alternative variant aSep08, mRNA.
<a href="#">Klh18</a>	<a href="#">Klh18.bSep08</a>	<a href="#">316012</a>	3303	883	2		
<a href="#">Klh122</a>	<a href="#">Klh122.bSep08</a>	<a href="#">303792</a>	20230	736	4	245	kelch-like 22 (Drosophila) (Klh122) alternative variant bSep08, mRNA.
<a href="#">Klh122</a>	<a href="#">Klh122.cSep08</a>	<a href="#">303792</a>	15957	643	4	125	kelch-like 22 (Drosophila) (Klh122) alternative variant cSep08, mRNA.
<a href="#">Klh123</a>	<a href="#">Klh123.bSep08</a>	<a href="#">311114</a>	2343	1208	2	166	kelch-like 23 (Drosophila) (Klh123) alternative variant bSep08, mRNA.
<a href="#">Klh124</a>	<a href="#">Klh124.aSep08</a>	<a href="#">303803</a>	24661	1754		535	DRE1 (Klh124) alternative variant aSep08, mRNA.
<a href="#">Klh124</a>	<a href="#">Klh124.bSep08</a>	<a href="#">303803</a>	10809	779	1	65	putative protein (Klh124) alternative variant bSep08, mRNA.
<a href="#">Klh125</a>	<a href="#">Klh125.bSep08</a>	<a href="#">293023</a>	19310	1650	1	550	kelch-like 25 (Drosophila) (Klh125) alternative variant bSep08, mRNA.
<a href="#">Klh129</a>	<a href="#">Klh129.bSep08</a>	<a href="#">298867</a>	172214	1562	6	335	kelch-like 29 (Drosophila) (Klh129) alternative variant bSep08, mRNA.
<a href="#">Klk1c10</a>	<a href="#">Klk1c10.aSep08</a>	<a href="#">292858</a>	1980	670		207	T-kininogenase (Klk1c10) mRNA.
<a href="#">Klk6</a>	<a href="#">Klk6.aSep08</a>	<a href="#">29245</a>	6473	1778	1	467	kallikrein related-peptidase 6 (Klk6) alternative variant aSep08, mRNA.

<a href="#">klobor</a>	<a href="#">klobor.aSep08</a>		2712	284		65	putative protein (7.3 kD) (klobor) alternative variant aSep08, mRNA.
<a href="#">klochy</a>	<a href="#">klochy.aSep08</a>		23806	504		39	putative protein (4.4 kD) (klochy) mRNA.
<a href="#">klodoy</a>	<a href="#">klodoy.aSep08</a>		5005	501		148	SMF protein like (klodoy) mRNA.
<a href="#">klofee</a>	<a href="#">klofee.aSep08</a>		3944	802		32	putative protein (3.8 kD) (klofee) mRNA.
<a href="#">kloflu</a>	<a href="#">kloflu.aSep08</a>		7071	383		127	trinucleotide repeat containing 6A (kloflu) mRNA.
<a href="#">klofly</a>	<a href="#">klofly.aSep08</a>		1038	926		50	putative protein (5.5 kD) (klofly) mRNA.
<a href="#">klogar</a>	<a href="#">klogar.aSep08</a>		3674	313	2	43	CRA a like (5.1 kD) (klogar) alternative variant aSep08, mRNA.
<a href="#">klogar</a>	<a href="#">klogar.bSep08</a>		3576	410	3	55	CRA a like (klogar) alternative variant bSep08, mRNA.
<a href="#">kloja</a>	<a href="#">kloja.aSep08</a>		34666	641		59	putative protein, with a coiled coil domain (6.9 kD) (kloja) mRNA.
<a href="#">klojey</a>	<a href="#">klojey.aSep08</a>		679	495		24	putative protein (2.6 kD) (klojey) mRNA.
<a href="#">klolo</a>	<a href="#">klolo.aSep08</a>		13707	432		96	putative protein (klolo) mRNA.
<a href="#">klomee</a>	<a href="#">klomee.aSep08</a>		2641	1648			
<a href="#">klopey</a>	<a href="#">klopey.aSep08</a>		21162	1404		467	dynein 2 heavy (klopey) mRNA.
<a href="#">klopor</a>	<a href="#">klopor.aSep08</a>		5542	994		118	oxysterol-binding protein-like protein 10 (klopor) mRNA.
<a href="#">klorbor</a>	<a href="#">klorbor.aSep08</a>		4606	1007		335	putative protein of vertebrate origin (klorbor) alternative variant aSep08, mRNA.
<a href="#">klorbor</a>	<a href="#">klorbor.bSep08</a>		4886	926		308	putative protein of vertebrate origin (klorbor) alternative variant bSep08, mRNA.
<a href="#">klorchy</a>	<a href="#">klorchy.aSep08</a>		961	308		31	CRA b like (klorchy) mRNA.
<a href="#">klordoy</a>	<a href="#">klordoy.bSep08</a>		24261	440	3	48	putative protein (5.7 kD) (klordoy) alternative variant bSep08, mRNA.
<a href="#">klorfee</a>	<a href="#">klorfee.aSep08</a>		10338	516		32	putative protein (3.6 kD) (klorfee) mRNA.
<a href="#">klorflu</a>	<a href="#">klorflu.aSep08</a>		569	386		128	CBP activator (klorflu) mRNA.
<a href="#">klorfly</a>	<a href="#">klorfly.aSep08</a>		1794	322		63	putative protein (klorfly) mRNA.
<a href="#">klogar</a>	<a href="#">klogar.aSep08</a>		7432	1163		317	putative protein (klogar) alternative variant aSep08, mRNA.
<a href="#">klogar</a>	<a href="#">klogar.bSep08</a>		6796	526		61	putative protein (6.7 kD) (klogar) alternative variant bSep08, mRNA.
<a href="#">klorja</a>	<a href="#">klorja.aSep08</a>		772	346		63	putative protein (klorja) mRNA.
<a href="#">klorjey</a>	<a href="#">klorjey.aSep08</a>		28848	655		40	putative protein (4.7 kD) (klorjey) mRNA.
<a href="#">klorlo</a>	<a href="#">klorlo.aSep08</a>		7342	381		47	putative protein (klorlo) mRNA.
<a href="#">klormee</a>	<a href="#">klormee.aSep08</a>		4343	722		240	gem associated protein 4 CRA b (klormee) mRNA.
<a href="#">kloroy</a>	<a href="#">kloroy.aSep08</a>		864	208		14	putative protein (kloroy) mRNA.
<a href="#">klorpey</a>	<a href="#">klorpey.bSep08</a>		1546	354	2	57	putative protein (6.7 kD) (klorpey) alternative variant bSep08, mRNA.
<a href="#">klorpor</a>	<a href="#">klorpor.aSep08</a>		5228	306		41	putative protein (klorpor) mRNA.
<a href="#">klorroy</a>	<a href="#">klorroy.aSep08</a>		4303	288		28	putative protein (3.4 kD) (klorroy) mRNA.
<a href="#">klorshaw</a>	<a href="#">klorshaw.aSep08</a>		6997	426	3	128	multiple C2 domains transmembrane 1 (klorshaw) alternative variant aSep08, mRNA.
<a href="#">klorshaw</a>	<a href="#">klorshaw.bSep08</a>		1904	742	1	36	putative protein (4.2 kD) (klorshaw) alternative variant bSep08, mRNA.

<a href="#">klorshee</a>	<a href="#">klorshee.aSep08</a>		1218	403		134	prematurely terminated mRNA decay factor-like (klorshee) mRNA.
<a href="#">klortu</a>	<a href="#">klortu.aSep08</a>		29214	428	5	83	spectrin repeat containing Nuclear envelope 2 (klortu) alternative variant aSep08, mRNA.
<a href="#">klorvo</a>	<a href="#">klorvo.aSep08</a>		2221	544	3	181	phosphatidylinositol glycan class O (klorvo) alternative variant aSep08, mRNA.
<a href="#">klorvo</a>	<a href="#">klorvo.bSep08</a>		1067	415	1	112	phosphatidylinositol glycan class O (klorvo) alternative variant bSep08, mRNA.
<a href="#">klorwer</a>	<a href="#">klorwer.aSep08</a>		16815	909		302	protein CRA c (klorwer) mRNA.
<a href="#">kloshee</a>	<a href="#">kloshee.aSep08</a>		3428	411		53	putative protein (kloshee) mRNA.
<a href="#">klotu</a>	<a href="#">klotu.aSep08</a>		3073	651		57	putative protein (klotu) mRNA.
<a href="#">klovo</a>	<a href="#">klovo.aSep08</a>		1279	738		46	putative protein (5.1 kD) (klovo) mRNA.
<a href="#">klower</a>	<a href="#">klower.aSep08</a>		11967	412	4	137	thrombospondin type-1 domain-containing protein 7A (klower) alternative variant aSep08, mRNA.
<a href="#">kloybor</a>	<a href="#">kloybor.aSep08</a>		685	411		137	putative protein of vertebrate origin (kloybor) mRNA.
<a href="#">kloychy</a>	<a href="#">kloychy.aSep08</a>		4657	564		82	putative protein (kloychy) mRNA.
<a href="#">kloydoy</a>	<a href="#">kloydoy.aSep08</a>		3090	533		177	CRA a (kloydoy) mRNA.
<a href="#">kloyfee</a>	<a href="#">kloyfee.aSep08</a>		1397	472		61	cathepsin 7 CRA b (kloyfee) mRNA.
<a href="#">kloyflu</a>	<a href="#">kloyflu.aSep08</a>		932	407		135	CBP activator (kloyflu) mRNA.
<a href="#">kloyfly</a>	<a href="#">kloyfly.aSep08</a>		11948	758	1	105	putative protein (11.5 kD) (kloyfly) alternative variant aSep08, mRNA.
<a href="#">kloyfly</a>	<a href="#">kloyfly.bSep08</a>		62891	788	1	104	putative protein (11.5 kD) (kloyfly) alternative variant bSep08, mRNA.
<a href="#">kloygar</a>	<a href="#">kloygar.aSep08</a>		893	725		84	putative protein (10.0 kD) (kloygar) mRNA.
<a href="#">kloyja</a>	<a href="#">kloyja.aSep08</a>		18413	685		228	diaphanous homolog 3 CRA b (kloyja) mRNA.
<a href="#">kloyjey</a>	<a href="#">kloyjey.bSep08</a>		712	473	2	129	putative protein (kloyjey) alternative variant bSep08, mRNA.
<a href="#">kloylo</a>	<a href="#">kloylo.aSep08</a>		624	299		90	putative protein (kloylo) mRNA.
<a href="#">kloymee</a>	<a href="#">kloymee.aSep08</a>		16987	698		232	putative protein of eukaryotic origin (kloymee) alternative variant aSep08, mRNA.
<a href="#">kloymee</a>	<a href="#">kloymee.bSep08</a>		5680	370		123	putative protein of ancient origin (kloymee) alternative variant bSep08, mRNA.
<a href="#">kloypey</a>	<a href="#">kloypey.aSep08</a>		19297	411		136	gtpase-activating protein 10-like (kloypey) mRNA.
<a href="#">kloypor</a>	<a href="#">kloypor.aSep08</a>		6932	389		25	putative protein (kloypor) mRNA.
<a href="#">kloyroy</a>	<a href="#">kloyroy.aSep08</a>		1985	444		55	putative protein (kloyroy) mRNA.
<a href="#">kloyshaw</a>	<a href="#">kloyshaw.aSep08</a>		14197	386		81	putative protein (kloyshaw) mRNA.
<a href="#">kloyshee</a>	<a href="#">kloyshee.aSep08</a>		427	245		23	putative protein (kloyshee) mRNA.
<a href="#">kloytu</a>	<a href="#">kloytu.aSep08</a>		14363	953		317	CRA a (kloytu) mRNA.
<a href="#">kloyvo</a>	<a href="#">kloyvo.aSep08</a>		1067	478		158	talín (kloyvo) mRNA.
<a href="#">kloywer</a>	<a href="#">kloywer.aSep08</a>		1692	825		274	protein CRA b (kloywer) mRNA.
<a href="#">Klra2</a>	<a href="#">Klra2.bSep08</a>	<a href="#">494194</a>	37003	3551	2	258	killer cell lectin-like receptor, subfamily A, member 2 (30.1 kD) (Klra2) alternative variant bSep08, mRNA.
<a href="#">Klre1</a>	<a href="#">Klre1.bSep08</a>	<a href="#">297645</a>	2370	585	1	151	killer cell lectin-like receptor, family E, member 1 (Klre1) alternative variant bSep08, mRNA.



<a href="#">Klre1</a>	<a href="#">Klre1.cSep08</a>	<a href="#">297645</a>	8779	811	4	99	killer cell lectin-like receptor, family E, member 1 (11.4 kD) (Klre1) alternative variant cSep08, mRNA.
<a href="#">Klrg1</a>	<a href="#">Klrg1.bSep08</a>	<a href="#">58975</a>	11230	545	1	77	killer cell lectin-like receptor subfamily G, member 1 (8.7 kD) (Klrg1) alternative variant bSep08, mRNA.
<a href="#">klubor</a>	<a href="#">klubor.aSep08</a>		2763	333		51	putative protein (klubor) mRNA.
<a href="#">kluchy</a>	<a href="#">kluchy.aSep08</a>		6723	667		194	core-binding factor runt domain alpha translocated 2 CRA a like (kluchy) mRNA.
<a href="#">kludoy</a>	<a href="#">kludoy.aSep08</a>		12289	690	3	230	protein SMF (kludoy) alternative variant aSep08, mRNA.
<a href="#">kludoy</a>	<a href="#">kludoy.bSep08</a>		7091	408	1	88	protein SMF (kludoy) alternative variant bSep08, mRNA.
<a href="#">klufee</a>	<a href="#">klufee.aSep08</a>		2161	701		48	putative protein (klufee) mRNA.
<a href="#">klufly</a>	<a href="#">klufly.aSep08</a>		1744	526		175	retinoblastoma binding protein 6 like (klufly) mRNA.
<a href="#">klufly</a>	<a href="#">klufly.aSep08</a>		9293	886		69	putative protein (klufly) mRNA.
<a href="#">klugar</a>	<a href="#">klugar.aSep08</a>		1177	794		72	putative protein (klugar) mRNA.
<a href="#">kluja</a>	<a href="#">kluja.aSep08</a>		41308	683		100	putative cytoplasmic protein (11.3 kD) (kluja) mRNA.
<a href="#">klujey</a>	<a href="#">klujey.aSep08</a>		1458	269		54	putative protein (klujey) mRNA.
<a href="#">klulo</a>	<a href="#">klulo.aSep08</a>		5508	850		283	putative protein of metazoan origin (klulo) mRNA.
<a href="#">klumee</a>	<a href="#">klumee.bSep08</a>		6023	235	2	28	putative protein (klumee) alternative variant bSep08, mRNA.
<a href="#">klupey</a>	<a href="#">klupey.aSep08</a>		46547	1309		281	dynein 2 heavy (klupey) mRNA.
<a href="#">klupor</a>	<a href="#">klupor.aSep08</a>		2799	378		48	putative protein (5.5 kD) (klupor) mRNA.
<a href="#">kluroy</a>	<a href="#">kluroy.aSep08</a>		4557	485		77	putative protein, with a coiled coil domain (8.6 kD) (kluroy) mRNA.
<a href="#">klushee</a>	<a href="#">klushee.aSep08</a>		11271	633		95	putative cytoplasmic protein (11.0 kD) (klushee) mRNA.
<a href="#">klutu</a>	<a href="#">klutu.aSep08</a>		480	324		57	putative protein (klutu) mRNA.
<a href="#">kluvo</a>	<a href="#">kluvo.aSep08</a>		841	531	2	84	CRA b like (kluvo) alternative variant aSep08, mRNA.
<a href="#">kluvo</a>	<a href="#">kluvo.bSep08</a>		632	286	2	70	putative protein (kluvo) alternative variant bSep08, mRNA.
<a href="#">kluwer</a>	<a href="#">kluwer.aSep08</a>		7130	1575	1	432	WD repeat-containing protein mio (kluwer) alternative variant aSep08, mRNA.
<a href="#">kluwer</a>	<a href="#">kluwer.bSep08</a>		5723	398	1	84	WD repeat-containing protein mio like (kluwer) alternative variant bSep08, mRNA.
<a href="#">klybor</a>	<a href="#">klybor.aSep08</a>		905	370		28	putative protein (klybor) mRNA.
<a href="#">klychy</a>	<a href="#">klychy.aSep08</a>		2065	421		140	core-binding factor runt domain alpha translocated 2 homolog CRA b like (klychy) mRNA.
<a href="#">klydoy</a>	<a href="#">klydoy.aSep08</a>		14417	1165		388	SMF protein (klydoy) mRNA.
<a href="#">klyfee</a>	<a href="#">klyfee.aSep08</a>		572	391		45	putative protein (klyfee) mRNA.
<a href="#">klyflu</a>	<a href="#">klyflu.aSep08</a>		4191	391		129	CRA d like (klyflu) mRNA.
<a href="#">klyfly</a>	<a href="#">klyfly.aSep08</a>		9830	886	1	69	putative protein (klyfly) alternative variant aSep08, mRNA.
<a href="#">klyfly</a>	<a href="#">klyfly.bSep08</a>		8982	785	1	122	putative protein (klyfly) alternative variant bSep08, mRNA.
<a href="#">klygar</a>	<a href="#">klygar.aSep08</a>		7188	351	2	78	CRA a like (klygar) alternative variant aSep08, mRNA.
<a href="#">klygar</a>	<a href="#">klygar.bSep08</a>		2030	609	3	42	putative protein (5.1 kD) (klygar) alternative variant bSep08, mRNA.
<a href="#">klyja</a>	<a href="#">klyja.aSep08</a>		4592	813		70	putative protein (7.8 kD) (klyja) mRNA.
<a href="#">klyjey</a>	<a href="#">klyjey.aSep08</a>		27151	751		155	putative protein (17.1 kD) (klyjey) mRNA.

<a href="#">klylo</a>	<a href="#">klylo.aSep08</a>		1016	404		134	putative protein of vertebrate origin (klylo) mRNA.
<a href="#">klymee</a>	<a href="#">klymee.aSep08</a>		2395	779		259	tsr1 20S rRNA accumulation (klymee) mRNA.
<a href="#">klypey</a>	<a href="#">klypey.aSep08</a>		3178	295		46	putative protein (klypey) mRNA.
<a href="#">klypor</a>	<a href="#">klypor.aSep08</a>		2248	1393		102	putative protein (11.5 kD) (klypor) mRNA.
<a href="#">klyroy</a>	<a href="#">klyroy.aSep08</a>		852	208		14	putative protein (klyroy) mRNA.
<a href="#">klyshee</a>	<a href="#">klyshee.aSep08</a>		8136	510		70	putative protein (klyshee) mRNA.
<a href="#">klytu</a>	<a href="#">klytu.aSep08</a>		28130	588		58	putative protein (klytu) mRNA.
<a href="#">klyvo</a>	<a href="#">klyvo.aSep08</a>		2330	289		81	dead box polypeptide (klyvo) mRNA.
<a href="#">klywer</a>	<a href="#">klywer.aSep08</a>		1665	398		132	collagen type XXVIII (klywer) mRNA.
<a href="#">Kmo</a>	<a href="#">Kmo.bSep08</a>	<a href="#">59113</a>	23779	906	9	249	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) (Kmo) alternative variant bSep08, mRNA.
<a href="#">Kmo</a>	<a href="#">Kmo.cSep08</a>	<a href="#">59113</a>	23686	743	8	247	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) (Kmo) alternative variant cSep08, mRNA.
<a href="#">Kntc1</a>	<a href="#">Kntc1.bSep08</a>	<a href="#">304477</a>	2173	699	4	162	kinetochore associated 1 (Kntc1) alternative variant bSep08, mRNA.
<a href="#">Kntc1</a>	<a href="#">Kntc1.cSep08</a>	<a href="#">304477</a>	639	327	2	69	kinetochore associated 1 (Kntc1) alternative variant cSep08, mRNA.
<a href="#">Kntc1</a>	<a href="#">Kntc1.dSep08</a>	<a href="#">304477</a>	2715	561	3	65	kinetochore associated 1 (Kntc1) alternative variant dSep08, mRNA.
<a href="#">koby</a>	<a href="#">koby.aSep08</a>		79944	376		116	putative protein (koby) mRNA.
<a href="#">kochy</a>	<a href="#">kochy.aSep08</a>		8474	734		53	putative protein (6.1 kD) (kochy) mRNA.
<a href="#">kodar</a>	<a href="#">kodar.aSep08</a>		1071	322		106	intestine specific homeobox like (kodar) mRNA.
<a href="#">kofer</a>	<a href="#">kofer.aSep08</a>		12074	960		319	supervillin CRA b (kofer) mRNA.
<a href="#">koflo</a>	<a href="#">koflo.aSep08</a>		37227	451		149	cytoplasmic polyadenylation element binding protein 3 like (koflo) mRNA.
<a href="#">koflu</a>	<a href="#">koflu.aSep08</a>		5721	327		72	putative protein (koflu) mRNA.
<a href="#">kojey</a>	<a href="#">kojey.aSep08</a>		1317	418		139	phosphatidylinositol glycan class G CRA c (kojey) mRNA.
<a href="#">kokee</a>	<a href="#">kokee.aSep08</a>		1375	687		190	protein tyrosine phosphatase (kokee) mRNA.
<a href="#">koloy</a>	<a href="#">koloy.aSep08</a>		9756	1435		478	specific peptidase 5 (koloy) mRNA.
<a href="#">komee</a>	<a href="#">komee.aSep08</a>		1033	447		38	putative protein (komee) mRNA.
<a href="#">komer</a>	<a href="#">komer.bSep08</a>		1919	594	3	53	putative protein (6.0 kD) (komer) alternative variant bSep08, mRNA.
<a href="#">konoy</a>	<a href="#">konoy.aSep08</a>		14811	390		129	heavy axonemal (konoy) mRNA.
<a href="#">kopor</a>	<a href="#">kopor.aSep08</a>		4887	594		101	putative protein (kopor) mRNA.
<a href="#">korby</a>	<a href="#">korby.aSep08</a>		3697	526		175	putative protein of metazoan origin (korby) mRNA.
<a href="#">korchy</a>	<a href="#">korchy.aSep08</a>		1337	949			
<a href="#">kordar</a>	<a href="#">kordar.aSep08</a>		5112	793		90	putative protein (9.8 kD) (kordar) mRNA.
<a href="#">korfer</a>	<a href="#">korfer.aSep08</a>		4545	515	4	171	enhancer of polycomb 1 (korfer) alternative variant aSep08, mRNA.
<a href="#">korflo</a>	<a href="#">korflo.aSep08</a>		2881	460		152	cytochrome (korflo) mRNA.
<a href="#">korflu</a>	<a href="#">korflu.aSep08</a>		5592	346		114	hermansky-Pudlak syndrome 5 like (korflu) mRNA.
<a href="#">korjey</a>	<a href="#">korjey.aSep08</a>		8239	527		174	ecotropic viral integration site 5 (korjey) mRNA.
<a href="#">korkee</a>	<a href="#">korkee.aSep08</a>		7890	496		43	putative protein (4.9 kD) (korkee) mRNA.

<a href="#">korloy</a>	<a href="#">korloy.aSep08</a>		4089	606		75	putative endoplasmic reticulum protein (8.4 kD) (korloy) mRNA.
<a href="#">kormee</a>	<a href="#">kormee.aSep08</a>		504	299		38	putative protein (kormee) mRNA.
<a href="#">kormer</a>	<a href="#">kormer.aSep08</a>		2515	746		107	putative cytoplasmic protein (12.3 kD) (kormer) mRNA.
<a href="#">kornoy</a>	<a href="#">kornoy.aSep08</a>		49619	424		86	putative protein (kornoy) mRNA.
<a href="#">korpor</a>	<a href="#">korpor.aSep08</a>		4512	575		191	promyelocytic leukemia like (korpor) mRNA.
<a href="#">korsa</a>	<a href="#">korsa.aSep08</a>		4582	722		21	putative protein (2.2 kD) (korsa) mRNA.
<a href="#">korshee</a>	<a href="#">korshee.aSep08</a>		39974	361	3	86	putative protein (korshee) alternative variant aSep08, mRNA.
<a href="#">korshee</a>	<a href="#">korshee.bSep08</a>		4004	679	2	59	putative protein (6.5 kD) (korshee) alternative variant bSep08, mRNA.
<a href="#">kortu</a>	<a href="#">kortu.aSep08</a>		6812	342		62	putative protein (kortu) alternative variant aSep08, mRNA.
<a href="#">kortu</a>	<a href="#">kortu.bSep08</a>		6823	407	1	54	putative protein (5.9 kD) (kortu) alternative variant bSep08, mRNA.
<a href="#">korvar</a>	<a href="#">korvar.aSep08</a>		908	449		149	microtubule-actin crosslinking factor 1 (korvar) mRNA.
<a href="#">korwey</a>	<a href="#">korwey.aSep08</a>		3858	336		112	leiomodoin 3 (korwey) mRNA.
<a href="#">kosa</a>	<a href="#">kosa.aSep08</a>		4374	695		94	putative protein of bilateral origin (kosa) mRNA.
<a href="#">koshee</a>	<a href="#">koshee.aSep08</a>		5016	483		67	putative protein (koshee) mRNA.
<a href="#">kotu</a>	<a href="#">kotu.aSep08</a>		4585	478	5	84	CRA a like (kotu) alternative variant aSep08, mRNA.
<a href="#">kotu</a>	<a href="#">kotu.bSep08</a>		4347	958	5	81	CRA a like (9.5 kD) (kotu) alternative variant bSep08, mRNA.
<a href="#">kotu</a>	<a href="#">kotu.dSep08</a>		2620	427	4	54	CRA a like (kotu) alternative variant dSep08, mRNA.
<a href="#">kovar</a>	<a href="#">kovar.bSep08</a>		2810	857	3	83	putative protein (9.0 kD) (kovar) alternative variant bSep08, mRNA.
<a href="#">kovar</a>	<a href="#">kovar.cSep08</a>		2714	740	3	52	putative protein (6.0 kD) (kovar) alternative variant cSep08, mRNA.
<a href="#">kowey</a>	<a href="#">kowey.aSep08</a>		5935	632		40	putative protein (4.7 kD) (kowey) mRNA.
<a href="#">koyby</a>	<a href="#">koyby.aSep08</a>		3572	831	3	113	putative protein (koyby) alternative variant aSep08, mRNA.
<a href="#">koyby</a>	<a href="#">koyby.bSep08</a>		3532	403	2	42	putative protein (koyby) alternative variant bSep08, mRNA.
<a href="#">koyby</a>	<a href="#">koyby.cSep08</a>		584	324	1	44	putative protein (koyby) alternative variant cSep08, mRNA.
<a href="#">koychy</a>	<a href="#">koychy.aSep08</a>		1243	1079		92	putative protein of mammalian origin (koychy) alternative variant aSep08, mRNA.
<a href="#">koydar</a>	<a href="#">koydar.aSep08</a>		622	512		21	putative protein (2.3 kD) (koydar) mRNA.
<a href="#">koyfer</a>	<a href="#">koyfer.aSep08</a>		2807	2694		78	mohawk (koyfer) mRNA.
<a href="#">koyflo</a>	<a href="#">koyflo.aSep08</a>		1312	967		148	polyprotein (17.1 kD) (koyflo) mRNA.
<a href="#">koyflu</a>	<a href="#">koyflu.aSep08</a>		1996	675		47	putative protein (5.3 kD) (koyflu) mRNA.
<a href="#">koyjey</a>	<a href="#">koyjey.aSep08</a>		20452	669		223	ecotropic viral integration site 5 (koyjey) mRNA.
<a href="#">koykee</a>	<a href="#">koykee.aSep08</a>		2226	758	2	36	putative protein (4.0 kD) (koykee) alternative variant aSep08, mRNA.
<a href="#">koyloy</a>	<a href="#">koyloy.aSep08</a>		1709	739		46	putative protein (5.2 kD) (koyloy) mRNA.
<a href="#">koymee</a>	<a href="#">koymee.aSep08</a>		1664	455		151	polycystic kidney disease 1 like (koymee) mRNA.
<a href="#">koymer</a>	<a href="#">koymer.aSep08</a>		1590	750		105	A member ATP-binding cassette sub-family 8 like (koymer) mRNA.

<a href="#">koynoy</a>	<a href="#">koynoy.aSep08</a>		1013	605	2	16	putative protein (1.9 kD) (koynoy) alternative variant aSep08, mRNA.
<a href="#">koypor</a>	<a href="#">koypor.aSep08</a>		3653	441		146	stomatin 1 (koypor) mRNA.
<a href="#">koysa</a>	<a href="#">koysa.aSep08</a>		17081	500		72	putative secreted or extracellular protein precursor (8.5 kD) (koysa) mRNA.
<a href="#">koysee</a>	<a href="#">koysee.aSep08</a>		10787	749	3	157	putative protein (koysee) alternative variant aSep08, mRNA.
<a href="#">koysee</a>	<a href="#">koysee.dSep08</a>		8819	1147	3	81	putative mitochondrial protein (9.3 kD) (koysee) alternative variant dSep08, mRNA.
<a href="#">koytu</a>	<a href="#">koytu.aSep08</a>		16958	713		21	putative protein (2.5 kD) (koytu) mRNA.
<a href="#">koyvar</a>	<a href="#">koyvar.aSep08</a>		4717	395		131	microtubule-actin crosslinking factor 1 (koyvar) mRNA.
<a href="#">koywey</a>	<a href="#">koywey.aSep08</a>		1802	830		36	putative protein (4.2 kD) (koywey) mRNA.
<a href="#">Kpna2</a>	<a href="#">Kpna2.aSep08</a>	<a href="#">85245</a>	12152	1966	5	529	karyopherin (importin) alpha 2 (57.8 kD) (Kpna2) alternative variant aSep08, complete mRNA.
<a href="#">Kpna2</a>	<a href="#">Kpna2.bSep08</a>	<a href="#">85245</a>	8401	1163	5	350	karyopherin (importin) alpha 2 (Kpna2) alternative variant bSep08, mRNA.
<a href="#">Kpna2</a>	<a href="#">Kpna2.cSep08</a>	<a href="#">85245</a>	1406	790	1	263	karyopherin (importin) alpha 2 (Kpna2) alternative variant cSep08, mRNA.
<a href="#">Kpna2</a>	<a href="#">Kpna2.dSep08</a>	<a href="#">85245</a>	1571	705	1	220	karyopherin (importin) alpha 2 (Kpna2) alternative variant dSep08, mRNA.
<a href="#">Kpna3</a>	<a href="#">Kpna3.bSep08</a>	<a href="#">361055</a>	11423	826	7	210	karyopherin (importin) alpha 3 (Kpna3) alternative variant bSep08, mRNA.
<a href="#">Kpna3</a>	<a href="#">Kpna3.cSep08</a>	<a href="#">361055</a>	9206	701	5	125	karyopherin (importin) alpha 3 (Kpna3) alternative variant cSep08, mRNA.
<a href="#">Kpna3</a>	<a href="#">Kpna3.dSep08</a>	<a href="#">361055</a>	49305	357	6	119	karyopherin (importin) alpha 3 (Kpna3) alternative variant dSep08, mRNA.
<a href="#">Kpna4</a>	<a href="#">Kpna4.bSep08</a>	<a href="#">361959</a>	29144	2278	1	486	karyopherin (importin) alpha 4 (Kpna4) alternative variant bSep08, mRNA.
<a href="#">Kpna6</a>	<a href="#">Kpna6.aSep08</a>	<a href="#">362607</a>	2421	526		175	karyopherin (importin) alpha 6 (Kpna6) mRNA.
<a href="#">Kpnb1</a>	<a href="#">Kpnb1.bSep08</a>	<a href="#">24917</a>	29063	1782	8	392	karyopherin (importin) beta 1 (Kpnb1) alternative variant bSep08, mRNA.
<a href="#">Kpnb1</a>	<a href="#">Kpnb1.cSep08</a>	<a href="#">24917</a>	7753	3675	7	240	karyopherin (importin) beta 1 (Kpnb1) alternative variant cSep08, mRNA.
<a href="#">Kpnb1</a>	<a href="#">Kpnb1.dSep08</a>	<a href="#">24917</a>	5162	657	7	218	karyopherin (importin) beta 1 (Kpnb1) alternative variant dSep08, mRNA.
<a href="#">Kpnb1</a>	<a href="#">Kpnb1.eSep08</a>	<a href="#">24917</a>	3130	683	3	104	karyopherin (importin) beta 1 (11.5 kD) (Kpnb1) alternative variant eSep08, mRNA.
<a href="#">Kptn</a>	<a href="#">Kptn.aSep08</a>	<a href="#">308107</a>	7697	1645	12	430	kaptin CRA a (47.5 kD) (Kptn) alternative variant aSep08, complete mRNA.
<a href="#">Kptn</a>	<a href="#">Kptn.bSep08</a>	<a href="#">308107</a>	3971	738	6	228	kaptin CRA a (Kptn) alternative variant bSep08, mRNA.
<a href="#">Kptn</a>	<a href="#">Kptn.cSep08</a>	<a href="#">308107</a>	3924	947	5	184	kaptin CRA a (Kptn) alternative variant cSep08, mRNA.
<a href="#">Kptn</a>	<a href="#">Kptn.dSep08</a>	<a href="#">308107</a>	1767	666	3	107	kaptin CRA a (Kptn) alternative variant dSep08, mRNA.
<a href="#">Kptn</a>	<a href="#">Kptn.gSep08</a>	<a href="#">308107</a>	3603	414	2	33	putative protein (Kptn) alternative variant gSep08, mRNA.
<a href="#">KRAB.0</a>	<a href="#">KRAB.0.aSep08</a>		1331	657	1	33	CRA b like (3.9 kD) (KRAB.0) alternative variant aSep08, mRNA.

<a href="#">KRAB.0</a>	<a href="#">KRAB.0.bSep08</a>		17877	658	2	42	KRAB box (5.1 kD) (KRAB.0) alternative variant bSep08, mRNA.
<a href="#">KRAB.1</a>	<a href="#">KRAB.1.aSep08</a>		18701	1066		108	KRAB box (KRAB.1) mRNA.
<a href="#">KRAB.2</a>	<a href="#">KRAB.2.aSep08</a>		25006	860	1	93	predicted gene like (11.4 kD) (KRAB.2) alternative variant aSep08, mRNA.
<a href="#">KRAB.2</a>	<a href="#">KRAB.2.bSep08</a>		24673	524	1	92	KRAB box (11.3 kD) (KRAB.2) alternative variant bSep08, mRNA.
<a href="#">KRAB.3</a>	<a href="#">KRAB.3.aSep08</a>		36245	1192		117	KRAB box (13.8 kD) (KRAB.3) mRNA.
<a href="#">KRAB.4</a>	<a href="#">KRAB.4.aSep08</a>		23106	485	2	118	KRAB box (14.1 kD) (KRAB.4) alternative variant aSep08, mRNA.
<a href="#">KRAB.4</a>	<a href="#">KRAB.4.bSep08</a>		85631	627	4	71	putative nuclear protein (8.1 kD) (KRAB.4) alternative variant bSep08, mRNA.
<a href="#">KRAB.5</a>	<a href="#">KRAB.5.aSep08</a>		23119	440		117	KRAB box (13.9 kD) (KRAB.5) mRNA.
<a href="#">KRAB.6</a>	<a href="#">KRAB.6.aSep08</a>		28659	1147		92	KRAB box (11.2 kD) (KRAB.6) mRNA.
<a href="#">KRAB.7</a>	<a href="#">KRAB.7.bSep08</a>		39302	739	3	92	predicted gene like (11.2 kD) (KRAB.7) alternative variant bSep08, mRNA.
<a href="#">KRAB.7</a>	<a href="#">KRAB.7.cSep08</a>		9596	546	2	91	predicted gene like (KRAB.7) alternative variant cSep08, mRNA.
<a href="#">KRAB.7</a>	<a href="#">KRAB.7.dSep08</a>		30224	517	3	89	predicted gene like (10.9 kD) (KRAB.7) alternative variant dSep08, mRNA.
<a href="#">KRAB.7</a>	<a href="#">KRAB.7.eSep08</a>		30087	514	4	58	putative protein of mammalian origin (7.1 kD) (KRAB.7) alternative variant eSep08, complete mRNA.
<a href="#">KRAB.8</a>	<a href="#">KRAB.8.aSep08</a>		20685	2710	2	423	KRAB box and zinc finger, C2H2-type (49.5 kD) (KRAB.8) alternative variant aSep08, mRNA.
<a href="#">KRAB.8</a>	<a href="#">KRAB.8.bSep08</a>		18827	725	1	217	zinc finger, C2H2-type (KRAB.8) alternative variant bSep08, mRNA.
<a href="#">KRAB.8</a>	<a href="#">KRAB.8.cSep08</a>		17246	295	1	43	putative protein (KRAB.8) alternative variant cSep08, mRNA.
<a href="#">KRAB.9</a>	<a href="#">KRAB.9.aSep08</a>		9757	990		330	KRAB box and zinc finger, C2H2-type (KRAB.9) mRNA.
<a href="#">KRAB.10</a>	<a href="#">KRAB.10.aSep08</a>		9971	1465	2	487	KRAB box and zinc finger, C2H2-type (KRAB.10) alternative variant aSep08, mRNA.
<a href="#">KRAB.10</a>	<a href="#">KRAB.10.bSep08</a>		8982	1122	1	102	CRA b (KRAB.10) alternative variant bSep08, mRNA.
<a href="#">KRAB.11</a>	<a href="#">KRAB.11.aSep08</a>		6925	1673		390	KRAB box and zinc finger, C2H2-type (KRAB.11) mRNA.
<a href="#">KRAB.12</a>	<a href="#">KRAB.12.aSep08</a>		27407	587		127	CRA a (KRAB.12) mRNA.
<a href="#">KRAB.13</a>	<a href="#">KRAB.13.aSep08</a>		17810	573		168	KRAB box (KRAB.13) mRNA.
<a href="#">KRAB.14</a>	<a href="#">KRAB.14.aSep08</a>		48771	538		74	CRA b (8.8 kD) (KRAB.14) mRNA.
<a href="#">KRAB.15</a>	<a href="#">KRAB.15.aSep08</a>		14996	1915		417	KRAB box and zinc finger, C2H2-type (48.3 kD) (KRAB.15) mRNA.
<a href="#">KRAB.16</a>	<a href="#">KRAB.16.aSep08</a>		5826	3196	2	463	CRA c (53.8 kD) (KRAB.16) alternative variant aSep08, mRNA.
<a href="#">KRAB.16</a>	<a href="#">KRAB.16.bSep08</a>		1885	1545	2	81	CRA b (KRAB.16) alternative variant bSep08, mRNA.
<a href="#">KRAB.17</a>	<a href="#">KRAB.17.aSep08</a>		5713	2793		497	KRAB box and zinc finger, C2H2-type (KRAB.17) mRNA.
<a href="#">KRAB.18</a>	<a href="#">KRAB.18.aSep08</a>		22796	608		64	CRA a (7.8 kD) (KRAB.18) mRNA.
<a href="#">KRAB.19</a>	<a href="#">KRAB.19.aSep08</a>		19135	732	2	211	KRAB box and zinc finger, C2H2-type (KRAB.19) alternative variant aSep08, mRNA.

<a href="#">KRAB.19</a>	<a href="#">KRAB.19.bSep08</a>		18260	650	2	100	CRA a (11.7 kD) (KRAB.19) alternative variant bSep08, mRNA.
<a href="#">KRAB.19</a>	<a href="#">KRAB.19.cSep08</a>		17400	342	1	64	CRA a (7.7 kD) (KRAB.19) alternative variant cSep08, mRNA.
<a href="#">KRAB.19</a>	<a href="#">KRAB.19.dSep08</a>		18643	509	2	64	CRA a (7.7 kD) (KRAB.19) alternative variant dSep08, mRNA.
<a href="#">KRAB.20</a>	<a href="#">KRAB.20.aSep08</a>		522	331		63	CRA b (KRAB.20) mRNA.
<a href="#">KRAB.21</a>	<a href="#">KRAB.21.aSep08</a>		34784	381		127	CRA a (KRAB.21) mRNA.
<a href="#">KRAB.22</a>	<a href="#">KRAB.22.aSep08</a>		492	301		55	CRA a (KRAB.22) mRNA.
<a href="#">KRAB.23</a>	<a href="#">KRAB.23.aSep08</a>		2192	890	1	296	KRAB box and zinc finger, C2H2-type (KRAB.23) alternative variant aSep08, mRNA.
<a href="#">KRAB.23</a>	<a href="#">KRAB.23.bSep08</a>		21041	586	2	133	CRA b (KRAB.23) alternative variant bSep08, mRNA.
<a href="#">KRAB.24</a>	<a href="#">KRAB.24.aSep08</a>		10014	927	1	255	CRA c (KRAB.24) alternative variant aSep08, mRNA.
<a href="#">KRAB.24</a>	<a href="#">KRAB.24.bSep08</a>		29355	391	1	129	CRA a (KRAB.24) alternative variant bSep08, mRNA.
<a href="#">KRAB.25</a>	<a href="#">KRAB.25.aSep08</a>		15014	390		80	CRA b (KRAB.25) mRNA.
<a href="#">KRAB.26</a>	<a href="#">KRAB.26.bSep08</a>		6440	443	4	53	putative protein (6.0 kD) (KRAB.26) alternative variant bSep08, mRNA.
<a href="#">KRAB.26</a>	<a href="#">KRAB.26.cSep08</a>		5643	445	4	40	zinc finger protein 560 like (4.7 kD) (KRAB.26) alternative variant cSep08, mRNA.
<a href="#">KRAB.26</a>	<a href="#">KRAB.26.dSep08</a>		5726	608	3	39	zinc finger protein 560 like (4.6 kD) (KRAB.26) alternative variant dSep08, mRNA.
<a href="#">KRAB.26</a>	<a href="#">KRAB.26.eSep08</a>		4591	407	2	44	putative protein (KRAB.26) alternative variant eSep08, mRNA.
<a href="#">KRAB.27</a>	<a href="#">KRAB.27.aSep08</a>		12087	443	2	103	CRA a (KRAB.27) alternative variant aSep08, mRNA.
<a href="#">KRAB.27</a>	<a href="#">KRAB.27.bSep08</a>		8298	446	1	76	CRA a (8.6 kD) (KRAB.27) alternative variant bSep08, mRNA.
<a href="#">KRAB.29</a>	<a href="#">KRAB.29.aSep08</a>		30881	872	3	78	CRA a like (8.9 kD) (KRAB.29) mRNA.
<a href="#">KRAB.30</a>	<a href="#">KRAB.30.aSep08</a>		79626	487		86	KRAB box (KRAB.30) mRNA.
<a href="#">KRAB.31</a>	<a href="#">KRAB.31.aSep08</a>		14797	712		208	CRA b (KRAB.31) mRNA.
<a href="#">KRAB.32</a>	<a href="#">KRAB.32.aSep08</a>		22766	570	1	190	CRA a (KRAB.32) alternative variant aSep08, mRNA.
<a href="#">KRAB.32</a>	<a href="#">KRAB.32.bSep08</a>		22799	403		97	CRA a (KRAB.32) alternative variant bSep08, mRNA.
<a href="#">KRAB.33</a>	<a href="#">KRAB.33.aSep08</a>		5391	405		134	CRA b (KRAB.33) mRNA.
<a href="#">KRAB.34</a>	<a href="#">KRAB.34.aSep08</a>		6679	309		57	zinc finger protein 625 CRA b like (KRAB.34) mRNA.
<a href="#">KRAB.35</a>	<a href="#">KRAB.35.aSep08</a>		126261	2882		545	zinc finger protein 422 related sequence 1 (KRAB.35) alternative variant aSep08, mRNA.
<a href="#">KRAB.36</a>	<a href="#">KRAB.36.aSep08</a>		4229	852	3	284	zinc finger protein 7 (KRAB.36) alternative variant aSep08, mRNA.
<a href="#">KRAB.36</a>	<a href="#">KRAB.36.cSep08</a>		2567	661	3	71	zinc finger protein 7 CRA b (8.0 kD) (KRAB.36) alternative variant cSep08, mRNA.
<a href="#">KRAB.36</a>	<a href="#">KRAB.36.dSep08</a>		770	280	2	45	zinc finger protein 7 (KRAB.36) alternative variant dSep08, mRNA.
<a href="#">KRAB.37</a>	<a href="#">KRAB.37.aSep08</a>		4491	389		67	finger protein zinc 4 (KRAB.37) mRNA.
<a href="#">KRAB.38</a>	<a href="#">KRAB.38.aSep08</a>		16846	1365		64	CRA a (7.7 kD) (KRAB.38) alternative variant aSep08, mRNA.
<a href="#">KRAB.40</a>	<a href="#">KRAB.40.aSep08</a>		1388	458		152	protein CRA d (KRAB.40) mRNA.

<a href="#">KRAB.41</a>	<a href="#">KRAB.41.aSep08</a>		8552	529	4	105	CRA a (KRAB.41) alternative variant aSep08, mRNA.
<a href="#">KRAB.42</a>	<a href="#">KRAB.42.aSep08</a>		22697	2753		585	zinc finger protein 248 CRA b (67.4 kD) (KRAB.42) alternative variant aSep08, mRNA.
<a href="#">KRAB.43</a>	<a href="#">KRAB.43.aSep08</a>		27298	819	6	85	KRAB box (10.0 kD) (KRAB.43) alternative variant aSep08, mRNA.
<a href="#">KRAB.43</a>	<a href="#">KRAB.43.bSep08</a>		28858	1904	4	70	KRAB box (8.4 kD) (KRAB.43) alternative variant bSep08, mRNA.
<a href="#">KRAB.43</a>	<a href="#">KRAB.43.cSep08</a>		10626	471	2	36	putative protein (KRAB.43) alternative variant cSep08, mRNA.
<a href="#">KRAB.44</a>	<a href="#">KRAB.44.aSep08</a>		14858	305	1	101	KRAB box (KRAB.44) alternative variant aSep08, mRNA.
<a href="#">KRAB.44</a>	<a href="#">KRAB.44.bSep08</a>		46208	1261	1	72	CRA b (8.4 kD) (KRAB.44) alternative variant bSep08, mRNA.
<a href="#">KRAB.45</a>	<a href="#">KRAB.45.aSep08</a>		23846	335		104	KRAB box (KRAB.45) mRNA.
<a href="#">KRAB.46</a>	<a href="#">KRAB.46.aSep08</a>		35867	726		150	KRAB box and zinc finger, C2H2-type (KRAB.46) mRNA.
<a href="#">KRAB.47</a>	<a href="#">KRAB.47.aSep08</a>		8391	512		115	KRAB box (KRAB.47) mRNA.
<a href="#">KRAB.49</a>	<a href="#">KRAB.49.aSep08</a>		10736	237		79	CRA b (KRAB.49) mRNA.
<a href="#">KRAB.50</a>	<a href="#">KRAB.50.aSep08</a>		521	330		63	CRA b (KRAB.50) mRNA.
<a href="#">KRAB.51</a>	<a href="#">KRAB.51.aSep08</a>		521	330		63	CRA b (KRAB.51) mRNA.
<a href="#">KRAB.52</a>	<a href="#">KRAB.52.aSep08</a>		8642	367		80	KRAB box (KRAB.52) mRNA.
<a href="#">KRAB.53</a>	<a href="#">KRAB.53.aSep08</a>		13857	386		89	KRAB box (KRAB.53) mRNA.
<a href="#">KRAB.54</a>	<a href="#">KRAB.54.aSep08</a>		1263	394		131	zinc finger protein 760 like (KRAB.54) mRNA.
<a href="#">KRAB.55</a>	<a href="#">KRAB.55.aSep08</a>		19241	2273	5	544	KRAB box and zinc finger, C2H2-type (65.1 kD) (KRAB.55) alternative variant aSep08, mRNA.
<a href="#">KRAB.55</a>	<a href="#">KRAB.55.bSep08</a>		31303	548	3	63	KRAB box (7.4 kD) (KRAB.55) alternative variant bSep08, mRNA.
<a href="#">KRAB.55</a>	<a href="#">KRAB.55.cSep08</a>		21782	486	2	63	KRAB box (7.4 kD) (KRAB.55) alternative variant cSep08, mRNA.
<a href="#">KRAB.56</a>	<a href="#">KRAB.56.aSep08</a>		3563	1655		336	CRA b (KRAB.56) alternative variant aSep08, mRNA.
<a href="#">KRAB.57</a>	<a href="#">KRAB.57.aSep08</a>		3378	376		46	finger protein zinc 41 like (KRAB.57) mRNA.
<a href="#">KRAB.58</a>	<a href="#">KRAB.58.aSep08</a>		22723	332	2	103	zinc finger protein ZNF140 (KRAB.58) alternative variant aSep08, mRNA.
<a href="#">KRAB.58</a>	<a href="#">KRAB.58.bSep08</a>		23278	595	2	42	putative protein (KRAB.58) alternative variant bSep08, mRNA.
<a href="#">KRAB.59</a>	<a href="#">KRAB.59.aSep08</a>		10563	401		106	KRAB box (KRAB.59) mRNA.
<a href="#">KRAB.60</a>	<a href="#">KRAB.60.aSep08</a>		5748	738	3	139	zinc finger protein 324 (KRAB.60) alternative variant aSep08, mRNA.
<a href="#">KRAB.60</a>	<a href="#">KRAB.60.bSep08</a>		5602	1457	3	115	zinc finger protein 324 (12.9 kD) (KRAB.60) alternative variant bSep08, mRNA.
<a href="#">KRAB.61</a>	<a href="#">KRAB.61.aSep08</a>		11917	449		143	zinc finger (KRAB.61) mRNA.
<a href="#">KRAB.62</a>	<a href="#">KRAB.62.aSep08</a>		12778	402		115	KRAB box (13.3 kD) (KRAB.62) mRNA.
<a href="#">KRAB.63</a>	<a href="#">KRAB.63.aSep08</a>		10314	729	5	104	KRAB box (12.0 kD) (KRAB.63) alternative variant aSep08, mRNA.
<a href="#">KRAB.63</a>	<a href="#">KRAB.63.bSep08</a>		8725	733	5	96	KRAB box (10.7 kD) (KRAB.63) alternative variant bSep08, mRNA.

<a href="#">KRAB.64</a>	<a href="#">KRAB.64.bSep08</a>		8452	513	1	73	CRA a like (8.6 kD) (KRAB.64) alternative variant bSep08, mRNA.
<a href="#">KRAB.65</a>	<a href="#">KRAB.65.aSep08</a>		25913	750		219	CRA c (KRAB.65) mRNA.
<a href="#">KRAB.66</a>	<a href="#">KRAB.66.aSep08</a>		17998	798	4	229	CRA b (KRAB.66) alternative variant aSep08, mRNA.
<a href="#">KRAB.67</a>	<a href="#">KRAB.67.aSep08</a>		19295	539		155	CRA c (KRAB.67) mRNA.
<a href="#">KRAB.68</a>	<a href="#">KRAB.68.aSep08</a>		10902	924	2	155	putative protein of mammalian origin (KRAB.68) alternative variant aSep08, mRNA.
<a href="#">KRAB.68</a>	<a href="#">KRAB.68.bSep08</a>		10918	812	2	123	putative protein of mammalian origin (KRAB.68) alternative variant bSep08, mRNA.
<a href="#">KRAB.69</a>	<a href="#">KRAB.69.aSep08</a>		13313	738	3	76	KRAB box (KRAB.69) alternative variant aSep08, mRNA.
<a href="#">KRAB.73</a>	<a href="#">KRAB.73.aSep08</a>		26817	772	4	233	CRA a (KRAB.73) alternative variant aSep08, mRNA.
<a href="#">Kras</a>	<a href="#">Kras.aSep08</a>	<a href="#">24525</a>	26389	1594	4	499	v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog (Kras) alternative variant aSep08, mRNA.
<a href="#">Kras</a>	<a href="#">Kras.cSep08</a>	<a href="#">24525</a>	29928	4545	3	78	v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog (9.0 kD) (Kras) alternative variant cSep08, mRNA.
<a href="#">Krba1</a>	<a href="#">Krba1.aSep08</a>	<a href="#">362358</a>	4821	1969	1	500	putative protein, with a coiled coil domain, of mammalian origin (Krba1) alternative variant aSep08, mRNA.
<a href="#">Krba1</a>	<a href="#">Krba1.bSep08</a>	<a href="#">362358</a>	2840	426	1	141	putative protein of mammalian origin (Krba1) alternative variant bSep08, mRNA.
<a href="#">Krcc1</a>	<a href="#">Krcc1.bSep08</a>	<a href="#">312437</a>	12309	773		215	lysine-rich coiled-coil 1 (Krcc1) alternative variant bSep08, mRNA.
<a href="#">Kremen1</a>	<a href="#">Kremen1.bSep08</a>	<a href="#">114107</a>	6612	2397	2	128	kringle containing transmembrane protein 1 (Kremen1) alternative variant bSep08, mRNA.
<a href="#">Krit1</a>	<a href="#">Krit1.bSep08</a>	<a href="#">362317</a>	5353	752	4	167	krev interaction trapped 1 (Krit1) alternative variant bSep08, mRNA.
<a href="#">Krit1</a>	<a href="#">Krit1.cSep08</a>	<a href="#">362317</a>	1160	733	2	101	krev interaction trapped 1 (Krit1) alternative variant cSep08, mRNA.
<a href="#">Krit1</a>	<a href="#">Krit1.dSep08</a>	<a href="#">362317</a>	4432	475	3	97	krev interaction trapped 1 (Krit1) alternative variant dSep08, mRNA.
<a href="#">Krt4</a>	<a href="#">Krt4.bSep08</a>	<a href="#">315323</a>	1979	1052	2	189	keratin 4 (Krt4) alternative variant bSep08, mRNA.
<a href="#">Krt7</a>	<a href="#">Krt7.aSep08</a>	<a href="#">300242</a>	16119	2061	6	457	keratin 7 (50.8 kD) (Krt7) alternative variant aSep08, mRNA.
<a href="#">Krt7</a>	<a href="#">Krt7.bSep08</a>	<a href="#">300242</a>	8396	840	1	279	keratin 7 (Krt7) alternative variant bSep08, mRNA.
<a href="#">Krt7</a>	<a href="#">Krt7.cSep08</a>	<a href="#">300242</a>	9559	1008	2	236	keratin 7 (26.6 kD) (Krt7) alternative variant cSep08, mRNA.
<a href="#">Krt8</a>	<a href="#">Krt8.bSep08</a>	<a href="#">25626</a>	6256	466	3	137	keratin 8 (Krt8) alternative variant bSep08, mRNA.
<a href="#">Krt10</a>	<a href="#">Krt10.bSep08</a>	<a href="#">450225</a>	2920	1929	6	290	keratin 10 (29.8 kD) (Krt10) alternative variant bSep08, mRNA.
<a href="#">Krt12</a>	<a href="#">Krt12.aSep08</a>	<a href="#">360625</a>	4172	918		305	keratin 12 (Meesmann corneal dystrophy) (Krt12) mRNA.
<a href="#">Krt15</a>	<a href="#">Krt15.bSep08</a>	<a href="#">287700</a>	750	486	1	142	keratin 15 (Krt15) alternative variant bSep08, mRNA.
<a href="#">Krt19</a>	<a href="#">Krt19.bSep08</a>	<a href="#">360626</a>	1189	919	3	189	keratin 19 (21.2 kD) (Krt19) alternative variant bSep08, mRNA.
<a href="#">Krt19</a>	<a href="#">Krt19.cSep08</a>	<a href="#">360626</a>	4500	256	2	85	keratin 19 (Krt19) alternative variant cSep08, mRNA.
<a href="#">Krt20</a>	<a href="#">Krt20.aSep08</a>	<a href="#">286912</a>	3915	837		186	keratin 20 (21.9 kD) (Krt20) alternative variant aSep08, mRNA.



<a href="#">Krt23</a>	<a href="#">Krt23.aSep08</a>	<a href="#">287678</a>	6075	795		166	keratin 23 (Krt23) mRNA.
<a href="#">Krt25</a>	<a href="#">Krt25.bSep08</a>	<a href="#">303519</a>	790	458	1	78	keratin 25 (8.8 kD) (Krt25) alternative variant bSep08, mRNA.
<a href="#">Krt26</a>	<a href="#">Krt26.aSep08</a>	<a href="#">407758</a>	4771	764		108	keratin 26 (Krt26) mRNA.
<a href="#">Krt39</a>	<a href="#">Krt39.bSep08</a>	<a href="#">303523</a>	2538	420	2	97	keratin 39 (11.0 kD) (Krt39) alternative variant bSep08, mRNA.
<a href="#">Krt76</a>	<a href="#">Krt76.bSep08</a>	<a href="#">407757</a>	4312	753	7	251	keratin 76 (Krt76) alternative variant bSep08, mRNA.
<a href="#">Krt76</a>	<a href="#">Krt76.cSep08</a>	<a href="#">407757</a>	2485	328	3	45	keratin 76 (Krt76) alternative variant cSep08, mRNA.
<a href="#">Krtcap2</a>	<a href="#">Krtcap2.aSep08</a>	<a href="#">295243</a>	3251	536	5	163	keratinocyte associated protein 2 (Krtcap2) alternative variant aSep08, mRNA.
<a href="#">Krtcap2</a>	<a href="#">Krtcap2.bSep08</a>	<a href="#">295243</a>	3811	915	5	136	keratinocyte associated protein 2 (14.7 kD) (Krtcap2) alternative variant bSep08, mRNA.
<a href="#">Krtcap2</a>	<a href="#">Krtcap2.cSep08</a>	<a href="#">295243</a>	3417	518	5	135	keratinocyte associated protein 2 (14.6 kD) (Krtcap2) alternative variant cSep08, complete mRNA.
<a href="#">Krtcap2</a>	<a href="#">Krtcap2.eSep08</a>	<a href="#">295243</a>	3418	455	4	117	keratinocyte associated protein 2 (12.9 kD) (Krtcap2) alternative variant eSep08, complete mRNA.
<a href="#">Krtcap2</a>	<a href="#">Krtcap2.fSep08</a>	<a href="#">295243</a>	3395	539	5	114	keratinocyte associated protein 2 (12.5 kD) (Krtcap2) alternative variant fSep08, mRNA.
<a href="#">Ktelc1</a>	<a href="#">Ktelc1.aSep08</a>	<a href="#">288091</a>	28899	2892	10	416	KTEL (Lys-Tyr-Glu-Leu) containing 1 (Ktelc1) alternative variant aSep08, mRNA.
<a href="#">Ktelc1</a>	<a href="#">Ktelc1.bSep08</a>	<a href="#">288091</a>	1645	909	1	53	KTEL (Lys-Tyr-Glu-Leu) containing 1 (6.2 kD) (Ktelc1) alternative variant bSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.aSep08</a>	<a href="#">361029</a>	71140	3928	41	1040	kinectin 1 (Ktn1) alternative variant aSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.bSep08</a>	<a href="#">361029</a>	55572	2055	20	656	kinectin 1 (Ktn1) alternative variant bSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.cSep08</a>	<a href="#">361029</a>	15130	1066	14	355	kinectin 1 (Ktn1) alternative variant cSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.dSep08</a>	<a href="#">361029</a>	35370	1343	16	311	kinectin (Ktn1) alternative variant dSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.eSep08</a>	<a href="#">361029</a>	15843	927	11	157	kinectin 1 (18.2 kD) (Ktn1) alternative variant eSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.fSep08</a>	<a href="#">361029</a>	22337	1468	9	114	kinectin (13.3 kD) (Ktn1) alternative variant fSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.gSep08</a>	<a href="#">361029</a>	8708	597	4	92	kinectin (Ktn1) alternative variant gSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.hSep08</a>	<a href="#">361029</a>	9066	714	4	76	kinectin (8.6 kD) (Ktn1) alternative variant hSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.iSep08</a>	<a href="#">361029</a>	5076	892	4	35	putative protein, with a coiled coil domain (4.0 kD) (Ktn1) alternative variant iSep08, mRNA.
<a href="#">Ktn1</a>	<a href="#">Ktn1.jSep08</a>	<a href="#">361029</a>	7518	528	9	23	putative protein (Ktn1) alternative variant jSep08, mRNA.
<a href="#">kuby</a>	<a href="#">kuby.aSep08</a>		1333	484		55	putative protein (kuby) alternative variant aSep08, mRNA.
<a href="#">kuchy</a>	<a href="#">kuchy.aSep08</a>		1192	413		90	enolase 1 like (kuchy) mRNA.
<a href="#">kudar</a>	<a href="#">kudar.aSep08</a>		459	346		24	putative protein (2.8 kD) (kudar) mRNA.
<a href="#">kufer</a>	<a href="#">kufer.aSep08</a>		36487	477	3	78	CRA a like (kufer) alternative variant aSep08, mRNA.
<a href="#">kufer</a>	<a href="#">kufer.bSep08</a>		37198	447	4	78	CRA a like (kufer) alternative variant bSep08, mRNA.
<a href="#">kufer</a>	<a href="#">kufer.cSep08</a>		82071	472	4	56	putative protein (kufer) alternative variant cSep08, mRNA.
<a href="#">kufer</a>	<a href="#">kufer.dSep08</a>		82089	984	7	60	putative protein (kufer) alternative variant dSep08, mRNA.
<a href="#">kufer</a>	<a href="#">kufer.eSep08</a>		3413	510	3	44	CRA b like (4.7 kD) (kufer) alternative variant eSep08, mRNA.

<a href="#">kufer</a>	<a href="#">kufer.fSep08</a>		8164	407	3	44	CRA b like (4.7 kD) (kufer) alternative variant fSep08, mRNA.
<a href="#">kuflo</a>	<a href="#">kuflo.aSep08</a>		2724	323		107	170 -associated factor (kuflo) mRNA.
<a href="#">kuflu</a>	<a href="#">kuflu.aSep08</a>		3522	736		81	putative protein (kuflu) mRNA.
<a href="#">kujey</a>	<a href="#">kujey.aSep08</a>		6836	753		88	CRA a (9.9 kD) (kujey) mRNA.
<a href="#">kukee</a>	<a href="#">kukee.aSep08</a>		37378	706		101	protein phosphatase 1 regulatory (kukee) mRNA.
<a href="#">kuloy</a>	<a href="#">kuloy.aSep08</a>		1673	767		61	putative protein (kuloy) mRNA.
<a href="#">kumee</a>	<a href="#">kumee.aSep08</a>		15645	531		87	putative protein (kumee) mRNA.
<a href="#">kumer</a>	<a href="#">kumer.aSep08</a>		9836	811		270	bromodomain PHD finger transcription factor (kumer) mRNA.
<a href="#">kunoy</a>	<a href="#">kunoy.aSep08</a>		36791	1008		255	heavy axonemal (29.1 kD) (kunoy) mRNA.
<a href="#">kupor</a>	<a href="#">kupor.aSep08</a>		1555	724		76	putative protein (kupor) mRNA.
<a href="#">kusa</a>	<a href="#">kusa.aSep08</a>		39356	690		161	putative protein of eukaryotic origin (kusa) mRNA.
<a href="#">kushee</a>	<a href="#">kushee.aSep08</a>		417	326		67	putative protein (kushee) mRNA.
<a href="#">kutu</a>	<a href="#">kutu.aSep08</a>		4534	388		107	CRA a (kutu) mRNA.
<a href="#">kuvar</a>	<a href="#">kuvar.aSep08</a>		1745	822		51	putative protein (5.5 kD) (kuvar) mRNA.
<a href="#">kuwey</a>	<a href="#">kuwey.aSep08</a>		1149	300		59	putative protein (kuwey) mRNA.
<a href="#">Ky</a>	<a href="#">Ky.bSep08</a>	<a href="#">315962</a>	19884	1087	8	315	kyphoscoliosis peptidase (Ky) alternative variant bSep08, mRNA.
<a href="#">Ky</a>	<a href="#">Ky.cSep08</a>	<a href="#">315962</a>	17378	917	7	234	kyphoscoliosis peptidase (Ky) alternative variant cSep08, mRNA.
<a href="#">kyby</a>	<a href="#">kyby.aSep08</a>		17093	627		84	putative protein of mammalian origin (10.0 kD) (kyby) mRNA.
<a href="#">kychy</a>	<a href="#">kychy.aSep08</a>		714	595		93	putative protein (10.6 kD) (kychy) mRNA.
<a href="#">kydar</a>	<a href="#">kydar.aSep08</a>		2941	286		76	putative protein (kydar) mRNA.
<a href="#">kyfer</a>	<a href="#">kyfer.aSep08</a>		522	343		114	putative protein (kyfer) mRNA.
<a href="#">kyflo</a>	<a href="#">kyflo.aSep08</a>		10762	403	4	86	BTAF1 RNA polymerase II B-TFIID transcription factor-associated (kyflo) alternative variant aSep08, mRNA.
<a href="#">kyflu</a>	<a href="#">kyflu.aSep08</a>		1381	411		136	tyrosine kinase (kyflu) mRNA.
<a href="#">kykee</a>	<a href="#">kykee.aSep08</a>		1932	382	2	123	protein phosphatase 1 regulatory (kykee) alternative variant aSep08, mRNA.
<a href="#">kykee</a>	<a href="#">kykee.bSep08</a>		5943	349	2	105	protein phosphatase 1 regulatory (kykee) alternative variant bSep08, mRNA.
<a href="#">kyloy</a>	<a href="#">kyloy.aSep08</a>		2940	374		37	putative protein (4.0 kD) (kyloy) mRNA.
<a href="#">kymee</a>	<a href="#">kymee.aSep08</a>		7560	674	2	139	flywch family member 2 (14.2 kD) (kymee) alternative variant aSep08, mRNA.
<a href="#">kymee</a>	<a href="#">kymee.bSep08</a>		443	320	1	64	flywch family member 2 (kymee) alternative variant bSep08, mRNA.
<a href="#">kymee</a>	<a href="#">kymee.cSep08</a>		2672	200	1	38	CRA b like (kymee) alternative variant cSep08, mRNA.
<a href="#">kymer</a>	<a href="#">kymer.aSep08</a>		1283	361	1	65	putative protein (7.1 kD) (kymer) alternative variant aSep08, mRNA.
<a href="#">kymer</a>	<a href="#">kymer.bSep08</a>		17762	1081	1	60	putative protein (6.5 kD) (kymer) alternative variant bSep08, mRNA.
<a href="#">kynoy</a>	<a href="#">kynoy.bSep08</a>		717	446	2	45	putative protein (kynoy) alternative variant bSep08, mRNA.

<a href="#">Kynu</a>	<a href="#">Kynu.bSep08</a>	<a href="#">116682</a>	8747	581	3	145	kynureninase (L-kynurenine hydrolase) (Kynu) alternative variant bSep08, mRNA.
<a href="#">kypor</a>	<a href="#">kypor.aSep08</a>		3558	507		168	septin 14 (kypor) mRNA.
<a href="#">kysa</a>	<a href="#">kysa.aSep08</a>		8558	517		62	putative protein (7.0 kD) (kysa) mRNA.
<a href="#">kyshee</a>	<a href="#">kyshee.aSep08</a>		29960	350	2	31	putative protein (kyshee) alternative variant aSep08, mRNA.
<a href="#">kyshee</a>	<a href="#">kyshee.bSep08</a>		32675	313	1	36	putative protein (kyshee) alternative variant bSep08, mRNA.
<a href="#">kytu</a>	<a href="#">kytu.aSep08</a>		826	703		65	putative protein (kytu) mRNA.
<a href="#">kyvar</a>	<a href="#">kyvar.aSep08</a>		536	415	2	35	putative protein (kyvar) alternative variant aSep08, mRNA.
<a href="#">kyvar</a>	<a href="#">kyvar.bSep08</a>		1140	390	2	35	putative protein (kyvar) alternative variant bSep08, mRNA.
<a href="#">kywey</a>	<a href="#">kywey.aSep08</a>		199253	762	2	71	putative protein (kywey) alternative variant aSep08, mRNA.
<a href="#">kywey</a>	<a href="#">kywey.bSep08</a>		254951	755	2	26	putative protein (2.9 kD) (kywey) alternative variant bSep08, mRNA.
<a href="#">L1cam</a>	<a href="#">L1cam.bSep08</a>	<a href="#">50687</a>	3106	1874	1	230	I1 cell adhesion molecule (L1cam) alternative variant bSep08, mRNA.
<a href="#">L2hgdh</a>	<a href="#">L2hgdh.bSep08</a>	<a href="#">314196</a>	7487	1980	1	84	L-2-hydroxyglutarate dehydrogenase (L2hgdh) alternative variant bSep08, mRNA.
<a href="#">L3mbtl</a>	<a href="#">L3mbtl.aSep08</a>	<a href="#">311613</a>	3610	551		108	I(3)mbt-like (Drosophila) (12.1 kD) (L3mbtl) mRNA.
<a href="#">L3mbtl2</a>	<a href="#">L3mbtl2.bSep08</a>	<a href="#">300320</a>	2912	526	1	157	I(3)mbt-like 2 (Drosophila) (17.4 kD) (L3mbtl2) alternative variant bSep08, mRNA.
<a href="#">L3mbtl3</a>	<a href="#">L3mbtl3.aSep08</a>	<a href="#">309550</a>	52881	1781	6	442	I(3)mbt-like 3 (Drosophila) (L3mbtl3) alternative variant aSep08, mRNA.
<a href="#">L3mbtl3</a>	<a href="#">L3mbtl3.cSep08</a>	<a href="#">309550</a>	5835	2384	3	244	I(3)mbt-like 3 (Drosophila) (L3mbtl3) alternative variant cSep08, mRNA.
<a href="#">L27_2.0</a>	<a href="#">L27_2.0.aSep08</a>		32591	406		98	InaD-like (L27_2.0) mRNA.
<a href="#">laby</a>	<a href="#">laby.aSep08</a>		2105	370		123	putative protein of vertebrate origin (laby) mRNA.
<a href="#">Lace1</a>	<a href="#">Lace1.aSep08</a>	<a href="#">502479</a>	6635	904	1	139	lactation elevated 1 (Lace1) alternative variant aSep08, mRNA.
<a href="#">lachy</a>	<a href="#">lachy.aSep08</a>		13014	846		57	putative protein (6.7 kD) (lachy) mRNA.
<a href="#">Lactb2</a>	<a href="#">Lactb2.bSep08</a>	<a href="#">297768</a>	13098	835	3	91	lactamase, beta 2 (Lactb2) alternative variant bSep08, mRNA.
<a href="#">ladar</a>	<a href="#">ladar.aSep08</a>		34068	872	8	193	lysophosphatidylcholine acyltransferase 2 (ladar) alternative variant aSep08, mRNA.
<a href="#">lafer</a>	<a href="#">lafer.aSep08</a>		9462	663		162	armadillo repeat containing 4 (18.3 kD) (lafer) mRNA.
<a href="#">laflo</a>	<a href="#">laflo.aSep08</a>		373	322		52	putative protein (6.0 kD) (laflo) mRNA.
<a href="#">laflu</a>	<a href="#">laflu.aSep08</a>		28664	437		21	putative protein (laflu) mRNA.
<a href="#">Lag3</a>	<a href="#">Lag3.bSep08</a>	<a href="#">297596</a>	6875	1294	7	350	lymphocyte-activation gene 3 (38.5 kD) (Lag3) alternative variant bSep08, mRNA.
<a href="#">Lag3</a>	<a href="#">Lag3.cSep08</a>	<a href="#">297596</a>	1678	1082	2	269	lymphocyte-activation gene 3 (Lag3) alternative variant cSep08, mRNA.
<a href="#">Lag3</a>	<a href="#">Lag3.dSep08</a>	<a href="#">297596</a>	1543	709	3	72	lymphocyte-activation gene 3 (8.2 kD) (Lag3) alternative variant dSep08, mRNA.
<a href="#">lajey</a>	<a href="#">lajey.aSep08</a>		5631	790		207	uncharacterized protein (lajey) mRNA.
<a href="#">lakee</a>	<a href="#">lakee.aSep08</a>		6776	718		108	putative protein (lakee) mRNA.

<a href="#">laloy</a>	<a href="#">laloy.aSep08</a>		4566	414		138	transmembrane protein 44 CRA j (laloy) mRNA.
<a href="#">Lama1</a>	<a href="#">Lama1.bSep08</a>	<a href="#">316758</a>	16505	1799	1	599	laminin, alpha 1 (Lama1) alternative variant bSep08, mRNA.
<a href="#">Lama2</a>	<a href="#">Lama2.aSep08</a>	<a href="#">309368</a>	84699	2478		826	laminin, alpha 2 (Lama2) mRNA.
<a href="#">Lama3</a>	<a href="#">Lama3.bSep08</a>	<a href="#">307582</a>	6762	616	1	175	laminin, alpha 3 (Lama3) alternative variant bSep08, mRNA.
<a href="#">Lama4</a>	<a href="#">Lama4.aSep08</a>	<a href="#">309816</a>	34218	2804	16	882	laminin, alpha 4 (Lama4) alternative variant aSep08, mRNA.
<a href="#">Lama4</a>	<a href="#">Lama4.bSep08</a>	<a href="#">309816</a>	2543	491	1	72	laminin, alpha 4 (Lama4) alternative variant bSep08, mRNA.
<a href="#">Lama5</a>	<a href="#">Lama5.aSep08</a>	<a href="#">140433</a>	17020	1852		616	laminin, alpha 5 (Lama5) alternative variant aSep08, mRNA.
<a href="#">Lama5</a>	<a href="#">Lama5.bSep08</a>	<a href="#">140433</a>	1389	431		143	laminin, alpha 5 (Lama5) alternative variant bSep08, mRNA.
<a href="#">Lamb1</a>	<a href="#">Lamb1.aSep08</a>	<a href="#">298941</a>	34645	3948	21	1265	laminin, beta 1 (Lamb1) alternative variant aSep08, mRNA.
<a href="#">Lamb1</a>	<a href="#">Lamb1.bSep08</a>	<a href="#">298941</a>	2963	615	3	204	laminin, beta 1 (Lamb1) alternative variant bSep08, mRNA.
<a href="#">Lamb1</a>	<a href="#">Lamb1.cSep08</a>	<a href="#">298941</a>	3073	641	4	203	laminin, beta 1 (Lamb1) alternative variant cSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.aSep08</a>	<a href="#">25473</a>	12169	5615	33	1801	laminin (196.5 kD) (Lamb2) alternative variant aSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.bSep08</a>	<a href="#">25473</a>	1146	993	3	266	laminin beta (Lamb2) alternative variant bSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.cSep08</a>	<a href="#">25473</a>	1208	1135	2	150	laminin (Lamb2) alternative variant cSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.dSep08</a>	<a href="#">25473</a>	736	433	4	122	laminin (Lamb2) alternative variant dSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.eSep08</a>	<a href="#">25473</a>	616	507	2	117	laminin (Lamb2) alternative variant eSep08, mRNA.
<a href="#">Lamb2</a>	<a href="#">Lamb2.fSep08</a>	<a href="#">25473</a>	2905	401	2	36	laminin (Lamb2) alternative variant fSep08, mRNA.
<a href="#">Lamb3</a>	<a href="#">Lamb3.aSep08</a>	<a href="#">305078</a>	8434	1575	6	397	laminin, beta 3 (Lamb3) alternative variant aSep08, mRNA.
<a href="#">Lamb3</a>	<a href="#">Lamb3.bSep08</a>	<a href="#">305078</a>	620	323	1	100	laminin, beta 3 (Lamb3) alternative variant bSep08, mRNA.
<a href="#">Lamc1</a>	<a href="#">Lamc1.aSep08</a>	<a href="#">117036</a>	32135	5440		957	laminin, gamma 1 (Lamc1) mRNA.
<a href="#">Lamc2</a>	<a href="#">Lamc2.aSep08</a>	<a href="#">192362</a>	9881	2400	2	373	laminin, gamma 2 (Lamc2) alternative variant aSep08, mRNA.
<a href="#">Lamc2</a>	<a href="#">Lamc2.bSep08</a>	<a href="#">192362</a>	2642	462	1	126	laminin, gamma 2 (Lamc2) alternative variant bSep08, mRNA.
<a href="#">lamee</a>	<a href="#">lamee.aSep08</a>		1282	260		86	polycystin 1 (lamee) mRNA.
<a href="#">lamer</a>	<a href="#">lamer.aSep08</a>		4796	578		192	A member ATP-binding cassette sub-family 8 like (lamer) mRNA.
<a href="#">Laminin_EGF.0</a>	<a href="#">Laminin_EGF.0.aSep08</a>		20328	442		147	laminin alpha 4 (Laminin_EGF.0) mRNA.
<a href="#">Laminin_EGF.1</a>	<a href="#">Laminin_EGF.1.aSep08</a>		2589	473		157	laminin (Laminin_EGF.1) mRNA.
<a href="#">Laminin_EGF.2</a>	<a href="#">Laminin_EGF.2.aSep08</a>		22723	403		134	laminin (Laminin_EGF.2) mRNA.
<a href="#">Laminin_EGF.3</a>	<a href="#">Laminin_EGF.3.aSep08</a>		1514	540		179	laminin (Laminin_EGF.3) mRNA.
<a href="#">Laminin_EGF.4</a>	<a href="#">Laminin_EGF.4.aSep08</a>		10888	689		228	laminin (Laminin_EGF.4) mRNA.

<a href="#">Laminin_EGF.5</a>	<a href="#">Laminin_EGF.5.aSep08</a>		3194	405		135	laminin beta 3 (Laminin_EGF.5) mRNA.
<a href="#">Laminin_EGF.6</a>	<a href="#">Laminin_EGF.6.aSep08</a>		1705	279		92	laminin (Laminin_EGF.6) mRNA.
<a href="#">Laminin_EGF.7</a>	<a href="#">Laminin_EGF.7.aSep08</a>		1520	349		115	heparan sulfate proteoglycan (Laminin_EGF.7) mRNA.
<a href="#">Laminin_EGF.8</a>	<a href="#">Laminin_EGF.8.aSep08</a>		3906	413		137	heparan sulfate proteoglycan (Laminin_EGF.8) mRNA.
<a href="#">Laminin_EGF.9</a>	<a href="#">Laminin_EGF.9.aSep08</a>		1320	690		224	laminin (Laminin_EGF.9) mRNA.
<a href="#">Laminin_EGF.10</a>	<a href="#">Laminin_EGF.10.aSep08</a>		5521	499		165	laminin (Laminin_EGF.10) mRNA.
<a href="#">Laminin_EGF.11</a>	<a href="#">Laminin_EGF.11.aSep08</a>		1187	410		136	multiple -like domain (Laminin_EGF.11) mRNA.
<a href="#">Laminin_EGF.12</a>	<a href="#">Laminin_EGF.12.aSep08</a>		1365	408		135	multiple -like domain (Laminin_EGF.12) mRNA.
<a href="#">Laminin_G_2.0</a>	<a href="#">Laminin_G_2.0.aSep08</a>		33543	548		182	neurexin 1 CRA b (Laminin_G_2.0) mRNA.
<a href="#">Laminin_G_2.1</a>	<a href="#">Laminin_G_2.1.aSep08</a>		4841	3130	15	987	laminin (Laminin_G_2.1) alternative variant aSep08, mRNA.
<a href="#">Laminin_G_2.1</a>	<a href="#">Laminin_G_2.1.bSep08</a>		1436	1026	2	305	laminin alpha (Laminin_G_2.1) alternative variant bSep08, mRNA.
<a href="#">Laminin_G_2.1</a>	<a href="#">Laminin_G_2.1.cSep08</a>		1544	990	7	193	laminin (Laminin_G_2.1) alternative variant cSep08, mRNA.
<a href="#">Laminin_G_2.2</a>	<a href="#">Laminin_G_2.2.aSep08</a>		6713	542		180	FAT 4 (Laminin_G_2.2) mRNA.
<a href="#">Laminin_G_2.3</a>	<a href="#">Laminin_G_2.3.aSep08</a>		72774	1451		403	type XI Collagen (Laminin_G_2.3) mRNA.
<a href="#">Laminin_G_2.4</a>	<a href="#">Laminin_G_2.4.aSep08</a>		3234	571		118	laminin (Laminin_G_2.4) mRNA.
<a href="#">Laminin_I.0</a>	<a href="#">Laminin_I.0.bSep08</a>		18267	715	6	237	laminin (Laminin_I.0) alternative variant bSep08, mRNA.
<a href="#">Laminin_I.1</a>	<a href="#">Laminin_I.1.aSep08</a>		5135	1838		612	laminin (Laminin_I.1) mRNA.
<a href="#">Laminin_II.0</a>	<a href="#">Laminin_II.0.aSep08</a>		2158	778	5	259	laminin (Laminin_II.0) alternative variant aSep08, mRNA.
<a href="#">Laminin_II.0</a>	<a href="#">Laminin_II.0.bSep08</a>		1056	764	1	118	putative protein (Laminin_II.0) alternative variant bSep08, mRNA.
<a href="#">Lamp1</a>	<a href="#">Lamp1.bSep08</a>	<a href="#">25328</a>	24883	4731	9	251	lysosomal-associated membrane protein 1 (Lamp1) alternative variant bSep08, mRNA.
<a href="#">Lamp1</a>	<a href="#">Lamp1.cSep08</a>	<a href="#">25328</a>	9252	458	3	152	lysosomal-associated membrane protein 1 (Lamp1) alternative variant cSep08, mRNA.
<a href="#">Lamp1</a>	<a href="#">Lamp1.dSep08</a>	<a href="#">25328</a>	2602	981	3	85	lysosomal-associated membrane protein 1 (Lamp1) alternative variant dSep08, mRNA.
<a href="#">Lamp2</a>	<a href="#">Lamp2.bSep08</a>	<a href="#">24944</a>	138128	618	3	125	lysosomal-associated membrane protein 2 (Lamp2) alternative variant bSep08, mRNA.
<a href="#">Lamp2</a>	<a href="#">Lamp2.cSep08</a>	<a href="#">24944</a>	15869	449	3	118	lysosomal-associated membrane protein 2 (Lamp2) alternative variant cSep08, mRNA.
<a href="#">Lamp3</a>	<a href="#">Lamp3.bSep08</a>	<a href="#">303801</a>	6482	366	1	58	lysosomal-associated membrane protein 3 (Lamp3) alternative variant bSep08, mRNA.
<a href="#">Lancl2</a>	<a href="#">Lancl2.bSep08</a>	<a href="#">362375</a>	25635	550	5	182	LanC (bacterial lantibiotic synthetase component C)-like 2 (Lancl2) alternative variant bSep08, mRNA.

<a href="#">Lancl2</a>	<a href="#">Lancl2.cSep08</a>	<a href="#">362375</a>	23812	470	3	126	LanC (bacterial lantibiotic synthetase component C)-like 2 (Lancl2) alternative variant cSep08, mRNA.
<a href="#">lanoy</a>	<a href="#">lanoy.aSep08</a>		2601	703		38	putative protein (4.5 kD) (lanoy) mRNA.
<a href="#">Lao1</a>	<a href="#">Lao1.bSep08</a>	<a href="#">298483</a>	8782	647		204	L-amino acid oxidase 1 (Lao1) alternative variant bSep08, mRNA.
<a href="#">Lap3</a>	<a href="#">Lap3.bSep08</a>	<a href="#">289668</a>	6818	739	1	151	leucine aminopeptidase 3 (Lap3) alternative variant bSep08, mRNA.
<a href="#">lapor</a>	<a href="#">lapor.aSep08</a>		590	285		44	putative protein (lapor) mRNA.
<a href="#">Laptm4a</a>	<a href="#">Laptm4a.bSep08</a>	<a href="#">298875</a>	1717	511	1	87	lysosomal-associated protein transmembrane 4A (Laptm4a) alternative variant bSep08, mRNA.
<a href="#">Laptm5</a>	<a href="#">Laptm5.bSep08</a>	<a href="#">89783</a>	4128	1515	2	110	lysosomal-associated protein transmembrane 5 (12.7 kD) (Laptm5) alternative variant bSep08, mRNA.
<a href="#">larby</a>	<a href="#">larby.bSep08</a>		2802	569		63	putative protein (larby) alternative variant bSep08, mRNA.
<a href="#">larchy</a>	<a href="#">larchy.aSep08</a>		1206	399		80	deafness autosomal recessive 59 (larchy) mRNA.
<a href="#">lardar</a>	<a href="#">lardar.aSep08</a>		2080	389	2	129	chromodomain helicase DNA binding protein 9 CRA c like (lardar) alternative variant aSep08, mRNA.
<a href="#">lardar</a>	<a href="#">lardar.bSep08</a>		1068	357	1	119	chromodomain helicase binding protein 9 like (lardar) alternative variant bSep08, mRNA.
<a href="#">larfer</a>	<a href="#">larfer.aSep08</a>		3863	753		121	putative protein of mammalian origin (larfer) mRNA.
<a href="#">larflo</a>	<a href="#">larflo.bSep08</a>		639	531	2	41	putative protein (larflo) alternative variant bSep08, mRNA.
<a href="#">larflu</a>	<a href="#">larflu.aSep08</a>		26877	436		144	neuron navigator 2 (larflu) mRNA.
<a href="#">Large</a>	<a href="#">Large.bSep08</a>	<a href="#">361368</a>	84386	1086	7	361	like-glycosyltransferase (Large) alternative variant bSep08, mRNA.
<a href="#">Large</a>	<a href="#">Large.cSep08</a>	<a href="#">361368</a>	224097	512	4	69	like-glycosyltransferase (Large) alternative variant cSep08, mRNA.
<a href="#">larjey</a>	<a href="#">larjey.aSep08</a>		2685	758		104	putative cytoplasmic protein of mammalian origin (11.5 kD) (larjey) mRNA.
<a href="#">larkee</a>	<a href="#">larkee.aSep08</a>		35831	605		67	putative protein (larkee) mRNA.
<a href="#">larloy</a>	<a href="#">larloy.aSep08</a>		4382	728	3	34	putative protein (larloy) alternative variant aSep08, mRNA.
<a href="#">larmee</a>	<a href="#">larmee.aSep08</a>		7353	533		39	putative protein (larmee) mRNA.
<a href="#">larmer</a>	<a href="#">larmer.aSep08</a>		2566	854		73	ATP-binding cassette sub-family A member like (larmer) mRNA.
<a href="#">larnoy</a>	<a href="#">larnoy.aSep08</a>		9389	275		39	putative protein (larnoy) mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.aSep08</a>	<a href="#">303158</a>	21128	5928	16	869	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant aSep08, mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.bSep08</a>	<a href="#">303158</a>	19511	1793	6	580	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant bSep08, mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.cSep08</a>	<a href="#">303158</a>	1583	744	3	247	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant cSep08, mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.dSep08</a>	<a href="#">303158</a>	2345	486	3	103	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant dSep08, mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.eSep08</a>	<a href="#">303158</a>	2317	487	3	91	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant eSep08, mRNA.
<a href="#">Larp1</a>	<a href="#">Larp1.gSep08</a>	<a href="#">303158</a>	736	600	2	23	la ribonucleoprotein domain family, member 1 (Larp1) alternative variant gSep08, mRNA.

<a href="#">Larp5</a>	<a href="#">Larp5.bSep08</a>	<a href="#">307070</a>	42908	674	1	124	la ribonucleoprotein domain family, member 5 (Larp5) alternative variant bSep08, mRNA.
<a href="#">larpor</a>	<a href="#">larpor.aSep08</a>		45627	575	2	88	CRA a like (larpor) alternative variant aSep08, mRNA.
<a href="#">larpor</a>	<a href="#">larpor.bSep08</a>		12241	842	2	56	CRA b like (larpor) alternative variant bSep08, mRNA.
<a href="#">larpor</a>	<a href="#">larpor.cSep08</a>		45622	442	1	47	putative protein, with a coiled coil domain (larpor) alternative variant cSep08, mRNA.
<a href="#">Lars2</a>	<a href="#">Lars2.bSep08</a>	<a href="#">363172</a>	26824	356	4	118	leucyl-tRNA synthetase, mitochondrial (Lars2) alternative variant bSep08, mRNA.
<a href="#">Lars2</a>	<a href="#">Lars2.cSep08</a>	<a href="#">363172</a>	4425	379	2	105	leucyl-tRNA synthetase, mitochondrial (Lars2) alternative variant cSep08, mRNA.
<a href="#">larsa</a>	<a href="#">larsa.aSep08</a>		9367	959		225	trafficking protein particle complex 9 (larsa) mRNA.
<a href="#">larshee</a>	<a href="#">larshee.aSep08</a>		9514	418		139	putative protein of metazoan origin (larshee) mRNA.
<a href="#">lartu</a>	<a href="#">lartu.bSep08</a>		18295	1378	2	31	metastasis associated 3 like (lartu) alternative variant bSep08, mRNA.
<a href="#">larvar</a>	<a href="#">larvar.aSep08</a>		696	594		108	putative protein (larvar) mRNA.
<a href="#">larwey</a>	<a href="#">larwey.aSep08</a>		1213	565		39	putative protein (4.8 kD) (larwey) mRNA.
<a href="#">Las1.0</a>	<a href="#">Las1.0.aSep08</a>		4022	525	4	174	CRA a (Las1.0) alternative variant aSep08, mRNA.
<a href="#">Las1l</a>	<a href="#">Las1l.aSep08</a>	<a href="#">296865</a>	14952	2380	6	438	putative protein, with a coiled coil domain, of vertebrate origin (49.7 kD) (Las1l) alternative variant aSep08, mRNA.
<a href="#">Las1l</a>	<a href="#">Las1l.bSep08</a>	<a href="#">296865</a>	14952	1318	8	262	putative protein (Las1l) alternative variant bSep08, mRNA.
<a href="#">Las1l</a>	<a href="#">Las1l.cSep08</a>	<a href="#">296865</a>	5865	599	3	133	CRA b (Las1l) alternative variant cSep08, mRNA.
<a href="#">Las1l</a>	<a href="#">Las1l.dSep08</a>	<a href="#">296865</a>	2321	596	2	65	CRA c like (7.2 kD) (Las1l) alternative variant dSep08, mRNA.
<a href="#">lasa</a>	<a href="#">lasa.aSep08</a>		16035	401		44	putative protein (5.0 kD) (lasa) mRNA.
<a href="#">lashee</a>	<a href="#">lashee.aSep08</a>		9776	377		125	putative protein of eukaryotic origin (lashee) mRNA.
<a href="#">Lasp1</a>	<a href="#">Lasp1.aSep08</a>	<a href="#">29278</a>	38521	1204		313	LIM and SH3 protein 1 (Lasp1) mRNA.
<a href="#">Lass2</a>	<a href="#">Lass2.bSep08</a>	<a href="#">310667</a>	9303	1729	11	407	LAG1 2 (Lass2) alternative variant bSep08, mRNA.
<a href="#">Lass2</a>	<a href="#">Lass2.cSep08</a>	<a href="#">310667</a>	6583	789	9	262	LAG1 homolog ceramide synthase 2 (Lass2) alternative variant cSep08, mRNA.
<a href="#">Lass2</a>	<a href="#">Lass2.dSep08</a>	<a href="#">310667</a>	1055	635	4	199	LAG1 2 (Lass2) alternative variant dSep08, mRNA.
<a href="#">Lass2</a>	<a href="#">Lass2.eSep08</a>	<a href="#">310667</a>	5027	384	2	87	LAG1 homolog ceramide synthase 2 (Lass2) alternative variant eSep08, mRNA.
<a href="#">Lass2</a>	<a href="#">Lass2.fSep08</a>	<a href="#">310667</a>	1628	259	2	57	LAG1 homolog ceramide synthase 2 (Lass2) alternative variant fSep08, mRNA.
<a href="#">Lass5</a>	<a href="#">Lass5.bSep08</a>	<a href="#">366984</a>	37861	1911	10	427	LAG1 homolog, ceramide synthase 5 (Lass5) alternative variant bSep08, mRNA.
<a href="#">Lass5</a>	<a href="#">Lass5.cSep08</a>	<a href="#">366984</a>	35373	936	7	311	LAG1 homolog, ceramide synthase 5 (Lass5) alternative variant cSep08, mRNA.
<a href="#">Lass5</a>	<a href="#">Lass5.dSep08</a>	<a href="#">366984</a>	10268	1991	7	231	LAG1 homolog, ceramide synthase 5 (Lass5) alternative variant dSep08, mRNA.
<a href="#">Lat</a>	<a href="#">Lat.bSep08</a>	<a href="#">81511</a>	4987	1217	10	137	linker for activation of T cells (15.1 kD) (Lat) alternative variant bSep08, complete mRNA.
<a href="#">Lat</a>	<a href="#">Lat.cSep08</a>	<a href="#">81511</a>	1435	785	4	123	linker for activation of T cells (13.4 kD) (Lat) alternative variant cSep08, mRNA.

<a href="#">Lat</a>	<a href="#">Lat.dSep08</a>	<a href="#">81511</a>	4091	1776	4	61	linker for activation of T cells (Lat) alternative variant dSep08, mRNA.
<a href="#">Lat2</a>	<a href="#">Lat2.bSep08</a>	<a href="#">317676</a>	4731	2331	3	89	linker for activation of T cells family, member 2 (9.7 kD) (Lat2) alternative variant bSep08, mRNA.
<a href="#">Lats2</a>	<a href="#">Lats2.bSep08</a>	<a href="#">305922</a>	1502	648	1	199	large tumor suppressor 2 (Lats2) alternative variant bSep08, mRNA.
<a href="#">latu</a>	<a href="#">latu.aSep08</a>		97410	893		157	putative cytoplasmic protein (17.8 kD) (latu) mRNA.
<a href="#">lavar</a>	<a href="#">lavar.aSep08</a>		4122	345		114	microtubule-actin crosslinking factor 1 CRA g (lavar) mRNA.
<a href="#">lawby</a>	<a href="#">lawby.aSep08</a>		27481	413	3	73	putative protein (lawby) alternative variant aSep08, mRNA.
<a href="#">lawby</a>	<a href="#">lawby.cSep08</a>		512	386	2	27	putative protein (lawby) alternative variant cSep08, mRNA.
<a href="#">lawchy</a>	<a href="#">lawchy.aSep08</a>		36302	435		78	putative protein (lawchy) mRNA.
<a href="#">lawdar</a>	<a href="#">lawdar.aSep08</a>		6810	797		265	chromodomain helicase DNA binding protein 9 CRA b like (lawdar) mRNA.
<a href="#">lawey</a>	<a href="#">lawey.aSep08</a>		795	591		74	putative protein (lawey) mRNA.
<a href="#">lawfer</a>	<a href="#">lawfer.aSep08</a>		28098	2602		151	cyclin fold protein 1 (lawfer) mRNA.
<a href="#">lawflo</a>	<a href="#">lawflo.aSep08</a>		2521	535		91	putative protein (10.3 kD) (lawflo) mRNA.
<a href="#">lawflu</a>	<a href="#">lawflu.aSep08</a>		10541	570		190	neuron navigator 2 (lawflu) mRNA.
<a href="#">lawjey</a>	<a href="#">lawjey.aSep08</a>		6096	1043		78	putative cytoplasmic protein (8.2 kD) (lawjey) mRNA.
<a href="#">lawkee</a>	<a href="#">lawkee.aSep08</a>		9536	878		21	putative protein (2.3 kD) (lawkee) mRNA.
<a href="#">lawloy</a>	<a href="#">lawloy.aSep08</a>		1964	484	1	44	putative protein (5.0 kD) (lawloy) alternative variant aSep08, mRNA.
<a href="#">lawloy</a>	<a href="#">lawloy.bSep08</a>		1923	345	1	86	putative protein (9.8 kD) (lawloy) alternative variant bSep08, mRNA.
<a href="#">lawmee</a>	<a href="#">lawmee.aSep08</a>		17876	677	4	225	intraflagellar transport 140 (lawmee) alternative variant aSep08, mRNA.
<a href="#">lawmee</a>	<a href="#">lawmee.bSep08</a>		4876	687	2	78	intraflagellar transport 140 (lawmee) alternative variant bSep08, mRNA.
<a href="#">lawmer</a>	<a href="#">lawmer.aSep08</a>		6110	658		218	ATP-binding cassette sub-family A member 6 like (lawmer) mRNA.
<a href="#">lawnoy</a>	<a href="#">lawnoy.aSep08</a>		7565	277		92	putative protein of mammalian origin (lawnoy) mRNA.
<a href="#">lawpor</a>	<a href="#">lawpor.aSep08</a>		2205	877			
<a href="#">lawsa</a>	<a href="#">lawsa.aSep08</a>		8877	715		238	putative protein of metazoan origin (lawsa) mRNA.
<a href="#">lawshee</a>	<a href="#">lawshee.aSep08</a>		2310	383		127	putative protein of metazoan origin (lawshee) mRNA.
<a href="#">lawtu</a>	<a href="#">lawtu.aSep08</a>		621	239		61	putative protein (lawtu) mRNA.
<a href="#">lawvar</a>	<a href="#">lawvar.aSep08</a>		1369	627		102	eukaryotic translation initiation factor 2C (lawvar) mRNA.
<a href="#">lawwey</a>	<a href="#">lawwey.aSep08</a>		17686	355		98	putative protein of metazoan origin (lawwey) mRNA.
<a href="#">Lbp</a>	<a href="#">Lbp.bSep08</a>	<a href="#">29469</a>	12502	662	8	220	lipopolysaccharide binding protein (Lbp) alternative variant bSep08, mRNA.
<a href="#">Lbp</a>	<a href="#">Lbp.cSep08</a>	<a href="#">29469</a>	5869	807	4	86	lipopolysaccharide binding protein (10.2 kD) (Lbp) alternative variant cSep08, complete mRNA.
<a href="#">Lca5</a>	<a href="#">Lca5.bSep08</a>	<a href="#">300866</a>	36437	2205	9	454	leber congenital amaurosis 5 (human) (52.1 kD) (Lca5) alternative variant bSep08, mRNA.
<a href="#">Lca5l</a>	<a href="#">Lca5l.bSep08</a>	<a href="#">498065</a>	4143	1280	1	375	leber congenital amaurosis 5-like (Lca5l) alternative variant bSep08, mRNA.



<a href="#">Lcat</a>	<a href="#">Lcat.bSep08</a>	<a href="#">24530</a>	2069	779	3	253	lecithin cholesterol acyltransferase (Lcat) alternative variant bSep08, mRNA.
<a href="#">Lcat</a>	<a href="#">Lcat.cSep08</a>	<a href="#">24530</a>	2144	845	4	218	lecithin cholesterol acyltransferase (Lcat) alternative variant cSep08, mRNA.
<a href="#">Lck</a>	<a href="#">Lck.bSep08</a>	<a href="#">313050</a>	1973	792	6	263	lymphocyte protein tyrosine kinase (Lck) alternative variant bSep08, mRNA.
<a href="#">Lck</a>	<a href="#">Lck.cSep08</a>	<a href="#">313050</a>	3299	756	5	251	lymphocyte protein tyrosine kinase (Lck) alternative variant cSep08, mRNA.
<a href="#">Lck</a>	<a href="#">Lck.dSep08</a>	<a href="#">313050</a>	1444	823	4	228	lymphocyte protein tyrosine kinase (Lck) alternative variant dSep08, mRNA.
<a href="#">Lck</a>	<a href="#">Lck.eSep08</a>	<a href="#">313050</a>	20175	906	7	219	lymphocyte protein tyrosine kinase (24.6 kD) (Lck) alternative variant eSep08, mRNA.
<a href="#">Lck</a>	<a href="#">Lck.fSep08</a>	<a href="#">313050</a>	2818	770	8	205	lymphocyte protein tyrosine kinase (Lck) alternative variant fSep08, mRNA.
<a href="#">Lcmt1</a>	<a href="#">Lcmt1.bSep08</a>	<a href="#">361643</a>	47411	994	8	277	leucine carboxyl methyltransferase 1 (31.8 kD) (Lcmt1) alternative variant bSep08, mRNA.
<a href="#">Lcmt1</a>	<a href="#">Lcmt1.cSep08</a>	<a href="#">361643</a>	2792	884	2	59	leucine carboxyl methyltransferase 1 (6.6 kD) (Lcmt1) alternative variant cSep08, mRNA.
<a href="#">Lcn2</a>	<a href="#">Lcn2.bSep08</a>	<a href="#">170496</a>	2419	711	2	127	lipocalin 2 (Lcn2) alternative variant bSep08, mRNA.
<a href="#">Lcn3</a>	<a href="#">Lcn3.aSep08</a>	<a href="#">502611</a>	1265	357		43	lipocalin 3 (4.8 kD) (Lcn3) mRNA.
<a href="#">Lcn5</a>	<a href="#">Lcn5.bSep08</a>	<a href="#">29552</a>	1215	286	1	57	lipocalin 5 (Lcn5) alternative variant bSep08, mRNA.
<a href="#">Lcor</a>	<a href="#">Lcor.aSep08</a>	<a href="#">365462</a>	52329	942		189	ligand dependent nuclear receptor corepressor (Lcor) mRNA.
<a href="#">Lcp2</a>	<a href="#">Lcp2.bSep08</a>	<a href="#">155918</a>	13945	748	1	128	lymphocyte cytosolic protein 2 (Lcp2) alternative variant bSep08, mRNA.
<a href="#">Lct</a>	<a href="#">Lct.aSep08</a>	<a href="#">116569</a>	6088	1014		337	lactase (Lct) mRNA.
<a href="#">Ldb1</a>	<a href="#">Ldb1.aSep08</a>	<a href="#">309447</a>	13068	2131	11	534	LIM domain binding 1 (Ldb1) alternative variant aSep08, mRNA.
<a href="#">Ldb1</a>	<a href="#">Ldb1.cSep08</a>	<a href="#">309447</a>	3273	2170	6	267	LIM domain binding 1 (Ldb1) alternative variant cSep08, mRNA.
<a href="#">Ldb2</a>	<a href="#">Ldb2.bSep08</a>	<a href="#">289664</a>	321913	2499	8	371	LIM domain binding 2 (42.5 kD) (Ldb2) alternative variant bSep08, mRNA.
<a href="#">Ldb2</a>	<a href="#">Ldb2.cSep08</a>	<a href="#">289664</a>	6066	584	3	74	LIM domain binding 2 (Ldb2) alternative variant cSep08, mRNA.
<a href="#">Ldb2</a>	<a href="#">Ldb2.dSep08</a>	<a href="#">289664</a>	3185	471	2	33	LIM domain binding 2 (3.7 kD) (Ldb2) alternative variant dSep08, mRNA.
<a href="#">Ldb3</a>	<a href="#">Ldb3.aSep08</a>	<a href="#">498587</a>	20513	3655	5	299	LIM domain binding 3 (Ldb3) alternative variant aSep08, mRNA.
<a href="#">Ldb3</a>	<a href="#">Ldb3.bSep08</a>	<a href="#">498587</a>	1664	455	2	151	LIM domain binding 3 (Ldb3) alternative variant bSep08, mRNA.
<a href="#">Ldb3</a>	<a href="#">Ldb3.cSep08</a>	<a href="#">498587</a>	9629	554	2	89	LIM domain binding 3 (Ldb3) alternative variant cSep08, mRNA.
<a href="#">Ldha</a>	<a href="#">Ldha.bSep08</a>	<a href="#">24533</a>	8178	2227	2	308	lactate dehydrogenase A (33.9 kD) (Ldha) alternative variant bSep08, mRNA.
<a href="#">Ldhb</a>	<a href="#">Ldhb.bSep08</a>	<a href="#">24534</a>	17966	1240	8	259	lactate dehydrogenase B (28.1 kD) (Ldhb) alternative variant bSep08, complete mRNA.

<a href="#">Ldhb</a>	<a href="#">Ldhb.cSep08</a>	<a href="#">24534</a>	2912	943	2	71	lactate dehydrogenase B (8.1 kD) (Ldhb) alternative variant cSep08, mRNA.
<a href="#">Ldhc</a>	<a href="#">Ldhc.aSep08</a>	<a href="#">29634</a>	16215	1093	1	340	lactate dehydrogenase C (Ldhc) alternative variant aSep08, mRNA.
<a href="#">Ldhc</a>	<a href="#">Ldhc.bSep08</a>	<a href="#">29634</a>	11777	729	2	185	lactate dehydrogenase C (Ldhc) alternative variant bSep08, mRNA.
<a href="#">Ldhd</a>	<a href="#">Ldhd.bSep08</a>	<a href="#">307858</a>	2016	884	6	294	lactate dehydrogenase D (Ldhd) alternative variant bSep08, mRNA.
<a href="#">Ldhd</a>	<a href="#">Ldhd.cSep08</a>	<a href="#">307858</a>	2800	2023	5	189	lactate dehydrogenase D (Ldhd) alternative variant cSep08, mRNA.
<a href="#">Ldhd</a>	<a href="#">Ldhd.dSep08</a>	<a href="#">307858</a>	593	355	3	82	lactate dehydrogenase D (Ldhd) alternative variant dSep08, mRNA.
<a href="#">Ldhd</a>	<a href="#">Ldhd.eSep08</a>	<a href="#">307858</a>	588	506	2	67	lactate dehydrogenase D (7.9 kD) (Ldhd) alternative variant eSep08, mRNA.
<a href="#">Ldlr</a>	<a href="#">Ldlr.bSep08</a>	<a href="#">300438</a>	2792	2116	2	87	low density lipoprotein receptor (18.5 kD) (Ldlr) alternative variant bSep08, mRNA.
<a href="#">Ldlrad3</a>	<a href="#">Ldlrad3.aSep08</a>	<a href="#">366138</a>	2228	498		116	putative protein of vertebrate origin (Ldlrad3) mRNA.
<a href="#">Ldl_recept_a.0</a>	<a href="#">Ldl_recept_a.0.aSep08</a>		11668	485		161	receptor protein (Ldl_recept_a.0) mRNA.
<a href="#">Ldl_recept_a.1</a>	<a href="#">Ldl_recept_a.1.aSep08</a>		3835	1045		348	low density lipoprotein-related protein 1B (Ldl_recept_a.1) mRNA.
<a href="#">Ldl_recept_a.2</a>	<a href="#">Ldl_recept_a.2.aSep08</a>		3483	1251		416	low density lipoprotein-related protein 1 CRA b (Ldl_recept_a.2) mRNA.
<a href="#">Ldl_recept_a.3</a>	<a href="#">Ldl_recept_a.3.aSep08</a>		7090	1182	8	394	low density lipoprotein-related protein 1 CRA a (Ldl_recept_a.3) alternative variant aSep08, mRNA.
<a href="#">Ldl_recept_a.4</a>	<a href="#">Ldl_recept_a.4.aSep08</a>		10213	386		128	low density lipoprotein-related protein 1 CRA a (Ldl_recept_a.4) mRNA.
<a href="#">Ldl_recept_a.6</a>	<a href="#">Ldl_recept_a.6.aSep08</a>		4491	740		219	low density lipoprotein-related protein 2 (Ldl_recept_a.6) mRNA.
<a href="#">Ldl_recept_a.8</a>	<a href="#">Ldl_recept_a.8.aSep08</a>		21680	734	5	221	complement component 7 CRA b (Ldl_recept_a.8) alternative variant aSep08, mRNA.
<a href="#">Ldl_recept_a.8</a>	<a href="#">Ldl_recept_a.8.bSep08</a>		7600	760	2	153	complement C7 (Ldl_recept_a.8) alternative variant bSep08, mRNA.
<a href="#">Ldl_recept_b.0</a>	<a href="#">Ldl_recept_b.0.aSep08</a>		6097	682		227	sortilin-related receptor (Ldl_recept_b.0) mRNA.
<a href="#">Ldl_recept_b.1</a>	<a href="#">Ldl_recept_b.1.aSep08</a>		2688	951	7	316	low density lipoprotein-related protein 1B (Ldl_recept_b.1) alternative variant aSep08, mRNA.
<a href="#">Leap2</a>	<a href="#">Leap2.bSep08</a>	<a href="#">497901</a>	1045	399	2	75	liver-expressed antimicrobial peptide 2 (Leap2) alternative variant bSep08, mRNA.
<a href="#">Lect1</a>	<a href="#">Lect1.bSep08</a>	<a href="#">81512</a>	25714	1330	7	332	leukocyte cell derived chemotaxin 1 (37.1 kD) (Lect1) alternative variant bSep08, mRNA.
<a href="#">Lect1</a>	<a href="#">Lect1.cSep08</a>	<a href="#">81512</a>	19777	1067	5	260	leukocyte cell derived chemotaxin 1 (Lect1) alternative variant cSep08, mRNA.
<a href="#">Lect1</a>	<a href="#">Lect1.dSep08</a>	<a href="#">81512</a>	6493	822	3	145	leukocyte cell derived chemotaxin 1 (15.5 kD) (Lect1) alternative variant dSep08, mRNA.
<a href="#">leeby</a>	<a href="#">leeby.aSep08</a>		10639	1784		18	putative protein (leeby) mRNA.
<a href="#">leechy</a>	<a href="#">leechy.aSep08</a>		384	291		96	titin CRA a (leechy) mRNA.

<a href="#">leedar</a>	<a href="#">leedar.aSep08</a>		6213	303		98	chromodomain helicase DNA binding protein 9 CRA c like (leedar) mRNA.
<a href="#">leefer</a>	<a href="#">leefer.bSep08</a>		1770	355		53	putative protein (leefer) alternative variant bSep08, mRNA.
<a href="#">leeflo</a>	<a href="#">leeflo.bSep08</a>		9451	617		41	putative protein (4.6 kD) (leeflo) alternative variant bSep08, mRNA.
<a href="#">leeflu</a>	<a href="#">leeflu.aSep08</a>		8850	802		267	neuron navigator 2 (leeflu) mRNA.
<a href="#">leejey</a>	<a href="#">leejey.aSep08</a>		1653	455		45	putative protein (5.3 kD) (leejey) mRNA.
<a href="#">leekee</a>	<a href="#">leekee.aSep08</a>		4101	486		114	putative protein (leekee) mRNA.
<a href="#">leeloy</a>	<a href="#">leeloy.aSep08</a>		12068	1563		91	putative protein (leeloy) mRNA.
<a href="#">leemee</a>	<a href="#">leemee.aSep08</a>		14857	820	6	273	intraflagellar transport 140 (leemee) alternative variant aSep08, mRNA.
<a href="#">leemee</a>	<a href="#">leemee.bSep08</a>		1175	377	1	77	intraflagellar transport 140 (leemee) alternative variant bSep08, mRNA.
<a href="#">leemer</a>	<a href="#">leemer.aSep08</a>		4002	431		143	ATP-binding cassette sub-family A member 6 like (leemer) mRNA.
<a href="#">leenoy</a>	<a href="#">leenoy.aSep08</a>		2600	333		79	putative protein, with a coiled coil domain, of mammalian origin (leenoy) mRNA.
<a href="#">leepor</a>	<a href="#">leepor.aSep08</a>		34157	787	3	85	putative protein (9.5 kD) (leepor) alternative variant aSep08, mRNA.
<a href="#">leepor</a>	<a href="#">leepor.bSep08</a>		2624	396	1	28	putative protein (leepor) alternative variant bSep08, mRNA.
<a href="#">leesa</a>	<a href="#">leesa.aSep08</a>		7437	3135	4	560	45 4 (leesa) alternative variant aSep08, mRNA.
<a href="#">leeshee</a>	<a href="#">leeshee.aSep08</a>		17695	373	5	124	putative protein of vertebrate origin (leeshee) alternative variant aSep08, mRNA.
<a href="#">leetu</a>	<a href="#">leetu.aSep08</a>		18740	1062		353	death receptor protein (leetu) mRNA.
<a href="#">leever</a>	<a href="#">leever.aSep08</a>		7827	411		137	eukaryotic translation initiation factor 2C 3 (leever) mRNA.
<a href="#">leewey</a>	<a href="#">leewey.aSep08</a>		1937	596		84	putative protein (9.7 kD) (leewey) mRNA.
<a href="#">Lef1</a>	<a href="#">Lef1.bSep08</a>	<a href="#">161452</a>	137824	2083	9	315	lymphoid enhancer binding factor 1 (Lef1) alternative variant bSep08, mRNA.
<a href="#">Lef1</a>	<a href="#">Lef1.cSep08</a>	<a href="#">161452</a>	31682	1152	8	208	lymphoid enhancer binding factor 1 (Lef1) alternative variant cSep08, mRNA.
<a href="#">Lef1</a>	<a href="#">Lef1.dSep08</a>	<a href="#">161452</a>	26349	741	5	160	lymphoid enhancer binding factor 1 (Lef1) alternative variant dSep08, mRNA.
<a href="#">Lef1</a>	<a href="#">Lef1.eSep08</a>	<a href="#">161452</a>	137169	761	3	101	lymphoid enhancer binding factor 1 (Lef1) alternative variant eSep08, mRNA.
<a href="#">LEM.0</a>	<a href="#">LEM.0.aSep08</a>		885	331		70	CRA c (LEM.0) mRNA.
<a href="#">Lemd2</a>	<a href="#">Lemd2.bSep08</a>	<a href="#">361807</a>	6622	1654	6	216	putative protein of metazoan origin (Lemd2) alternative variant bSep08, mRNA.
<a href="#">Lemd2</a>	<a href="#">Lemd2.cSep08</a>	<a href="#">361807</a>	2292	389	4	97	putative protein of vertebrate origin (Lemd2) alternative variant cSep08, mRNA.
<a href="#">Lemd3</a>	<a href="#">Lemd3.aSep08</a>	<a href="#">680066</a>	49945	3723	1	577	inner nuclear membrane protein (Lemd3) alternative variant aSep08, mRNA.
<a href="#">Lemd3</a>	<a href="#">Lemd3.bSep08</a>	<a href="#">680066</a>	41745	442		48	inner nuclear membrane protein like (Lemd3) alternative variant bSep08, mRNA.
<a href="#">Lenep</a>	<a href="#">Lenep.bSep08</a>	<a href="#">113917</a>	8342	1736	8	474	lens epithelial protein (52.6 kD) (Lenep) alternative variant bSep08, complete mRNA.

<a href="#">Lenep</a>	<a href="#">Lenep.cSep08</a>	<a href="#">113917</a>	5183	1254	5	332	lens epithelial protein (Lenep) alternative variant cSep08, mRNA.
<a href="#">Lenep</a>	<a href="#">Lenep.dSep08</a>	<a href="#">113917</a>	4225	1312	2	258	lens epithelial protein (28.3 kD) (Lenep) alternative variant dSep08, mRNA.
<a href="#">Lenep</a>	<a href="#">Lenep.eSep08</a>	<a href="#">113917</a>	3626	751	2	250	lens epithelial protein (Lenep) alternative variant eSep08, mRNA.
<a href="#">Leng8</a>	<a href="#">Leng8.bSep08</a>	<a href="#">361506</a>	1890	881	6	293	leukocyte receptor cluster (LRC) member 8 (Leng8) alternative variant bSep08, mRNA.
<a href="#">Leng8</a>	<a href="#">Leng8.cSep08</a>	<a href="#">361506</a>	3518	502	5	129	leukocyte receptor cluster (LRC) member 8 (Leng8) alternative variant cSep08, mRNA.
<a href="#">Leng8</a>	<a href="#">Leng8.dSep08</a>	<a href="#">361506</a>	2071	648	2	114	leukocyte receptor cluster (LRC) member 8 (Leng8) alternative variant dSep08, mRNA.
<a href="#">Leo1</a>	<a href="#">Leo1.bSep08</a>	<a href="#">300837</a>	5999	1304	2	93	leo1, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (10.3 kD) (Leo1) alternative variant bSep08, mRNA.
<a href="#">Lepre1</a>	<a href="#">Lepre1.bSep08</a>	<a href="#">114200</a>	15181	2763	14	411	leprecan 1 (46.8 kD) (Lepre1) alternative variant bSep08, complete mRNA.
<a href="#">Leprel2</a>	<a href="#">Leprel2.aSep08</a>	<a href="#">297595</a>	15856	2571	15	735	leprecan-like 2 (Leprel2) alternative variant aSep08, mRNA.
<a href="#">Leprel2</a>	<a href="#">Leprel2.bSep08</a>	<a href="#">297595</a>	1148	1033	2	208	leprecan-like 2 (23.2 kD) (Leprel2) alternative variant bSep08, mRNA.
<a href="#">Leprel2</a>	<a href="#">Leprel2.dSep08</a>	<a href="#">297595</a>	5665	406	5	104	leprecan-like 2 (Leprel2) alternative variant dSep08, mRNA.
<a href="#">Leprel2</a>	<a href="#">Leprel2.fSep08</a>	<a href="#">297595</a>	2872	667	3	90	leprecan-like 2 (Leprel2) alternative variant fSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.bSep08</a>	<a href="#">24536</a>	37394	1782	10	380	leptin receptor (Leprot) alternative variant bSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.bSep08</a>	<a href="#">56766</a>	37394	1782	10	380	leptin receptor (Leprot) alternative variant bSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.cSep08</a>	<a href="#">24536</a>	9670	605	5	188	leptin receptor (Leprot) alternative variant cSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.cSep08</a>	<a href="#">56766</a>	9670	605	5	188	leptin receptor (Leprot) alternative variant cSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.eSep08</a>	<a href="#">24536</a>	9328	1231	3	113	leptin receptor (12.2 kD) (Leprot) alternative variant eSep08, mRNA.
<a href="#">Leprot</a>	<a href="#">Leprot.eSep08</a>	<a href="#">56766</a>	9328	1231	3	113	leptin receptor (12.2 kD) (Leprot) alternative variant eSep08, mRNA.
<a href="#">Ierby</a>	<a href="#">Ierby.aSep08</a>		11512	533		177	WWC family member 3 (Ierby) mRNA.
<a href="#">Ierchy</a>	<a href="#">Ierchy.aSep08</a>		4329	258		85	titin (Ierchy) mRNA.
<a href="#">Ierdar</a>	<a href="#">Ierdar.aSep08</a>		15446	652		41	putative protein (4.5 kD) (Ierdar) mRNA.
<a href="#">Ierfer</a>	<a href="#">Ierfer.aSep08</a>		1475	331	1	62	putative protein (Ierfer) alternative variant aSep08, mRNA.
<a href="#">Ierfer</a>	<a href="#">Ierfer.bSep08</a>		1228	753	1	54	putative protein (6.2 kD) (Ierfer) alternative variant bSep08, mRNA.
<a href="#">Ierflo</a>	<a href="#">Ierflo.aSep08</a>		2046	610		98	putative protein (Ierflo) mRNA.
<a href="#">Ierflu</a>	<a href="#">Ierflu.aSep08</a>		41697	404		134	NEL-like 1 (Ierflu) mRNA.
<a href="#">Iergar</a>	<a href="#">Iergar.aSep08</a>		2576	516		69	putative protein (Iergar) mRNA.
<a href="#">Ierjey</a>	<a href="#">Ierjey.aSep08</a>		1632	220		72	polycystic kidney disease like (Ierjey) mRNA.
<a href="#">Ierkee</a>	<a href="#">Ierkee.aSep08</a>		5152	263		87	complement factor H-related protein (Ierkee) mRNA.
<a href="#">Ierloy</a>	<a href="#">Ierloy.aSep08</a>		865	372		39	putative protein (Ierloy) mRNA.

<a href="#">lermee</a>	<a href="#">lermee.aSep08</a>		3745	1023	7	340	intraflagellar transport 140 (lermee) alternative variant aSep08, mRNA.
<a href="#">lermee</a>	<a href="#">lermee.bSep08</a>		837	396	2	114	intraflagellar transport 140 (lermee) alternative variant bSep08, mRNA.
<a href="#">lermer</a>	<a href="#">lermer.aSep08</a>		1138	393		29	putative protein (lermer) mRNA.
<a href="#">lernoy</a>	<a href="#">lernoy.aSep08</a>		6876	1829	1	567	bone protein receptor type II (lernoy) alternative variant aSep08, mRNA.
<a href="#">lernoy</a>	<a href="#">lernoy.bSep08</a>		1847	386	1	105	bone protein receptor type II (lernoy) alternative variant bSep08, mRNA.
<a href="#">lerpor</a>	<a href="#">lerpor.aSep08</a>		4093	467		76	putative membrane protein (8.6 kD) (lerpor) mRNA.
<a href="#">lersa</a>	<a href="#">lersa.aSep08</a>		2987	276		28	putative protein (lersa) mRNA.
<a href="#">lershee</a>	<a href="#">lershee.aSep08</a>		7023	943	5	314	putative protein of metazoan origin (lershee) alternative variant aSep08, mRNA.
<a href="#">lershee</a>	<a href="#">lershee.bSep08</a>		2480	393	2	131	putative protein of bilateral origin (lershee) alternative variant bSep08, mRNA.
<a href="#">lershee</a>	<a href="#">lershee.cSep08</a>		2997	364	3	121	putative protein of metazoan origin (lershee) alternative variant cSep08, mRNA.
<a href="#">lertu</a>	<a href="#">lertu.aSep08</a>		9039	405		134	adenoma-associated protein homolog (lertu) mRNA.
<a href="#">lervar</a>	<a href="#">lervar.aSep08</a>		33896	696		60	putative protein (6.5 kD) (lervar) mRNA.
<a href="#">lerwey</a>	<a href="#">lerwey.aSep08</a>		12283	686		77	putative protein (lerwey) mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.aSep08</a>	<a href="#">361169</a>	5611	1853	4	529	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant aSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.cSep08</a>	<a href="#">361169</a>	17804	1217	8	405	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant cSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.dSep08</a>	<a href="#">361169</a>	4223	624	3	181	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant dSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.fSep08</a>	<a href="#">361169</a>	3753	310	3	103	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant fSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.gSep08</a>	<a href="#">361169</a>	1778	401	3	55	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant gSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.iSep08</a>	<a href="#">361169</a>	1099	672	2	108	leucine zipper-EF-hand containing transmembrane protein 2 (10.9 kD) (Letm2) alternative variant iSep08, mRNA.
<a href="#">Letm2</a>	<a href="#">Letm2.lSep08</a>	<a href="#">361169</a>	837	326	2	36	leucine zipper-EF-hand containing transmembrane protein 2 (Letm2) alternative variant lSep08, mRNA.
<a href="#">Letmd1</a>	<a href="#">Letmd1.aSep08</a>	<a href="#">681352</a>	13422	1344	9	371	LETM1-like (43.1 kD) (Letmd1) alternative variant aSep08, complete mRNA.
<a href="#">Letmd1</a>	<a href="#">Letmd1.cSep08</a>	<a href="#">681352</a>	2978	688	3	217	CRA c (24.8 kD) (Letmd1) alternative variant cSep08, mRNA.
<a href="#">Letmd1</a>	<a href="#">Letmd1.dSep08</a>	<a href="#">681352</a>	4393	2114	7	145	putative protein of vertebrate origin (Letmd1) alternative variant dSep08, mRNA.
<a href="#">Letmd1</a>	<a href="#">Letmd1.eSep08</a>	<a href="#">681352</a>	1039	593	2	106	putative protein of vertebrate origin (Letmd1) alternative variant eSep08, mRNA.
<a href="#">leyby</a>	<a href="#">leyby.aSep08</a>		3902	602		200	WWC family member 3 (leyby) mRNA.
<a href="#">leychy</a>	<a href="#">leychy.aSep08</a>		3443	313		104	titin like (leychy) mRNA.
<a href="#">leydar</a>	<a href="#">leydar.aSep08</a>		66570	708		76	putative protein (leydar) mRNA.
<a href="#">leyfer</a>	<a href="#">leyfer.aSep08</a>		34333	750		33	putative protein (leyfer) mRNA.

<a href="#">leyflo</a>	<a href="#">leyflo.aSep08</a>		1245	1129		16	putative protein (1.9 kD) (leyflo) alternative variant aSep08, mRNA.
<a href="#">leyflu</a>	<a href="#">leyflu.aSep08</a>		19624	668		39	putative protein (4.6 kD) (leyflu) mRNA.
<a href="#">leygar</a>	<a href="#">leygar.aSep08</a>		1133	530		60	putative protein (leygar) mRNA.
<a href="#">leyjey</a>	<a href="#">leyjey.aSep08</a>		3272	462		153	polycystic kidney disease like (leyjey) mRNA.
<a href="#">leykee</a>	<a href="#">leykee.aSep08</a>		22001	785		152	potassium channel subfamily T member 2 CRA a (leykee) mRNA.
<a href="#">leyloy</a>	<a href="#">leyloy.aSep08</a>		1170	701		30	putative protein (leyloy) mRNA.
<a href="#">leymee</a>	<a href="#">leymee.aSep08</a>		4536	884		111	intraflagellar transport 140 (leymee) mRNA.
<a href="#">leymer</a>	<a href="#">leymer.aSep08</a>		16167	601		55	putative protein (6.0 kD) (leymer) mRNA.
<a href="#">leynoy</a>	<a href="#">leynoy.aSep08</a>		928	713		46	putative protein (leynoy) mRNA.
<a href="#">leypor</a>	<a href="#">leypor.aSep08</a>		988	692		54	putative protein (5.5 kD) (leypor) mRNA.
<a href="#">leysa</a>	<a href="#">leysa.aSep08</a>		2586	689		110	putative nuclear protein of mammalian origin (12.3 kD) (leysa) mRNA.
<a href="#">leyshee</a>	<a href="#">leyshee.aSep08</a>		1248	230		76	putative protein of vertebrate origin (leyshee) mRNA.
<a href="#">leytu</a>	<a href="#">leytu.aSep08</a>		34186	252		84	putative protein, with a coiled coil domain, of vertebrate origin (leytu) mRNA.
<a href="#">leyvar</a>	<a href="#">leyvar.aSep08</a>		76329	1898		632	polycystic kidney disease 1-like (leyvar) mRNA.
<a href="#">leywey</a>	<a href="#">leywey.aSep08</a>		2143	334		110	inositol 1 receptor (leywey) mRNA.
<a href="#">Lfng</a>	<a href="#">Lfng.aSep08</a>	<a href="#">170905</a>	1540	1409	1	414	putative protein (Lfng) alternative variant aSep08, mRNA.
<a href="#">Lgals1</a>	<a href="#">Lgals1.cSep08</a>	<a href="#">56646</a>	1353	1154	2	101	lectin, galactose binding, soluble 1 (Lgals1) alternative variant cSep08, mRNA.
<a href="#">Lgals2</a>	<a href="#">Lgals2.bSep08</a>	<a href="#">171134</a>	2382	1398	3	119	lectin, galactoside-binding, soluble 2 (13.4 kD) (Lgals2) alternative variant bSep08, mRNA.
<a href="#">Lgals2</a>	<a href="#">Lgals2.cSep08</a>	<a href="#">171134</a>	7473	1431	4	41	lectin, galactoside-binding, soluble 2 (4.6 kD) (Lgals2) alternative variant cSep08, mRNA.
<a href="#">Lgals3</a>	<a href="#">Lgals3.aSep08</a>	<a href="#">83781</a>	11935	947	6	262	lectin, galactose binding, soluble 3 (27.2 kD) (Lgals3) alternative variant aSep08, complete mRNA.
<a href="#">Lgals3</a>	<a href="#">Lgals3.cSep08</a>	<a href="#">83781</a>	3947	390	3	129	lectin, galactose binding, soluble 3 (Lgals3) alternative variant cSep08, mRNA.
<a href="#">Lgals3</a>	<a href="#">Lgals3.dSep08</a>	<a href="#">83781</a>	2498	1682	2	104	lectin, galactose binding, soluble 3 (12.1 kD) (Lgals3) alternative variant dSep08, mRNA.
<a href="#">Lgals3bp</a>	<a href="#">Lgals3bp.bSep08</a>	<a href="#">245955</a>	5866	2109	2	160	lectin, galactoside-binding, soluble, 3 binding protein (17.5 kD) (Lgals3bp) alternative variant bSep08, mRNA.
<a href="#">Lgals7</a>	<a href="#">Lgals7.aSep08</a>	<a href="#">29518</a>	2413	1502	3	426	lectin, galactose binding, soluble 7 (Lgals7) alternative variant aSep08, mRNA.
<a href="#">Lgals7</a>	<a href="#">Lgals7.cSep08</a>	<a href="#">29518</a>	868	525	1	100	lectin, galactose binding, soluble 7 (11.4 kD) (Lgals7) alternative variant cSep08, mRNA.
<a href="#">Lgals8</a>	<a href="#">Lgals8.bSep08</a>	<a href="#">116641</a>	8266	2121	2	123	lectin, galactose binding, soluble 8 (14.0 kD) (Lgals8) alternative variant bSep08, mRNA.
<a href="#">Lgals9</a>	<a href="#">Lgals9.bSep08</a>	<a href="#">25476</a>	20406	916	10	291	lectin, galactose binding, soluble 9 (Lgals9) alternative variant bSep08, mRNA.
<a href="#">Lgals9</a>	<a href="#">Lgals9.cSep08</a>	<a href="#">25476</a>	20395	773	8	248	lectin, galactose binding, soluble 9 (Lgals9) alternative variant cSep08, mRNA.

<a href="#">Lgals9</a>	<a href="#">Lgals9.eSep08</a>	<a href="#">25476</a>	2494	1756	3	76	lectin, galactose binding, soluble 9 (8.0 kD) (Lgals9) alternative variant eSep08, mRNA.
<a href="#">Lgi2</a>	<a href="#">Lgi2.aSep08</a>	<a href="#">305417</a>	27483	1758	8	551	leucine-rich repeat LGI family, member 2 (Lgi2) alternative variant aSep08, mRNA.
<a href="#">Lgi3</a>	<a href="#">Lgi3.bSep08</a>	<a href="#">306013</a>	3142	1092	5	241	leucine-rich repeat LGI family member 3 CRA a (Lgi3) alternative variant bSep08, mRNA.
<a href="#">Lgi3</a>	<a href="#">Lgi3.cSep08</a>	<a href="#">306013</a>	3001	1579	3	208	leucine-rich repeat Lgi family member 3 CRA a (Lgi3) alternative variant cSep08, mRNA.
<a href="#">Lgi3</a>	<a href="#">Lgi3.dSep08</a>	<a href="#">306013</a>	853	696	2	127	leucine-rich repeat LGI family member 3 (Lgi3) alternative variant dSep08, mRNA.
<a href="#">Lgi4</a>	<a href="#">Lgi4.bSep08</a>	<a href="#">361549</a>	8430	693	5	93	leucine-rich repeat LGI family, member 4 (Lgi4) alternative variant bSep08, mRNA.
<a href="#">Lgmn</a>	<a href="#">Lgmn.bSep08</a>	<a href="#">63865</a>	31714	858	9	240	legumain (Lgmn) alternative variant bSep08, mRNA.
<a href="#">Lgtn</a>	<a href="#">Lgtn.bSep08</a>	<a href="#">498225</a>	1327	755	2	130	ligatin (Lgtn) alternative variant bSep08, mRNA.
<a href="#">Lgtn</a>	<a href="#">Lgtn.cSep08</a>	<a href="#">498225</a>	4910	425	3	107	ligatin (Lgtn) alternative variant cSep08, mRNA.
<a href="#">Lgtn</a>	<a href="#">Lgtn.dSep08</a>	<a href="#">498225</a>	11321	709	4	90	ligatin (Lgtn) alternative variant dSep08, mRNA.
<a href="#">Lgtn</a>	<a href="#">Lgtn.eSep08</a>	<a href="#">498225</a>	947	615	2	77	ligatin (Lgtn) alternative variant eSep08, mRNA.
<a href="#">Lhpp</a>	<a href="#">Lhpp.bSep08</a>	<a href="#">361663</a>	17317	707	4	159	phospholysine phosphohistidine inorganic pyrophosphate phosphatase (Lhpp) alternative variant bSep08, mRNA.
<a href="#">Lhpp</a>	<a href="#">Lhpp.cSep08</a>	<a href="#">361663</a>	9475	316	2	105	phospholysine phosphohistidine inorganic pyrophosphate phosphatase (Lhpp) alternative variant cSep08, mRNA.
<a href="#">Lhx2</a>	<a href="#">Lhx2.bSep08</a>	<a href="#">296706</a>	1529	659	1	165	LIM homeobox protein 2 (Lhx2) alternative variant bSep08, mRNA.
<a href="#">Lhx3</a>	<a href="#">Lhx3.aSep08</a>	<a href="#">170671</a>	1171	492		164	LIM homeobox protein 3 (Lhx3) mRNA.
<a href="#">Lhx4</a>	<a href="#">Lhx4.bSep08</a>	<a href="#">360858</a>	2595	512	2	146	LIM homeobox protein 4 (Lhx4) alternative variant bSep08, mRNA.
<a href="#">Lhx4</a>	<a href="#">Lhx4.cSep08</a>	<a href="#">360858</a>	29978	408	2	127	LIM homeobox protein 4 (Lhx4) alternative variant cSep08, mRNA.
<a href="#">Lhx4</a>	<a href="#">Lhx4.dSep08</a>	<a href="#">360858</a>	27114	529	2	94	LIM homeobox protein 4 (Lhx4) alternative variant dSep08, mRNA.
<a href="#">Lhx9</a>	<a href="#">Lhx9.bSep08</a>	<a href="#">289048</a>	18783	1787	6	315	LIM homeobox 9 (Lhx9) alternative variant bSep08, mRNA.
<a href="#">Lias</a>	<a href="#">Lias.bSep08</a>	<a href="#">305348</a>	4735	735	1	54	lipoic acid synthetase (Lias) alternative variant bSep08, mRNA.
<a href="#">Lifr</a>	<a href="#">Lifr.bSep08</a>	<a href="#">81680</a>	29426	449	4	128	leukemia inhibitory factor receptor (Lifr) alternative variant bSep08, mRNA.
<a href="#">Lig1</a>	<a href="#">Lig1.bSep08</a>	<a href="#">81513</a>	1304	1208	1	48	ligase I, DNA, ATP-dependent (5.5 kD) (Lig1) alternative variant bSep08, mRNA.
<a href="#">Lig1</a>	<a href="#">Lig1.cSep08</a>	<a href="#">81513</a>	1080	428	2	24	ligase I, DNA, ATP-dependent (Lig1) alternative variant cSep08, mRNA.
<a href="#">Lig3</a>	<a href="#">Lig3.bSep08</a>	<a href="#">303369</a>	4029	3014	1	93	ligase III, DNA, ATP-dependent (Lig3) alternative variant bSep08, mRNA.
<a href="#">Lig4</a>	<a href="#">Lig4.aSep08</a>	<a href="#">290907</a>	8273	4764	2	952	ligase IV, DNA, ATP-dependent (Lig4) alternative variant aSep08, mRNA.
<a href="#">Lilrb3</a>	<a href="#">Lilrb3.bSep08</a>	<a href="#">361493</a>	42459	564	2	136	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 (Lilrb3) alternative variant bSep08, mRNA.

<a href="#">Lilrb4</a>	<a href="#">Lilrb4.bSep08</a>	<a href="#">292594</a>	5417	867	6	204	leukocyte immunoglobulin-like receptor, subfamily B, member 4 (Lilrb4) alternative variant bSep08, mRNA.
<a href="#">Lilrb4</a>	<a href="#">Lilrb4.cSep08</a>	<a href="#">292594</a>	2454	653	5	131	leukocyte immunoglobulin-like receptor, subfamily B, member 4 (Lilrb4) alternative variant cSep08, mRNA.
<a href="#">Lilrb4</a>	<a href="#">Lilrb4.dSep08</a>	<a href="#">292594</a>	2159	890	2	70	leukocyte immunoglobulin-like receptor, subfamily B, member 4 (8.1 kD) (Lilrb4) alternative variant dSep08, mRNA.
<a href="#">Lilrc1</a>	<a href="#">Lilrc1.bSep08</a>	<a href="#">691533</a>	3527	511	1	89	leukocyte immunoglobulin-like receptor (Lilrc1) alternative variant bSep08, mRNA.
<a href="#">Lilrc1</a>	<a href="#">Lilrc1.cSep08</a>	<a href="#">691533</a>	3672	501	2	85	leukocyte immunoglobulin-like receptor (Lilrc1) alternative variant cSep08, mRNA.
<a href="#">LIM.0</a>	<a href="#">LIM.0.aSep08</a>		8646	288		63	zinc finger protein 185 (LIM.0) mRNA.
<a href="#">LIM.1</a>	<a href="#">LIM.1.aSep08</a>		4190	765		234	lim domain actin binding 1 like (LIM.1) mRNA.
<a href="#">LIM.2</a>	<a href="#">LIM.2.aSep08</a>		16124	750	5	249	LIM, zinc-binding (LIM.2) alternative variant aSep08, mRNA.
<a href="#">LIM.2</a>	<a href="#">LIM.2.bSep08</a>		4647	688	5	229	putative protein of mammalian origin (LIM.2) alternative variant bSep08, mRNA.
<a href="#">LIM.2</a>	<a href="#">LIM.2.cSep08</a>		7868	604	5	169	LIM, zinc-binding (LIM.2) alternative variant cSep08, mRNA.
<a href="#">LIM.2</a>	<a href="#">LIM.2.dSep08</a>		13015	493	5	163	putative protein of vertebrate origin (LIM.2) alternative variant dSep08, mRNA.
<a href="#">LIM.3</a>	<a href="#">LIM.3.aSep08</a>		86628	683	5	227	actin-binding LIM protein 1 like (LIM.3) alternative variant aSep08, mRNA.
<a href="#">LIM.3</a>	<a href="#">LIM.3.bSep08</a>		1372	597	1	62	putative protein (LIM.3) alternative variant bSep08, mRNA.
<a href="#">Lim2</a>	<a href="#">Lim2.bSep08</a>	<a href="#">114903</a>	3210	503	1	88	lens intrinsic membrane protein 2 (Lim2) alternative variant bSep08, mRNA.
<a href="#">Lima1</a>	<a href="#">Lima1.aSep08</a>	<a href="#">300228</a>	59749	617		205	LIM domain and actin binding 1 (Lima1) mRNA.
<a href="#">Limd2</a>	<a href="#">Limd2.aSep08</a>	<a href="#">360646</a>	3146	970	6	128	LIM, zinc-binding (14.2 kD) (Limd2) alternative variant aSep08, mRNA.
<a href="#">Lime1</a>	<a href="#">Lime1.bSep08</a>	<a href="#">362289</a>	5692	4340	5	99	lck interacting transmembrane adaptor 1 (11.0 kD) (Lime1) alternative variant bSep08, mRNA.
<a href="#">Lime1</a>	<a href="#">Lime1.cSep08</a>	<a href="#">362289</a>	1316	1038	3	54	lck interacting transmembrane adaptor 1 (6.1 kD) (Lime1) alternative variant cSep08, mRNA.
<a href="#">Lime1</a>	<a href="#">Lime1.dSep08</a>	<a href="#">362289</a>	1398	749	3	43	lck interacting transmembrane adaptor 1 (Lime1) alternative variant dSep08, mRNA.
<a href="#">Limk1</a>	<a href="#">Limk1.aSep08</a>	<a href="#">65172</a>	26934	1556	11	518	LIM domain kinase 1 (Limk1) alternative variant aSep08, mRNA.
<a href="#">Limk1</a>	<a href="#">Limk1.bSep08</a>	<a href="#">65172</a>	6534	1026	5	200	LIM domain kinase 1 (23.2 kD) (Limk1) alternative variant bSep08, mRNA.
<a href="#">Limk2</a>	<a href="#">Limk2.bSep08</a>	<a href="#">29524</a>	15240	938	5	197	LIM motif-containing protein kinase 2 (Limk2) alternative variant bSep08, mRNA.
<a href="#">Limk2</a>	<a href="#">Limk2.cSep08</a>	<a href="#">29524</a>	3077	747	5	171	LIM motif-containing protein kinase 2 (Limk2) alternative variant cSep08, mRNA.
<a href="#">Lims1</a>	<a href="#">Lims1.aSep08</a>	<a href="#">499443</a>	105703	991	7	330	LIM and senescent cell antigen-like domains 1 (Lims1) alternative variant aSep08, mRNA.
<a href="#">Lims1</a>	<a href="#">Lims1.bSep08</a>	<a href="#">499443</a>	6063	318	1	81	LIM and senescent cell antigen-like domains 1 (Lims1) alternative variant bSep08, mRNA.



<a href="#">Lims1</a>	<a href="#">Lims1.cSep08</a>	<a href="#">499443</a>	51408	302	2	22	LIM and senescent cell antigen-like domains 1 (Lims1) alternative variant cSep08, mRNA.
<a href="#">Lims2</a>	<a href="#">Lims2.aSep08</a>	<a href="#">361303</a>	28243	1426	10	341	LIM and senescent cell antigen like domains 2 (39.0 kD) (Lims2) alternative variant aSep08, mRNA.
<a href="#">Lims2</a>	<a href="#">Lims2.dSep08</a>	<a href="#">361303</a>	1307	654	3	58	LIM and senescent cell antigen like domains 2 (Lims2) alternative variant dSep08, mRNA.
<a href="#">Lin7a</a>	<a href="#">Lin7a.bSep08</a>	<a href="#">85327</a>	43487	698	1	125	lin-7 homolog A (Lin7a) alternative variant bSep08, mRNA.
<a href="#">Lin7b</a>	<a href="#">Lin7b.aSep08</a>	<a href="#">60377</a>	2600	1576	2	212	lin-7 homolog C (22.7 kD) (Lin7b) alternative variant aSep08, mRNA.
<a href="#">Lin7b</a>	<a href="#">Lin7b.cSep08</a>	<a href="#">60377</a>	1316	384	2	52	lin-7 homolog B (Lin7b) alternative variant cSep08, mRNA.
<a href="#">Lin9</a>	<a href="#">Lin9.aSep08</a>	<a href="#">689523</a>	8952	348		116	lin-9 homolog (Lin9) mRNA.
<a href="#">Lin28</a>	<a href="#">Lin28.bSep08</a>	<a href="#">500562</a>	2191	406	1	94	lin-28 homolog (Lin28) alternative variant bSep08, mRNA.
<a href="#">Lin37</a>	<a href="#">Lin37.bSep08</a>	<a href="#">292787</a>	1429	570	2	163	lin-37 homolog (Lin37) alternative variant bSep08, mRNA.
<a href="#">Lin37</a>	<a href="#">Lin37.cSep08</a>	<a href="#">292787</a>	954	789	1	107	putative nuclear protein (11.7 kD) (Lin37) alternative variant cSep08, mRNA.
<a href="#">Lin52</a>	<a href="#">Lin52.aSep08</a>	<a href="#">362763</a>	85841	2119		111	lin-52 homolog (Lin52) mRNA.
<a href="#">Lingo1</a>	<a href="#">Lingo1.aSep08</a>	<a href="#">315691</a>	201756	2874	4	614	leucine-rich repeat, cysteine-rich flanking region, N-terminal and Leucine-rich repeat containing protein and immunoglobulin I-set (69.1 kD) (Lingo1) alternative variant aSep08, mRNA.
<a href="#">Lingo1</a>	<a href="#">Lingo1.bSep08</a>	<a href="#">315691</a>	14606	933	3	111	putative protein (11.5 kD) (Lingo1) alternative variant bSep08, mRNA.
<a href="#">Lingo1</a>	<a href="#">Lingo1.cSep08</a>	<a href="#">315691</a>	14308	543	2	103	leucine-rich repeat, cysteine-rich flanking region, N-terminal (Lingo1) alternative variant cSep08, mRNA.
<a href="#">Lingo1</a>	<a href="#">Lingo1.dSep08</a>	<a href="#">315691</a>	109454	400	3	74	putative protein (Lingo1) alternative variant dSep08, mRNA.
<a href="#">Lipa</a>	<a href="#">Lipa.bSep08</a>	<a href="#">25055</a>	19257	723	5	177	lysosomal acid lipase A (Lipa) alternative variant bSep08, mRNA.
<a href="#">Lipa</a>	<a href="#">Lipa.dSep08</a>	<a href="#">25055</a>	1590	775	2	67	lysosomal acid lipase A (7.9 kD) (Lipa) alternative variant dSep08, mRNA.
<a href="#">Lipc</a>	<a href="#">Lipc.bSep08</a>	<a href="#">24538</a>	107282	769	1	236	lipase, hepatic (Lipc) alternative variant bSep08, mRNA.
<a href="#">Lipe</a>	<a href="#">Lipe.bSep08</a>	<a href="#">25330</a>	4011	1117	3	240	lipase, hormone sensitive (Lipe) alternative variant bSep08, mRNA.
<a href="#">Lipe</a>	<a href="#">Lipe.cSep08</a>	<a href="#">25330</a>	7625	872	2	215	lipase, hormone sensitive (Lipe) alternative variant cSep08, mRNA.
<a href="#">Lipn</a>	<a href="#">Lipn.aSep08</a>	<a href="#">499345</a>	15390	975		325	lipase, family member N (Lipn) mRNA.
<a href="#">Lipocalin.0</a>	<a href="#">Lipocalin.0.aSep08</a>		3293	653		177	lipocalin 11 precursor (20.0 kD) (Lipocalin.0) mRNA.
<a href="#">Lix1</a>	<a href="#">Lix1.bSep08</a>	<a href="#">292381</a>	54219	1042	2	198	limb expression 1 homolog (chicken) (22.3 kD) (Lix1) alternative variant bSep08, mRNA.
<a href="#">Lix1</a>	<a href="#">Lix1.cSep08</a>	<a href="#">292381</a>	34786	982	2	172	limb expression 1 homolog (chicken) (Lix1) alternative variant cSep08, mRNA.
<a href="#">Lkap</a>	<a href="#">Lkap.bSep08</a>	<a href="#">170946</a>	3677	911	3	151	limkain b1 (Lkap) alternative variant bSep08, mRNA.
<a href="#">Lkap</a>	<a href="#">Lkap.cSep08</a>	<a href="#">170946</a>	610	439	2	131	limkain b1 (Lkap) alternative variant cSep08, mRNA.
<a href="#">Llgl1</a>	<a href="#">Llgl1.bSep08</a>	<a href="#">54265</a>	3047	2013	6	321	lethal giant larvae homolog 1 (Drosophila) (Llgl1) alternative variant bSep08, mRNA.

<a href="#">Llgl1</a>	<a href="#">Llgl1.cSep08</a>	<a href="#">54265</a>	1101	769	3	82	lethal giant larvae homolog 1 (Drosophila) (Llgl1) alternative variant cSep08, mRNA.
<a href="#">Llgl2</a>	<a href="#">Llgl2.bSep08</a>	<a href="#">360661</a>	1984	783	3	214	lethal giant larvae homolog 2 CRA c (Llgl2) alternative variant bSep08, mRNA.
<a href="#">Llgl2</a>	<a href="#">Llgl2.cSep08</a>	<a href="#">360661</a>	1968	679	5	211	lethal giant larvae homolog 2 CRA b (Llgl2) alternative variant cSep08, mRNA.
<a href="#">Llgl2</a>	<a href="#">Llgl2.dSep08</a>	<a href="#">360661</a>	22299	426	5	141	lethal giant larvae homolog 2 (Llgl2) alternative variant dSep08, mRNA.
<a href="#">Llgl2</a>	<a href="#">Llgl2.eSep08</a>	<a href="#">360661</a>	1550	612	6	116	lethal giant larvae homolog 2 CRA b (Llgl2) alternative variant eSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.bSep08</a>	<a href="#">116666</a>	18701	1777	6	514	lectin mannose-binding 1 CRA a like (Lman1) alternative variant bSep08, complete mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.cSep08</a>	<a href="#">116666</a>	9467	928	6	235	lectin mannose-binding 1 CRA a like (Lman1) alternative variant cSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.dSep08</a>	<a href="#">116666</a>	12158	2513	7	198	lectin mannose-binding 1 CRA a like (Lman1) alternative variant dSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.eSep08</a>	<a href="#">116666</a>	4757	354	4	84	lectin mannose-binding 1 like (Lman1) alternative variant eSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.fSep08</a>	<a href="#">116666</a>	2149	415	2	76	lectin mannose-binding 1 CRA a like (Lman1) alternative variant fSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.gSep08</a>	<a href="#">116666</a>	1688	303	3	54	putative protein (5.9 kD) (Lman1) alternative variant gSep08, mRNA.
<a href="#">Lman1</a>	<a href="#">Lman1.hSep08</a>	<a href="#">116666</a>	375	275	2	32	lectin mannose-binding 1 CRA a like (Lman1) alternative variant hSep08, mRNA.
<a href="#">Lman1l</a>	<a href="#">Lman1l.bSep08</a>	<a href="#">300743</a>	3734	629	1	172	lectin, mannose-binding, 1 like (Lman1l) alternative variant bSep08, mRNA.
<a href="#">Lman2</a>	<a href="#">Lman2.bSep08</a>	<a href="#">290994</a>	1730	528	2	106	lectin, mannose-binding 2 (12.1 kD) (Lman2) alternative variant bSep08, mRNA.
<a href="#">Lman2l</a>	<a href="#">Lman2l.aSep08</a>	<a href="#">301343</a>	19257	806	6	261	lectin mannose-binding 2-like (Lman2l) alternative variant aSep08, mRNA.
<a href="#">Lman2l</a>	<a href="#">Lman2l.bSep08</a>	<a href="#">301343</a>	6072	2477	4	171	lectin mannose-binding 2-like CRA c (19.8 kD) (Lman2l) alternative variant bSep08, mRNA.
<a href="#">Lman2l</a>	<a href="#">Lman2l.cSep08</a>	<a href="#">301343</a>	24166	1810	5	133	putative mitochondrial protein of metazoan origin (Lman2l) alternative variant cSep08, complete mRNA.
<a href="#">Lmbr1</a>	<a href="#">Lmbr1.aSep08</a>	<a href="#">362295</a>	98993	4281	13	367	limb region 1 (Lmbr1) alternative variant aSep08, mRNA.
<a href="#">Lmbr1</a>	<a href="#">Lmbr1.bSep08</a>	<a href="#">362295</a>	101390	622	4	158	limb region 1 (Lmbr1) alternative variant bSep08, mRNA.
<a href="#">LMBR1.1</a>	<a href="#">LMBR1.1.aSep08</a>		4531	695	8	231	limb region 1 like CRA b (LMBR1.1) alternative variant aSep08, mRNA.
<a href="#">Lmbr1l</a>	<a href="#">Lmbr1l.bSep08</a>	<a href="#">300215</a>	2960	716	3	88	limb region 1-like homolog (mouse) (9.9 kD) (Lmbr1l) alternative variant bSep08, mRNA.
<a href="#">Lmbrd1</a>	<a href="#">Lmbrd1.bSep08</a>	<a href="#">246046</a>	8387	415	1	54	putative protein (6.0 kD) (Lmbrd1) alternative variant bSep08, mRNA.
<a href="#">Lmbrd1</a>	<a href="#">Lmbrd1.cSep08</a>	<a href="#">246046</a>	31536	541	2		
<a href="#">Lmcd1</a>	<a href="#">Lmcd1.bSep08</a>	<a href="#">494021</a>	58811	831	2	144	LIM and cysteine-rich domains 1 (16.1 kD) (Lmcd1) alternative variant bSep08, mRNA.
<a href="#">Lmf1</a>	<a href="#">Lmf1.aSep08</a>	<a href="#">360495</a>	69476	805		51	lipase maturation factor 1 (Lmf1) mRNA.

<a href="#">Lmf2</a>	<a href="#">Lmf2.bSep08</a>	<a href="#">315218</a>	1143	743	6	247	lipase maturation factor 2 (Lmf2) alternative variant bSep08, mRNA.
<a href="#">Lmf2</a>	<a href="#">Lmf2.cSep08</a>	<a href="#">315218</a>	1029	388	3	129	lipase maturation factor 2 (Lmf2) alternative variant cSep08, mRNA.
<a href="#">Lmf2</a>	<a href="#">Lmf2.dSep08</a>	<a href="#">315218</a>	1022	498	2	127	lipase maturation factor 2 (Lmf2) alternative variant dSep08, mRNA.
<a href="#">Lmf2</a>	<a href="#">Lmf2.fSep08</a>	<a href="#">315218</a>	749	504	2	88	lipase maturation factor 2 (Lmf2) alternative variant fSep08, mRNA.
<a href="#">Lmna</a>	<a href="#">Lmna.bSep08</a>	<a href="#">60374</a>	3374	1353	6	277	lamin A (Lmna) alternative variant bSep08, mRNA.
<a href="#">Lmna</a>	<a href="#">Lmna.dSep08</a>	<a href="#">60374</a>	754	505	2	98	lamin A (Lmna) alternative variant dSep08, mRNA.
<a href="#">Lmnb1</a>	<a href="#">Lmnb1.bSep08</a>	<a href="#">116685</a>	1578	706	2	116	lamin B1 (Lmnb1) alternative variant bSep08, mRNA.
<a href="#">Lmnb2</a>	<a href="#">Lmnb2.aSep08</a>	<a href="#">299625</a>	16104	3632	4	616	lamin B2 (68.8 kD) (Lmnb2) alternative variant aSep08, mRNA.
<a href="#">Lmnb2</a>	<a href="#">Lmnb2.bSep08</a>	<a href="#">299625</a>	1514	699	1	233	lamin B2 (Lmnb2) alternative variant bSep08, mRNA.
<a href="#">Lmo2</a>	<a href="#">Lmo2.aSep08</a>	<a href="#">362176</a>	10623	714	3	237	LIM domain only 2 (Lmo2) alternative variant aSep08, mRNA.
<a href="#">Lmo2</a>	<a href="#">Lmo2.bSep08</a>	<a href="#">362176</a>	22034	829	3	232	LIM domain only 2 (25.7 kD) (Lmo2) alternative variant bSep08, mRNA.
<a href="#">Lmo2</a>	<a href="#">Lmo2.dSep08</a>	<a href="#">362176</a>	11459	1782	2	158	LIM domain only 2 (18.3 kD) (Lmo2) alternative variant dSep08, mRNA.
<a href="#">Lmo2</a>	<a href="#">Lmo2.eSep08</a>	<a href="#">362176</a>	11766	1393	3	138	LIM domain only 2 (Lmo2) alternative variant eSep08, mRNA.
<a href="#">Lmo2</a>	<a href="#">Lmo2.fSep08</a>	<a href="#">362176</a>	5119	695	1	100	LIM domain only 2 (Lmo2) alternative variant fSep08, mRNA.
<a href="#">Lmo4</a>	<a href="#">Lmo4.aSep08</a>	<a href="#">362051</a>	12281	905	4	200	LIM domain (Lmo4) alternative variant aSep08, mRNA.
<a href="#">Lmo4</a>	<a href="#">Lmo4.cSep08</a>	<a href="#">362051</a>	14056	1183	5	165	LIM domain (18.0 kD) (Lmo4) alternative variant cSep08, mRNA.
<a href="#">Lmo4</a>	<a href="#">Lmo4.dSep08</a>	<a href="#">362051</a>	1115	455	3	81	LIM domain (8.8 kD) (Lmo4) alternative variant dSep08, mRNA.
<a href="#">Lmo4</a>	<a href="#">Lmo4.eSep08</a>	<a href="#">362051</a>	2228	1064	2	63	putative protein (7.1 kD) (Lmo4) alternative variant eSep08, mRNA.
<a href="#">Lmo7</a>	<a href="#">Lmo7.bSep08</a>	<a href="#">361084</a>	44132	785	5	187	LIM domain only 7 (Lmo7) alternative variant bSep08, mRNA.
<a href="#">Lmo7</a>	<a href="#">Lmo7.cSep08</a>	<a href="#">361084</a>	10889	738	3	140	LIM domain only 7 (Lmo7) alternative variant cSep08, mRNA.
<a href="#">Lmod1</a>	<a href="#">Lmod1.aSep08</a>	<a href="#">304816</a>	38920	679		179	leiomodoin 1 (smooth muscle) (Lmod1) mRNA.
<a href="#">Lmod3</a>	<a href="#">Lmod3.aSep08</a>	<a href="#">500267</a>	10359	1847		441	leiomodoin 3 (fetal) (Lmod3) mRNA.
<a href="#">Lmtk2</a>	<a href="#">Lmtk2.aSep08</a>	<a href="#">304286</a>	32792	437		82	lemur tyrosine kinase 2 (Lmtk2) mRNA.
<a href="#">Lmx1a</a>	<a href="#">Lmx1a.bSep08</a>	<a href="#">289201</a>	108387	572	3	190	LIM homeobox transcription factor 1 alpha (Lmx1a) alternative variant bSep08, mRNA.
<a href="#">Lmx1a</a>	<a href="#">Lmx1a.cSep08</a>	<a href="#">289201</a>	96885	475	2	46	LIM homeobox transcription factor 1 alpha (5.2 kD) (Lmx1a) alternative variant cSep08, mRNA.
<a href="#">Lmx1b</a>	<a href="#">Lmx1b.aSep08</a>	<a href="#">114501</a>	78061	769		255	LIM homeobox transcription factor 1 beta (Lmx1b) mRNA.
<a href="#">Lnp</a>	<a href="#">Lnp.bSep08</a>	<a href="#">362151</a>	8686	1056	2	102	limb and neural patterns (Lnp) alternative variant bSep08, mRNA.

<a href="#">Lnp</a>	<a href="#">Lnp.cSep08</a>	<a href="#">362151</a>	9786	383	4	75	limb and neural patterns (Lnp) alternative variant cSep08, mRNA.
<a href="#">Lnx1</a>	<a href="#">Lnx1.bSep08</a>	<a href="#">360926</a>	29474	1200	6	354	ligand of numb-protein X 1 (Lnx1) alternative variant bSep08, mRNA.
<a href="#">Lnx1</a>	<a href="#">Lnx1.cSep08</a>	<a href="#">360926</a>	48607	536	2	103	ligand of numb-protein X 1 (Lnx1) alternative variant cSep08, mRNA.
<a href="#">Lnx1</a>	<a href="#">Lnx1.dSep08</a>	<a href="#">360926</a>	1772	1426	2	43	ligand of numb-protein X 1 (4.9 kD) (Lnx1) alternative variant dSep08, mRNA.
<a href="#">Lnx2</a>	<a href="#">Lnx2.bSep08</a>	<a href="#">360761</a>	36046	402	1	73	ligand of numb-protein X 2 (Lnx2) alternative variant bSep08, mRNA.
<a href="#">loby</a>	<a href="#">loby.aSep08</a>		2805	1749	1	333	motif Sec7 domain-containing protein 2 like (loby) mRNA.
<a href="#">LOC56764</a>	<a href="#">LOC56764.bSep08</a>	<a href="#">56764</a>	37394	1497	3	162	dnaj-like protein (LOC56764) alternative variant bSep08, mRNA.
<a href="#">LOC224733I-4</a>	<a href="#">LOC224733I-4.aSep08</a>	<a href="#">415070</a>	1186	686		117	LOC224733I-4 pseudogene (12.7 kD) (LOC224733I-4) mRNA.
<a href="#">LOC224733I-6</a>	<a href="#">LOC224733I-6.aSep08</a>	<a href="#">415076</a>	20984	564		25	LOC224733I-6 pseudogene (3.2 kD) (LOC224733I-6) mRNA.
<a href="#">LOC245925</a>	<a href="#">LOC245925.aSep08</a>	<a href="#">245925</a>	3096	2024		607	CTD-binding SR-like protein rA9 (LOC245925) alternative variant aSep08, mRNA.
<a href="#">LOC245925</a>	<a href="#">LOC245925.bSep08</a>	<a href="#">245925</a>	4219	1776		537	CTD-binding SR-like protein rA9 (LOC245925) alternative variant bSep08, mRNA.
<a href="#">LOC246295</a>	<a href="#">LOC246295.aSep08</a>	<a href="#">246295</a>	14313	1797		517	glycine-, glutamate-, thienylcyclohexylpiperidine-binding protein (LOC246295) alternative variant aSep08, mRNA.
<a href="#">LOC246295</a>	<a href="#">LOC246295.bSep08</a>	<a href="#">246295</a>	19261	3365	2	390	glycine-, glutamate-, thienylcyclohexylpiperidine-binding protein (LOC246295) alternative variant bSep08, mRNA.
<a href="#">LOC246295</a>	<a href="#">LOC246295.cSep08</a>	<a href="#">246295</a>	10551	952	3	173	glycine-, glutamate-, thienylcyclohexylpiperidine-binding protein (LOC246295) alternative variant cSep08, mRNA.
<a href="#">LOC246295</a>	<a href="#">LOC246295.dSep08</a>	<a href="#">246295</a>	6606	779	4	128	glycine-, glutamate-, thienylcyclohexylpiperidine-binding protein (LOC246295) alternative variant dSep08, mRNA.
<a href="#">LOC246295</a>	<a href="#">LOC246295.eSep08</a>	<a href="#">246295</a>	9674	992		147	glycine-, glutamate-, thienylcyclohexylpiperidine-binding protein (LOC246295) alternative variant eSep08, complete mRNA.
<a href="#">LOC252890</a>	<a href="#">LOC252890.aSep08</a>	<a href="#">252890</a>	1870	1124	2	63	z39 small nucleolar RNA (7.4 kD) (LOC252890) alternative variant aSep08, mRNA.
<a href="#">LOC257643</a>	<a href="#">LOC257643.bSep08</a>	<a href="#">257643</a>	1858	317	1	38	cystatin SC (LOC257643) alternative variant bSep08, mRNA.
<a href="#">LOC257650</a>	<a href="#">LOC257650.aSep08</a>	<a href="#">257650</a>	12472	741	2	140	hippyrgranin (LOC257650) alternative variant aSep08, mRNA.
<a href="#">LOC257650</a>	<a href="#">LOC257650.bSep08</a>	<a href="#">257650</a>	12170	533	3	97	hippyrgranin (LOC257650) alternative variant bSep08, mRNA.
<a href="#">LOC257650</a>	<a href="#">LOC257650.cSep08</a>	<a href="#">257650</a>	24459	1351	5	140	hippyrgranin (LOC257650) alternative variant cSep08, mRNA.
<a href="#">LOC259246</a>	<a href="#">LOC259246.aSep08</a>	<a href="#">259246</a>	3205	768	2	188	globulin PGCL1 (LOC259246) alternative variant aSep08, mRNA.
<a href="#">LOC259246</a>	<a href="#">LOC259246.aSep08</a>	<a href="#">298116</a>	3205	768	2	188	globulin PGCL1 (LOC259246) alternative variant aSep08, mRNA.

<a href="#">LOC259246</a>	<a href="#">LOC259246.dSep08</a>	<a href="#">259246</a>	3147	676	2	181	globulin PGCL1 precursor (20.7 kD) (LOC259246) alternative variant dSep08, mRNA.
<a href="#">LOC259246</a>	<a href="#">LOC259246.dSep08</a>	<a href="#">298116</a>	3147	676	2	181	globulin PGCL1 precursor (20.7 kD) (LOC259246) alternative variant dSep08, mRNA.
<a href="#">LOC287228</a>	<a href="#">LOC287228.aSep08</a>	<a href="#">287228</a>	2762	814		173	similar to GTPase activating protein testicular GAP1 (LOC287228) mRNA.
<a href="#">LOC287274</a>	<a href="#">LOC287274.aSep08</a>	<a href="#">287274</a>	4166	974	1	140	sedlin-like (16.5 kD) (LOC287274) alternative variant aSep08, complete mRNA.
<a href="#">LOC287274</a>	<a href="#">LOC287274.bSep08</a>	<a href="#">287274</a>	4254	864	1	140	sedlin-like (16.5 kD) (LOC287274) alternative variant bSep08, mRNA.
<a href="#">LOC288526</a>	<a href="#">LOC288526.bSep08</a>	<a href="#">288526</a>	4017	1596	4	307	putative protein, with 3 coiled coil domains (LOC288526) alternative variant bSep08, mRNA.
<a href="#">LOC288913</a>	<a href="#">LOC288913.bSep08</a>	<a href="#">288913</a>	3023	847	1	77	similar to LEYDIG CELL TUMOR 10 KD PROTEIN (LOC288913) alternative variant bSep08, mRNA.
<a href="#">LOC288978</a>	<a href="#">LOC288978.aSep08</a>	<a href="#">288978</a>	65584	515	6	144	hypothetical LOC288978 (LOC288978) alternative variant aSep08, mRNA.
<a href="#">LOC288978</a>	<a href="#">LOC288978.bSep08</a>	<a href="#">288978</a>	64667	527	6	126	hypothetical LOC288978 (14.7 kD) (LOC288978) alternative variant bSep08, mRNA.
<a href="#">LOC288978</a>	<a href="#">LOC288978.dSep08</a>	<a href="#">288978</a>	24295	400	5	105	hypothetical LOC288978 (LOC288978) alternative variant dSep08, mRNA.
<a href="#">LOC288978</a>	<a href="#">LOC288978.eSep08</a>	<a href="#">288978</a>	12204	256	3	85	hypothetical LOC288978 (LOC288978) alternative variant eSep08, mRNA.
<a href="#">LOC288978</a>	<a href="#">LOC288978.fSep08</a>	<a href="#">288978</a>	35324	316	3	79	hypothetical LOC288978 (LOC288978) alternative variant fSep08, mRNA.
<a href="#">LOC289035</a>	<a href="#">LOC289035.bSep08</a>	<a href="#">289035</a>	2904	328	1	40	similar to UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-acetylgluco (LOC289035) alternative variant bSep08, mRNA.
<a href="#">LOC289378</a>	<a href="#">LOC289378.bSep08</a>	<a href="#">289378</a>	6843	964	2	117	similar to B0432.8 (12.9 kD) (LOC289378) alternative variant bSep08, mRNA.
<a href="#">LOC289471</a>	<a href="#">LOC289471.aSep08</a>	<a href="#">289471</a>	4527	714		128	similar to DNA helicase HEL308 (LOC289471) mRNA.
<a href="#">LOC289606</a>	<a href="#">LOC289606.aSep08</a>	<a href="#">289606</a>	94196	776		258	similar to Gamma-aminobutyric-acid receptor alpha-2 subunit precursor (GABA(A) receptor) (LOC289606) mRNA.
<a href="#">LOC289740</a>	<a href="#">LOC289740.bSep08</a>	<a href="#">289740</a>	8883	921	6	239	pescadillo (27.5 kD) (LOC289740) alternative variant bSep08, mRNA.
<a href="#">LOC289740</a>	<a href="#">LOC289740.cSep08</a>	<a href="#">289740</a>	5847	755	4	184	pescadillo (LOC289740) alternative variant cSep08, mRNA.
<a href="#">LOC289740</a>	<a href="#">LOC289740.dSep08</a>	<a href="#">289740</a>	1674	959	2	77	pescadillo homolog 1 containing BRCT domain (9.2 kD) (LOC289740) alternative variant dSep08, mRNA.
<a href="#">LOC290577</a>	<a href="#">LOC290577.aSep08</a>	<a href="#">290577</a>	14807	2156	9	333	wings apart-like homolog CRA b (37.2 kD) (LOC290577) alternative variant aSep08, mRNA.
<a href="#">LOC290577</a>	<a href="#">LOC290577.bSep08</a>	<a href="#">290577</a>	11653	1183	7	332	wings apart-like homolog (LOC290577) alternative variant bSep08, mRNA.
<a href="#">LOC290577</a>	<a href="#">LOC290577.cSep08</a>	<a href="#">290577</a>	33610	1011	6	133	wings apart-like homolog (LOC290577) alternative variant cSep08, mRNA.
<a href="#">LOC290577</a>	<a href="#">LOC290577.eSep08</a>	<a href="#">290577</a>	6619	470	3	64	wings apart-like homolog (LOC290577) alternative variant eSep08, mRNA.

<a href="#">LOC290763</a>	<a href="#">LOC290763.aSep08</a>	<a href="#">290763</a>	2549	784		162	similar to testase 4 (LOC290763) mRNA.
<a href="#">LOC290876</a>	<a href="#">LOC290876.bSep08</a>	<a href="#">290876</a>	8010	547	7	170	similar to RIKEN cDNA 1700029H14 (LOC290876) alternative variant bSep08, mRNA.
<a href="#">LOC290876</a>	<a href="#">LOC290876.dSep08</a>	<a href="#">290876</a>	8198	739	7	90	similar to RIKEN cDNA 1700029H14 (LOC290876) alternative variant dSep08, mRNA.
<a href="#">LOC291249</a>	<a href="#">LOC291249.aSep08</a>	<a href="#">291249</a>	6877	732		244	testis-specific transcript (LOC291249) mRNA.
<a href="#">LOC291863</a>	<a href="#">LOC291863.bSep08</a>	<a href="#">291863</a>	9531	742	6	247	carboxylesterase (LOC291863) alternative variant bSep08, mRNA.
<a href="#">LOC291863</a>	<a href="#">LOC291863.cSep08</a>	<a href="#">291863</a>	3524	359	2	86	carboxylesterase precursor (9.4 kD) (LOC291863) alternative variant cSep08, mRNA.
<a href="#">LOC291863</a>	<a href="#">LOC291863.dSep08</a>	<a href="#">291863</a>	3529	876	2	73	carboxylesterase (8.4 kD) (LOC291863) alternative variant dSep08, mRNA.
<a href="#">LOC291863</a>	<a href="#">LOC291863.eSep08</a>	<a href="#">291863</a>	4332	658	3	51	putative protein (5.9 kD) (LOC291863) alternative variant eSep08, mRNA.
<a href="#">LOC292069</a>	<a href="#">LOC292069.bSep08</a>	<a href="#">292069</a>	1342	727	3	148	similar to RIKEN cDNA 2310061F22 (LOC292069) alternative variant bSep08, mRNA.
<a href="#">LOC292069</a>	<a href="#">LOC292069.cSep08</a>	<a href="#">292069</a>	865	761	1	123	similar to RIKEN cDNA 2310061F22 (LOC292069) alternative variant cSep08, mRNA.
<a href="#">LOC292282</a>	<a href="#">LOC292282.aSep08</a>	<a href="#">292282</a>	884	406	2	128	similar to USE1-like protein (Hematopoietic stem/progenitor cells protein MDS032) (Putative MAPK-activating protein PM26) (Protein p31) (LOC292282) alternative variant aSep08, mRNA.
<a href="#">LOC292282</a>	<a href="#">LOC292282.bSep08</a>	<a href="#">292282</a>	971	749	1	84	similar to USE1-like protein (Hematopoietic stem/progenitor cells protein MDS032) (Putative MAPK-activating protein PM26) (Protein p31) (9.7 kD) (LOC292282) alternative variant bSep08, mRNA.
<a href="#">LOC292666</a>	<a href="#">LOC292666.aSep08</a>	<a href="#">292666</a>	16268	1783	4	541	similar to pregnancy-specific beta 1-glycoprotein (LOC292666) alternative variant aSep08, mRNA.
<a href="#">LOC292666</a>	<a href="#">LOC292666.cSep08</a>	<a href="#">292666</a>	16202	682	2	203	similar to pregnancy-specific beta 1-glycoprotein (LOC292666) alternative variant cSep08, mRNA.
<a href="#">LOC292722</a>	<a href="#">LOC292722.aSep08</a>	<a href="#">292722</a>	4993	1244	13	414	similar to C15A7.2 (LOC292722) alternative variant aSep08, mRNA.
<a href="#">LOC292722</a>	<a href="#">LOC292722.bSep08</a>	<a href="#">292722</a>	1034	414	5	138	similar to C15A7.2 (LOC292722) alternative variant bSep08, mRNA.
<a href="#">LOC293103</a>	<a href="#">LOC293103.aSep08</a>	<a href="#">293103</a>	18462	1775	3	412	similar to RIKEN cDNA 0610007P06 (LOC293103) alternative variant aSep08, mRNA.
<a href="#">LOC293103</a>	<a href="#">LOC293103.bSep08</a>	<a href="#">293103</a>	23947	1573	3	197	similar to RIKEN cDNA 0610007P06 (21.6 kD) (LOC293103) alternative variant bSep08, mRNA.
<a href="#">LOC293103</a>	<a href="#">LOC293103.cSep08</a>	<a href="#">293103</a>	21956	736	2	180	similar to RIKEN cDNA 0610007P06 (19.4 kD) (LOC293103) alternative variant cSep08, mRNA.
<a href="#">LOC293989</a>	<a href="#">LOC293989.bSep08</a>	<a href="#">293989</a>	20439	416	3	138	p450 2c2 (LOC293989) alternative variant bSep08, mRNA.
<a href="#">LOC293989</a>	<a href="#">LOC293989.cSep08</a>	<a href="#">293989</a>	3302	558	3	123	CRA a like (LOC293989) alternative variant cSep08, mRNA.
<a href="#">LOC293989</a>	<a href="#">LOC293989.dSep08</a>	<a href="#">293989</a>	7947	836	4	107	putative cytoplasmic protein of vertebrate origin (12.5 kD) (LOC293989) alternative variant dSep08, mRNA.
<a href="#">LOC293989</a>	<a href="#">LOC293989.eSep08</a>	<a href="#">293989</a>	17350	263	2	77	cytochrome p450 family 2 subfamily c polypeptide (LOC293989) alternative variant eSep08, mRNA.

<a href="#">LOC294154</a>	<a href="#">LOC294154.aSep08</a>	<a href="#">294154</a>	66261	2007	1	291	putative protein of ancient origin (LOC294154) alternative variant aSep08, mRNA.
<a href="#">LOC294154</a>	<a href="#">LOC294154.bSep08</a>	<a href="#">294154</a>	65764	1181		244	putative protein of ancient origin (LOC294154) alternative variant bSep08, mRNA.
<a href="#">LOC294887</a>	<a href="#">LOC294887.aSep08</a>	<a href="#">294887</a>	1215	275		91	similar to GTPase activating protein testicular GAP1 (LOC294887) mRNA.
<a href="#">LOC295452</a>	<a href="#">LOC295452.aSep08</a>	<a href="#">295452</a>	67948	919		66	similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (LOC295452) mRNA.
<a href="#">LOC295528</a>	<a href="#">LOC295528.aSep08</a>	<a href="#">295528</a>	17133	689		229	similar to T06D8.1a (LOC295528) mRNA.
<a href="#">LOC296637</a>	<a href="#">LOC296637.bSep08</a>	<a href="#">296637</a>	948	392	2	130	similar to HLA-B associated transcript-2 isoform a (LOC296637) alternative variant bSep08, mRNA.
<a href="#">LOC298018</a>	<a href="#">LOC298018.aSep08</a>	<a href="#">298018</a>	6029	2997		210	hypothetical LOC298018 (LOC298018) mRNA.
<a href="#">LOC298109</a>	<a href="#">LOC298109.aSep08</a>	<a href="#">298109</a>	3256	823	1	181	alpha-2u globulin PGCL2 (20.7 kD) (LOC298109) alternative variant aSep08, mRNA.
<a href="#">LOC298138</a>	<a href="#">LOC298138.aSep08</a>	<a href="#">298138</a>	31893	1261	3	420	putative protein, with a coiled coil domain, of metazoan origin (LOC298138) alternative variant aSep08, mRNA.
<a href="#">LOC298138</a>	<a href="#">LOC298138.bSep08</a>	<a href="#">298138</a>	31534	1236	3	194	putative nuclear protein, with a coiled coil domain, of metazoan origin (22.7 kD) (LOC298138) alternative variant bSep08, mRNA.
<a href="#">LOC298250</a>	<a href="#">LOC298250.bSep08</a>	<a href="#">298250</a>	52380	630	5	185	similar to hypothetical protein FLJ10986 (LOC298250) alternative variant bSep08, mRNA.
<a href="#">LOC298442</a>	<a href="#">LOC298442.bSep08</a>	<a href="#">298442</a>	4226	718	6	158	putative protein, with 2 coiled coil domains, of mammalian origin (LOC298442) alternative variant bSep08, mRNA.
<a href="#">LOC298442</a>	<a href="#">LOC298442.cSep08</a>	<a href="#">298442</a>	390	285	2	94	putative protein of mammalian origin (LOC298442) alternative variant cSep08, mRNA.
<a href="#">LOC298442</a>	<a href="#">LOC298442.dSep08</a>	<a href="#">298442</a>	1390	1093	2	63	putative protein (LOC298442) alternative variant dSep08, mRNA.
<a href="#">LOC298442</a>	<a href="#">LOC298442.eSep08</a>	<a href="#">298442</a>	1120	437	2	62	putative protein of mammalian origin (7.0 kD) (LOC298442) alternative variant eSep08, mRNA.
<a href="#">LOC298977</a>	<a href="#">LOC298977.aSep08</a>	<a href="#">298977</a>	47378	1285		361	similar to zinc finger protein 277 isoform 1 (LOC298977) mRNA.
<a href="#">LOC299282</a>	<a href="#">LOC299282.bSep08</a>	<a href="#">299282</a>	7357	1668	5	413	serine protease inhibitor (46.3 kD) (LOC299282) alternative variant bSep08, complete mRNA.
<a href="#">LOC299282</a>	<a href="#">LOC299282.cSep08</a>	<a href="#">299282</a>	4705	930	3	273	serine protease inhibitor (LOC299282) alternative variant cSep08, mRNA.
<a href="#">LOC299282</a>	<a href="#">LOC299282.dSep08</a>	<a href="#">299282</a>	4458	815	4	216	serine protease inhibitor (LOC299282) alternative variant dSep08, mRNA.
<a href="#">LOC299282</a>	<a href="#">LOC299282.fSep08</a>	<a href="#">299282</a>	1498	246	3	81	serine protease inhibitor (LOC299282) alternative variant fSep08, mRNA.
<a href="#">LOC300024</a>	<a href="#">LOC300024.aSep08</a>	<a href="#">300024</a>	2427	653		103	similar to Ly6-B antigen gene (LOC300024) mRNA.
<a href="#">LOC300191</a>	<a href="#">LOC300191.bSep08</a>	<a href="#">300191</a>	3039	2517	2	89	similar to RIKEN cDNA 4930570C03 (10.6 kD) (LOC300191) alternative variant bSep08, mRNA.
<a href="#">LOC300225</a>	<a href="#">LOC300225.aSep08</a>	<a href="#">300225</a>	38417	672		200	similar to formin-like 3 protein (LOC300225) mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.aSep08</a>	<a href="#">300308</a>	205409	630	2	84	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant aSep08, mRNA.

<a href="#">LOC300308</a>	<a href="#">LOC300308.aSep08</a>	<a href="#">500815</a>	205409	630	2	84	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant aSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.cSep08</a>	<a href="#">300308</a>	6648	1474	5	183	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant cSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.cSep08</a>	<a href="#">500815</a>	6648	1474	5	183	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant cSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.dSep08</a>	<a href="#">300308</a>	5066	752	2	35	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (4.1 kD) (LOC300308) alternative variant dSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.dSep08</a>	<a href="#">500815</a>	5066	752	2	35	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (4.1 kD) (LOC300308) alternative variant dSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.eSep08</a>	<a href="#">300308</a>	4242	272	2	39	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant eSep08, mRNA.
<a href="#">LOC300308</a>	<a href="#">LOC300308.eSep08</a>	<a href="#">500815</a>	4242	272	2	39	similar to hypothetical protein 4930509O22 and similar to par-1 CG8201-PO, isoform O (LOC300308) alternative variant eSep08, mRNA.
<a href="#">LOC300314</a>	<a href="#">LOC300314.aSep08</a>	<a href="#">300314</a>	7778	934		277	similar to Protein KIAA0196 (LOC300314) mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.bSep08</a>	<a href="#">300963</a>	43332	2241	13	538	similar to centrosome protein Cep63 (62.7 kD) (LOC300963) alternative variant bSep08, complete mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.cSep08</a>	<a href="#">300963</a>	39633	1588	13	516	similar to centrosome protein Cep63 (LOC300963) alternative variant cSep08, mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.dSep08</a>	<a href="#">300963</a>	11350	955	4	213	similar to centrosome protein Cep63 (LOC300963) alternative variant dSep08, mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.eSep08</a>	<a href="#">300963</a>	22659	766	6	183	similar to centrosome protein Cep63 (LOC300963) alternative variant eSep08, mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.fSep08</a>	<a href="#">300963</a>	27749	799	6	140	similar to centrosome protein Cep63 (LOC300963) alternative variant fSep08, mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.gSep08</a>	<a href="#">300963</a>	2193	515	2	82	similar to centrosome protein Cep63 (LOC300963) alternative variant gSep08, mRNA.
<a href="#">LOC300963</a>	<a href="#">LOC300963.iSep08</a>	<a href="#">300963</a>	633	279	2	82	similar to centrosome protein Cep63 (LOC300963) alternative variant iSep08, mRNA.
<a href="#">LOC301124</a>	<a href="#">LOC301124.aSep08</a>	<a href="#">301124</a>	2076	1156	3	130	hypothetical LOC301124 (14.3 kD) (LOC301124) alternative variant aSep08, mRNA.
<a href="#">LOC301124</a>	<a href="#">LOC301124.cSep08</a>	<a href="#">301124</a>	1977	496	5	98	hypothetical LOC301124 (11.2 kD) (LOC301124) alternative variant cSep08, mRNA.
<a href="#">LOC301124</a>	<a href="#">LOC301124.dSep08</a>	<a href="#">301124</a>	1926	1428	2	71	hypothetical LOC301124 (7.9 kD) (LOC301124) alternative variant dSep08, complete mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.aSep08</a>	<a href="#">301126</a>	21114	2988	18	923	scaffold attachment factor B2 (LOC301126) alternative variant aSep08, mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.bSep08</a>	<a href="#">301126</a>	11978	1913	13	492	scaffold attachment factor B2 (58.5 kD) (LOC301126) alternative variant bSep08, mRNA.



<a href="#">LOC301126</a>	<a href="#">LOC301126.cSep08</a>	<a href="#">301126</a>	4193	1128	5	257	scaffold attachment factor B2 (31.5 kD) (LOC301126) alternative variant cSep08, mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.eSep08</a>	<a href="#">301126</a>	1662	691	3	148	CRA b like (LOC301126) alternative variant eSep08, mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.fSep08</a>	<a href="#">301126</a>	2382	1216	2	103	scaffold attachment factor B2 (11.2 kD) (LOC301126) alternative variant fSep08, mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.gSep08</a>	<a href="#">301126</a>	4741	391	3	90	scaffold attachment factor B2 CRA b (LOC301126) alternative variant gSep08, mRNA.
<a href="#">LOC301126</a>	<a href="#">LOC301126.hSep08</a>	<a href="#">301126</a>	1564	566	3	82	scaffold attachment factor B2 CRA b (LOC301126) alternative variant hSep08, mRNA.
<a href="#">LOC301128</a>	<a href="#">LOC301128.bSep08</a>	<a href="#">301128</a>	3225	777	5	112	putative protein (LOC301128) alternative variant bSep08, mRNA.
<a href="#">LOC301128</a>	<a href="#">LOC301128.cSep08</a>	<a href="#">301128</a>	1863	414	3	58	putative protein of metazoan origin (LOC301128) alternative variant cSep08, mRNA.
<a href="#">LOC301165</a>	<a href="#">LOC301165.aSep08</a>	<a href="#">301165</a>	2203	1769	2	512	similar to hypothetical protein 4932415M13 (57.3 kD) (LOC301165) alternative variant aSep08, mRNA.
<a href="#">LOC301455</a>	<a href="#">LOC301455.aSep08</a>	<a href="#">301455</a>	3016	394		130	par-3 Partitioning defective 3 homolog b (LOC301455) mRNA.
<a href="#">LOC301977</a>	<a href="#">LOC301977.aSep08</a>	<a href="#">301977</a>	4668	838		174	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC301977) mRNA.
<a href="#">LOC302192</a>	<a href="#">LOC302192.aSep08</a>	<a href="#">302192</a>	7638	1391		202	similar to RIKEN cDNA 1700001E04 (24.1 kD) (LOC302192) complete mRNA.
<a href="#">LOC302199</a>	<a href="#">LOC302199.aSep08</a>	<a href="#">302199</a>	1946	468		119	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC302199) mRNA.
<a href="#">LOC302473</a>	<a href="#">LOC302473.bSep08</a>	<a href="#">302473</a>	4053	302	1	100	similar to SLIT and NTRK-like family, member 4 (LOC302473) alternative variant bSep08, mRNA.
<a href="#">LOC302640</a>	<a href="#">LOC302640.bSep08</a>	<a href="#">302640</a>	49063	871	1	212	similar to acyl-CoA thioesterase (LOC302640) alternative variant bSep08, mRNA.
<a href="#">LOC302680</a>	<a href="#">LOC302680.bSep08</a>	<a href="#">302680</a>	19811	839	5	271	similar to CXORF15 (LOC302680) alternative variant bSep08, mRNA.
<a href="#">LOC303059</a>	<a href="#">LOC303059.bSep08</a>	<a href="#">303059</a>	15916	2622	1	77	similar to novel protein (8.5 kD) (LOC303059) alternative variant bSep08, mRNA.
<a href="#">LOC303140</a>	<a href="#">LOC303140.aSep08</a>	<a href="#">303140</a>	5496	1152		214	up-regulator of carnitine transporter, OCTN2 (LOC303140) mRNA.
<a href="#">LOC303259</a>	<a href="#">LOC303259.aSep08</a>	<a href="#">303259</a>	40742	709	6	178	similar to Map4k6-pending protein (LOC303259) alternative variant aSep08, mRNA.
<a href="#">LOC303259</a>	<a href="#">LOC303259.bSep08</a>	<a href="#">303259</a>	35952	988	1	65	similar to Map4k6-pending protein (LOC303259) alternative variant bSep08, mRNA.
<a href="#">LOC303448</a>	<a href="#">LOC303448.aSep08</a>	<a href="#">303448</a>	6054	823		239	similar to glyceraldehyde-3-phosphate dehydrogenase (LOC303448) mRNA.
<a href="#">LOC303823</a>	<a href="#">LOC303823.bSep08</a>	<a href="#">303823</a>	33567	759	3	190	similar to mitogen-activated protein kinase kinase kinase 13; leucine zipper-bearing kinase (LOC303823) alternative variant bSep08, mRNA.
<a href="#">LOC303823</a>	<a href="#">LOC303823.cSep08</a>	<a href="#">303823</a>	15761	1429	3	51	similar to mitogen-activated protein kinase kinase kinase 13; leucine zipper-bearing kinase (LOC303823) alternative variant cSep08, mRNA.

<a href="#">LOC304000</a>	<a href="#">LOC304000.bSep08</a>	<a href="#">304000</a>	21782	1308	2	227	cell adhesion molecule JCAM (24.6 kD) (LOC304000) alternative variant bSep08, mRNA.
<a href="#">LOC304000</a>	<a href="#">LOC304000.cSep08</a>	<a href="#">304000</a>	21208	756	2	140	cell adhesion molecule JCAM (15.3 kD) (LOC304000) alternative variant cSep08, mRNA.
<a href="#">LOC304239</a>	<a href="#">LOC304239.aSep08</a>	<a href="#">304239</a>	46258	3759		1194	similar to RaIA binding protein 1 and similar to GTPase activating protein testicular GAP1 (LOC304239) mRNA.
<a href="#">LOC304239</a>	<a href="#">LOC304239.aSep08</a>	<a href="#">689661</a>	46258	3759		1194	similar to RaIA binding protein 1 and similar to GTPase activating protein testicular GAP1 (LOC304239) mRNA.
<a href="#">LOC304396</a>	<a href="#">LOC304396.aSep08</a>	<a href="#">304396</a>	12179	1622	10	512	similar to hypothetical protein DKFZp434K1815 (LOC304396) alternative variant aSep08, mRNA.
<a href="#">LOC304396</a>	<a href="#">LOC304396.cSep08</a>	<a href="#">304396</a>	2051	971	2	168	similar to hypothetical protein DKFZp434K1815 (18.1 kD) (LOC304396) alternative variant cSep08, mRNA.
<a href="#">LOC304558</a>	<a href="#">LOC304558.aSep08</a>	<a href="#">304558</a>	109544	504	1	67	similar to TPR repeat-containing protein KIAA1043 (7.4 kD) (LOC304558) alternative variant aSep08, mRNA.
<a href="#">LOC304558</a>	<a href="#">LOC304558.bSep08</a>	<a href="#">304558</a>	55178	562	2	53	similar to TPR repeat-containing protein KIAA1043 (5.8 kD) (LOC304558) alternative variant bSep08, mRNA.
<a href="#">LOC304725</a>	<a href="#">LOC304725.aSep08</a>	<a href="#">304725</a>	282524	473	2	135	similar to contactin associated protein-like 5 isoform 1 (15.0 kD) (LOC304725) alternative variant aSep08, mRNA.
<a href="#">LOC304725</a>	<a href="#">LOC304725.bSep08</a>	<a href="#">304725</a>	36615	392	1	48	similar to contactin associated protein-like 5 isoform 1 (LOC304725) alternative variant bSep08, mRNA.
<a href="#">LOC305076</a>	<a href="#">LOC305076.bSep08</a>	<a href="#">305076</a>	6612	1309	6	131	digestive-organ expansion factor homolog (LOC305076) alternative variant bSep08, mRNA.
<a href="#">LOC305076</a>	<a href="#">LOC305076.cSep08</a>	<a href="#">305076</a>	8185	957	4	120	putative protein (13.2 kD) (LOC305076) alternative variant cSep08, mRNA.
<a href="#">LOC305633</a>	<a href="#">LOC305633.aSep08</a>	<a href="#">305633</a>	95393	2547	8	221	similar to Antxr2 protein (LOC305633) alternative variant aSep08, mRNA.
<a href="#">LOC305691</a>	<a href="#">LOC305691.bSep08</a>	<a href="#">305691</a>	18908	266	1	64	similar to hypothetical protein FLJ22419 (LOC305691) alternative variant bSep08, mRNA.
<a href="#">LOC305771</a>	<a href="#">LOC305771.aSep08</a>	<a href="#">305771</a>	47869	538	2	108	similar to nidogen 2 and hypothetical LOC501227 (11.8 kD) (LOC305771) alternative variant aSep08, mRNA.
<a href="#">LOC305771</a>	<a href="#">LOC305771.aSep08</a>	<a href="#">501227</a>	47869	538	2	108	similar to nidogen 2 and hypothetical LOC501227 (11.8 kD) (LOC305771) alternative variant aSep08, mRNA.
<a href="#">LOC305771</a>	<a href="#">LOC305771.bSep08</a>	<a href="#">305771</a>	49436	821	3	56	similar to nidogen 2 and hypothetical LOC501227 (6.5 kD) (LOC305771) alternative variant bSep08, mRNA.
<a href="#">LOC305771</a>	<a href="#">LOC305771.bSep08</a>	<a href="#">501227</a>	49436	821	3	56	similar to nidogen 2 and hypothetical LOC501227 (6.5 kD) (LOC305771) alternative variant bSep08, mRNA.
<a href="#">LOC306096</a>	<a href="#">LOC306096.aSep08</a>	<a href="#">306096</a>	379727	1917	9	220	similar to Dachshund homolog 1 (Dach1) (24.8 kD) (LOC306096) alternative variant aSep08, mRNA.
<a href="#">LOC306312</a>	<a href="#">LOC306312.aSep08</a>	<a href="#">306312</a>	1504	1161		340	similar to RNA polymerase II transcription factor SIII subunit A2 (Elongin A2) (EloA2) (Transcription elongation factor B polypeptide 3B) (LOC306312) mRNA.
<a href="#">LOC306365</a>	<a href="#">LOC306365.aSep08</a>	<a href="#">306365</a>	18164	568		189	ankyrin (LOC306365) mRNA.
<a href="#">LOC306766</a>	<a href="#">LOC306766.bSep08</a>	<a href="#">306766</a>	13095	2345	5	160	hypothetical LOC306766 (17.3 kD) (LOC306766) alternative variant bSep08, complete mRNA.
<a href="#">LOC306766</a>	<a href="#">LOC306766.cSep08</a>	<a href="#">306766</a>	10869	916	7	83	hypothetical LOC306766 (LOC306766) alternative variant cSep08, mRNA.

<a href="#">LOC306766</a>	<a href="#">LOC306766.dSep08</a>	<a href="#">306766</a>	10841	702	5	53	hypothetical LOC306766 (LOC306766) alternative variant dSep08, mRNA.
<a href="#">LOC306766</a>	<a href="#">LOC306766.fSep08</a>	<a href="#">306766</a>	946	377	2	35	hypothetical LOC306766 (LOC306766) alternative variant fSep08, mRNA.
<a href="#">LOC306766</a>	<a href="#">LOC306766.gSep08</a>	<a href="#">306766</a>	741	350	2	52	hypothetical LOC306766 (LOC306766) alternative variant gSep08, mRNA.
<a href="#">LOC307347</a>	<a href="#">LOC307347.aSep08</a>	<a href="#">307347</a>	37091	2486	17	734	hypothetical protein LOC307347 (LOC307347) alternative variant aSep08, mRNA.
<a href="#">LOC307347</a>	<a href="#">LOC307347.bSep08</a>	<a href="#">307347</a>	4893	413	3	137	hypothetical protein LOC307347 (LOC307347) alternative variant bSep08, mRNA.
<a href="#">LOC308320</a>	<a href="#">LOC308320.bSep08</a>	<a href="#">308320</a>	2472	748	1	106	similar to 5730403M16Rik protein (11.8 kD) (LOC308320) alternative variant bSep08, mRNA.
<a href="#">LOC308320</a>	<a href="#">LOC308320.cSep08</a>	<a href="#">308320</a>	4761	753	4	78	similar to 5730403M16Rik protein (8.7 kD) (LOC308320) alternative variant cSep08, mRNA.
<a href="#">LOC308398</a>	<a href="#">LOC308398.aSep08</a>	<a href="#">308398</a>	28517	1392	3	333	similar to F28C1.3a (LOC308398) alternative variant aSep08, mRNA.
<a href="#">LOC308398</a>	<a href="#">LOC308398.bSep08</a>	<a href="#">308398</a>	6917	629	1	180	similar to F28C1.3a (LOC308398) alternative variant bSep08, mRNA.
<a href="#">LOC308670</a>	<a href="#">LOC308670.cSep08</a>	<a href="#">308670</a>	3821	397	3	67	putative protein (LOC308670) alternative variant cSep08, mRNA.
<a href="#">LOC308670</a>	<a href="#">LOC308670.cSep08</a>	<a href="#">691462</a>	3821	397	3	67	putative protein (LOC308670) alternative variant cSep08, mRNA.
<a href="#">LOC308954</a>	<a href="#">LOC308954.bSep08</a>	<a href="#">308954</a>	11804	672	1	206	similar to hypothetical protein MGC50721 (LOC308954) alternative variant bSep08, mRNA.
<a href="#">LOC308990</a>	<a href="#">LOC308990.aSep08</a>	<a href="#">308990</a>	2531	843	3	232	hypothetical protein LOC308990 (25.4 kD) (LOC308990) alternative variant aSep08, mRNA.
<a href="#">LOC308990</a>	<a href="#">LOC308990.cSep08</a>	<a href="#">308990</a>	1500	778	2	225	hypothetical protein LOC308990 (LOC308990) alternative variant cSep08, mRNA.
<a href="#">LOC308990</a>	<a href="#">LOC308990.dSep08</a>	<a href="#">308990</a>	2263	1583	2	225	hypothetical protein LOC308990 (24.5 kD) (LOC308990) alternative variant dSep08, mRNA.
<a href="#">LOC308990</a>	<a href="#">LOC308990.eSep08</a>	<a href="#">308990</a>	1274	735	2	97	hypothetical protein LOC308990 (LOC308990) alternative variant eSep08, mRNA.
<a href="#">LOC309378</a>	<a href="#">LOC309378.aSep08</a>	<a href="#">309378</a>	18516	798	6	235	similar to golgi autoantigen, golgin subfamily a, 7 (LOC309378) alternative variant aSep08, mRNA.
<a href="#">LOC309378</a>	<a href="#">LOC309378.bSep08</a>	<a href="#">309378</a>	6977	550	2	182	similar to golgi autoantigen, golgin subfamily a, 7 (LOC309378) alternative variant bSep08, mRNA.
<a href="#">LOC309378</a>	<a href="#">LOC309378.cSep08</a>	<a href="#">309378</a>	1522	379	2	78	similar to golgi autoantigen, golgin subfamily a, 7 (LOC309378) alternative variant cSep08, mRNA.
<a href="#">LOC309433</a>	<a href="#">LOC309433.aSep08</a>	<a href="#">309433</a>	7081	1784	4	586	similar to <i>S. cerevisiae</i> SEC31-like 2 isoform a (LOC309433) alternative variant aSep08, mRNA.
<a href="#">LOC309433</a>	<a href="#">LOC309433.bSep08</a>	<a href="#">309433</a>	5855	1039	7	345	similar to <i>S. cerevisiae</i> SEC31-like 2 isoform a (LOC309433) alternative variant bSep08, mRNA.
<a href="#">LOC309433</a>	<a href="#">LOC309433.cSep08</a>	<a href="#">309433</a>	2036	321	5	107	similar to <i>S. cerevisiae</i> SEC31-like 2 isoform a (LOC309433) alternative variant cSep08, mRNA.
<a href="#">LOC309433</a>	<a href="#">LOC309433.dSep08</a>	<a href="#">309433</a>	4137	435	4	21	similar to <i>S. cerevisiae</i> SEC31-like 2 isoform a (2.4 kD) (LOC309433) alternative variant dSep08, mRNA.

<a href="#">LOC309433</a>	<a href="#">LOC309433.eSep08</a>	<a href="#">309433</a>	1054	367	3	47	similar to <i>S. cerevisiae</i> SEC31-like 2 isoform a (LOC309433) alternative variant eSep08, mRNA.
<a href="#">LOC309692</a>	<a href="#">LOC309692.aSep08</a>	<a href="#">309692</a>	13410	1785		444	similar to pericentrin (LOC309692) mRNA.
<a href="#">LOC310013</a>	<a href="#">LOC310013.aSep08</a>	<a href="#">310013</a>	9882	2017		266	hypothetical LOC310013 (LOC310013) mRNA.
<a href="#">LOC310902</a>	<a href="#">LOC310902.aSep08</a>	<a href="#">310902</a>	8248	576		104	similar to Alcohol dehydrogenase 1A (Alcohol dehydrogenase alpha subunit) (LOC310902) mRNA.
<a href="#">LOC310926</a>	<a href="#">LOC310926.cSep08</a>	<a href="#">310926</a>	1503	1337	2	257	hypothetical protein LOC310926 (LOC310926) alternative variant cSep08, mRNA.
<a href="#">LOC310958</a>	<a href="#">LOC310958.bSep08</a>	<a href="#">310958</a>	6773	888	3	113	putative protein of eukaryotic origin (LOC310958) alternative variant bSep08, mRNA.
<a href="#">LOC311026</a>	<a href="#">LOC311026.aSep08</a>	<a href="#">311026</a>	30754	1104	2	155	similar to mKIAA1461 protein (LOC311026) alternative variant aSep08, mRNA.
<a href="#">LOC311026</a>	<a href="#">LOC311026.bSep08</a>	<a href="#">311026</a>	30410	893	1	85	similar to mKIAA1461 protein (9.7 kD) (LOC311026) alternative variant bSep08, mRNA.
<a href="#">LOC311134</a>	<a href="#">LOC311134.aSep08</a>	<a href="#">311134</a>	95663	2158		547	hypothetical LOC311134 (LOC311134) mRNA.
<a href="#">LOC311352</a>	<a href="#">LOC311352.bSep08</a>	<a href="#">311352</a>	5218	737	4	160	similar to Adenosine deaminase CG11994-PA (LOC311352) alternative variant bSep08, mRNA.
<a href="#">LOC311984</a>	<a href="#">LOC311984.bSep08</a>	<a href="#">311984</a>	17706	1665	12	288	similar to RIKEN cDNA A530088I07 gene (33.3 kD) (LOC311984) alternative variant bSep08, mRNA.
<a href="#">LOC311984</a>	<a href="#">LOC311984.cSep08</a>	<a href="#">311984</a>	1518	718	2	46	similar to RIKEN cDNA A530088I07 gene (LOC311984) alternative variant cSep08, mRNA.
<a href="#">LOC312502</a>	<a href="#">LOC312502.aSep08</a>	<a href="#">312502</a>	26266	968		322	similar to RAB11 family interacting protein 5 (class I) isoform 1 (LOC312502) mRNA.
<a href="#">LOC312678</a>	<a href="#">LOC312678.aSep08</a>	<a href="#">312678</a>	34643	1803	11	601	similar to Retinoblastoma-binding protein 2 (RBBP-2) (LOC312678) alternative variant aSep08, mRNA.
<a href="#">LOC312678</a>	<a href="#">LOC312678.bSep08</a>	<a href="#">312678</a>	33221	2315	12	579	similar to Retinoblastoma-binding protein 2 (RBBP-2) (66.1 kD) (LOC312678) alternative variant bSep08, mRNA.
<a href="#">LOC312678</a>	<a href="#">LOC312678.cSep08</a>	<a href="#">312678</a>	17450	1416	9	471	similar to Retinoblastoma-binding protein 2 (RBBP-2) (LOC312678) alternative variant cSep08, mRNA.
<a href="#">LOC312678</a>	<a href="#">LOC312678.dSep08</a>	<a href="#">312678</a>	6728	1028	4	342	similar to Retinoblastoma-binding protein 2 (RBBP-2) (LOC312678) alternative variant dSep08, mRNA.
<a href="#">LOC312678</a>	<a href="#">LOC312678.eSep08</a>	<a href="#">312678</a>	9031	716	4	188	similar to Retinoblastoma-binding protein 2 (RBBP-2) (LOC312678) alternative variant eSep08, mRNA.
<a href="#">LOC312831</a>	<a href="#">LOC312831.bSep08</a>	<a href="#">312831</a>	29565	818	4	207	similar to SRY (sex determining region Y)-box 5 isoform a (LOC312831) alternative variant bSep08, mRNA.
<a href="#">LOC312831</a>	<a href="#">LOC312831.cSep08</a>	<a href="#">312831</a>	34919	760	5	187	similar to SRY (sex determining region Y)-box 5 isoform a (LOC312831) alternative variant cSep08, mRNA.
<a href="#">LOC313149</a>	<a href="#">LOC313149.aSep08</a>	<a href="#">313149</a>	6529	490		162	hypothetical LOC313149 (LOC313149) mRNA.
<a href="#">LOC313519.1</a>	<a href="#">LOC313519.1.aSep08</a>	<a href="#">313519</a>	40898	3499	10	475	similar to RIKEN cDNA 5330440M15 (52.4 kD) (LOC313519.1) alternative variant aSep08, complete mRNA.
<a href="#">LOC313519.1</a>	<a href="#">LOC313519.1.bSep08</a>	<a href="#">313519</a>	2521	1059	1	196	similar to RIKEN cDNA 5330440M15 (22.0 kD) (LOC313519.1) alternative variant bSep08, mRNA.
<a href="#">LOC313519.1</a>	<a href="#">LOC313519.1.cSep08</a>	<a href="#">313519</a>	13342	386	2	58	similar to RIKEN cDNA 5330440M15 (LOC313519.1) alternative variant cSep08, mRNA.
<a href="#">LOC313641</a>	<a href="#">LOC313641.aSep08</a>	<a href="#">313641</a>	19344	4886	26	1168	perlecan (125.2 kD) (LOC313641) alternative variant aSep08, mRNA.

<a href="#">LOC313641</a>	<a href="#">LOC313641.bSep08</a>	<a href="#">313641</a>	2501	633	5	211	perlecan (LOC313641) alternative variant bSep08, mRNA.
<a href="#">LOC313641</a>	<a href="#">LOC313641.cSep08</a>	<a href="#">313641</a>	3031	528	6	175	perlecan (LOC313641) alternative variant cSep08, mRNA.
<a href="#">LOC313672</a>	<a href="#">LOC313672.aSep08</a>	<a href="#">313672</a>	91763	1796	2	598	similar to CG11206-PA (LOC313672) alternative variant aSep08, mRNA.
<a href="#">LOC313707</a>	<a href="#">LOC313707.aSep08</a>	<a href="#">313707</a>	23616	2759	25	537	exosome component 10 CRA b (61.3 kD) (LOC313707) alternative variant aSep08, mRNA.
<a href="#">LOC313707</a>	<a href="#">LOC313707.cSep08</a>	<a href="#">313707</a>	2534	844	4	115	exosome component 10 (12.9 kD) (LOC313707) alternative variant cSep08, mRNA.
<a href="#">LOC313707</a>	<a href="#">LOC313707.dSep08</a>	<a href="#">313707</a>	3243	560	3	108	exosome component 10 (LOC313707) alternative variant dSep08, mRNA.
<a href="#">LOC313707</a>	<a href="#">LOC313707.eSep08</a>	<a href="#">313707</a>	13811	473	6	80	exosome component 10 (8.8 kD) (LOC313707) alternative variant eSep08, mRNA.
<a href="#">LOC313707</a>	<a href="#">LOC313707.fSep08</a>	<a href="#">313707</a>	1804	1298	2	112	putative protein (11.9 kD) (LOC313707) alternative variant fSep08, mRNA.
<a href="#">LOC313936</a>	<a href="#">LOC313936.bSep08</a>	<a href="#">313936</a>	13564	741	1	143	hypothetical LOC313936 (LOC313936) alternative variant bSep08, mRNA.
<a href="#">LOC314140</a>	<a href="#">LOC314140.cSep08</a>	<a href="#">314140</a>	6869	375	2	83	putative protein (LOC314140) alternative variant cSep08, mRNA.
<a href="#">LOC314140</a>	<a href="#">LOC314140.dSep08</a>	<a href="#">314140</a>	6997	479	2	26	putative protein (LOC314140) alternative variant dSep08, mRNA.
<a href="#">LOC314328</a>	<a href="#">LOC314328.aSep08</a>	<a href="#">314328</a>	9876	792	2	185	uncharacterized protein (LOC314328) alternative variant aSep08, mRNA.
<a href="#">LOC314328</a>	<a href="#">LOC314328.bSep08</a>	<a href="#">314328</a>	9880	765	2	169	uncharacterized protein (19.1 kD) (LOC314328) alternative variant bSep08, complete mRNA.
<a href="#">LOC314328</a>	<a href="#">LOC314328.cSep08</a>	<a href="#">314328</a>	9851	723	2	128	putative protein of metazoan origin (14.2 kD) (LOC314328) alternative variant cSep08, complete mRNA.
<a href="#">LOC314655</a>	<a href="#">LOC314655.aSep08</a>	<a href="#">314655</a>	22702	3525	21	705	similar to a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 10 (77.0 kD) (LOC314655) alternative variant aSep08, mRNA.
<a href="#">LOC314655</a>	<a href="#">LOC314655.cSep08</a>	<a href="#">314655</a>	5560	741	6	189	similar to a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 10 (LOC314655) alternative variant cSep08, mRNA.
<a href="#">LOC314655</a>	<a href="#">LOC314655.dSep08</a>	<a href="#">314655</a>	4456	663	3	145	similar to a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 10 (LOC314655) alternative variant dSep08, mRNA.
<a href="#">LOC314655</a>	<a href="#">LOC314655.eSep08</a>	<a href="#">314655</a>	889	748	2	144	similar to a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 10 (LOC314655) alternative variant eSep08, mRNA.
<a href="#">LOC314655</a>	<a href="#">LOC314655.fSep08</a>	<a href="#">314655</a>	618	536	2	106	similar to a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 10 (LOC314655) alternative variant fSep08, mRNA.
<a href="#">LOC314776</a>	<a href="#">LOC314776.aSep08</a>	<a href="#">314776</a>	37193	598		69	similar to Peptidyl-prolyl cis-trans isomerase A (PPIase) (Rotamase) (Cyclophilin A) (Cyclosporin A-binding protein) (P31) (p1B15) (8.0 kD) (LOC314776) mRNA.
<a href="#">LOC314942</a>	<a href="#">LOC314942.aSep08</a>	<a href="#">314942</a>	18110	2708		301	similar to CUB and Sushi multiple domains 3 isoform 1 (LOC314942) mRNA.
<a href="#">LOC315766</a>	<a href="#">LOC315766.aSep08</a>	<a href="#">315766</a>	1540	392		130	similar to zinc finger protein 609 (LOC315766) mRNA.

<a href="#">LOC315970</a>	<a href="#">LOC315970.bSep08</a>	<a href="#">315970</a>	1816	776	1	65	similar to CDV-3B (LOC315970) alternative variant bSep08, mRNA.
<a href="#">LOC316124</a>	<a href="#">LOC316124.bSep08</a>	<a href="#">316124</a>	1957	696	4	231	similar to gonadotropin-regulated long chain acyl-CoA synthetase (LOC316124) alternative variant bSep08, mRNA.
<a href="#">LOC316124</a>	<a href="#">LOC316124.cSep08</a>	<a href="#">316124</a>	5633	677	6	225	similar to gonadotropin-regulated long chain acyl-CoA synthetase (LOC316124) alternative variant cSep08, mRNA.
<a href="#">LOC316124</a>	<a href="#">LOC316124.dSep08</a>	<a href="#">316124</a>	817	314	2	53	similar to gonadotropin-regulated long chain acyl-CoA synthetase (LOC316124) alternative variant dSep08, mRNA.
<a href="#">LOC316919</a>	<a href="#">LOC316919.bSep08</a>	<a href="#">316919</a>	508	385	2	10	similar to spermatogenesis associated glutamate (E)-rich protein 4d (1.1 kD) (LOC316919) alternative variant bSep08, mRNA.
<a href="#">LOC317416</a>	<a href="#">LOC317416.aSep08</a>	<a href="#">317416</a>	21282	897		260	similar to inter alpha-trypsin inhibitor, heavy chain 4 (29.4 kD) (LOC317416) mRNA.
<a href="#">LOC360479</a>	<a href="#">LOC360479.bSep08</a>	<a href="#">360479</a>	5114	483	4	105	similar to hypothetical protein (LOC360479) alternative variant bSep08, mRNA.
<a href="#">LOC360504</a>	<a href="#">LOC360504.bSep08</a>	<a href="#">360504</a>	13923	366	3	93	hemoglobin alpha 2 chain (LOC360504) alternative variant bSep08, mRNA.
<a href="#">LOC360684</a>	<a href="#">LOC360684.aSep08</a>	<a href="#">360684</a>	13834	783		228	similar to eyes absent 4 isoform a (LOC360684) mRNA.
<a href="#">LOC360721</a>	<a href="#">LOC360721.bSep08</a>	<a href="#">360721</a>	27070	993	5	155	growth and transformation-dependent protein (17.8 kD) (LOC360721) alternative variant bSep08, mRNA.
<a href="#">LOC360721</a>	<a href="#">LOC360721.cSep08</a>	<a href="#">360721</a>	27315	832	5	126	growth and transformation-dependent protein (14.5 kD) (LOC360721) alternative variant cSep08, mRNA.
<a href="#">LOC360721</a>	<a href="#">LOC360721.fSep08</a>	<a href="#">360721</a>	5214	589	2	19	growth and transformation-dependent protein (LOC360721) alternative variant fSep08, complete mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.aSep08</a>	<a href="#">360807</a>	17876	1260	9	419	arginine serine-rich coiled-coil 2 (LOC360807) alternative variant aSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.cSep08</a>	<a href="#">360807</a>	9973	1718	6	293	putative protein (LOC360807) alternative variant cSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.dSep08</a>	<a href="#">360807</a>	10886	765	6	255	arginine serine-rich coiled-coil 2 (LOC360807) alternative variant dSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.eSep08</a>	<a href="#">360807</a>	11869	743	6	233	arginine serine-rich coiled-coil 2 (LOC360807) alternative variant eSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.fSep08</a>	<a href="#">360807</a>	3242	872	3	179	arginine serine-rich coiled-coil 2 (LOC360807) alternative variant fSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.gSep08</a>	<a href="#">360807</a>	2879	984	2	156	arginine serine-rich coiled-coil 2 (17.2 kD) (LOC360807) alternative variant gSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.hSep08</a>	<a href="#">360807</a>	2340	640	2	102	CRA b like (LOC360807) alternative variant hSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.iSep08</a>	<a href="#">360807</a>	3135	770	3	101	putative protein (LOC360807) alternative variant iSep08, mRNA.
<a href="#">LOC360807</a>	<a href="#">LOC360807.jSep08</a>	<a href="#">360807</a>	5528	370	3	98	arginine serine-rich coiled-coil 2 (LOC360807) alternative variant jSep08, mRNA.
<a href="#">LOC360824</a>	<a href="#">LOC360824.aSep08</a>	<a href="#">360824</a>	20157	991	3	282	similar to CG9164-PA, isoform A (LOC360824) alternative variant aSep08, mRNA.

<a href="#">LOC360824</a>	<a href="#">LOC360824.bSep08</a>	<a href="#">360824</a>	7711	783	1	172	similar to CG9164-PA, isoform A (LOC360824) alternative variant bSep08, mRNA.
<a href="#">LOC360919</a>	<a href="#">LOC360919.aSep08</a>	<a href="#">360919</a>	22976	858	7	286	similar to alpha-fetoprotein (LOC360919) alternative variant aSep08, mRNA.
<a href="#">LOC360919</a>	<a href="#">LOC360919.cSep08</a>	<a href="#">360919</a>	6589	527	4	175	similar to alpha-fetoprotein (LOC360919) alternative variant cSep08, mRNA.
<a href="#">LOC360919</a>	<a href="#">LOC360919.dSep08</a>	<a href="#">360919</a>	5215	1180	2	51	similar to alpha-fetoprotein (5.7 kD) (LOC360919) alternative variant dSep08, mRNA.
<a href="#">LOC360919</a>	<a href="#">LOC360919.eSep08</a>	<a href="#">360919</a>	943	451	2	30	similar to alpha-fetoprotein (LOC360919) alternative variant eSep08, mRNA.
<a href="#">LOC360990</a>	<a href="#">LOC360990.aSep08</a>	<a href="#">360990</a>	23128	2708	13	715	similar to ubiquitin specific protease 34 (LOC360990) alternative variant aSep08, mRNA.
<a href="#">LOC360990</a>	<a href="#">LOC360990.bSep08</a>	<a href="#">360990</a>	17845	739	8	236	similar to ubiquitin specific protease 34 (LOC360990) alternative variant bSep08, mRNA.
<a href="#">LOC360990</a>	<a href="#">LOC360990.cSep08</a>	<a href="#">360990</a>	14544	698	6	232	similar to ubiquitin specific protease 34 (LOC360990) alternative variant cSep08, mRNA.
<a href="#">LOC360990</a>	<a href="#">LOC360990.dSep08</a>	<a href="#">360990</a>	4075	1068	3	87	similar to ubiquitin specific protease 34 (10.1 kD) (LOC360990) alternative variant dSep08, mRNA.
<a href="#">LOC360990</a>	<a href="#">LOC360990.eSep08</a>	<a href="#">360990</a>	13221	663	4	85	similar to ubiquitin specific protease 34 (LOC360990) alternative variant eSep08, mRNA.
<a href="#">LOC361016</a>	<a href="#">LOC361016.aSep08</a>	<a href="#">361016</a>	50337	888	7	295	similar to RIKEN cDNA 4933406L09 (LOC361016) alternative variant aSep08, mRNA.
<a href="#">LOC361016</a>	<a href="#">LOC361016.bSep08</a>	<a href="#">361016</a>	47831	717	5	239	similar to RIKEN cDNA 4933406L09 (LOC361016) alternative variant bSep08, mRNA.
<a href="#">LOC361016</a>	<a href="#">LOC361016.cSep08</a>	<a href="#">361016</a>	66720	945	4	137	similar to RIKEN cDNA 4933406L09 (LOC361016) alternative variant cSep08, mRNA.
<a href="#">LOC361016</a>	<a href="#">LOC361016.dSep08</a>	<a href="#">361016</a>	2966	604	3	121	similar to RIKEN cDNA 4933406L09 (LOC361016) alternative variant dSep08, mRNA.
<a href="#">LOC361041</a>	<a href="#">LOC361041.aSep08</a>	<a href="#">361041</a>	5960	1678	7	559	similar to hypothetical protein (LOC361041) alternative variant aSep08, mRNA.
<a href="#">LOC361041</a>	<a href="#">LOC361041.bSep08</a>	<a href="#">361041</a>	1276	425	3	122	similar to hypothetical protein (LOC361041) alternative variant bSep08, mRNA.
<a href="#">LOC361041</a>	<a href="#">LOC361041.cSep08</a>	<a href="#">361041</a>	825	384	1	83	similar to hypothetical protein (LOC361041) alternative variant cSep08, mRNA.
<a href="#">LOC361098</a>	<a href="#">LOC361098.aSep08</a>	<a href="#">361098</a>	5382	386		128	similar to SGT1 protein homolog (Ecdysoneless homolog) (LOC361098) mRNA.
<a href="#">LOC361100</a>	<a href="#">LOC361100.aSep08</a>	<a href="#">361100</a>	12308	1619		431	similar to topoisomerase (DNA) II beta (LOC361100) mRNA.
<a href="#">LOC361128</a>	<a href="#">LOC361128.bSep08</a>	<a href="#">361128</a>	2267	1139	5	140	similar to TR4 orphan receptor associated protein TRA16 (16.1 kD) (LOC361128) alternative variant bSep08, mRNA.
<a href="#">LOC361128</a>	<a href="#">LOC361128.cSep08</a>	<a href="#">361128</a>	613	405	2	92	similar to TR4 orphan receptor associated protein TRA16 (LOC361128) alternative variant cSep08, mRNA.
<a href="#">LOC361128</a>	<a href="#">LOC361128.dSep08</a>	<a href="#">361128</a>	1806	750	4	83	similar to TR4 orphan receptor associated protein TRA16 (9.6 kD) (LOC361128) alternative variant dSep08, mRNA.
<a href="#">LOC361187</a>	<a href="#">LOC361187.aSep08</a>	<a href="#">361187</a>	5324	1780	1	538	ankyrin (LOC361187) alternative variant aSep08, mRNA.
<a href="#">LOC361188</a>	<a href="#">LOC361188.aSep08</a>	<a href="#">361188</a>	10504	655		159	similar to WD repeat domain 17 (LOC361188) mRNA.
<a href="#">LOC361230</a>	<a href="#">LOC361230.aSep08</a>	<a href="#">361230</a>	31855	893		116	ac1258 (LOC361230) mRNA.

<a href="#">LOC361346</a>	<a href="#">LOC361346.bSep08</a>	<a href="#">361346</a>	4338	981	4	158	CRA a like (LOC361346) alternative variant bSep08, mRNA.
<a href="#">LOC361346</a>	<a href="#">LOC361346.cSep08</a>	<a href="#">361346</a>	4199	873	4	146	CRA f like (LOC361346) alternative variant cSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.bSep08</a>	<a href="#">361399</a>	3383	2042	13	680	enhancer of mRNA decapping 4 CRA b (LOC361399) alternative variant bSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.cSep08</a>	<a href="#">361399</a>	1682	837	8	278	enhancer of mRNA decapping 4 CRA a (LOC361399) alternative variant cSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.dSep08</a>	<a href="#">361399</a>	1877	1288	7	224	enhancer of mRNA decapping 4 CRA a (LOC361399) alternative variant dSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.eSep08</a>	<a href="#">361399</a>	1647	1015	3	180	enhancer of mRNA decapping 4 CRA a (LOC361399) alternative variant eSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.gSep08</a>	<a href="#">361399</a>	697	558	2	106	neurtin 1-like (11.0 kD) (LOC361399) alternative variant gSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.hSep08</a>	<a href="#">361399</a>	457	382	2	105	enhancer of mRNA decapping 4 CRA a (LOC361399) alternative variant hSep08, mRNA.
<a href="#">LOC361399</a>	<a href="#">LOC361399.iSep08</a>	<a href="#">361399</a>	355	274	2	91	putative protein (LOC361399) alternative variant iSep08, mRNA.
<a href="#">LOC361414</a>	<a href="#">LOC361414.aSep08</a>	<a href="#">361414</a>	102473	1765	4	473	similar to Synaptic vesicle membrane protein VAT-1 homolog (LOC361414) alternative variant aSep08, mRNA.
<a href="#">LOC361414</a>	<a href="#">LOC361414.bSep08</a>	<a href="#">361414</a>	100493	962	4	179	similar to Synaptic vesicle membrane protein VAT-1 homolog (LOC361414) alternative variant bSep08, mRNA.
<a href="#">LOC361414</a>	<a href="#">LOC361414.cSep08</a>	<a href="#">361414</a>	2320	657	1	75	similar to Synaptic vesicle membrane protein VAT-1 homolog (LOC361414) alternative variant cSep08, mRNA.
<a href="#">LOC361635</a>	<a href="#">LOC361635.bSep08</a>	<a href="#">361635</a>	51240	1229	14	409	similar to RIKEN cDNA 9030624J02 (LOC361635) alternative variant bSep08, mRNA.
<a href="#">LOC361646</a>	<a href="#">LOC361646.bSep08</a>	<a href="#">361646</a>	29492	975	5	324	similar to K04F10.2 (LOC361646) alternative variant bSep08, mRNA.
<a href="#">LOC361646</a>	<a href="#">LOC361646.cSep08</a>	<a href="#">361646</a>	86610	767	6	241	similar to K04F10.2 (LOC361646) alternative variant cSep08, mRNA.
<a href="#">LOC361776</a>	<a href="#">LOC361776.aSep08</a>	<a href="#">361776</a>	63686	1328	8	356	LOC361776 (41.3 kD) (LOC361776) alternative variant aSep08, mRNA.
<a href="#">LOC361776</a>	<a href="#">LOC361776.cSep08</a>	<a href="#">361776</a>	15461	759	4	253	LOC361776 (LOC361776) alternative variant cSep08, mRNA.
<a href="#">LOC361780</a>	<a href="#">LOC361780.aSep08</a>	<a href="#">361780</a>	1831	530	4	142	similar to CG3295-PA (LOC361780) alternative variant aSep08, mRNA.
<a href="#">LOC361780</a>	<a href="#">LOC361780.bSep08</a>	<a href="#">361780</a>	2073	1994	2	66	similar to CG3295-PA (7.5 kD) (LOC361780) alternative variant bSep08, mRNA.
<a href="#">LOC361781</a>	<a href="#">LOC361781.aSep08</a>	<a href="#">361781</a>	7001	694		230	similar to helicase, lymphoid specific (LOC361781) mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.aSep08</a>	<a href="#">361990</a>	10491	1981	6	249	similar to DKFZP547E1010 protein (26.6 kD) (LOC361990) alternative variant aSep08, complete mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.bSep08</a>	<a href="#">361990</a>	6973	1268	4	223	similar to DKFZP547E1010 protein (23.8 kD) (LOC361990) alternative variant bSep08, mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.cSep08</a>	<a href="#">361990</a>	9429	916	6	220	similar to DKFZP547E1010 protein (LOC361990) alternative variant cSep08, mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.eSep08</a>	<a href="#">361990</a>	9335	687	5	147	similar to DKFZP547E1010 protein (LOC361990) alternative variant eSep08, mRNA.



<a href="#">LOC361990</a>	<a href="#">LOC361990.fSep08</a>	<a href="#">361990</a>	2814	852	2	141	similar to DKFZP547E1010 protein (LOC361990) alternative variant fSep08, mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.gSep08</a>	<a href="#">361990</a>	3680	647	2	72	similar to DKFZP547E1010 protein (7.8 kD) (LOC361990) alternative variant gSep08, mRNA.
<a href="#">LOC361990</a>	<a href="#">LOC361990.hSep08</a>	<a href="#">361990</a>	2554	1046	2	61	similar to DKFZP547E1010 protein (6.7 kD) (LOC361990) alternative variant hSep08, mRNA.
<a href="#">LOC362464</a>	<a href="#">LOC362464.aSep08</a>	<a href="#">362464</a>	11056	544		100	similar to aryl hydrocarbon receptor nuclear translocator-like 2 (LOC362464) mRNA.
<a href="#">LOC362526</a>	<a href="#">LOC362526.bSep08</a>	<a href="#">362526</a>	20158	1685	2	238	hypothetical protein LOC362526 (28.9 kD) (LOC362526) alternative variant bSep08, mRNA.
<a href="#">LOC362564</a>	<a href="#">LOC362564.bSep08</a>	<a href="#">362564</a>	18030	613	2	118	hypothetical LOC362564 (LOC362564) alternative variant bSep08, mRNA.
<a href="#">LOC362564</a>	<a href="#">LOC362564.cSep08</a>	<a href="#">362564</a>	5343	390	2	91	hypothetical LOC362564 (LOC362564) alternative variant cSep08, mRNA.
<a href="#">LOC362587</a>	<a href="#">LOC362587.aSep08</a>	<a href="#">362587</a>	31827	4903	24	1200	similar to microfilament and actin filament cross-linker protein isoform b (LOC362587) alternative variant aSep08, mRNA.
<a href="#">LOC362587</a>	<a href="#">LOC362587.bSep08</a>	<a href="#">362587</a>	19968	1369	10	456	similar to microfilament and actin filament cross-linker protein isoform b (LOC362587) alternative variant bSep08, mRNA.
<a href="#">LOC362587</a>	<a href="#">LOC362587.cSep08</a>	<a href="#">362587</a>	14527	1272	6	268	similar to microfilament and actin filament cross-linker protein isoform b (LOC362587) alternative variant cSep08, mRNA.
<a href="#">LOC362587</a>	<a href="#">LOC362587.dSep08</a>	<a href="#">362587</a>	1349	661	2	121	similar to microfilament and actin filament cross-linker protein isoform b (12.4 kD) (LOC362587) alternative variant dSep08, mRNA.
<a href="#">LOC362710</a>	<a href="#">LOC362710.bSep08</a>	<a href="#">362710</a>	2703	1691	3	173	hypothetical LOC362710 (LOC362710) alternative variant bSep08, mRNA.
<a href="#">LOC362710</a>	<a href="#">LOC362710.cSep08</a>	<a href="#">362710</a>	1780	503	3	114	hypothetical LOC362710 (LOC362710) alternative variant cSep08, mRNA.
<a href="#">LOC362845</a>	<a href="#">LOC362845.aSep08</a>	<a href="#">362845</a>	13660	1793	2	443	similar to zinc finger protein 709 (LOC362845) alternative variant aSep08, mRNA.
<a href="#">LOC362845</a>	<a href="#">LOC362845.cSep08</a>	<a href="#">362845</a>	11919	630	1	167	similar to zinc finger protein 709 (LOC362845) alternative variant cSep08, mRNA.
<a href="#">LOC362855</a>	<a href="#">LOC362855.bSep08</a>	<a href="#">362855</a>	4915	962	1	281	p55 (LOC362855) alternative variant bSep08, mRNA.
<a href="#">LOC363060</a>	<a href="#">LOC363060.bSep08</a>	<a href="#">363060</a>	11410	1351	2	254	similar to RIKEN cDNA 1600029D21 (26.7 kD) (LOC363060) alternative variant bSep08, complete mRNA.
<a href="#">LOC363060</a>	<a href="#">LOC363060.cSep08</a>	<a href="#">363060</a>	6950	412	1	95	similar to RIKEN cDNA 1600029D21 (LOC363060) alternative variant cSep08, mRNA.
<a href="#">LOC363181</a>	<a href="#">LOC363181.aSep08</a>	<a href="#">363181</a>	7328	842		204	similar to RIKEN cDNA 1700001E04 (LOC363181) mRNA.
<a href="#">LOC363188</a>	<a href="#">LOC363188.aSep08</a>	<a href="#">363188</a>	81905	1792	3	546	similar to ubiquitin protein ligase E3 component n-recognin 2 (LOC363188) alternative variant aSep08, mRNA.
<a href="#">LOC363188</a>	<a href="#">LOC363188.bSep08</a>	<a href="#">363188</a>	5125	1345	3	214	similar to ubiquitin protein ligase E3 component n-recognin 2 (LOC363188) alternative variant bSep08, mRNA.
<a href="#">LOC363188</a>	<a href="#">LOC363188.cSep08</a>	<a href="#">363188</a>	8893	448	2	149	similar to ubiquitin protein ligase E3 component n-recognin 2 (LOC363188) alternative variant cSep08, mRNA.

<a href="#">LOC363265</a>	<a href="#">LOC363265.aSep08</a>	<a href="#">363265</a>	58052	800	3	84	similar to procollagen, type IV, alpha 3 (8.7 kD) (LOC363265) alternative variant aSep08, mRNA.
<a href="#">LOC363265</a>	<a href="#">LOC363265.bSep08</a>	<a href="#">363265</a>	22359	508	1	58	similar to procollagen, type IV, alpha 3 (6.1 kD) (LOC363265) alternative variant bSep08, mRNA.
<a href="#">LOC363266</a>	<a href="#">LOC363266.aSep08</a>	<a href="#">363266</a>	53288	3042	3	495	similar to HIV-1 Rev binding protein (LOC363266) alternative variant aSep08, mRNA.
<a href="#">LOC363266</a>	<a href="#">LOC363266.bSep08</a>	<a href="#">363266</a>	60572	2496	4	331	similar to HIV-1 Rev binding protein (33.3 kD) (LOC363266) alternative variant bSep08, mRNA.
<a href="#">LOC363266</a>	<a href="#">LOC363266.cSep08</a>	<a href="#">363266</a>	16774	1040	2	328	similar to HIV-1 Rev binding protein (LOC363266) alternative variant cSep08, mRNA.
<a href="#">LOC363267</a>	<a href="#">LOC363267.bSep08</a>	<a href="#">363267</a>	22134	715	5	219	hypothetical protein LOC363267 (LOC363267) alternative variant bSep08, mRNA.
<a href="#">LOC363267</a>	<a href="#">LOC363267.cSep08</a>	<a href="#">363267</a>	26433	378	1	125	hypothetical protein LOC363267 (LOC363267) alternative variant cSep08, mRNA.
<a href="#">LOC363301</a>	<a href="#">LOC363301.bSep08</a>	<a href="#">363301</a>	4849	500	4	84	hypothetical LOC363301 (LOC363301) alternative variant bSep08, mRNA.
<a href="#">LOC363306</a>	<a href="#">LOC363306.aSep08</a>	<a href="#">363306</a>	8293	1419	3	206	hypothetical protein LOC363306 and similar to Reticulocalbin-1 precursor (LOC363306) alternative variant aSep08, mRNA.
<a href="#">LOC363306</a>	<a href="#">LOC363306.aSep08</a>	<a href="#">688719</a>	8293	1419	3	206	hypothetical protein LOC363306 and similar to Reticulocalbin-1 precursor (LOC363306) alternative variant aSep08, mRNA.
<a href="#">LOC363306</a>	<a href="#">LOC363306.bSep08</a>	<a href="#">363306</a>	21263	495	1	102	hypothetical protein LOC363306 and similar to Reticulocalbin-1 precursor (LOC363306) alternative variant bSep08, mRNA.
<a href="#">LOC363306</a>	<a href="#">LOC363306.bSep08</a>	<a href="#">688719</a>	21263	495	1	102	hypothetical protein LOC363306 and similar to Reticulocalbin-1 precursor (LOC363306) alternative variant bSep08, mRNA.
<a href="#">LOC363313</a>	<a href="#">LOC363313.aSep08</a>	<a href="#">363313</a>	2782	846		177	hypothetical LOC363313 (LOC363313) mRNA.
<a href="#">LOC363326</a>	<a href="#">LOC363326.bSep08</a>	<a href="#">363326</a>	776	587	3	195	hypothetical LOC363326 (LOC363326) alternative variant bSep08, mRNA.
<a href="#">LOC363326</a>	<a href="#">LOC363326.cSep08</a>	<a href="#">363326</a>	467	383	2	55	hypothetical LOC363326 (LOC363326) alternative variant cSep08, mRNA.
<a href="#">LOC363336</a>	<a href="#">LOC363336.aSep08</a>	<a href="#">363336</a>	7710	485			
<a href="#">LOC363337</a>	<a href="#">LOC363337.aSep08</a>	<a href="#">363337</a>	2703	776	2	154	similar to RIKEN cDNA 1700081O22 (LOC363337) alternative variant aSep08, mRNA.
<a href="#">LOC363410</a>	<a href="#">LOC363410.aSep08</a>	<a href="#">363410</a>	1870	732		92	similar to Ras-related protein Rab-27B (LOC363410) mRNA.
<a href="#">LOC363458</a>	<a href="#">LOC363458.aSep08</a>	<a href="#">363458</a>	1015	318		105	similar to procollagen, type IV, alpha 6 (LOC363458) alternative variant aSep08, mRNA.
<a href="#">LOC363458</a>	<a href="#">LOC363458.bSep08</a>	<a href="#">363458</a>	970	378	1	63	similar to procollagen, type IV, alpha 6 (LOC363458) alternative variant bSep08, mRNA.
<a href="#">LOC363711</a>	<a href="#">LOC363711.aSep08</a>	<a href="#">363711</a>	7869	402		133	similar to Ciliary dynein heavy chain 9 (Axonemal beta dynein heavy chain 9) (LOC363711) mRNA.
<a href="#">LOC363849</a>	<a href="#">LOC363849.aSep08</a>	<a href="#">363849</a>	51466	2354	11	532	similar to histone cell cycle regulation defective homolog A isoform 1 (LOC363849) alternative variant aSep08, mRNA.

<a href="#">LOC363849</a>	<a href="#">LOC363849.bSep08</a>	<a href="#">363849</a>	8483	889	8	274	similar to histone cell cycle regulation defective homolog A isoform 1 (LOC363849) alternative variant bSep08, mRNA.
<a href="#">LOC363849</a>	<a href="#">LOC363849.cSep08</a>	<a href="#">363849</a>	57650	840	3	151	similar to histone cell cycle regulation defective homolog A isoform 1 (LOC363849) alternative variant cSep08, mRNA.
<a href="#">LOC363915</a>	<a href="#">LOC363915.aSep08</a>	<a href="#">363915</a>	3113	709		236	similar to CG14446-PA (LOC363915) mRNA.
<a href="#">LOC364084</a>	<a href="#">LOC364084.aSep08</a>	<a href="#">364084</a>	1437	321		106	putative protein of ancient origin (LOC364084) mRNA.
<a href="#">LOC364224</a>	<a href="#">LOC364224.aSep08</a>	<a href="#">364224</a>	47020	467		107	similar to Acylphosphatase, muscle type isozyme (Acylphosphate phosphohydrolase) (LOC364224) mRNA.
<a href="#">LOC364427</a>	<a href="#">LOC364427.bSep08</a>	<a href="#">364427</a>	4750	749	1	174	similar to CG5407-PA, isoform A (LOC364427) alternative variant bSep08, mRNA.
<a href="#">LOC364558</a>	<a href="#">LOC364558.aSep08</a>	<a href="#">364558</a>	25749	718	2	154	similar to palladin; CGI-151 protein (LOC364558) alternative variant aSep08, mRNA.
<a href="#">LOC364558</a>	<a href="#">LOC364558.bSep08</a>	<a href="#">364558</a>	2200	316	1	36	similar to palladin; CGI-151 protein (LOC364558) alternative variant bSep08, mRNA.
<a href="#">LOC364653</a>	<a href="#">LOC364653.aSep08</a>	<a href="#">364653</a>	1769	414		138	similar to WD repeat domain 17 (LOC364653) mRNA.
<a href="#">LOC364773</a>	<a href="#">LOC364773.bSep08</a>	<a href="#">364773</a>	4941	522	1	168	similar to liver regeneration-related protein LRRG07 (19.2 kD) (LOC364773) alternative variant bSep08, mRNA.
<a href="#">LOC364802</a>	<a href="#">LOC364802.aSep08</a>	<a href="#">364802</a>	1787	1585	2	221	similar to CG8043-PA (LOC364802) alternative variant aSep08, mRNA.
<a href="#">LOC365052</a>	<a href="#">LOC365052.aSep08</a>	<a href="#">365052</a>	41017	400		31	similar to ribosomal protein L10a (LOC365052) mRNA.
<a href="#">LOC365084</a>	<a href="#">LOC365084.aSep08</a>	<a href="#">365084</a>	42822	599		63	similar to nidogen 2 and similar to MIC2 like 1 (LOC365084) mRNA.
<a href="#">LOC365084</a>	<a href="#">LOC365084.aSep08</a>	<a href="#">679721</a>	42822	599		63	similar to nidogen 2 and similar to MIC2 like 1 (LOC365084) mRNA.
<a href="#">LOC365476</a>	<a href="#">LOC365476.aSep08</a>	<a href="#">365476</a>	19419	2277		542	wd repeat-containing protein (LOC365476) mRNA.
<a href="#">LOC365601</a>	<a href="#">LOC365601.aSep08</a>	<a href="#">365601</a>	91536	1033	7	344	similar to autophagy 5-like (LOC365601) alternative variant aSep08, mRNA.
<a href="#">LOC365601</a>	<a href="#">LOC365601.bSep08</a>	<a href="#">365601</a>	74341	1935	6	241	similar to autophagy 5-like (LOC365601) alternative variant bSep08, mRNA.
<a href="#">LOC365601</a>	<a href="#">LOC365601.cSep08</a>	<a href="#">365601</a>	24978	587	3	162	similar to autophagy 5-like (LOC365601) alternative variant cSep08, mRNA.
<a href="#">LOC365601</a>	<a href="#">LOC365601.dSep08</a>	<a href="#">365601</a>	52573	714	4	151	similar to autophagy 5-like (LOC365601) alternative variant dSep08, mRNA.
<a href="#">LOC365778</a>	<a href="#">LOC365778.aSep08</a>	<a href="#">365778</a>	2092	1206	2	176	similar to RIKEN cDNA 1700034I23 (19.4 kD) (LOC365778) alternative variant aSep08, complete mRNA.
<a href="#">LOC365948</a>	<a href="#">LOC365948.aSep08</a>	<a href="#">365948</a>	160847	2100	11	442	putative protein, with a coiled coil domain, of bilateral origin (LOC365948) alternative variant aSep08, mRNA.
<a href="#">LOC365948</a>	<a href="#">LOC365948.bSep08</a>	<a href="#">365948</a>	43711	960	6	319	putative protein, with a coiled coil domain (LOC365948) alternative variant bSep08, mRNA.
<a href="#">LOC365948</a>	<a href="#">LOC365948.cSep08</a>	<a href="#">365948</a>	109222	762	4	142	putative protein of bilateral origin (LOC365948) alternative variant cSep08, mRNA.
<a href="#">LOC365948</a>	<a href="#">LOC365948.dSep08</a>	<a href="#">365948</a>	34219	589	7	135	putative protein, with a coiled coil domain (LOC365948) alternative variant dSep08, mRNA.
<a href="#">LOC365985</a>	<a href="#">LOC365985.bSep08</a>	<a href="#">365985</a>	21510	1516		156	similar to adenylate kinase 5 isoform 1 (LOC365985) alternative variant bSep08, mRNA.

<a href="#">LOC366057</a>	<a href="#">LOC366057.aSep08</a>	<a href="#">366057</a>	1898	1451		256	similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (27.1 kD) (LOC366057) complete mRNA.
<a href="#">LOC366300</a>	<a href="#">LOC366300.aSep08</a>	<a href="#">366300</a>	71869	4150	7	372	hypothetical LOC366300 (42.3 kD) (LOC366300) alternative variant aSep08, mRNA.
<a href="#">LOC366300</a>	<a href="#">LOC366300.bSep08</a>	<a href="#">366300</a>	22833	813	5	133	hypothetical LOC366300 (LOC366300) alternative variant bSep08, mRNA.
<a href="#">LOC366300</a>	<a href="#">LOC366300.cSep08</a>	<a href="#">366300</a>	16613	304	3	101	hypothetical LOC366300 (LOC366300) alternative variant cSep08, mRNA.
<a href="#">LOC366300</a>	<a href="#">LOC366300.eSep08</a>	<a href="#">366300</a>	8578	473	4	25	hypothetical LOC366300 (LOC366300) alternative variant eSep08, mRNA.
<a href="#">LOC366431</a>	<a href="#">LOC366431.cSep08</a>	<a href="#">366431</a>	31728	658	1	66	putative protein (LOC366431) alternative variant cSep08, mRNA.
<a href="#">LOC366473</a>	<a href="#">LOC366473.aSep08</a>	<a href="#">366473</a>	2689	460		83	similar to ornithine decarboxylase-like protein (LOC366473) mRNA.
<a href="#">LOC366515</a>	<a href="#">LOC366515.aSep08</a>	<a href="#">366515</a>	6052	447		140	similar to Wdr8 protein (LOC366515) mRNA.
<a href="#">LOC366669</a>	<a href="#">LOC366669.aSep08</a>	<a href="#">366669</a>	11797	2200		437	similar to mKIAA1011 protein (LOC366669) mRNA.
<a href="#">LOC366772</a>	<a href="#">LOC366772.aSep08</a>	<a href="#">366772</a>	5634	1191	3	303	similar to immunoglobulin heavy chain (32.9 kD) (LOC366772) alternative variant aSep08, mRNA.
<a href="#">LOC367196</a>	<a href="#">LOC367196.aSep08</a>	<a href="#">367196</a>	15130	770	6	232	similar to Acyl-CoA dehydrogenase family member 8, mitochondrial precursor (ACAD-8) (Isobutyryl-CoA dehydrogenase) (LOC367196) alternative variant aSep08, mRNA.
<a href="#">LOC367196</a>	<a href="#">LOC367196.bSep08</a>	<a href="#">367196</a>	8516	468	2	112	similar to Acyl-CoA dehydrogenase family member 8, mitochondrial precursor (ACAD-8) (Isobutyryl-CoA dehydrogenase) (12.3 kD) (LOC367196) alternative variant bSep08, mRNA.
<a href="#">LOC367196</a>	<a href="#">LOC367196.cSep08</a>	<a href="#">367196</a>	5338	719	2	92	similar to Acyl-CoA dehydrogenase family member 8, mitochondrial precursor (ACAD-8) (Isobutyryl-CoA dehydrogenase) (10.5 kD) (LOC367196) alternative variant cSep08, mRNA.
<a href="#">LOC367198</a>	<a href="#">LOC367198.aSep08</a>	<a href="#">367198</a>	7752	649		215	similar to Serine/threonine-protein kinase ATR (Ataxia telangiectasia and Rad3-related protein) (LOC367198) mRNA.
<a href="#">LOC367289</a>	<a href="#">LOC367289.aSep08</a>	<a href="#">367289</a>	73412	1446	7	481	similar to CG4699-PA, isoform A (LOC367289) alternative variant aSep08, mRNA.
<a href="#">LOC367289</a>	<a href="#">LOC367289.bSep08</a>	<a href="#">367289</a>	46986	693	5	230	similar to CG4699-PA, isoform A (LOC367289) alternative variant bSep08, mRNA.
<a href="#">LOC367289</a>	<a href="#">LOC367289.cSep08</a>	<a href="#">367289</a>	2789	499	2	102	similar to CG4699-PA, isoform A (11.7 kD) (LOC367289) alternative variant cSep08, mRNA.
<a href="#">LOC367409</a>	<a href="#">LOC367409.aSep08</a>	<a href="#">367409</a>	15364	1158		93	similar to Reticulocalbin-1 precursor and similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (10.8 kD) (LOC367409) mRNA.
<a href="#">LOC367409</a>	<a href="#">LOC367409.aSep08</a>	<a href="#">690474</a>	15364	1158		93	similar to Reticulocalbin-1 precursor and similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (10.8 kD) (LOC367409) mRNA.
<a href="#">LOC367597</a>	<a href="#">LOC367597.aSep08</a>	<a href="#">367597</a>	10415	759	1	79	similar to nidogen 2 (8.6 kD) (LOC367597) alternative variant aSep08, mRNA.

<a href="#">LOC367597</a>	<a href="#">LOC367597.bSep08</a>	<a href="#">367597</a>	9622	691	1	79	similar to nidogen 2 (8.6 kD) (LOC367597) alternative variant bSep08, mRNA.
<a href="#">LOC367653</a>	<a href="#">LOC367653.aSep08</a>	<a href="#">367653</a>	4704	807		141	similar to nidogen 2 (LOC367653) mRNA.
<a href="#">LOC367746</a>	<a href="#">LOC367746.aSep08</a>	<a href="#">367746</a>	1778	1067		335	similar to Spindlin-like protein 2 (SPIN-2) (LOC367746) mRNA.
<a href="#">LOC367808</a>	<a href="#">LOC367808.bSep08</a>	<a href="#">367808</a>	1280	366	3	38	similar to Sid3177p (4.2 kD) (LOC367808) alternative variant bSep08, mRNA.
<a href="#">LOC367858</a>	<a href="#">LOC367858.aSep08</a>	<a href="#">367858</a>	547	401		46	similar to GTPase HRas precursor (Transforming protein p21) (p21ras) (H-Ras-1) (c-H-ras) (5.2 kD) (LOC367858) mRNA.
<a href="#">LOC367976</a>	<a href="#">LOC367976.aSep08</a>	<a href="#">367976</a>	10406	2176	1	559	similar to DNA replication licensing factor MCM3 (DNA polymerase alpha holoenzyme-associated protein P1) (P1-MCM3) (63.1 kD) (LOC367976) alternative variant aSep08, mRNA.
<a href="#">LOC367976</a>	<a href="#">LOC367976.bSep08</a>	<a href="#">367976</a>	3678	710	2	236	similar to DNA replication licensing factor MCM3 (DNA polymerase alpha holoenzyme-associated protein P1) (P1-MCM3) (LOC367976) alternative variant bSep08, mRNA.
<a href="#">LOC367994</a>	<a href="#">LOC367994.bSep08</a>	<a href="#">367994</a>	2618	508	1	111	similar to uroplakin 3B isoform b (12.1 kD) (LOC367994) alternative variant bSep08, mRNA.
<a href="#">LOC368084</a>	<a href="#">LOC368084.aSep08</a>	<a href="#">368084</a>	11193	1258	12	281	methyltransferase type 11 and methyltransferase type 12 (31.6 kD) (LOC368084) alternative variant aSep08, complete mRNA.
<a href="#">LOC368084</a>	<a href="#">LOC368084.bSep08</a>	<a href="#">368084</a>	927	842	2	88	putative nuclear protein of mammalian origin (10.5 kD) (LOC368084) alternative variant bSep08, mRNA.
<a href="#">LOC368084</a>	<a href="#">LOC368084.cSep08</a>	<a href="#">368084</a>	2987	1555	4	77	putative protein of eukaryotic origin (LOC368084) alternative variant cSep08, mRNA.
<a href="#">LOC474147</a>	<a href="#">LOC474147.aSep08</a>	<a href="#">474147</a>	10161	671		138	RBSC-skeletrophin (LOC474147) mRNA.
<a href="#">LOC494499</a>	<a href="#">LOC494499.bSep08</a>	<a href="#">494499</a>	2907	730	2	98	LOC494499 protein (11.4 kD) (LOC494499) alternative variant bSep08, mRNA.
<a href="#">LOC497842</a>	<a href="#">LOC497842.aSep08</a>	<a href="#">497842</a>	49287	335		111	similar to guanylate cyclase 1, soluble, alpha 2 (LOC497842) mRNA.
<a href="#">LOC497940</a>	<a href="#">LOC497940.bSep08</a>	<a href="#">497940</a>	2114	976	2	75	similar to RIKEN cDNA 2810408A11 (LOC497940) alternative variant bSep08, mRNA.
<a href="#">LOC497959</a>	<a href="#">LOC497959.bSep08</a>	<a href="#">497959</a>	13769	1526	5	263	similar to novel protein (29.6 kD) (LOC497959) alternative variant bSep08, mRNA.
<a href="#">LOC497959</a>	<a href="#">LOC497959.cSep08</a>	<a href="#">497959</a>	13249	746	3	75	similar to novel protein (8.6 kD) (LOC497959) alternative variant cSep08, mRNA.
<a href="#">LOC497959</a>	<a href="#">LOC497959.dSep08</a>	<a href="#">497959</a>	2991	738	2	89	similar to novel protein (9.3 kD) (LOC497959) alternative variant dSep08, complete mRNA.
<a href="#">LOC497978</a>	<a href="#">LOC497978.bSep08</a>	<a href="#">497978</a>	12458	3224	3	230	similar to diacylglycerol kinase epsilon (26.0 kD) (LOC497978) alternative variant bSep08, mRNA.
<a href="#">LOC498029</a>	<a href="#">LOC498029.aSep08</a>	<a href="#">498029</a>	16323	1822	3	337	similar to RIKEN cDNA A730011L01 gene (LOC498029) alternative variant aSep08, mRNA.
<a href="#">LOC498029</a>	<a href="#">LOC498029.bSep08</a>	<a href="#">498029</a>	15840	1127	1	96	similar to RIKEN cDNA A730011L01 gene (15.8 kD) (LOC498029) alternative variant bSep08, mRNA.
<a href="#">LOC498029</a>	<a href="#">LOC498029.cSep08</a>	<a href="#">498029</a>	14590	775	1	96	similar to RIKEN cDNA A730011L01 gene (11.0 kD) (LOC498029) alternative variant cSep08, mRNA.

<a href="#">LOC498154</a>	<a href="#">LOC498154.bSep08</a>	<a href="#">498154</a>	14111	1363	4	176	hypothetical protein LOC498154 (LOC498154) alternative variant bSep08, mRNA.
<a href="#">LOC498154</a>	<a href="#">LOC498154.cSep08</a>	<a href="#">498154</a>	35715	621	4	137	hypothetical protein LOC498154 (15.9 kD) (LOC498154) alternative variant cSep08, mRNA.
<a href="#">LOC498154</a>	<a href="#">LOC498154.fSep08</a>	<a href="#">498154</a>	8845	393	2	51	hypothetical protein LOC498154 (LOC498154) alternative variant fSep08, mRNA.
<a href="#">LOC498201</a>	<a href="#">LOC498201.aSep08</a>	<a href="#">498201</a>	10107	793		166	hypothetical LOC498201 (LOC498201) mRNA.
<a href="#">LOC498236</a>	<a href="#">LOC498236.bSep08</a>	<a href="#">498236</a>	13510	765	6	254	LRRGT00186 (LOC498236) alternative variant bSep08, mRNA.
<a href="#">LOC498236</a>	<a href="#">LOC498236.cSep08</a>	<a href="#">498236</a>	2580	604	4	159	LRRGT00186 (LOC498236) alternative variant cSep08, mRNA.
<a href="#">LOC498276</a>	<a href="#">LOC498276.aSep08</a>	<a href="#">498276</a>	10400	1404	6	267	fc gamma receptor II beta (30.3 kD) (LOC498276) alternative variant aSep08, mRNA.
<a href="#">LOC498276</a>	<a href="#">LOC498276.bSep08</a>	<a href="#">498276</a>	8490	763	5	254	fc gamma receptor II beta (LOC498276) alternative variant bSep08, mRNA.
<a href="#">LOC498276</a>	<a href="#">LOC498276.cSep08</a>	<a href="#">498276</a>	8961	961	4	182	fc gamma receptor II beta (20.4 kD) (LOC498276) alternative variant cSep08, mRNA.
<a href="#">LOC498330</a>	<a href="#">LOC498330.aSep08</a>	<a href="#">498330</a>	7094	749	1	117	similar to hypothetical protein MGC26744 (13.3 kD) (LOC498330) alternative variant aSep08, mRNA.
<a href="#">LOC498350</a>	<a href="#">LOC498350.bSep08</a>	<a href="#">498350</a>	21996	793	1	199	CRA c (LOC498350) alternative variant bSep08, mRNA.
<a href="#">LOC498353</a>	<a href="#">LOC498353.bSep08</a>	<a href="#">498353</a>	75136	562	5	150	putative protein of eukaryotic origin (LOC498353) alternative variant bSep08, mRNA.
<a href="#">LOC498353</a>	<a href="#">LOC498353.dSep08</a>	<a href="#">498353</a>	8613	753	3	44	putative protein of vertebrate origin (4.9 kD) (LOC498353) alternative variant dSep08, mRNA.
<a href="#">LOC498400</a>	<a href="#">LOC498400.aSep08</a>	<a href="#">498400</a>	67458	1765	3	342	putative protein of eukaryotic origin (LOC498400) alternative variant aSep08, mRNA.
<a href="#">LOC498400</a>	<a href="#">LOC498400.cSep08</a>	<a href="#">498400</a>	59214	1136	1	64	putative protein of vertebrate origin (7.2 kD) (LOC498400) alternative variant cSep08, mRNA.
<a href="#">LOC498435</a>	<a href="#">LOC498435.aSep08</a>	<a href="#">498435</a>	8555	1211		170	similar to ATP-binding cassette sub-family G member 3 (19.4 kD) (LOC498435) mRNA.
<a href="#">LOC498465</a>	<a href="#">LOC498465.aSep08</a>	<a href="#">498465</a>	3335	756		204	similar to RIKEN cDNA 1700001F09 (23.9 kD) (LOC498465) mRNA.
<a href="#">LOC498601</a>	<a href="#">LOC498601.aSep08</a>	<a href="#">498601</a>	4214	458		124	similar to cyclin B2 (LOC498601) alternative variant aSep08, mRNA.
<a href="#">LOC498601</a>	<a href="#">LOC498601.bSep08</a>	<a href="#">498601</a>	4275	619	1	94	similar to cyclin B2 (LOC498601) alternative variant bSep08, mRNA.
<a href="#">LOC498604</a>	<a href="#">LOC498604.aSep08</a>	<a href="#">498604</a>	2913	750		249	similar to B-cell novel protein 1 (LOC498604) mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.aSep08</a>	<a href="#">498606</a>	13677	1226	5	227	CRA a (LOC498606) alternative variant aSep08, mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.cSep08</a>	<a href="#">498606</a>	9102	1096	6	177	CRA a (19.9 kD) (LOC498606) alternative variant cSep08, mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.dSep08</a>	<a href="#">498606</a>	8859	817	5	173	CRA a (LOC498606) alternative variant dSep08, mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.eSep08</a>	<a href="#">498606</a>	6480	1651	3	145	protein CRA a (LOC498606) alternative variant eSep08, mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.fSep08</a>	<a href="#">498606</a>	6189	1599	5	108	protein CRA a (LOC498606) alternative variant fSep08, mRNA.

<a href="#">LOC498606</a>	<a href="#">LOC498606.gSep08</a>	<a href="#">498606</a>	9393	1491	4	96	protein CRA a (11.4 kD) (LOC498606) alternative variant gSep08, complete mRNA.
<a href="#">LOC498606</a>	<a href="#">LOC498606.hSep08</a>	<a href="#">498606</a>	1906	818	2	91	protein CRA a (9.8 kD) (LOC498606) alternative variant hSep08, mRNA.
<a href="#">LOC498662</a>	<a href="#">LOC498662.aSep08</a>	<a href="#">498662</a>	28475	2626		182	similar to RIKEN cDNA 2610019F03 (20.2 kD) (LOC498662) alternative variant aSep08, complete mRNA.
<a href="#">LOC498664</a>	<a href="#">LOC498664.bSep08</a>	<a href="#">498664</a>	13162	349	4	69	similar to hypothetical protein MGC35169 (LOC498664) alternative variant bSep08, mRNA.
<a href="#">LOC498664</a>	<a href="#">LOC498664.cSep08</a>	<a href="#">498664</a>	430	314	1	67	similar to hypothetical protein MGC35169 (LOC498664) alternative variant cSep08, mRNA.
<a href="#">LOC498685</a>	<a href="#">LOC498685.aSep08</a>	<a href="#">498685</a>	17487	767	6	195	similar to UPF0308 protein C9orf21 (LOC498685) alternative variant aSep08, mRNA.
<a href="#">LOC498685</a>	<a href="#">LOC498685.bSep08</a>	<a href="#">498685</a>	17677	825	5	151	similar to UPF0308 protein C9orf21 (LOC498685) alternative variant bSep08, mRNA.
<a href="#">LOC498735</a>	<a href="#">LOC498735.aSep08</a>	<a href="#">498735</a>	10284	988		240	similar to hypothetical protein MGC43581 (LOC498735) alternative variant aSep08, mRNA.
<a href="#">LOC498750</a>	<a href="#">LOC498750.aSep08</a>	<a href="#">498750</a>	11957	4033	5	229	putative protein, with 2 coiled coil domains, of metazoan origin (27.1 kD) (LOC498750) alternative variant aSep08, mRNA.
<a href="#">LOC498750</a>	<a href="#">LOC498750.bSep08</a>	<a href="#">498750</a>	2772	800	2	90	putative endoplasmic reticulum protein (10.2 kD) (LOC498750) alternative variant bSep08, mRNA.
<a href="#">LOC498781</a>	<a href="#">LOC498781.aSep08</a>	<a href="#">498781</a>	6800	782		191	similar to serine/threonine kinase (LOC498781) mRNA.
<a href="#">LOC498793</a>	<a href="#">LOC498793.aSep08</a>	<a href="#">498793</a>	36911	3070	19	995	similar to inter-alpha-inhibitor H2 chain (LOC498793) alternative variant aSep08, mRNA.
<a href="#">LOC498909</a>	<a href="#">LOC498909.aSep08</a>	<a href="#">498909</a>	3363	1798	3	371	similar to RIKEN cDNA 2310005O14 (LOC498909) alternative variant aSep08, mRNA.
<a href="#">LOC498909</a>	<a href="#">LOC498909.cSep08</a>	<a href="#">498909</a>	8796	673	5	163	similar to RIKEN cDNA 2310005O14 (LOC498909) alternative variant cSep08, mRNA.
<a href="#">LOC498909</a>	<a href="#">LOC498909.fSep08</a>	<a href="#">498909</a>	3985	561	4	87	similar to RIKEN cDNA 2310005O14 (LOC498909) alternative variant fSep08, mRNA.
<a href="#">LOC498951</a>	<a href="#">LOC498951.aSep08</a>	<a href="#">498951</a>	48682	1653	6	235	similar to 2010004A03Rik protein (26.9 kD) (LOC498951) alternative variant aSep08, complete mRNA.
<a href="#">LOC498951</a>	<a href="#">LOC498951.cSep08</a>	<a href="#">498951</a>	5816	302	2	100	similar to 2010004A03Rik protein (LOC498951) alternative variant cSep08, mRNA.
<a href="#">LOC498957</a>	<a href="#">LOC498957.bSep08</a>	<a href="#">498957</a>	7970	712	1	182	kelch-like 36 (LOC498957) alternative variant bSep08, mRNA.
<a href="#">LOC498972</a>	<a href="#">LOC498972.aSep08</a>	<a href="#">498972</a>	7338	929		176	similar to copine II (LOC498972) mRNA.
<a href="#">LOC498974</a>	<a href="#">LOC498974.aSep08</a>	<a href="#">498974</a>	1417	1165		158	similar to kinesin family member 7 (LOC498974) mRNA.
<a href="#">LOC499089</a>	<a href="#">LOC499089.aSep08</a>	<a href="#">499089</a>	8411	1074	4	342	similar to pregnancy-specific glycoprotein 25 (LOC499089) alternative variant aSep08, mRNA.
<a href="#">LOC499089</a>	<a href="#">LOC499089.bSep08</a>	<a href="#">499089</a>	2659	624	2	135	similar to pregnancy-specific glycoprotein 25 (15.2 kD) (LOC499089) alternative variant bSep08, mRNA.
<a href="#">LOC499110</a>	<a href="#">LOC499110.aSep08</a>	<a href="#">499110</a>	16398	571		173	similar to Zinc finger protein 354A (Transcription factor 17) (Renal transcription factor Kid-1) (Kidney, ischemia, and developmentally regulated protein 1) (LOC499110) mRNA.
<a href="#">LOC499120</a>	<a href="#">LOC499120.aSep08</a>	<a href="#">499120</a>	12434	2481	5	474	hypothetical protein LOC499120 (54.4 kD) (LOC499120) alternative variant aSep08, mRNA.

<a href="#">LOC499120</a>	<a href="#">LOC499120.bSep08</a>	<a href="#">499120</a>	9004	781	3	191	hypothetical protein LOC499120 (LOC499120) alternative variant bSep08, mRNA.
<a href="#">LOC499120</a>	<a href="#">LOC499120.cSep08</a>	<a href="#">499120</a>	8467	397	2	132	hypothetical protein LOC499120 (LOC499120) alternative variant cSep08, mRNA.
<a href="#">LOC499234</a>	<a href="#">LOC499234.aSep08</a>	<a href="#">499234</a>	5108	762		177	similar to NACHT, leucine rich repeat and PYD containing 14-like (LOC499234) mRNA.
<a href="#">LOC499276</a>	<a href="#">LOC499276.aSep08</a>	<a href="#">499276</a>	2464	738		245	similar to RIKEN cDNA 1700022C21 (LOC499276) mRNA.
<a href="#">LOC499279</a>	<a href="#">LOC499279.bSep08</a>	<a href="#">499279</a>	1687	269	1	35	hypothetical gene supported by BC079265 (LOC499279) alternative variant bSep08, mRNA.
<a href="#">LOC499315</a>	<a href="#">LOC499315.aSep08</a>	<a href="#">499315</a>	68331	981	6	186	AHNAK 1 (20.1 kD) (LOC499315) alternative variant aSep08, mRNA.
<a href="#">LOC499315</a>	<a href="#">LOC499315.bSep08</a>	<a href="#">499315</a>	1058	206	2	68	AHNAK 1 (LOC499315) alternative variant bSep08, mRNA.
<a href="#">LOC499315</a>	<a href="#">LOC499315.cSep08</a>	<a href="#">499315</a>	42362	742	2	41	AHNAK 1 (LOC499315) alternative variant cSep08, mRNA.
<a href="#">LOC499330</a>	<a href="#">LOC499330.aSep08</a>	<a href="#">499330</a>	25069	836	1	224	similar to Nicotinamide riboside kinase 1 (LOC499330) alternative variant aSep08, mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.bSep08</a>	<a href="#">499339</a>	3549	458	3	152	hypothetical protein LOC499339 (LOC499339) alternative variant bSep08, mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.cSep08</a>	<a href="#">499339</a>	36896	749	5	108	hypothetical protein LOC499339 (11.8 kD) (LOC499339) alternative variant cSep08, complete mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.dSep08</a>	<a href="#">499339</a>	5768	557	3	55	hypothetical protein LOC499339 (6.4 kD) (LOC499339) alternative variant dSep08, mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.eSep08</a>	<a href="#">499339</a>	6625	904	3	62	hypothetical protein LOC499339 (6.7 kD) (LOC499339) alternative variant eSep08, mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.fSep08</a>	<a href="#">499339</a>	4371	654	2	62	hypothetical protein LOC499339 (6.7 kD) (LOC499339) alternative variant fSep08, mRNA.
<a href="#">LOC499339</a>	<a href="#">LOC499339.hSep08</a>	<a href="#">499339</a>	3634	275	3	51	hypothetical protein LOC499339 (LOC499339) alternative variant hSep08, mRNA.
<a href="#">LOC499376</a>	<a href="#">LOC499376.aSep08</a>	<a href="#">499376</a>	582	309		70	hypothetical LOC499376 (LOC499376) mRNA.
<a href="#">LOC499391</a>	<a href="#">LOC499391.bSep08</a>	<a href="#">499391</a>	9139	595	2	129	similar to gem (nuclear organelle) associated protein 7 (14.3 kD) (LOC499391) alternative variant bSep08, mRNA.
<a href="#">LOC499587</a>	<a href="#">LOC499587.aSep08</a>	<a href="#">499587</a>	60258	532		78	similar to solute carrier family 7, member 14 (LOC499587) mRNA.
<a href="#">LOC499602</a>	<a href="#">LOC499602.aSep08</a>	<a href="#">499602</a>	15950	1300	9	366	hypothetical protein LOC499602 (LOC499602) alternative variant aSep08, mRNA.
<a href="#">LOC499602</a>	<a href="#">LOC499602.bSep08</a>	<a href="#">499602</a>	1585	615	2	79	hypothetical protein LOC499602 (LOC499602) alternative variant bSep08, mRNA.
<a href="#">LOC499602</a>	<a href="#">LOC499602.dSep08</a>	<a href="#">499602</a>	8488	1157	5	61	hypothetical protein LOC499602 (7.5 kD) (LOC499602) alternative variant dSep08, mRNA.
<a href="#">LOC499602</a>	<a href="#">LOC499602.eSep08</a>	<a href="#">499602</a>	14009	530	2	59	hypothetical protein LOC499602 (LOC499602) alternative variant eSep08, mRNA.
<a href="#">LOC499618</a>	<a href="#">LOC499618.bSep08</a>	<a href="#">499618</a>	20664	711	7	191	CRA a (LOC499618) alternative variant bSep08, mRNA.
<a href="#">LOC499618</a>	<a href="#">LOC499618.cSep08</a>	<a href="#">499618</a>	1518	352	3	43	CRA b like (4.8 kD) (LOC499618) alternative variant cSep08, mRNA.
<a href="#">LOC499653</a>	<a href="#">LOC499653.bSep08</a>	<a href="#">499653</a>	19923	700	1	233	dingo protein (LOC499653) alternative variant bSep08, mRNA.



<a href="#">LOC499677</a>	<a href="#">LOC499677.bSep08</a>	<a href="#">499677</a>	13433	3182	5	240	similar to Lix1 homolog (mouse) like (LOC499677) alternative variant bSep08, mRNA.
<a href="#">LOC499677</a>	<a href="#">LOC499677.cSep08</a>	<a href="#">499677</a>	11170	778	4	193	similar to Lix1 homolog (mouse) like (LOC499677) alternative variant cSep08, mRNA.
<a href="#">LOC499746</a>	<a href="#">LOC499746.aSep08</a>	<a href="#">499746</a>	1295	752	2	250	similar to hypothetical gene supported by AK097565; BC033939 (LOC499746) alternative variant aSep08, mRNA.
<a href="#">LOC499746</a>	<a href="#">LOC499746.bSep08</a>	<a href="#">499746</a>	6479	1673	3	151	similar to hypothetical gene supported by AK097565; BC033939 (17.1 kD) (LOC499746) alternative variant bSep08, mRNA.
<a href="#">LOC499749</a>	<a href="#">LOC499749.bSep08</a>	<a href="#">499749</a>	1181	675	2	103	similar to RIKEN cDNA C430004E15 (11.1 kD) (LOC499749) alternative variant bSep08, mRNA.
<a href="#">LOC499770</a>	<a href="#">LOC499770.bSep08</a>	<a href="#">499770</a>	3746	760	7	174	similar to LOC495800 protein (LOC499770) alternative variant bSep08, mRNA.
<a href="#">LOC499770</a>	<a href="#">LOC499770.dSep08</a>	<a href="#">499770</a>	3252	378	5	125	similar to LOC495800 protein (LOC499770) alternative variant dSep08, mRNA.
<a href="#">LOC499770</a>	<a href="#">LOC499770.eSep08</a>	<a href="#">499770</a>	1811	1118	4	90	similar to LOC495800 protein (LOC499770) alternative variant eSep08, mRNA.
<a href="#">LOC499779</a>	<a href="#">LOC499779.aSep08</a>	<a href="#">499779</a>	4117	724	1	89	similar to RIKEN cDNA 2900010J23 (10.3 kD) (LOC499779) alternative variant aSep08, mRNA.
<a href="#">LOC499779</a>	<a href="#">LOC499779.bSep08</a>	<a href="#">499779</a>	9132	748	2	89	similar to RIKEN cDNA 2900010J23 (10.3 kD) (LOC499779) alternative variant bSep08, mRNA.
<a href="#">LOC499781</a>	<a href="#">LOC499781.aSep08</a>	<a href="#">499781</a>	4751	740		246	similar to CG17122-PA (LOC499781) mRNA.
<a href="#">LOC499782</a>	<a href="#">LOC499782.bSep08</a>	<a href="#">499782</a>	2342	839	7	138	ribosomal protein L11 (15.0 kD) (LOC499782) alternative variant bSep08, complete mRNA.
<a href="#">LOC499782</a>	<a href="#">LOC499782.cSep08</a>	<a href="#">499782</a>	2407	1971	2	127	putative protein (LOC499782) alternative variant cSep08, mRNA.
<a href="#">LOC499782</a>	<a href="#">LOC499782.dSep08</a>	<a href="#">499782</a>	2349	835	7	118	ribosomal protein L11 (12.9 kD) (LOC499782) alternative variant dSep08, complete mRNA.
<a href="#">LOC499782</a>	<a href="#">LOC499782.fSep08</a>	<a href="#">499782</a>	1808	724	4	80	ribosomal protein L11 (LOC499782) alternative variant fSep08, mRNA.
<a href="#">LOC499806</a>	<a href="#">LOC499806.bSep08</a>	<a href="#">499806</a>	34884	1304	11	336	similar to RIKEN cDNA 4933404M02 (LOC499806) alternative variant bSep08, mRNA.
<a href="#">LOC499806</a>	<a href="#">LOC499806.cSep08</a>	<a href="#">499806</a>	36633	2057	11	252	similar to RIKEN cDNA 4933404M02 (26.8 kD) (LOC499806) alternative variant cSep08, mRNA.
<a href="#">LOC499806</a>	<a href="#">LOC499806.dSep08</a>	<a href="#">499806</a>	11104	447	3	79	similar to RIKEN cDNA 4933404M02 (LOC499806) alternative variant dSep08, mRNA.
<a href="#">LOC499806</a>	<a href="#">LOC499806.eSep08</a>	<a href="#">499806</a>	13164	556	3		
<a href="#">LOC500007</a>	<a href="#">LOC500007.aSep08</a>	<a href="#">500007</a>	6369	296		31	similar to EF hand calcium binding domain 1 (3.6 kD) (LOC500007) mRNA.
<a href="#">LOC500034</a>	<a href="#">LOC500034.aSep08</a>	<a href="#">500034</a>	65879	1798	4	454	similar to CG3570-PA (LOC500034) alternative variant aSep08, mRNA.
<a href="#">LOC500034</a>	<a href="#">LOC500034.cSep08</a>	<a href="#">500034</a>	21994	2711	3	314	similar to CG3570-PA (LOC500034) alternative variant cSep08, mRNA.
<a href="#">LOC500034</a>	<a href="#">LOC500034.eSep08</a>	<a href="#">500034</a>	44816	417	2	138	similar to CG3570-PA (LOC500034) alternative variant eSep08, mRNA.

<a href="#">LOC500046</a>	<a href="#">LOC500046.aSep08</a>	<a href="#">500046</a>	31794	1323		145	similar to hypothetical protein FLJ21986 (LOC500046) mRNA.
<a href="#">LOC500054</a>	<a href="#">LOC500054.bSep08</a>	<a href="#">500054</a>	46529	1156	8	241	protection of telomeres 1A (27.0 kD) (LOC500054) alternative variant bSep08, mRNA.
<a href="#">LOC500054</a>	<a href="#">LOC500054.cSep08</a>	<a href="#">500054</a>	7260	513	6	170	protection of telomeres 1A (LOC500054) alternative variant cSep08, mRNA.
<a href="#">LOC500054</a>	<a href="#">LOC500054.dSep08</a>	<a href="#">500054</a>	3437	881	3	90	protection of telomeres 1A (LOC500054) alternative variant dSep08, mRNA.
<a href="#">LOC500054</a>	<a href="#">LOC500054.eSep08</a>	<a href="#">500054</a>	5772	708	2	97	putative protein (LOC500054) alternative variant eSep08, mRNA.
<a href="#">LOC500066</a>	<a href="#">LOC500066.aSep08</a>	<a href="#">500066</a>	1443	332		110	similar to Protein FAM40B (LOC500066) mRNA.
<a href="#">LOC500105</a>	<a href="#">LOC500105.aSep08</a>	<a href="#">500105</a>	283582	622		191	similar to contactin associated protein-like 2 isoform a (LOC500105) mRNA.
<a href="#">LOC500118</a>	<a href="#">LOC500118.aSep08</a>	<a href="#">500118</a>	14601	776	6	258	similar to RIKEN cDNA D330028D13 (LOC500118) alternative variant aSep08, mRNA.
<a href="#">LOC500118</a>	<a href="#">LOC500118.cSep08</a>	<a href="#">500118</a>	21248	1131	6	206	similar to RIKEN cDNA D330028D13 (LOC500118) alternative variant cSep08, mRNA.
<a href="#">LOC500118</a>	<a href="#">LOC500118.dSep08</a>	<a href="#">500118</a>	6743	752	2	51	similar to RIKEN cDNA D330028D13 (5.8 kD) (LOC500118) alternative variant dSep08, mRNA.
<a href="#">LOC500124</a>	<a href="#">LOC500124.bSep08</a>	<a href="#">500124</a>	16034	799	4	198	similar to RIKEN cDNA 4921507P07 (LOC500124) alternative variant bSep08, mRNA.
<a href="#">LOC500124</a>	<a href="#">LOC500124.cSep08</a>	<a href="#">500124</a>	13320	736	5	182	similar to RIKEN cDNA 4921507P07 (LOC500124) alternative variant cSep08, mRNA.
<a href="#">LOC500124</a>	<a href="#">LOC500124.dSep08</a>	<a href="#">500124</a>	13289	691	4	141	similar to RIKEN cDNA 4921507P07 (16.5 kD) (LOC500124) alternative variant dSep08, mRNA.
<a href="#">LOC500227</a>	<a href="#">LOC500227.bSep08</a>	<a href="#">500227</a>	1290	1171	1	314	hypothetical gene supported by BC079424 (LOC500227) alternative variant bSep08, mRNA.
<a href="#">LOC500227</a>	<a href="#">LOC500227.cSep08</a>	<a href="#">500227</a>	5748	1226	2	222	hypothetical gene supported by BC079424 (LOC500227) alternative variant cSep08, mRNA.
<a href="#">LOC500251</a>	<a href="#">LOC500251.bSep08</a>	<a href="#">500251</a>	3474	801	2	145	hypothetical protein LOC500251 (LOC500251) alternative variant bSep08, mRNA.
<a href="#">LOC500251</a>	<a href="#">LOC500251.cSep08</a>	<a href="#">500251</a>	3457	421	1	109	hypothetical protein LOC500251 (LOC500251) alternative variant cSep08, mRNA.
<a href="#">LOC500378</a>	<a href="#">LOC500378.bSep08</a>	<a href="#">500378</a>	829	691	2	208	similar to Protein C1orf77 homolog (LOC500378) alternative variant bSep08, mRNA.
<a href="#">LOC500392</a>	<a href="#">LOC500392.bSep08</a>	<a href="#">500392</a>	1119	512	4	102	similar to hypothetical protein FLJ25692 (11.5 kD) (LOC500392) alternative variant bSep08, mRNA.
<a href="#">LOC500413</a>	<a href="#">LOC500413.aSep08</a>	<a href="#">500413</a>	72855	873	2	86	hypothetical protein LOC500413 (9.7 kD) (LOC500413) alternative variant aSep08, mRNA.
<a href="#">LOC500413</a>	<a href="#">LOC500413.bSep08</a>	<a href="#">500413</a>	73509	1532	2	86	hypothetical protein LOC500413 (9.7 kD) (LOC500413) alternative variant bSep08, mRNA.
<a href="#">LOC500413</a>	<a href="#">LOC500413.cSep08</a>	<a href="#">500413</a>	72415	435	2	58	hypothetical protein LOC500413 (LOC500413) alternative variant cSep08, mRNA.
<a href="#">LOC500415</a>	<a href="#">LOC500415.bSep08</a>	<a href="#">500415</a>	76605	748	1	188	similar to Leucine-rich repeat protein SHOC-2 (Ras-binding protein Sur-8) (LOC500415) alternative variant bSep08, mRNA.

<a href="#">LOC500420</a>	<a href="#">LOC500420.aSep08</a>	<a href="#">500420</a>	9442	1512	2	157	similar to CG12279-PA (17.5 kD) (LOC500420) alternative variant aSep08, complete mRNA.
<a href="#">LOC500420</a>	<a href="#">LOC500420.bSep08</a>	<a href="#">500420</a>	7546	462	1	132	similar to CG12279-PA (LOC500420) alternative variant bSep08, mRNA.
<a href="#">LOC500445</a>	<a href="#">LOC500445.bSep08</a>	<a href="#">500445</a>	5823	1654	5	248	similar to hypothetical protein 4931430D02 (27.5 kD) (LOC500445) alternative variant bSep08, complete mRNA.
<a href="#">LOC500475</a>	<a href="#">LOC500475.bSep08</a>	<a href="#">500475</a>	1745	316	1	47	similar to hypothetical protein 4933430I17 (LOC500475) alternative variant bSep08, mRNA.
<a href="#">LOC500502</a>	<a href="#">LOC500502.aSep08</a>	<a href="#">500502</a>	1368	343		103	similar to RIKEN cDNA 4930579C15 (LOC500502) mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.aSep08</a>	<a href="#">500532</a>	17692	1895	11	282	ring finger protein 220 (31.8 kD) (LOC500532) alternative variant aSep08, complete mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.bSep08</a>	<a href="#">500532</a>	11718	705	6	178	finger protein 220 (LOC500532) alternative variant bSep08, mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.cSep08</a>	<a href="#">500532</a>	11856	741	7	160	ring finger protein 220 (LOC500532) alternative variant cSep08, mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.dSep08</a>	<a href="#">500532</a>	12068	764	8	160	ring finger protein 220 (LOC500532) alternative variant dSep08, mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.eSep08</a>	<a href="#">500532</a>	4291	2095	3	149	putative protein (LOC500532) alternative variant eSep08, mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.fSep08</a>	<a href="#">500532</a>	1295	486	3	105	ring finger protein 220 (LOC500532) alternative variant fSep08, mRNA.
<a href="#">LOC500532</a>	<a href="#">LOC500532.hSep08</a>	<a href="#">500532</a>	3787	335	3	47	ring finger protein 220 (LOC500532) alternative variant hSep08, mRNA.
<a href="#">LOC500584</a>	<a href="#">LOC500584.aSep08</a>	<a href="#">500584</a>	2369	1366	3	125	similar to casein kinase 1, gamma 3 isoform 2 (LOC500584) alternative variant aSep08, mRNA.
<a href="#">LOC500584</a>	<a href="#">LOC500584.bSep08</a>	<a href="#">500584</a>	21710	2204	5	78	similar to casein kinase 1, gamma 3 isoform 2 (9.0 kD) (LOC500584) alternative variant bSep08, complete mRNA.
<a href="#">LOC500591</a>	<a href="#">LOC500591.aSep08</a>	<a href="#">500591</a>	58141	1470		264	similar to calmodulin-binding transcription activator 1 (LOC500591) mRNA.
<a href="#">LOC500598</a>	<a href="#">LOC500598.bSep08</a>	<a href="#">500598</a>	682	618	1	114	tumor necrosis factor receptor superfamily member 18 CRA c like (12.9 kD) (LOC500598) alternative variant bSep08, mRNA.
<a href="#">LOC500668</a>	<a href="#">LOC500668.aSep08</a>	<a href="#">500668</a>	26656	626		155	similar to Centromeric protein E (CENP-E protein) (LOC500668) mRNA.
<a href="#">LOC500700</a>	<a href="#">LOC500700.aSep08</a>	<a href="#">500700</a>	42096	699		205	CRA b (LOC500700) mRNA.
<a href="#">LOC500705</a>	<a href="#">LOC500705.aSep08</a>	<a href="#">500705</a>	33833	475		158	calcium-binding EF-hand containing protein (LOC500705) mRNA.
<a href="#">LOC500726</a>	<a href="#">LOC500726.aSep08</a>	<a href="#">500726</a>	3779	2111		703	similar to KARP-1 binding protein 1 (LOC500726) mRNA.
<a href="#">LOC500797</a>	<a href="#">LOC500797.bSep08</a>	<a href="#">500797</a>	1508	265	1	88	hypothetical LOC500797 (LOC500797) alternative variant bSep08, mRNA.
<a href="#">LOC500797</a>	<a href="#">LOC500797.cSep08</a>	<a href="#">500797</a>	5160	341	2	37	hypothetical LOC500797 (LOC500797) alternative variant cSep08, mRNA.
<a href="#">LOC500827</a>	<a href="#">LOC500827.bSep08</a>	<a href="#">500827</a>	23224	780	5	259	similar to hypothetical protein FLJ35821 (LOC500827) alternative variant bSep08, mRNA.
<a href="#">LOC500827</a>	<a href="#">LOC500827.cSep08</a>	<a href="#">500827</a>	15092	652	2	97	similar to hypothetical protein FLJ35821 (LOC500827) alternative variant cSep08, mRNA.

<a href="#">LOC500947</a>	<a href="#">LOC500947.aSep08</a>	<a href="#">500947</a>	20236	1863		199	hypothetical gene supported by BC088439 (22.5 kD) (LOC500947) mRNA.
<a href="#">LOC500956</a>	<a href="#">LOC500956.bSep08</a>	<a href="#">500956</a>	3983	794	3	191	hypothetical protein LOC500956 (LOC500956) alternative variant bSep08, mRNA.
<a href="#">LOC500956</a>	<a href="#">LOC500956.dSep08</a>	<a href="#">500956</a>	1996	547	1	91	hypothetical protein LOC500956 (LOC500956) alternative variant dSep08, mRNA.
<a href="#">LOC500974</a>	<a href="#">LOC500974.aSep08</a>	<a href="#">500974</a>	26174	344		114	putative protein of eukaryotic origin (LOC500974) mRNA.
<a href="#">LOC500990</a>	<a href="#">LOC500990.bSep08</a>	<a href="#">500990</a>	9217	703	3	205	similar to RIKEN cDNA 4931429L15 (LOC500990) alternative variant bSep08, mRNA.
<a href="#">LOC500990</a>	<a href="#">LOC500990.cSep08</a>	<a href="#">500990</a>	10328	807	3	179	similar to RIKEN cDNA 4931429L15 (LOC500990) alternative variant cSep08, mRNA.
<a href="#">LOC501046</a>	<a href="#">LOC501046.aSep08</a>	<a href="#">501046</a>	29194	820	4	194	similar to Phakinin (Beaded filament structural protein 2) (Lens fiber cell beaded filament protein CP 49) (CP49) (49 kDa cytoskeletal protein) (LOC501046) alternative variant aSep08, mRNA.
<a href="#">LOC501046</a>	<a href="#">LOC501046.bSep08</a>	<a href="#">501046</a>	7811	712	1	179	similar to Phakinin (Beaded filament structural protein 2) (Lens fiber cell beaded filament protein CP 49) (CP49) (49 kDa cytoskeletal protein) (LOC501046) alternative variant bSep08, mRNA.
<a href="#">LOC501091</a>	<a href="#">LOC501091.aSep08</a>	<a href="#">501091</a>	17763	1044	1	243	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC501091) alternative variant aSep08, mRNA.
<a href="#">LOC501091</a>	<a href="#">LOC501091.bSep08</a>	<a href="#">501091</a>	23921	1026	1	157	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (18.7 kD) (LOC501091) alternative variant bSep08, mRNA.
<a href="#">LOC501092</a>	<a href="#">LOC501092.aSep08</a>	<a href="#">501092</a>	19649	453	2	108	hypothetical LOC501092 (LOC501092) alternative variant aSep08, mRNA.
<a href="#">LOC501092</a>	<a href="#">LOC501092.bSep08</a>	<a href="#">501092</a>	3865	306	1	41	hypothetical LOC501092 (LOC501092) alternative variant bSep08, mRNA.
<a href="#">LOC501098</a>	<a href="#">LOC501098.aSep08</a>	<a href="#">501098</a>	3424	1537		108	similar to ubiquitin specific protease 49 (LOC501098) mRNA.
<a href="#">LOC501110</a>	<a href="#">LOC501110.bSep08</a>	<a href="#">501110</a>	51852	719	6	78	similar to Glutathione S-transferase A1 (GTH1) (HA subunit 1) (GST-epsilon) (GSTA1-1) (GST class-alpha) (LOC501110) alternative variant bSep08, mRNA.
<a href="#">LOC501110</a>	<a href="#">LOC501110.cSep08</a>	<a href="#">501110</a>	20659	669	2	63	similar to Glutathione S-transferase A1 (GTH1) (HA subunit 1) (GST-epsilon) (GSTA1-1) (GST class-alpha) (7.3 kD) (LOC501110) alternative variant cSep08, mRNA.
<a href="#">LOC501126</a>	<a href="#">LOC501126.aSep08</a>	<a href="#">501126</a>	67002	2019	14	611	putative protein of metazoan origin (LOC501126) alternative variant aSep08, mRNA.
<a href="#">LOC501126</a>	<a href="#">LOC501126.bSep08</a>	<a href="#">501126</a>	21780	1026	8	266	putative nuclear protein of metazoan origin (30.1 kD) (LOC501126) alternative variant bSep08, mRNA.
<a href="#">LOC501126</a>	<a href="#">LOC501126.cSep08</a>	<a href="#">501126</a>	27438	988	6	224	putative protein of metazoan origin (25.5 kD) (LOC501126) alternative variant cSep08, mRNA.
<a href="#">LOC501126</a>	<a href="#">LOC501126.dSep08</a>	<a href="#">501126</a>	26376	747	6	212	putative protein of metazoan origin (LOC501126) alternative variant dSep08, mRNA.
<a href="#">LOC501126</a>	<a href="#">LOC501126.eSep08</a>	<a href="#">501126</a>	26360	728	6	206	putative protein of metazoan origin (LOC501126) alternative variant eSep08, mRNA.

<a href="#">LOC501126</a>	<a href="#">LOC501126.fSep08</a>	<a href="#">501126</a>	26376	721	6	203	putative protein of metazoan origin (LOC501126) alternative variant fSep08, mRNA.
<a href="#">LOC501194</a>	<a href="#">LOC501194.bSep08</a>	<a href="#">501194</a>	3436	648	3	163	similar to hypothetical protein D330021B20 and hypothetical protein LOC685917 (LOC501194) alternative variant bSep08, mRNA.
<a href="#">LOC501194</a>	<a href="#">LOC501194.bSep08</a>	<a href="#">685917</a>	3436	648	3	163	similar to hypothetical protein D330021B20 and hypothetical protein LOC685917 (LOC501194) alternative variant bSep08, mRNA.
<a href="#">LOC501212</a>	<a href="#">LOC501212.aSep08</a>	<a href="#">501212</a>	8884	341		113	hypothetical LOC501212 (LOC501212) mRNA.
<a href="#">LOC501221</a>	<a href="#">LOC501221.aSep08</a>	<a href="#">501221</a>	16045	689	2	81	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (8.7 kD) (LOC501221) alternative variant aSep08, mRNA.
<a href="#">LOC501221</a>	<a href="#">LOC501221.bSep08</a>	<a href="#">501221</a>	20903	492	2	36	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (4.0 kD) (LOC501221) alternative variant bSep08, mRNA.
<a href="#">LOC501223</a>	<a href="#">LOC501223.aSep08</a>	<a href="#">501223</a>	4656	881		35	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC501223) mRNA.
<a href="#">LOC501226</a>	<a href="#">LOC501226.aSep08</a>	<a href="#">501226</a>	2492	437		145	hypothetical LOC501226 (LOC501226) mRNA.
<a href="#">LOC501256</a>	<a href="#">LOC501256.aSep08</a>	<a href="#">501256</a>	17005	761		91	hypothetical LOC501256 (LOC501256) mRNA.
<a href="#">LOC501271</a>	<a href="#">LOC501271.aSep08</a>	<a href="#">501271</a>	87575	469		79	hypothetical LOC501271 (8.6 kD) (LOC501271) mRNA.
<a href="#">LOC501282</a>	<a href="#">LOC501282.aSep08</a>	<a href="#">501282</a>	8113	1714	6	121	similar to lymphocyte antigen 6 complex, locus E ligand (13.0 kD) (LOC501282) alternative variant aSep08, complete mRNA.
<a href="#">LOC501282</a>	<a href="#">LOC501282.bSep08</a>	<a href="#">501282</a>	7132	1411	6	98	similar to lymphocyte antigen 6 complex, locus E ligand (LOC501282) alternative variant bSep08, mRNA.
<a href="#">LOC501283</a>	<a href="#">LOC501283.bSep08</a>	<a href="#">501283</a>	2224	722	1	198	similar to lipid droplet associated protein (LOC501283) alternative variant bSep08, mRNA.
<a href="#">LOC501306</a>	<a href="#">LOC501306.aSep08</a>	<a href="#">501306</a>	2331	836	2	144	hypothetical gene supported by BC082068 (17.2 kD) (LOC501306) alternative variant aSep08, mRNA.
<a href="#">LOC501306</a>	<a href="#">LOC501306.bSep08</a>	<a href="#">501306</a>	18182	896	5	128	hypothetical gene supported by BC082068 (15.2 kD) (LOC501306) alternative variant bSep08, mRNA.
<a href="#">LOC501307</a>	<a href="#">LOC501307.aSep08</a>	<a href="#">501307</a>	6026	766		114	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A (LOC501307) mRNA.
<a href="#">LOC501307</a>	<a href="#">LOC501307.aSep08</a>	<a href="#">503337</a>	6026	766		114	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A (LOC501307) mRNA.
<a href="#">LOC501327</a>	<a href="#">LOC501327.aSep08</a>	<a href="#">501327</a>	71357	737		42	hypothetical gene supported by BC078986 (4.8 kD) (LOC501327) mRNA.
<a href="#">LOC501329</a>	<a href="#">LOC501329.aSep08</a>	<a href="#">501329</a>	271090	1120	5	226	hypothetical gene supported by BC082068 and hypothetical protein LOC690822 (26.4 kD) (LOC501329) alternative variant aSep08, mRNA.
<a href="#">LOC501329</a>	<a href="#">LOC501329.aSep08</a>	<a href="#">690822</a>	271090	1120	5	226	hypothetical gene supported by BC082068 and hypothetical protein LOC690822 (26.4 kD) (LOC501329) alternative variant aSep08, mRNA.

<a href="#">LOC501329</a>	<a href="#">LOC501329.bSep08</a>	<a href="#">501329</a>	18717	715	2	83	hypothetical gene supported by BC082068 and hypothetical protein LOC690822 (9.4 kD) (LOC501329) alternative variant bSep08, mRNA.
<a href="#">LOC501329</a>	<a href="#">LOC501329.bSep08</a>	<a href="#">690822</a>	18717	715	2	83	hypothetical gene supported by BC082068 and hypothetical protein LOC690822 (9.4 kD) (LOC501329) alternative variant bSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.aSep08</a>	<a href="#">501339</a>	9093	389	4	90	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (10.4 kD) (LOC501339) alternative variant aSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.aSep08</a>	<a href="#">690518</a>	9093	389	4	90	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (10.4 kD) (LOC501339) alternative variant aSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.bSep08</a>	<a href="#">501339</a>	41715	596	4	81	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (LOC501339) alternative variant bSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.bSep08</a>	<a href="#">690518</a>	41715	596	4	81	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (LOC501339) alternative variant bSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.cSep08</a>	<a href="#">501339</a>	29066	423	2	49	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (LOC501339) alternative variant cSep08, mRNA.
<a href="#">LOC501339</a>	<a href="#">LOC501339.cSep08</a>	<a href="#">690518</a>	29066	423	2	49	similar to spermatogenesis associated glutamate (E)-rich protein 4d and similar to zinc finger CCCH type, antiviral 1 (LOC501339) alternative variant cSep08, mRNA.
<a href="#">LOC501344</a>	<a href="#">LOC501344.aSep08</a>	<a href="#">501344</a>	2142	668		117	hypothetical gene supported by BC082068 (LOC501344) mRNA.
<a href="#">LOC501350</a>	<a href="#">LOC501350.aSep08</a>	<a href="#">501350</a>	1215	213		58	hypothetical gene supported by BC082068 (LOC501350) mRNA.
<a href="#">LOC501355</a>	<a href="#">LOC501355.aSep08</a>	<a href="#">501355</a>	68288	573		122	hypothetical gene supported by BC078986 (LOC501355) mRNA.
<a href="#">LOC501358</a>	<a href="#">LOC501358.aSep08</a>	<a href="#">501358</a>	2143	668		117	hypothetical gene supported by BC082068 (LOC501358) mRNA.
<a href="#">LOC501362</a>	<a href="#">LOC501362.aSep08</a>	<a href="#">501362</a>	4753	666		117	hypothetical gene supported by BC082068 and hypothetical protein LOC689545 (LOC501362) mRNA.
<a href="#">LOC501362</a>	<a href="#">LOC501362.aSep08</a>	<a href="#">689545</a>	4753	666		117	hypothetical gene supported by BC082068 and hypothetical protein LOC689545 (LOC501362) mRNA.
<a href="#">LOC501391</a>	<a href="#">LOC501391.aSep08</a>	<a href="#">501391</a>	26739	725	2	95	hypothetical LOC501391 (10.1 kD) (LOC501391) alternative variant aSep08, mRNA.
<a href="#">LOC501391</a>	<a href="#">LOC501391.bSep08</a>	<a href="#">501391</a>	31399	810	3	51	hypothetical LOC501391 (5.4 kD) (LOC501391) alternative variant bSep08, mRNA.
<a href="#">LOC501391</a>	<a href="#">LOC501391.cSep08</a>	<a href="#">501391</a>	28451	325	2	51	hypothetical LOC501391 (5.4 kD) (LOC501391) alternative variant cSep08, mRNA.
<a href="#">LOC501399</a>	<a href="#">LOC501399.aSep08</a>	<a href="#">501399</a>	7919	522		161	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC501399) mRNA.
<a href="#">LOC501463</a>	<a href="#">LOC501463.aSep08</a>	<a href="#">501463</a>	1229	703		108	similar to Iroquois-class homeodomain protein IRX-1 (Iroquois homeobox protein 1) (Homeodomain protein IRXA1) (LOC501463) mRNA.

<a href="#">LOC501477</a>	<a href="#">LOC501477.aSep08</a>	<a href="#">501477</a>	2703	776		154	hypothetical gene supported by BC059164; BC082068 (LOC501477) mRNA.
<a href="#">LOC501488</a>	<a href="#">LOC501488.aSep08</a>	<a href="#">501488</a>	2783	846		177	hypothetical gene supported by BC082068 (LOC501488) mRNA.
<a href="#">LOC501515</a>	<a href="#">LOC501515.aSep08</a>	<a href="#">501515</a>	19200	2038	15	671	similar to Zinc finger CCHC domain-containing protein 6 (LOC501515) alternative variant aSep08, mRNA.
<a href="#">LOC501515</a>	<a href="#">LOC501515.bSep08</a>	<a href="#">501515</a>	5798	514	4	171	similar to Zinc finger CCHC domain-containing protein 6 (LOC501515) alternative variant bSep08, mRNA.
<a href="#">LOC501515</a>	<a href="#">LOC501515.cSep08</a>	<a href="#">501515</a>	13914	726	5	165	similar to Zinc finger CCHC domain-containing protein 6 (LOC501515) alternative variant cSep08, mRNA.
<a href="#">LOC501515</a>	<a href="#">LOC501515.dSep08</a>	<a href="#">501515</a>	14080	1627	3	132	similar to Zinc finger CCHC domain-containing protein 6 (LOC501515) alternative variant dSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.aSep08</a>	<a href="#">501546</a>	14046	3716	15	1067	hypothetical protein LOC501546 (LOC501546) alternative variant aSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.bSep08</a>	<a href="#">501546</a>	13032	1784	8	555	hypothetical protein LOC501546 (LOC501546) alternative variant bSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.cSep08</a>	<a href="#">501546</a>	7690	1370	7	456	hypothetical protein LOC501546 (LOC501546) alternative variant cSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.dSep08</a>	<a href="#">501546</a>	5718	1067	4	329	hypothetical protein LOC501546 (LOC501546) alternative variant dSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.eSep08</a>	<a href="#">501546</a>	5800	943	5	314	hypothetical protein LOC501546 (LOC501546) alternative variant eSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.fSep08</a>	<a href="#">501546</a>	2860	843	5	281	hypothetical protein LOC501546 (LOC501546) alternative variant fSep08, mRNA.
<a href="#">LOC501546</a>	<a href="#">LOC501546.gSep08</a>	<a href="#">501546</a>	1940	709	3	191	hypothetical protein LOC501546 (LOC501546) alternative variant gSep08, mRNA.
<a href="#">LOC501736</a>	<a href="#">LOC501736.aSep08</a>	<a href="#">501736</a>	2808	982		102	similar to CD300A antigen (LOC501736) mRNA.
<a href="#">LOC501810</a>	<a href="#">LOC501810.aSep08</a>	<a href="#">501810</a>	4957	1553		216	similar to Leukosialin precursor (Leucocyte sialoglycoprotein) (Sialophorin) (Ly-48) (B cell differentiation antigen LP-3) (CD43 antigen) (23.0 kD) (LOC501810) mRNA.
<a href="#">LOC502201</a>	<a href="#">LOC502201.aSep08</a>	<a href="#">502201</a>	15151	2306	7	607	similar to CG8272-PA (LOC502201) alternative variant aSep08, mRNA.
<a href="#">LOC502371</a>	<a href="#">LOC502371.aSep08</a>	<a href="#">502371</a>	1151	1042		347	similar to 40S ribosomal protein S21 (LOC502371) mRNA.
<a href="#">LOC502374</a>	<a href="#">LOC502374.aSep08</a>	<a href="#">502374</a>	2016	382		67	hypothetical protein LOC502374 (7.6 kD) (LOC502374) complete mRNA.
<a href="#">LOC502684</a>	<a href="#">LOC502684.aSep08</a>	<a href="#">502684</a>	15601	794	6	264	hypothetical protein LOC502684 (LOC502684) alternative variant aSep08, mRNA.
<a href="#">LOC502684</a>	<a href="#">LOC502684.cSep08</a>	<a href="#">502684</a>	2107	357	2	52	hypothetical protein LOC502684 (LOC502684) alternative variant cSep08, mRNA.
<a href="#">LOC502684</a>	<a href="#">LOC502684.dSep08</a>	<a href="#">502684</a>	6960	1714	5	61	hypothetical protein LOC502684 (LOC502684) alternative variant dSep08, mRNA.
<a href="#">LOC502684</a>	<a href="#">LOC502684.eSep08</a>	<a href="#">502684</a>	29753	475	4	38	hypothetical protein LOC502684 (LOC502684) alternative variant eSep08, mRNA.

<a href="#">LOC502710</a>	<a href="#">LOC502710.aSep08</a>	<a href="#">502710</a>	88070	502	1	144	similar to Myeloid/lymphoid or mixed-lineage leukemia protein 3 homolog (Histone-lysine N-methyltransferase, H3 lysine-4 specific MLL3) (LOC502710) alternative variant aSep08, mRNA.
<a href="#">LOC502710</a>	<a href="#">LOC502710.bSep08</a>	<a href="#">502710</a>	88074	503	1	144	similar to Myeloid/lymphoid or mixed-lineage leukemia protein 3 homolog (Histone-lysine N-methyltransferase, H3 lysine-4 specific MLL3) (LOC502710) alternative variant bSep08, mRNA.
<a href="#">LOC502894</a>	<a href="#">LOC502894.aSep08</a>	<a href="#">502894</a>	8864	844	2	257	hypothetical protein LOC502894 (LOC502894) alternative variant aSep08, mRNA.
<a href="#">LOC503175</a>	<a href="#">LOC503175.aSep08</a>	<a href="#">503175</a>	2904	806		91	similar to Protein KIAA0280 (LOC503175) mRNA.
<a href="#">LOC503391</a>	<a href="#">LOC503391.aSep08</a>	<a href="#">503391</a>	14065	727		71	similar to nidogen 2 (LOC503391) mRNA.
<a href="#">LOC619573</a>	<a href="#">LOC619573.bSep08</a>	<a href="#">619573</a>	6718	398	1	105	hypothetical protein LOC619573 (LOC619573) alternative variant bSep08, mRNA.
<a href="#">LOC619574</a>	<a href="#">LOC619574.bSep08</a>	<a href="#">619574</a>	2638	666	3	103	hypothetical protein LOC619574 (LOC619574) alternative variant bSep08, mRNA.
<a href="#">LOC619574</a>	<a href="#">LOC619574.dSep08</a>	<a href="#">619574</a>	2183	466	3	49	hypothetical protein LOC619574 (5.4 kD) (LOC619574) alternative variant dSep08, mRNA.
<a href="#">LOC654482</a>	<a href="#">LOC654482.bSep08</a>	<a href="#">654482</a>	489	325	2	108	hypothetical protein LOC654482 (LOC654482) alternative variant bSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.aSep08</a>	<a href="#">678701</a>	10515	1553	5	461	hypothetical protein LOC678701 (50.6 kD) (LOC678701) alternative variant aSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.bSep08</a>	<a href="#">678701</a>	4334	986	4	328	hypothetical protein LOC678701 (LOC678701) alternative variant bSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.cSep08</a>	<a href="#">678701</a>	9243	775	4	257	hypothetical protein LOC678701 (LOC678701) alternative variant cSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.dSep08</a>	<a href="#">678701</a>	3393	740	4	206	hypothetical protein LOC678701 (LOC678701) alternative variant dSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.eSep08</a>	<a href="#">678701</a>	5113	567	3	153	hypothetical protein LOC678701 (LOC678701) alternative variant eSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.gSep08</a>	<a href="#">678701</a>	2623	725	3	151	hypothetical protein LOC678701 (LOC678701) alternative variant gSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.hSep08</a>	<a href="#">678701</a>	1315	1196	2	144	hypothetical protein LOC678701 (LOC678701) alternative variant hSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.iSep08</a>	<a href="#">678701</a>	9104	340	2	113	hypothetical protein LOC678701 (LOC678701) alternative variant iSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.kSep08</a>	<a href="#">678701</a>	13945	216	2	71	hypothetical protein LOC678701 (LOC678701) alternative variant kSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.lSep08</a>	<a href="#">678701</a>	14599	298	2	39	hypothetical protein LOC678701 (LOC678701) alternative variant lSep08, mRNA.
<a href="#">LOC678701</a>	<a href="#">LOC678701.mSep08</a>	<a href="#">678701</a>	2302	288	2	33	hypothetical protein LOC678701 (LOC678701) alternative variant mSep08, mRNA.
<a href="#">LOC678704</a>	<a href="#">LOC678704.bSep08</a>	<a href="#">678704</a>	1095	451	1	69	putative protein (LOC678704) alternative variant bSep08, mRNA.
<a href="#">LOC678704</a>	<a href="#">LOC678704.cSep08</a>	<a href="#">678704</a>	27543	1795	3	66	putative protein (LOC678704) alternative variant cSep08, mRNA.



<a href="#">LOC678738</a>	<a href="#">LOC678738.aSep08</a>	<a href="#">678738</a>	1873	926		108	similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) (12.4 kD) (LOC678738) mRNA.
<a href="#">LOC678741</a>	<a href="#">LOC678741.aSep08</a>	<a href="#">678741</a>	20011	875		291	zinc finger ccch domain-containing protein 4 like (LOC678741) mRNA.
<a href="#">LOC678756</a>	<a href="#">LOC678756.aSep08</a>	<a href="#">678756</a>	12763	705		146	similar to GTPase activating protein testicular GAP1 (LOC678756) mRNA.
<a href="#">LOC679532</a>	<a href="#">LOC679532.bSep08</a>	<a href="#">679532</a>	4280	938	8	270	similar to Elongation of very long chain fatty acids protein 1 (LOC679532) alternative variant bSep08, mRNA.
<a href="#">LOC679532</a>	<a href="#">LOC679532.cSep08</a>	<a href="#">679532</a>	2188	1028	8	233	similar to Elongation of very long chain fatty acids protein 1 (LOC679532) alternative variant cSep08, mRNA.
<a href="#">LOC679532</a>	<a href="#">LOC679532.dSep08</a>	<a href="#">679532</a>	2242	811	8	229	similar to Elongation of very long chain fatty acids protein 1 (LOC679532) alternative variant dSep08, mRNA.
<a href="#">LOC679532</a>	<a href="#">LOC679532.eSep08</a>	<a href="#">679532</a>	1210	751	5	125	similar to Elongation of very long chain fatty acids protein 1 (LOC679532) alternative variant eSep08, mRNA.
<a href="#">LOC679532</a>	<a href="#">LOC679532.fSep08</a>	<a href="#">679532</a>	4103	1132	5	111	similar to Elongation of very long chain fatty acids protein 1 (LOC679532) alternative variant fSep08, mRNA.
<a href="#">LOC679534</a>	<a href="#">LOC679534.aSep08</a>	<a href="#">679534</a>	56333	386		128	hypothetical protein LOC679534 (LOC679534) mRNA.
<a href="#">LOC679536</a>	<a href="#">LOC679536.aSep08</a>	<a href="#">679536</a>	3467	371	2	123	hypothetical protein LOC679536 (LOC679536) alternative variant aSep08, mRNA.
<a href="#">LOC679536</a>	<a href="#">LOC679536.bSep08</a>	<a href="#">679536</a>	901	380	2	77	hypothetical protein LOC679536 (LOC679536) alternative variant bSep08, mRNA.
<a href="#">LOC679543</a>	<a href="#">LOC679543.aSep08</a>	<a href="#">679543</a>	71930	472		49	similar to GTPase activating protein testicular GAP1 (5.6 kD) (LOC679543) mRNA.
<a href="#">LOC679551</a>	<a href="#">LOC679551.aSep08</a>	<a href="#">679551</a>	6829	399		91	similar to MIC2 like 1 (LOC679551) mRNA.
<a href="#">LOC679554</a>	<a href="#">LOC679554.aSep08</a>	<a href="#">679554</a>	57710	923		103	hypothetical protein LOC679554 (11.5 kD) (LOC679554) mRNA.
<a href="#">LOC679572</a>	<a href="#">LOC679572.aSep08</a>	<a href="#">679572</a>	1451	433	2	79	similar to CG6878-PA (8.2 kD) (LOC679572) alternative variant aSep08, mRNA.
<a href="#">LOC679572</a>	<a href="#">LOC679572.bSep08</a>	<a href="#">679572</a>	1594	397	3	79	similar to CG6878-PA (8.2 kD) (LOC679572) alternative variant bSep08, mRNA.
<a href="#">LOC679585</a>	<a href="#">LOC679585.cSep08</a>	<a href="#">679585</a>	13782	415	5	126	similar to acid phosphatase 1 isoform b (LOC679585) alternative variant cSep08, mRNA.
<a href="#">LOC679585</a>	<a href="#">LOC679585.dSep08</a>	<a href="#">679585</a>	1830	735	2	36	similar to acid phosphatase 1 isoform b (LOC679585) alternative variant dSep08, mRNA.
<a href="#">LOC679587</a>	<a href="#">LOC679587.aSep08</a>	<a href="#">679587</a>	64934	618	3	163	hypothetical protein LOC679587 and hypothetical protein LOC679608 (LOC679587) alternative variant aSep08, mRNA.
<a href="#">LOC679587</a>	<a href="#">LOC679587.aSep08</a>	<a href="#">679608</a>	64934	618	3	163	hypothetical protein LOC679587 and hypothetical protein LOC679608 (LOC679587) alternative variant aSep08, mRNA.
<a href="#">LOC679587</a>	<a href="#">LOC679587.bSep08</a>	<a href="#">679587</a>	29399	747	3	118	hypothetical protein LOC679587 and hypothetical protein LOC679608 (12.8 kD) (LOC679587) alternative variant bSep08, mRNA.
<a href="#">LOC679587</a>	<a href="#">LOC679587.bSep08</a>	<a href="#">679608</a>	29399	747	3	118	hypothetical protein LOC679587 and hypothetical protein LOC679608 (12.8 kD) (LOC679587) alternative variant bSep08, mRNA.

<a href="#">LOC679587</a>	<a href="#">LOC679587.cSep08</a>	<a href="#">679587</a>	6597	444	2	61	hypothetical protein LOC679587 and hypothetical protein LOC679608 (7.0 kD) (LOC679587) alternative variant cSep08, mRNA.
<a href="#">LOC679587</a>	<a href="#">LOC679587.cSep08</a>	<a href="#">679608</a>	6597	444	2	61	hypothetical protein LOC679587 and hypothetical protein LOC679608 (7.0 kD) (LOC679587) alternative variant cSep08, mRNA.
<a href="#">LOC679610</a>	<a href="#">LOC679610.aSep08</a>	<a href="#">679610</a>	34371	761	5	202	CRA c (LOC679610) alternative variant aSep08, mRNA.
<a href="#">LOC679610</a>	<a href="#">LOC679610.bSep08</a>	<a href="#">679610</a>	38353	882	4	140	CRA c (15.7 kD) (LOC679610) alternative variant bSep08, mRNA.
<a href="#">LOC679610</a>	<a href="#">LOC679610.cSep08</a>	<a href="#">679610</a>	13624	661	1	100	CRA c like (11.3 kD) (LOC679610) alternative variant cSep08, mRNA.
<a href="#">LOC679620</a>	<a href="#">LOC679620.aSep08</a>	<a href="#">679620</a>	18591	648		213	similar to CG4329-PA, isoform A (LOC679620) mRNA.
<a href="#">LOC679623</a>	<a href="#">LOC679623.aSep08</a>	<a href="#">679623</a>	18809	1055	6	166	similar to Nuclear autoantigen Sp-100 (Speckled 100 kDa) (Nuclear dot-associated Sp100 protein) (18.8 kD) (LOC679623) alternative variant aSep08, mRNA.
<a href="#">LOC679623</a>	<a href="#">LOC679623.bSep08</a>	<a href="#">679623</a>	1434	870	2	56	similar to Nuclear autoantigen Sp-100 (Speckled 100 kDa) (Nuclear dot-associated Sp100 protein) (6.6 kD) (LOC679623) alternative variant bSep08, mRNA.
<a href="#">LOC679651</a>	<a href="#">LOC679651.aSep08</a>	<a href="#">679651</a>	38128	357		115	hypothetical protein LOC679651 (LOC679651) mRNA.
<a href="#">LOC679693</a>	<a href="#">LOC679693.aSep08</a>	<a href="#">679693</a>	3738	1654	7	405	similar to Mediator of RNA polymerase II transcription subunit 12 (Thyroid hormone receptor-associated protein complex 230 kDa component) (Trap230) (Activator-recruited cofactor 240 kDa component) (ARC240) (CAG repeat protein 45) (OPA-containing pr (43.7 kD) (LOC679693) alternative variant aSep08, mRNA.
<a href="#">LOC679693</a>	<a href="#">LOC679693.bSep08</a>	<a href="#">679693</a>	555	382	2	127	similar to Mediator of RNA polymerase II transcription subunit 12 (Thyroid hormone receptor-associated protein complex 230 kDa component) (Trap230) (Activator-recruited cofactor 240 kDa component) (ARC240) (CAG repeat protein 45) (OPA-containing pr (LOC679693) alternative variant bSep08, mRNA.
<a href="#">LOC679693</a>	<a href="#">LOC679693.cSep08</a>	<a href="#">679693</a>	1443	273	3	91	similar to Mediator of RNA polymerase II transcription subunit 12 (Thyroid hormone receptor-associated protein complex 230 kDa component) (Trap230) (Activator-recruited cofactor 240 kDa component) (ARC240) (CAG repeat protein 45) (OPA-containing pr (LOC679693) alternative variant cSep08, mRNA.
<a href="#">LOC679714</a>	<a href="#">LOC679714.aSep08</a>	<a href="#">679714</a>	11572	1253		72	similar to CG7220-PA, isoform A (LOC679714) mRNA.
<a href="#">LOC679725</a>	<a href="#">LOC679725.aSep08</a>	<a href="#">679725</a>	14110	3515	9	1023	CRA a (LOC679725) alternative variant aSep08, mRNA.
<a href="#">LOC679725</a>	<a href="#">LOC679725.bSep08</a>	<a href="#">679725</a>	17642	2934	8	869	CRA a (LOC679725) alternative variant bSep08, mRNA.
<a href="#">LOC679725</a>	<a href="#">LOC679725.cSep08</a>	<a href="#">679725</a>	10648	949	2	210	CRA a (22.1 kD) (LOC679725) alternative variant cSep08, mRNA.
<a href="#">LOC679725</a>	<a href="#">LOC679725.dSep08</a>	<a href="#">679725</a>	1747	634	3	111	eukaryotic translation initiation factor 4E binding protein like (LOC679725) alternative variant dSep08, mRNA.
<a href="#">LOC679725</a>	<a href="#">LOC679725.eSep08</a>	<a href="#">679725</a>	758	642	2	83	CRA c like (9.1 kD) (LOC679725) alternative variant eSep08, mRNA.

<a href="#">LOC679739</a>	<a href="#">LOC679739.aSep08</a>	<a href="#">679739</a>	8555	487	4	120	NADH dehydrogenase (ubiquinone) Fe-S protein 6 (LOC679739) alternative variant aSep08, mRNA.
<a href="#">LOC679739</a>	<a href="#">LOC679739.bSep08</a>	<a href="#">679739</a>	8554	547	4	85	NADH dehydrogenase (ubiquinone) Fe-S protein 6 (9.8 kD) (LOC679739) alternative variant bSep08, mRNA.
<a href="#">LOC679751</a>	<a href="#">LOC679751.aSep08</a>	<a href="#">679751</a>	2152	254		60	hypothetical protein LOC679751 (LOC679751) mRNA.
<a href="#">LOC679769</a>	<a href="#">LOC679769.aSep08</a>	<a href="#">679769</a>	36199	670		41	similar to nidogen 2 (4.5 kD) (LOC679769) mRNA.
<a href="#">LOC679801</a>	<a href="#">LOC679801.aSep08</a>	<a href="#">679801</a>	1211	388		129	similar to Pre-B-cell leukemia transcription factor 3 (Homeobox protein PBX3) (LOC679801) mRNA.
<a href="#">LOC679811</a>	<a href="#">LOC679811.aSep08</a>	<a href="#">679811</a>	30937	3469	16	934	similar to RIKEN cDNA D930015E06 (LOC679811) alternative variant aSep08, mRNA.
<a href="#">LOC679811</a>	<a href="#">LOC679811.bSep08</a>	<a href="#">679811</a>	10431	767	4	223	similar to RIKEN cDNA D930015E06 (LOC679811) alternative variant bSep08, mRNA.
<a href="#">LOC679811</a>	<a href="#">LOC679811.cSep08</a>	<a href="#">679811</a>	12620	444	5	147	similar to RIKEN cDNA D930015E06 (LOC679811) alternative variant cSep08, mRNA.
<a href="#">LOC679811</a>	<a href="#">LOC679811.dSep08</a>	<a href="#">679811</a>	1269	729	2	85	similar to RIKEN cDNA D930015E06 (LOC679811) alternative variant dSep08, mRNA.
<a href="#">LOC679818</a>	<a href="#">LOC679818.aSep08</a>	<a href="#">679818</a>	10175	506		168	similar to Maltase-glucoamylase, intestinal (LOC679818) mRNA.
<a href="#">LOC679824</a>	<a href="#">LOC679824.aSep08</a>	<a href="#">679824</a>	5600	508	1	120	hypothetical protein LOC679824 (13.1 kD) (LOC679824) alternative variant aSep08, mRNA.
<a href="#">LOC679824</a>	<a href="#">LOC679824.bSep08</a>	<a href="#">679824</a>	3831	380	1	80	hypothetical protein LOC679824 (LOC679824) alternative variant bSep08, mRNA.
<a href="#">LOC679824</a>	<a href="#">LOC679824.cSep08</a>	<a href="#">679824</a>	14430	515	1	59	hypothetical protein LOC679824 (6.3 kD) (LOC679824) alternative variant cSep08, mRNA.
<a href="#">LOC679834</a>	<a href="#">LOC679834.aSep08</a>	<a href="#">679834</a>	12439	550	1	88	similar to Meiosis expressed protein 1 (10.8 kD) (LOC679834) alternative variant aSep08, mRNA.
<a href="#">LOC679834</a>	<a href="#">LOC679834.cSep08</a>	<a href="#">679834</a>	11923	508	1	88	similar to Meiosis expressed protein 1 (10.8 kD) (LOC679834) alternative variant cSep08, mRNA.
<a href="#">LOC679858</a>	<a href="#">LOC679858.aSep08</a>	<a href="#">679858</a>	7153	493		163	hypothetical protein LOC679858 (LOC679858) mRNA.
<a href="#">LOC679860</a>	<a href="#">LOC679860.aSep08</a>	<a href="#">679860</a>	4701	360		60	hypothetical protein LOC679860 (6.9 kD) (LOC679860) mRNA.
<a href="#">LOC679861</a>	<a href="#">LOC679861.aSep08</a>	<a href="#">679861</a>	6772	498	3	92	similar to MIC2 like 1 (LOC679861) alternative variant aSep08, mRNA.
<a href="#">LOC679863</a>	<a href="#">LOC679863.aSep08</a>	<a href="#">679863</a>	1080	329		82	similar to N-myc downstream regulated gene 3 (LOC679863) mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.aSep08</a>	<a href="#">679869</a>	178055	1788	3	595	CRA f like (LOC679869) alternative variant aSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.bSep08</a>	<a href="#">679869</a>	13979	724	4	241	transcription factor 7-like 2 CRA d (LOC679869) alternative variant bSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.cSep08</a>	<a href="#">679869</a>	177261	724	7	240	transcription factor 7-like 2 (LOC679869) alternative variant cSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.dSep08</a>	<a href="#">679869</a>	171779	513	6	170	T-cell Transcription C-terminal like (LOC679869) alternative variant dSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.eSep08</a>	<a href="#">679869</a>	6081	815	4	157	transcription factor 7-like 2 CRA a (LOC679869) alternative variant eSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.fSep08</a>	<a href="#">679869</a>	15873	2545	4	151	transcription factor 7-like 2 CRA h (LOC679869) alternative variant fSep08, mRNA.

<a href="#">LOC679869</a>	<a href="#">LOC679869.hSep08</a>	<a href="#">679869</a>	5138	422	3	40	transcription factor 7-like 2 CRA a (4.5 kD) (LOC679869) alternative variant hSep08, mRNA.
<a href="#">LOC679869</a>	<a href="#">LOC679869.jSep08</a>	<a href="#">679869</a>	5026	329	3	6	putative protein (0.7 kD) (LOC679869) alternative variant jSep08, mRNA.
<a href="#">LOC679873</a>	<a href="#">LOC679873.aSep08</a>	<a href="#">679873</a>	1139	731		218	similar to retinoic acid early transcript 1L (LOC679873) mRNA.
<a href="#">LOC679885</a>	<a href="#">LOC679885.aSep08</a>	<a href="#">679885</a>	21185	689	3	38	similar to N-myc downstream regulated gene 3 (4.2 kD) (LOC679885) alternative variant aSep08, mRNA.
<a href="#">LOC679896</a>	<a href="#">LOC679896.aSep08</a>	<a href="#">679896</a>	8507	930		234	hypothetical protein LOC679896 (24.9 kD) (LOC679896) mRNA.
<a href="#">LOC679898</a>	<a href="#">LOC679898.aSep08</a>	<a href="#">679898</a>	4633	1806	6	198	similar to zinc finger, matrin type 2 (LOC679898) alternative variant aSep08, mRNA.
<a href="#">LOC679902</a>	<a href="#">LOC679902.aSep08</a>	<a href="#">679902</a>	13266	770		73	hypothetical protein LOC679902 (LOC679902) mRNA.
<a href="#">LOC679924</a>	<a href="#">LOC679924.aSep08</a>	<a href="#">679924</a>	2596	680		210	similar to Kif19A CG9913-PB, isoform B (LOC679924) mRNA.
<a href="#">LOC679934</a>	<a href="#">LOC679934.aSep08</a>	<a href="#">679934</a>	15237	1106		75	probable protein (8.8 kD) (LOC679934) complete mRNA.
<a href="#">LOC679937</a>	<a href="#">LOC679937.aSep08</a>	<a href="#">679937</a>	37061	2743		204	similar to CG4025-PA (LOC679937) alternative variant aSep08, mRNA.
<a href="#">LOC679937</a>	<a href="#">LOC679937.bSep08</a>	<a href="#">679937</a>	2604	2052		45	similar to CG4025-PA (LOC679937) alternative variant bSep08, mRNA.
<a href="#">LOC679949</a>	<a href="#">LOC679949.aSep08</a>	<a href="#">679949</a>	38782	1022		145	similar to CG4502-PA, isoform A (LOC679949) mRNA.
<a href="#">LOC679989</a>	<a href="#">LOC679989.aSep08</a>	<a href="#">679989</a>	4619	623		141	similar to t-complex 11 protein (LOC679989) mRNA.
<a href="#">LOC680025</a>	<a href="#">LOC680025.aSep08</a>	<a href="#">680025</a>	1035	686	2	72	similar to nuclear RNA export factor 2 (LOC680025) alternative variant aSep08, mRNA.
<a href="#">LOC680025</a>	<a href="#">LOC680025.bSep08</a>	<a href="#">680025</a>	12270	684	3	62	similar to nuclear RNA export factor 2 (LOC680025) alternative variant bSep08, mRNA.
<a href="#">LOC680025</a>	<a href="#">LOC680025.cSep08</a>	<a href="#">680025</a>	3239	1494	4	291	similar to nuclear RNA export factor 2 (LOC680025) alternative variant cSep08, mRNA.
<a href="#">LOC680025</a>	<a href="#">LOC680025.eSep08</a>	<a href="#">680025</a>	24656	791	7	57	similar to nuclear RNA export factor 2 (LOC680025) alternative variant eSep08, mRNA.
<a href="#">LOC680036</a>	<a href="#">LOC680036.aSep08</a>	<a href="#">680036</a>	50038	418		138	similar to methionine sulfoxide reductase B3 isoform 2 (LOC680036) mRNA.
<a href="#">LOC680039</a>	<a href="#">LOC680039.bSep08</a>	<a href="#">680039</a>	15555	2610	2	474	hypothetical protein LOC680039 (LOC680039) alternative variant bSep08, mRNA.
<a href="#">LOC680039</a>	<a href="#">LOC680039.cSep08</a>	<a href="#">680039</a>	1953	237	2	36	hypothetical protein LOC680039 (LOC680039) alternative variant cSep08, mRNA.
<a href="#">LOC680045</a>	<a href="#">LOC680045.bSep08</a>	<a href="#">680045</a>	7398	410	1	136	hypothetical protein LOC680045 (LOC680045) alternative variant bSep08, mRNA.
<a href="#">LOC680045</a>	<a href="#">LOC680045.cSep08</a>	<a href="#">680045</a>	7569	682	1	121	hypothetical protein LOC680045 (LOC680045) alternative variant cSep08, mRNA.
<a href="#">LOC680069</a>	<a href="#">LOC680069.aSep08</a>	<a href="#">680069</a>	58276	317		105	similar to Pappalysin-2 precursor (Pregnancy-associated plasma protein-A2) (PAPP-A2) (Pregnancy-associated plasma protein-E1) (PAPP-E) (LOC680069) mRNA.
<a href="#">LOC680074</a>	<a href="#">LOC680074.aSep08</a>	<a href="#">680074</a>	12390	264		40	hypothetical protein LOC680074 (LOC680074) mRNA.
<a href="#">LOC680079</a>	<a href="#">LOC680079.aSep08</a>	<a href="#">680079</a>	3592	829	4	275	similar to Chloride channel protein 6 (CIC-6) (LOC680079) alternative variant aSep08, mRNA.

<a href="#">LOC680079</a>	<a href="#">LOC680079.bSep08</a>	<a href="#">680079</a>	2674	973	1	149	similar to Chloride channel protein 6 (ClC-6) (LOC680079) alternative variant bSep08, mRNA.
<a href="#">LOC680080</a>	<a href="#">LOC680080.aSep08</a>	<a href="#">680080</a>	2836	636	2	51	hypothetical protein LOC680080 (6.0 kD) (LOC680080) alternative variant aSep08, complete mRNA.
<a href="#">LOC680096</a>	<a href="#">LOC680096.aSep08</a>	<a href="#">680096</a>	123729	501		76	similar to ribosomal protein L31 (LOC680096) mRNA.
<a href="#">LOC680117</a>	<a href="#">LOC680117.aSep08</a>	<a href="#">680117</a>	1932	801	2	161	similar to T-cell leukemia homeobox protein 2 (Homeobox protein Hox-11L1) (Homeobox TLX-2) (PMUR10F) (LOC680117) alternative variant aSep08, mRNA.
<a href="#">LOC680117</a>	<a href="#">LOC680117.bSep08</a>	<a href="#">680117</a>	1698	463	2	126	similar to T-cell leukemia homeobox protein 2 (Homeobox protein Hox-11L1) (Homeobox TLX-2) (PMUR10F) (LOC680117) alternative variant bSep08, mRNA.
<a href="#">LOC680117</a>	<a href="#">LOC680117.cSep08</a>	<a href="#">680117</a>	1704	474	2	123	similar to T-cell leukemia homeobox protein 2 (Homeobox protein Hox-11L1) (Homeobox TLX-2) (PMUR10F) (LOC680117) alternative variant cSep08, mRNA.
<a href="#">LOC680117</a>	<a href="#">LOC680117.dSep08</a>	<a href="#">680117</a>	671	288	1	90	similar to T-cell leukemia homeobox protein 2 (Homeobox protein Hox-11L1) (Homeobox TLX-2) (PMUR10F) (LOC680117) alternative variant dSep08, mRNA.
<a href="#">LOC680117</a>	<a href="#">LOC680117.eSep08</a>	<a href="#">680117</a>	1718	592	2	90	similar to T-cell leukemia homeobox protein 2 (Homeobox protein Hox-11L1) (Homeobox TLX-2) (PMUR10F) (LOC680117) alternative variant eSep08, mRNA.
<a href="#">LOC680133</a>	<a href="#">LOC680133.aSep08</a>	<a href="#">680133</a>	19004	377		125	similar to TBC1 domain family, member 8 (LOC680133) mRNA.
<a href="#">LOC680149</a>	<a href="#">LOC680149.aSep08</a>	<a href="#">680149</a>	2914	443	1	74	similar to Guanine nucleotide-binding protein G(T) gamma-T1 subunit precursor (Transducin gamma chain) (8.6 kD) (LOC680149) alternative variant aSep08, mRNA.
<a href="#">LOC680149</a>	<a href="#">LOC680149.bSep08</a>	<a href="#">680149</a>	2985	498	1	74	similar to Guanine nucleotide-binding protein G(T) gamma-T1 subunit precursor (Transducin gamma chain) (8.6 kD) (LOC680149) alternative variant bSep08, mRNA.
<a href="#">LOC680149</a>	<a href="#">LOC680149.cSep08</a>	<a href="#">680149</a>	2871	527	1	56	similar to Guanine nucleotide-binding protein G(T) gamma-T1 subunit precursor (Transducin gamma chain) (LOC680149) alternative variant cSep08, mRNA.
<a href="#">LOC680155</a>	<a href="#">LOC680155.aSep08</a>	<a href="#">680155</a>	16317	825		260	hypothetical protein LOC680155 (LOC680155) mRNA.
<a href="#">LOC680157</a>	<a href="#">LOC680157.aSep08</a>	<a href="#">680157</a>	4382	725	2	87	hypothetical protein LOC680157 (9.8 kD) (LOC680157) alternative variant aSep08, mRNA.
<a href="#">LOC680157</a>	<a href="#">LOC680157.bSep08</a>	<a href="#">680157</a>	2845	269	1	64	hypothetical protein LOC680157 (LOC680157) alternative variant bSep08, mRNA.
<a href="#">LOC680162</a>	<a href="#">LOC680162.aSep08</a>	<a href="#">680162</a>	3047	1958		429	similar to Nucleosome binding protein 1 (Nucleosome binding protein 45) (NBP-45) (GARP45 protein) and similar to Nucleosome binding protein 1 (Nucleosome binding protein 45) (NBP-45) (GARP45 protein) (48.6 kD) (LOC680162) mRNA.
<a href="#">LOC680162</a>	<a href="#">LOC680162.aSep08</a>	<a href="#">680182</a>	3047	1958		429	similar to Nucleosome binding protein 1 (Nucleosome binding protein 45) (NBP-45) (GARP45 protein) and similar to Nucleosome binding protein 1 (Nucleosome binding protein 45) (NBP-45) (GARP45 protein) (48.6 kD) (LOC680162) mRNA.
<a href="#">LOC680164</a>	<a href="#">LOC680164.aSep08</a>	<a href="#">680164</a>	5745	1344		413	hypothetical protein LOC680164 (LOC680164) mRNA.

<a href="#">LOC680166</a>	<a href="#">LOC680166.aSep08</a>	<a href="#">680166</a>	1351	522		69	hypothetical protein LOC680166 (LOC680166) mRNA.
<a href="#">LOC680168</a>	<a href="#">LOC680168.aSep08</a>	<a href="#">680168</a>	5338	1139	2	379	similar to retinoblastoma-associated protein 140 (LOC680168) alternative variant aSep08, mRNA.
<a href="#">LOC680168</a>	<a href="#">LOC680168.bSep08</a>	<a href="#">680168</a>	1723	572	2	94	similar to retinoblastoma-associated protein 140 (LOC680168) alternative variant bSep08, mRNA.
<a href="#">LOC680185</a>	<a href="#">LOC680185.aSep08</a>	<a href="#">680185</a>	1275	559		38	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC680185) mRNA.
<a href="#">LOC680199</a>	<a href="#">LOC680199.aSep08</a>	<a href="#">680199</a>	1108	696		133	hypothetical protein LOC680199 (14.9 kD) (LOC680199) mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.aSep08</a>	<a href="#">680200</a>	17216	1513	4	470	zinc finger protein 455 (55.4 kD) (LOC680200) alternative variant aSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.aSep08</a>	<a href="#">680222</a>	17216	1513	4	470	zinc finger protein 455 (55.4 kD) (LOC680200) alternative variant aSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.bSep08</a>	<a href="#">680200</a>	16801	1578	4	470	zinc finger protein 455 (55.4 kD) (LOC680200) alternative variant bSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.bSep08</a>	<a href="#">680222</a>	16801	1578	4	470	zinc finger protein 455 (55.4 kD) (LOC680200) alternative variant bSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.cSep08</a>	<a href="#">680200</a>	11347	1057	4	171	finger protein zinc 45 (LOC680200) alternative variant cSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.cSep08</a>	<a href="#">680222</a>	11347	1057	4	171	finger protein zinc 45 (LOC680200) alternative variant cSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.dSep08</a>	<a href="#">680200</a>	9062	567	4	58	KRAB box (7.1 kD) (LOC680200) alternative variant dSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.dSep08</a>	<a href="#">680222</a>	9062	567	4	58	KRAB box (7.1 kD) (LOC680200) alternative variant dSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.eSep08</a>	<a href="#">680200</a>	55933	346	4	54	KRAB box (6.3 kD) (LOC680200) alternative variant eSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.eSep08</a>	<a href="#">680222</a>	55933	346	4	54	KRAB box (6.3 kD) (LOC680200) alternative variant eSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.fSep08</a>	<a href="#">680200</a>	10282	1676	3	53	KRAB box (6.4 kD) (LOC680200) alternative variant fSep08, mRNA.
<a href="#">LOC680200</a>	<a href="#">LOC680200.fSep08</a>	<a href="#">680222</a>	10282	1676	3	53	KRAB box (6.4 kD) (LOC680200) alternative variant fSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.aSep08</a>	<a href="#">680230</a>	38179	658	5	98	hypothetical protein LOC680230 (LOC680230) alternative variant aSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.bSep08</a>	<a href="#">680230</a>	17617	517	4	89	hypothetical protein LOC680230 (LOC680230) alternative variant bSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.cSep08</a>	<a href="#">680230</a>	17603	593	4	87	hypothetical protein LOC680230 (LOC680230) alternative variant cSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.dSep08</a>	<a href="#">680230</a>	13880	458	3	79	hypothetical protein LOC680230 (LOC680230) alternative variant dSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.eSep08</a>	<a href="#">680230</a>	38044	331	4	78	hypothetical protein LOC680230 (LOC680230) alternative variant eSep08, mRNA.
<a href="#">LOC680230</a>	<a href="#">LOC680230.fSep08</a>	<a href="#">680230</a>	38263	529	3	54	hypothetical protein LOC680230 (6.2 kD) (LOC680230) alternative variant fSep08, mRNA.
<a href="#">LOC680231</a>	<a href="#">LOC680231.aSep08</a>	<a href="#">680231</a>	17274	1263	9	328	similar to chromodomain helicase DNA binding protein 9 (LOC680231) alternative variant aSep08, mRNA.

<a href="#">LOC680231</a>	<a href="#">LOC680231.bSep08</a>	<a href="#">680231</a>	19607	511	3	161	similar to chromodomain helicase DNA binding protein 9 (LOC680231) alternative variant bSep08, mRNA.
<a href="#">LOC680235</a>	<a href="#">LOC680235.aSep08</a>	<a href="#">680235</a>	10435	606			
<a href="#">LOC680241</a>	<a href="#">LOC680241.aSep08</a>	<a href="#">680241</a>	4066	1703		536	similar to zinc finger protein 87 (LOC680241) mRNA.
<a href="#">LOC680252</a>	<a href="#">LOC680252.aSep08</a>	<a href="#">680252</a>	6251	2037	5	533	similar to Dystrophia myotonica WD repeat-containing protein (Dystrophia myotonica-containing WD repeat motif protein) (DMR-N9 protein) (LOC680252) alternative variant aSep08, mRNA.
<a href="#">LOC680252</a>	<a href="#">LOC680252.cSep08</a>	<a href="#">680252</a>	1927	820	2	135	similar to Dystrophia myotonica WD repeat-containing protein (Dystrophia myotonica-containing WD repeat motif protein) (DMR-N9 protein) (14.5 kD) (LOC680252) alternative variant cSep08, mRNA.
<a href="#">LOC680259</a>	<a href="#">LOC680259.aSep08</a>	<a href="#">680259</a>	82787	557		185	hypothetical protein LOC680259 (LOC680259) mRNA.
<a href="#">LOC680270</a>	<a href="#">LOC680270.aSep08</a>	<a href="#">680270</a>	4956	544		159	hypothetical protein LOC680270 (LOC680270) mRNA.
<a href="#">LOC680282</a>	<a href="#">LOC680282.aSep08</a>	<a href="#">680282</a>	2247	493	1	76	hypothetical protein LOC680282 (LOC680282) alternative variant aSep08, mRNA.
<a href="#">LOC680282</a>	<a href="#">LOC680282.bSep08</a>	<a href="#">680282</a>	2260	418		32	hypothetical protein LOC680282 (LOC680282) alternative variant bSep08, mRNA.
<a href="#">LOC680290</a>	<a href="#">LOC680290.aSep08</a>	<a href="#">680290</a>	34286	729	5	103	similar to Zinc finger protein 208 (LOC680290) alternative variant aSep08, mRNA.
<a href="#">LOC680290</a>	<a href="#">LOC680290.bSep08</a>	<a href="#">680290</a>	29311	432	5	99	similar to Zinc finger protein 208 (LOC680290) alternative variant bSep08, mRNA.
<a href="#">LOC680290</a>	<a href="#">LOC680290.dSep08</a>	<a href="#">680290</a>	34259	798	6	62	similar to Zinc finger protein 208 (LOC680290) alternative variant dSep08, mRNA.
<a href="#">LOC680308</a>	<a href="#">LOC680308.bSep08</a>	<a href="#">680308</a>	5743	509	1	169	similar to Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase, mitochondrial precursor (LOC680308) alternative variant bSep08, mRNA.
<a href="#">LOC680344</a>	<a href="#">LOC680344.aSep08</a>	<a href="#">680344</a>	147621	627	4	208	similar to Protein KIAA0574 (LOC680344) alternative variant aSep08, mRNA.
<a href="#">LOC680344</a>	<a href="#">LOC680344.bSep08</a>	<a href="#">680344</a>	28941	655	1	105	similar to Protein KIAA0574 (LOC680344) alternative variant bSep08, mRNA.
<a href="#">LOC680367</a>	<a href="#">LOC680367.aSep08</a>	<a href="#">680367</a>	3605	554		101	similar to Urinary protein 3 precursor (RUP-3) (11.1 kD) (LOC680367) mRNA.
<a href="#">LOC680392</a>	<a href="#">LOC680392.aSep08</a>	<a href="#">680392</a>	19232	1044	2	243	similar to Reticulocalbin-1 precursor (LOC680392) alternative variant aSep08, mRNA.
<a href="#">LOC680392</a>	<a href="#">LOC680392.bSep08</a>	<a href="#">680392</a>	6891	756	2	208	similar to Reticulocalbin-1 precursor (LOC680392) alternative variant bSep08, mRNA.
<a href="#">LOC680392</a>	<a href="#">LOC680392.cSep08</a>	<a href="#">680392</a>	23462	830	2	156	similar to Reticulocalbin-1 precursor (18.6 kD) (LOC680392) alternative variant cSep08, mRNA.
<a href="#">LOC680406</a>	<a href="#">LOC680406.aSep08</a>	<a href="#">680406</a>	579	118		39	similar to Urinary protein 2 precursor (RUP-2) (LOC680406) mRNA.
<a href="#">LOC680409</a>	<a href="#">LOC680409.aSep08</a>	<a href="#">680409</a>	17010	2022	12	554	similar to Proline oxidase, mitochondrial precursor (Proline dehydrogenase) (LOC680409) alternative variant aSep08, mRNA.
<a href="#">LOC680409</a>	<a href="#">LOC680409.bSep08</a>	<a href="#">680409</a>	1414	550	1	107	similar to Proline oxidase, mitochondrial precursor (Proline dehydrogenase) (12.1 kD) (LOC680409) alternative variant bSep08, mRNA.

<a href="#">LOC680415</a>	<a href="#">LOC680415.aSep08</a>	<a href="#">680415</a>	22012	643		214	similar to pappalysin 2 isoform 1 (LOC680415) mRNA.
<a href="#">LOC680433</a>	<a href="#">LOC680433.aSep08</a>	<a href="#">680433</a>	2335	743		161	hypothetical protein LOC680433 (LOC680433) mRNA.
<a href="#">LOC680445</a>	<a href="#">LOC680445.bSep08</a>	<a href="#">680445</a>	24076	1492	2	98	muscleblind-like 2 CRA d (11.0 kD) (LOC680445) alternative variant bSep08, mRNA.
<a href="#">LOC680445</a>	<a href="#">LOC680445.cSep08</a>	<a href="#">680445</a>	38685	1136	3	94	muscleblind-like 2 (10.4 kD) (LOC680445) alternative variant cSep08, mRNA.
<a href="#">LOC680445</a>	<a href="#">LOC680445.dSep08</a>	<a href="#">680445</a>	38445	991	4	88	muscleblind-like 2 (9.2 kD) (LOC680445) alternative variant dSep08, complete mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.aSep08</a>	<a href="#">680451</a>	4697	1700	17	489	similar to nuclear receptor binding protein (LOC680451) alternative variant aSep08, mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.bSep08</a>	<a href="#">680451</a>	4434	1593	16	295	similar to nuclear receptor binding protein (33.5 kD) (LOC680451) alternative variant bSep08, mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.cSep08</a>	<a href="#">680451</a>	5365	3357	11	258	similar to nuclear receptor binding protein (29.6 kD) (LOC680451) alternative variant cSep08, mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.dSep08</a>	<a href="#">680451</a>	2734	1247	5	100	similar to nuclear receptor binding protein (11.1 kD) (LOC680451) alternative variant dSep08, mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.eSep08</a>	<a href="#">680451</a>	1667	918	6	88	similar to nuclear receptor binding protein (9.0 kD) (LOC680451) alternative variant eSep08, mRNA.
<a href="#">LOC680451</a>	<a href="#">LOC680451.fSep08</a>	<a href="#">680451</a>	758	419	2	37	similar to nuclear receptor binding protein (LOC680451) alternative variant fSep08, mRNA.
<a href="#">LOC680453</a>	<a href="#">LOC680453.aSep08</a>	<a href="#">680453</a>	1958	781		102	hypothetical protein LOC680453 (LOC680453) mRNA.
<a href="#">LOC680485</a>	<a href="#">LOC680485.bSep08</a>	<a href="#">680485</a>	3622	489	3	45	hypothetical protein LOC680485 (LOC680485) alternative variant bSep08, mRNA.
<a href="#">LOC680494</a>	<a href="#">LOC680494.aSep08</a>	<a href="#">680494</a>	16690	1136		67	hypothetical protein LOC680494 (7.4 kD) (LOC680494) mRNA.
<a href="#">LOC680513</a>	<a href="#">LOC680513.aSep08</a>	<a href="#">680513</a>	1122	402		79	similar to Alpha-N-acetylgalactosaminide alpha-2,6-sialyltransferase 5 (GalNAc alpha-2,6-sialyltransferase V) (ST6GalNAc V) (GD1 alpha synthase) (Sialyltransferase 7E) (LOC680513) mRNA.
<a href="#">LOC680545</a>	<a href="#">LOC680545.aSep08</a>	<a href="#">680545</a>	4857	4592		1373	hypothetical protein LOC680545 and hypothetical protein LOC680559 (LOC680545) mRNA.
<a href="#">LOC680545</a>	<a href="#">LOC680545.aSep08</a>	<a href="#">680559</a>	4857	4592		1373	hypothetical protein LOC680545 and hypothetical protein LOC680559 (LOC680545) mRNA.
<a href="#">LOC680549</a>	<a href="#">LOC680549.aSep08</a>	<a href="#">680549</a>	24352	386	3	128	similar to Pbx/knotted 1 homeobox 2 (LOC680549) alternative variant aSep08, mRNA.
<a href="#">LOC680549</a>	<a href="#">LOC680549.bSep08</a>	<a href="#">680549</a>	159036	529	5	77	similar to Pbx/knotted 1 homeobox 2 (8.1 kD) (LOC680549) alternative variant bSep08, mRNA.
<a href="#">LOC680551</a>	<a href="#">LOC680551.bSep08</a>	<a href="#">680551</a>	4202	1025	1	70	similar to Apolipoprotein C-IV precursor (Apo-CIV) (ApoC-IV) (Apolipoprotein C2-linked) (ACL) (8.2 kD) (LOC680551) alternative variant bSep08, complete mRNA.
<a href="#">LOC680558</a>	<a href="#">LOC680558.aSep08</a>	<a href="#">680558</a>	28238	397	5	120	similar to solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1 (LOC680558) alternative variant aSep08, mRNA.
<a href="#">LOC680586</a>	<a href="#">LOC680586.aSep08</a>	<a href="#">680586</a>	38229	813		87	hypothetical protein LOC680586 (9.6 kD) (LOC680586) mRNA.



<a href="#">LOC680590</a>	<a href="#">LOC680590.aSep08</a>	<a href="#">680590</a>	5990	546	4	95	hypothetical protein LOC680590 (10.6 kD) (LOC680590) alternative variant aSep08, mRNA.
<a href="#">LOC680590</a>	<a href="#">LOC680590.bSep08</a>	<a href="#">680590</a>	19529	747	8	72	hypothetical protein LOC680590 (8.3 kD) (LOC680590) alternative variant bSep08, mRNA.
<a href="#">LOC680590</a>	<a href="#">LOC680590.cSep08</a>	<a href="#">680590</a>	5397	308	2	50	hypothetical protein LOC680590 (LOC680590) alternative variant cSep08, mRNA.
<a href="#">LOC680590</a>	<a href="#">LOC680590.dSep08</a>	<a href="#">680590</a>	15150	491	5	45	hypothetical protein LOC680590 (LOC680590) alternative variant dSep08, mRNA.
<a href="#">LOC680606</a>	<a href="#">LOC680606.aSep08</a>	<a href="#">680606</a>	12594	272		90	hypothetical protein LOC680606 (LOC680606) mRNA.
<a href="#">LOC680614</a>	<a href="#">LOC680614.aSep08</a>	<a href="#">680614</a>	1579	466		137	similar to ETAA16 protein (LOC680614) mRNA.
<a href="#">LOC680633</a>	<a href="#">LOC680633.aSep08</a>	<a href="#">680633</a>	11026	693		54	hypothetical protein LOC680633 (LOC680633) mRNA.
<a href="#">LOC680666</a>	<a href="#">LOC680666.aSep08</a>	<a href="#">680666</a>	5823	723		76	hypothetical protein LOC680666 (LOC680666) mRNA.
<a href="#">LOC680682</a>	<a href="#">LOC680682.aSep08</a>	<a href="#">680682</a>	12946	618	4	163	hypothetical protein LOC680682 (LOC680682) alternative variant aSep08, mRNA.
<a href="#">LOC680682</a>	<a href="#">LOC680682.bSep08</a>	<a href="#">680682</a>	12052	833	2	120	hypothetical protein LOC680682 (14.0 kD) (LOC680682) alternative variant bSep08, mRNA.
<a href="#">LOC680682</a>	<a href="#">LOC680682.cSep08</a>	<a href="#">680682</a>	1725	694	2	114	hypothetical protein LOC680682 (13.5 kD) (LOC680682) alternative variant cSep08, mRNA.
<a href="#">LOC680682</a>	<a href="#">LOC680682.dSep08</a>	<a href="#">680682</a>	28173	709	2	116	hypothetical protein LOC680682 (LOC680682) alternative variant dSep08, mRNA.
<a href="#">LOC680687</a>	<a href="#">LOC680687.aSep08</a>	<a href="#">680687</a>	5818	1221	4	41	hypothetical protein LOC680687 (LOC680687) alternative variant aSep08, mRNA.
<a href="#">LOC680687</a>	<a href="#">LOC680687.cSep08</a>	<a href="#">680687</a>	2478	415	2	72	hypothetical protein LOC680687 (LOC680687) alternative variant cSep08, mRNA.
<a href="#">LOC680689</a>	<a href="#">LOC680689.aSep08</a>	<a href="#">680689</a>	5608	672		106	hypothetical protein LOC680689 (LOC680689) mRNA.
<a href="#">LOC680691</a>	<a href="#">LOC680691.aSep08</a>	<a href="#">680691</a>	3651	353		83	hypothetical protein LOC680691 (LOC680691) mRNA.
<a href="#">LOC680692</a>	<a href="#">LOC680692.aSep08</a>	<a href="#">680692</a>	36311	1944	1	483	similar to Golgi phosphoprotein 2 (Golgi membrane protein GP73) (54.3 kD) (LOC680692) alternative variant aSep08, complete mRNA.
<a href="#">LOC680692</a>	<a href="#">LOC680692.bSep08</a>	<a href="#">680692</a>	25306	486	1	121	similar to Golgi phosphoprotein 2 (Golgi membrane protein GP73) (LOC680692) alternative variant bSep08, mRNA.
<a href="#">LOC680693</a>	<a href="#">LOC680693.aSep08</a>	<a href="#">680693</a>	6113	956	6	180	similar to Sperm flagellar protein 1 (20.7 kD) (LOC680693) alternative variant aSep08, mRNA.
<a href="#">LOC680693</a>	<a href="#">LOC680693.bSep08</a>	<a href="#">680693</a>	1588	827	2	112	similar to Sperm flagellar protein 1 (LOC680693) alternative variant bSep08, mRNA.
<a href="#">LOC680711</a>	<a href="#">LOC680711.aSep08</a>	<a href="#">680711</a>	11603	1112		252	hypothetical protein LOC680711 (28.2 kD) (LOC680711) mRNA.
<a href="#">LOC680722</a>	<a href="#">LOC680722.aSep08</a>	<a href="#">680722</a>	2910	423		123	hypothetical protein LOC680722 (LOC680722) mRNA.
<a href="#">LOC680724</a>	<a href="#">LOC680724.aSep08</a>	<a href="#">680724</a>	5235	540		139	similar to NS1-associated protein 1 isoform 2 (LOC680724) mRNA.
<a href="#">LOC680726</a>	<a href="#">LOC680726.aSep08</a>	<a href="#">680726</a>	324910	1421	5	269	similar to RNA binding motif, single stranded interacting protein 3 isoform 1 (29.2 kD) (LOC680726) alternative variant aSep08, mRNA.
<a href="#">LOC680726</a>	<a href="#">LOC680726.bSep08</a>	<a href="#">680726</a>	150860	714	1	178	similar to RNA binding motif, single stranded interacting protein 3 isoform 1 (LOC680726) alternative variant bSep08, mRNA.

<a href="#">LOC680728</a>	<a href="#">LOC680728.aSep08</a>	<a href="#">680728</a>	2224	542		120	similar to Caspase-14 precursor (CASP-14) (LOC680728) alternative variant aSep08, mRNA.
<a href="#">LOC680745</a>	<a href="#">LOC680745.aSep08</a>	<a href="#">680745</a>	1585	326		88	hypothetical protein LOC680745 (LOC680745) mRNA.
<a href="#">LOC680752</a>	<a href="#">LOC680752.aSep08</a>	<a href="#">680752</a>	582	416		93	hypothetical protein LOC680752 (LOC680752) mRNA.
<a href="#">LOC680770</a>	<a href="#">LOC680770.bSep08</a>	<a href="#">680770</a>	5882	938	3	52	similar to dachshund b (LOC680770) alternative variant bSep08, mRNA.
<a href="#">LOC680782</a>	<a href="#">LOC680782.aSep08</a>	<a href="#">680782</a>	8140	3468	5	220	similar to signal peptidase complex subunit 3 homolog (LOC680782) alternative variant aSep08, mRNA.
<a href="#">LOC680787</a>	<a href="#">LOC680787.aSep08</a>	<a href="#">680787</a>	5324	619		124	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC680787) mRNA.
<a href="#">LOC680796</a>	<a href="#">LOC680796.aSep08</a>	<a href="#">680796</a>	17300	475		75	hypothetical protein LOC680796 (LOC680796) mRNA.
<a href="#">LOC680802</a>	<a href="#">LOC680802.aSep08</a>	<a href="#">680802</a>	2985	1639		91	similar to Zinc finger protein 45 (BRC1744) (10.1 kD) (LOC680802) mRNA.
<a href="#">LOC680815</a>	<a href="#">LOC680815.aSep08</a>	<a href="#">680815</a>	1743	1244	1	265	hypothetical protein LOC680815 (29.8 kD) (LOC680815) alternative variant aSep08, mRNA.
<a href="#">LOC680815</a>	<a href="#">LOC680815.cSep08</a>	<a href="#">680815</a>	1310	474		92	hypothetical protein LOC680815 (LOC680815) alternative variant cSep08, mRNA.
<a href="#">LOC680833</a>	<a href="#">LOC680833.aSep08</a>	<a href="#">680833</a>	6895	934		311	novel protein containing fibronectin type 3 FN3 domains (LOC680833) mRNA.
<a href="#">LOC680835</a>	<a href="#">LOC680835.aSep08</a>	<a href="#">680835</a>	8339	3224	19	1047	similar to cullin 7 (LOC680835) alternative variant aSep08, mRNA.
<a href="#">LOC680835</a>	<a href="#">LOC680835.bSep08</a>	<a href="#">680835</a>	1126	686	4	228	similar to cullin 7 (LOC680835) alternative variant bSep08, mRNA.
<a href="#">LOC680875</a>	<a href="#">LOC680875.aSep08</a>	<a href="#">680875</a>	6771	1789	4	491	similar to dystonin isoform 1 (LOC680875) alternative variant aSep08, mRNA.
<a href="#">LOC680875</a>	<a href="#">LOC680875.bSep08</a>	<a href="#">680875</a>	1341	787	4	211	similar to dystonin isoform 1 (LOC680875) alternative variant bSep08, mRNA.
<a href="#">LOC680875</a>	<a href="#">LOC680875.cSep08</a>	<a href="#">680875</a>	1200	800	4	194	similar to dystonin isoform 1 (LOC680875) alternative variant cSep08, mRNA.
<a href="#">LOC680889</a>	<a href="#">LOC680889.aSep08</a>	<a href="#">680889</a>	1801	827	2	65	hypothetical protein LOC680889 (7.0 kD) (LOC680889) alternative variant aSep08, mRNA.
<a href="#">LOC680889</a>	<a href="#">LOC680889.bSep08</a>	<a href="#">680889</a>	1546	542	2	41	hypothetical protein LOC680889 (4.7 kD) (LOC680889) alternative variant bSep08, mRNA.
<a href="#">LOC680893</a>	<a href="#">LOC680893.aSep08</a>	<a href="#">680893</a>	16720	752		83	hypothetical protein LOC680893 (LOC680893) mRNA.
<a href="#">LOC680897</a>	<a href="#">LOC680897.aSep08</a>	<a href="#">680897</a>	85151	287		95	similar to protein tyrosine phosphatase, receptor type, T (LOC680897) mRNA.
<a href="#">LOC680899</a>	<a href="#">LOC680899.aSep08</a>	<a href="#">680899</a>	3786	373		110	hypothetical protein LOC680899 (LOC680899) mRNA.
<a href="#">LOC680906</a>	<a href="#">LOC680906.aSep08</a>	<a href="#">680906</a>	5613	709		50	hypothetical protein LOC680906 (LOC680906) mRNA.
<a href="#">LOC680910</a>	<a href="#">LOC680910.aSep08</a>	<a href="#">680910</a>	3003	1029	2	50	similar to paired immunoglobulin-like type 2 receptor beta (LOC680910) alternative variant aSep08, mRNA.
<a href="#">LOC680910</a>	<a href="#">LOC680910.bSep08</a>	<a href="#">680910</a>	885	328	2	46	similar to paired immunoglobulin-like type 2 receptor beta (5.1 kD) (LOC680910) alternative variant bSep08, mRNA.
<a href="#">LOC680918</a>	<a href="#">LOC680918.bSep08</a>	<a href="#">680918</a>	16046	720	1	92	similar to spermatogenesis and centriole associated 1 (10.7 kD) (LOC680918) alternative variant bSep08, mRNA.
<a href="#">LOC680919</a>	<a href="#">LOC680919.aSep08</a>	<a href="#">680919</a>	6928	522		75	hypothetical protein LOC680919 (LOC680919) mRNA.

<a href="#">LOC680923</a>	<a href="#">LOC680923.aSep08</a>	<a href="#">680923</a>	869	625		208	similar to paired immunoglobulin-like type 2 receptor beta (LOC680923) mRNA.
<a href="#">LOC680929</a>	<a href="#">LOC680929.aSep08</a>	<a href="#">680929</a>	8361	399	4	49	similar to serine/threonine kinase (5.2 kD) (LOC680929) alternative variant aSep08, mRNA.
<a href="#">LOC680929</a>	<a href="#">LOC680929.bSep08</a>	<a href="#">680929</a>	5788	634	4	31	similar to serine/threonine kinase (3.5 kD) (LOC680929) alternative variant bSep08, mRNA.
<a href="#">LOC680929</a>	<a href="#">LOC680929.dSep08</a>	<a href="#">680929</a>	10755	545	3	44	similar to serine/threonine kinase (4.8 kD) (LOC680929) alternative variant dSep08, mRNA.
<a href="#">LOC680961</a>	<a href="#">LOC680961.aSep08</a>	<a href="#">680961</a>	54717	952	4	246	hypothetical protein LOC680961 (LOC680961) alternative variant aSep08, mRNA.
<a href="#">LOC680961</a>	<a href="#">LOC680961.bSep08</a>	<a href="#">680961</a>	54556	1413	3	246	hypothetical protein LOC680961 (LOC680961) alternative variant bSep08, mRNA.
<a href="#">LOC680961</a>	<a href="#">LOC680961.cSep08</a>	<a href="#">680961</a>	52835	1222	1	54	hypothetical protein LOC680961 (5.9 kD) (LOC680961) alternative variant cSep08, mRNA.
<a href="#">LOC680961</a>	<a href="#">LOC680961.dSep08</a>	<a href="#">680961</a>	1625	739	2	54	hypothetical protein LOC680961 (5.9 kD) (LOC680961) alternative variant dSep08, mRNA.
<a href="#">LOC680969</a>	<a href="#">LOC680969.aSep08</a>	<a href="#">680969</a>	2633	523		111	similar to uroplakin 3B isoform b (12.1 kD) (LOC680969) mRNA.
<a href="#">LOC681004</a>	<a href="#">LOC681004.aSep08</a>	<a href="#">681004</a>	19313	1786	7	429	similar to GC-rich sequence DNA-binding factor homolog (LOC681004) alternative variant aSep08, mRNA.
<a href="#">LOC681004</a>	<a href="#">LOC681004.bSep08</a>	<a href="#">681004</a>	19166	2437	12	411	similar to GC-rich sequence DNA-binding factor homolog (47.9 kD) (LOC681004) alternative variant bSep08, mRNA.
<a href="#">LOC681004</a>	<a href="#">LOC681004.cSep08</a>	<a href="#">681004</a>	4705	1516	6	227	similar to GC-rich sequence DNA-binding factor homolog (26.6 kD) (LOC681004) alternative variant cSep08, mRNA.
<a href="#">LOC681018</a>	<a href="#">LOC681018.aSep08</a>	<a href="#">681018</a>	964	455	1	111	hypothetical protein LOC681018 (13.0 kD) (LOC681018) alternative variant aSep08, mRNA.
<a href="#">LOC681018</a>	<a href="#">LOC681018.bSep08</a>	<a href="#">681018</a>	1004	378		104	hypothetical protein LOC681018 (10.4 kD) (LOC681018) alternative variant bSep08, mRNA.
<a href="#">LOC681025</a>	<a href="#">LOC681025.aSep08</a>	<a href="#">681025</a>	36767	750		250	hypothetical protein LOC681025 and similar to bile acid Coenzyme A: amino acid N-acyltransferase (LOC681025) mRNA.
<a href="#">LOC681025</a>	<a href="#">LOC681025.aSep08</a>	<a href="#">681043</a>	36767	750		250	hypothetical protein LOC681025 and similar to bile acid Coenzyme A: amino acid N-acyltransferase (LOC681025) mRNA.
<a href="#">LOC681027</a>	<a href="#">LOC681027.aSep08</a>	<a href="#">681027</a>	12770	625		104	hypothetical protein LOC681027 (11.1 kD) (LOC681027) mRNA.
<a href="#">LOC681031</a>	<a href="#">LOC681031.aSep08</a>	<a href="#">681031</a>	7468	335	4	76	similar to small nuclear ribonucleoprotein polypeptide G (8.5 kD) (LOC681031) alternative variant aSep08, mRNA.
<a href="#">LOC681031</a>	<a href="#">LOC681031.cSep08</a>	<a href="#">681031</a>	3616	727	3	64	similar to small nuclear ribonucleoprotein polypeptide G (7.1 kD) (LOC681031) alternative variant cSep08, mRNA.
<a href="#">LOC681035</a>	<a href="#">LOC681035.aSep08</a>	<a href="#">681035</a>	3827	397		97	similar to paired immunoglobulin-like type 2 receptor beta (LOC681035) mRNA.
<a href="#">LOC681048</a>	<a href="#">LOC681048.aSep08</a>	<a href="#">681048</a>	5915	694		90	similar to glutaredoxin cysteine-rich 1 protein (LOC681048) mRNA.
<a href="#">LOC681069</a>	<a href="#">LOC681069.aSep08</a>	<a href="#">681069</a>	7731	1320		212	similar to paired immunoglobulin-like type 2 receptor beta (23.9 kD) (LOC681069) mRNA.

<a href="#">LOC681086</a>	<a href="#">LOC681086.aSep08</a>	<a href="#">681086</a>	28055	968	7	181	similar to Phosphatidylinositol-glycan biosynthesis, class F protein (PIG-F) (20.6 kD) (LOC681086) alternative variant aSep08, mRNA.
<a href="#">LOC681086</a>	<a href="#">LOC681086.bSep08</a>	<a href="#">681086</a>	24477	613	4	137	similar to Phosphatidylinositol-glycan biosynthesis, class F protein (PIG-F) (LOC681086) alternative variant bSep08, mRNA.
<a href="#">LOC681086</a>	<a href="#">LOC681086.cSep08</a>	<a href="#">681086</a>	22874	591	4	88	similar to Phosphatidylinositol-glycan biosynthesis, class F protein (PIG-F) (LOC681086) alternative variant cSep08, mRNA.
<a href="#">LOC681086</a>	<a href="#">LOC681086.dSep08</a>	<a href="#">681086</a>	12325	829	2	44	similar to Phosphatidylinositol-glycan biosynthesis, class F protein (PIG-F) (4.9 kD) (LOC681086) alternative variant dSep08, mRNA.
<a href="#">LOC681166</a>	<a href="#">LOC681166.aSep08</a>	<a href="#">681166</a>	6205	310		62	similar to MIC2 like 1 (LOC681166) mRNA.
<a href="#">LOC681185</a>	<a href="#">LOC681185.aSep08</a>	<a href="#">681185</a>	13289	856	6	129	similar to centromere autoantigen H (LOC681185) alternative variant aSep08, mRNA.
<a href="#">LOC681185</a>	<a href="#">LOC681185.cSep08</a>	<a href="#">681185</a>	16086	1778	6	112	similar to centromere autoantigen H (LOC681185) alternative variant cSep08, mRNA.
<a href="#">LOC681185</a>	<a href="#">LOC681185.dSep08</a>	<a href="#">681185</a>	2055	663	2	61	similar to centromere autoantigen H (LOC681185) alternative variant dSep08, mRNA.
<a href="#">LOC681185</a>	<a href="#">LOC681185.eSep08</a>	<a href="#">681185</a>	3926	672	1	44	similar to centromere autoantigen H (LOC681185) alternative variant eSep08, mRNA.
<a href="#">LOC681186</a>	<a href="#">LOC681186.aSep08</a>	<a href="#">681186</a>	2755	540	3	125	hypothetical protein LOC681186 (LOC681186) alternative variant aSep08, mRNA.
<a href="#">LOC681186</a>	<a href="#">LOC681186.bSep08</a>	<a href="#">681186</a>	2222	441	1	92	hypothetical protein LOC681186 (LOC681186) alternative variant bSep08, mRNA.
<a href="#">LOC681200</a>	<a href="#">LOC681200.aSep08</a>	<a href="#">681200</a>	33530	708		230	similar to Ladybird homeobox corepressor 1 (LOC681200) mRNA.
<a href="#">LOC681205</a>	<a href="#">LOC681205.aSep08</a>	<a href="#">681205</a>	4235	440		75	hypothetical protein LOC681205 (LOC681205) mRNA.
<a href="#">LOC681219</a>	<a href="#">LOC681219.aSep08</a>	<a href="#">681219</a>	2428	852		71	hypothetical protein LOC681219 (7.9 kD) (LOC681219) mRNA.
<a href="#">LOC681237</a>	<a href="#">LOC681237.aSep08</a>	<a href="#">681237</a>	1689	264		65	hypothetical protein LOC681237 (LOC681237) mRNA.
<a href="#">LOC681239</a>	<a href="#">LOC681239.aSep08</a>	<a href="#">681239</a>	1208	945		114	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (12.6 kD) (LOC681239) mRNA.
<a href="#">LOC681251</a>	<a href="#">LOC681251.aSep08</a>	<a href="#">681251</a>	1674	561		186	hypothetical protein LOC681251 (LOC681251) mRNA.
<a href="#">LOC681252</a>	<a href="#">LOC681252.aSep08</a>	<a href="#">681252</a>	5705	4187		309	similar to Myristoylated alanine-rich C-kinase substrate (MARCKS) (Protein kinase C substrate 80 kDa protein) (29.8 kD) (LOC681252) mRNA.
<a href="#">LOC681258</a>	<a href="#">LOC681258.aSep08</a>	<a href="#">681258</a>	7601	355		28	similar to La-related protein 4 (La ribonucleoprotein domain family member 4) (LOC681258) mRNA.
<a href="#">LOC681272</a>	<a href="#">LOC681272.aSep08</a>	<a href="#">681272</a>	4576	287		53	hypothetical protein LOC681272 (LOC681272) mRNA.
<a href="#">LOC681282</a>	<a href="#">LOC681282.aSep08</a>	<a href="#">681282</a>	1989	1078		164	hypothetical protein LOC681282 (18.4 kD) (LOC681282) mRNA.
<a href="#">LOC681302</a>	<a href="#">LOC681302.aSep08</a>	<a href="#">681302</a>	28710	457	5	152	similar to Multifunctional protein ADE2 (LOC681302) alternative variant aSep08, mRNA.
<a href="#">LOC681303</a>	<a href="#">LOC681303.aSep08</a>	<a href="#">681303</a>	11008	690		229	similar to tumor endothelial marker 8 isoform 1 precursor (LOC681303) mRNA.

<a href="#">LOC681309</a>	<a href="#">LOC681309.aSep08</a>	<a href="#">681309</a>	5654	1683		476	similar to Thrombospondin-3 precursor (LOC681309) mRNA.
<a href="#">LOC681343</a>	<a href="#">LOC681343.aSep08</a>	<a href="#">681343</a>	25208	327		81	hypothetical protein LOC681343 (LOC681343) mRNA.
<a href="#">LOC681351</a>	<a href="#">LOC681351.aSep08</a>	<a href="#">681351</a>	8646	1034		284	similar to apolipoprotein L, 3 (LOC681351) mRNA.
<a href="#">LOC681370</a>	<a href="#">LOC681370.aSep08</a>	<a href="#">681370</a>	445	349		77	hypothetical protein LOC681370 (LOC681370) mRNA.
<a href="#">LOC681381</a>	<a href="#">LOC681381.aSep08</a>	<a href="#">681381</a>	62540	582	6	152	similar to Zinc finger CW-type PWWP domain protein 1 homolog (LOC681381) alternative variant aSep08, mRNA.
<a href="#">LOC681381</a>	<a href="#">LOC681381.bSep08</a>	<a href="#">681381</a>	79438	487	4	108	similar to Zinc finger CW-type PWWP domain protein 1 homolog (LOC681381) alternative variant bSep08, mRNA.
<a href="#">LOC681381</a>	<a href="#">LOC681381.cSep08</a>	<a href="#">681381</a>	36425	470	2	89	similar to Zinc finger CW-type PWWP domain protein 1 homolog (LOC681381) alternative variant cSep08, mRNA.
<a href="#">LOC681395</a>	<a href="#">LOC681395.bSep08</a>	<a href="#">681395</a>	1797	793	1	24	similar to ring finger protein 133 (LOC681395) alternative variant bSep08, mRNA.
<a href="#">LOC681397</a>	<a href="#">LOC681397.bSep08</a>	<a href="#">681397</a>	1069	238		48	fgfr1 oncogene partner (LOC681397) alternative variant bSep08, mRNA.
<a href="#">LOC681397</a>	<a href="#">LOC681397.cSep08</a>	<a href="#">681397</a>	2489	265		30	putative protein (LOC681397) alternative variant cSep08, mRNA.
<a href="#">LOC681398</a>	<a href="#">LOC681398.aSep08</a>	<a href="#">681398</a>	53341	679		153	hypothetical protein LOC681398 (16.2 kD) (LOC681398) mRNA.
<a href="#">LOC681405</a>	<a href="#">LOC681405.aSep08</a>	<a href="#">681405</a>	2010	534		93	similar to RIKEN cDNA 5031410I06 (LOC681405) mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.bSep08</a>	<a href="#">681578</a>	127122	730	7	203	similar to ring finger protein 13 (LOC681578) alternative variant bSep08, mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.cSep08</a>	<a href="#">681578</a>	127101	709	7	202	similar to ring finger protein 13 (LOC681578) alternative variant cSep08, mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.dSep08</a>	<a href="#">681578</a>	13769	1817	3	148	similar to ring finger protein 13 (16.4 kD) (LOC681578) alternative variant dSep08, mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.eSep08</a>	<a href="#">681578</a>	127119	636	6	141	similar to ring finger protein 13 (15.5 kD) (LOC681578) alternative variant eSep08, mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.fSep08</a>	<a href="#">681578</a>	119432	548	6	138	similar to ring finger protein 13 (LOC681578) alternative variant fSep08, mRNA.
<a href="#">LOC681578</a>	<a href="#">LOC681578.hSep08</a>	<a href="#">681578</a>	127081	573	6	21	similar to ring finger protein 13 (2.6 kD) (LOC681578) alternative variant hSep08, mRNA.
<a href="#">LOC683514</a>	<a href="#">LOC683514.bSep08</a>	<a href="#">683514</a>	18700	602	1	67	hypothetical protein LOC683514 (LOC683514) alternative variant bSep08, mRNA.
<a href="#">LOC684322</a>	<a href="#">LOC684322.bSep08</a>	<a href="#">684322</a>	6309	765	2	144	similar to potassium channel modulatory factor 1 (LOC684322) alternative variant bSep08, mRNA.
<a href="#">LOC684964</a>	<a href="#">LOC684964.aSep08</a>	<a href="#">684964</a>	110324	723		46	hypothetical protein LOC684964 (5.3 kD) (LOC684964) mRNA.
<a href="#">LOC684989</a>	<a href="#">LOC684989.aSep08</a>	<a href="#">684989</a>	2004	1121		96	hypothetical protein LOC684989 (LOC684989) mRNA.
<a href="#">LOC684993</a>	<a href="#">LOC684993.bSep08</a>	<a href="#">684993</a>	4812	1389	3	178	hypothetical protein LOC684993 (LOC684993) alternative variant bSep08, mRNA.
<a href="#">LOC684993</a>	<a href="#">LOC684993.dSep08</a>	<a href="#">684993</a>	623	315	2	40	hypothetical protein LOC684993 (LOC684993) alternative variant dSep08, mRNA.
<a href="#">LOC684994</a>	<a href="#">LOC684994.aSep08</a>	<a href="#">684994</a>	5959	398	1	110	similar to retinoic acid receptor, beta (LOC684994) alternative variant aSep08, mRNA.

<a href="#">LOC684994</a>	<a href="#">LOC684994.bSep08</a>	<a href="#">684994</a>	27082	683	1	102	similar to retinoic acid receptor, beta (LOC684994) alternative variant bSep08, mRNA.
<a href="#">LOC684996</a>	<a href="#">LOC684996.aSep08</a>	<a href="#">684996</a>	3176	2003	3	178	putative protein of vertebrate origin (19.1 kD) (LOC684996) alternative variant aSep08, mRNA.
<a href="#">LOC684998</a>	<a href="#">LOC684998.bSep08</a>	<a href="#">684998</a>	54565	443	4	36	hypothetical protein LOC684998 (LOC684998) alternative variant bSep08, mRNA.
<a href="#">LOC685001</a>	<a href="#">LOC685001.aSep08</a>	<a href="#">685001</a>	9237	3647	4	134	similar to MIR-interacting saposin-like protein precursor (Transmembrane protein 4) (Putative secreted protein ZSIG9) (15.6 kD) (LOC685001) alternative variant aSep08, mRNA.
<a href="#">LOC685002</a>	<a href="#">LOC685002.aSep08</a>	<a href="#">685002</a>	773	446		51	hypothetical protein LOC685002 (5.6 kD) (LOC685002) mRNA.
<a href="#">LOC685009</a>	<a href="#">LOC685009.aSep08</a>	<a href="#">685009</a>	9733	410		107	hypothetical protein LOC685009 (LOC685009) mRNA.
<a href="#">LOC685019</a>	<a href="#">LOC685019.aSep08</a>	<a href="#">685019</a>	6081	690		104	similar to serine/threonine kinase (LOC685019) mRNA.
<a href="#">LOC685020</a>	<a href="#">LOC685020.aSep08</a>	<a href="#">685020</a>	10706	1204		214	similar to paired immunoglobulin-like type 2 receptor alpha (LOC685020) mRNA.
<a href="#">LOC685029</a>	<a href="#">LOC685029.aSep08</a>	<a href="#">685029</a>	7966	523	3	74	similar to anaphase promoting complex subunit 13 (8.4 kD) (LOC685029) alternative variant aSep08, mRNA.
<a href="#">LOC685029</a>	<a href="#">LOC685029.bSep08</a>	<a href="#">685029</a>	8459	658	3	74	similar to anaphase promoting complex subunit 13 (8.4 kD) (LOC685029) alternative variant bSep08, mRNA.
<a href="#">LOC685030</a>	<a href="#">LOC685030.aSep08</a>	<a href="#">685030</a>	3215	735		191	similar to paired immunoglobulin-like type 2 receptor beta (LOC685030) mRNA.
<a href="#">LOC685055</a>	<a href="#">LOC685055.aSep08</a>	<a href="#">685055</a>	4129	631		210	similar to Serine/threonine-protein kinase ATR (Ataxia telangiectasia and Rad3-related protein) (LOC685055) mRNA.
<a href="#">LOC685067</a>	<a href="#">LOC685067.aSep08</a>	<a href="#">685067</a>	9624	774		257	similar to guanylate binding protein family, member 6 (LOC685067) mRNA.
<a href="#">LOC685072</a>	<a href="#">LOC685072.aSep08</a>	<a href="#">685072</a>	10879	1759	6	485	similar to Retinoic acid receptor gamma-A (RAR-gamma-A) (53.7 kD) (LOC685072) alternative variant aSep08, mRNA.
<a href="#">LOC685072</a>	<a href="#">LOC685072.bSep08</a>	<a href="#">685072</a>	13416	613	3	204	similar to Retinoic acid receptor gamma-A (RAR-gamma-A) (LOC685072) alternative variant bSep08, mRNA.
<a href="#">LOC685072</a>	<a href="#">LOC685072.cSep08</a>	<a href="#">685072</a>	1471	265	1	59	similar to Retinoic acid receptor gamma-A (RAR-gamma-A) (LOC685072) alternative variant cSep08, mRNA.
<a href="#">LOC685074</a>	<a href="#">LOC685074.aSep08</a>	<a href="#">685074</a>	10785	1115		84	hypothetical protein LOC685074 (9.4 kD) (LOC685074) mRNA.
<a href="#">LOC685079</a>	<a href="#">LOC685079.aSep08</a>	<a href="#">685079</a>	4505	1756	4	259	similar to Protein SYS1 homolog (LOC685079) mRNA.
<a href="#">LOC685080</a>	<a href="#">LOC685080.aSep08</a>	<a href="#">685080</a>	7919	747		248	similar to nipsnap homolog 3A (LOC685080) mRNA.
<a href="#">LOC685081</a>	<a href="#">LOC685081.aSep08</a>	<a href="#">685081</a>	1390	786	2	165	similar to solute carrier family 22 (organic cation transporter), member 13 (LOC685081) alternative variant aSep08, mRNA.
<a href="#">LOC685081</a>	<a href="#">LOC685081.bSep08</a>	<a href="#">685081</a>	2528	755	2	99	similar to solute carrier family 22 (organic cation transporter), member 13 (LOC685081) alternative variant bSep08, mRNA.
<a href="#">LOC685095</a>	<a href="#">LOC685095.aSep08</a>	<a href="#">685095</a>	33433	1353		114	hypothetical protein LOC685095 (13.0 kD) (LOC685095) mRNA.
<a href="#">LOC685099</a>	<a href="#">LOC685099.bSep08</a>	<a href="#">685099</a>	3818	784	1	89	similar to holdem CG15329-PA (LOC685099) alternative variant bSep08, mRNA.

<a href="#">LOC685114</a>	<a href="#">LOC685114.aSep08</a>	<a href="#">685114</a>	2558	602	3	144	hypothetical protein LOC685114 (LOC685114) alternative variant aSep08, mRNA.
<a href="#">LOC685114</a>	<a href="#">LOC685114.bSep08</a>	<a href="#">685114</a>	2378	430	3	122	hypothetical protein LOC685114 (LOC685114) alternative variant bSep08, mRNA.
<a href="#">LOC685128</a>	<a href="#">LOC685128.aSep08</a>	<a href="#">685128</a>	4082	604		145	similar to gastroke 1 (LOC685128) mRNA.
<a href="#">LOC685136</a>	<a href="#">LOC685136.aSep08</a>	<a href="#">685136</a>	2332	452		98	similar to alpha 3 type VI collagen isoform 1 precursor (LOC685136) mRNA.
<a href="#">LOC685144</a>	<a href="#">LOC685144.bSep08</a>	<a href="#">685144</a>	21908	4460	23	1095	protein transport sec24c C (118.5 kD) (LOC685144) alternative variant bSep08, mRNA.
<a href="#">LOC685144</a>	<a href="#">LOC685144.cSep08</a>	<a href="#">685144</a>	1000	763	3	111	sec24 related gene family member C (12.3 kD) (LOC685144) alternative variant cSep08, mRNA.
<a href="#">LOC685144</a>	<a href="#">LOC685144.dSep08</a>	<a href="#">685144</a>	717	400	3	107	sec24 related gene family member C CRA b (LOC685144) alternative variant dSep08, mRNA.
<a href="#">LOC685144</a>	<a href="#">LOC685144.eSep08</a>	<a href="#">685144</a>	1821	447	2	71	putative protein (LOC685144) alternative variant eSep08, mRNA.
<a href="#">LOC685152</a>	<a href="#">LOC685152.aSep08</a>	<a href="#">685152</a>	5755	2779	7	375	ATPase class I type 8B member 2 (LOC685152) alternative variant aSep08, mRNA.
<a href="#">LOC685152</a>	<a href="#">LOC685152.bSep08</a>	<a href="#">685152</a>	1259	877	3	153	phospholipid-transporting ATPase ID (LOC685152) alternative variant bSep08, mRNA.
<a href="#">LOC685152</a>	<a href="#">LOC685152.cSep08</a>	<a href="#">685152</a>	1026	783	2	112	phospholipid-transporting ATPase ID (12.6 kD) (LOC685152) alternative variant cSep08, mRNA.
<a href="#">LOC685158</a>	<a href="#">LOC685158.aSep08</a>	<a href="#">685158</a>	520777	864		233	similar to CG8138-PA (LOC685158) mRNA.
<a href="#">LOC685160</a>	<a href="#">LOC685160.aSep08</a>	<a href="#">685160</a>	2067	409		47	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC685160) mRNA.
<a href="#">LOC685172</a>	<a href="#">LOC685172.aSep08</a>	<a href="#">685172</a>	66155	661		110	hypothetical protein LOC685172 (LOC685172) mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.aSep08</a>	<a href="#">685179</a>	24446	3167	26	987	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant aSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.bSep08</a>	<a href="#">685179</a>	25795	1977	17	637	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant bSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.cSep08</a>	<a href="#">685179</a>	13897	1460	13	486	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant cSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.dSep08</a>	<a href="#">685179</a>	5336	2356	5	355	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant dSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.fSep08</a>	<a href="#">685179</a>	7775	410	5	136	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant fSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.gSep08</a>	<a href="#">685179</a>	1190	380	4	126	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant gSep08, mRNA.
<a href="#">LOC685179</a>	<a href="#">LOC685179.hSep08</a>	<a href="#">685179</a>	1390	515	2	112	similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2 (LOC685179) alternative variant hSep08, mRNA.

<a href="#">LOC685187</a>	<a href="#">LOC685187.aSep08</a>	<a href="#">685187</a>	1280	758		252	similar to ADAM 21 precursor (A disintegrin and metalloproteinase domain 21) (ADAM 31) (LOC685187) mRNA.
<a href="#">LOC685195</a>	<a href="#">LOC685195.aSep08</a>	<a href="#">685195</a>	1534	931	3	137	hypothetical protein LOC685195 (LOC685195) alternative variant aSep08, mRNA.
<a href="#">LOC685210</a>	<a href="#">LOC685210.aSep08</a>	<a href="#">685210</a>	3050	648		215	similar to RAN binding protein 5 (LOC685210) mRNA.
<a href="#">LOC685215</a>	<a href="#">LOC685215.aSep08</a>	<a href="#">685215</a>	12829	738		245	putative protein (LOC685215) mRNA.
<a href="#">LOC685227</a>	<a href="#">LOC685227.aSep08</a>	<a href="#">685227</a>	2272	266		83	similar to WAP four-disulfide core domain 11 precursor (LOC685227) mRNA.
<a href="#">LOC685233</a>	<a href="#">LOC685233.aSep08</a>	<a href="#">685233</a>	39814	445		60	hypothetical protein LOC685233 (LOC685233) mRNA.
<a href="#">LOC685249</a>	<a href="#">LOC685249.aSep08</a>	<a href="#">685249</a>	2105	704	2	110	hypothetical protein LOC685249 (12.2 kD) (LOC685249) alternative variant aSep08, mRNA.
<a href="#">LOC685249</a>	<a href="#">LOC685249.bSep08</a>	<a href="#">685249</a>	6275	622	3	81	hypothetical protein LOC685249 (LOC685249) alternative variant bSep08, mRNA.
<a href="#">LOC685249</a>	<a href="#">LOC685249.cSep08</a>	<a href="#">685249</a>	4332	382	2	63	hypothetical protein LOC685249 (LOC685249) alternative variant cSep08, mRNA.
<a href="#">LOC685253</a>	<a href="#">LOC685253.aSep08</a>	<a href="#">685253</a>	2735	510		57	hypothetical protein LOC685253 (LOC685253) mRNA.
<a href="#">LOC685269</a>	<a href="#">LOC685269.aSep08</a>	<a href="#">685269</a>	3894	958		314	similar to Integrin alpha-IIb precursor (Platelet membrane glycoprotein IIb) (GPalpha IIb) (GPIIb) (CD41 antigen) (LOC685269) mRNA.
<a href="#">LOC685276</a>	<a href="#">LOC685276.aSep08</a>	<a href="#">685276</a>	23203	265		77	hypothetical protein LOC685276 (LOC685276) mRNA.
<a href="#">LOC685293</a>	<a href="#">LOC685293.aSep08</a>	<a href="#">685293</a>	2225	1025		161	similar to RT1 class Ia, locus A2 (18.1 kD) (LOC685293) mRNA.
<a href="#">LOC685294</a>	<a href="#">LOC685294.aSep08</a>	<a href="#">685294</a>	22558	743		132	similar to RalA-binding protein 1 (RalBP1) (Ral-interacting protein 1) (Cytocentrin) (Dinitrophenyl S-glutathione ATPase) (DNP-SG ATPase) (LOC685294) mRNA.
<a href="#">LOC685297</a>	<a href="#">LOC685297.aSep08</a>	<a href="#">685297</a>	13611	2121		483	similar to BAI1-associated protein 3 (BAP3) (LOC685297) alternative variant aSep08, mRNA.
<a href="#">LOC685297</a>	<a href="#">LOC685297.bSep08</a>	<a href="#">685297</a>	3341	2358		448	similar to BAI1-associated protein 3 (BAP3) (LOC685297) alternative variant bSep08, mRNA.
<a href="#">LOC685317</a>	<a href="#">LOC685317.aSep08</a>	<a href="#">685317</a>	78500	491		72	hypothetical protein LOC685317 (8.1 kD) (LOC685317) mRNA.
<a href="#">LOC685336</a>	<a href="#">LOC685336.aSep08</a>	<a href="#">685336</a>	22610	616		72	hypothetical protein LOC685336 (8.0 kD) (LOC685336) mRNA.
<a href="#">LOC685385</a>	<a href="#">LOC685385.aSep08</a>	<a href="#">685385</a>	1853	1096		199	similar to S100 calcium binding protein A14 (LOC685385) alternative variant aSep08, mRNA.
<a href="#">LOC685405</a>	<a href="#">LOC685405.aSep08</a>	<a href="#">685405</a>	5125	3405		230	similar to family 53, member C protein (LOC685405) mRNA.
<a href="#">LOC685440</a>	<a href="#">LOC685440.aSep08</a>	<a href="#">685440</a>	4181	470		95	similar to NIMA (never in mitosis gene a)-related expressed kinase 5 (LOC685440) alternative variant aSep08, mRNA.
<a href="#">LOC685444</a>	<a href="#">LOC685444.aSep08</a>	<a href="#">685444</a>	9180	1567		77	hypothetical protein LOC685444 (LOC685444) alternative variant aSep08, mRNA.
<a href="#">LOC685454</a>	<a href="#">LOC685454.aSep08</a>	<a href="#">685454</a>	16512	231		35	hypothetical protein LOC685454 (LOC685454) mRNA.
<a href="#">LOC685487</a>	<a href="#">LOC685487.aSep08</a>	<a href="#">685487</a>	47890	355		115	similar to ring finger protein 43 (LOC685487) mRNA.
<a href="#">LOC685527</a>	<a href="#">LOC685527.aSep08</a>	<a href="#">685527</a>	3832	395	1	59	hypothetical protein LOC685527 (LOC685527) alternative variant aSep08, mRNA.



<a href="#">LOC685527</a>	<a href="#">LOC685527.bSep08</a>	<a href="#">685527</a>	4064	548		47	hypothetical protein LOC685527 (4.8 kD) (LOC685527) alternative variant bSep08, mRNA.
<a href="#">LOC685569</a>	<a href="#">LOC685569.aSep08</a>	<a href="#">685569</a>	2247	530	2	124	hypothetical protein LOC685569 (LOC685569) alternative variant aSep08, mRNA.
<a href="#">LOC685574</a>	<a href="#">LOC685574.aSep08</a>	<a href="#">685574</a>	7021	344		63	similar to zinc finger protein 334 (LOC685574) mRNA.
<a href="#">LOC685576</a>	<a href="#">LOC685576.aSep08</a>	<a href="#">685576</a>	1324	567		58	hypothetical protein LOC685576 (LOC685576) mRNA.
<a href="#">LOC685580</a>	<a href="#">LOC685580.aSep08</a>	<a href="#">685580</a>	4405	733		244	hypothetical protein LOC685580 (LOC685580) mRNA.
<a href="#">LOC685584</a>	<a href="#">LOC685584.aSep08</a>	<a href="#">685584</a>	1151	392		109	hypothetical protein LOC685584 (LOC685584) mRNA.
<a href="#">LOC685601</a>	<a href="#">LOC685601.aSep08</a>	<a href="#">685601</a>	22174	754		250	similar to MICAL CG33208-PB, isoform B (LOC685601) mRNA.
<a href="#">LOC685608</a>	<a href="#">LOC685608.aSep08</a>	<a href="#">685608</a>	3600	603	4	137	hypothetical protein LOC685608 (14.4 kD) (LOC685608) alternative variant aSep08, mRNA.
<a href="#">LOC685609</a>	<a href="#">LOC685609.aSep08</a>	<a href="#">685609</a>	1784	755		118	hypothetical protein LOC685609 (14.0 kD) (LOC685609) mRNA.
<a href="#">LOC685612</a>	<a href="#">LOC685612.aSep08</a>	<a href="#">685612</a>	10194	1703		144	similar to Emu2 (LOC685612) mRNA.
<a href="#">LOC685619</a>	<a href="#">LOC685619.aSep08</a>	<a href="#">685619</a>	2260	901	2	246	similar to TP53-regulating kinase (p53-related protein kinase) (Nori-2) (27.6 kD) (LOC685619) alternative variant aSep08, mRNA.
<a href="#">LOC685619</a>	<a href="#">LOC685619.bSep08</a>	<a href="#">685619</a>	3035	1878	2	245	similar to TP53-regulating kinase (p53-related protein kinase) (Nori-2) (27.0 kD) (LOC685619) alternative variant bSep08, mRNA.
<a href="#">LOC685632</a>	<a href="#">LOC685632.aSep08</a>	<a href="#">685632</a>	1413	698		87	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC685632) mRNA.
<a href="#">LOC685634</a>	<a href="#">LOC685634.aSep08</a>	<a href="#">685634</a>	3811	981	2	89	hypothetical protein LOC685634 (LOC685634) alternative variant aSep08, mRNA.
<a href="#">LOC685634</a>	<a href="#">LOC685634.bSep08</a>	<a href="#">685634</a>	1751	486	2	83	hypothetical protein LOC685634 (LOC685634) alternative variant bSep08, mRNA.
<a href="#">LOC685645</a>	<a href="#">LOC685645.aSep08</a>	<a href="#">685645</a>	1978	635		143	similar to carboxylesterase isoenzyme gene (LOC685645) mRNA.
<a href="#">LOC685652</a>	<a href="#">LOC685652.aSep08</a>	<a href="#">685652</a>	1345	678		90	similar to keratin 6L (LOC685652) mRNA.
<a href="#">LOC685661</a>	<a href="#">LOC685661.aSep08</a>	<a href="#">685661</a>	2495	437		145	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC685661) mRNA.
<a href="#">LOC685663</a>	<a href="#">LOC685663.aSep08</a>	<a href="#">685663</a>	4500	599		188	hypothetical protein LOC685663 (LOC685663) mRNA.
<a href="#">LOC685671</a>	<a href="#">LOC685671.aSep08</a>	<a href="#">685671</a>	7706	293		97	similar to myocyte enhancer factor 2C (LOC685671) mRNA.
<a href="#">LOC685694</a>	<a href="#">LOC685694.aSep08</a>	<a href="#">685694</a>	5605	499	2	153	similar to TruB pseudouridine (psi) synthase homolog 2 (LOC685694) alternative variant aSep08, mRNA.
<a href="#">LOC685694</a>	<a href="#">LOC685694.bSep08</a>	<a href="#">685694</a>	4890	672	1	150	similar to TruB pseudouridine (psi) synthase homolog 2 (LOC685694) alternative variant bSep08, mRNA.
<a href="#">LOC685700</a>	<a href="#">LOC685700.aSep08</a>	<a href="#">685700</a>	5532	635		73	hypothetical protein LOC685700 (LOC685700) mRNA.
<a href="#">LOC685702</a>	<a href="#">LOC685702.cSep08</a>	<a href="#">685702</a>	3587	648	2	61	similar to C44B7.7 (LOC685702) alternative variant cSep08, mRNA.
<a href="#">LOC685707</a>	<a href="#">LOC685707.aSep08</a>	<a href="#">685707</a>	49014	405		134	similar to neuron navigator 1 (LOC685707) mRNA.
<a href="#">LOC685716</a>	<a href="#">LOC685716.aSep08</a>	<a href="#">685716</a>	2541	384		127	similar to OX-2 membrane glycoprotein precursor (MRC OX-2 antigen) (CD200 antigen) (LOC685716) mRNA.

<a href="#">LOC685729</a>	<a href="#">LOC685729.aSep08</a>	<a href="#">685729</a>	13005	732		62	similar to dimerization cofactor of hepatocyte nuclear factor 1 (HNF1) from muscle (LOC685729) mRNA.
<a href="#">LOC685734</a>	<a href="#">LOC685734.aSep08</a>	<a href="#">685734</a>	13930	721		76	similar to erythrocyte membrane protein band 4.1 like 4B isoform 2 (LOC685734) mRNA.
<a href="#">LOC685742</a>	<a href="#">LOC685742.aSep08</a>	<a href="#">685742</a>	3010	423	2	71	hypothetical protein LOC685742 (8.2 kD) (LOC685742) alternative variant aSep08, mRNA.
<a href="#">LOC685746</a>	<a href="#">LOC685746.aSep08</a>	<a href="#">685746</a>	2131	666		117	similar to RIKEN cDNA 5031410I06 (LOC685746) mRNA.
<a href="#">LOC685767</a>	<a href="#">LOC685767.aSep08</a>	<a href="#">685767</a>	20004	731		178	similar to OX-2 membrane glycoprotein precursor (MRC OX-2 antigen) (CD200 antigen) (LOC685767) mRNA.
<a href="#">LOC685779</a>	<a href="#">LOC685779.aSep08</a>	<a href="#">685779</a>	40804	739		125	hypothetical protein LOC685779 (LOC685779) mRNA.
<a href="#">LOC685785</a>	<a href="#">LOC685785.aSep08</a>	<a href="#">685785</a>	90757	775		172	similar to solute carrier organic anion transporter family, member 6c1 (19.1 kD) (LOC685785) mRNA.
<a href="#">LOC685792</a>	<a href="#">LOC685792.aSep08</a>	<a href="#">685792</a>	2490	437		145	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC685792) mRNA.
<a href="#">LOC685808</a>	<a href="#">LOC685808.aSep08</a>	<a href="#">685808</a>	2403	677		64	similar to transmembrane NK cell receptor 2B4 (LOC685808) mRNA.
<a href="#">LOC685817</a>	<a href="#">LOC685817.aSep08</a>	<a href="#">685817</a>	485	362		10	similar to Myosin-9B (Myosin IXb) (Unconventional myosin-9b) (1.1 kD) (LOC685817) mRNA.
<a href="#">LOC685834</a>	<a href="#">LOC685834.aSep08</a>	<a href="#">685834</a>	4426	871	2	90	similar to zinc finger protein 740 (LOC685834) alternative variant aSep08, mRNA.
<a href="#">LOC685834</a>	<a href="#">LOC685834.bSep08</a>	<a href="#">685834</a>	4380	586	2	63	similar to zinc finger protein 740 (LOC685834) alternative variant bSep08, mRNA.
<a href="#">LOC685849</a>	<a href="#">LOC685849.aSep08</a>	<a href="#">685849</a>	5536	1292	1	233	hypothetical protein LOC685849 (25.7 kD) (LOC685849) alternative variant aSep08, mRNA.
<a href="#">LOC685849</a>	<a href="#">LOC685849.bSep08</a>	<a href="#">685849</a>	4969	728	1	105	hypothetical protein LOC685849 (LOC685849) alternative variant bSep08, mRNA.
<a href="#">LOC685868</a>	<a href="#">LOC685868.aSep08</a>	<a href="#">685868</a>	45777	688		229	similar to otogelin (LOC685868) mRNA.
<a href="#">LOC685888</a>	<a href="#">LOC685888.aSep08</a>	<a href="#">685888</a>	8132	832	3	65	hypothetical protein LOC685888 (LOC685888) alternative variant aSep08, mRNA.
<a href="#">LOC685888</a>	<a href="#">LOC685888.bSep08</a>	<a href="#">685888</a>	8162	695	3	32	hypothetical protein LOC685888 (LOC685888) alternative variant bSep08, mRNA.
<a href="#">LOC685888</a>	<a href="#">LOC685888.cSep08</a>	<a href="#">685888</a>	2249	540	2	67	hypothetical protein LOC685888 (LOC685888) alternative variant cSep08, mRNA.
<a href="#">LOC685890</a>	<a href="#">LOC685890.aSep08</a>	<a href="#">685890</a>	4396	688		140	hypothetical protein LOC685890 (LOC685890) mRNA.
<a href="#">LOC685906</a>	<a href="#">LOC685906.aSep08</a>	<a href="#">685906</a>	56893	656	5	218	similar to splicing coactivator subunit SRm300 (LOC685906) alternative variant aSep08, mRNA.
<a href="#">LOC685908</a>	<a href="#">LOC685908.aSep08</a>	<a href="#">685908</a>	10257	588		125	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (13.4 kD) (LOC685908) mRNA.
<a href="#">LOC685909</a>	<a href="#">LOC685909.bSep08</a>	<a href="#">685909</a>	6076	623	3	107	similar to H2A histone family, member V isoform 1 (11.6 kD) (LOC685909) alternative variant bSep08, mRNA.
<a href="#">LOC685923</a>	<a href="#">LOC685923.bSep08</a>	<a href="#">685923</a>	1744	653	1	148	similar to cerebral cavernous malformation 2 homolog (LOC685923) alternative variant bSep08, mRNA.
<a href="#">LOC685925</a>	<a href="#">LOC685925.aSep08</a>	<a href="#">685925</a>	20707	599		163	similar to zinc finger protein 455 (LOC685925) mRNA.
<a href="#">LOC685931</a>	<a href="#">LOC685931.aSep08</a>	<a href="#">685931</a>	1975	880		87	similar to serine/threonine kinase (LOC685931) mRNA.

<a href="#">LOC685961</a>	<a href="#">LOC685961.aSep08</a>	<a href="#">685961</a>	69433	756		140	similar to SIRP beta 1 isoform 3 (15.6 kD) (LOC685961) mRNA.
<a href="#">LOC685964</a>	<a href="#">LOC685964.bSep08</a>	<a href="#">685964</a>	4470	423	1	110	hypothetical protein LOC685964 (LOC685964) alternative variant bSep08, mRNA.
<a href="#">LOC685974</a>	<a href="#">LOC685974.aSep08</a>	<a href="#">685974</a>	3611	730		243	similar to spermatogenesis associated glutamate (E)-rich protein 4b (LOC685974) mRNA.
<a href="#">LOC685988</a>	<a href="#">LOC685988.aSep08</a>	<a href="#">685988</a>	3026	667		117	hypothetical protein LOC685988 (LOC685988) mRNA.
<a href="#">LOC686006</a>	<a href="#">LOC686006.aSep08</a>	<a href="#">686006</a>	1062	757		97	hypothetical protein LOC686006 (10.9 kD) (LOC686006) mRNA.
<a href="#">LOC686032</a>	<a href="#">LOC686032.aSep08</a>	<a href="#">686032</a>	36728	562		68	hypothetical protein LOC686032 (7.7 kD) (LOC686032) mRNA.
<a href="#">LOC686059</a>	<a href="#">LOC686059.aSep08</a>	<a href="#">686059</a>	8029	913		304	similar to ryanodine receptor 1 (skeletal) (LOC686059) mRNA.
<a href="#">LOC686083</a>	<a href="#">LOC686083.aSep08</a>	<a href="#">686083</a>	11326	376		77	similar to transducer of regulated CREB protein 3 (LOC686083) mRNA.
<a href="#">LOC686084</a>	<a href="#">LOC686084.aSep08</a>	<a href="#">686084</a>	15164	710		236	similar to CG32580-PA (LOC686084) mRNA.
<a href="#">LOC686088</a>	<a href="#">LOC686088.aSep08</a>	<a href="#">686088</a>	18269	623	2	207	fshd region gene 1 (LOC686088) alternative variant aSep08, mRNA.
<a href="#">LOC686088</a>	<a href="#">LOC686088.aSep08</a>	<a href="#">686103</a>	18269	623	2	207	fshd region gene 1 (LOC686088) alternative variant aSep08, mRNA.
<a href="#">LOC686088</a>	<a href="#">LOC686088.bSep08</a>	<a href="#">686088</a>	7341	600	1	191	fshd region gene 1 (LOC686088) alternative variant bSep08, mRNA.
<a href="#">LOC686088</a>	<a href="#">LOC686088.bSep08</a>	<a href="#">686103</a>	7341	600	1	191	fshd region gene 1 (LOC686088) alternative variant bSep08, mRNA.
<a href="#">LOC686091</a>	<a href="#">LOC686091.aSep08</a>	<a href="#">686091</a>	21354	1493		464	putative protein, with 5 coiled coil domains, of eukaryotic origin (LOC686091) mRNA.
<a href="#">LOC686120</a>	<a href="#">LOC686120.aSep08</a>	<a href="#">686120</a>	965	455	1	111	hypothetical protein LOC686120 (13.0 kD) (LOC686120) alternative variant aSep08, mRNA.
<a href="#">LOC686120</a>	<a href="#">LOC686120.bSep08</a>	<a href="#">686120</a>	1005	378		104	hypothetical protein LOC686120 (10.4 kD) (LOC686120) alternative variant bSep08, mRNA.
<a href="#">LOC686123</a>	<a href="#">LOC686123.aSep08</a>	<a href="#">686123</a>	2738	519		103	putative protein of mammalian origin (LOC686123) mRNA.
<a href="#">LOC686139</a>	<a href="#">LOC686139.aSep08</a>	<a href="#">686139</a>	2729	1159	6	321	PGAP1-like and alpha/beta hydrolase fold-1 (LOC686139) alternative variant aSep08, mRNA.
<a href="#">LOC686139</a>	<a href="#">LOC686139.bSep08</a>	<a href="#">686139</a>	1697	663	3	157	putative protein of ancient origin (LOC686139) alternative variant bSep08, mRNA.
<a href="#">LOC686139</a>	<a href="#">LOC686139.dSep08</a>	<a href="#">686139</a>	2784	994	5	127	putative protein of ancient origin (13.5 kD) (LOC686139) alternative variant dSep08, mRNA.
<a href="#">LOC686139</a>	<a href="#">LOC686139.eSep08</a>	<a href="#">686139</a>	2003	1424	3	39	putative protein (LOC686139) alternative variant eSep08, mRNA.
<a href="#">LOC686141</a>	<a href="#">LOC686141.aSep08</a>	<a href="#">686141</a>	5874	1115		204	similar to Spetex-2C protein (23.9 kD) (LOC686141) mRNA.
<a href="#">LOC686149</a>	<a href="#">LOC686149.aSep08</a>	<a href="#">686149</a>	22960	801		190	similar to Actin, aortic smooth muscle (Alpha-actin-2) (LOC686149) alternative variant aSep08, mRNA.
<a href="#">LOC686149</a>	<a href="#">LOC686149.bSep08</a>	<a href="#">686149</a>	4727	462		85	similar to Actin, aortic smooth muscle (Alpha-actin-2) (LOC686149) alternative variant bSep08, mRNA.
<a href="#">LOC686841</a>	<a href="#">LOC686841.bSep08</a>	<a href="#">686841</a>	458	271	2	67	similar to Protein EAN57 (LOC686841) alternative variant bSep08, mRNA.

<a href="#">LOC686841</a>	<a href="#">LOC686841.cSep08</a>	<a href="#">686841</a>	46159	1740	2	58	similar to Protein EAN57 (LOC686841) alternative variant cSep08, mRNA.
<a href="#">LOC688400</a>	<a href="#">LOC688400.aSep08</a>	<a href="#">688400</a>	272916	780		155	similar to EGF-like repeats and discoidin I-like domains-containing protein 3 (LOC688400) mRNA.
<a href="#">LOC688433</a>	<a href="#">LOC688433.aSep08</a>	<a href="#">688433</a>	4273	304		44	putative protein of vertebrate origin (LOC688433) mRNA.
<a href="#">LOC688435</a>	<a href="#">LOC688435.aSep08</a>	<a href="#">688435</a>	30620	730		79	similar to 40S ribosomal protein S2 (8.7 kD) (LOC688435) mRNA.
<a href="#">LOC688442</a>	<a href="#">LOC688442.aSep08</a>	<a href="#">688442</a>	22561	688		228	similar to limkain b1 (LOC688442) mRNA.
<a href="#">LOC688448</a>	<a href="#">LOC688448.aSep08</a>	<a href="#">688448</a>	8191	848		203	glycine n-acyltransferase-like protein 3 (LOC688448) mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.aSep08</a>	<a href="#">688452</a>	9982	728	5	153	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant aSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.aSep08</a>	<a href="#">690129</a>	9982	728	5	153	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant aSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.bSep08</a>	<a href="#">688452</a>	8163	693	3	124	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant bSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.bSep08</a>	<a href="#">690129</a>	8163	693	3	124	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant bSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.cSep08</a>	<a href="#">688452</a>	8553	1890	5	96	hypothetical protein LOC688452 and hypothetical protein LOC690129 (10.4 kD) (LOC688452) alternative variant cSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.cSep08</a>	<a href="#">690129</a>	8553	1890	5	96	hypothetical protein LOC688452 and hypothetical protein LOC690129 (10.4 kD) (LOC688452) alternative variant cSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.dSep08</a>	<a href="#">688452</a>	17918	854	6	94	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant dSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.dSep08</a>	<a href="#">690129</a>	17918	854	6	94	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant dSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.eSep08</a>	<a href="#">688452</a>	1610	737	2	68	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant eSep08, mRNA.
<a href="#">LOC688452</a>	<a href="#">LOC688452.eSep08</a>	<a href="#">690129</a>	1610	737	2	68	hypothetical protein LOC688452 and hypothetical protein LOC690129 (LOC688452) alternative variant eSep08, mRNA.
<a href="#">LOC688463</a>	<a href="#">LOC688463.aSep08</a>	<a href="#">688463</a>	7025	706		137	hypothetical protein LOC688463 (LOC688463) mRNA.
<a href="#">LOC688464</a>	<a href="#">LOC688464.aSep08</a>	<a href="#">688464</a>	13195	764	1	130	hypothetical protein LOC688464 (LOC688464) alternative variant aSep08, mRNA.
<a href="#">LOC688464</a>	<a href="#">LOC688464.bSep08</a>	<a href="#">688464</a>	12674	770	1	104	hypothetical protein LOC688464 (11.5 kD) (LOC688464) alternative variant bSep08, mRNA.

<a href="#">LOC688466</a>	<a href="#">LOC688466.aSep08</a>	<a href="#">688466</a>	1740	949	2	249	similar to Ubiquitin carboxyl-terminal hydrolase 21 (Ubiquitin thiolesterase 21) (Ubiquitin-specific processing protease 21) (Deubiquitinating enzyme 21) (LOC688466) alternative variant aSep08, mRNA.
<a href="#">LOC688466</a>	<a href="#">LOC688466.bSep08</a>	<a href="#">688466</a>	49851	558	3	185	similar to Ubiquitin carboxyl-terminal hydrolase 21 (Ubiquitin thiolesterase 21) (Ubiquitin-specific processing protease 21) (Deubiquitinating enzyme 21) (LOC688466) alternative variant bSep08, mRNA.
<a href="#">LOC688466</a>	<a href="#">LOC688466.dSep08</a>	<a href="#">688466</a>	1145	846	3	111	similar to Ubiquitin carboxyl-terminal hydrolase 21 (Ubiquitin thiolesterase 21) (Ubiquitin-specific processing protease 21) (Deubiquitinating enzyme 21) (12.5 kD) (LOC688466) alternative variant dSep08, mRNA.
<a href="#">LOC688479</a>	<a href="#">LOC688479.aSep08</a>	<a href="#">688479</a>	5362	631	1	209	region containing hypothetical protein LOC501396; similar to RIKEN cDNA 1700001E04 (LOC688479) alternative variant aSep08, mRNA.
<a href="#">LOC688479</a>	<a href="#">LOC688479.bSep08</a>	<a href="#">688479</a>	21321	858	1	123	region containing hypothetical protein LOC501396; similar to RIKEN cDNA 1700001E04 (LOC688479) alternative variant bSep08, mRNA.
<a href="#">LOC688488</a>	<a href="#">LOC688488.aSep08</a>	<a href="#">688488</a>	2818	374		86	hypothetical protein LOC688488 (LOC688488) mRNA.
<a href="#">LOC688495</a>	<a href="#">LOC688495.aSep08</a>	<a href="#">688495</a>	3991	1367		108	hypothetical protein LOC688495 (LOC688495) mRNA.
<a href="#">LOC688499</a>	<a href="#">LOC688499.aSep08</a>	<a href="#">688499</a>	1397	390		129	hypothetical protein LOC688499 (LOC688499) mRNA.
<a href="#">LOC688502</a>	<a href="#">LOC688502.aSep08</a>	<a href="#">688502</a>	14827	1297		360	similar to Protein arginine N-methyltransferase 4 (Heterogeneous nuclear ribonucleoprotein methyltransferase-like protein 4) (LOC688502) mRNA.
<a href="#">LOC688504</a>	<a href="#">LOC688504.aSep08</a>	<a href="#">688504</a>	2126	586		84	similar to 60S ribosomal protein L28 (9.3 kD) (LOC688504) mRNA.
<a href="#">LOC688514</a>	<a href="#">LOC688514.aSep08</a>	<a href="#">688514</a>	1139	426	1	82	similar to major urinary protein 4 (LOC688514) alternative variant aSep08, mRNA.
<a href="#">LOC688514</a>	<a href="#">LOC688514.bSep08</a>	<a href="#">688514</a>	1138	379	1	67	similar to major urinary protein 4 (LOC688514) alternative variant bSep08, mRNA.
<a href="#">LOC688534</a>	<a href="#">LOC688534.bSep08</a>	<a href="#">688534</a>	371	284	2	11	similar to Mitochondrial import inner membrane translocase subunit Tim9 B (TIMM10B) (Tim10b) (Fracture callus protein 1) (FxC1) (LOC688534) alternative variant bSep08, mRNA.
<a href="#">LOC688535</a>	<a href="#">LOC688535.aSep08</a>	<a href="#">688535</a>	4099	954		58	hypothetical protein LOC688535 (LOC688535) mRNA.
<a href="#">LOC688536</a>	<a href="#">LOC688536.aSep08</a>	<a href="#">688536</a>	9110	848		203	glycine n-acyltransferase-like protein 3 (LOC688536) mRNA.
<a href="#">LOC688542</a>	<a href="#">LOC688542.aSep08</a>	<a href="#">688542</a>	1118	428		67	similar to carboxylesterase 5 (8.0 kD) (LOC688542) mRNA.
<a href="#">LOC688548</a>	<a href="#">LOC688548.bSep08</a>	<a href="#">688548</a>	10836	1206	2	181	hypothetical protein LOC688548 (LOC688548) alternative variant bSep08, mRNA.
<a href="#">LOC688553</a>	<a href="#">LOC688553.aSep08</a>	<a href="#">688553</a>	19158	1503	6	292	hypothetical protein LOC688553 (32.8 kD) (LOC688553) alternative variant aSep08, mRNA.
<a href="#">LOC688553</a>	<a href="#">LOC688553.bSep08</a>	<a href="#">688553</a>	1574	326	1	49	hypothetical protein LOC688553 (LOC688553) alternative variant bSep08, mRNA.
<a href="#">LOC688568</a>	<a href="#">LOC688568.aSep08</a>	<a href="#">688568</a>	16894	495		74	similar to similar to RIKEN cDNA 1700001E04 (8.3 kD) (LOC688568) mRNA.

<a href="#">LOC688579</a>	<a href="#">LOC688579.aSep08</a>	<a href="#">688579</a>	21543	737	2	111	hypothetical protein LOC688579 (LOC688579) alternative variant aSep08, mRNA.
<a href="#">LOC688579</a>	<a href="#">LOC688579.bSep08</a>	<a href="#">688579</a>	127192	393	2	104	hypothetical protein LOC688579 (LOC688579) alternative variant bSep08, mRNA.
<a href="#">LOC688582</a>	<a href="#">LOC688582.aSep08</a>	<a href="#">688582</a>	2418	344		114	similar to hemicentin 1 (LOC688582) mRNA.
<a href="#">LOC688585</a>	<a href="#">LOC688585.aSep08</a>	<a href="#">688585</a>	139719	454		31	hypothetical protein LOC688585 (3.7 kD) (LOC688585) mRNA.
<a href="#">LOC688587</a>	<a href="#">LOC688587.aSep08</a>	<a href="#">688587</a>	1776	429		97	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC688587) mRNA.
<a href="#">LOC688588</a>	<a href="#">LOC688588.aSep08</a>	<a href="#">688588</a>	17151	831		157	similar to MIC2 like 1 (18.1 kD) (LOC688588) mRNA.
<a href="#">LOC688599</a>	<a href="#">LOC688599.aSep08</a>	<a href="#">688599</a>	1772	403		133	similar to hemicentin 1 (LOC688599) mRNA.
<a href="#">LOC688603</a>	<a href="#">LOC688603.aSep08</a>	<a href="#">688603</a>	1777	430	3	97	similar to RIKEN cDNA 5031410I06 (LOC688603) alternative variant aSep08, mRNA.
<a href="#">LOC688631</a>	<a href="#">LOC688631.aSep08</a>	<a href="#">688631</a>	317776	773		257	similar to DNA repair endonuclease XPF (DNA excision repair protein ERCC-4) (LOC688631) mRNA.
<a href="#">LOC688634</a>	<a href="#">LOC688634.aSep08</a>	<a href="#">688634</a>	117077	1297		319	similar to Histone acetyltransferase MYST4 (MYST protein 4) (MOZ, YBF2/SAS3, SAS2 and TIP60 protein 4) (Querkopf protein) and hypothetical protein LOC688653 (LOC688634) mRNA.
<a href="#">LOC688634</a>	<a href="#">LOC688634.aSep08</a>	<a href="#">688653</a>	117077	1297		319	similar to Histone acetyltransferase MYST4 (MYST protein 4) (MOZ, YBF2/SAS3, SAS2 and TIP60 protein 4) (Querkopf protein) and hypothetical protein LOC688653 (LOC688634) mRNA.
<a href="#">LOC688637</a>	<a href="#">LOC688637.aSep08</a>	<a href="#">688637</a>	28092	3116	23	922	similar to WD repeat domain 36 (LOC688637) alternative variant aSep08, mRNA.
<a href="#">LOC688637</a>	<a href="#">LOC688637.bSep08</a>	<a href="#">688637</a>	11679	716	8	238	similar to WD repeat domain 36 (LOC688637) alternative variant bSep08, mRNA.
<a href="#">LOC688662</a>	<a href="#">LOC688662.aSep08</a>	<a href="#">688662</a>	17698	412	2	92	similar to Reticulocalbin-1 precursor (LOC688662) alternative variant aSep08, mRNA.
<a href="#">LOC688662</a>	<a href="#">LOC688662.bSep08</a>	<a href="#">688662</a>	10605	427	2	39	similar to Reticulocalbin-1 precursor (LOC688662) alternative variant bSep08, mRNA.
<a href="#">LOC688673</a>	<a href="#">LOC688673.bSep08</a>	<a href="#">688673</a>	5125	684	6	97	similar to Protein O-mannosyl-transferase 2 (Dolichyl-phosphate-mannose--protein mannosyltransferase 2) (LOC688673) alternative variant bSep08, mRNA.
<a href="#">LOC688673</a>	<a href="#">LOC688673.cSep08</a>	<a href="#">688673</a>	6838	342	3	18	similar to Protein O-mannosyl-transferase 2 (Dolichyl-phosphate-mannose--protein mannosyltransferase 2) (LOC688673) alternative variant cSep08, mRNA.
<a href="#">LOC688676</a>	<a href="#">LOC688676.aSep08</a>	<a href="#">688676</a>	593	371		30	hypothetical protein LOC688676 (LOC688676) mRNA.
<a href="#">LOC688681</a>	<a href="#">LOC688681.aSep08</a>	<a href="#">688681</a>	949	868	2	189	similar to transforming growth factor, beta receptor III (betaglycan, 300kDa) (20.6 kD) (LOC688681) alternative variant aSep08, mRNA.
<a href="#">LOC688681</a>	<a href="#">LOC688681.bSep08</a>	<a href="#">688681</a>	1486	445	3	148	similar to transforming growth factor, beta receptor III (betaglycan, 300kDa) (LOC688681) alternative variant bSep08, mRNA.
<a href="#">LOC688682</a>	<a href="#">LOC688682.bSep08</a>	<a href="#">688682</a>	448	346	2	73	similar to small nuclear ribonucleoparticle-associated protein (LOC688682) alternative variant bSep08, mRNA.

<a href="#">LOC688686</a>	<a href="#">LOC688686.aSep08</a>	<a href="#">688686</a>	5120	621		77	hypothetical protein LOC688686 (9.1 kD) (LOC688686) mRNA.
<a href="#">LOC688749</a>	<a href="#">LOC688749.aSep08</a>	<a href="#">688749</a>	36414	690		229	similar to Tubby-related protein 3 (Tubby-like protein 3) (LOC688749) mRNA.
<a href="#">LOC688773</a>	<a href="#">LOC688773.bSep08</a>	<a href="#">688773</a>	12834	1520	3	104	similar to nidogen 2 (11.6 kD) (LOC688773) alternative variant bSep08, mRNA.
<a href="#">LOC688773</a>	<a href="#">LOC688773.cSep08</a>	<a href="#">688773</a>	11898	773	3	94	similar to nidogen 2 (LOC688773) alternative variant cSep08, mRNA.
<a href="#">LOC688773</a>	<a href="#">LOC688773.dSep08</a>	<a href="#">688773</a>	1479	750	2	93	similar to nidogen 2 (10.4 kD) (LOC688773) alternative variant dSep08, mRNA.
<a href="#">LOC688778</a>	<a href="#">LOC688778.aSep08</a>	<a href="#">688778</a>	1385	676	2	225	similar to fatty aldehyde dehydrogenase-like (LOC688778) alternative variant aSep08, mRNA.
<a href="#">LOC688778</a>	<a href="#">LOC688778.bSep08</a>	<a href="#">688778</a>	916	382	1	73	similar to fatty aldehyde dehydrogenase-like (LOC688778) alternative variant bSep08, mRNA.
<a href="#">LOC688801</a>	<a href="#">LOC688801.aSep08</a>	<a href="#">688801</a>	6849	636		89	hypothetical protein LOC688801 (LOC688801) mRNA.
<a href="#">LOC688807</a>	<a href="#">LOC688807.aSep08</a>	<a href="#">688807</a>	3707	701		137	hypothetical protein LOC688807 (14.4 kD) (LOC688807) mRNA.
<a href="#">LOC688808</a>	<a href="#">LOC688808.aSep08</a>	<a href="#">688808</a>	27464	532	3	19	similar to RIKEN cDNA 5031410I06 (2.1 kD) (LOC688808) alternative variant aSep08, mRNA.
<a href="#">LOC688808</a>	<a href="#">LOC688808.bSep08</a>	<a href="#">688808</a>	4423	380	2	70	similar to RIKEN cDNA 5031410I06 (LOC688808) alternative variant bSep08, mRNA.
<a href="#">LOC688832</a>	<a href="#">LOC688832.aSep08</a>	<a href="#">688832</a>	1699	590		71	hypothetical protein LOC688832 (7.8 kD) (LOC688832) mRNA.
<a href="#">LOC688833</a>	<a href="#">LOC688833.aSep08</a>	<a href="#">688833</a>	768	234		60	hypothetical protein LOC688833 (LOC688833) mRNA.
<a href="#">LOC688864</a>	<a href="#">LOC688864.aSep08</a>	<a href="#">688864</a>	9796	655		84	similar to transmembrane protein 61 (9.4 kD) (LOC688864) mRNA.
<a href="#">LOC688869</a>	<a href="#">LOC688869.bSep08</a>	<a href="#">688869</a>	7300	437	3	86	similar to cytochrome c oxidase, subunit VIb polypeptide 1 (10.1 kD) (LOC688869) alternative variant bSep08, complete mRNA.
<a href="#">LOC688872</a>	<a href="#">LOC688872.bSep08</a>	<a href="#">688872</a>	14599	1390	6	296	similar to fetal Alzheimer antigen isoform 2 (LOC688872) alternative variant bSep08, mRNA.
<a href="#">LOC688872</a>	<a href="#">LOC688872.cSep08</a>	<a href="#">688872</a>	11511	1797	6	227	similar to fetal Alzheimer antigen isoform 2 (LOC688872) alternative variant cSep08, mRNA.
<a href="#">LOC688872</a>	<a href="#">LOC688872.dSep08</a>	<a href="#">688872</a>	1449	313	2	104	similar to fetal Alzheimer antigen isoform 2 (LOC688872) alternative variant dSep08, mRNA.
<a href="#">LOC688877</a>	<a href="#">LOC688877.aSep08</a>	<a href="#">688877</a>	12946	662		121	similar to nidogen 2 (13.0 kD) (LOC688877) mRNA.
<a href="#">LOC688893</a>	<a href="#">LOC688893.aSep08</a>	<a href="#">688893</a>	1373	388		54	hypothetical protein LOC688893 (LOC688893) mRNA.
<a href="#">LOC688903</a>	<a href="#">LOC688903.aSep08</a>	<a href="#">688903</a>	11020	767	2	113	similar to Nucleoside diphosphate kinase homolog 5 (NDK-H 5) (NDP kinase homolog 5) (nm23-M5) (LOC688903) alternative variant aSep08, mRNA.
<a href="#">LOC688903</a>	<a href="#">LOC688903.bSep08</a>	<a href="#">688903</a>	16761	833	1	46	similar to Nucleoside diphosphate kinase homolog 5 (NDK-H 5) (NDP kinase homolog 5) (nm23-M5) (LOC688903) alternative variant bSep08, mRNA.
<a href="#">LOC688905</a>	<a href="#">LOC688905.aSep08</a>	<a href="#">688905</a>	2655	527		75	similar to metaxin 3 (LOC688905) mRNA.
<a href="#">LOC688914</a>	<a href="#">LOC688914.aSep08</a>	<a href="#">688914</a>	72487	402		122	hypothetical protein LOC688914 (LOC688914) mRNA.

<a href="#">LOC688915</a>	<a href="#">LOC688915.aSep08</a>	<a href="#">688915</a>	26587	1008	1	310	similar to cardiomyopathy associated 5 (LOC688915) alternative variant aSep08, mRNA.
<a href="#">LOC688915</a>	<a href="#">LOC688915.bSep08</a>	<a href="#">688915</a>	25484	1529	1	221	similar to cardiomyopathy associated 5 (24.6 kD) (LOC688915) alternative variant bSep08, mRNA.
<a href="#">LOC688916</a>	<a href="#">LOC688916.aSep08</a>	<a href="#">688916</a>	69395	582		108	hypothetical protein LOC688916 (LOC688916) mRNA.
<a href="#">LOC688968</a>	<a href="#">LOC688968.aSep08</a>	<a href="#">688968</a>	14748	1322	2	348	similar to SGT1 protein homolog (Ecdysoneless homolog) (LOC688968) alternative variant aSep08, mRNA.
<a href="#">LOC688968</a>	<a href="#">LOC688968.bSep08</a>	<a href="#">688968</a>	11167	737	1	154	similar to SGT1 protein homolog (Ecdysoneless homolog) (LOC688968) alternative variant bSep08, mRNA.
<a href="#">LOC688972</a>	<a href="#">LOC688972.aSep08</a>	<a href="#">688972</a>	17859	1257	7	132	similar to Glycophorin (14.0 kD) (LOC688972) alternative variant aSep08, complete mRNA.
<a href="#">LOC688990</a>	<a href="#">LOC688990.aSep08</a>	<a href="#">688990</a>	1777	812		78	hypothetical protein LOC688990 (LOC688990) mRNA.
<a href="#">LOC689010</a>	<a href="#">LOC689010.aSep08</a>	<a href="#">689010</a>	52114	574		45	hypothetical protein LOC689010 (5.3 kD) (LOC689010) mRNA.
<a href="#">LOC689012</a>	<a href="#">LOC689012.aSep08</a>	<a href="#">689012</a>	12343	425		34	hypothetical protein LOC689012 (LOC689012) mRNA.
<a href="#">LOC689019</a>	<a href="#">LOC689019.aSep08</a>	<a href="#">689019</a>	1842	407		99	hypothetical protein LOC689019 (LOC689019) mRNA.
<a href="#">LOC689027</a>	<a href="#">LOC689027.bSep08</a>	<a href="#">689027</a>	15986	746	8	248	similar to Probable polypeptide N-acetylgalactosaminyltransferase 8 (Protein-UDP acetylgalactosaminyltransferase 8) (UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 8) (Polypeptide GalNAc transferase 8) (GalNAc-T8) (pp-GaNTase 8) (LOC689027) alternative variant bSep08, mRNA.
<a href="#">LOC689035</a>	<a href="#">LOC689035.aSep08</a>	<a href="#">689035</a>	13730	858		285	similar to CG5946-PB, isoform B (LOC689035) mRNA.
<a href="#">LOC689043</a>	<a href="#">LOC689043.bSep08</a>	<a href="#">689043</a>	244814	1781	1	67	similar to TFAA4 protein (LOC689043) alternative variant bSep08, mRNA.
<a href="#">LOC689044</a>	<a href="#">LOC689044.bSep08</a>	<a href="#">689044</a>	78187	388	1	76	hypothetical protein LOC689044 (LOC689044) alternative variant bSep08, mRNA.
<a href="#">LOC689074</a>	<a href="#">LOC689074.aSep08</a>	<a href="#">689074</a>	18430	1057		351	similar to Protein KIAA1543 (LOC689074) mRNA.
<a href="#">LOC689084</a>	<a href="#">LOC689084.aSep08</a>	<a href="#">689084</a>	3140	701		144	hypothetical protein LOC689084 (17.2 kD) (LOC689084) mRNA.
<a href="#">LOC689088</a>	<a href="#">LOC689088.aSep08</a>	<a href="#">689088</a>	5752	1510		158	predicted gene (LOC689088) mRNA.
<a href="#">LOC689095</a>	<a href="#">LOC689095.aSep08</a>	<a href="#">689095</a>	14266	290	2	96	similar to stearoyl-Coenzyme A desaturase 1 (LOC689095) alternative variant aSep08, mRNA.
<a href="#">LOC689103</a>	<a href="#">LOC689103.aSep08</a>	<a href="#">689103</a>	553	441		120	similar to vitelliform macular dystrophy 2-like 2 (LOC689103) mRNA.
<a href="#">LOC689116</a>	<a href="#">LOC689116.aSep08</a>	<a href="#">689116</a>	3798	1023	3	254	similar to nuclear cap binding protein subunit 2 (LOC689116) alternative variant aSep08, mRNA.
<a href="#">LOC689116</a>	<a href="#">LOC689116.bSep08</a>	<a href="#">689116</a>	3591	819	3	226	similar to nuclear cap binding protein subunit 2 (LOC689116) alternative variant bSep08, mRNA.
<a href="#">LOC689116</a>	<a href="#">LOC689116.cSep08</a>	<a href="#">689116</a>	29057	647	5	181	similar to nuclear cap binding protein subunit 2 (LOC689116) alternative variant cSep08, mRNA.
<a href="#">LOC689116</a>	<a href="#">LOC689116.eSep08</a>	<a href="#">689116</a>	8137	805	5	156	similar to nuclear cap binding protein subunit 2 (18.0 kD) (LOC689116) alternative variant eSep08, mRNA.
<a href="#">LOC689116</a>	<a href="#">LOC689116.fSep08</a>	<a href="#">689116</a>	7830	425	3	110	similar to nuclear cap binding protein subunit 2 (LOC689116) alternative variant fSep08, mRNA.



<a href="#">LOC689116</a>	<a href="#">LOC689116.gSep08</a>	<a href="#">689116</a>	838	738	2	40	similar to nuclear cap binding protein subunit 2 (LOC689116) alternative variant gSep08, mRNA.
<a href="#">LOC689124</a>	<a href="#">LOC689124.aSep08</a>	<a href="#">689124</a>	617	385		10	similar to RIKEN cDNA 5031410I06 (1.1 kD) (LOC689124) mRNA.
<a href="#">LOC689133</a>	<a href="#">LOC689133.aSep08</a>	<a href="#">689133</a>	1763	404		81	similar to Extracellular peptidase inhibitor precursor (Protein WDNM1) (LOC689133) mRNA.
<a href="#">LOC689134</a>	<a href="#">LOC689134.aSep08</a>	<a href="#">689134</a>	6273	368	4	75	similar to Protein transport protein SEC61 gamma subunit (LOC689134) alternative variant aSep08, mRNA.
<a href="#">LOC689134</a>	<a href="#">LOC689134.bSep08</a>	<a href="#">689134</a>	6409	477	4	68	similar to Protein transport protein SEC61 gamma subunit (7.7 kD) (LOC689134) alternative variant bSep08, complete mRNA.
<a href="#">LOC689147</a>	<a href="#">LOC689147.aSep08</a>	<a href="#">689147</a>	25440	877	2	131	hypothetical protein LOC689147 (14.9 kD) (LOC689147) alternative variant aSep08, mRNA.
<a href="#">LOC689147</a>	<a href="#">LOC689147.bSep08</a>	<a href="#">689147</a>	19501	713	1	108	hypothetical protein LOC689147 (LOC689147) alternative variant bSep08, mRNA.
<a href="#">LOC689168</a>	<a href="#">LOC689168.aSep08</a>	<a href="#">689168</a>	5864	626	1	84	similar to Cystatin S precursor (LM protein) (9.4 kD) (LOC689168) alternative variant aSep08, mRNA.
<a href="#">LOC689168</a>	<a href="#">LOC689168.bSep08</a>	<a href="#">689168</a>	12181	648	1	82	similar to Cystatin S precursor (LM protein) (LOC689168) alternative variant bSep08, mRNA.
<a href="#">LOC689176</a>	<a href="#">LOC689176.aSep08</a>	<a href="#">689176</a>	16347	784		260	similar to transmembrane protein 64 (LOC689176) mRNA.
<a href="#">LOC689178</a>	<a href="#">LOC689178.aSep08</a>	<a href="#">689178</a>	2767	473		98	hypothetical protein LOC689178 (LOC689178) mRNA.
<a href="#">LOC689212</a>	<a href="#">LOC689212.aSep08</a>	<a href="#">689212</a>	1480	435		65	similar to Cystatin S precursor (LM protein) (LOC689212) mRNA.
<a href="#">LOC689226</a>	<a href="#">LOC689226.aSep08</a>	<a href="#">689226</a>	60141	1690	1	235	similar to ubiquitin-conjugating enzyme E2R 2 (LOC689226) alternative variant aSep08, mRNA.
<a href="#">LOC689240</a>	<a href="#">LOC689240.aSep08</a>	<a href="#">689240</a>	1070	646		90	neurobeachin like 1 (LOC689240) mRNA.
<a href="#">LOC689252</a>	<a href="#">LOC689252.aSep08</a>	<a href="#">689252</a>	86410	817		119	hypothetical protein LOC689252 (12.8 kD) (LOC689252) mRNA.
<a href="#">LOC689275</a>	<a href="#">LOC689275.aSep08</a>	<a href="#">689275</a>	91182	1943		69	hypothetical protein LOC689275 (7.8 kD) (LOC689275) mRNA.
<a href="#">LOC689276</a>	<a href="#">LOC689276.aSep08</a>	<a href="#">689276</a>	1982	388		45	hypothetical protein LOC689276 (LOC689276) mRNA.
<a href="#">LOC689288</a>	<a href="#">LOC689288.aSep08</a>	<a href="#">689288</a>	5310	1784	2	419	similar to copine II (LOC689288) alternative variant aSep08, mRNA.
<a href="#">LOC689288</a>	<a href="#">LOC689288.bSep08</a>	<a href="#">689288</a>	8359	983	3	87	similar to copine II (LOC689288) alternative variant bSep08, mRNA.
<a href="#">LOC689289</a>	<a href="#">LOC689289.bSep08</a>	<a href="#">689289</a>	2289	570	3	143	similar to Y73F8A.5 (LOC689289) alternative variant bSep08, mRNA.
<a href="#">LOC689294</a>	<a href="#">LOC689294.aSep08</a>	<a href="#">689294</a>	28537	312		101	similar to ADAM 22 precursor (A disintegrin and metalloproteinase domain 22) (LOC689294) mRNA.
<a href="#">LOC689303</a>	<a href="#">LOC689303.aSep08</a>	<a href="#">689303</a>	14692	327		94	similar to vitamin A-deficient testicular protein 11-like (LOC689303) mRNA.
<a href="#">LOC689314</a>	<a href="#">LOC689314.aSep08</a>	<a href="#">689314</a>	4146	2125	11	314	similar to mitogen-activated protein kinase 11 (LOC689314) alternative variant aSep08, mRNA.
<a href="#">LOC689316</a>	<a href="#">LOC689316.aSep08</a>	<a href="#">689316</a>	3076	884		196	hypothetical protein LOC689316 (21.8 kD) (LOC689316) mRNA.
<a href="#">LOC689329</a>	<a href="#">LOC689329.aSep08</a>	<a href="#">689329</a>	7278	651		103	hypothetical protein LOC689329 (12.0 kD) (LOC689329) mRNA.

<a href="#">LOC689340</a>	<a href="#">LOC689340.aSep08</a>	<a href="#">689340</a>	36467	582		39	similar to 40S ribosomal protein S2 (4.3 kD) (LOC689340) mRNA.
<a href="#">LOC689372</a>	<a href="#">LOC689372.aSep08</a>	<a href="#">689372</a>	1308	465		68	similar to spermatogenesis associated glutamate (E)-rich protein 4d (8.3 kD) (LOC689372) mRNA.
<a href="#">LOC689396</a>	<a href="#">LOC689396.aSep08</a>	<a href="#">689396</a>	8016	769	2	234	similar to RNA polymerase II elongation factor ELL2 (LOC689396) alternative variant aSep08, mRNA.
<a href="#">LOC689399</a>	<a href="#">LOC689399.aSep08</a>	<a href="#">689399</a>	4938	1600		86	hypothetical protein LOC689399 (9.8 kD) (LOC689399) mRNA.
<a href="#">LOC689418</a>	<a href="#">LOC689418.aSep08</a>	<a href="#">689418</a>	2932	400		133	hypothetical protein LOC689418 (LOC689418) mRNA.
<a href="#">LOC689443</a>	<a href="#">LOC689443.aSep08</a>	<a href="#">689443</a>	197243	2307		250	similar to TBC1 domain family member 22A (LOC689443) mRNA.
<a href="#">LOC689445</a>	<a href="#">LOC689445.aSep08</a>	<a href="#">689445</a>	6483	341	1	113	hypothetical protein LOC689445 (LOC689445) alternative variant aSep08, mRNA.
<a href="#">LOC689445</a>	<a href="#">LOC689445.bSep08</a>	<a href="#">689445</a>	4247	499	2	61	hypothetical protein LOC689445 (7.0 kD) (LOC689445) alternative variant bSep08, mRNA.
<a href="#">LOC689464</a>	<a href="#">LOC689464.aSep08</a>	<a href="#">689464</a>	5442	1014	3	76	similar to 40S ribosomal protein SA (p40) (34/67 kDa laminin receptor) (LOC689464) alternative variant aSep08, mRNA.
<a href="#">LOC689479</a>	<a href="#">LOC689479.aSep08</a>	<a href="#">689479</a>	8359	802	3	206	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC689479) alternative variant aSep08, mRNA.
<a href="#">LOC689479</a>	<a href="#">LOC689479.bSep08</a>	<a href="#">689479</a>	2760	732	1	122	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC689479) alternative variant bSep08, mRNA.
<a href="#">LOC689486</a>	<a href="#">LOC689486.aSep08</a>	<a href="#">689486</a>	31248	377		93	hypothetical protein LOC689486 (LOC689486) mRNA.
<a href="#">LOC689541</a>	<a href="#">LOC689541.aSep08</a>	<a href="#">689541</a>	2007	533		93	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC689541) mRNA.
<a href="#">LOC689570</a>	<a href="#">LOC689570.bSep08</a>	<a href="#">689570</a>	5536	448	2	72	similar to Kazal type serine protease inhibitor 4 (LOC689570) alternative variant bSep08, mRNA.
<a href="#">LOC689574</a>	<a href="#">LOC689574.aSep08</a>	<a href="#">689574</a>	9955	854	3	115	hypothetical protein LOC689574 (LOC689574) alternative variant aSep08, mRNA.
<a href="#">LOC689574</a>	<a href="#">LOC689574.bSep08</a>	<a href="#">689574</a>	7152	289	3	96	hypothetical protein LOC689574 (LOC689574) alternative variant bSep08, mRNA.
<a href="#">LOC689574</a>	<a href="#">LOC689574.dSep08</a>	<a href="#">689574</a>	1201	547	2	50	hypothetical protein LOC689574 (LOC689574) alternative variant dSep08, mRNA.
<a href="#">LOC689589</a>	<a href="#">LOC689589.aSep08</a>	<a href="#">689589</a>	20840	883		87	hypothetical protein LOC689589 (10.0 kD) (LOC689589) mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.aSep08</a>	<a href="#">689593</a>	7696	1795	3	499	CRA g (LOC689593) alternative variant aSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.bSep08</a>	<a href="#">689593</a>	7536	2793	8	335	DnaJ homolog subfamily B member (LOC689593) alternative variant bSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.cSep08</a>	<a href="#">689593</a>	8212	1879	10	277	DnaJ homolog subfamily B member (30.7 kD) (LOC689593) alternative variant cSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.eSep08</a>	<a href="#">689593</a>	3673	864	5	223	DnaJ homolog subfamily B member (LOC689593) alternative variant eSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.fSep08</a>	<a href="#">689593</a>	4454	937	5	169	DnaJ homolog subfamily B member (19.1 kD) (LOC689593) alternative variant fSep08, mRNA.

<a href="#">LOC689593</a>	<a href="#">LOC689593.gSep08</a>	<a href="#">689593</a>	4569	562	7	156	DnaJ homolog subfamily B member (LOC689593) alternative variant gSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.hSep08</a>	<a href="#">689593</a>	1078	574	2	138	CRA g (15.6 kD) (LOC689593) alternative variant hSep08, mRNA.
<a href="#">LOC689593</a>	<a href="#">LOC689593.iSep08</a>	<a href="#">689593</a>	930	733	2	86	putative protein (LOC689593) alternative variant iSep08, mRNA.
<a href="#">LOC689601</a>	<a href="#">LOC689601.aSep08</a>	<a href="#">689601</a>	22476	1003	7	230	similar to density-regulated protein (LOC689601) alternative variant aSep08, mRNA.
<a href="#">LOC689601</a>	<a href="#">LOC689601.cSep08</a>	<a href="#">689601</a>	3004	619	3	102	similar to density-regulated protein (LOC689601) alternative variant cSep08, mRNA.
<a href="#">LOC689601</a>	<a href="#">LOC689601.dSep08</a>	<a href="#">689601</a>	773	662	2		
<a href="#">LOC689617</a>	<a href="#">LOC689617.aSep08</a>	<a href="#">689617</a>	4614	424		141	similar to GTPase activating protein testicular GAP1 (LOC689617) mRNA.
<a href="#">LOC689619</a>	<a href="#">LOC689619.aSep08</a>	<a href="#">689619</a>	16450	659	2	184	similar to nuclear receptor interacting protein 2 (LOC689619) alternative variant aSep08, mRNA.
<a href="#">LOC689619</a>	<a href="#">LOC689619.bSep08</a>	<a href="#">689619</a>	7833	429	1	69	similar to nuclear receptor interacting protein 2 (LOC689619) alternative variant bSep08, mRNA.
<a href="#">LOC689636</a>	<a href="#">LOC689636.aSep08</a>	<a href="#">689636</a>	35263	393	2	86	hypothetical protein LOC689636 (LOC689636) alternative variant aSep08, mRNA.
<a href="#">LOC689636</a>	<a href="#">LOC689636.bSep08</a>	<a href="#">689636</a>	43293	597	3	81	hypothetical protein LOC689636 (LOC689636) alternative variant bSep08, mRNA.
<a href="#">LOC689663</a>	<a href="#">LOC689663.aSep08</a>	<a href="#">689663</a>	1853	473	1	157	hypothetical protein LOC689663 (LOC689663) alternative variant aSep08, mRNA.
<a href="#">LOC689663</a>	<a href="#">LOC689663.bSep08</a>	<a href="#">689663</a>	6045	373	2	80	hypothetical protein LOC689663 (LOC689663) alternative variant bSep08, mRNA.
<a href="#">LOC689696</a>	<a href="#">LOC689696.aSep08</a>	<a href="#">689696</a>	214974	401		81	similar to Receptor-type tyrosine-protein phosphatase delta precursor (Protein-tyrosine phosphatase delta) (R-PTP-delta) (LOC689696) mRNA.
<a href="#">LOC689717</a>	<a href="#">LOC689717.aSep08</a>	<a href="#">689717</a>	66828	682	3	69	similar to carboxypeptidase O (7.7 kD) (LOC689717) alternative variant aSep08, mRNA.
<a href="#">LOC689727</a>	<a href="#">LOC689727.aSep08</a>	<a href="#">689727</a>	8878	556		61	hypothetical protein LOC689727 (7.0 kD) (LOC689727) mRNA.
<a href="#">LOC689732</a>	<a href="#">LOC689732.aSep08</a>	<a href="#">689732</a>	24564	229		76	hypothetical protein LOC689732 (LOC689732) mRNA.
<a href="#">LOC689748</a>	<a href="#">LOC689748.aSep08</a>	<a href="#">689748</a>	5896	497		127	hypothetical protein LOC689748 (LOC689748) mRNA.
<a href="#">LOC689750</a>	<a href="#">LOC689750.aSep08</a>	<a href="#">689750</a>	1308	465		68	similar to spermatogenesis associated glutamate (E)-rich protein 4d (8.3 kD) (LOC689750) mRNA.
<a href="#">LOC689755</a>	<a href="#">LOC689755.aSep08</a>	<a href="#">689755</a>	20140	816	3	182	hypothetical protein LOC689755 (LOC689755) alternative variant aSep08, mRNA.
<a href="#">LOC689755</a>	<a href="#">LOC689755.cSep08</a>	<a href="#">689755</a>	14022	341	2	32	hypothetical protein LOC689755 (LOC689755) alternative variant cSep08, mRNA.
<a href="#">LOC689770</a>	<a href="#">LOC689770.aSep08</a>	<a href="#">689770</a>	12590	676		182	similar to osteoclast inhibitory lectin (LOC689770) mRNA.
<a href="#">LOC689775</a>	<a href="#">LOC689775.aSep08</a>	<a href="#">689775</a>	8325	418		83	hypothetical protein LOC689775 (LOC689775) mRNA.
<a href="#">LOC689791</a>	<a href="#">LOC689791.aSep08</a>	<a href="#">689791</a>	10263	334	1	92	hypothetical protein LOC689791 (LOC689791) alternative variant aSep08, mRNA.
<a href="#">LOC689791</a>	<a href="#">LOC689791.bSep08</a>	<a href="#">689791</a>	10771	702		52	hypothetical protein LOC689791 (LOC689791) alternative variant bSep08, mRNA.

<a href="#">LOC689795</a>	<a href="#">LOC689795.aSep08</a>	<a href="#">689795</a>	10180	463		76	hypothetical protein LOC689795 (8.5 kD) (LOC689795) alternative variant aSep08, mRNA.
<a href="#">LOC689795</a>	<a href="#">LOC689795.bSep08</a>	<a href="#">689795</a>	10426	708		76	hypothetical protein LOC689795 (8.5 kD) (LOC689795) alternative variant bSep08, mRNA.
<a href="#">LOC689797</a>	<a href="#">LOC689797.aSep08</a>	<a href="#">689797</a>	3352	655		83	hypothetical protein LOC689797 (9.2 kD) (LOC689797) mRNA.
<a href="#">LOC689799</a>	<a href="#">LOC689799.aSep08</a>	<a href="#">689799</a>	21254	979		228	similar to proteoglycan 4 (LOC689799) mRNA.
<a href="#">LOC689801</a>	<a href="#">LOC689801.aSep08</a>	<a href="#">689801</a>	27642	1202		377	similar to butyrophilin-like 8 (42.8 kD) (LOC689801) mRNA.
<a href="#">LOC689820</a>	<a href="#">LOC689820.aSep08</a>	<a href="#">689820</a>	19220	2780	1	384	similar to SET domain-containing protein (LOC689820) alternative variant aSep08, mRNA.
<a href="#">LOC689820</a>	<a href="#">LOC689820.bSep08</a>	<a href="#">689820</a>	5779	519	1	101	similar to SET domain-containing protein (LOC689820) alternative variant bSep08, mRNA.
<a href="#">LOC689822</a>	<a href="#">LOC689822.aSep08</a>	<a href="#">689822</a>	6487	1426		181	similar to Nucleolar GTP-binding protein 1 (Chronic renal failure gene protein) (GTP-binding protein NGB) (LOC689822) mRNA.
<a href="#">LOC689823</a>	<a href="#">LOC689823.aSep08</a>	<a href="#">689823</a>	3360	433		144	similar to RAD21 homolog (LOC689823) mRNA.
<a href="#">LOC689842</a>	<a href="#">LOC689842.aSep08</a>	<a href="#">689842</a>	5759	1633		129	similar to Nucleolar GTP-binding protein 1 (Chronic renal failure gene protein) (GTP-binding protein NGB) (14.8 kD) (LOC689842) mRNA.
<a href="#">LOC689852</a>	<a href="#">LOC689852.aSep08</a>	<a href="#">689852</a>	13948	821	5	222	similar to Proteasome inhibitor PI31 subunit (LOC689852) alternative variant aSep08, mRNA.
<a href="#">LOC689852</a>	<a href="#">LOC689852.bSep08</a>	<a href="#">689852</a>	12109	720	4	178	similar to Proteasome inhibitor PI31 subunit (LOC689852) alternative variant bSep08, mRNA.
<a href="#">LOC689852</a>	<a href="#">LOC689852.dSep08</a>	<a href="#">689852</a>	22040	1010	7	178	similar to Proteasome inhibitor PI31 subunit (LOC689852) alternative variant dSep08, mRNA.
<a href="#">LOC689852</a>	<a href="#">LOC689852.eSep08</a>	<a href="#">689852</a>	16311	224	3	52	similar to Proteasome inhibitor PI31 subunit (LOC689852) alternative variant eSep08, mRNA.
<a href="#">LOC689852</a>	<a href="#">LOC689852.fSep08</a>	<a href="#">689852</a>	2085	305	2	30	similar to Proteasome inhibitor PI31 subunit (LOC689852) alternative variant fSep08, mRNA.
<a href="#">LOC689920</a>	<a href="#">LOC689920.aSep08</a>	<a href="#">689920</a>	5149	1031		129	similar to G protein-binding protein CRFG (14.9 kD) (LOC689920) mRNA.
<a href="#">LOC689925</a>	<a href="#">LOC689925.aSep08</a>	<a href="#">689925</a>	17066	724		76	similar to xenotropic and polytropic retrovirus receptor 1 (8.7 kD) (LOC689925) mRNA.
<a href="#">LOC689926</a>	<a href="#">LOC689926.aSep08</a>	<a href="#">689926</a>	5732	847	1	110	hypothetical protein LOC689926 (LOC689926) alternative variant aSep08, mRNA.
<a href="#">LOC689926</a>	<a href="#">LOC689926.bSep08</a>	<a href="#">689926</a>	4807	462	1	97	hypothetical protein LOC689926 (LOC689926) alternative variant bSep08, mRNA.
<a href="#">LOC689933</a>	<a href="#">LOC689933.aSep08</a>	<a href="#">689933</a>	26356	644		66	hypothetical protein LOC689933 (LOC689933) mRNA.
<a href="#">LOC689953</a>	<a href="#">LOC689953.bSep08</a>	<a href="#">689953</a>	15079	1261	2	79	hypothetical protein LOC689953 (LOC689953) alternative variant bSep08, mRNA.
<a href="#">LOC689959</a>	<a href="#">LOC689959.bSep08</a>	<a href="#">689959</a>	5901	227	1	75	hypothetical protein LOC689959 (LOC689959) alternative variant bSep08, mRNA.
<a href="#">LOC689963</a>	<a href="#">LOC689963.bSep08</a>	<a href="#">689963</a>	4825	777	2	142	hypothetical protein LOC689963 (LOC689963) alternative variant bSep08, mRNA.
<a href="#">LOC689982</a>	<a href="#">LOC689982.aSep08</a>	<a href="#">689982</a>	4184	919		219	similar to CG18437-PA (LOC689982) mRNA.
<a href="#">LOC689985</a>	<a href="#">LOC689985.aSep08</a>	<a href="#">689985</a>	4071	834		175	hypothetical protein LOC689985 (19.8 kD) (LOC689985) mRNA.

<a href="#">LOC689991</a>	<a href="#">LOC689991.aSep08</a>	<a href="#">689991</a>	28929	1766	10	588	similar to CG18437-PA and similar to CG18437-PA (LOC689991) alternative variant aSep08, mRNA.
<a href="#">LOC689991</a>	<a href="#">LOC689991.aSep08</a>	<a href="#">689998</a>	28929	1766	10	588	similar to CG18437-PA and similar to CG18437-PA (LOC689991) alternative variant aSep08, mRNA.
<a href="#">LOC689991</a>	<a href="#">LOC689991.bSep08</a>	<a href="#">689991</a>	9122	414	2	138	similar to CG18437-PA and similar to CG18437-PA (LOC689991) alternative variant bSep08, mRNA.
<a href="#">LOC689991</a>	<a href="#">LOC689991.bSep08</a>	<a href="#">689998</a>	9122	414	2	138	similar to CG18437-PA and similar to CG18437-PA (LOC689991) alternative variant bSep08, mRNA.
<a href="#">LOC689996</a>	<a href="#">LOC689996.aSep08</a>	<a href="#">689996</a>	4876	765	1	68	similar to translocase of outer mitochondrial membrane 20 homolog (7.8 kD) (LOC689996) alternative variant aSep08, mRNA.
<a href="#">LOC689996</a>	<a href="#">LOC689996.bSep08</a>	<a href="#">689996</a>	5377	279	2	66	similar to translocase of outer mitochondrial membrane 20 homolog (LOC689996) alternative variant bSep08, mRNA.
<a href="#">LOC690000</a>	<a href="#">LOC690000.aSep08</a>	<a href="#">690000</a>	13518	3189	2	141	similar to CG3740-PA (15.1 kD) (LOC690000) alternative variant aSep08, mRNA.
<a href="#">LOC690000</a>	<a href="#">LOC690000.bSep08</a>	<a href="#">690000</a>	10921	455	2	130	similar to CG3740-PA (LOC690000) alternative variant bSep08, mRNA.
<a href="#">LOC690000</a>	<a href="#">LOC690000.cSep08</a>	<a href="#">690000</a>	1714	441	1	48	similar to CG3740-PA (LOC690000) alternative variant cSep08, mRNA.
<a href="#">LOC690012</a>	<a href="#">LOC690012.aSep08</a>	<a href="#">690012</a>	10918	798	6	118	similar to High mobility group protein 2 (HMG-2) (13.0 kD) (LOC690012) alternative variant aSep08, mRNA.
<a href="#">LOC690020</a>	<a href="#">LOC690020.aSep08</a>	<a href="#">690020</a>	9857	601		57	similar to killer cell lectin-like receptor, subfamily A, member 17 (LOC690020) mRNA.
<a href="#">LOC690032</a>	<a href="#">LOC690032.aSep08</a>	<a href="#">690032</a>	16198	1916	9	428	similar to WD repeat domain 4 (LOC690032) alternative variant aSep08, mRNA.
<a href="#">LOC690032</a>	<a href="#">LOC690032.bSep08</a>	<a href="#">690032</a>	18942	1112	3	105	similar to WD repeat domain 4 (LOC690032) alternative variant bSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.aSep08</a>	<a href="#">299123</a>	82893	1783	12	593	CRA a (LOC690035) alternative variant aSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.aSep08</a>	<a href="#">690035</a>	82893	1783	12	593	CRA a (LOC690035) alternative variant aSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.bSep08</a>	<a href="#">299123</a>	58178	2006	12	343	CRA a (37.2 kD) (LOC690035) alternative variant bSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.bSep08</a>	<a href="#">690035</a>	58178	2006	12	343	CRA a (37.2 kD) (LOC690035) alternative variant bSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.cSep08</a>	<a href="#">299123</a>	41779	674	4	165	CRA a like (LOC690035) alternative variant cSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.cSep08</a>	<a href="#">690035</a>	41779	674	4	165	CRA a like (LOC690035) alternative variant cSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.dSep08</a>	<a href="#">299123</a>	2005	625	2	86	CRA a (LOC690035) alternative variant dSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.dSep08</a>	<a href="#">690035</a>	2005	625	2	86	CRA a (LOC690035) alternative variant dSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.eSep08</a>	<a href="#">299123</a>	63857	466	4	72	CRA a like (LOC690035) alternative variant eSep08, mRNA.
<a href="#">LOC690035</a>	<a href="#">LOC690035.eSep08</a>	<a href="#">690035</a>	63857	466	4	72	CRA a like (LOC690035) alternative variant eSep08, mRNA.
<a href="#">LOC690040</a>	<a href="#">LOC690040.aSep08</a>	<a href="#">690040</a>	18447	679	5	183	similar to protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b (LOC690040) alternative variant aSep08, mRNA.

<a href="#">LOC690042</a>	<a href="#">LOC690042.aSep08</a>	<a href="#">690042</a>	10288	942	4	261	hypothetical protein LOC690042 (LOC690042) alternative variant aSep08, mRNA.
<a href="#">LOC690042</a>	<a href="#">LOC690042.bSep08</a>	<a href="#">690042</a>	10129	618	3	163	hypothetical protein LOC690042 (LOC690042) alternative variant bSep08, mRNA.
<a href="#">LOC690042</a>	<a href="#">LOC690042.cSep08</a>	<a href="#">690042</a>	5316	444	2	61	hypothetical protein LOC690042 (7.0 kD) (LOC690042) alternative variant cSep08, mRNA.
<a href="#">LOC690043</a>	<a href="#">LOC690043.bSep08</a>	<a href="#">690043</a>	16329	2885	2	316	similar to ring finger protein 168 (35.6 kD) (LOC690043) alternative variant bSep08, mRNA.
<a href="#">LOC690045</a>	<a href="#">LOC690045.aSep08</a>	<a href="#">690045</a>	21215	859		193	similar to immunoreceptor Ly49si1 (LOC690045) mRNA.
<a href="#">LOC690068</a>	<a href="#">LOC690068.aSep08</a>	<a href="#">690068</a>	22317	1749	8	100	putative protein of ancient origin (LOC690068) alternative variant aSep08, mRNA.
<a href="#">LOC690068</a>	<a href="#">LOC690068.bSep08</a>	<a href="#">690068</a>	9671	519	3	92	putative protein of eukaryotic origin (LOC690068) alternative variant bSep08, mRNA.
<a href="#">LOC690068</a>	<a href="#">LOC690068.cSep08</a>	<a href="#">690068</a>	3565	244	3	52	putative protein (LOC690068) alternative variant cSep08, mRNA.
<a href="#">LOC690068</a>	<a href="#">LOC690068.dSep08</a>	<a href="#">690068</a>	5023	561	3	32	putative protein (3.6 kD) (LOC690068) alternative variant dSep08, mRNA.
<a href="#">LOC690068</a>	<a href="#">LOC690068.eSep08</a>	<a href="#">690068</a>	22330	411	5	43	putative protein of eukaryotic origin (LOC690068) alternative variant eSep08, mRNA.
<a href="#">LOC690073</a>	<a href="#">LOC690073.aSep08</a>	<a href="#">690073</a>	102821	2023	9	561	similar to neuroblastoma-amplified protein (LOC690073) alternative variant aSep08, mRNA.
<a href="#">LOC690073</a>	<a href="#">LOC690073.bSep08</a>	<a href="#">690073</a>	36428	932	5	241	similar to neuroblastoma-amplified protein (27.2 kD) (LOC690073) alternative variant bSep08, mRNA.
<a href="#">LOC690073</a>	<a href="#">LOC690073.cSep08</a>	<a href="#">690073</a>	31846	766	4	207	similar to neuroblastoma-amplified protein (LOC690073) alternative variant cSep08, mRNA.
<a href="#">LOC690073</a>	<a href="#">LOC690073.dSep08</a>	<a href="#">690073</a>	25487	759	3	198	similar to neuroblastoma-amplified protein (22.1 kD) (LOC690073) alternative variant dSep08, mRNA.
<a href="#">LOC690073</a>	<a href="#">LOC690073.eSep08</a>	<a href="#">690073</a>	21019	753	3	103	similar to neuroblastoma-amplified protein (LOC690073) alternative variant eSep08, mRNA.
<a href="#">LOC690082</a>	<a href="#">LOC690082.aSep08</a>	<a href="#">690082</a>	4326	1085		115	similar to melanoma ubiquitous mutated protein (12.6 kD) (LOC690082) mRNA.
<a href="#">LOC690085</a>	<a href="#">LOC690085.aSep08</a>	<a href="#">690085</a>	30186	1974		550	similar to B-cell CLL/lymphoma 7A (LOC690085) alternative variant aSep08, mRNA.
<a href="#">LOC690097</a>	<a href="#">LOC690097.aSep08</a>	<a href="#">690097</a>	11168	813		50	similar to immunoreceptor Ly49si3 (LOC690097) mRNA.
<a href="#">LOC690101</a>	<a href="#">LOC690101.aSep08</a>	<a href="#">690101</a>	617	263		87	hypothetical protein LOC690101 (LOC690101) mRNA.
<a href="#">LOC690116</a>	<a href="#">LOC690116.aSep08</a>	<a href="#">690116</a>	22980	781		243	similar to regulator of G-protein signalling like 2 (LOC690116) alternative variant aSep08, mRNA.
<a href="#">LOC690116</a>	<a href="#">LOC690116.bSep08</a>	<a href="#">690116</a>	8236	563		187	similar to regulator of G-protein signalling like 2 (LOC690116) alternative variant bSep08, mRNA.
<a href="#">LOC690135</a>	<a href="#">LOC690135.aSep08</a>	<a href="#">690135</a>	53341	618		163	hypothetical protein LOC690135 (LOC690135) mRNA.
<a href="#">LOC690139</a>	<a href="#">LOC690139.aSep08</a>	<a href="#">690139</a>	9179	1073	2	180	similar to RNA binding motif protein 24 (LOC690139) alternative variant aSep08, mRNA.
<a href="#">LOC690139</a>	<a href="#">LOC690139.bSep08</a>	<a href="#">690139</a>	9934	851	4	152	similar to RNA binding motif protein 24 (LOC690139) alternative variant bSep08, mRNA.
<a href="#">LOC690139</a>	<a href="#">LOC690139.cSep08</a>	<a href="#">690139</a>	3165	694	2	146	similar to RNA binding motif protein 24 (LOC690139) alternative variant cSep08, mRNA.

<a href="#">LOC690155</a>	<a href="#">LOC690155.aSep08</a>	<a href="#">690155</a>	7830	1949		208	similar to phosducin-like (23.8 kD) (LOC690155) mRNA.
<a href="#">LOC690171</a>	<a href="#">LOC690171.bSep08</a>	<a href="#">690171</a>	724	392	2	125	similar to H3 histone, family 3B (LOC690171) alternative variant bSep08, mRNA.
<a href="#">LOC690188</a>	<a href="#">LOC690188.aSep08</a>	<a href="#">690188</a>	15983	341		113	similar to Protein KIAA0280 (LOC690188) mRNA.
<a href="#">LOC690190</a>	<a href="#">LOC690190.aSep08</a>	<a href="#">690190</a>	1100	995		31	hypothetical protein LOC690190 (12.7 kD) (LOC690190) mRNA.
<a href="#">LOC690208</a>	<a href="#">LOC690208.aSep08</a>	<a href="#">690208</a>	23560	1165	4	307	hypothetical protein LOC690208 (LOC690208) alternative variant aSep08, mRNA.
<a href="#">LOC690208</a>	<a href="#">LOC690208.bSep08</a>	<a href="#">690208</a>	9704	1254	1	201	hypothetical protein LOC690208 (LOC690208) alternative variant bSep08, mRNA.
<a href="#">LOC690214</a>	<a href="#">LOC690214.aSep08</a>	<a href="#">690214</a>	2326	1688	2	231	similar to peptide deformylase-like protein (LOC690214) alternative variant aSep08, complete mRNA.
<a href="#">LOC690217</a>	<a href="#">LOC690217.aSep08</a>	<a href="#">690217</a>	7002	1001		147	similar to B0511.12 (LOC690217) mRNA.
<a href="#">LOC690226</a>	<a href="#">LOC690226.aSep08</a>	<a href="#">690226</a>	10394	1617	5	318	similar to dehydrogenase/reductase (SDR family) member 7 (LOC690226) alternative variant aSep08, mRNA.
<a href="#">LOC690226</a>	<a href="#">LOC690226.bSep08</a>	<a href="#">690226</a>	1448	841	1	64	similar to dehydrogenase/reductase (SDR family) member 7 (7.6 kD) (LOC690226) alternative variant bSep08, mRNA.
<a href="#">LOC690236</a>	<a href="#">LOC690236.aSep08</a>	<a href="#">690236</a>	52575	306		68	hypothetical protein LOC690236 (LOC690236) mRNA.
<a href="#">LOC690243</a>	<a href="#">LOC690243.aSep08</a>	<a href="#">690243</a>	2282	748	8	57	CRA a like (6.3 kD) (LOC690243) alternative variant aSep08, mRNA.
<a href="#">LOC690243</a>	<a href="#">LOC690243.bSep08</a>	<a href="#">690243</a>	1897	1176	4	56	CRA d like (6.4 kD) (LOC690243) alternative variant bSep08, mRNA.
<a href="#">LOC690243</a>	<a href="#">LOC690243.cSep08</a>	<a href="#">690243</a>	1417	727	4	40	CRA d like (LOC690243) alternative variant cSep08, mRNA.
<a href="#">LOC690243</a>	<a href="#">LOC690243.dSep08</a>	<a href="#">690243</a>	2459	1214	6	56	CRA d like (6.4 kD) (LOC690243) alternative variant dSep08, mRNA.
<a href="#">LOC690243</a>	<a href="#">LOC690243.eSep08</a>	<a href="#">690243</a>	2197	749	7	40	CRA d like (LOC690243) alternative variant eSep08, complete mRNA.
<a href="#">LOC690274</a>	<a href="#">LOC690274.aSep08</a>	<a href="#">690274</a>	2257	555	1	89	hypothetical protein LOC690274 (LOC690274) alternative variant aSep08, mRNA.
<a href="#">LOC690274</a>	<a href="#">LOC690274.bSep08</a>	<a href="#">690274</a>	965	525	1	71	hypothetical protein LOC690274 (8.0 kD) (LOC690274) alternative variant bSep08, mRNA.
<a href="#">LOC690276</a>	<a href="#">LOC690276.bSep08</a>	<a href="#">690276</a>	3749	586	2	132	hypothetical protein LOC690276 (LOC690276) alternative variant bSep08, mRNA.
<a href="#">LOC690280</a>	<a href="#">LOC690280.aSep08</a>	<a href="#">690280</a>	2495	437		145	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC690280) mRNA.
<a href="#">LOC690286</a>	<a href="#">LOC690286.aSep08</a>	<a href="#">690286</a>	4851	276		92	similar to hepatic leukemia factor (LOC690286) mRNA.
<a href="#">LOC690295</a>	<a href="#">LOC690295.aSep08</a>	<a href="#">690295</a>	632	305		94	similar to dynein, axonemal, heavy polypeptide 1 (LOC690295) mRNA.
<a href="#">LOC690298</a>	<a href="#">LOC690298.aSep08</a>	<a href="#">690298</a>	20502	648		93	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) and similar to Reticulocalbin-1 precursor (10.9 kD) (LOC690298) mRNA.
<a href="#">LOC690298</a>	<a href="#">LOC690298.aSep08</a>	<a href="#">690306</a>	20502	648		93	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) and similar to Reticulocalbin-1 precursor (10.9 kD) (LOC690298) mRNA.

<a href="#">LOC690323</a>	<a href="#">LOC690323.aSep08</a>	<a href="#">690323</a>	5826	2732	11	157	similar to Myosin-15 (Myosin XV) (Unconventional myosin-15) (17.8 kD) (LOC690323) alternative variant aSep08, mRNA.
<a href="#">LOC690323</a>	<a href="#">LOC690323.bSep08</a>	<a href="#">690323</a>	8779	375	4	66	similar to Myosin-15 (Myosin XV) (Unconventional myosin-15) (LOC690323) alternative variant bSep08, mRNA.
<a href="#">LOC690340</a>	<a href="#">LOC690340.aSep08</a>	<a href="#">690340</a>	3732	593		142	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC690340) mRNA.
<a href="#">LOC690343</a>	<a href="#">LOC690343.aSep08</a>	<a href="#">690343</a>	29999	658		112	similar to zinc finger protein 420 (LOC690343) mRNA.
<a href="#">LOC690344</a>	<a href="#">LOC690344.aSep08</a>	<a href="#">690344</a>	796	707	2	89	similar to Protein UNQ655/PRO1286 homolog precursor (9.7 kD) (LOC690344) alternative variant aSep08, mRNA.
<a href="#">LOC690347</a>	<a href="#">LOC690347.aSep08</a>	<a href="#">690347</a>	870	287	1	86	hypothetical protein LOC690347 (LOC690347) alternative variant aSep08, mRNA.
<a href="#">LOC690347</a>	<a href="#">LOC690347.bSep08</a>	<a href="#">690347</a>	9050	313	2	52	hypothetical protein LOC690347 (LOC690347) alternative variant bSep08, mRNA.
<a href="#">LOC690358</a>	<a href="#">LOC690358.aSep08</a>	<a href="#">690358</a>	51824	749		185	putative protein of metazoan origin (20.7 kD) (LOC690358) mRNA.
<a href="#">LOC690372</a>	<a href="#">LOC690372.aSep08</a>	<a href="#">690372</a>	17569	1452	12	471	similar to U2 (RNU2) small nuclear RNA auxiliary factor 2 isoform b (53.1 kD) (LOC690372) alternative variant aSep08, mRNA.
<a href="#">LOC690372</a>	<a href="#">LOC690372.bSep08</a>	<a href="#">690372</a>	10185	759	7	252	similar to U2 (RNU2) small nuclear RNA auxiliary factor 2 isoform b (LOC690372) alternative variant bSep08, mRNA.
<a href="#">LOC690372</a>	<a href="#">LOC690372.cSep08</a>	<a href="#">690372</a>	6808	848	6	223	similar to U2 (RNU2) small nuclear RNA auxiliary factor 2 isoform b (LOC690372) alternative variant cSep08, mRNA.
<a href="#">LOC690402</a>	<a href="#">LOC690402.bSep08</a>	<a href="#">690402</a>	1171	333	2	14	similar to Short palate, lung and nasal epithelium carcinoma-associated protein 3 homolog precursor (LOC690402) alternative variant bSep08, mRNA.
<a href="#">LOC690414</a>	<a href="#">LOC690414.aSep08</a>	<a href="#">690414</a>	9785	1475		66	hypothetical protein LOC690414 (6.8 kD) (LOC690414) mRNA.
<a href="#">LOC690419</a>	<a href="#">LOC690419.aSep08</a>	<a href="#">690419</a>	16886	352		100	similar to zinc finger protein 709 (LOC690419) mRNA.
<a href="#">LOC690422</a>	<a href="#">LOC690422.aSep08</a>	<a href="#">690422</a>	1275	558	4	185	hypothetical protein LOC690422 (LOC690422) alternative variant aSep08, mRNA.
<a href="#">LOC690422</a>	<a href="#">LOC690422.bSep08</a>	<a href="#">690422</a>	4665	740	6	183	hypothetical protein LOC690422 (LOC690422) alternative variant bSep08, mRNA.
<a href="#">LOC690422</a>	<a href="#">LOC690422.cSep08</a>	<a href="#">690422</a>	4260	727	4	112	hypothetical protein LOC690422 (LOC690422) alternative variant cSep08, mRNA.
<a href="#">LOC690437</a>	<a href="#">LOC690437.aSep08</a>	<a href="#">690437</a>	1014	581	1	110	hypothetical protein LOC690437 (4.0 kD) (LOC690437) alternative variant aSep08, mRNA.
<a href="#">LOC690437</a>	<a href="#">LOC690437.bSep08</a>	<a href="#">690437</a>	2219	432	3	84	hypothetical protein LOC690437 (LOC690437) alternative variant bSep08, mRNA.
<a href="#">LOC690470</a>	<a href="#">LOC690470.bSep08</a>	<a href="#">690470</a>	28493	758	1	135	similar to 1-aminocyclopropane-1-carboxylate synthase (LOC690470) alternative variant bSep08, mRNA.
<a href="#">LOC690479</a>	<a href="#">LOC690479.aSep08</a>	<a href="#">690479</a>	5252	335		35	hypothetical protein LOC690479 (LOC690479) mRNA.
<a href="#">LOC690482</a>	<a href="#">LOC690482.aSep08</a>	<a href="#">690482</a>	2008	533		93	hypothetical protein LOC690482 (LOC690482) mRNA.
<a href="#">LOC690483</a>	<a href="#">LOC690483.aSep08</a>	<a href="#">690483</a>	4087	659		209	hypothetical protein LOC690483 (LOC690483) mRNA.
<a href="#">LOC690492</a>	<a href="#">LOC690492.aSep08</a>	<a href="#">690492</a>	6136	1773	1	502	similar to CD33 antigen (LOC690492) alternative variant aSep08, mRNA.



<a href="#">LOC690492</a>	<a href="#">LOC690492.bSep08</a>	<a href="#">690492</a>	1961	804	3	65	similar to CD33 antigen (LOC690492) alternative variant bSep08, mRNA.
<a href="#">LOC690492</a>	<a href="#">LOC690492.dSep08</a>	<a href="#">690492</a>	3794	675	3	8	similar to CD33 antigen (0.9 kD) (LOC690492) alternative variant dSep08, mRNA.
<a href="#">LOC690492</a>	<a href="#">LOC690492.eSep08</a>	<a href="#">690492</a>	550	293			
<a href="#">LOC690502</a>	<a href="#">LOC690502.aSep08</a>	<a href="#">690502</a>	26229	820	2	153	similar to zinc finger protein 420 (LOC690502) alternative variant aSep08, mRNA.
<a href="#">LOC690502</a>	<a href="#">LOC690502.bSep08</a>	<a href="#">690502</a>	24163	536	1	114	similar to zinc finger protein 420 (LOC690502) alternative variant bSep08, mRNA.
<a href="#">LOC690502</a>	<a href="#">LOC690502.cSep08</a>	<a href="#">690502</a>	19065	462	2	65	similar to zinc finger protein 420 (LOC690502) alternative variant cSep08, mRNA.
<a href="#">LOC690507</a>	<a href="#">LOC690507.aSep08</a>	<a href="#">690507</a>	2641	295		98	similar to Vomeromodulin (LOC690507) mRNA.
<a href="#">LOC690538</a>	<a href="#">LOC690538.aSep08</a>	<a href="#">690538</a>	73203	725		150	similar to Protein C9orf126 homolog (LOC690538) mRNA.
<a href="#">LOC690543</a>	<a href="#">LOC690543.aSep08</a>	<a href="#">690543</a>	14324	1075		235	similar to zinc finger protein 679 (LOC690543) mRNA.
<a href="#">LOC690559</a>	<a href="#">LOC690559.aSep08</a>	<a href="#">690559</a>	45319	595		131	similar to reduced expression 2 (LOC690559) mRNA.
<a href="#">LOC690576</a>	<a href="#">LOC690576.aSep08</a>	<a href="#">690576</a>	18345	714	1	213	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC690576) alternative variant aSep08, mRNA.
<a href="#">LOC690576</a>	<a href="#">LOC690576.bSep08</a>	<a href="#">690576</a>	18400	609		164	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC690576) alternative variant bSep08, mRNA.
<a href="#">LOC690629</a>	<a href="#">LOC690629.aSep08</a>	<a href="#">690629</a>	5587	1784		515	hypothetical protein LOC690629 (LOC690629) mRNA.
<a href="#">LOC690665</a>	<a href="#">LOC690665.aSep08</a>	<a href="#">690665</a>	13782	997		156	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (18.5 kD) (LOC690665) mRNA.
<a href="#">LOC690692</a>	<a href="#">LOC690692.aSep08</a>	<a href="#">690692</a>	55275	528		56	hypothetical protein LOC690692 (6.2 kD) (LOC690692) mRNA.
<a href="#">LOC690693</a>	<a href="#">LOC690693.aSep08</a>	<a href="#">690693</a>	18497	1128	1	334	similar to DDX19 homolog (LOC690693) alternative variant aSep08, mRNA.
<a href="#">LOC690693</a>	<a href="#">LOC690693.bSep08</a>	<a href="#">690693</a>	20871	708	1	235	similar to DDX19 homolog (LOC690693) alternative variant bSep08, mRNA.
<a href="#">LOC690727</a>	<a href="#">LOC690727.aSep08</a>	<a href="#">690727</a>	1045	924		235	similar to nucleolar protein 1 (LOC690727) mRNA.
<a href="#">LOC690728</a>	<a href="#">LOC690728.bSep08</a>	<a href="#">690728</a>	13993	2070	9	524	similar to Protein C12orf11 (Sarcoma antigen NY-SAR-95) (LOC690728) alternative variant bSep08, mRNA.
<a href="#">LOC690728</a>	<a href="#">LOC690728.cSep08</a>	<a href="#">690728</a>	5253	990	5	218	similar to Protein C12orf11 (Sarcoma antigen NY-SAR-95) (LOC690728) alternative variant cSep08, mRNA.
<a href="#">LOC690728</a>	<a href="#">LOC690728.dSep08</a>	<a href="#">690728</a>	1217	337	3	100	similar to Protein C12orf11 (Sarcoma antigen NY-SAR-95) (LOC690728) alternative variant dSep08, mRNA.
<a href="#">LOC690739</a>	<a href="#">LOC690739.aSep08</a>	<a href="#">690739</a>	18424	711	3	147	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC690739) alternative variant aSep08, mRNA.
<a href="#">LOC690739</a>	<a href="#">LOC690739.bSep08</a>	<a href="#">690739</a>	2482	437	2	145	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC690739) alternative variant bSep08, mRNA.

<a href="#">LOC690739</a>	<a href="#">LOC690739.cSep08</a>	<a href="#">690739</a>	23750	1476	4	93	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (10.8 kD) (LOC690739) alternative variant cSep08, mRNA.
<a href="#">LOC690743</a>	<a href="#">LOC690743.aSep08</a>	<a href="#">690743</a>	11416	717		239	similar to mixed lineage kinase domain-like (LOC690743) mRNA.
<a href="#">LOC690745</a>	<a href="#">LOC690745.aSep08</a>	<a href="#">690745</a>	54986	1571		204	MOSC (LOC690745) mRNA.
<a href="#">LOC690746</a>	<a href="#">LOC690746.aSep08</a>	<a href="#">690746</a>	70670	385		128	similar to projection protein PF6 (LOC690746) mRNA.
<a href="#">LOC690751</a>	<a href="#">LOC690751.aSep08</a>	<a href="#">690751</a>	26318	1489	2	496	similar to WD repeat domain 59 (LOC690751) alternative variant aSep08, mRNA.
<a href="#">LOC690751</a>	<a href="#">LOC690751.bSep08</a>	<a href="#">690751</a>	1667	259	1	85	similar to WD repeat domain 59 (LOC690751) alternative variant bSep08, mRNA.
<a href="#">LOC690768</a>	<a href="#">LOC690768.aSep08</a>	<a href="#">690768</a>	15329	690		53	hypothetical protein LOC690768 (5.7 kD) (LOC690768) mRNA.
<a href="#">LOC690769</a>	<a href="#">LOC690769.aSep08</a>	<a href="#">690769</a>	85097	2418	5	326	similar to zinc ring finger protein 1 (35.0 kD) (LOC690769) alternative variant aSep08, mRNA.
<a href="#">LOC690769</a>	<a href="#">LOC690769.bSep08</a>	<a href="#">690769</a>	83683	682	5	120	similar to zinc ring finger protein 1 (LOC690769) alternative variant bSep08, mRNA.
<a href="#">LOC690769</a>	<a href="#">LOC690769.cSep08</a>	<a href="#">690769</a>	32981	640	5	71	similar to zinc ring finger protein 1 (8.1 kD) (LOC690769) alternative variant cSep08, mRNA.
<a href="#">LOC690771</a>	<a href="#">LOC690771.aSep08</a>	<a href="#">690771</a>	594	302		100	hypothetical protein LOC690771 (LOC690771) mRNA.
<a href="#">LOC690774</a>	<a href="#">LOC690774.aSep08</a>	<a href="#">690774</a>	426	231		45	hypothetical protein LOC690774 (LOC690774) mRNA.
<a href="#">LOC690776</a>	<a href="#">LOC690776.aSep08</a>	<a href="#">690776</a>	30668	1211		97	hypothetical protein LOC690776 (LOC690776) mRNA.
<a href="#">LOC690777</a>	<a href="#">LOC690777.aSep08</a>	<a href="#">690777</a>	12395	741		223	similar to RUN and FYVE domain-containing 2 (LOC690777) mRNA.
<a href="#">LOC690785</a>	<a href="#">LOC690785.aSep08</a>	<a href="#">690785</a>	7309	477		100	hypothetical protein LOC690785 (11.5 kD) (LOC690785) alternative variant aSep08, mRNA.
<a href="#">LOC690785</a>	<a href="#">LOC690785.bSep08</a>	<a href="#">690785</a>	7431	688	1	100	hypothetical protein LOC690785 (11.5 kD) (LOC690785) alternative variant bSep08, mRNA.
<a href="#">LOC690789</a>	<a href="#">LOC690789.aSep08</a>	<a href="#">690789</a>	3527	1594		173	similar to Ornithine decarboxylase antizyme 2 (ODC-Az 2) (AZ2) (LOC690789) mRNA.
<a href="#">LOC690806</a>	<a href="#">LOC690806.aSep08</a>	<a href="#">690806</a>	14555	746		88	hypothetical protein LOC690806 (10.0 kD) (LOC690806) mRNA.
<a href="#">LOC690818</a>	<a href="#">LOC690818.aSep08</a>	<a href="#">690818</a>	10415	759	1	79	similar to nidogen 2 (8.6 kD) (LOC690818) alternative variant aSep08, mRNA.
<a href="#">LOC690818</a>	<a href="#">LOC690818.bSep08</a>	<a href="#">690818</a>	9622	691	1	79	similar to nidogen 2 (8.6 kD) (LOC690818) alternative variant bSep08, mRNA.
<a href="#">LOC690825</a>	<a href="#">LOC690825.aSep08</a>	<a href="#">690825</a>	2643	595	2	86	similar to guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2 (LOC690825) alternative variant aSep08, mRNA.
<a href="#">LOC690825</a>	<a href="#">LOC690825.bSep08</a>	<a href="#">690825</a>	2083	501	1	69	similar to guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2 (7.8 kD) (LOC690825) alternative variant bSep08, mRNA.
<a href="#">LOC690825</a>	<a href="#">LOC690825.cSep08</a>	<a href="#">690825</a>	3087	498	2	69	similar to guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2 (7.8 kD) (LOC690825) alternative variant cSep08, mRNA.
<a href="#">LOC690846</a>	<a href="#">LOC690846.bSep08</a>	<a href="#">690846</a>	51868	808	1	268	similar to resistance to inhibitors of cholinesterase 3 homolog (LOC690846) alternative variant bSep08, mRNA.

<a href="#">LOC690851</a>	<a href="#">LOC690851.aSep08</a>	<a href="#">690851</a>	7945	261		65	hypothetical protein LOC690851 (LOC690851) mRNA.
<a href="#">LOC690871</a>	<a href="#">LOC690871.aSep08</a>	<a href="#">690871</a>	2149	450		86	hypothetical protein LOC690871 (LOC690871) mRNA.
<a href="#">LOC690883</a>	<a href="#">LOC690883.aSep08</a>	<a href="#">690883</a>	6070	1196		74	hypothetical protein LOC690883 (8.0 kD) (LOC690883) mRNA.
<a href="#">LOC690895</a>	<a href="#">LOC690895.bSep08</a>	<a href="#">690895</a>	29886	764	8	133	similar to zinc finger protein 426 (LOC690895) alternative variant bSep08, mRNA.
<a href="#">LOC690895</a>	<a href="#">LOC690895.cSep08</a>	<a href="#">690895</a>	10892	365	4	67	similar to zinc finger protein 426 (LOC690895) alternative variant cSep08, mRNA.
<a href="#">LOC690911</a>	<a href="#">LOC690911.aSep08</a>	<a href="#">690911</a>	2490	1799	4	253	similar to Msx2-interacting protein (SPEN homolog) (SMART/HDAC1-associated repressor protein) (LOC690911) alternative variant aSep08, mRNA.
<a href="#">LOC690930</a>	<a href="#">LOC690930.aSep08</a>	<a href="#">690930</a>	5402	615		123	similar to membrane-spanning 4-domains, subfamily A, member 6B (LOC690930) mRNA.
<a href="#">LOC690945</a>	<a href="#">LOC690945.aSep08</a>	<a href="#">690945</a>	79695	1739	8	345	similar to Putative deoxyribose-phosphate aldolase (Phosphodeoxyriboaldolase) (Deoxyriboaldolase) (DERA) (LOC690945) alternative variant aSep08, mRNA.
<a href="#">LOC690945</a>	<a href="#">LOC690945.bSep08</a>	<a href="#">690945</a>	79170	1342	6	300	similar to Putative deoxyribose-phosphate aldolase (Phosphodeoxyriboaldolase) (Deoxyriboaldolase) (DERA) (LOC690945) alternative variant bSep08, mRNA.
<a href="#">LOC690945</a>	<a href="#">LOC690945.cSep08</a>	<a href="#">690945</a>	79255	694	3	116	similar to Putative deoxyribose-phosphate aldolase (Phosphodeoxyriboaldolase) (Deoxyriboaldolase) (DERA) (12.9 kD) (LOC690945) alternative variant cSep08, complete mRNA.
<a href="#">LOC690945</a>	<a href="#">LOC690945.dSep08</a>	<a href="#">690945</a>	11995	1278	3	38	similar to Putative deoxyribose-phosphate aldolase (Phosphodeoxyriboaldolase) (Deoxyriboaldolase) (DERA) (LOC690945) alternative variant dSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.aSep08</a>	<a href="#">690948</a>	4668	1299	9	388	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant aSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.aSep08</a>	<a href="#">690955</a>	4668	1299	9	388	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant aSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.bSep08</a>	<a href="#">690948</a>	44343	1083	5	361	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant bSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.bSep08</a>	<a href="#">690955</a>	44343	1083	5	361	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant bSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.cSep08</a>	<a href="#">690948</a>	3457	689	7	186	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant cSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.cSep08</a>	<a href="#">690955</a>	3457	689	7	186	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (LOC690948) alternative variant cSep08, mRNA.
<a href="#">LOC690948</a>	<a href="#">LOC690948.dSep08</a>	<a href="#">690948</a>	1522	592	2	79	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (8.4 kD) (LOC690948) alternative variant dSep08, mRNA.

<a href="#">LOC690948</a>	<a href="#">LOC690948.dSep08</a>	<a href="#">690955</a>	1522	592	2	79	similar to paired-Ig-like receptor A11 and similar to paired-Ig-like receptor B (8.4 kD) (LOC690948) alternative variant dSep08, mRNA.
<a href="#">LOC690965</a>	<a href="#">LOC690965.aSep08</a>	<a href="#">690965</a>	4943	500		108	similar to CG17265-PA (11.8 kD) (LOC690965) mRNA.
<a href="#">LOC690987</a>	<a href="#">LOC690987.bSep08</a>	<a href="#">690987</a>	6923	752	1	181	similar to glycogen synthase 1, muscle (LOC690987) alternative variant bSep08, mRNA.
<a href="#">LOC690989</a>	<a href="#">LOC690989.aSep08</a>	<a href="#">690989</a>	4117	387	1	109	similar to Docking protein 5 (Downstream of tyrosine kinase 5) (Protein dok-5) (LOC690989) alternative variant aSep08, mRNA.
<a href="#">LOC690989</a>	<a href="#">LOC690989.bSep08</a>	<a href="#">690989</a>	1243	298	1	79	similar to Docking protein 5 (Downstream of tyrosine kinase 5) (Protein dok-5) (LOC690989) alternative variant bSep08, mRNA.
<a href="#">LOC691000</a>	<a href="#">LOC691000.aSep08</a>	<a href="#">691000</a>	5203	619		79	hypothetical protein LOC691000 (8.7 kD) (LOC691000) mRNA.
<a href="#">LOC691024</a>	<a href="#">LOC691024.bSep08</a>	<a href="#">691024</a>	46292	424	3	54	similar to Protein C9orf25 homolog (LOC691024) alternative variant bSep08, mRNA.
<a href="#">LOC691031</a>	<a href="#">LOC691031.aSep08</a>	<a href="#">691031</a>	38616	2781	4	83	hypothetical protein LOC691031 (9.1 kD) (LOC691031) alternative variant aSep08, mRNA.
<a href="#">LOC691031</a>	<a href="#">LOC691031.bSep08</a>	<a href="#">691031</a>	65690	698	4	61	hypothetical protein LOC691031 (LOC691031) alternative variant bSep08, mRNA.
<a href="#">LOC691031</a>	<a href="#">LOC691031.dSep08</a>	<a href="#">691031</a>	47378	773	2	52	hypothetical protein LOC691031 (5.8 kD) (LOC691031) alternative variant dSep08, mRNA.
<a href="#">LOC691031</a>	<a href="#">LOC691031.eSep08</a>	<a href="#">691031</a>	36793	656	3	57	hypothetical protein LOC691031 (5.9 kD) (LOC691031) alternative variant eSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.aSep08</a>	<a href="#">691036</a>	380074	6933	34	1068	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant aSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.aSep08</a>	<a href="#">691042</a>	380074	6933	34	1068	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant aSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.bSep08</a>	<a href="#">691036</a>	26288	994	6	307	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant bSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.bSep08</a>	<a href="#">691042</a>	26288	994	6	307	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant bSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.cSep08</a>	<a href="#">691036</a>	4231	455	3	84	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant cSep08, mRNA.
<a href="#">LOC691036</a>	<a href="#">LOC691036.cSep08</a>	<a href="#">691042</a>	4231	455	3	84	similar to SET binding factor 2 and similar to SET binding factor 2 (LOC691036) alternative variant cSep08, mRNA.
<a href="#">LOC691056</a>	<a href="#">LOC691056.aSep08</a>	<a href="#">691056</a>	9101	2669	5	812	similar to 5E5 antigen (LOC691056) alternative variant aSep08, mRNA.
<a href="#">LOC691056</a>	<a href="#">LOC691056.bSep08</a>	<a href="#">691056</a>	5161	1041	4	347	similar to 5E5 antigen (LOC691056) alternative variant bSep08, mRNA.
<a href="#">LOC691059</a>	<a href="#">LOC691059.aSep08</a>	<a href="#">691059</a>	10863	732	4	185	similar to GTPase activating protein testicular GAP1 (LOC691059) alternative variant aSep08, mRNA.
<a href="#">LOC691059</a>	<a href="#">LOC691059.bSep08</a>	<a href="#">691059</a>	899	771	2	68	similar to GTPase activating protein testicular GAP1 (LOC691059) alternative variant bSep08, mRNA.
<a href="#">LOC691059</a>	<a href="#">LOC691059.cSep08</a>	<a href="#">691059</a>	5770	705	4	98	similar to GTPase activating protein testicular GAP1 (LOC691059) alternative variant cSep08, mRNA.

<a href="#">LOC691059</a>	<a href="#">LOC691059.dSep08</a>	<a href="#">691059</a>	7038	693	4	77	similar to GTPase activating protein testicular GAP1 (LOC691059) alternative variant dSep08, mRNA.
<a href="#">LOC691083</a>	<a href="#">LOC691083.aSep08</a>	<a href="#">691083</a>	89565	776	1	199	hypothetical protein LOC691083 (22.4 kD) (LOC691083) alternative variant aSep08, mRNA.
<a href="#">LOC691083</a>	<a href="#">LOC691083.bSep08</a>	<a href="#">691083</a>	89161	460	2	139	hypothetical protein LOC691083 (LOC691083) alternative variant bSep08, mRNA.
<a href="#">LOC691083</a>	<a href="#">LOC691083.cSep08</a>	<a href="#">691083</a>	159008	883	1	138	hypothetical protein LOC691083 (15.8 kD) (LOC691083) alternative variant cSep08, mRNA.
<a href="#">LOC691092</a>	<a href="#">LOC691092.aSep08</a>	<a href="#">691092</a>	1909	701		233	similar to major histocompatibility complex class II integral membrane alpha chain gene (LOC691092) mRNA.
<a href="#">LOC691099</a>	<a href="#">LOC691099.bSep08</a>	<a href="#">691099</a>	1099	769	1	70	putative protein (LOC691099) alternative variant bSep08, mRNA.
<a href="#">LOC691107</a>	<a href="#">LOC691107.aSep08</a>	<a href="#">691107</a>	11534	794	3	87	hypothetical protein LOC691107 (9.8 kD) (LOC691107) alternative variant aSep08, mRNA.
<a href="#">LOC691107</a>	<a href="#">LOC691107.bSep08</a>	<a href="#">691107</a>	23954	758	4	33	hypothetical protein LOC691107 (3.9 kD) (LOC691107) alternative variant bSep08, mRNA.
<a href="#">LOC691110</a>	<a href="#">LOC691110.aSep08</a>	<a href="#">691110</a>	2017	792		82	similar to taste receptor protein 1 (8.9 kD) (LOC691110) mRNA.
<a href="#">LOC691124</a>	<a href="#">LOC691124.aSep08</a>	<a href="#">691124</a>	84815	552	2	63	hypothetical protein LOC691124 (LOC691124) alternative variant aSep08, mRNA.
<a href="#">LOC691124</a>	<a href="#">LOC691124.bSep08</a>	<a href="#">691124</a>	119746	1800	2	35	hypothetical protein LOC691124 (4.0 kD) (LOC691124) alternative variant bSep08, mRNA.
<a href="#">LOC691125</a>	<a href="#">LOC691125.aSep08</a>	<a href="#">691125</a>	20151	1785		569	similar to Probable phospholipid-transporting ATPase ID (ATPase class I type 8B member 2) (LOC691125) alternative variant aSep08, mRNA.
<a href="#">LOC691125</a>	<a href="#">LOC691125.bSep08</a>	<a href="#">691125</a>	14740	398		132	similar to Probable phospholipid-transporting ATPase ID (ATPase class I type 8B member 2) (LOC691125) alternative variant bSep08, mRNA.
<a href="#">LOC691135</a>	<a href="#">LOC691135.aSep08</a>	<a href="#">691135</a>	9021	760		222	similar to zinc finger protein 418 (LOC691135) mRNA.
<a href="#">LOC691141</a>	<a href="#">LOC691141.bSep08</a>	<a href="#">691141</a>	18305	804	6	224	hypothetical protein LOC691141 (LOC691141) alternative variant bSep08, mRNA.
<a href="#">LOC691142</a>	<a href="#">LOC691142.aSep08</a>	<a href="#">691142</a>	7039	306		102	hypothetical protein LOC691142 (LOC691142) mRNA.
<a href="#">LOC691143</a>	<a href="#">LOC691143.aSep08</a>	<a href="#">691143</a>	17983	1836	9	474	similar to Serum amyloid A-3 protein precursor (LOC691143) alternative variant aSep08, mRNA.
<a href="#">LOC691143</a>	<a href="#">LOC691143.bSep08</a>	<a href="#">691143</a>	8221	548	2	182	similar to Serum amyloid A-3 protein precursor (LOC691143) alternative variant bSep08, mRNA.
<a href="#">LOC691143</a>	<a href="#">LOC691143.cSep08</a>	<a href="#">691143</a>	7647	715	3	168	similar to Serum amyloid A-3 protein precursor (LOC691143) alternative variant cSep08, mRNA.
<a href="#">LOC691149</a>	<a href="#">LOC691149.bSep08</a>	<a href="#">691149</a>	1202	879	2	54	putative protein (6.1 kD) (LOC691149) alternative variant bSep08, mRNA.
<a href="#">LOC691149</a>	<a href="#">LOC691149.cSep08</a>	<a href="#">691149</a>	25599	874	3	40	protein actin (LOC691149) alternative variant cSep08, mRNA.
<a href="#">LOC691161</a>	<a href="#">LOC691161.aSep08</a>	<a href="#">691161</a>	4162	782		202	hypothetical protein LOC691161 (22.4 kD) (LOC691161) mRNA.
<a href="#">LOC691169</a>	<a href="#">LOC691169.aSep08</a>	<a href="#">691169</a>	22194	1778	5	133	hypothetical protein LOC691169 (LOC691169) alternative variant aSep08, mRNA.

<a href="#">LOC691169</a>	<a href="#">LOC691169.bSep08</a>	<a href="#">691169</a>	9762	970	3	128	hypothetical protein LOC691169 (LOC691169) alternative variant bSep08, mRNA.
<a href="#">LOC691169</a>	<a href="#">LOC691169.cSep08</a>	<a href="#">691169</a>	24710	865	6	95	hypothetical protein LOC691169 (10.5 kD) (LOC691169) alternative variant cSep08, mRNA.
<a href="#">LOC691169</a>	<a href="#">LOC691169.dSep08</a>	<a href="#">691169</a>	8210	362	3	77	hypothetical protein LOC691169 (8.3 kD) (LOC691169) alternative variant dSep08, mRNA.
<a href="#">LOC691178</a>	<a href="#">LOC691178.aSep08</a>	<a href="#">691178</a>	59084	639		118	similar to glutamate receptor, ionotropic, AMPA1 (alpha 1) (LOC691178) mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.aSep08</a>	<a href="#">691221</a>	24036	1561	7	326	similar to CG1998-PA (37.5 kD) (LOC691221) alternative variant aSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.bSep08</a>	<a href="#">691221</a>	22331	1117	8	312	similar to CG1998-PA (LOC691221) alternative variant bSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.cSep08</a>	<a href="#">691221</a>	24841	1974	6	256	similar to CG1998-PA (LOC691221) alternative variant cSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.dSep08</a>	<a href="#">691221</a>	7312	667	5	222	similar to CG1998-PA (LOC691221) alternative variant dSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.eSep08</a>	<a href="#">691221</a>	1808	373	2	123	similar to CG1998-PA (LOC691221) alternative variant eSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.fSep08</a>	<a href="#">691221</a>	22297	521	4	122	similar to CG1998-PA (LOC691221) alternative variant fSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.hSep08</a>	<a href="#">691221</a>	3083	736	2	93	similar to CG1998-PA (9.8 kD) (LOC691221) alternative variant hSep08, mRNA.
<a href="#">LOC691221</a>	<a href="#">LOC691221.iSep08</a>	<a href="#">691221</a>	3103	611	3	42	similar to CG1998-PA (LOC691221) alternative variant iSep08, mRNA.
<a href="#">LOC691222</a>	<a href="#">LOC691222.aSep08</a>	<a href="#">691222</a>	3463	1785		170	similar to ubiquitin carboxyl-terminal hydrolase CYLD (LOC691222) mRNA.
<a href="#">LOC691223</a>	<a href="#">LOC691223.aSep08</a>	<a href="#">691223</a>	3447	298		38	hypothetical protein LOC691223 (LOC691223) mRNA.
<a href="#">LOC691231</a>	<a href="#">LOC691231.aSep08</a>	<a href="#">691231</a>	23557	1904	10	634	similar to gem (nuclear organelle) associated protein 5 (LOC691231) alternative variant aSep08, mRNA.
<a href="#">LOC691231</a>	<a href="#">LOC691231.bSep08</a>	<a href="#">691231</a>	8497	827	7	275	similar to gem (nuclear organelle) associated protein 5 (LOC691231) alternative variant bSep08, mRNA.
<a href="#">LOC691231</a>	<a href="#">LOC691231.cSep08</a>	<a href="#">691231</a>	4704	550	1	120	similar to gem (nuclear organelle) associated protein 5 (LOC691231) alternative variant cSep08, mRNA.
<a href="#">LOC691254</a>	<a href="#">LOC691254.aSep08</a>	<a href="#">691254</a>	29546	549		118	hypothetical protein LOC691254 (LOC691254) mRNA.
<a href="#">LOC691257</a>	<a href="#">LOC691257.aSep08</a>	<a href="#">691257</a>	7193	456	1	142	similar to Zinc finger protein 267 (Zinc finger protein HZF2) (LOC691257) alternative variant aSep08, mRNA.
<a href="#">LOC691257</a>	<a href="#">LOC691257.bSep08</a>	<a href="#">691257</a>	17240	523	4	109	similar to Zinc finger protein 267 (Zinc finger protein HZF2) (LOC691257) alternative variant bSep08, mRNA.
<a href="#">LOC691259</a>	<a href="#">LOC691259.aSep08</a>	<a href="#">691259</a>	10835	1000	5	249	hypothetical protein LOC691259 (26.6 kD) (LOC691259) alternative variant aSep08, mRNA.
<a href="#">LOC691259</a>	<a href="#">LOC691259.cSep08</a>	<a href="#">691259</a>	8041	521	1	38	hypothetical protein LOC691259 (LOC691259) alternative variant cSep08, mRNA.
<a href="#">LOC691289</a>	<a href="#">LOC691289.aSep08</a>	<a href="#">691289</a>	3982	935		33	similar to serine (or cysteine) proteinase inhibitor, clade B, member 1a (LOC691289) mRNA.
<a href="#">LOC691293</a>	<a href="#">LOC691293.aSep08</a>	<a href="#">691293</a>	3035	304		68	reproductive homeobox (LOC691293) mRNA.
<a href="#">LOC691297</a>	<a href="#">LOC691297.aSep08</a>	<a href="#">691297</a>	1508	641		125	hypothetical protein LOC691297 (13.5 kD) (LOC691297) mRNA.

<a href="#">LOC691300</a>	<a href="#">LOC691300.aSep08</a>	<a href="#">691300</a>	53790	450	2	149	hypothetical protein LOC691300 (LOC691300) alternative variant aSep08, mRNA.
<a href="#">LOC691345</a>	<a href="#">LOC691345.aSep08</a>	<a href="#">691345</a>	2232	1032		142	hypothetical protein LOC691345 (15.7 kD) (LOC691345) mRNA.
<a href="#">LOC691354</a>	<a href="#">LOC691354.aSep08</a>	<a href="#">691354</a>	8449	507	3	161	hypothetical protein LOC691354 (LOC691354) alternative variant aSep08, mRNA.
<a href="#">LOC691358</a>	<a href="#">LOC691358.aSep08</a>	<a href="#">691358</a>	1096	399		74	hypothetical protein LOC691358 (LOC691358) mRNA.
<a href="#">LOC691375</a>	<a href="#">LOC691375.aSep08</a>	<a href="#">691375</a>	19481	644		168	similar to serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2 (LOC691375) mRNA.
<a href="#">LOC691396</a>	<a href="#">LOC691396.aSep08</a>	<a href="#">691396</a>	2456	980	1	171	similar to Zinc finger protein 551 (Zinc finger protein KOX23) (LOC691396) alternative variant aSep08, mRNA.
<a href="#">LOC691396</a>	<a href="#">LOC691396.bSep08</a>	<a href="#">691396</a>	12343	542	1	33	similar to Zinc finger protein 551 (Zinc finger protein KOX23) (LOC691396) alternative variant bSep08, mRNA.
<a href="#">LOC691422</a>	<a href="#">LOC691422.aSep08</a>	<a href="#">691422</a>	1314	1122		31	similar to zinc finger protein 101 (LOC691422) mRNA.
<a href="#">LOC691426</a>	<a href="#">LOC691426.aSep08</a>	<a href="#">691426</a>	3734	710	4	236	similar to CG7744-PA (LOC691426) alternative variant aSep08, mRNA.
<a href="#">LOC691427</a>	<a href="#">LOC691427.aSep08</a>	<a href="#">691427</a>	5783	649	4	60	similar to 6.8 kDa mitochondrial proteolipid (6.9 kD) (LOC691427) alternative variant aSep08, mRNA.
<a href="#">LOC691431</a>	<a href="#">LOC691431.bSep08</a>	<a href="#">691431</a>	6810	844	5	215	similar to mitochondrial carrier protein MGC4399 (LOC691431) alternative variant bSep08, mRNA.
<a href="#">LOC691431</a>	<a href="#">LOC691431.cSep08</a>	<a href="#">691431</a>	6592	666	5	212	similar to mitochondrial carrier protein MGC4399 (LOC691431) alternative variant cSep08, mRNA.
<a href="#">LOC691431</a>	<a href="#">LOC691431.dSep08</a>	<a href="#">691431</a>	2460	1103	3	168	similar to mitochondrial carrier protein MGC4399 (LOC691431) alternative variant dSep08, mRNA.
<a href="#">LOC691452</a>	<a href="#">LOC691452.aSep08</a>	<a href="#">691452</a>	64761	710	3	236	hypothetical protein LOC691452 (LOC691452) alternative variant aSep08, mRNA.
<a href="#">LOC691452</a>	<a href="#">LOC691452.bSep08</a>	<a href="#">691452</a>	64741	681	3	226	hypothetical protein LOC691452 (LOC691452) alternative variant bSep08, mRNA.
<a href="#">LOC691452</a>	<a href="#">LOC691452.cSep08</a>	<a href="#">691452</a>	30397	665	1	105	hypothetical protein LOC691452 (LOC691452) alternative variant cSep08, mRNA.
<a href="#">LOC691468</a>	<a href="#">LOC691468.aSep08</a>	<a href="#">691468</a>	12592	710		236	similar to Zinc finger protein 84 (Zinc finger protein HPF2) (LOC691468) mRNA.
<a href="#">LOC691504</a>	<a href="#">LOC691504.aSep08</a>	<a href="#">691504</a>	52310	579		182	similar to Zinc finger protein ZFPM1 (Zinc finger protein multitype 1) (Friend of GATA protein 1) (Friend of GATA-1) (FOG-1) (LOC691504) mRNA.
<a href="#">LOC691509</a>	<a href="#">LOC691509.aSep08</a>	<a href="#">691509</a>	22977	1206		142	hypothetical protein LOC691509 (16.0 kD) (LOC691509) mRNA.
<a href="#">LOC691519</a>	<a href="#">LOC691519.aSep08</a>	<a href="#">691519</a>	8276	451		70	putative protein, with 2 coiled coil domains (LOC691519) mRNA.
<a href="#">LOC691521</a>	<a href="#">LOC691521.aSep08</a>	<a href="#">691521</a>	10645	543	5	180	similar to MEGF11 protein (LOC691521) alternative variant aSep08, mRNA.
<a href="#">LOC691521</a>	<a href="#">LOC691521.bSep08</a>	<a href="#">691521</a>	9441	935	5	163	similar to MEGF11 protein (LOC691521) alternative variant bSep08, mRNA.
<a href="#">LOC691551</a>	<a href="#">LOC691551.aSep08</a>	<a href="#">691551</a>	1511	298		52	similar to F28B3.5a (LOC691551) mRNA.
<a href="#">LOC691556</a>	<a href="#">LOC691556.bSep08</a>	<a href="#">691556</a>	2533	1928	1	385	similar to zinc finger protein 238 (LOC691556) alternative variant bSep08, mRNA.

<a href="#">LOC691565</a>	<a href="#">LOC691565.aSep08</a>	<a href="#">691565</a>	33750	388		84	hypothetical protein LOC691565 (LOC691565) mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.aSep08</a>	<a href="#">691572</a>	11621	818	4	177	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (LOC691572) alternative variant aSep08, mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.aSep08</a>	<a href="#">691574</a>	11621	818	4	177	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (LOC691572) alternative variant aSep08, mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.bSep08</a>	<a href="#">691572</a>	9942	650	3	162	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (LOC691572) alternative variant bSep08, mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.bSep08</a>	<a href="#">691574</a>	9942	650	3	162	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (LOC691572) alternative variant bSep08, mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.cSep08</a>	<a href="#">691572</a>	989	672	2	54	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (6.4 kD) (LOC691572) alternative variant cSep08, mRNA.
<a href="#">LOC691572</a>	<a href="#">LOC691572.cSep08</a>	<a href="#">691574</a>	989	672	2	54	hypothetical protein LOC691572 and similar to ubiquitin-associated protein 1 (6.4 kD) (LOC691572) alternative variant cSep08, mRNA.
<a href="#">LOC691586</a>	<a href="#">LOC691586.aSep08</a>	<a href="#">691586</a>	2111	635		105	similar to Integrin alpha-6 precursor (VLA-6) (CD49f antigen) (LOC691586) mRNA.
<a href="#">LOC691595</a>	<a href="#">LOC691595.aSep08</a>	<a href="#">691595</a>	2137	666		117	hypothetical protein LOC691595 (LOC691595) mRNA.
<a href="#">LOC691600</a>	<a href="#">LOC691600.aSep08</a>	<a href="#">691600</a>	940	271		89	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC691600) mRNA.
<a href="#">LOC691608</a>	<a href="#">LOC691608.aSep08</a>	<a href="#">691608</a>	10981	407	4	72	hypothetical protein LOC691608 (LOC691608) alternative variant aSep08, mRNA.
<a href="#">LOC691608</a>	<a href="#">LOC691608.bSep08</a>	<a href="#">691608</a>	6543	512	2	70	hypothetical protein LOC691608 (LOC691608) alternative variant bSep08, mRNA.
<a href="#">LOC691608</a>	<a href="#">LOC691608.cSep08</a>	<a href="#">691608</a>	3369	384	1	66	hypothetical protein LOC691608 (LOC691608) alternative variant cSep08, mRNA.
<a href="#">LOC691608</a>	<a href="#">LOC691608.dSep08</a>	<a href="#">691608</a>	19555	670	5	69	hypothetical protein LOC691608 (LOC691608) alternative variant dSep08, mRNA.
<a href="#">LOC691627</a>	<a href="#">LOC691627.aSep08</a>	<a href="#">691627</a>	1281	565		38	similar to spermatogenesis associated glutamate (E)-rich protein 4d (LOC691627) mRNA.
<a href="#">LOC691631</a>	<a href="#">LOC691631.aSep08</a>	<a href="#">691631</a>	47742	1866	2	545	hypothetical protein LOC691631 (LOC691631) alternative variant aSep08, mRNA.
<a href="#">LOC691632</a>	<a href="#">LOC691632.aSep08</a>	<a href="#">691632</a>	6999	402		92	similar to MIC2 like 1 (LOC691632) mRNA.
<a href="#">LOC691636</a>	<a href="#">LOC691636.aSep08</a>	<a href="#">691636</a>	13107	545		163	putative protein of mammalian origin (LOC691636) mRNA.
<a href="#">LOC691649</a>	<a href="#">LOC691649.aSep08</a>	<a href="#">691649</a>	120105	2008	6	282	hypothetical protein LOC691649 (31.2 kD) (LOC691649) alternative variant aSep08, mRNA.
<a href="#">LOC691652</a>	<a href="#">LOC691652.aSep08</a>	<a href="#">691652</a>	17914	624		74	hypothetical protein LOC691652 (LOC691652) mRNA.
<a href="#">LOC691653</a>	<a href="#">LOC691653.aSep08</a>	<a href="#">691653</a>	20150	941	4	101	hypothetical protein LOC691653 and hypothetical protein LOC691659 (11.4 kD) (LOC691653) alternative variant aSep08, mRNA.
<a href="#">LOC691653</a>	<a href="#">LOC691653.aSep08</a>	<a href="#">691659</a>	20150	941	4	101	hypothetical protein LOC691653 and hypothetical protein LOC691659 (11.4 kD) (LOC691653) alternative variant aSep08, mRNA.



<a href="#">LOC691653</a>	<a href="#">LOC691653.bSep08</a>	<a href="#">691653</a>	4329	308	1	50	hypothetical protein LOC691653 and hypothetical protein LOC691659 (LOC691653) alternative variant bSep08, mRNA.
<a href="#">LOC691653</a>	<a href="#">LOC691653.bSep08</a>	<a href="#">691659</a>	4329	308	1	50	hypothetical protein LOC691653 and hypothetical protein LOC691659 (LOC691653) alternative variant bSep08, mRNA.
<a href="#">LOC691657</a>	<a href="#">LOC691657.aSep08</a>	<a href="#">691657</a>	7872	463	2	115	similar to Cysteine-rich protein 1 (Cysteine-rich intestinal protein) (CRIP) (LOC691657) alternative variant aSep08, mRNA.
<a href="#">LOC691657</a>	<a href="#">LOC691657.bSep08</a>	<a href="#">691657</a>	1933	539	1	77	similar to Cysteine-rich protein 1 (Cysteine-rich intestinal protein) (CRIP) (8.5 kD) (LOC691657) alternative variant bSep08, mRNA.
<a href="#">LOC691658</a>	<a href="#">LOC691658.aSep08</a>	<a href="#">691658</a>	17578	1013		300	similar to anterior pharynx defective 1b homolog (LOC691658) alternative variant aSep08, mRNA.
<a href="#">LOC691658</a>	<a href="#">LOC691658.bSep08</a>	<a href="#">691658</a>	3788	1098		96	similar to anterior pharynx defective 1b homolog (11.1 kD) (LOC691658) alternative variant bSep08, mRNA.
<a href="#">LOC691661</a>	<a href="#">LOC691661.aSep08</a>	<a href="#">691661</a>	33362	768		42	putative protein (LOC691661) mRNA.
<a href="#">LOC691672</a>	<a href="#">LOC691672.aSep08</a>	<a href="#">691672</a>	5838	1017		77	similar to Discs large homolog 5 (Placenta and prostate DLG) (Discs large protein P-dlg) (LOC691672) mRNA.
<a href="#">LOC691673</a>	<a href="#">LOC691673.aSep08</a>	<a href="#">691673</a>	18646	618		163	hypothetical protein LOC691673 (LOC691673) mRNA.
<a href="#">LOC691687</a>	<a href="#">LOC691687.bSep08</a>	<a href="#">691687</a>	15663	765	3	102	hypothetical protein LOC691687 (10.3 kD) (LOC691687) alternative variant bSep08, mRNA.
<a href="#">LOC691693</a>	<a href="#">LOC691693.aSep08</a>	<a href="#">691693</a>	23114	250		54	ankyrin (LOC691693) mRNA.
<a href="#">LOC691722</a>	<a href="#">LOC691722.aSep08</a>	<a href="#">691722</a>	32576	752	8	69	hypothetical protein LOC691722 (LOC691722) alternative variant aSep08, mRNA.
<a href="#">LOC691742</a>	<a href="#">LOC691742.aSep08</a>	<a href="#">691742</a>	18907	1091		52	hypothetical protein LOC691742 (LOC691742) mRNA.
<a href="#">LOC691750</a>	<a href="#">LOC691750.aSep08</a>	<a href="#">691750</a>	2106	789	5	145	hypothetical protein LOC691750 (LOC691750) alternative variant aSep08, mRNA.
<a href="#">LOC691750</a>	<a href="#">LOC691750.cSep08</a>	<a href="#">691750</a>	1425	417	2	61	hypothetical protein LOC691750 (LOC691750) alternative variant cSep08, mRNA.
<a href="#">LOC691759</a>	<a href="#">LOC691759.aSep08</a>	<a href="#">691759</a>	2150	420	3	77	hypothetical protein LOC691759 (8.0 kD) (LOC691759) alternative variant aSep08, mRNA.
<a href="#">LOC691759</a>	<a href="#">LOC691759.bSep08</a>	<a href="#">691759</a>	2053	270	2	67	hypothetical protein LOC691759 (LOC691759) alternative variant bSep08, mRNA.
<a href="#">LOC691762</a>	<a href="#">LOC691762.aSep08</a>	<a href="#">691762</a>	398137	618		163	hypothetical protein LOC691762 and hypothetical protein LOC691795 (LOC691762) mRNA.
<a href="#">LOC691762</a>	<a href="#">LOC691762.aSep08</a>	<a href="#">691795</a>	398137	618		163	hypothetical protein LOC691762 and hypothetical protein LOC691795 (LOC691762) mRNA.
<a href="#">LOC691773</a>	<a href="#">LOC691773.aSep08</a>	<a href="#">691773</a>	2764	838		174	hypothetical protein LOC691773 (LOC691773) mRNA.
<a href="#">LOC691777</a>	<a href="#">LOC691777.aSep08</a>	<a href="#">691777</a>	13441	328		109	hypothetical protein LOC691777 (LOC691777) mRNA.
<a href="#">LOC691809</a>	<a href="#">LOC691809.aSep08</a>	<a href="#">691809</a>	63096	413		84	hypothetical protein LOC691809 (LOC691809) mRNA.
<a href="#">LOC691813</a>	<a href="#">LOC691813.aSep08</a>	<a href="#">691813</a>	5489	305		101	similar to phospholipase A2, group IVC (cytosolic, calcium-independent) (LOC691813) mRNA.
<a href="#">LOC691817</a>	<a href="#">LOC691817.aSep08</a>	<a href="#">691817</a>	22984	782		88	hypothetical protein LOC691817 (LOC691817) mRNA.

<a href="#">LOC691842</a>	<a href="#">LOC691842.bSep08</a>	<a href="#">691842</a>	3036	1437	5	267	similar to Myc-associated zinc finger protein (MAZI) (Purine-binding transcription factor) (Pur-1) (LOC691842) alternative variant bSep08, mRNA.
<a href="#">LOC691842</a>	<a href="#">LOC691842.cSep08</a>	<a href="#">691842</a>	3542	789	4	140	similar to Myc-associated zinc finger protein (MAZI) (Purine-binding transcription factor) (Pur-1) (LOC691842) alternative variant cSep08, mRNA.
<a href="#">LOC691842</a>	<a href="#">LOC691842.eSep08</a>	<a href="#">691842</a>	3550	2101	3	78	similar to Myc-associated zinc finger protein (MAZI) (Purine-binding transcription factor) (Pur-1) (7.6 kD) (LOC691842) alternative variant eSep08, mRNA.
<a href="#">LOC691849</a>	<a href="#">LOC691849.aSep08</a>	<a href="#">691849</a>	5328	1598		123	hypothetical protein LOC691849 (13.8 kD) (LOC691849) mRNA.
<a href="#">LOC691853</a>	<a href="#">LOC691853.aSep08</a>	<a href="#">691853</a>	28450	495		164	similar to COX10 homolog, cytochrome c oxidase assembly protein, heme A: farnesyltransferase (LOC691853) mRNA.
<a href="#">LOC691862</a>	<a href="#">LOC691862.aSep08</a>	<a href="#">691862</a>	7110	1279		27	hypothetical protein LOC691862 (LOC691862) mRNA.
<a href="#">LOC691887</a>	<a href="#">LOC691887.aSep08</a>	<a href="#">691887</a>	1152	493		164	similar to zinc finger protein HIT-39 (LOC691887) mRNA.
<a href="#">LOC691889</a>	<a href="#">LOC691889.aSep08</a>	<a href="#">691889</a>	126868	692		230	similar to ATPase, aminophospholipid transporter-like, class I, type 8A, member 2 (LOC691889) mRNA.
<a href="#">LOC691904</a>	<a href="#">LOC691904.bSep08</a>	<a href="#">691904</a>	3799	400		99	hypothetical protein LOC691904 (LOC691904) alternative variant bSep08, mRNA.
<a href="#">LOC691911</a>	<a href="#">LOC691911.aSep08</a>	<a href="#">691911</a>	7869	402		133	similar to Ciliary dynein heavy chain 9 (Axonemal beta dynein heavy chain 9) (LOC691911) mRNA.
<a href="#">LOC691914</a>	<a href="#">LOC691914.aSep08</a>	<a href="#">691914</a>	8289	652		216	similar to Leo1, Paf1/RNA polymerase II complex component, homolog (LOC691914) mRNA.
<a href="#">LOC691921</a>	<a href="#">LOC691921.aSep08</a>	<a href="#">691921</a>	3085	337	3	85	hypothetical protein LOC691921 (LOC691921) alternative variant aSep08, mRNA.
<a href="#">LOC691921</a>	<a href="#">LOC691921.bSep08</a>	<a href="#">691921</a>	2916	428	2	67	hypothetical protein LOC691921 (7.6 kD) (LOC691921) alternative variant bSep08, mRNA.
<a href="#">LOC691923</a>	<a href="#">LOC691923.aSep08</a>	<a href="#">691923</a>	52312	653		65	hypothetical protein LOC691923 (LOC691923) mRNA.
<a href="#">LOC691926</a>	<a href="#">LOC691926.aSep08</a>	<a href="#">691926</a>	9173	853		284	similar to Stabilin-2 precursor (Hyaluronan receptor for endocytosis) (LOC691926) mRNA.
<a href="#">LOC691928</a>	<a href="#">LOC691928.aSep08</a>	<a href="#">691928</a>	39293	662	5	33	hypothetical protein LOC691928 (3.8 kD) (LOC691928) alternative variant aSep08, mRNA.
<a href="#">LOC691932</a>	<a href="#">LOC691932.aSep08</a>	<a href="#">691932</a>	697	419		72	similar to transglutaminase 7 (LOC691932) mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.aSep08</a>	<a href="#">691933</a>	30939	1452	3	484	similar to Protein C6orf142 homolog (LOC691933) alternative variant aSep08, mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.bSep08</a>	<a href="#">691933</a>	101129	947	8	315	similar to Protein C6orf142 homolog (LOC691933) alternative variant bSep08, mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.cSep08</a>	<a href="#">691933</a>	82319	666	4	221	similar to Protein C6orf142 homolog (LOC691933) alternative variant cSep08, mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.dSep08</a>	<a href="#">691933</a>	74471	591	4	196	similar to Protein C6orf142 homolog (LOC691933) alternative variant dSep08, mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.eSep08</a>	<a href="#">691933</a>	71900	586	5	195	similar to Protein C6orf142 homolog (LOC691933) alternative variant eSep08, mRNA.
<a href="#">LOC691933</a>	<a href="#">LOC691933.fSep08</a>	<a href="#">691933</a>	101147	683	5	167	similar to Protein C6orf142 homolog (LOC691933) alternative variant fSep08, mRNA.

<a href="#">LOC691933</a>	<a href="#">LOC691933.gSep08</a>	<a href="#">691933</a>	27083	420	6	140	similar to Protein C6orf142 homolog (LOC691933) alternative variant gSep08, mRNA.
<a href="#">LOC691951</a>	<a href="#">LOC691951.aSep08</a>	<a href="#">691951</a>	36290	404	1	70	hypothetical protein LOC691951 (LOC691951) alternative variant aSep08, mRNA.
<a href="#">LOC691951</a>	<a href="#">LOC691951.bSep08</a>	<a href="#">691951</a>	8607	557	1	47	hypothetical protein LOC691951 (LOC691951) alternative variant bSep08, mRNA.
<a href="#">LOC691960</a>	<a href="#">LOC691960.aSep08</a>	<a href="#">691960</a>	18664	1817	14	536	similar to solute carrier family 28, member 2 (LOC691960) alternative variant aSep08, mRNA.
<a href="#">LOC691962</a>	<a href="#">LOC691962.aSep08</a>	<a href="#">691962</a>	72455	764		150	hypothetical protein LOC691962 (17.2 kD) (LOC691962) mRNA.
<a href="#">LOC691966</a>	<a href="#">LOC691966.aSep08</a>	<a href="#">691966</a>	8977	1107		297	similar to Sulfide:quinone oxidoreductase, mitochondrial precursor (LOC691966) mRNA.
<a href="#">LOC691969</a>	<a href="#">LOC691969.aSep08</a>	<a href="#">691969</a>	9592	1476		344	similar to deleted in malignant brain tumors 1 isoform a precursor (38.7 kD) (LOC691969) mRNA.
<a href="#">LOC691979</a>	<a href="#">LOC691979.aSep08</a>	<a href="#">691979</a>	16224	2109		619	similar to N-acetyltransferase ESCO2 (Establishment of cohesion 1 homolog 2) (ECO1 homolog 2) (LOC691979) mRNA.
<a href="#">LOC691984</a>	<a href="#">LOC691984.aSep08</a>	<a href="#">691984</a>	421847	697	3	232	similar to Glypican-6 precursor (LOC691984) alternative variant aSep08, mRNA.
<a href="#">LOC691988</a>	<a href="#">LOC691988.aSep08</a>	<a href="#">691988</a>	19875	446		116	similar to a disintegrin and metalloproteinase domain 28 (LOC691988) mRNA.
<a href="#">LOC692032</a>	<a href="#">LOC692032.aSep08</a>	<a href="#">692032</a>	10575	2576	1	78	hypothetical protein LOC692032 (8.1 kD) (LOC692032) alternative variant aSep08, mRNA.
<a href="#">LOC692032</a>	<a href="#">LOC692032.bSep08</a>	<a href="#">692032</a>	6911	761	1	59	hypothetical protein LOC692032 (LOC692032) alternative variant bSep08, mRNA.
<a href="#">LOC692032</a>	<a href="#">LOC692032.cSep08</a>	<a href="#">692032</a>	8719	701	1	41	hypothetical protein LOC692032 (4.4 kD) (LOC692032) alternative variant cSep08, mRNA.
<a href="#">LOC100125368</a>	<a href="#">LOC100125368.aSep08</a>	<a href="#">100125368</a>	22365	1960		501	zinc finger protein LOC100125368 (58.2 kD) (LOC100125368) mRNA.
<a href="#">LOC100125371</a>	<a href="#">LOC100125371.aSep08</a>	<a href="#">100125371</a>	5349	2817	3	73	hypothetical LOC100125371 (8.3 kD) (LOC100125371) alternative variant aSep08, mRNA.
<a href="#">LOC100125386</a>	<a href="#">LOC100125386.aSep08</a>	<a href="#">100125386</a>	10099	1324		134	hypothetical LOC100125386 (15.3 kD) (LOC100125386) mRNA.
<a href="#">LOC100151767</a>	<a href="#">LOC100151767.aSep08</a>	<a href="#">100151767</a>	11143	3549	6	291	hypothetical LOC100151767 (33.7 kD) (LOC100151767) alternative variant aSep08, complete mRNA.
<a href="#">LOC100151767</a>	<a href="#">LOC100151767.cSep08</a>	<a href="#">100151767</a>	5977	398	3	98	hypothetical LOC100151767 (LOC100151767) alternative variant cSep08, mRNA.
<a href="#">LOC100158225</a>	<a href="#">LOC100158225.bSep08</a>	<a href="#">100158225</a>	1820	1015	5	195	hypothetical protein LOC100158225 (LOC100158225) alternative variant bSep08, mRNA.
<a href="#">LOC100174909</a>	<a href="#">LOC100174909.bSep08</a>	<a href="#">100174909</a>	9841	490	3	98	hypothetical LOC100174909 (LOC100174909) alternative variant bSep08, mRNA.
<a href="#">LOC100174909</a>	<a href="#">LOC100174909.cSep08</a>	<a href="#">100174909</a>	5698	521	2	91	hypothetical LOC100174909 (LOC100174909) alternative variant cSep08, mRNA.
<a href="#">LOC100174909</a>	<a href="#">LOC100174909.dSep08</a>	<a href="#">100174909</a>	6845	1614	2	55	hypothetical LOC100174909 (6.5 kD) (LOC100174909) alternative variant dSep08, mRNA.

<a href="#">LOC100188932</a>	<a href="#">LOC100188932.aSep08</a>	<a href="#">100188932</a>	3506	1164	3	110	dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 4 (12.9 kD) (LOC100188932) alternative variant aSep08, mRNA.
<a href="#">LOC100188933</a>	<a href="#">LOC100188933.aSep08</a>	<a href="#">100188933</a>	11687	449	1	80	hypothetical protein LOC100188933 (8.8 kD) (LOC100188933) alternative variant aSep08, mRNA.
<a href="#">LOC100188936</a>	<a href="#">LOC100188936.aSep08</a>	<a href="#">100188936</a>	4992	697	3	214	hypothetical protein LOC100188936 (LOC100188936) alternative variant aSep08, mRNA.
<a href="#">lochy</a>	<a href="#">lochy.aSep08</a>		13815	772		27	putative protein (lochy) mRNA.
<a href="#">lodar</a>	<a href="#">lodar.aSep08</a>		1518	450		42	putative protein (lodar) mRNA.
<a href="#">lofer</a>	<a href="#">lofer.aSep08</a>		3247	199		63	gag protein like (lofer) mRNA.
<a href="#">loflo</a>	<a href="#">loflo.aSep08</a>		38674	680		67	putative protein (7.7 kD) (loflo) mRNA.
<a href="#">loflu</a>	<a href="#">loflu.aSep08</a>		8630	719		239	neuron navigator 2 CRA c (loflu) mRNA.
<a href="#">Loh11cr2a</a>	<a href="#">Loh11cr2a.bSep08</a>	<a href="#">301097</a>	7104	2309	7	288	loss of heterozygosity, 11, chromosomal region 2, gene A homolog (human) (Loh11cr2a) alternative variant bSep08, mRNA.
<a href="#">Loh11cr2a</a>	<a href="#">Loh11cr2a.cSep08</a>	<a href="#">301097</a>	4393	525	5	116	loss of heterozygosity, 11, chromosomal region 2, gene A homolog (human) (Loh11cr2a) alternative variant cSep08, mRNA.
<a href="#">Loh11cr2a</a>	<a href="#">Loh11cr2a.eSep08</a>	<a href="#">301097</a>	1290	390	2	55	loss of heterozygosity, 11, chromosomal region 2, gene A homolog (human) (Loh11cr2a) alternative variant eSep08, mRNA.
<a href="#">Loh12cr1</a>	<a href="#">Loh12cr1.cSep08</a>	<a href="#">362452</a>	59837	1047	5	91	loss of heterozygosity, 12, chromosomal region 1 homolog (human) (10.1 kD) (Loh12cr1) alternative variant cSep08, mRNA.
<a href="#">lojey</a>	<a href="#">lojey.aSep08</a>		4663	628		36	putative protein (4.1 kD) (lojey) mRNA.
<a href="#">lokee</a>	<a href="#">lokee.aSep08</a>		2798	400		133	putative protein (lokee) mRNA.
<a href="#">loloy</a>	<a href="#">loloy.aSep08</a>		2605	417		97	putative protein (loloy) mRNA.
<a href="#">lomee</a>	<a href="#">lomee.aSep08</a>		2516	708		41	putative protein (4.3 kD) (lomee) mRNA.
<a href="#">lomer</a>	<a href="#">lomer.aSep08</a>		697	248		38	ATP-binding cassette sub-family A member like (lomer) mRNA.
<a href="#">lonoy</a>	<a href="#">lonoy.aSep08</a>		677	419		135	putative protein of mammalian origin (lonoy) mRNA.
<a href="#">Lonp1</a>	<a href="#">Lonp1.bSep08</a>	<a href="#">170916</a>	2050	721	5	204	lon peptidase 1, mitochondrial (Lonp1) alternative variant bSep08, mRNA.
<a href="#">Lonp1</a>	<a href="#">Lonp1.cSep08</a>	<a href="#">170916</a>	404	320	2	106	lon peptidase 1, mitochondrial (Lonp1) alternative variant cSep08, mRNA.
<a href="#">Lonp2</a>	<a href="#">Lonp2.aSep08</a>	<a href="#">291922</a>	92901	2829	15	852	peroxisomal lon (94.4 kD) (Lonp2) alternative variant aSep08, mRNA.
<a href="#">Lonp2</a>	<a href="#">Lonp2.cSep08</a>	<a href="#">291922</a>	7575	757	4	211	CRA c (Lonp2) alternative variant cSep08, mRNA.
<a href="#">Lonrf1</a>	<a href="#">Lonrf1.aSep08</a>	<a href="#">306505</a>	15971	444	2	147	LON peptidase N-terminal domain and ring finger 1 (Lonrf1) alternative variant aSep08, mRNA.
<a href="#">Lonrf1</a>	<a href="#">Lonrf1.bSep08</a>	<a href="#">306505</a>	5304	346	1	115	LON peptidase N-terminal domain and ring finger 1 (Lonrf1) alternative variant bSep08, mRNA.
<a href="#">Lonrf2</a>	<a href="#">Lonrf2.aSep08</a>	<a href="#">301361</a>	3746	432		110	LON peptidase N-terminal domain and ring finger 2 (Lonrf2) mRNA.
<a href="#">Lonrf3</a>	<a href="#">Lonrf3.aSep08</a>	<a href="#">298322</a>	15464	770		256	LON peptidase N-terminal domain and ring finger 3 (Lonrf3) mRNA.

<a href="#">lopor</a>	<a href="#">lopor.aSep08</a>		6829	698		142	putative protein (15.5 kD) (lopor) mRNA.
<a href="#">lorby</a>	<a href="#">lorby.aSep08</a>		9165	3243		253	WWC family member 3 like (lorby) mRNA.
<a href="#">lorchy</a>	<a href="#">lorchy.aSep08</a>		4514	278		92	titin (lorchy) mRNA.
<a href="#">lordar</a>	<a href="#">lordar.aSep08</a>		936	381		19	putative protein (lordar) mRNA.
<a href="#">lorfer</a>	<a href="#">lorfer.aSep08</a>		24384	329		109	nidogen (lorfer) mRNA.
<a href="#">lorflo</a>	<a href="#">lorflo.aSep08</a>		1179	694		55	putative protein (6.1 kD) (lorflo) mRNA.
<a href="#">lorflu</a>	<a href="#">lorflu.aSep08</a>		12599	464		89	putative protein (lorflu) mRNA.
<a href="#">lorgar</a>	<a href="#">lorgar.aSep08</a>		72054	397	1	48	putative protein (lorgar) alternative variant aSep08, mRNA.
<a href="#">lorgar</a>	<a href="#">lorgar.bSep08</a>		12493	327	1	42	putative protein (4.4 kD) (lorgar) alternative variant bSep08, mRNA.
<a href="#">lorjey</a>	<a href="#">lorjey.aSep08</a>		12176	624		207	polycystin (lorjey) mRNA.
<a href="#">lorkee</a>	<a href="#">lorkee.aSep08</a>		113831	442		99	putative nuclear protein (11.3 kD) (lorkee) mRNA.
<a href="#">lorloy</a>	<a href="#">lorloy.aSep08</a>		2774	1120		101	ATPase 13a5 (11.8 kD) (lorloy) mRNA.
<a href="#">lormee</a>	<a href="#">lormee.aSep08</a>		1369	613		192	putative protein, with a coiled coil domain, of mammalian origin (lormee) mRNA.
<a href="#">lormer</a>	<a href="#">lormer.aSep08</a>		2013	556		36	putative protein (4.0 kD) (lormer) mRNA.
<a href="#">lornoy</a>	<a href="#">lornoy.aSep08</a>		37241	325		62	putative protein (lornoy) mRNA.
<a href="#">lorpor</a>	<a href="#">lorpor.aSep08</a>		1946	309		63	putative protein (lorpor) mRNA.
<a href="#">lorsa</a>	<a href="#">lorsa.aSep08</a>		33336	459		21	putative protein (lorsa) mRNA.
<a href="#">lorshee</a>	<a href="#">lorshee.aSep08</a>		2260	660		38	putative protein (lorshee) mRNA.
<a href="#">lortu</a>	<a href="#">lortu.aSep08</a>		962	339		87	putative protein (lortu) mRNA.
<a href="#">lorvar</a>	<a href="#">lorvar.aSep08</a>		15435	1464	10	452	polycystic kidney disease like (lorvar) alternative variant aSep08, mRNA.
<a href="#">lorvar</a>	<a href="#">lorvar.bSep08</a>		2579	1318	3	156	polycystic kidney disease 1-like (17.3 kD) (lorvar) alternative variant bSep08, mRNA.
<a href="#">lorvar</a>	<a href="#">lorvar.cSep08</a>		6105	888	4	109	polycystic kidney disease like (lorvar) alternative variant cSep08, mRNA.
<a href="#">lorvar</a>	<a href="#">lorvar.dSep08</a>		2767	438	4	95	polycystic kidney disease 1-like (lorvar) alternative variant dSep08, mRNA.
<a href="#">lorwey</a>	<a href="#">lorwey.aSep08</a>		57096	904		300	inositol 1 receptor CRA a (lorwey) mRNA.
<a href="#">losa</a>	<a href="#">losa.aSep08</a>		83806	1281	8	426	trafficking protein particle complex 9 (losa) alternative variant aSep08, mRNA.
<a href="#">losa</a>	<a href="#">losa.bSep08</a>		16798	555	5	184	trafficking protein particle complex 9 (losa) alternative variant bSep08, mRNA.
<a href="#">losa</a>	<a href="#">losa.cSep08</a>		16720	720	4	143	trafficking protein particle complex 9 (losa) alternative variant cSep08, mRNA.
<a href="#">loshee</a>	<a href="#">loshee.aSep08</a>		4042	494		164	putative protein of metazoan origin (loshee) mRNA.
<a href="#">lotu</a>	<a href="#">lotu.aSep08</a>		1046	452		32	putative protein (3.5 kD) (lotu) mRNA.
<a href="#">lovar</a>	<a href="#">lovar.aSep08</a>		665	377		61	putative protein (lovar) mRNA.
<a href="#">lowey</a>	<a href="#">lowey.aSep08</a>		3946	385		23	putative protein (lowey) mRNA.
<a href="#">Loxl1</a>	<a href="#">Loxl1.bSep08</a>	<a href="#">315714</a>	7840	768	4	42	lysyl oxidase-like 1 (4.8 kD) (Loxl1) alternative variant bSep08, mRNA.
<a href="#">Loxl2</a>	<a href="#">Loxl2.aSep08</a>	<a href="#">290350</a>	89019	4784	14	750	lysyl oxidase-like 2 CRA b (Loxl2) alternative variant aSep08, mRNA.

<a href="#">Loxl2</a>	<a href="#">Loxl2.bSep08</a>	<a href="#">290350</a>	33906	1953	12	466	lysyl oxidase-like 2 CRA b (Loxl2) alternative variant bSep08, mRNA.
<a href="#">Loxl2</a>	<a href="#">Loxl2.dSep08</a>	<a href="#">290350</a>	26486	393	2	70	putative protein (Loxl2) alternative variant dSep08, mRNA.
<a href="#">Loxl2</a>	<a href="#">Loxl2.eSep08</a>	<a href="#">290350</a>	1127	517	2	45	lysyl oxidase-like 2 (5.4 kD) (Loxl2) alternative variant eSep08, mRNA.
<a href="#">Loxl3</a>	<a href="#">Loxl3.bSep08</a>	<a href="#">312478</a>	800	613	3	133	lysyl oxidase-like 3 (Loxl3) alternative variant bSep08, mRNA.
<a href="#">Loxl3</a>	<a href="#">Loxl3.cSep08</a>	<a href="#">312478</a>	1483	951	4	111	lysyl oxidase-like 3 (12.9 kD) (Loxl3) alternative variant cSep08, mRNA.
<a href="#">loyby</a>	<a href="#">loyby.aSep08</a>		489485	852	1	283	rho GTPase activating protein 6 (loyby) alternative variant aSep08, mRNA.
<a href="#">loyby</a>	<a href="#">loyby.bSep08</a>		216582	622	1	181	rho GTPase activating protein 6 (loyby) alternative variant bSep08, mRNA.
<a href="#">loychy</a>	<a href="#">loychy.aSep08</a>		888	246		82	titin (loychy) mRNA.
<a href="#">loydar</a>	<a href="#">loydar.aSep08</a>		9076	558		40	putative protein (4.4 kD) (loydar) mRNA.
<a href="#">loyfer</a>	<a href="#">loyfer.cSep08</a>		1587	761	2	65	putative protein (7.2 kD) (loyfer) alternative variant cSep08, mRNA.
<a href="#">loyflo</a>	<a href="#">loyflo.aSep08</a>		2897	443		77	putative protein (loyflo) mRNA.
<a href="#">loyflu</a>	<a href="#">loyflu.aSep08</a>		21287	559		105	putative mitochondrial protein (11.4 kD) (loyflu) mRNA.
<a href="#">loygar</a>	<a href="#">loygar.aSep08</a>		5889	757	3	118	putative protein (loygar) alternative variant aSep08, mRNA.
<a href="#">loyjey</a>	<a href="#">loyjey.aSep08</a>		670	401	1	118	putative protein (loyjey) alternative variant aSep08, mRNA.
<a href="#">loyjey</a>	<a href="#">loyjey.bSep08</a>		11371	754	2	48	putative protein (5.2 kD) (loyjey) alternative variant bSep08, mRNA.
<a href="#">loykee</a>	<a href="#">loykee.aSep08</a>		14459	1314		42	putative protein (4.8 kD) (loykee) alternative variant aSep08, mRNA.
<a href="#">loyloy</a>	<a href="#">loyloy.aSep08</a>		1213	313		70	putative protein (loyloy) mRNA.
<a href="#">loymee</a>	<a href="#">loymee.aSep08</a>		556	482		87	putative protein of bilateral origin (loymee) mRNA.
<a href="#">loymer</a>	<a href="#">loymer.aSep08</a>		10956	827		41	putative protein (loymer) mRNA.
<a href="#">loynoy</a>	<a href="#">loynoy.aSep08</a>		3892	798		24	putative protein (loynoy) mRNA.
<a href="#">loypor</a>	<a href="#">loypor.aSep08</a>		10510	550		111	putative protein (loypor) mRNA.
<a href="#">loysa</a>	<a href="#">loysa.aSep08</a>		2098	557		88	CRA b like (9.8 kD) (loysa) alternative variant aSep08, mRNA.
<a href="#">loysa</a>	<a href="#">loysa.bSep08</a>		1862	500	1	60	CRA a like (loysa) alternative variant bSep08, mRNA.
<a href="#">loyshee</a>	<a href="#">loyshee.aSep08</a>		50915	334	1	103	putative protein (loyshee) alternative variant aSep08, mRNA.
<a href="#">loyshee</a>	<a href="#">loyshee.bSep08</a>		93397	343	1	96	putative protein (10.2 kD) (loyshee) alternative variant bSep08, mRNA.
<a href="#">loytu</a>	<a href="#">loytu.aSep08</a>		4993	467		109	putative protein, with a coiled coil domain, of vertebrate origin (loytu) mRNA.
<a href="#">loyvar</a>	<a href="#">loyvar.aSep08</a>		412	306		51	putative protein (loyvar) mRNA.
<a href="#">loywey</a>	<a href="#">loywey.aSep08</a>		16259	1008	6	268	inositol 1 receptor CRA a (loywey) alternative variant aSep08, mRNA.
<a href="#">loywey</a>	<a href="#">loywey.bSep08</a>		1517	329	2	69	inositol 1 receptor CRA a (loywey) alternative variant bSep08, mRNA.

<a href="#">Lpar1</a>	<a href="#">Lpar1.bSep08</a>	<a href="#">116744</a>	74908	620	2	158	lysophosphatidic acid receptor 1 (Lpar1) alternative variant bSep08, mRNA.
<a href="#">Lpar1</a>	<a href="#">Lpar1.cSep08</a>	<a href="#">116744</a>	74792	441	2	60	lysophosphatidic acid receptor 1 (Lpar1) alternative variant cSep08, mRNA.
<a href="#">Lpar1</a>	<a href="#">Lpar1.dSep08</a>	<a href="#">116744</a>	12954	411	1	65	lysophosphatidic acid receptor 1 (Lpar1) alternative variant dSep08, mRNA.
<a href="#">Lpar1</a>	<a href="#">Lpar1.eSep08</a>	<a href="#">116744</a>	71872	351	2	54	lysophosphatidic acid receptor 1 (Lpar1) alternative variant eSep08, mRNA.
<a href="#">Lpcat1</a>	<a href="#">Lpcat1.aSep08</a>	<a href="#">361467</a>	50893	3607	14	579	lysophosphatidylcholine acyltransferase 1 (Lpcat1) alternative variant aSep08, mRNA.
<a href="#">Lphn1</a>	<a href="#">Lphn1.bSep08</a>	<a href="#">65096</a>	5032	1573	10	437	latrophilin 1 (Lphn1) alternative variant bSep08, mRNA.
<a href="#">Lphn1</a>	<a href="#">Lphn1.dSep08</a>	<a href="#">65096</a>	1487	477	1	139	latrophilin 1 (Lphn1) alternative variant dSep08, mRNA.
<a href="#">Lphn2</a>	<a href="#">Lphn2.bSep08</a>	<a href="#">171447</a>	20836	2687	7	448	latrophilin 2 (Lphn2) alternative variant bSep08, mRNA.
<a href="#">Lphn2</a>	<a href="#">Lphn2.cSep08</a>	<a href="#">171447</a>	5176	403	4	63	latrophilin 2 (Lphn2) alternative variant cSep08, mRNA.
<a href="#">Lphn2</a>	<a href="#">Lphn2.dSep08</a>	<a href="#">171447</a>	5875	773	2	45	latrophilin 2 (Lphn2) alternative variant dSep08, mRNA.
<a href="#">Lphn2</a>	<a href="#">Lphn2.eSep08</a>	<a href="#">171447</a>	2603	382	3	45	latrophilin 2 (Lphn2) alternative variant eSep08, mRNA.
<a href="#">Lphn3</a>	<a href="#">Lphn3.bSep08</a>	<a href="#">170641</a>	69696	813	3	176	latrophilin 3 (Lphn3) alternative variant bSep08, mRNA.
<a href="#">Lpin1</a>	<a href="#">Lpin1.bSep08</a>	<a href="#">313977</a>	11943	755	5	251	lipin 1 (Lpin1) alternative variant bSep08, mRNA.
<a href="#">Lpin1</a>	<a href="#">Lpin1.cSep08</a>	<a href="#">313977</a>	9965	916	7	251	lipin 1 (Lpin1) alternative variant cSep08, mRNA.
<a href="#">Lpin1</a>	<a href="#">Lpin1.dSep08</a>	<a href="#">313977</a>	9571	850	5	245	lipin 1 (Lpin1) alternative variant dSep08, mRNA.
<a href="#">Lpin2</a>	<a href="#">Lpin2.bSep08</a>	<a href="#">316737</a>	7119	675	3	187	lipin 2 (Lpin2) alternative variant bSep08, mRNA.
<a href="#">Lpin3</a>	<a href="#">Lpin3.bSep08</a>	<a href="#">362261</a>	7507	815	5	185	lipin 3 CRA b (20.2 kD) (Lpin3) alternative variant bSep08, mRNA.
<a href="#">Lpin3</a>	<a href="#">Lpin3.cSep08</a>	<a href="#">362261</a>	2355	768	4	111	lipin 3 (Lpin3) alternative variant cSep08, mRNA.
<a href="#">Lpp</a>	<a href="#">Lpp.bSep08</a>	<a href="#">288010</a>	196027	1362	6	263	LIM, zinc-binding (Lpp) alternative variant bSep08, mRNA.
<a href="#">Lpp</a>	<a href="#">Lpp.cSep08</a>	<a href="#">288010</a>	43720	757	3	217	putative protein of vertebrate origin (Lpp) alternative variant cSep08, mRNA.
<a href="#">Lpp</a>	<a href="#">Lpp.eSep08</a>	<a href="#">288010</a>	1856	342	3	40	putative protein (4.5 kD) (Lpp) alternative variant eSep08, mRNA.
<a href="#">Lpp</a>	<a href="#">Lpp.fSep08</a>	<a href="#">288010</a>	1040	272	2	48	putative protein (Lpp) alternative variant fSep08, mRNA.
<a href="#">Lpxn</a>	<a href="#">Lpxn.aSep08</a>	<a href="#">293783</a>	36511	1799	6	483	leupaxin (Lpxn) alternative variant aSep08, mRNA.
<a href="#">Lpxn</a>	<a href="#">Lpxn.cSep08</a>	<a href="#">293783</a>	29704	686	6	218	leupaxin (Lpxn) alternative variant cSep08, mRNA.
<a href="#">Lpxn</a>	<a href="#">Lpxn.dSep08</a>	<a href="#">293783</a>	7337	757	5	173	leupaxin (Lpxn) alternative variant dSep08, mRNA.
<a href="#">Lpxn</a>	<a href="#">Lpxn.eSep08</a>	<a href="#">293783</a>	29379	701	5	173	leupaxin (Lpxn) alternative variant eSep08, mRNA.
<a href="#">Lrba</a>	<a href="#">Lrba.bSep08</a>	<a href="#">361975</a>	193242	2828	14	596	LPS-responsive beige-like anchor (Lrba) alternative variant bSep08, mRNA.
<a href="#">Lrba</a>	<a href="#">Lrba.cSep08</a>	<a href="#">361975</a>	6637	985	4	311	LPS-responsive beige-like anchor (Lrba) alternative variant cSep08, mRNA.
<a href="#">Lrba</a>	<a href="#">Lrba.dSep08</a>	<a href="#">361975</a>	13901	712	6	217	LPS-responsive beige-like anchor (Lrba) alternative variant dSep08, mRNA.
<a href="#">Lrba</a>	<a href="#">Lrba.eSep08</a>	<a href="#">361975</a>	106615	557	5	185	LPS-responsive beige-like anchor (Lrba) alternative variant eSep08, mRNA.
<a href="#">Lrch1</a>	<a href="#">Lrch1.bSep08</a>	<a href="#">502020</a>	30488	746	5	210	calponin-like actin-binding (Lrch1) alternative variant bSep08, mRNA.

<a href="#">Lrch1</a>	<a href="#">Lrch1.cSep08</a>	<a href="#">502020</a>	35718	779	7	199	leucine-rich repeat calponin homology domain-containing protein 1 (22.5 kD) (Lrch1) alternative variant cSep08, mRNA.
<a href="#">Lrch1</a>	<a href="#">Lrch1.dSep08</a>	<a href="#">502020</a>	3480	525	2	154	putative protein of bilateral origin (Lrch1) alternative variant dSep08, mRNA.
<a href="#">Lrch2</a>	<a href="#">Lrch2.aSep08</a>	<a href="#">680591</a>	3646	499		165	putative protein of vertebrate origin (Lrch2) mRNA.
<a href="#">Lrch4</a>	<a href="#">Lrch4.bSep08</a>	<a href="#">360779</a>	14939	2007	18	542	leucine-rich repeat containing protein and calponin-like actin-binding (59.1 kD) (Lrch4) alternative variant bSep08, mRNA.
<a href="#">Lrch4</a>	<a href="#">Lrch4.cSep08</a>	<a href="#">360779</a>	3459	729	8	242	leucine-rich repeat containing protein (Lrch4) alternative variant cSep08, mRNA.
<a href="#">Lrch4</a>	<a href="#">Lrch4.dSep08</a>	<a href="#">360779</a>	772	493	4	122	putative protein of mammalian origin (Lrch4) alternative variant dSep08, mRNA.
<a href="#">Lrch4</a>	<a href="#">Lrch4.eSep08</a>	<a href="#">360779</a>	31612	393	2	87	putative protein of vertebrate origin (Lrch4) alternative variant eSep08, mRNA.
<a href="#">Lrch4</a>	<a href="#">Lrch4.fSep08</a>	<a href="#">360779</a>	20028	596	2	43	putative protein (5.0 kD) (Lrch4) alternative variant fSep08, mRNA.
<a href="#">Lrdd</a>	<a href="#">Lrdd.aSep08</a>	<a href="#">293625</a>	1271	840		279	putative protein (Lrdd) mRNA.
<a href="#">Lrig1</a>	<a href="#">Lrig1.aSep08</a>	<a href="#">312574</a>	38293	492	5	163	leucine-rich repeats and immunoglobulin-like domains 1 (Lrig1) alternative variant aSep08, mRNA.
<a href="#">Lrig3</a>	<a href="#">Lrig3.aSep08</a>	<a href="#">299830</a>	12761	2268	8	649	leucine-rich repeats immunoglobulin-like domains 3 (Lrig3) alternative variant aSep08, mRNA.
<a href="#">Lrig3</a>	<a href="#">Lrig3.bSep08</a>	<a href="#">299830</a>	50597	5100	17	526	leucine-rich repeats immunoglobulin-like domains 3 (Lrig3) alternative variant bSep08, mRNA.
<a href="#">Lrig3</a>	<a href="#">Lrig3.cSep08</a>	<a href="#">299830</a>	19754	788	4	204	leucine-rich repeats immunoglobulin-like domains 3 (Lrig3) alternative variant cSep08, mRNA.
<a href="#">Lrig3</a>	<a href="#">Lrig3.dSep08</a>	<a href="#">299830</a>	2047	858	2	193	leucine-rich repeats immunoglobulin-like domains 3 precursor (21.7 kD) (Lrig3) alternative variant dSep08, mRNA.
<a href="#">Lrit3</a>	<a href="#">Lrit3.aSep08</a>	<a href="#">502596</a>	3551	675		222	leucine-rich repeat, immunoglobulin-like and transmembrane domains 3 (Lrit3) mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.aSep08</a>	<a href="#">500361</a>	22623	907	10	301	lymphoid-restricted membrane protein (Lrmp) alternative variant aSep08, mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.bSep08</a>	<a href="#">500361</a>	11410	864	10	287	lymphoid-restricted membrane protein (Lrmp) alternative variant bSep08, mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.cSep08</a>	<a href="#">500361</a>	2934	710	3	161	lymphoid-restricted membrane protein (17.5 kD) (Lrmp) alternative variant cSep08, mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.eSep08</a>	<a href="#">500361</a>	1733	1174	2	126	lymphoid-restricted membrane protein (13.9 kD) (Lrmp) alternative variant eSep08, mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.fSep08</a>	<a href="#">500361</a>	7764	715	8	118	lymphoid-restricted membrane protein (Lrmp) alternative variant fSep08, mRNA.
<a href="#">Lrmp</a>	<a href="#">Lrmp.gSep08</a>	<a href="#">500361</a>	2711	667	4	88	lymphoid-restricted membrane protein (Lrmp) alternative variant gSep08, mRNA.
<a href="#">Lrp1</a>	<a href="#">Lrp1.aSep08</a>	<a href="#">299858</a>	8026	3232	18	819	low density lipoprotein receptor-related protein 1 (Lrp1) alternative variant aSep08, mRNA.
<a href="#">Lrp1</a>	<a href="#">Lrp1.bSep08</a>	<a href="#">299858</a>	4306	783	7	260	low density lipoprotein receptor-related protein 1 (Lrp1) alternative variant bSep08, mRNA.



<a href="#">Lrp1</a>	<a href="#">Lrp1.cSep08</a>	<a href="#">299858</a>	10807	631	3	209	low density lipoprotein receptor-related protein 1 (Lrp1) alternative variant cSep08, mRNA.
<a href="#">Lrp1</a>	<a href="#">Lrp1.dSep08</a>	<a href="#">299858</a>	68305	571	4	171	low density lipoprotein receptor-related protein 1 (Lrp1) alternative variant dSep08, mRNA.
<a href="#">Lrp1</a>	<a href="#">Lrp1.eSep08</a>	<a href="#">299858</a>	1631	1027	5	107	low density lipoprotein receptor-related protein 1 (Lrp1) alternative variant eSep08, mRNA.
<a href="#">Lrp2</a>	<a href="#">Lrp2.aSep08</a>	<a href="#">29216</a>	28434	3791		817	low density lipoprotein receptor-related protein 2 (Lrp2) alternative variant aSep08, mRNA.
<a href="#">Lrp2</a>	<a href="#">Lrp2.bSep08</a>	<a href="#">29216</a>	2077	605	1	126	low density lipoprotein receptor-related protein 2 (Lrp2) alternative variant bSep08, mRNA.
<a href="#">Lrp3</a>	<a href="#">Lrp3.bSep08</a>	<a href="#">89787</a>	12125	704	7	179	low density lipoprotein receptor-related protein 3 (Lrp3) alternative variant bSep08, mRNA.
<a href="#">Lrp6</a>	<a href="#">Lrp6.bSep08</a>	<a href="#">312781</a>	8759	1067	5	276	low density lipoprotein receptor-related protein 6 (Lrp6) alternative variant bSep08, mRNA.
<a href="#">Lrp8</a>	<a href="#">Lrp8.aSep08</a>	<a href="#">362558</a>	15785	474		157	low density lipoprotein receptor-related protein 8, apolipoprotein e receptor (Lrp8) mRNA.
<a href="#">Lrp10</a>	<a href="#">Lrp10.bSep08</a>	<a href="#">305880</a>	4758	1920	5	499	low-density lipoprotein receptor-related protein 10 (Lrp10) alternative variant bSep08, mRNA.
<a href="#">Lrp10</a>	<a href="#">Lrp10.cSep08</a>	<a href="#">305880</a>	1479	383	3	62	low-density lipoprotein receptor-related protein 10 (Lrp10) alternative variant cSep08, mRNA.
<a href="#">Lrp11</a>	<a href="#">Lrp11.bSep08</a>	<a href="#">292462</a>	17560	1075	5	357	low density lipoprotein receptor-related protein 11 (Lrp11) alternative variant bSep08, mRNA.
<a href="#">Lrp11</a>	<a href="#">Lrp11.cSep08</a>	<a href="#">292462</a>	57909	1102	7	243	low density lipoprotein receptor-related protein 11 (26.3 kD) (Lrp11) alternative variant cSep08, mRNA.
<a href="#">Lrpap1</a>	<a href="#">Lrpap1.aSep08</a>	<a href="#">116565</a>	14038	3442	8	367	low density lipoprotein receptor-related protein associated 1 (Lrpap1) alternative variant aSep08, mRNA.
<a href="#">Lrpap1</a>	<a href="#">Lrpap1.bSep08</a>	<a href="#">116565</a>	3899	743	2	147	low density lipoprotein receptor-related protein associated 1 (Lrpap1) alternative variant bSep08, mRNA.
<a href="#">Lrpap1</a>	<a href="#">Lrpap1.cSep08</a>	<a href="#">116565</a>	2478	525	3	99	low density lipoprotein receptor-related protein associated 1 CRA c (Lrpap1) alternative variant cSep08, mRNA.
<a href="#">Lrpprc</a>	<a href="#">Lrpprc.bSep08</a>	<a href="#">313867</a>	65216	3527	32	1131	leucine-rich PPR-motif containing (Lrpprc) alternative variant bSep08, mRNA.
<a href="#">Lrpprc</a>	<a href="#">Lrpprc.cSep08</a>	<a href="#">313867</a>	14049	330	4	109	leucine-rich PPR-motif containing (Lrpprc) alternative variant cSep08, mRNA.
<a href="#">Lrpprc</a>	<a href="#">Lrpprc.dSep08</a>	<a href="#">313867</a>	3280	665	4	70	leucine-rich PPR-motif containing (8.0 kD) (Lrpprc) alternative variant dSep08, mRNA.
<a href="#">Lrrc1</a>	<a href="#">Lrrc1.bSep08</a>	<a href="#">367113</a>	110560	683	6	227	leucine rich repeat containing 1 (Lrrc1) alternative variant bSep08, mRNA.
<a href="#">Lrrc1</a>	<a href="#">Lrrc1.cSep08</a>	<a href="#">367113</a>	8192	1012	2	67	leucine rich repeat containing 1 (10.2 kD) (Lrrc1) alternative variant cSep08, mRNA.
<a href="#">Lrrc2</a>	<a href="#">Lrrc2.bSep08</a>	<a href="#">301033</a>	15892	769	5	209	leucine rich repeat containing 2 (Lrrc2) alternative variant bSep08, mRNA.
<a href="#">Lrrc2</a>	<a href="#">Lrrc2.cSep08</a>	<a href="#">301033</a>	15729	803	5	183	leucine rich repeat containing 2 (Lrrc2) alternative variant cSep08, mRNA.
<a href="#">Lrrc2</a>	<a href="#">Lrrc2.dSep08</a>	<a href="#">301033</a>	6003	661	4	159	leucine rich repeat containing 2 (Lrrc2) alternative variant dSep08, mRNA.

<a href="#">Lrrc3b</a>	<a href="#">Lrrc3b.aSep08</a>	<a href="#">305705</a>	77833	1357		259	leucine rich repeat containing 3B (29.3 kD) (Lrrc3b) complete mRNA.
<a href="#">Lrrc4b</a>	<a href="#">Lrrc4b.aSep08</a>	<a href="#">308571</a>	16709	1917	2	638	leucine rich repeat containing 4B (Lrrc4b) alternative variant aSep08, mRNA.
<a href="#">Lrrc8a</a>	<a href="#">Lrrc8a.cSep08</a>	<a href="#">311846</a>	18662	388	2	116	leucine rich repeat containing 8A (Lrrc8a) alternative variant cSep08, mRNA.
<a href="#">Lrrc8a</a>	<a href="#">Lrrc8a.dSep08</a>	<a href="#">311846</a>	3613	374	1	41	leucine rich repeat containing 8A (Lrrc8a) alternative variant dSep08, mRNA.
<a href="#">Lrrc8c</a>	<a href="#">Lrrc8c.bSep08</a>	<a href="#">289443</a>	59717	551	3	87	leucine rich repeat containing 8 family, member C (9.7 kD) (Lrrc8c) alternative variant bSep08, mRNA.
<a href="#">Lrrc8c</a>	<a href="#">Lrrc8c.cSep08</a>	<a href="#">289443</a>	22600	396	1	30	leucine rich repeat containing 8 family, member C (3.6 kD) (Lrrc8c) alternative variant cSep08, mRNA.
<a href="#">Lrrc8d</a>	<a href="#">Lrrc8d.bSep08</a>	<a href="#">305131</a>	106863	3802	3	858	leucine rich repeat containing 8D (97.9 kD) (Lrrc8d) alternative variant bSep08, mRNA.
<a href="#">Lrrc8d</a>	<a href="#">Lrrc8d.cSep08</a>	<a href="#">305131</a>	74801	401	3	76	leucine rich repeat containing 8D (Lrrc8d) alternative variant cSep08, mRNA.
<a href="#">Lrrc9</a>	<a href="#">Lrrc9.aSep08</a>	<a href="#">299129</a>	25839	474		152	leucine rich repeat containing 9 (Lrrc9) mRNA.
<a href="#">Lrrc14</a>	<a href="#">Lrrc14.bSep08</a>	<a href="#">500900</a>	5742	1333	2	245	leucine rich repeat containing 14 (Lrrc14) alternative variant bSep08, mRNA.
<a href="#">Lrrc16a</a>	<a href="#">Lrrc16a.aSep08</a>	<a href="#">306941</a>	17742	1759	5	291	leucine rich repeat containing 16A (Lrrc16a) alternative variant aSep08, mRNA.
<a href="#">Lrrc18</a>	<a href="#">Lrrc18.bSep08</a>	<a href="#">306278</a>	18811	1647	3	255	leucine rich repeat containing 18 (28.8 kD) (Lrrc18) alternative variant bSep08, mRNA.
<a href="#">Lrrc18</a>	<a href="#">Lrrc18.cSep08</a>	<a href="#">306278</a>	19515	764	3	12	leucine rich repeat containing 18 (1.4 kD) (Lrrc18) alternative variant cSep08, mRNA.
<a href="#">Lrrc23</a>	<a href="#">Lrrc23.aSep08</a>	<a href="#">312707</a>	4316	1782	4	444	leucine rich repeat containing 23 (Lrrc23) alternative variant aSep08, mRNA.
<a href="#">Lrrc23</a>	<a href="#">Lrrc23.bSep08</a>	<a href="#">312707</a>	10190	1598	8	345	leucine rich repeat containing 23 (40.0 kD) (Lrrc23) alternative variant bSep08, mRNA.
<a href="#">Lrrc23</a>	<a href="#">Lrrc23.dSep08</a>	<a href="#">312707</a>	4294	1111	5	230	leucine rich repeat containing 23 (25.9 kD) (Lrrc23) alternative variant dSep08, complete mRNA.
<a href="#">Lrrc23</a>	<a href="#">Lrrc23.eSep08</a>	<a href="#">312707</a>	847	765	2	57	leucine rich repeat containing 23 (Lrrc23) alternative variant eSep08, mRNA.
<a href="#">Lrrc23</a>	<a href="#">Lrrc23.fSep08</a>	<a href="#">312707</a>	2090	1651	2	88	leucine rich repeat containing 23 (10.4 kD) (Lrrc23) alternative variant fSep08, mRNA.
<a href="#">Lrrc25</a>	<a href="#">Lrrc25.aSep08</a>	<a href="#">498605</a>	2931	1099		169	leucine rich repeat containing 25 (Lrrc25) mRNA.
<a href="#">Lrrc27</a>	<a href="#">Lrrc27.aSep08</a>	<a href="#">499281</a>	30292	1870	11	513	leucine rich repeat containing 27 (59.1 kD) (Lrrc27) alternative variant aSep08, mRNA.
<a href="#">Lrrc27</a>	<a href="#">Lrrc27.cSep08</a>	<a href="#">499281</a>	13755	838	6	239	leucine rich repeat containing 27 (Lrrc27) alternative variant cSep08, mRNA.
<a href="#">Lrrc27</a>	<a href="#">Lrrc27.dSep08</a>	<a href="#">499281</a>	11535	723	5	182	leucine rich repeat containing 27 (Lrrc27) alternative variant dSep08, mRNA.
<a href="#">Lrrc27</a>	<a href="#">Lrrc27.eSep08</a>	<a href="#">499281</a>	13800	821	5	146	leucine rich repeat containing 27 (17.1 kD) (Lrrc27) alternative variant eSep08, mRNA.
<a href="#">Lrrc28</a>	<a href="#">Lrrc28.bSep08</a>	<a href="#">361588</a>	77115	576	6	151	leucine rich repeat containing 28 (17.0 kD) (Lrrc28) alternative variant bSep08, mRNA.

<a href="#">Lrrc28</a>	<a href="#">Lrrc28.cSep08</a>	<a href="#">361588</a>	100085	805	8	151	leucine rich repeat containing 28 (17.0 kD) (Lrrc28) alternative variant cSep08, mRNA.
<a href="#">Lrrc28</a>	<a href="#">Lrrc28.dSep08</a>	<a href="#">361588</a>	90508	758	7	103	leucine rich repeat containing 28 (Lrrc28) alternative variant dSep08, mRNA.
<a href="#">Lrrc28</a>	<a href="#">Lrrc28.eSep08</a>	<a href="#">361588</a>	30330	692	4	46	leucine rich repeat containing 28 (5.3 kD) (Lrrc28) alternative variant eSep08, mRNA.
<a href="#">Lrrc33</a>	<a href="#">Lrrc33.bSep08</a>	<a href="#">303875</a>	15639	764	2	254	leucine rich repeat containing 33 (Lrrc33) alternative variant bSep08, mRNA.
<a href="#">Lrrc33</a>	<a href="#">Lrrc33.cSep08</a>	<a href="#">303875</a>	3901	871	1	245	leucine rich repeat containing 33 (Lrrc33) alternative variant cSep08, mRNA.
<a href="#">Lrrc36</a>	<a href="#">Lrrc36.bSep08</a>	<a href="#">361394</a>	8310	1124	1	286	leucine rich repeat containing 36 (31.6 kD) (Lrrc36) alternative variant bSep08, mRNA.
<a href="#">Lrrc37a</a>	<a href="#">Lrrc37a.aSep08</a>	<a href="#">303556</a>	5711	751		189	leucine rich repeat containing 37A (Lrrc37a) mRNA.
<a href="#">Lrrc39</a>	<a href="#">Lrrc39.bSep08</a>	<a href="#">691307</a>	13641	510	5	170	leucine rich repeat containing 39 (Lrrc39) alternative variant bSep08, mRNA.
<a href="#">Lrrc39</a>	<a href="#">Lrrc39.cSep08</a>	<a href="#">691307</a>	11308	1113	5	150	leucine rich repeat containing 39 (17.2 kD) (Lrrc39) alternative variant cSep08, mRNA.
<a href="#">Lrrc39</a>	<a href="#">Lrrc39.dSep08</a>	<a href="#">691307</a>	3735	405	3	51	leucine rich repeat containing 39 (Lrrc39) alternative variant dSep08, mRNA.
<a href="#">Lrrc40</a>	<a href="#">Lrrc40.bSep08</a>	<a href="#">310946</a>	6655	2182	4	153	leucine rich repeat containing 40 (17.2 kD) (Lrrc40) alternative variant bSep08, mRNA.
<a href="#">Lrrc40</a>	<a href="#">Lrrc40.cSep08</a>	<a href="#">310946</a>	5000	521	3	132	leucine rich repeat containing 40 (Lrrc40) alternative variant cSep08, mRNA.
<a href="#">Lrrc41</a>	<a href="#">Lrrc41.bSep08</a>	<a href="#">362566</a>	12337	748	1	133	leucine rich repeat containing 41 (Lrrc41) alternative variant bSep08, mRNA.
<a href="#">Lrrc42</a>	<a href="#">Lrrc42.bSep08</a>	<a href="#">298309</a>	5515	401	1	75	leucine rich repeat containing 42 and hypothetical protein LOC689122 (Lrrc42) alternative variant bSep08, mRNA.
<a href="#">Lrrc42</a>	<a href="#">Lrrc42.bSep08</a>	<a href="#">689122</a>	5515	401	1	75	leucine rich repeat containing 42 and hypothetical protein LOC689122 (Lrrc42) alternative variant bSep08, mRNA.
<a href="#">Lrrc43</a>	<a href="#">Lrrc43.aSep08</a>	<a href="#">288751</a>	10535	756		251	leucine rich repeat containing 43 (Lrrc43) mRNA.
<a href="#">Lrrc44</a>	<a href="#">Lrrc44.bSep08</a>	<a href="#">499732</a>	8307	751	1	198	leucine rich repeat containing 44 (Lrrc44) alternative variant bSep08, mRNA.
<a href="#">Lrrc46</a>	<a href="#">Lrrc46.bSep08</a>	<a href="#">287653</a>	11410	800	2	182	leucine rich repeat containing 46 (Lrrc46) alternative variant bSep08, mRNA.
<a href="#">Lrrc47</a>	<a href="#">Lrrc47.aSep08</a>	<a href="#">362672</a>	9433	2888	7	579	leucine rich repeat containing 47 (Lrrc47) alternative variant aSep08, mRNA.
<a href="#">Lrrc47</a>	<a href="#">Lrrc47.cSep08</a>	<a href="#">362672</a>	926	794	2	87	leucine rich repeat containing 47 (9.4 kD) (Lrrc47) alternative variant cSep08, mRNA.
<a href="#">Lrrc48</a>	<a href="#">Lrrc48.bSep08</a>	<a href="#">287371</a>	13841	709	6	117	leucine rich repeat containing 48 (Lrrc48) alternative variant bSep08, mRNA.
<a href="#">Lrrc48</a>	<a href="#">Lrrc48.cSep08</a>	<a href="#">287371</a>	751	444	2	40	leucine rich repeat containing 48 (Lrrc48) alternative variant cSep08, mRNA.
<a href="#">Lrrc49</a>	<a href="#">Lrrc49.bSep08</a>	<a href="#">300763</a>	7980	755	5	164	leucine rich repeat containing 49 (Lrrc49) alternative variant bSep08, mRNA.
<a href="#">Lrrc49</a>	<a href="#">Lrrc49.cSep08</a>	<a href="#">300763</a>	38851	638	4	76	leucine rich repeat containing 49 (Lrrc49) alternative variant cSep08, mRNA.

<a href="#">Lrrc50</a>	<a href="#">Lrrc50.bSep08</a>	<a href="#">361419</a>	3623	633	1	163	leucine rich repeat containing 50 (Lrrc50) alternative variant bSep08, mRNA.
<a href="#">Lrrc56</a>	<a href="#">Lrrc56.bSep08</a>	<a href="#">365389</a>	3865	844	1	126	leucine rich repeat containing 56 (13.9 kD) (Lrrc56) alternative variant bSep08, complete mRNA.
<a href="#">Lrrc57</a>	<a href="#">Lrrc57.bSep08</a>	<a href="#">311346</a>	1945	1026	2	175	leucine rich repeat containing 57 (19.5 kD) (Lrrc57) alternative variant bSep08, mRNA.
<a href="#">Lrrc57</a>	<a href="#">Lrrc57.cSep08</a>	<a href="#">311346</a>	1010	716	1	97	leucine rich repeat containing 57 (10.7 kD) (Lrrc57) alternative variant cSep08, mRNA.
<a href="#">Lrrc58</a>	<a href="#">Lrrc58.aSep08</a>	<a href="#">303919</a>	9021	722		240	leucine rich repeat containing 58 (Lrrc58) mRNA.
<a href="#">Lrrcc1</a>	<a href="#">Lrrcc1.aSep08</a>	<a href="#">266808</a>	11859	988		328	putative protein, with 3 coiled coil domains, of eukaryotic origin (Lrrcc1) mRNA.
<a href="#">LRRCT.0</a>	<a href="#">LRRCT.0.aSep08</a>		4667	347		115	leucine-rich repeats immunoglobulin-like domains 1 (LRRCT.0) mRNA.
<a href="#">Lrrfip1</a>	<a href="#">Lrrfip1.bSep08</a>	<a href="#">367314</a>	62502	1952	11	402	leucine rich repeat (in FLII) interacting protein 1 (Lrrfip1) alternative variant bSep08, mRNA.
<a href="#">Lrrfip1</a>	<a href="#">Lrrfip1.cSep08</a>	<a href="#">367314</a>	57892	783	6	199	leucine rich repeat (in FLII) interacting protein 1 (Lrrfip1) alternative variant cSep08, mRNA.
<a href="#">Lrrfip1</a>	<a href="#">Lrrfip1.dSep08</a>	<a href="#">367314</a>	95954	718	12	144	leucine rich repeat (in FLII) interacting protein 1 (Lrrfip1) alternative variant dSep08, mRNA.
<a href="#">Lrrfip1</a>	<a href="#">Lrrfip1.eSep08</a>	<a href="#">367314</a>	104866	735	12	143	leucine rich repeat (in FLII) interacting protein 1 (Lrrfip1) alternative variant eSep08, mRNA.
<a href="#">Lrrfip2</a>	<a href="#">Lrrfip2.bSep08</a>	<a href="#">301035</a>	101054	1453	13	342	leucine rich repeat (in FLII) interacting protein 2 (Lrrfip2) alternative variant bSep08, mRNA.
<a href="#">Lrrfip2</a>	<a href="#">Lrrfip2.cSep08</a>	<a href="#">301035</a>	41961	778	8	239	leucine rich repeat (in FLII) interacting protein 2 (Lrrfip2) alternative variant cSep08, mRNA.
<a href="#">Lrrfip2</a>	<a href="#">Lrrfip2.dSep08</a>	<a href="#">301035</a>	11564	1899	7	218	leucine rich repeat (in FLII) interacting protein 2 (Lrrfip2) alternative variant dSep08, mRNA.
<a href="#">Lrriq1</a>	<a href="#">Lrriq1.aSep08</a>	<a href="#">299759</a>	25733	675		224	leucine-rich repeats and IQ motif containing 1 (Lrriq1) mRNA.
<a href="#">Lrrk1</a>	<a href="#">Lrrk1.aSep08</a>	<a href="#">308703</a>	10376	990		330	leucine-rich repeat kinase 1 (Lrrk1) mRNA.
<a href="#">Lrrk2</a>	<a href="#">Lrrk2.aSep08</a>	<a href="#">300160</a>	53473	2507		681	leucine-rich repeat kinase 2 (Lrrk2) mRNA.
<a href="#">Lrrn2</a>	<a href="#">Lrrn2.aSep08</a>	<a href="#">289020</a>	58028	630		113	leucine rich repeat protein 2, neuronal (Lrrn2) mRNA.
<a href="#">LRRNT.0</a>	<a href="#">LRRNT.0.aSep08</a>		10298	588		154	slit homolog 2 (LRRNT.0) mRNA.
<a href="#">LRRNT.1</a>	<a href="#">LRRNT.1.aSep08</a>		3057	389		129	leucine-rich repeat, cysteine-rich flanking region, N-terminal (LRRNT.1) mRNA.
<a href="#">LRRNT.3</a>	<a href="#">LRRNT.3.aSep08</a>		1045	459		153	CRA a (LRRNT.3) mRNA.
<a href="#">LRR_1.1</a>	<a href="#">LRR_1.1.aSep08</a>		15257	1315		422	extracellular matrix protein 2 (LRR_1.1) mRNA.
<a href="#">LRR_1.2</a>	<a href="#">LRR_1.2.aSep08</a>		45700	582	6	194	G protein-coupled receptor 125 (LRR_1.2) alternative variant aSep08, mRNA.
<a href="#">LRR_1.2</a>	<a href="#">LRR_1.2.bSep08</a>		55583	957	8	169	G protein-coupled receptor 125 (LRR_1.2) alternative variant bSep08, mRNA.
<a href="#">LRR_1.2</a>	<a href="#">LRR_1.2.eSep08</a>		3457	548	2	39	putative protein (4.6 kD) (LRR_1.2) alternative variant eSep08, mRNA.
<a href="#">LRR_1.3</a>	<a href="#">LRR_1.3.aSep08</a>		7485	595	4	197	leucine-rich repeat-containing protein homolog (LRR_1.3) alternative variant aSep08, mRNA.
<a href="#">LRR_1.4</a>	<a href="#">LRR_1.4.aSep08</a>		808	378		125	scribbled homolog CRA a (LRR_1.4) mRNA.

<a href="#">LRR_1.5</a>	<a href="#">LRR_1.5.aSep08</a>		5668	399		132	leucine-rich repeat kinase 2 (LRR_1.5) mRNA.
<a href="#">LRR_1.6</a>	<a href="#">LRR_1.6.aSep08</a>		7761	2154		412	podocan (LRR_1.6) mRNA.
<a href="#">LRR_1.7</a>	<a href="#">LRR_1.7.aSep08</a>		8292	788		109	leucine-rich repeats immunoglobulin-like domains 1 (LRR_1.7) mRNA.
<a href="#">LRR_1.8</a>	<a href="#">LRR_1.8.aSep08</a>		2128	589		162	leucine rich neuronal (LRR_1.8) mRNA.
<a href="#">LRR_1.9</a>	<a href="#">LRR_1.9.aSep08</a>		18640	593		181	erbb2 interacting protein (LRR_1.9) mRNA.
<a href="#">LRR_1.10</a>	<a href="#">LRR_1.10.aSep08</a>		3389	425		141	leucine-rich repeat containing protein (LRR_1.10) mRNA.
<a href="#">LRR_1.11</a>	<a href="#">LRR_1.11.aSep08</a>		2846	1264		322	leucine-rich repeats IQ motif containing 4 (LRR_1.11) mRNA.
<a href="#">LRR_1.12</a>	<a href="#">LRR_1.12.aSep08</a>		11015	785		261	leucine-rich repeat kinase 1 (LRR_1.12) mRNA.
<a href="#">Lrsam1</a>	<a href="#">Lrsam1.bSep08</a>	<a href="#">311866</a>	4061	642	3	119	leucine rich repeat and sterile alpha motif containing 1 (Lrsam1) alternative variant bSep08, mRNA.
<a href="#">Lrsam1</a>	<a href="#">Lrsam1.cSep08</a>	<a href="#">311866</a>	1727	405	2	50	leucine rich repeat and sterile alpha motif containing 1 (Lrsam1) alternative variant cSep08, mRNA.
<a href="#">Lrsam1</a>	<a href="#">Lrsam1.dSep08</a>	<a href="#">311866</a>	4093	343	5	58	leucine rich repeat and sterile alpha motif containing 1 (Lrsam1) alternative variant dSep08, mRNA.
<a href="#">Lsamp</a>	<a href="#">Lsamp.bSep08</a>	<a href="#">29561</a>	257256	1188	7	194	limbic system-associated membrane protein (Lsamp) alternative variant bSep08, mRNA.
<a href="#">Lsg1</a>	<a href="#">Lsg1.bSep08</a>	<a href="#">288029</a>	5698	647	5	174	large subunit GTPase 1 homolog (S. cerevisiae) (20.8 kD) (Lsg1) alternative variant bSep08, complete mRNA.
<a href="#">Lsg1</a>	<a href="#">Lsg1.dSep08</a>	<a href="#">288029</a>	2963	318	2	65	large subunit GTPase 1 homolog (S. cerevisiae) (Lsg1) alternative variant dSep08, mRNA.
<a href="#">LSM.0</a>	<a href="#">LSM.0.bSep08</a>		2502	604	2	80	like-Sm ribonucleoprotein, core (9.1 kD) (LSM.0) alternative variant bSep08, mRNA.
<a href="#">LSM.1</a>	<a href="#">LSM.1.aSep08</a>		6199	374	4	82	small nuclear ribonucleoprotein E CRA a (LSM.1) alternative variant aSep08, mRNA.
<a href="#">LSM.1</a>	<a href="#">LSM.1.bSep08</a>		6238	417	4	67	small nuclear ribonucleoprotein E CRA a (7.9 kD) (LSM.1) alternative variant bSep08, mRNA.
<a href="#">LSM.1</a>	<a href="#">LSM.1.cSep08</a>		6190	307	2	61	small nuclear ribonucleoprotein e (LSM.1) alternative variant cSep08, mRNA.
<a href="#">LSM.1</a>	<a href="#">LSM.1.dSep08</a>		3360	364	2	59	small nuclear ribonucleoprotein e CRA a (LSM.1) alternative variant dSep08, mRNA.
<a href="#">LSM.1</a>	<a href="#">LSM.1.eSep08</a>		6292	898	3	52	small nuclear ribonucleoprotein e (5.9 kD) (LSM.1) alternative variant eSep08, mRNA.
<a href="#">Lsm4</a>	<a href="#">Lsm4.cSep08</a>	<a href="#">290647</a>	5301	812	5	103	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae) (11.3 kD) (Lsm4) alternative variant cSep08, mRNA.
<a href="#">Lsm4</a>	<a href="#">Lsm4.dSep08</a>	<a href="#">290647</a>	4526	653	4	40	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae) (Lsm4) alternative variant dSep08, mRNA.
<a href="#">Lsm5</a>	<a href="#">Lsm5.aSep08</a>	<a href="#">306222</a>	3138	460	4	83	LSM5 homolog, U6 small nuclear RNA associated (S. cerevisiae) (Lsm5) alternative variant aSep08, mRNA.
<a href="#">Lsm5</a>	<a href="#">Lsm5.bSep08</a>	<a href="#">306222</a>	1587	418	1	59	LSM5 homolog, U6 small nuclear RNA associated (S. cerevisiae) (Lsm5) alternative variant bSep08, mRNA.
<a href="#">Lsm6</a>	<a href="#">Lsm6.bSep08</a>	<a href="#">498934</a>	2328	456	2	87	LSM6 homolog, U6 small nuclear RNA associated (S. cerevisiae) (Lsm6) alternative variant bSep08, mRNA.

<a href="#">Lsm6</a>	<a href="#">Lsm6.dSep08</a>	<a href="#">498934</a>	2346	448	2	63	LSM6 homolog, U6 small nuclear RNA associated ( <i>S. cerevisiae</i> ) (Lsm6) alternative variant dSep08, mRNA.
<a href="#">Lsm7</a>	<a href="#">Lsm7.bSep08</a>	<a href="#">362829</a>	2356	587	4	40	LSM7 homolog, U6 small nuclear RNA associated ( <i>S. cerevisiae</i> ) (4.6 kD) (Lsm7) alternative variant bSep08, mRNA.
<a href="#">Lsm8</a>	<a href="#">Lsm8.bSep08</a>	<a href="#">296913</a>	5772	697	4	88	LSM8 homolog, U6 small nuclear RNA associated ( <i>S. cerevisiae</i> ) (Lsm8) alternative variant bSep08, mRNA.
<a href="#">Lsm8</a>	<a href="#">Lsm8.cSep08</a>	<a href="#">296913</a>	3931	769	3	67	LSM8 homolog, U6 small nuclear RNA associated ( <i>S. cerevisiae</i> ) (7.4 kD) (Lsm8) alternative variant cSep08, mRNA.
<a href="#">Lsm10</a>	<a href="#">Lsm10.bSep08</a>	<a href="#">366468</a>	1910	819	2	122	u7 snRNP-specific Sm-like protein LSM10 (13.9 kD) (Lsm10) alternative variant bSep08, complete mRNA.
<a href="#">Lsm10</a>	<a href="#">Lsm10.cSep08</a>	<a href="#">366468</a>	529	367	2	103	u7 snRNP-specific Sm-like protein LSM10 (Lsm10) alternative variant cSep08, mRNA.
<a href="#">Lsm11</a>	<a href="#">Lsm11.aSep08</a>	<a href="#">501688</a>	15613	2390		362	u7 snRNP-specific Sm-like protein LSM11 (Lsm11) alternative variant aSep08, mRNA.
<a href="#">Lsm14a</a>	<a href="#">Lsm14a.bSep08</a>	<a href="#">361554</a>	44259	2464	9	373	LSM14 homolog A (SCD6, <i>S. cerevisiae</i> ) (Lsm14a) alternative variant bSep08, mRNA.
<a href="#">Lsm14a</a>	<a href="#">Lsm14a.cSep08</a>	<a href="#">361554</a>	13788	1580	7	283	LSM14 homolog A (SCD6, <i>S. cerevisiae</i> ) (Lsm14a) alternative variant cSep08, mRNA.
<a href="#">Lsm16</a>	<a href="#">Lsm16.aSep08</a>	<a href="#">315708</a>	9459	1199		193	LSM16 homolog (EDC3, <i>S. cerevisiae</i> ) (Lsm16) mRNA.
<a href="#">Lsm1</a>	<a href="#">Lsm1.bSep08</a>	<a href="#">287429</a>	541	382	2	73	LSM domain-containing 1 (7.9 kD) (Lsm1) alternative variant bSep08, mRNA.
<a href="#">Lsm1</a>	<a href="#">Lsm1.cSep08</a>	<a href="#">287429</a>	1182	528	3	73	LSM domain-containing 1 (7.9 kD) (Lsm1) alternative variant cSep08, mRNA.
<a href="#">Lsp1</a>	<a href="#">Lsp1.bSep08</a>	<a href="#">361680</a>	22445	1513	11	329	lymphocyte specific 1 (36.4 kD) (Lsp1) alternative variant bSep08, mRNA.
<a href="#">Lsp1</a>	<a href="#">Lsp1.cSep08</a>	<a href="#">361680</a>	8317	1057	9	224	lymphocyte specific 1 (Lsp1) alternative variant cSep08, mRNA.
<a href="#">Lsp1</a>	<a href="#">Lsp1.dSep08</a>	<a href="#">361680</a>	13248	644	4	184	lymphocyte specific 1 (Lsp1) alternative variant dSep08, mRNA.
<a href="#">Lsr</a>	<a href="#">Lsr.bSep08</a>	<a href="#">64355</a>	15206	1799	8	528	lipolysis stimulated lipoprotein receptor (Lsr) alternative variant bSep08, mRNA.
<a href="#">Lsr</a>	<a href="#">Lsr.cSep08</a>	<a href="#">64355</a>	13910	726	4	241	lipolysis stimulated lipoprotein receptor (Lsr) alternative variant cSep08, mRNA.
<a href="#">Lsr</a>	<a href="#">Lsr.dSep08</a>	<a href="#">64355</a>	1008	888	2	102	lipolysis stimulated lipoprotein receptor (Lsr) alternative variant dSep08, mRNA.
<a href="#">Lss</a>	<a href="#">Lss.aSep08</a>	<a href="#">81681</a>	19453	1726		575	lanosterol synthase (Lss) mRNA.
<a href="#">Lta4h</a>	<a href="#">Lta4h.aSep08</a>	<a href="#">299732</a>	32861	2031	19	611	leukotriene A4 hydrolase and similar to Proteasome subunit beta type 3 (Proteasome theta chain) (Proteasome chain 13) (Proteasome component C10-II) (69.1 kD) (Lta4h) alternative variant aSep08, mRNA.
<a href="#">Lta4h</a>	<a href="#">Lta4h.aSep08</a>	<a href="#">680680</a>	32861	2031	19	611	leukotriene A4 hydrolase and similar to Proteasome subunit beta type 3 (Proteasome theta chain) (Proteasome chain 13) (Proteasome component C10-II) (69.1 kD) (Lta4h) alternative variant aSep08, mRNA.

<a href="#">Lta4h</a>	<a href="#">Lta4h.bSep08</a>	<a href="#">299732</a>	9807	479	5	77	leukotriene A4 hydrolase and similar to Proteasome subunit beta type 3 (Proteasome theta chain) (Proteasome chain 13) (Proteasome component C10-II) (Lta4h) alternative variant bSep08, mRNA.
<a href="#">Lta4h</a>	<a href="#">Lta4h.bSep08</a>	<a href="#">680680</a>	9807	479	5	77	leukotriene A4 hydrolase and similar to Proteasome subunit beta type 3 (Proteasome theta chain) (Proteasome chain 13) (Proteasome component C10-II) (Lta4h) alternative variant bSep08, mRNA.
<a href="#">Ltb</a>	<a href="#">Ltb.bSep08</a>	<a href="#">361795</a>	968	644	1	183	lymphotoxin B (Ltb) alternative variant bSep08, mRNA.
<a href="#">Ltb4dh</a>	<a href="#">Ltb4dh.bSep08</a>	<a href="#">192227</a>	4730	572	1	108	leukotriene B4 12-hydroxydehydrogenase (Ltb4dh) alternative variant bSep08, mRNA.
<a href="#">Ltbp1</a>	<a href="#">Ltbp1.aSep08</a>	<a href="#">59107</a>	42639	2179	10	463	latent transforming growth factor beta binding protein 1 (Ltbp1) alternative variant aSep08, mRNA.
<a href="#">Ltbp2</a>	<a href="#">Ltbp2.bSep08</a>	<a href="#">59106</a>	3152	1945	6	303	latent transforming growth factor beta binding protein 2 (Ltbp2) alternative variant bSep08, mRNA.
<a href="#">Ltbp4</a>	<a href="#">Ltbp4.aSep08</a>	<a href="#">292734</a>	9186	2137	16	712	latent transforming growth factor beta binding protein 4 (Ltbp4) alternative variant aSep08, mRNA.
<a href="#">Ltbp4</a>	<a href="#">Ltbp4.bSep08</a>	<a href="#">292734</a>	16733	2421	13	674	latent transforming growth factor beta binding protein 4 (Ltbp4) alternative variant bSep08, mRNA.
<a href="#">Ltbp4</a>	<a href="#">Ltbp4.cSep08</a>	<a href="#">292734</a>	20818	1786	9	509	latent transforming growth factor beta binding protein 4 (Ltbp4) alternative variant cSep08, mRNA.
<a href="#">Ltbp4</a>	<a href="#">Ltbp4.dSep08</a>	<a href="#">292734</a>	2781	1084	4	273	latent transforming growth factor beta binding protein 4 (Ltbp4) alternative variant dSep08, mRNA.
<a href="#">Ltbp4</a>	<a href="#">Ltbp4.eSep08</a>	<a href="#">292734</a>	1859	990	3	243	latent transforming growth factor beta binding protein 4 (Ltbp4) alternative variant eSep08, mRNA.
<a href="#">Ltbr</a>	<a href="#">Ltbr.bSep08</a>	<a href="#">297604</a>	1375	702	4	159	lymphotoxin B receptor (Ltbr) alternative variant bSep08, mRNA.
<a href="#">Ltc4s</a>	<a href="#">Ltc4s.aSep08</a>	<a href="#">114097</a>	2012	724	4	240	leukotriene C4 synthase (Ltc4s) alternative variant aSep08, mRNA.
<a href="#">Ltv1</a>	<a href="#">Ltv1.bSep08</a>	<a href="#">361452</a>	2124	1602	1	74	LTV1 homolog (S. cerevisiae) (8.8 kD) (Ltv1) alternative variant bSep08, mRNA.
<a href="#">luby</a>	<a href="#">luby.aSep08</a>		3573	1220	4	406	putative protein of eukaryotic origin (luby) mRNA.
<a href="#">Luc7l</a>	<a href="#">Luc7l.aSep08</a>	<a href="#">360503</a>	29743	1500	11	318	CRA b (38.1 kD) (Luc7l) alternative variant aSep08, complete mRNA.
<a href="#">Luc7l</a>	<a href="#">Luc7l.cSep08</a>	<a href="#">360503</a>	25951	801	7	238	luc7 homolog -like (Luc7l) alternative variant cSep08, mRNA.
<a href="#">Luc7l</a>	<a href="#">Luc7l.dSep08</a>	<a href="#">360503</a>	1201	812	2	102	putative nuclear protein (11.3 kD) (Luc7l) alternative variant dSep08, complete mRNA.
<a href="#">Luc7l</a>	<a href="#">Luc7l.fSep08</a>	<a href="#">360503</a>	6072	534	4	79	putative protein (8.6 kD) (Luc7l) alternative variant fSep08, complete mRNA.
<a href="#">Luc7l</a>	<a href="#">Luc7l.gSep08</a>	<a href="#">360503</a>	6786	422	2	70	CRA a (Luc7l) alternative variant gSep08, mRNA.
<a href="#">Luc7l2</a>	<a href="#">Luc7l2.bSep08</a>	<a href="#">312251</a>	27773	1365	7	330	LUC7-like 2 (S. cerevisiae) and hypothetical protein LOC689861 (Luc7l2) alternative variant bSep08, mRNA.
<a href="#">Luc7l2</a>	<a href="#">Luc7l2.bSep08</a>	<a href="#">689861</a>	27773	1365	7	330	LUC7-like 2 (S. cerevisiae) and hypothetical protein LOC689861 (Luc7l2) alternative variant bSep08, mRNA.
<a href="#">Luc7l2</a>	<a href="#">Luc7l2.cSep08</a>	<a href="#">312251</a>	22739	1870	6	248	LUC7-like 2 (S. cerevisiae) and hypothetical protein LOC689861 (Luc7l2) alternative variant cSep08, mRNA.

<a href="#">Luc7l2</a>	<a href="#">Luc7l2.cSep08</a>	<a href="#">689861</a>	22739	1870	6	248	LUC7-like 2 ( <i>S. cerevisiae</i> ) and hypothetical protein LOC689861 (Luc7l2) alternative variant cSep08, mRNA.
<a href="#">Luc7l2</a>	<a href="#">Luc7l2.dSep08</a>	<a href="#">312251</a>	19929	884	4	214	LUC7-like 2 ( <i>S. cerevisiae</i> ) and hypothetical protein LOC689861 (Luc7l2) alternative variant dSep08, mRNA.
<a href="#">Luc7l2</a>	<a href="#">Luc7l2.dSep08</a>	<a href="#">689861</a>	19929	884	4	214	LUC7-like 2 ( <i>S. cerevisiae</i> ) and hypothetical protein LOC689861 (Luc7l2) alternative variant dSep08, mRNA.
<a href="#">luchy</a>	<a href="#">luchy.aSep08</a>		1210	731		34	putative protein (3.8 kD) (luchy) mRNA.
<a href="#">ludar</a>	<a href="#">ludar.aSep08</a>		1761	505		44	putative protein (4.7 kD) (ludar) mRNA.
<a href="#">lufer</a>	<a href="#">lufer.aSep08</a>		6263	994	6	187	wac mouse WW domain-containing adapter protein with coiled-coil like (20.1 kD) (lufer) alternative variant aSep08, mRNA.
<a href="#">lufer</a>	<a href="#">lufer.bSep08</a>		5686	767	5	142	putative protein of vertebrate origin (lufer) alternative variant bSep08, mRNA.
<a href="#">lufer</a>	<a href="#">lufer.cSep08</a>		2663	982	2	67	CRA a (lufer) alternative variant cSep08, mRNA.
<a href="#">luflo</a>	<a href="#">luflo.aSep08</a>		533	384		106	tectonic family member 3 like (luflo) mRNA.
<a href="#">lufly</a>	<a href="#">lufly.aSep08</a>		19536	431		66	putative protein, with a coiled coil domain (7.3 kD) (lufly) mRNA.
<a href="#">luje</a>	<a href="#">luje.aSep08</a>		18980	932	4	173	brdt mouse testis-specific protein like (luje) mRNA.
<a href="#">lukee</a>	<a href="#">lukee.aSep08</a>		477	327		53	putative protein (5.9 kD) (lukee) mRNA.
<a href="#">luloy</a>	<a href="#">luloy.aSep08</a>		18717	850		229	ATPase type 13a3 (luloy) mRNA.
<a href="#">lumee</a>	<a href="#">lumee.bSep08</a>		441	206	2	37	putative protein (lumee) alternative variant bSep08, mRNA.
<a href="#">lumer</a>	<a href="#">lumer.aSep08</a>		3046	401		133	A member cassette sub-family 8 (lumer) mRNA.
<a href="#">Lung_7-TM_R.0</a>	<a href="#">Lung_7-TM_R.0.aSep08</a>		18987	1127		352	transmembrane protein 87B (Lung_7-TM_R.0) mRNA.
<a href="#">lunoy</a>	<a href="#">lunoy.aSep08</a>		13606	1060	4	78	putative protein of vertebrate origin (8.9 kD) (lunoy) alternative variant aSep08, mRNA.
<a href="#">lunoy</a>	<a href="#">lunoy.bSep08</a>		3849	743	1	51	putative protein of mammalian origin (5.8 kD) (lunoy) alternative variant bSep08, mRNA.
<a href="#">lupor</a>	<a href="#">lupor.aSep08</a>		10218	469		54	putative protein (lupor) mRNA.
<a href="#">lusa</a>	<a href="#">lusa.aSep08</a>		31137	388		128	collagen type xxii alpha 1 (lusa) mRNA.
<a href="#">lushee</a>	<a href="#">lushee.aSep08</a>		11401	744		247	putative protein of vertebrate origin (lushee) mRNA.
<a href="#">lutu</a>	<a href="#">lutu.aSep08</a>		684	551		79	putative protein (8.6 kD) (lutu) mRNA.
<a href="#">luvar</a>	<a href="#">luvar.aSep08</a>		1919	1303		36	putative protein (3.8 kD) (luvar) mRNA.
<a href="#">luwey</a>	<a href="#">luwey.aSep08</a>		1185	449		40	putative protein (4.4 kD) (luwey) mRNA.
<a href="#">Luzp5</a>	<a href="#">Luzp5.aSep08</a>	<a href="#">362798</a>	8500	286		95	leucine zipper protein 5 (Luzp5) mRNA.
<a href="#">Lxn</a>	<a href="#">Lxn.bSep08</a>	<a href="#">59073</a>	2749	616	3	146	latexin (16.4 kD) (Lxn) alternative variant bSep08, mRNA.
<a href="#">Ly6b</a>	<a href="#">Ly6b.cSep08</a>	<a href="#">246138</a>	2591	768	3	56	lymphocyte antigen 6 complex, locus B (Ly6b) alternative variant cSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.aSep08</a>	<a href="#">362934</a>	2930	852	5	179	lymphocyte antigen 6 complex, locus E (18.8 kD) (Ly6e) alternative variant aSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.bSep08</a>	<a href="#">362934</a>	3451	681	5	160	lymphocyte antigen 6 complex, locus E (Ly6e) alternative variant bSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.cSep08</a>	<a href="#">362934</a>	3894	1083	4	136	lymphocyte antigen 6 complex, locus E (14.3 kD) (Ly6e) alternative variant cSep08, mRNA.



<a href="#">Ly6e</a>	<a href="#">Ly6e.dSep08</a>	<a href="#">362934</a>	3366	955	6	136	lymphocyte antigen 6 complex, locus E (14.3 kD) (Ly6e) alternative variant dSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.eSep08</a>	<a href="#">362934</a>	3366	599	5	136	lymphocyte antigen 6 complex, locus E (14.3 kD) (Ly6e) alternative variant eSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.fSep08</a>	<a href="#">362934</a>	3704	1270	4	136	lymphocyte antigen 6 complex, locus E (14.3 kD) (Ly6e) alternative variant fSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.hSep08</a>	<a href="#">362934</a>	2737	758	5	120	lymphocyte antigen 6 complex, locus E (Ly6e) alternative variant hSep08, mRNA.
<a href="#">Ly6e</a>	<a href="#">Ly6e.iSep08</a>	<a href="#">362934</a>	3063	1107	4	110	lymphocyte antigen 6 complex, locus E (11.7 kD) (Ly6e) alternative variant iSep08, mRNA.
<a href="#">Ly6g6f</a>	<a href="#">Ly6g6f.bSep08</a>	<a href="#">309609</a>	718	417	2	72	lymphocyte antigen 6 complex, locus G6F (Ly6g6f) alternative variant bSep08, mRNA.
<a href="#">Ly6h</a>	<a href="#">Ly6h.bSep08</a>	<a href="#">300025</a>	1918	910	2	181	lymphocyte antigen 6 complex, locus H (Ly6h) alternative variant bSep08, mRNA.
<a href="#">Ly6h</a>	<a href="#">Ly6h.cSep08</a>	<a href="#">300025</a>	1722	543	2	149	lymphocyte antigen 6 complex, locus H (Ly6h) alternative variant cSep08, mRNA.
<a href="#">Ly6h</a>	<a href="#">Ly6h.dSep08</a>	<a href="#">300025</a>	1602	401	2	78	lymphocyte antigen 6 complex, locus H (Ly6h) alternative variant dSep08, mRNA.
<a href="#">Ly9</a>	<a href="#">Ly9.aSep08</a>	<a href="#">289227</a>	1766	575	2	191	lymphocyte antigen 9 (Ly9) alternative variant aSep08, mRNA.
<a href="#">Ly49s6</a>	<a href="#">Ly49s6.aSep08</a>	<a href="#">494196</a>	23583	684		64	ly49 stimulatory receptor 6 (7.2 kD) (Ly49s6) mRNA.
<a href="#">Ly75</a>	<a href="#">Ly75.aSep08</a>	<a href="#">499800</a>	4955	327		108	lymphocyte antigen 75 (Ly75) mRNA.
<a href="#">Lyar</a>	<a href="#">Lyar.bSep08</a>	<a href="#">289707</a>	10010	935	6	253	ly1 antibody reactive clone (Lyar) alternative variant bSep08, mRNA.
<a href="#">Lyar</a>	<a href="#">Lyar.cSep08</a>	<a href="#">289707</a>	3346	450	3	59	ly1 antibody reactive clone (Lyar) alternative variant cSep08, mRNA.
<a href="#">lyby</a>	<a href="#">lyby.aSep08</a>		2464	1526	4	366	putative protein of vertebrate origin (lyby) alternative variant aSep08, mRNA.
<a href="#">lyby</a>	<a href="#">lyby.bSep08</a>		2627	635	4	211	putative protein of vertebrate origin (lyby) alternative variant bSep08, mRNA.
<a href="#">Lycat</a>	<a href="#">Lycat.aSep08</a>	<a href="#">362702</a>	154201	1190	2	346	lysocardiolipin acyltransferase (Lycat) alternative variant aSep08, mRNA.
<a href="#">lychy</a>	<a href="#">lychy.aSep08</a>		6536	779		67	putative protein (lychy) mRNA.
<a href="#">lydar</a>	<a href="#">lydar.aSep08</a>		52616	439		55	putative protein (lydar) mRNA.
<a href="#">lyfer</a>	<a href="#">lyfer.aSep08</a>		3959	199	2	63	gag protein like (lyfer) alternative variant aSep08, mRNA.
<a href="#">lyflo</a>	<a href="#">lyflo.aSep08</a>		1607	273		26	putative protein (lyflo) mRNA.
<a href="#">lyflu</a>	<a href="#">lyflu.aSep08</a>		6715	629		51	putative protein (5.4 kD) (lyflu) mRNA.
<a href="#">lyjey</a>	<a href="#">lyjey.aSep08</a>		6106	1535		106	bromodomain testis-specific CRA a (lyjey) mRNA.
<a href="#">Lyk5</a>	<a href="#">Lyk5.aSep08</a>	<a href="#">303605</a>	28120	1373	5	416	protein kinase LYK5 (Lyk5) alternative variant aSep08, mRNA.
<a href="#">Lyk5</a>	<a href="#">Lyk5.cSep08</a>	<a href="#">303605</a>	24584	830	1	248	protein kinase LYK5 (Lyk5) alternative variant cSep08, mRNA.
<a href="#">Lyk5</a>	<a href="#">Lyk5.dSep08</a>	<a href="#">303605</a>	28985	2129	4	235	protein kinase LYK5 (26.1 kD) (Lyk5) alternative variant dSep08, complete mRNA.
<a href="#">Lyk5</a>	<a href="#">Lyk5.eSep08</a>	<a href="#">303605</a>	24091	848	2	192	protein kinase LYK5 (Lyk5) alternative variant eSep08, mRNA.

<a href="#">lykee</a>	<a href="#">lykee.aSep08</a>		7602	338		44	putative protein (lykee) mRNA.
<a href="#">lyloy</a>	<a href="#">lyloy.aSep08</a>		3083	542		60	putative protein (6.8 kD) (lyloy) mRNA.
<a href="#">lymee</a>	<a href="#">lymee.aSep08</a>		821	369		122	polycystic kidney disease 1 like (lymee) mRNA.
<a href="#">lymer</a>	<a href="#">lymer.aSep08</a>		5176	414		137	member 8a (lymer) mRNA.
<a href="#">lynoy</a>	<a href="#">lynoy.aSep08</a>		7571	632		97	putative mitochondrial protein (11.2 kD) (lynoy) mRNA.
<a href="#">Lypd2</a>	<a href="#">Lypd2.bSep08</a>	<a href="#">300017</a>	1795	446	1	121	CD59 antigen (Lypd2) alternative variant bSep08, mRNA.
<a href="#">Lypd3</a>	<a href="#">Lypd3.bSep08</a>	<a href="#">60378</a>	1689	749	1	249	CD59 antigen (Lypd3) alternative variant bSep08, mRNA.
<a href="#">Lypla1</a>	<a href="#">Lypla1.bSep08</a>	<a href="#">25514</a>	27697	786	8	198	lysophospholipase 1 (Lypla1) alternative variant bSep08, mRNA.
<a href="#">Lypla1</a>	<a href="#">Lypla1.cSep08</a>	<a href="#">25514</a>	3758	762	3	80	lysophospholipase 1 (Lypla1) alternative variant cSep08, mRNA.
<a href="#">Lypla2</a>	<a href="#">Lypla2.bSep08</a>	<a href="#">83510</a>	3565	602	4	182	lysophospholipase 2 (Lypla2) alternative variant bSep08, mRNA.
<a href="#">Lypla2</a>	<a href="#">Lypla2.cSep08</a>	<a href="#">83510</a>	4463	1491	4	164	lysophospholipase 2 (17.7 kD) (Lypla2) alternative variant cSep08, mRNA.
<a href="#">Lypla2</a>	<a href="#">Lypla2.dSep08</a>	<a href="#">83510</a>	3535	578	4	163	lysophospholipase 2 (Lypla2) alternative variant dSep08, mRNA.
<a href="#">Lyplal1</a>	<a href="#">Lyplal1.cSep08</a>	<a href="#">289357</a>	14748	281	2	29	lysophospholipase-like 1 (3.1 kD) (Lyplal1) alternative variant cSep08, mRNA.
<a href="#">lypor</a>	<a href="#">lypor.aSep08</a>		3160	590	1	103	putative protein (11.8 kD) (lypor) alternative variant aSep08, mRNA.
<a href="#">lypor</a>	<a href="#">lypor.bSep08</a>		34686	523	1	19	putative protein (11.8 kD) (lypor) alternative variant bSep08, mRNA.
<a href="#">Lym1</a>	<a href="#">Lym1.bSep08</a>	<a href="#">365361</a>	20007	1242	3	122	LYR motif containing 1 (14.2 kD) (Lym1) alternative variant bSep08, mRNA.
<a href="#">lysa</a>	<a href="#">lysa.aSep08</a>		1716	1108		83	putative protein (9.4 kD) (lysa) mRNA.
<a href="#">lyshee</a>	<a href="#">lyshee.aSep08</a>		1914	400		133	putative protein of bilateral origin (lyshee) mRNA.
<a href="#">Lysmd2</a>	<a href="#">Lysmd2.bSep08</a>	<a href="#">300839</a>	7076	1550	3	149	putative protein of vertebrate origin (16.9 kD) (Lysmd2) alternative variant bSep08, mRNA.
<a href="#">Lysmd2</a>	<a href="#">Lysmd2.cSep08</a>	<a href="#">300839</a>	12129	180	2	59	putative protein (Lysmd2) alternative variant cSep08, mRNA.
<a href="#">Lysmd3</a>	<a href="#">Lysmd3.bSep08</a>	<a href="#">315923</a>	7728	2534	3	162	putative protein of vertebrate origin (Lysmd3) alternative variant bSep08, mRNA.
<a href="#">Lyst</a>	<a href="#">Lyst.aSep08</a>	<a href="#">85419</a>	51554	2248		654	lysosomal trafficking regulator (Lyst) alternative variant aSep08, mRNA.
<a href="#">lytu</a>	<a href="#">lytu.aSep08</a>		1359	410		91	putative protein (10.0 kD) (lytu) mRNA.
<a href="#">lyvar</a>	<a href="#">lyvar.aSep08</a>		1162	773		51	putative protein (lyvar) mRNA.
<a href="#">lywey</a>	<a href="#">lywey.aSep08</a>		69890	219		30	putative protein (lywey) mRNA.
<a href="#">Lyz2</a>	<a href="#">Lyz2.bSep08</a>	<a href="#">25211</a>	1915	387	2	121	lysozyme 2 (Lyz2) alternative variant bSep08, mRNA.
<a href="#">Lyzl4</a>	<a href="#">Lyzl4.aSep08</a>	<a href="#">363168</a>	4352	505		108	lysozyme-like 4 (Lyzl4) mRNA.
<a href="#">Lyzl6</a>	<a href="#">Lyzl6.aSep08</a>	<a href="#">287751</a>	5492	807		149	lysozyme-like 6 (17.2 kD) (Lyzl6) mRNA.
<a href="#">Lzic</a>	<a href="#">Lzic.cSep08</a>	<a href="#">366507</a>	3235	296	4	98	putative protein, with a coiled coil domain, of eukaryotic origin (Lzic) alternative variant cSep08, mRNA.
<a href="#">Lztf1</a>	<a href="#">Lztf1.bSep08</a>	<a href="#">316102</a>	12394	776	9	134	leucine zipper transcription factor-like 1 (15.5 kD) (Lztf1) alternative variant bSep08, mRNA.

<a href="#">Lztr1</a>	<a href="#">Lztr1.bSep08</a>	<a href="#">360745</a>	4462	1524	11	508	leucine-zipper-like transcription regulator 1 (Lztr1) alternative variant bSep08, mRNA.
<a href="#">Lztr1</a>	<a href="#">Lztr1.cSep08</a>	<a href="#">360745</a>	3575	868	7	249	leucine-zipper-like transcription regulator 1 (Lztr1) alternative variant cSep08, mRNA.
<a href="#">Lztr1</a>	<a href="#">Lztr1.dSep08</a>	<a href="#">360745</a>	3435	2260	6	209	leucine-zipper-like transcription regulator 1 (23.1 kD) (Lztr1) alternative variant dSep08, mRNA.
<a href="#">Lztr1</a>	<a href="#">Lztr1.fSep08</a>	<a href="#">360745</a>	1451	359	3	107	leucine-zipper-like transcription regulator 1 (Lztr1) alternative variant fSep08, mRNA.
<a href="#">Lzts2</a>	<a href="#">Lzts2.aSep08</a>	<a href="#">365468</a>	11764	3245	5	670	leucine zipper, putative tumor suppressor 2 (72.6 kD) (Lzts2) alternative variant aSep08, mRNA.
<a href="#">Lzts2</a>	<a href="#">Lzts2.cSep08</a>	<a href="#">365468</a>	7739	539	3	154	leucine zipper, putative tumor suppressor 2 (Lzts2) alternative variant cSep08, mRNA.
<a href="#">M-rip</a>	<a href="#">M-rip.cSep08</a>	<a href="#">116504</a>	16672	3565	9	520	myosin phosphatase-Rho interacting protein (M-rip) alternative variant cSep08, mRNA.
<a href="#">M-rip</a>	<a href="#">M-rip.dSep08</a>	<a href="#">116504</a>	32823	1845	6	397	myosin phosphatase-Rho interacting protein (M-rip) alternative variant dSep08, mRNA.
<a href="#">M-rip</a>	<a href="#">M-rip.eSep08</a>	<a href="#">116504</a>	8949	874	6	158	myosin phosphatase-Rho interacting protein (M-rip) alternative variant eSep08, mRNA.
<a href="#">M-rip</a>	<a href="#">M-rip.fSep08</a>	<a href="#">116504</a>	51150	421	4	102	myosin phosphatase-Rho interacting protein (11.8 kD) (M-rip) alternative variant fSep08, mRNA.
<a href="#">M-rip</a>	<a href="#">M-rip.gSep08</a>	<a href="#">116504</a>	5879	940	4	70	myosin phosphatase-Rho interacting protein (M-rip) alternative variant gSep08, mRNA.
<a href="#">M6pr</a>	<a href="#">M6pr.bSep08</a>	<a href="#">312689</a>	4647	708	3	107	mannose-6-phosphate receptor, cation dependent (M6pr) alternative variant bSep08, mRNA.
<a href="#">M6pr</a>	<a href="#">M6pr.eSep08</a>	<a href="#">312689</a>	3590	390	3	62	mannose-6-phosphate receptor, cation dependent (M6pr) alternative variant eSep08, mRNA.
<a href="#">M6prbp1</a>	<a href="#">M6prbp1.aSep08</a>	<a href="#">316130</a>	11870	1937	8	438	mannose-6-phosphate receptor binding protein 1 (47.3 kD) (M6prbp1) alternative variant aSep08, complete mRNA.
<a href="#">M6prbp1</a>	<a href="#">M6prbp1.bSep08</a>	<a href="#">316130</a>	5672	730	3	154	mannose-6-phosphate receptor binding protein 1 (16.4 kD) (M6prbp1) alternative variant bSep08, complete mRNA.
<a href="#">maby</a>	<a href="#">maby.aSep08</a>		5959	498		70	putative protein (maby) mRNA.
<a href="#">machy</a>	<a href="#">machy.aSep08</a>		6124	415		137	CRA a (machy) mRNA.
<a href="#">Macrod1</a>	<a href="#">Macrod1.bSep08</a>	<a href="#">246233</a>	132067	716	7	197	putative protein of ancient origin (Macrod1) alternative variant bSep08, mRNA.
<a href="#">Macrod1</a>	<a href="#">Macrod1.cSep08</a>	<a href="#">246233</a>	10907	664	3	186	putative protein of fungal and metazoan origin (20.1 kD) (Macrod1) alternative variant cSep08, mRNA.
<a href="#">Mad2l1</a>	<a href="#">Mad2l1.bSep08</a>	<a href="#">297176</a>	5448	1556	1	172	MAD2 (mitotic arrest deficient, homolog)-like 1 (yeast) (Mad2l1) alternative variant bSep08, mRNA.
<a href="#">Mad2l1</a>	<a href="#">Mad2l1.cSep08</a>	<a href="#">297176</a>	10507	707	1	162	MAD2 (mitotic arrest deficient, homolog)-like 1 (yeast) (Mad2l1) alternative variant cSep08, mRNA.
<a href="#">Mad2l2</a>	<a href="#">Mad2l2.bSep08</a>	<a href="#">313702</a>	4614	1158	8	211	MAD2 mitotic arrest deficient-like 2 (yeast) (24.4 kD) (Mad2l2) alternative variant bSep08, mRNA.
<a href="#">Mad2l2</a>	<a href="#">Mad2l2.cSep08</a>	<a href="#">313702</a>	2973	699	5	141	MAD2 mitotic arrest deficient-like 2 (yeast) (Mad2l2) alternative variant cSep08, mRNA.
<a href="#">Mad2l2</a>	<a href="#">Mad2l2.dSep08</a>	<a href="#">313702</a>	10515	717	6	83	MAD2 mitotic arrest deficient-like 2 (yeast) (9.3 kD) (Mad2l2) alternative variant dSep08, mRNA.
<a href="#">madar</a>	<a href="#">madar.aSep08</a>		4109	597		57	putative protein (madar) mRNA.

<a href="#">Madd</a>	<a href="#">Madd.bSep08</a>	<a href="#">94193</a>	6686	708	7	236	MAP-kinase activating death domain (Madd) alternative variant bSep08, mRNA.
<a href="#">Madd</a>	<a href="#">Madd.dSep08</a>	<a href="#">94193</a>	8920	1440	6	134	MAP-kinase activating death domain (Madd) alternative variant dSep08, mRNA.
<a href="#">Maea</a>	<a href="#">Maea.bSep08</a>	<a href="#">298982</a>	19918	402	3	95	macrophage erythroblast attacher (Maea) alternative variant bSep08, mRNA.
<a href="#">Mael</a>	<a href="#">Mael.bSep08</a>	<a href="#">364039</a>	36000	772	1	256	maelstrom homolog (Drosophila) (Mael) alternative variant bSep08, mRNA.
<a href="#">Maf</a>	<a href="#">Maf.bSep08</a>	<a href="#">54267</a>	3038	2236	2	206	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) (Maf) alternative variant bSep08, mRNA.
<a href="#">Maf</a>	<a href="#">Maf.cSep08</a>	<a href="#">54267</a>	368439	523	3	113	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) (Maf) alternative variant cSep08, mRNA.
<a href="#">Maf</a>	<a href="#">Maf.dSep08</a>	<a href="#">54267</a>	13127	402	2	62	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) (Maf) alternative variant dSep08, mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.aSep08</a>	<a href="#">315093</a>	3052	1804	7	262	MAF1 homolog (S. cerevisiae) (28.9 kD) (Maf1) alternative variant aSep08, complete mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.cSep08</a>	<a href="#">315093</a>	1054	753	4	224	MAF1 homolog (S. cerevisiae) (Maf1) alternative variant cSep08, mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.dSep08</a>	<a href="#">315093</a>	2879	1329	7	213	MAF1 homolog (S. cerevisiae) (24.3 kD) (Maf1) alternative variant dSep08, mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.eSep08</a>	<a href="#">315093</a>	1249	661	2	81	MAF1 homolog (S. cerevisiae) (Maf1) alternative variant eSep08, mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.gSep08</a>	<a href="#">315093</a>	1346	575	2	51	MAF1 homolog (S. cerevisiae) (Maf1) alternative variant gSep08, mRNA.
<a href="#">Maf1</a>	<a href="#">Maf1.hSep08</a>	<a href="#">315093</a>	1412	335	3	55	MAF1 homolog (S. cerevisiae) (Maf1) alternative variant hSep08, mRNA.
<a href="#">mafer</a>	<a href="#">mafer.aSep08</a>		16623	752		54	putative protein (5.7 kD) (mafer) mRNA.
<a href="#">Mafg</a>	<a href="#">Mafg.aSep08</a>	<a href="#">64188</a>	2224	1543	3	233	v-maf musculoaponeurotic fibrosarcoma oncogene family, protein G (avian) (Mafg) alternative variant aSep08, mRNA.
<a href="#">Mafg</a>	<a href="#">Mafg.cSep08</a>	<a href="#">64188</a>	763	682	2	52	v-maf musculoaponeurotic fibrosarcoma oncogene family, protein G (avian) (Mafg) alternative variant cSep08, mRNA.
<a href="#">maflo</a>	<a href="#">maflo.aSep08</a>		2315	590		85	putative protein (maflo) mRNA.
<a href="#">maflu</a>	<a href="#">maflu.aSep08</a>		11082	402		133	pink-eyed dilution CRA a (maflu) mRNA.
<a href="#">Mag</a>	<a href="#">Mag.bSep08</a>	<a href="#">29409</a>	9641	1826	2	339	myelin-associated glycoprotein (37.2 kD) (Mag) alternative variant bSep08, mRNA.
<a href="#">Mag</a>	<a href="#">Mag.cSep08</a>	<a href="#">29409</a>	2687	991	3	153	myelin-associated glycoprotein (Mag) alternative variant cSep08, mRNA.
<a href="#">magar</a>	<a href="#">magar.aSep08</a>		1196	727		33	putative protein (magar) alternative variant aSep08, mRNA.
<a href="#">Maged1</a>	<a href="#">Maged1.bSep08</a>	<a href="#">84469</a>	3498	1875	8	207	melanoma antigen family d 1 like (23.8 kD) (Maged1) alternative variant bSep08, mRNA.
<a href="#">Maged1</a>	<a href="#">Maged1.dSep08</a>	<a href="#">84469</a>	1044	516	3	43	melanoma antigen family D 1 like (5.3 kD) (Maged1) alternative variant dSep08, mRNA.
<a href="#">Maged2</a>	<a href="#">Maged2.bSep08</a>	<a href="#">113947</a>	2807	927	4	277	melanoma antigen, family D, 2 (Maged2) alternative variant bSep08, mRNA.
<a href="#">Maged2</a>	<a href="#">Maged2.cSep08</a>	<a href="#">113947</a>	2797	829	5	229	melanoma antigen, family D, 2 (Maged2) alternative variant cSep08, mRNA.

<a href="#">Maged2</a>	<a href="#">Maged2.dSep08</a>	<a href="#">113947</a>	2619	764	3	184	melanoma antigen, family D, 2 (19.1 kD) (Maged2) alternative variant dSep08, mRNA.
<a href="#">Maged2</a>	<a href="#">Maged2.eSep08</a>	<a href="#">113947</a>	1145	785	2	179	melanoma antigen, family D, 2 (Maged2) alternative variant eSep08, mRNA.
<a href="#">Maged2</a>	<a href="#">Maged2.fSep08</a>	<a href="#">113947</a>	5862	583	3	147	melanoma antigen, family D, 2 (Maged2) alternative variant fSep08, mRNA.
<a href="#">Magel2</a>	<a href="#">Magel2.aSep08</a>	<a href="#">679875</a>	4090	2964		359	melanoma antigen, family L, 2 (35.8 kD) (Magel2) mRNA.
<a href="#">Magi1</a>	<a href="#">Magi1.bSep08</a>	<a href="#">500261</a>	13812	2366	5	223	PDZ/DHR/GLGF (24.5 kD) (Magi1) alternative variant bSep08, mRNA.
<a href="#">Magi1</a>	<a href="#">Magi1.cSep08</a>	<a href="#">500261</a>	129388	1006	4	157	putative protein of metazoan origin (17.2 kD) (Magi1) alternative variant cSep08, mRNA.
<a href="#">Magi1</a>	<a href="#">Magi1.dSep08</a>	<a href="#">500261</a>	12688	343	3	114	putative protein of vertebrate origin (Magi1) alternative variant dSep08, mRNA.
<a href="#">Magi1</a>	<a href="#">Magi1.eSep08</a>	<a href="#">500261</a>	12688	340	3	113	putative protein of vertebrate origin (Magi1) alternative variant eSep08, mRNA.
<a href="#">Magi1</a>	<a href="#">Magi1.gSep08</a>	<a href="#">500261</a>	9305	400	2	50	putative protein (Magi1) alternative variant gSep08, mRNA.
<a href="#">Magi2</a>	<a href="#">Magi2.bSep08</a>	<a href="#">113970</a>	29160	757	6	240	membrane associated guanylate kinase 2 (Magi2) alternative variant bSep08, mRNA.
<a href="#">Magi2</a>	<a href="#">Magi2.cSep08</a>	<a href="#">113970</a>	29614	722	6	240	membrane associated guanylate kinase 2 (Magi2) alternative variant cSep08, mRNA.
<a href="#">Magi2</a>	<a href="#">Magi2.dSep08</a>	<a href="#">113970</a>	56285	410	3	135	PDZ/DHR/GLGF (Magi2) alternative variant dSep08, mRNA.
<a href="#">Magi3</a>	<a href="#">Magi3.bSep08</a>	<a href="#">245903</a>	25292	1101	8	306	PDZ/DHR/GLGF (Magi3) alternative variant bSep08, mRNA.
<a href="#">Magi3</a>	<a href="#">Magi3.cSep08</a>	<a href="#">245903</a>	8207	627	5	172	PDZ/DHR/GLGF (Magi3) alternative variant cSep08, mRNA.
<a href="#">Magmas</a>	<a href="#">Magmas.aSep08</a>	<a href="#">679907</a>	8128	986	4	180	mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction (19.4 kD) (Magmas) alternative variant aSep08, mRNA.
<a href="#">Magmas</a>	<a href="#">Magmas.bSep08</a>	<a href="#">679907</a>	1583	531	4	160	mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction (Magmas) alternative variant bSep08, mRNA.
<a href="#">Magmas</a>	<a href="#">Magmas.dSep08</a>	<a href="#">679907</a>	794	382	3	103	mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction (Magmas) alternative variant dSep08, mRNA.
<a href="#">Magoh</a>	<a href="#">Magoh.aSep08</a>	<a href="#">298385</a>	7451	656		146	mago-nashi homolog, proliferation-associated (Drosophila) (17.2 kD) (Magoh) complete mRNA.
<a href="#">Magt1</a>	<a href="#">Magt1.bSep08</a>	<a href="#">116967</a>	23514	797	1	156	magnesium transporter 1 (Magt1) alternative variant bSep08, mRNA.
<a href="#">majey</a>	<a href="#">majey.aSep08</a>		1655	262		25	putative protein (majey) mRNA.
<a href="#">Mak10</a>	<a href="#">Mak10.bSep08</a>	<a href="#">64472</a>	9370	843	8	280	corneal wound healing-related protein (Mak10) alternative variant bSep08, mRNA.
<a href="#">Mak10</a>	<a href="#">Mak10.cSep08</a>	<a href="#">64472</a>	4550	521	4	172	corneal wound healing-related protein (Mak10) alternative variant cSep08, mRNA.
<a href="#">Mak10</a>	<a href="#">Mak10.dSep08</a>	<a href="#">64472</a>	22072	561	7	161	mak10 homolog amino-acid N-acetyltransferase CRA b (Mak10) alternative variant dSep08, mRNA.

<a href="#">Mak10</a>	<a href="#">Mak10.eSep08</a>	<a href="#">64472</a>	12934	414	5	136	mak10 homolog amino-acid N-acetyltransferase CRA b (Mak10) alternative variant eSep08, mRNA.
<a href="#">Mak10</a>	<a href="#">Mak10.fSep08</a>	<a href="#">64472</a>	3819	546	4	133	corneal wound healing-related protein (15.3 kD) (Mak10) alternative variant fSep08, mRNA.
<a href="#">Mak10</a>	<a href="#">Mak10.gSep08</a>	<a href="#">64472</a>	19624	791	7	63	mak10 homolog amino-acid N-acetyltransferase CRA b (7.4 kD) (Mak10) alternative variant gSep08, mRNA.
<a href="#">makee</a>	<a href="#">makee.aSep08</a>		5078	574		62	putative protein (7.2 kD) (makee) mRNA.
<a href="#">maloy</a>	<a href="#">maloy.bSep08</a>		1013	302	3	46	putative protein (maloy) alternative variant bSep08, mRNA.
<a href="#">Malt1</a>	<a href="#">Malt1.aSep08</a>	<a href="#">307366</a>	16114	1474	4	380	mucosa associated lymphoid tissue lymphoma translocation gene 1 (Malt1) alternative variant aSep08, mRNA.
<a href="#">Malt1</a>	<a href="#">Malt1.bSep08</a>	<a href="#">307366</a>	20083	764	5	138	mucosa associated lymphoid tissue lymphoma translocation gene 1 (Malt1) alternative variant bSep08, mRNA.
<a href="#">MAM.0</a>	<a href="#">MAM.0.aSep08</a>		96051	326		108	protein tyrosine phosphatase Receptor (MAM.0) mRNA.
<a href="#">MAM.1</a>	<a href="#">MAM.1.aSep08</a>		85198	608		202	anaplastic lymphoma kinase like (MAM.1) mRNA.
<a href="#">Mamdc2</a>	<a href="#">Mamdc2.aSep08</a>	<a href="#">309410</a>	56222	2039		270	MAM (Mamdc2) mRNA.
<a href="#">mamee</a>	<a href="#">mamee.aSep08</a>		1177	529		162	putative protein of mammalian origin (mamee) mRNA.
<a href="#">mamer</a>	<a href="#">mamer.aSep08</a>		1934	385		76	putative protein (mamer) mRNA.
<a href="#">Maml1</a>	<a href="#">Maml1.aSep08</a>	<a href="#">303101</a>	10857	3518	4	446	mastermind like 1 (Drosophila) (Maml1) alternative variant aSep08, mRNA.
<a href="#">Maml2</a>	<a href="#">Maml2.aSep08</a>	<a href="#">689844</a>	8611	604		200	mastermind like 2 (Drosophila) (Maml2) mRNA.
<a href="#">Man1a</a>	<a href="#">Man1a.bSep08</a>	<a href="#">294410</a>	21951	3427	2	262	mannosidase 1, alpha (Man1a) alternative variant bSep08, mRNA.
<a href="#">Man1a2</a>	<a href="#">Man1a2.bSep08</a>	<a href="#">295319</a>	83638	717	8	239	mannosidase, alpha, class 1A, member 2 (Man1a2) alternative variant bSep08, mRNA.
<a href="#">Man1a2</a>	<a href="#">Man1a2.cSep08</a>	<a href="#">295319</a>	16213	3706	3	125	mannosidase, alpha, class 1A, member 2 (Man1a2) alternative variant cSep08, mRNA.
<a href="#">Man1b1</a>	<a href="#">Man1b1.aSep08</a>	<a href="#">499751</a>	21112	3725	13	692	mannosidase, alpha, class 1B, member 1 (Man1b1) alternative variant aSep08, mRNA.
<a href="#">Man1b1</a>	<a href="#">Man1b1.bSep08</a>	<a href="#">499751</a>	6728	758	4	252	mannosidase, alpha, class 1B, member 1 (Man1b1) alternative variant bSep08, mRNA.
<a href="#">Man1b1</a>	<a href="#">Man1b1.cSep08</a>	<a href="#">499751</a>	2433	641	2	169	mannosidase, alpha, class 1B, member 1 (Man1b1) alternative variant cSep08, mRNA.
<a href="#">Man1b1</a>	<a href="#">Man1b1.dSep08</a>	<a href="#">499751</a>	5103	367	3	93	mannosidase, alpha, class 1B, member 1 (Man1b1) alternative variant dSep08, mRNA.
<a href="#">Man2a1</a>	<a href="#">Man2a1.bSep08</a>	<a href="#">25478</a>	4372	923	1	145	mannosidase 2, alpha 1 (Man2a1) alternative variant bSep08, mRNA.
<a href="#">Man2a2</a>	<a href="#">Man2a2.bSep08</a>	<a href="#">308757</a>	3975	2025	7	382	mannosidase 2, alpha 2 (Man2a2) alternative variant bSep08, mRNA.
<a href="#">Man2a2</a>	<a href="#">Man2a2.cSep08</a>	<a href="#">308757</a>	2890	519	4	167	mannosidase 2, alpha 2 (Man2a2) alternative variant cSep08, mRNA.
<a href="#">Man2a2</a>	<a href="#">Man2a2.dSep08</a>	<a href="#">308757</a>	4236	2956	3	149	mannosidase 2, alpha 2 (Man2a2) alternative variant dSep08, mRNA.
<a href="#">Man2a2</a>	<a href="#">Man2a2.eSep08</a>	<a href="#">308757</a>	2870	397	3	132	mannosidase 2, alpha 2 (Man2a2) alternative variant eSep08, mRNA.

<a href="#">Man2b1</a>	<a href="#">Man2b1.bSep08</a>	<a href="#">361378</a>	2149	645	4	214	mannosidase 2, alpha B1 (Man2b1) alternative variant bSep08, mRNA.
<a href="#">Man2b1</a>	<a href="#">Man2b1.cSep08</a>	<a href="#">361378</a>	2355	1385	6	189	mannosidase 2, alpha B1 (Man2b1) alternative variant cSep08, mRNA.
<a href="#">Man2b1</a>	<a href="#">Man2b1.dSep08</a>	<a href="#">361378</a>	1817	724	4	155	mannosidase 2, alpha B1 (Man2b1) alternative variant dSep08, mRNA.
<a href="#">Man2b1</a>	<a href="#">Man2b1.eSep08</a>	<a href="#">361378</a>	1213	888	3	121	mannosidase 2, alpha B1 (Man2b1) alternative variant eSep08, mRNA.
<a href="#">Man2c1</a>	<a href="#">Man2c1.bSep08</a>	<a href="#">246136</a>	1017	693	4	202	mannosidase alpha class 2C member 1 (Man2c1) alternative variant bSep08, mRNA.
<a href="#">Man2c1</a>	<a href="#">Man2c1.cSep08</a>	<a href="#">246136</a>	3513	2441	7	199	mannosidase alpha class 2C member 1 (Man2c1) alternative variant cSep08, mRNA.
<a href="#">Man2c1</a>	<a href="#">Man2c1.dSep08</a>	<a href="#">246136</a>	1243	901	4	168	alpha mannosidase 6A8B (Man2c1) alternative variant dSep08, mRNA.
<a href="#">Man2c1</a>	<a href="#">Man2c1.eSep08</a>	<a href="#">246136</a>	1048	778	4	139	mannosidase alpha class 2C member 1 CRA c (15.4 kD) (Man2c1) alternative variant eSep08, mRNA.
<a href="#">Man2c1</a>	<a href="#">Man2c1.fSep08</a>	<a href="#">246136</a>	964	806	2	110	mannosidase alpha class 2C member 1 (Man2c1) alternative variant fSep08, mRNA.
<a href="#">Manba</a>	<a href="#">Manba.bSep08</a>	<a href="#">310864</a>	28335	1131	1	288	mannosidase, beta A, lysosomal (33.6 kD) (Manba) alternative variant bSep08, mRNA.
<a href="#">Manbal</a>	<a href="#">Manbal.aSep08</a>	<a href="#">499934</a>	17842	1062		85	mannosidase, beta A, lysosomal-like (9.5 kD) (Manbal) mRNA.
<a href="#">Manea</a>	<a href="#">Manea.bSep08</a>	<a href="#">140808</a>	20283	2784	5	229	mannosidase, endo-alpha (26.6 kD) (Manea) alternative variant bSep08, mRNA.
<a href="#">Manea</a>	<a href="#">Manea.cSep08</a>	<a href="#">140808</a>	6726	276	2	91	mannosidase, endo-alpha (Manea) alternative variant cSep08, mRNA.
<a href="#">manoy</a>	<a href="#">manoy.aSep08</a>		876	666		73	putative protein (manoy) mRNA.
<a href="#">Maoa</a>	<a href="#">Maoa.aSep08</a>	<a href="#">29253</a>	68365	3499		537	monoamine oxidase A (Maoa) mRNA.
<a href="#">Maob</a>	<a href="#">Maob.bSep08</a>	<a href="#">25750</a>	77567	731	5	211	monoamine oxidase B (23.8 kD) (Maob) alternative variant bSep08, mRNA.
<a href="#">Maob</a>	<a href="#">Maob.cSep08</a>	<a href="#">25750</a>	66625	349	2	116	monoamine oxidase B (Maob) alternative variant cSep08, mRNA.
<a href="#">Map1lc3a</a>	<a href="#">Map1lc3a.aSep08</a>	<a href="#">362245</a>	1083	714	3	121	microtubule-associated protein 1 light chain 3 alpha (Map1lc3a) alternative variant aSep08, mRNA.
<a href="#">Map1lc3a</a>	<a href="#">Map1lc3a.cSep08</a>	<a href="#">362245</a>	451	291	2	75	microtubule-associated protein 1 light chain 3 alpha (Map1lc3a) alternative variant cSep08, mRNA.
<a href="#">Map1lc3b</a>	<a href="#">Map1lc3b.bSep08</a>	<a href="#">64862</a>	3678	408	1	100	microtubule-associated protein 1 light chain 3 beta (11.0 kD) (Map1lc3b) alternative variant bSep08, mRNA.
<a href="#">Map1s</a>	<a href="#">Map1s.bSep08</a>	<a href="#">290640</a>	4767	363	1	108	microtubule-associated protein 1S (Map1s) alternative variant bSep08, mRNA.
<a href="#">Map2</a>	<a href="#">Map2.bSep08</a>	<a href="#">25595</a>	110348	813	7	217	microtubule-associated protein 2 (Map2) alternative variant bSep08, mRNA.
<a href="#">Map2</a>	<a href="#">Map2.dSep08</a>	<a href="#">25595</a>	723	518	2	55	microtubule-associated protein 2 (6.5 kD) (Map2) alternative variant dSep08, mRNA.
<a href="#">Map2</a>	<a href="#">Map2.eSep08</a>	<a href="#">25595</a>	13791	410	3	56	microtubule-associated protein 2 (Map2) alternative variant eSep08, mRNA.

<a href="#">Map2k1</a>	<a href="#">Map2k1.bSep08</a>	<a href="#">170851</a>	8686	687	5	211	mitogen activated protein kinase kinase 1 (Map2k1) alternative variant bSep08, mRNA.
<a href="#">Map2k1</a>	<a href="#">Map2k1.cSep08</a>	<a href="#">170851</a>	28533	630	5	208	mitogen activated protein kinase kinase 1 (Map2k1) alternative variant cSep08, mRNA.
<a href="#">Map2k1</a>	<a href="#">Map2k1.dSep08</a>	<a href="#">170851</a>	2045	374	1	94	mitogen activated protein kinase kinase 1 (Map2k1) alternative variant dSep08, mRNA.
<a href="#">Map2k1ip1</a>	<a href="#">Map2k1ip1.aSep08</a>	<a href="#">362045</a>	11969	934	6	124	mitogen-activated protein kinase kinase 1 interacting protein 1 (13.6 kD) (Map2k1ip1) alternative variant aSep08, mRNA.
<a href="#">Map2k1ip1</a>	<a href="#">Map2k1ip1.bSep08</a>	<a href="#">362045</a>	11962	739	7	124	mitogen-activated protein kinase kinase 1 interacting protein 1 (13.6 kD) (Map2k1ip1) alternative variant bSep08, mRNA.
<a href="#">Map2k1ip1</a>	<a href="#">Map2k1ip1.dSep08</a>	<a href="#">362045</a>	12529	1251	7	124	mitogen-activated protein kinase kinase 1 interacting protein 1 (13.6 kD) (Map2k1ip1) alternative variant dSep08, complete mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.bSep08</a>	<a href="#">58960</a>	9572	1728	9	265	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant bSep08, mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.cSep08</a>	<a href="#">58960</a>	3488	913	4	258	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant cSep08, mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.dSep08</a>	<a href="#">58960</a>	1587	701	4	233	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant dSep08, mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.eSep08</a>	<a href="#">58960</a>	2269	886	3	220	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant eSep08, mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.fSep08</a>	<a href="#">58960</a>	2430	635	3	63	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant fSep08, mRNA.
<a href="#">Map2k2</a>	<a href="#">Map2k2.hSep08</a>	<a href="#">58960</a>	2929	206	2	58	mitogen activated protein kinase kinase 2 (Map2k2) alternative variant hSep08, mRNA.
<a href="#">Map2k5</a>	<a href="#">Map2k5.aSep08</a>	<a href="#">29568</a>	227048	2111	22	592	mitogen activated protein kinase kinase 5 (Map2k5) alternative variant aSep08, mRNA.
<a href="#">Map2k5</a>	<a href="#">Map2k5.bSep08</a>	<a href="#">29568</a>	70499	619	5	95	mitogen activated protein kinase kinase 5 (Map2k5) alternative variant bSep08, mRNA.
<a href="#">Map2k5</a>	<a href="#">Map2k5.cSep08</a>	<a href="#">29568</a>	1196	884	2	37	mitogen activated protein kinase kinase 5 (4.4 kD) (Map2k5) alternative variant cSep08, mRNA.
<a href="#">Map2k7</a>	<a href="#">Map2k7.bSep08</a>	<a href="#">363855</a>	5171	593	6	187	mitogen activated protein kinase kinase 7 (Map2k7) alternative variant bSep08, mRNA.
<a href="#">Map3k1</a>	<a href="#">Map3k1.bSep08</a>	<a href="#">116667</a>	10677	2595	6	714	mitogen activated protein kinase kinase kinase 1 (Map3k1) alternative variant bSep08, mRNA.
<a href="#">Map3k1</a>	<a href="#">Map3k1.cSep08</a>	<a href="#">116667</a>	40860	1794	5	525	mitogen activated protein kinase kinase kinase 1 (Map3k1) alternative variant cSep08, mRNA.
<a href="#">Map3k2</a>	<a href="#">Map3k2.bSep08</a>	<a href="#">171492</a>	35760	320		71	mitogen activated protein kinase kinase kinase 2 (Map3k2) alternative variant bSep08, mRNA.
<a href="#">Map3k4</a>	<a href="#">Map3k4.bSep08</a>	<a href="#">308106</a>	7005	573	6	191	mitogen activated protein kinase kinase kinase 4 (Map3k4) alternative variant bSep08, mRNA.
<a href="#">Map3k4</a>	<a href="#">Map3k4.cSep08</a>	<a href="#">308106</a>	7035	539	6	179	mitogen activated protein kinase kinase kinase 4 (Map3k4) alternative variant cSep08, mRNA.
<a href="#">Map3k4</a>	<a href="#">Map3k4.dSep08</a>	<a href="#">308106</a>	1684	577	2	157	mitogen activated protein kinase kinase kinase 4 (Map3k4) alternative variant dSep08, mRNA.



<a href="#">Map3k6</a>	<a href="#">Map3k6.bSep08</a>	<a href="#">313022</a>	1298	853	4	187	mitogen-activated protein kinase kinase kinase 6 (Map3k6) alternative variant bSep08, mRNA.
<a href="#">Map3k6</a>	<a href="#">Map3k6.cSep08</a>	<a href="#">313022</a>	989	760	2	101	mitogen-activated protein kinase kinase kinase 6 (Map3k6) alternative variant cSep08, mRNA.
<a href="#">Map3k7</a>	<a href="#">Map3k7.bSep08</a>	<a href="#">313121</a>	18024	396	5	131	mitogen activated protein kinase kinase kinase 7 (Map3k7) alternative variant bSep08, mRNA.
<a href="#">Map3k7</a>	<a href="#">Map3k7.cSep08</a>	<a href="#">313121</a>	22319	392	5	130	mitogen activated protein kinase kinase kinase 7 (Map3k7) alternative variant cSep08, mRNA.
<a href="#">Map3k7ip1</a>	<a href="#">Map3k7ip1.bSep08</a>	<a href="#">315139</a>	2947	1775	1	44	mitogen-activated protein kinase kinase kinase 7 interacting protein 1 (Map3k7ip1) alternative variant bSep08, mRNA.
<a href="#">Map3k7ip2</a>	<a href="#">Map3k7ip2.bSep08</a>	<a href="#">308267</a>	14893	3651	5	614	mitogen-activated protein kinase kinase kinase 7 interacting protein 2 (Map3k7ip2) alternative variant bSep08, mRNA.
<a href="#">Map3k7ip2</a>	<a href="#">Map3k7ip2.cSep08</a>	<a href="#">308267</a>	853	761	2	70	mitogen-activated protein kinase kinase kinase 7 interacting protein 2 (Map3k7ip2) alternative variant cSep08, mRNA.
<a href="#">Map3k8</a>	<a href="#">Map3k8.bSep08</a>	<a href="#">116596</a>	3863	468	2	156	mitogen-activated protein kinase kinase kinase 8 (Map3k8) alternative variant bSep08, mRNA.
<a href="#">Map3k9</a>	<a href="#">Map3k9.aSep08</a>	<a href="#">500690</a>	6363	675		225	mitogen-activated protein kinase kinase kinase 9 (Map3k9) mRNA.
<a href="#">Map3k10</a>	<a href="#">Map3k10.aSep08</a>	<a href="#">308463</a>	6194	863		287	mitogen activated protein kinase kinase kinase 10 (Map3k10) mRNA.
<a href="#">Map3k12</a>	<a href="#">Map3k12.bSep08</a>	<a href="#">25579</a>	2561	1982	2	113	mitogen activated protein kinase kinase kinase 12 (13.1 kD) (Map3k12) alternative variant bSep08, mRNA.
<a href="#">Map3k14</a>	<a href="#">Map3k14.bSep08</a>	<a href="#">360640</a>	994	675	1	76	mitogen-activated protein kinase kinase kinase 14 (Map3k14) alternative variant bSep08, mRNA.
<a href="#">Map4</a>	<a href="#">Map4.bSep08</a>	<a href="#">367171</a>	29724	1797	5	587	microtubule-associated protein 4 (Map4) alternative variant bSep08, mRNA.
<a href="#">Map4</a>	<a href="#">Map4.cSep08</a>	<a href="#">367171</a>	20809	3966	4	501	microtubule-associated protein 4 (51.0 kD) (Map4) alternative variant cSep08, mRNA.
<a href="#">Map4</a>	<a href="#">Map4.dSep08</a>	<a href="#">367171</a>	9634	1162	3	161	microtubule-associated protein 4 (Map4) alternative variant dSep08, mRNA.
<a href="#">Map4</a>	<a href="#">Map4.eSep08</a>	<a href="#">367171</a>	8964	395		131	microtubule-associated protein 4 (Map4) alternative variant eSep08, mRNA.
<a href="#">Map4</a>	<a href="#">Map4.fSep08</a>	<a href="#">367171</a>	2601	768	2	111	microtubule-associated protein 4 (12.1 kD) (Map4) alternative variant fSep08, mRNA.
<a href="#">Map4k1</a>	<a href="#">Map4k1.bSep08</a>	<a href="#">292763</a>	3021	704	7	197	mitogen-activated protein kinase 1 (Map4k1) alternative variant bSep08, mRNA.
<a href="#">Map4k1</a>	<a href="#">Map4k1.cSep08</a>	<a href="#">292763</a>	9173	777	10	138	mitogen activated protein kinase 1 CRA c (15.3 kD) (Map4k1) alternative variant cSep08, mRNA.
<a href="#">Map4k1</a>	<a href="#">Map4k1.dSep08</a>	<a href="#">292763</a>	5448	418	5	114	mitogen activated protein kinase 1 CRA c (13.3 kD) (Map4k1) alternative variant dSep08, mRNA.
<a href="#">Map4k1</a>	<a href="#">Map4k1.eSep08</a>	<a href="#">292763</a>	809	724	2	113	mitogen activated protein kinase 1 CRA a (Map4k1) alternative variant eSep08, mRNA.
<a href="#">Map4k1</a>	<a href="#">Map4k1.fSep08</a>	<a href="#">292763</a>	1867	485	6	102	mitogen-activated protein kinase 1 CRA c (11.3 kD) (Map4k1) alternative variant fSep08, mRNA.

<a href="#">Map4k2</a>	<a href="#">Map4k2.bSep08</a>	<a href="#">293694</a>	2165	631	9	203	mitogen activated protein kinase kinase kinase kinase 2 (Map4k2) alternative variant bSep08, mRNA.
<a href="#">Map4k3</a>	<a href="#">Map4k3.aSep08</a>	<a href="#">170920</a>	139912	1558	20	474	mitogen-activated protein kinase kinase kinase kinase 3 (Map4k3) alternative variant aSep08, mRNA.
<a href="#">Map4k3</a>	<a href="#">Map4k3.bSep08</a>	<a href="#">170920</a>	57337	3227	23	351	mitogen-activated protein kinase kinase kinase kinase 3 (40.2 kD) (Map4k3) alternative variant bSep08, mRNA.
<a href="#">Map4k3</a>	<a href="#">Map4k3.cSep08</a>	<a href="#">170920</a>	8205	477	5	156	mitogen-activated protein kinase kinase kinase kinase 3 (Map4k3) alternative variant cSep08, mRNA.
<a href="#">Map4k3</a>	<a href="#">Map4k3.dSep08</a>	<a href="#">170920</a>	6442	576	3	112	mitogen-activated protein kinase kinase kinase kinase 3 (Map4k3) alternative variant dSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.bSep08</a>	<a href="#">301363</a>	18685	1809	11	516	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant bSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.cSep08</a>	<a href="#">301363</a>	32759	1785	10	508	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant cSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.dSep08</a>	<a href="#">301363</a>	4912	669	5	222	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant dSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.eSep08</a>	<a href="#">301363</a>	7680	557	4	185	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant eSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.fSep08</a>	<a href="#">301363</a>	2223	793	4	155	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant fSep08, mRNA.
<a href="#">Map4k4</a>	<a href="#">Map4k4.hSep08</a>	<a href="#">301363</a>	990	342	2	69	mitogen-activated protein kinase kinase kinase kinase 4 (Map4k4) alternative variant hSep08, mRNA.
<a href="#">Map4k5</a>	<a href="#">Map4k5.aSep08</a>	<a href="#">503027</a>	23987	1280	12	426	mitogen-activated protein kinase kinase kinase kinase 5 (Map4k5) alternative variant aSep08, mRNA.
<a href="#">Map4k5</a>	<a href="#">Map4k5.bSep08</a>	<a href="#">503027</a>	20606	1011	10	336	mitogen-activated protein kinase kinase kinase kinase 5 (Map4k5) alternative variant bSep08, mRNA.
<a href="#">Map4k5</a>	<a href="#">Map4k5.cSep08</a>	<a href="#">503027</a>	17579	1596	9	276	mitogen-activated protein kinase kinase kinase kinase 5 (Map4k5) alternative variant cSep08, mRNA.
<a href="#">Map4k5</a>	<a href="#">Map4k5.dSep08</a>	<a href="#">503027</a>	6079	633	4	210	mitogen-activated protein kinase kinase kinase kinase 5 (Map4k5) alternative variant dSep08, mRNA.
<a href="#">Map4k5</a>	<a href="#">Map4k5.fSep08</a>	<a href="#">503027</a>	1543	376	2	80	mitogen-activated protein kinase kinase kinase kinase 5 (Map4k5) alternative variant fSep08, mRNA.
<a href="#">Map7</a>	<a href="#">Map7.aSep08</a>	<a href="#">293016</a>	46010	2019	15	626	microtubule-associated protein 7 (Map7) alternative variant aSep08, mRNA.
<a href="#">Map7</a>	<a href="#">Map7.cSep08</a>	<a href="#">293016</a>	23773	336	4	111	microtubule-associated protein 7 (Map7) alternative variant cSep08, mRNA.
<a href="#">Map7</a>	<a href="#">Map7.eSep08</a>	<a href="#">293016</a>	4178	724	3	64	microtubule-associated protein 7 (7.0 kD) (Map7) alternative variant eSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.aSep08</a>	<a href="#">681287</a>	24077	2675	16	846	arginine proline rich coiled-coil 1 (Map7d1) alternative variant aSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.bSep08</a>	<a href="#">681287</a>	5167	841	5	280	arginine proline rich coiled-coil 1 (Map7d1) alternative variant bSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.cSep08</a>	<a href="#">681287</a>	1631	704	7	234	arginine proline rich coiled-coil 1 CRA a (Map7d1) alternative variant cSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.dSep08</a>	<a href="#">681287</a>	2062	1230	7	215	arginine proline rich coiled-coil 1 (Map7d1) alternative variant dSep08, mRNA.

<a href="#">Map7d1</a>	<a href="#">Map7d1.eSep08</a>	<a href="#">681287</a>	4849	591	5	197	arginine proline rich coiled-coil 1 (Map7d1) alternative variant eSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.gSep08</a>	<a href="#">681287</a>	2332	446	4	148	arginine proline rich coiled-coil 1 (Map7d1) alternative variant gSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.hSep08</a>	<a href="#">681287</a>	1017	383	3	127	arginine proline rich coiled-coil 1 (Map7d1) alternative variant hSep08, mRNA.
<a href="#">Map7d1</a>	<a href="#">Map7d1.iSep08</a>	<a href="#">681287</a>	1254	705	5	105	arginine proline rich coiled-coil 1 CRA a (11.1 kD) (Map7d1) alternative variant iSep08, mRNA.
<a href="#">Map7d2</a>	<a href="#">Map7d2.aSep08</a>	<a href="#">317508</a>	115875	3970	6	763	putative protein, with 2 coiled coil domains, of vertebrate origin (84.9 kD) (Map7d2) alternative variant aSep08, mRNA.
<a href="#">Map7d2</a>	<a href="#">Map7d2.bSep08</a>	<a href="#">317508</a>	28747	943	1	227	putative protein of vertebrate origin (Map7d2) alternative variant bSep08, mRNA.
<a href="#">Map7d2</a>	<a href="#">Map7d2.cSep08</a>	<a href="#">317508</a>	10852	622	2	146	putative protein of vertebrate origin (Map7d2) alternative variant cSep08, mRNA.
<a href="#">Mapbpip</a>	<a href="#">Mapbpip.aSep08</a>	<a href="#">295234</a>	4023	521	4	134	mitogen-activated protein binding protein interacting protein (Mapbpip) alternative variant aSep08, mRNA.
<a href="#">Mapbpip</a>	<a href="#">Mapbpip.cSep08</a>	<a href="#">295234</a>	3322	517	4	124	mitogen-activated protein binding protein interacting protein (13.4 kD) (Mapbpip) alternative variant cSep08, complete mRNA.
<a href="#">Mapbpip</a>	<a href="#">Mapbpip.eSep08</a>	<a href="#">295234</a>	1339	635	2	65	mitogen-activated protein binding protein interacting protein (6.6 kD) (Mapbpip) alternative variant eSep08, mRNA.
<a href="#">Mapbpip</a>	<a href="#">Mapbpip.fSep08</a>	<a href="#">295234</a>	2315	514	2	54	mitogen-activated protein binding protein interacting protein (5.9 kD) (Mapbpip) alternative variant fSep08, mRNA.
<a href="#">Mapk1</a>	<a href="#">Mapk1.bSep08</a>	<a href="#">116590</a>	21281	1310	6	162	mitogen activated protein kinase 1 (18.7 kD) (Mapk1) alternative variant bSep08, mRNA.
<a href="#">Mapk1</a>	<a href="#">Mapk1.cSep08</a>	<a href="#">116590</a>	18168	1577	5	136	mitogen activated protein kinase 1 (Mapk1) alternative variant cSep08, mRNA.
<a href="#">Mapk1</a>	<a href="#">Mapk1.eSep08</a>	<a href="#">116590</a>	4193	315	3	52	mitogen activated protein kinase 1 (Mapk1) alternative variant eSep08, mRNA.
<a href="#">Mapk1ip1</a>	<a href="#">Mapk1ip1.aSep08</a>	<a href="#">499280</a>	11057	1461	2	263	putative protein (11.6 kD) (Mapk1ip1) alternative variant aSep08, complete mRNA.
<a href="#">Mapk1ip1</a>	<a href="#">Mapk1ip1.cSep08</a>	<a href="#">499280</a>	10289	755	3	110	putative protein (11.7 kD) (Mapk1ip1) alternative variant cSep08, mRNA.
<a href="#">Mapk1ip1</a>	<a href="#">Mapk1ip1.dSep08</a>	<a href="#">499280</a>	10242	767	3	110	putative protein (11.7 kD) (Mapk1ip1) alternative variant dSep08, mRNA.
<a href="#">Mapk1ip1</a>	<a href="#">Mapk1ip1.eSep08</a>	<a href="#">499280</a>	10105	753	4	29	putative protein (3.2 kD) (Mapk1ip1) alternative variant eSep08, mRNA.
<a href="#">Mapk1ip1l</a>	<a href="#">Mapk1ip1l.aSep08</a>	<a href="#">361028</a>	26562	3528	4	240	mitogen-activated protein kinase 1 interacting protein 1-like (23.7 kD) (Mapk1ip1l) alternative variant aSep08, complete mRNA.
<a href="#">Mapk3</a>	<a href="#">Mapk3.bSep08</a>	<a href="#">50689</a>	5205	1778	7	370	protein kinase 3 (42.3 kD) (Mapk3) alternative variant bSep08, mRNA.
<a href="#">Mapk3</a>	<a href="#">Mapk3.cSep08</a>	<a href="#">50689</a>	4196	753	5	233	protein kinase 3 (Mapk3) alternative variant cSep08, mRNA.
<a href="#">Mapk3</a>	<a href="#">Mapk3.dSep08</a>	<a href="#">50689</a>	1607	700	5	233	protein kinase 3 (Mapk3) alternative variant dSep08, mRNA.

<a href="#">Mapk3</a>	<a href="#">Mapk3.fSep08</a>	<a href="#">50689</a>	3020	753	6	123	mitogen-activated protein kinase 3 (Mapk3) alternative variant fSep08, mRNA.
<a href="#">Mapk3</a>	<a href="#">Mapk3.gSep08</a>	<a href="#">50689</a>	1262	345	2	93	protein kinase 3 (Mapk3) alternative variant gSep08, mRNA.
<a href="#">Mapk4</a>	<a href="#">Mapk4.aSep08</a>	<a href="#">54268</a>	3945	1396		283	mitogen-activated protein kinase 4 (Mapk4) mRNA.
<a href="#">Mapk8</a>	<a href="#">Mapk8.aSep08</a>	<a href="#">116554</a>	58085	488	3	120	mitogen-activated protein kinase 8 (Mapk8) alternative variant aSep08, mRNA.
<a href="#">Mapk8</a>	<a href="#">Mapk8.bSep08</a>	<a href="#">116554</a>	52585	448	1	87	mitogen-activated protein kinase 8 (9.8 kD) (Mapk8) alternative variant bSep08, complete mRNA.
<a href="#">Mapk8ip1</a>	<a href="#">Mapk8ip1.bSep08</a>	<a href="#">116457</a>	2375	821	3	219	mitogen-activated protein kinase 8 interacting protein 1 (Mapk8ip1) alternative variant bSep08, mRNA.
<a href="#">Mapk8ip1</a>	<a href="#">Mapk8ip1.dSep08</a>	<a href="#">116457</a>	1287	546	4	75	mitogen-activated protein kinase 8 interacting protein 1 (Mapk8ip1) alternative variant dSep08, mRNA.
<a href="#">Mapk8ip2</a>	<a href="#">Mapk8ip2.aSep08</a>	<a href="#">315220</a>	3746	1764	6	345	mitogen-activated protein kinase 8 interacting protein 2 (Mapk8ip2) alternative variant aSep08, mRNA.
<a href="#">Mapk8ip2</a>	<a href="#">Mapk8ip2.bSep08</a>	<a href="#">315220</a>	1212	740	1	71	mitogen-activated protein kinase 8 interacting protein 2 (8.3 kD) (Mapk8ip2) alternative variant bSep08, mRNA.
<a href="#">Mapk8ip3</a>	<a href="#">Mapk8ip3.aSep08</a>	<a href="#">302983</a>	8823	3874	20	826	mitogen-activated protein kinase 8 interacting protein 3 (Mapk8ip3) alternative variant aSep08, mRNA.
<a href="#">Mapk8ip3</a>	<a href="#">Mapk8ip3.bSep08</a>	<a href="#">302983</a>	927	571	5	147	mitogen-activated protein kinase 8 interacting protein 3 (Mapk8ip3) alternative variant bSep08, mRNA.
<a href="#">Mapk8ip3</a>	<a href="#">Mapk8ip3.cSep08</a>	<a href="#">302983</a>	802	677	2	60	mitogen-activated protein kinase 8 interacting protein 3 (Mapk8ip3) alternative variant cSep08, mRNA.
<a href="#">Mapk8ip3</a>	<a href="#">Mapk8ip3.dSep08</a>	<a href="#">302983</a>	490	404	2	56	mitogen-activated protein kinase 8 interacting protein 3 (Mapk8ip3) alternative variant dSep08, mRNA.
<a href="#">Mapk9</a>	<a href="#">Mapk9.bSep08</a>	<a href="#">50658</a>	6927	2662	5	175	mitogen-activated protein kinase 9 (20.0 kD) (Mapk9) alternative variant bSep08, mRNA.
<a href="#">Mapk9</a>	<a href="#">Mapk9.cSep08</a>	<a href="#">50658</a>	28016	745	1	128	mitogen-activated protein kinase 9 (Mapk9) alternative variant cSep08, mRNA.
<a href="#">Mapk9</a>	<a href="#">Mapk9.dSep08</a>	<a href="#">50658</a>	4652	392	5	122	mitogen-activated protein kinase 9 (Mapk9) alternative variant dSep08, mRNA.
<a href="#">Mapk10</a>	<a href="#">Mapk10.bSep08</a>	<a href="#">25272</a>	125130	1281	11	426	mitogen activated protein kinase 10 (Mapk10) alternative variant bSep08, mRNA.
<a href="#">Mapk10</a>	<a href="#">Mapk10.cSep08</a>	<a href="#">25272</a>	52000	1289	5	106	mitogen activated protein kinase 10 (Mapk10) alternative variant cSep08, mRNA.
<a href="#">Mapk12</a>	<a href="#">Mapk12.cSep08</a>	<a href="#">60352</a>	2876	1158	4	106	mitogen-activated protein kinase 12 (12.1 kD) (Mapk12) alternative variant cSep08, mRNA.
<a href="#">Mapk12</a>	<a href="#">Mapk12.dSep08</a>	<a href="#">60352</a>	2338	718	3	100	mitogen-activated protein kinase 12 (Mapk12) alternative variant dSep08, mRNA.
<a href="#">Mapk14</a>	<a href="#">Mapk14.bSep08</a>	<a href="#">81649</a>	18513	1254	1	155	mitogen activated protein kinase 14 (Mapk14) alternative variant bSep08, mRNA.
<a href="#">Mapkap1</a>	<a href="#">Mapkap1.bSep08</a>	<a href="#">296648</a>	199289	1502	10	445	mitogen-activated protein kinase associated protein 1 (Mapkap1) alternative variant bSep08, mRNA.
<a href="#">Mapkap1</a>	<a href="#">Mapkap1.cSep08</a>	<a href="#">296648</a>	86865	761	5	200	mitogen-activated protein kinase associated protein 1 (Mapkap1) alternative variant cSep08, mRNA.
<a href="#">Mapkap1</a>	<a href="#">Mapkap1.dSep08</a>	<a href="#">296648</a>	103239	770	4	184	mitogen-activated protein kinase associated protein 1 (Mapkap1) alternative variant dSep08, mRNA.

<a href="#">Mapkap1</a>	<a href="#">Mapkap1.eSep08</a>	<a href="#">296648</a>	5359	1932	2	118	mitogen-activated protein kinase associated protein 1 (13.4 kD) (Mapkap1) alternative variant eSep08, mRNA.
<a href="#">Mapkap1</a>	<a href="#">Mapkap1.fSep08</a>	<a href="#">296648</a>	37447	505	3	90	mitogen-activated protein kinase associated protein 1 (Mapkap1) alternative variant fSep08, mRNA.
<a href="#">Mapkapk2</a>	<a href="#">Mapkapk2.bSep08</a>	<a href="#">289014</a>	1393	327	4	105	MAP kinase-activated protein kinase 2 (Mapkapk2) alternative variant bSep08, mRNA.
<a href="#">Mapkapk3</a>	<a href="#">Mapkapk3.bSep08</a>	<a href="#">315994</a>	27290	754	5	158	mitogen-activated protein kinase-activated protein kinase 3 (Mapkapk3) alternative variant bSep08, mRNA.
<a href="#">Mapkapk5</a>	<a href="#">Mapkapk5.bSep08</a>	<a href="#">498183</a>	13095	905	9	301	MAP kinase-activated protein kinase 5 (Mapkapk5) alternative variant bSep08, mRNA.
<a href="#">Mapkapk5</a>	<a href="#">Mapkapk5.cSep08</a>	<a href="#">498183</a>	2938	658	5	187	MAP kinase-activated protein kinase 5 (Mapkapk5) alternative variant cSep08, mRNA.
<a href="#">Mapkapk5</a>	<a href="#">Mapkapk5.dSep08</a>	<a href="#">498183</a>	4141	440	4	146	MAP kinase-activated protein kinase 5 (Mapkapk5) alternative variant dSep08, mRNA.
<a href="#">Mapkapk5</a>	<a href="#">Mapkapk5.eSep08</a>	<a href="#">498183</a>	1075	732	2	80	MAP kinase-activated protein kinase 5 (9.4 kD) (Mapkapk5) alternative variant eSep08, mRNA.
<a href="#">Mapkapk5</a>	<a href="#">Mapkapk5.hSep08</a>	<a href="#">498183</a>	2828	423	4	55	MAP kinase-activated protein kinase 5 (6.6 kD) (Mapkapk5) alternative variant hSep08, mRNA.
<a href="#">mapor</a>	<a href="#">mapor.aSep08</a>		3580	455		42	putative protein (mapor) mRNA.
<a href="#">Mapre1</a>	<a href="#">Mapre1.bSep08</a>	<a href="#">114764</a>	24919	852	6	249	microtubule-associated protein, RP/EB family, member 1 (Mapre1) alternative variant bSep08, mRNA.
<a href="#">Mapre1</a>	<a href="#">Mapre1.cSep08</a>	<a href="#">114764</a>	24602	745	6	248	microtubule-associated protein, RP/EB family, member 1 (Mapre1) alternative variant cSep08, mRNA.
<a href="#">Mapre3</a>	<a href="#">Mapre3.bSep08</a>	<a href="#">298848</a>	44325	892	6	243	microtubule-associated protein, RP/EB family, member 3 (Mapre3) alternative variant bSep08, mRNA.
<a href="#">Mapre3</a>	<a href="#">Mapre3.cSep08</a>	<a href="#">298848</a>	44636	1026	7	220	microtubule-associated protein, RP/EB family, member 3 (24.8 kD) (Mapre3) alternative variant cSep08, mRNA.
<a href="#">Mapre3</a>	<a href="#">Mapre3.dSep08</a>	<a href="#">298848</a>	1444	1195	3	157	microtubule-associated protein, RP/EB family, member 3 (17.7 kD) (Mapre3) alternative variant dSep08, mRNA.
<a href="#">Mapre3</a>	<a href="#">Mapre3.eSep08</a>	<a href="#">298848</a>	6675	394	3	48	microtubule-associated protein, RP/EB family, member 3 (Mapre3) alternative variant eSep08, mRNA.
<a href="#">Mapt</a>	<a href="#">Mapt.aSep08</a>	<a href="#">29477</a>	64085	1316	6	397	microtubule-associated protein tau (Mapt) alternative variant aSep08, mRNA.
<a href="#">Mapt</a>	<a href="#">Mapt.bSep08</a>	<a href="#">29477</a>	94002	1428	9	343	microtubule-associated protein tau (35.9 kD) (Mapt) alternative variant bSep08, mRNA.
<a href="#">Mapt</a>	<a href="#">Mapt.cSep08</a>	<a href="#">29477</a>	42367	5250	10	330	microtubule-associated protein tau (34.3 kD) (Mapt) alternative variant cSep08, mRNA.
<a href="#">Mapt</a>	<a href="#">Mapt.dSep08</a>	<a href="#">29477</a>	53219	551	3	138	protein tau (14.6 kD) (Mapt) alternative variant dSep08, mRNA.
<a href="#">marby</a>	<a href="#">marby.aSep08</a>		4439	550		49	putative protein (marby) mRNA.
<a href="#">1-Mar</a>	<a href="#">March1.aSep08</a>	<a href="#">361135</a>	72108	993		277	membrane-associated ring finger (C3HC4) 1 (March1) mRNA.
<a href="#">2-Mar</a>	<a href="#">March2.bSep08</a>	<a href="#">362849</a>	25199	2515	4	232	membrane-associated ring finger (C3HC4) 2 (25.1 kD) (March2) alternative variant bSep08, mRNA.
<a href="#">2-Mar</a>	<a href="#">March2.cSep08</a>	<a href="#">362849</a>	13236	737	1	60	membrane-associated ring finger (C3HC4) 2 (6.2 kD) (March2) alternative variant cSep08, mRNA.

<a href="#">3-Mar</a>	<a href="#">March3.bSep08</a>	<a href="#">364878</a>	164055	780	5	175	glutaredoxin-like protein (19.0 kD) (March3) alternative variant bSep08, mRNA.
<a href="#">3-Mar</a>	<a href="#">March3.dSep08</a>	<a href="#">364878</a>	15991	1037	3	96	glutaredoxin-like protein (11.2 kD) (March3) alternative variant dSep08, complete mRNA.
<a href="#">3-Mar</a>	<a href="#">March3.eSep08</a>	<a href="#">364878</a>	204313	2868	8	64	glutaredoxin-like protein (7.1 kD) (March3) alternative variant eSep08, mRNA.
<a href="#">3-Mar</a>	<a href="#">March3.gSep08</a>	<a href="#">364878</a>	151977	840	4	67	putative protein (March3) alternative variant gSep08, mRNA.
<a href="#">5-Mar</a>	<a href="#">March5.bSep08</a>	<a href="#">294079</a>	602	456	2	65	membrane-associated ring finger (C3HC4) 5 (March5) alternative variant bSep08, mRNA.
<a href="#">6-Mar</a>	<a href="#">March6.aSep08</a>	<a href="#">294862</a>	28949	4514		395	membrane-associated ring finger (C3HC4) 6 (March6) mRNA.
<a href="#">7-Mar</a>	<a href="#">March7.aSep08</a>	<a href="#">311059</a>	38487	3538	9	745	membrane-associated ring finger (C3HC4) 7 (March7) alternative variant aSep08, mRNA.
<a href="#">7-Mar</a>	<a href="#">March7.cSep08</a>	<a href="#">311059</a>	6878	839	3	134	membrane-associated ring finger (C3HC4) 7 (March7) alternative variant cSep08, mRNA.
<a href="#">7-Mar</a>	<a href="#">March7.eSep08</a>	<a href="#">311059</a>	18621	603	2	100	membrane-associated ring finger (C3HC4) 7 (March7) alternative variant eSep08, mRNA.
<a href="#">7-Mar</a>	<a href="#">March7.fSep08</a>	<a href="#">311059</a>	4768	706	3	48	membrane-associated ring finger (C3HC4) 7 (March7) alternative variant fSep08, mRNA.
<a href="#">marchy</a>	<a href="#">marchy.aSep08</a>		34268	345		30	putative protein (3.3 kD) (marchy) mRNA.
<a href="#">Marcksl1</a>	<a href="#">Marcksl1.bSep08</a>	<a href="#">81520</a>	1109	850	1	194	MARCKS-like 1 (Marcksl1) alternative variant bSep08, mRNA.
<a href="#">mardar</a>	<a href="#">mardar.aSep08</a>		14654	673		55	putative protein (6.4 kD) (mardar) alternative variant aSep08, mRNA.
<a href="#">mardar</a>	<a href="#">mardar.bSep08</a>		14257	491	1	88	putative nuclear protein (9.7 kD) (mardar) alternative variant bSep08, mRNA.
<a href="#">Mare</a>	<a href="#">Mare.bSep08</a>	<a href="#">360505</a>	10320	1374	7	352	alpha globin regulatory element containing (Mare) alternative variant bSep08, mRNA.
<a href="#">Mare</a>	<a href="#">Mare.cSep08</a>	<a href="#">360505</a>	24350	961	8	272	alpha globin regulatory element containing (Mare) alternative variant cSep08, mRNA.
<a href="#">Mare</a>	<a href="#">Mare.dSep08</a>	<a href="#">360505</a>	1819	715	3	114	alpha globin regulatory element containing (13.0 kD) (Mare) alternative variant dSep08, mRNA.
<a href="#">Mare</a>	<a href="#">Mare.eSep08</a>	<a href="#">360505</a>	4001	404	2	42	putative protein (Mare) alternative variant eSep08, mRNA.
<a href="#">marfer</a>	<a href="#">marfer.aSep08</a>		2052	527		97	putative protein (marfer) mRNA.
<a href="#">marflo</a>	<a href="#">marflo.aSep08</a>		1783	458		32	putative protein (3.7 kD) (marflo) mRNA.
<a href="#">marflu</a>	<a href="#">marflu.aSep08</a>		57401	1091	3	363	ubiquitin protein ligase E3A (marflu) alternative variant aSep08, mRNA.
<a href="#">marflu</a>	<a href="#">marflu.bSep08</a>		53033	534	4	107	ubiquitin protein ligase E3A (marflu) alternative variant bSep08, mRNA.
<a href="#">marflu</a>	<a href="#">marflu.cSep08</a>		19879	529	4	39	putative protein (marflu) alternative variant cSep08, mRNA.
<a href="#">margar</a>	<a href="#">margar.aSep08</a>		9362	803		54	putative protein (6.3 kD) (margar) mRNA.
<a href="#">marjey</a>	<a href="#">marjey.aSep08</a>		29548	1170	7	187	protein Tyrosine phosphatase non-receptor type 13 (marjey) mRNA.
<a href="#">Mark1</a>	<a href="#">Mark1.bSep08</a>	<a href="#">117016</a>	16991	2365	8	637	MAP/microtubule affinity-regulating kinase 1 (Mark1) alternative variant bSep08, mRNA.

<a href="#">Mark2</a>	<a href="#">Mark2.bSep08</a>	<a href="#">60328</a>	9402	3219	10	489	MAP/microtubule affinity-regulating kinase 2 (Mark2) alternative variant bSep08, mRNA.
<a href="#">Mark2</a>	<a href="#">Mark2.cSep08</a>	<a href="#">60328</a>	5626	1602	4	272	MAP/microtubule affinity-regulating kinase 2 (29.8 kD) (Mark2) alternative variant cSep08, mRNA.
<a href="#">Mark2</a>	<a href="#">Mark2.dSep08</a>	<a href="#">60328</a>	1672	1078	2	167	MAP/microtubule affinity-regulating kinase 2 (Mark2) alternative variant dSep08, mRNA.
<a href="#">Mark3</a>	<a href="#">Mark3.bSep08</a>	<a href="#">170577</a>	33069	1686	6	361	MAP/microtubule affinity-regulating kinase 3 (Mark3) alternative variant bSep08, mRNA.
<a href="#">Mark3</a>	<a href="#">Mark3.cSep08</a>	<a href="#">170577</a>	56021	544	7	155	MAP/microtubule affinity-regulating kinase 3 (Mark3) alternative variant cSep08, mRNA.
<a href="#">Mark3</a>	<a href="#">Mark3.dSep08</a>	<a href="#">170577</a>	2241	781	2	152	MAP/microtubule affinity-regulating kinase 3 (17.7 kD) (Mark3) alternative variant dSep08, mRNA.
<a href="#">Mark4</a>	<a href="#">Mark4.aSep08</a>	<a href="#">680407</a>	14477	1497	9	498	MAP/microtubule affinity-regulating kinase 4 (Mark4) alternative variant aSep08, mRNA.
<a href="#">Mark4</a>	<a href="#">Mark4.bSep08</a>	<a href="#">680407</a>	7523	855	6	285	MAP/microtubule affinity-regulating kinase 4 (Mark4) alternative variant bSep08, mRNA.
<a href="#">Mark4</a>	<a href="#">Mark4.cSep08</a>	<a href="#">680407</a>	2996	340	3	91	MAP/microtubule affinity-regulating kinase 4 (Mark4) alternative variant cSep08, mRNA.
<a href="#">Mark4</a>	<a href="#">Mark4.dSep08</a>	<a href="#">680407</a>	2073	503	1	71	MAP/microtubule affinity-regulating kinase 4 (Mark4) alternative variant dSep08, mRNA.
<a href="#">Mark4</a>	<a href="#">Mark4.eSep08</a>	<a href="#">680407</a>	2756	339	3	60	MAP/microtubule affinity-regulating kinase 4 (Mark4) alternative variant eSep08, mRNA.
<a href="#">markee</a>	<a href="#">markee.aSep08</a>		6728	695		155	putative protein of eukaryotic origin (markee) mRNA.
<a href="#">marloy</a>	<a href="#">marloy.aSep08</a>		21748	679		41	putative protein (marloy) mRNA.
<a href="#">marmee</a>	<a href="#">marmee.aSep08</a>		1708	780		259	CRA b (marmee) mRNA.
<a href="#">marmer</a>	<a href="#">marmer.aSep08</a>		757	603		104	putative protein (11.9 kD) (marmer) mRNA.
<a href="#">marnoy</a>	<a href="#">marnoy.aSep08</a>		424	268		38	putative protein (marnoy) mRNA.
<a href="#">marpor</a>	<a href="#">marpor.aSep08</a>		2496	653	1	101	putative protein (10.9 kD) (marpor) alternative variant aSep08, mRNA.
<a href="#">marpor</a>	<a href="#">marpor.bSep08</a>		2370	483		101	putative protein (10.9 kD) (marpor) alternative variant bSep08, mRNA.
<a href="#">Mars</a>	<a href="#">Mars.bSep08</a>	<a href="#">299851</a>	3621	1427	8	295	methionine-tRNA synthetase (Mars) alternative variant bSep08, mRNA.
<a href="#">Mars</a>	<a href="#">Mars.cSep08</a>	<a href="#">299851</a>	555	433	2	50	methionine-tRNA synthetase (5.9 kD) (Mars) alternative variant cSep08, mRNA.
<a href="#">Mars</a>	<a href="#">Mars.dSep08</a>	<a href="#">299851</a>	3274	303	2	33	methionine-tRNA synthetase (Mars) alternative variant dSep08, mRNA.
<a href="#">marsa</a>	<a href="#">marsa.aSep08</a>		8069	766		72	putative protein (8.4 kD) (marsa) mRNA.
<a href="#">marshee</a>	<a href="#">marshee.aSep08</a>		2042	717		46	putative protein (marshee) mRNA.
<a href="#">martu</a>	<a href="#">martu.aSep08</a>		2600	731		126	putative protein (14.2 kD) (martu) mRNA.
<a href="#">marvar</a>	<a href="#">marvar.aSep08</a>		7800	548		77	putative protein (marvar) mRNA.
<a href="#">marwey</a>	<a href="#">marwey.aSep08</a>		19900	817		107	putative protein (marwey) mRNA.
<a href="#">masa</a>	<a href="#">masa.aSep08</a>		2863	426		142	putative protein of mammalian origin (masa) mRNA.
<a href="#">mashee</a>	<a href="#">mashee.bSep08</a>		5590	681		19	putative protein (mashee) alternative variant bSep08, mRNA.

<a href="#">Masp1</a>	<a href="#">Masp1.bSep08</a>	<a href="#">64023</a>	4644	676	1	51	mannan-binding lectin serine peptidase 1 (Masp1) alternative variant bSep08, mRNA.
<a href="#">Masp2</a>	<a href="#">Masp2.bSep08</a>	<a href="#">64459</a>	1797	625	3	112	mannan-binding lectin serine 2 like (Masp2) alternative variant bSep08, mRNA.
<a href="#">Masp2</a>	<a href="#">Masp2.cSep08</a>	<a href="#">64459</a>	1962	829	2	88	putative protein (10.0 kD) (Masp2) alternative variant cSep08, mRNA.
<a href="#">MAST1</a>	<a href="#">MAST1.bSep08</a>	<a href="#">353118</a>	1713	272	3	90	microtubule associated serine/threonine kinase 1 (MAST1) alternative variant bSep08, mRNA.
<a href="#">Mast2</a>	<a href="#">Mast2.bSep08</a>	<a href="#">313819</a>	3366	2573	4	653	microtubule associated serine/threonine kinase 2 (Mast2) alternative variant bSep08, mRNA.
<a href="#">Mast2</a>	<a href="#">Mast2.cSep08</a>	<a href="#">313819</a>	816	732	2	117	microtubule associated serine/threonine kinase 2 (Mast2) alternative variant cSep08, mRNA.
<a href="#">Mast3</a>	<a href="#">Mast3.bSep08</a>	<a href="#">688540</a>	2230	1019	3	132	microtubule associated serine/threonine kinase 3 (Mast3) alternative variant bSep08, mRNA.
<a href="#">Mast4</a>	<a href="#">Mast4.aSep08</a>	<a href="#">310040</a>	325007	701		180	microtubule associated serine/threonine kinase family member 4 (Mast4) mRNA.
<a href="#">Mastl</a>	<a href="#">Mastl.bSep08</a>	<a href="#">307169</a>	17400	1513	1	409	microtubule associated serine/threonine kinase-like (Mastl) alternative variant bSep08, mRNA.
<a href="#">Mat1a</a>	<a href="#">Mat1a.bSep08</a>	<a href="#">25331</a>	2969	2393	1	65	methionine adenosyltransferase I, alpha (Mat1a) alternative variant bSep08, mRNA.
<a href="#">Mat2a</a>	<a href="#">Mat2a.bSep08</a>	<a href="#">171347</a>	2870	2222	5	236	methionine adenosyltransferase II, alpha (Mat2a) alternative variant bSep08, mRNA.
<a href="#">Mat2a</a>	<a href="#">Mat2a.cSep08</a>	<a href="#">171347</a>	2568	2235	3	199	methionine adenosyltransferase II, alpha (22.0 kD) (Mat2a) alternative variant cSep08, mRNA.
<a href="#">Mat2a</a>	<a href="#">Mat2a.eSep08</a>	<a href="#">171347</a>	4885	369	3	95	methionine adenosyltransferase II, alpha (Mat2a) alternative variant eSep08, mRNA.
<a href="#">Mat2b</a>	<a href="#">Mat2b.bSep08</a>	<a href="#">689330</a>	5233	1352	4	214	methionine adenosyltransferase II, beta (Mat2b) alternative variant bSep08, mRNA.
<a href="#">Matk</a>	<a href="#">Matk.bSep08</a>	<a href="#">60450</a>	2975	1080	6	217	megakaryocyte-associated tyrosine kinase (24.3 kD) (Matk) alternative variant bSep08, mRNA.
<a href="#">Matn1</a>	<a href="#">Matn1.bSep08</a>	<a href="#">297894</a>	2475	723	4	240	matrilin 1, cartilage matrix protein (Matn1) alternative variant bSep08, mRNA.
<a href="#">Matn2</a>	<a href="#">Matn2.aSep08</a>	<a href="#">299996</a>	54611	2250	13	562	matrilin 2 (Matn2) alternative variant aSep08, mRNA.
<a href="#">Matn2</a>	<a href="#">Matn2.bSep08</a>	<a href="#">299996</a>	4129	1457	2	149	matrilin 2 (Matn2) alternative variant bSep08, mRNA.
<a href="#">Matn2</a>	<a href="#">Matn2.cSep08</a>	<a href="#">299996</a>	3117	837	4	118	matrilin 2 (Matn2) alternative variant cSep08, mRNA.
<a href="#">Matn2</a>	<a href="#">Matn2.dSep08</a>	<a href="#">299996</a>	19944	278	3	92	matrilin 2 (Matn2) alternative variant dSep08, mRNA.
<a href="#">Matn3</a>	<a href="#">Matn3.bSep08</a>	<a href="#">313954</a>	7843	1317	3	302	matrilin 3 (Matn3) alternative variant bSep08, mRNA.
<a href="#">Matn3</a>	<a href="#">Matn3.cSep08</a>	<a href="#">313954</a>	14178	1134	7	248	matrilin 3 (Matn3) alternative variant cSep08, mRNA.
<a href="#">Matn3</a>	<a href="#">Matn3.dSep08</a>	<a href="#">313954</a>	8586	1204	5	248	matrilin 3 (Matn3) alternative variant dSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.bSep08</a>	<a href="#">29150</a>	19525	1680	9	481	matrin 3 (Matr3) alternative variant bSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.cSep08</a>	<a href="#">29150</a>	20022	870	9	290	matrin 3 (Matr3) alternative variant cSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.dSep08</a>	<a href="#">29150</a>	3579	758	2	165	matrin 3 (Matr3) alternative variant dSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.eSep08</a>	<a href="#">29150</a>	10187	707	2	148	matrin 3 (Matr3) alternative variant eSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.fSep08</a>	<a href="#">29150</a>	647	356	2	68	matrin 3 (Matr3) alternative variant fSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.gSep08</a>	<a href="#">29150</a>	1172	923	2	60	matrin 3 like (Matr3) alternative variant gSep08, mRNA.



<a href="#">Matr3</a>	<a href="#">Matr3.hSep08</a>	<a href="#">29150</a>	15635	367	2	39	matrin 3 (Matr3) alternative variant hSep08, mRNA.
<a href="#">Matr3</a>	<a href="#">Matr3.jSep08</a>	<a href="#">29150</a>	9718	519	2	88	matrin 3 (Matr3) alternative variant jSep08, mRNA.
<a href="#">matu</a>	<a href="#">matu.aSep08</a>		3290	269		77	putative protein of vertebrate origin (matu) mRNA.
<a href="#">mavar</a>	<a href="#">mavar.aSep08</a>		15396	304		40	putative protein (mavar) alternative variant aSep08, mRNA.
<a href="#">mavar</a>	<a href="#">mavar.bSep08</a>		15948	857		22	putative protein (mavar) alternative variant bSep08, mRNA.
<a href="#">mawby</a>	<a href="#">mawby.aSep08</a>		976	618		39	putative protein (3.9 kD) (mawby) mRNA.
<a href="#">mawchy</a>	<a href="#">mawchy.bSep08</a>		880	471	2	46	putative protein (5.5 kD) (mawchy) alternative variant bSep08, mRNA.
<a href="#">mawdar</a>	<a href="#">mawdar.bSep08</a>		815	251	2	59	putative protein (mawdar) alternative variant bSep08, mRNA.
<a href="#">mawey</a>	<a href="#">mawey.aSep08</a>		31167	1986	5	237	inositol 1 receptor CRA a (mawey) alternative variant aSep08, mRNA.
<a href="#">mawey</a>	<a href="#">mawey.bSep08</a>		98936	1015	7	93	putative protein (10.4 kD) (mawey) alternative variant bSep08, mRNA.
<a href="#">mawfer</a>	<a href="#">mawfer.aSep08</a>		9126	1989	2	439	peripheral benzodiazepine receptor associated protein (mawfer) alternative variant aSep08, mRNA.
<a href="#">mawflo</a>	<a href="#">mawflo.bSep08</a>		3132	400	2	102	putative protein (mawflo) alternative variant bSep08, mRNA.
<a href="#">mawflu</a>	<a href="#">mawflu.aSep08</a>		11684	353		46	putative protein (mawflu) mRNA.
<a href="#">mawgar</a>	<a href="#">mawgar.aSep08</a>		24916	297		93	putative protein (mawgar) mRNA.
<a href="#">mawjey</a>	<a href="#">mawjey.aSep08</a>		6502	1901		38	putative protein (mawjey) alternative variant aSep08, mRNA.
<a href="#">mawkee</a>	<a href="#">mawkee.aSep08</a>		26257	566		72	putative protein (8.4 kD) (mawkee) mRNA.
<a href="#">mawloy</a>	<a href="#">mawloy.bSep08</a>		2876	312	2	66	putative protein (mawloy) alternative variant bSep08, mRNA.
<a href="#">mawloy</a>	<a href="#">mawloy.cSep08</a>		1625	303	2	61	putative protein (mawloy) alternative variant cSep08, mRNA.
<a href="#">mawmee</a>	<a href="#">mawmee.aSep08</a>		2098	276		78	putative protein (mawmee) mRNA.
<a href="#">mawmer</a>	<a href="#">mawmer.aSep08</a>		1610	406	1	130	putative protein (mawmer) alternative variant aSep08, mRNA.
<a href="#">mawmer</a>	<a href="#">mawmer.bSep08</a>		1544	479	1	36	putative protein (mawmer) alternative variant bSep08, mRNA.
<a href="#">mawnoy</a>	<a href="#">mawnoy.aSep08</a>		27044	566	3	136	putative protein (mawnoy) alternative variant aSep08, mRNA.
<a href="#">mawnoy</a>	<a href="#">mawnoy.bSep08</a>		555	453	2	123	putative protein (mawnoy) alternative variant bSep08, mRNA.
<a href="#">mawnoy</a>	<a href="#">mawnoy.cSep08</a>		12334	916	3	117	putative protein (13.0 kD) (mawnoy) alternative variant cSep08, mRNA.
<a href="#">mawpor</a>	<a href="#">mawpor.aSep08</a>		7819	2170		95	coronin actin binding protein 2B like (mawpor) mRNA.
<a href="#">mawsa</a>	<a href="#">mawsa.aSep08</a>		3237	227		62	putative protein (7.0 kD) (mawsa) mRNA.
<a href="#">mawshee</a>	<a href="#">mawshee.aSep08</a>		12540	492	2	48	CRA a like (mawshee) alternative variant aSep08, mRNA.
<a href="#">mawshee</a>	<a href="#">mawshee.bSep08</a>		2283	383	1	48	CRA a like (mawshee) alternative variant bSep08, mRNA.
<a href="#">mawtu</a>	<a href="#">mawtu.aSep08</a>		33271	381		64	putative protein (mawtu) mRNA.
<a href="#">mawvar</a>	<a href="#">mawvar.aSep08</a>		10820	533			
<a href="#">mawwey</a>	<a href="#">mawwey.aSep08</a>		31028	457		57	putative protein (mawwey) mRNA.

<a href="#">Max</a>	<a href="#">Max.cSep08</a>	<a href="#">60661</a>	25392	1901	4	151	max protein (17.2 kD) (Max) alternative variant cSep08, complete mRNA.
<a href="#">Max</a>	<a href="#">Max.dSep08</a>	<a href="#">60661</a>	25371	1992	6	103	max protein (12.1 kD) (Max) alternative variant dSep08, complete mRNA.
<a href="#">Max</a>	<a href="#">Max.eSep08</a>	<a href="#">60661</a>	24133	727	5	94	max protein (11.0 kD) (Max) alternative variant eSep08, mRNA.
<a href="#">Max</a>	<a href="#">Max.fSep08</a>	<a href="#">60661</a>	24116	726	5	94	max protein (11.0 kD) (Max) alternative variant fSep08, complete mRNA.
<a href="#">Mbd1</a>	<a href="#">Mbd1.bSep08</a>	<a href="#">291439</a>	1293	463	3	154	methyl-CpG binding domain protein 1 (Mbd1) alternative variant bSep08, mRNA.
<a href="#">Mbd1</a>	<a href="#">Mbd1.cSep08</a>	<a href="#">291439</a>	1204	406	4	134	methyl-CpG binding domain protein 1 (Mbd1) alternative variant cSep08, mRNA.
<a href="#">Mbd1</a>	<a href="#">Mbd1.dSep08</a>	<a href="#">291439</a>	4349	3119	2	124	methyl-CpG binding domain protein 1 (Mbd1) alternative variant dSep08, mRNA.
<a href="#">Mbd1</a>	<a href="#">Mbd1.eSep08</a>	<a href="#">291439</a>	1391	313	2	93	methyl-CpG binding domain protein 1 (Mbd1) alternative variant eSep08, mRNA.
<a href="#">Mbd1</a>	<a href="#">Mbd1.gSep08</a>	<a href="#">291439</a>	2741	1412	2	51	methyl-CpG binding domain protein 1 (5.5 kD) (Mbd1) alternative variant gSep08, mRNA.
<a href="#">Mbd2</a>	<a href="#">Mbd2.bSep08</a>	<a href="#">680172</a>	54401	693	6	201	methyl-CpG binding domain protein 2 (Mbd2) alternative variant bSep08, mRNA.
<a href="#">MBD3</a>	<a href="#">MBD3.bSep08</a>	<a href="#">362834</a>	7197	1566	5	352	methyl-CpG binding domain protein 3 (MBD3) alternative variant bSep08, mRNA.
<a href="#">MBD3</a>	<a href="#">MBD3.cSep08</a>	<a href="#">362834</a>	6916	1405	6	345	methyl-CpG binding domain protein 3 (MBD3) alternative variant cSep08, mRNA.
<a href="#">MBD3</a>	<a href="#">MBD3.dSep08</a>	<a href="#">362834</a>	5670	751	4	250	methyl-CpG binding domain protein 3 (MBD3) alternative variant dSep08, mRNA.
<a href="#">MBD3</a>	<a href="#">MBD3.eSep08</a>	<a href="#">362834</a>	1181	416	1	138	methyl-CpG binding domain protein 3 (MBD3) alternative variant eSep08, mRNA.
<a href="#">Mbd4</a>	<a href="#">Mbd4.aSep08</a>	<a href="#">680915</a>	3699	1404		178	methyl-CpG binding domain protein 4 (Mbd4) alternative variant aSep08, mRNA.
<a href="#">Mbd4</a>	<a href="#">Mbd4.bSep08</a>	<a href="#">680915</a>	9898	1619		169	methyl-CpG binding domain protein 4 (20.2 kD) (Mbd4) alternative variant bSep08, mRNA.
<a href="#">Mbd6</a>	<a href="#">Mbd6.aSep08</a>	<a href="#">362892</a>	3739	2643	1	730	methyl-CpG binding domain protein 6 (Mbd6) alternative variant aSep08, mRNA.
<a href="#">Mbd6</a>	<a href="#">Mbd6.bSep08</a>	<a href="#">362892</a>	1825	1371	1	104	methyl-CpG binding domain protein 6 (Mbd6) alternative variant bSep08, mRNA.
<a href="#">Mbip</a>	<a href="#">Mbip.bSep08</a>	<a href="#">362740</a>	7220	659	3	78	MAP3K12 binding inhibitory protein 1 (Mbip) alternative variant bSep08, mRNA.
<a href="#">Mbl2</a>	<a href="#">Mbl2.aSep08</a>	<a href="#">64668</a>	7127	1072	5	244	mannose binding lectin 2 (protein C) (26.0 kD) (Mbl2) alternative variant aSep08, mRNA.
<a href="#">Mbl2</a>	<a href="#">Mbl2.cSep08</a>	<a href="#">64668</a>	6818	694	4	152	mannose binding lectin 2 (protein C) (Mbl2) alternative variant cSep08, mRNA.
<a href="#">Mbl2</a>	<a href="#">Mbl2.dSep08</a>	<a href="#">64668</a>	38634	498	2	37	mannose binding lectin 2 (protein C) (Mbl2) alternative variant dSep08, mRNA.
<a href="#">Mbnl1</a>	<a href="#">Mbnl1.aSep08</a>	<a href="#">282635</a>	139746	1106	6	287	muscleblind-like 1 (Drosophila) (Mbnl1) alternative variant aSep08, mRNA.

<a href="#">Mbnl1</a>	<a href="#">Mbnl1.bSep08</a>	<a href="#">282635</a>	29244	863	3	271	muscleblind-like 1 (Drosophila) (Mbnl1) alternative variant bSep08, mRNA.
<a href="#">Mbnl1</a>	<a href="#">Mbnl1.cSep08</a>	<a href="#">282635</a>	19783	718	3	239	muscleblind-like 1 (Drosophila) (Mbnl1) alternative variant cSep08, mRNA.
<a href="#">Mbnl1</a>	<a href="#">Mbnl1.dSep08</a>	<a href="#">282635</a>	8068	380	1	126	muscleblind-like 1 (Drosophila) (Mbnl1) alternative variant dSep08, mRNA.
<a href="#">Mbnl3</a>	<a href="#">Mbnl3.bSep08</a>	<a href="#">302492</a>	80774	801	2	266	muscleblind-like 3 (Drosophila) (Mbnl3) alternative variant bSep08, mRNA.
<a href="#">Mbnl3</a>	<a href="#">Mbnl3.cSep08</a>	<a href="#">302492</a>	24058	738	3	245	muscleblind-like 3 (Drosophila) (Mbnl3) alternative variant cSep08, mRNA.
<a href="#">Mbnl3</a>	<a href="#">Mbnl3.dSep08</a>	<a href="#">302492</a>	19833	716	2	238	muscleblind-like 3 (Drosophila) (Mbnl3) alternative variant dSep08, mRNA.
<a href="#">Mbnl3</a>	<a href="#">Mbnl3.eSep08</a>	<a href="#">302492</a>	67436	605	1	171	muscleblind-like 3 (Drosophila) (Mbnl3) alternative variant eSep08, mRNA.
<a href="#">MBOAT.0</a>	<a href="#">MBOAT.0.aSep08</a>		14335	2345	8	473	leukocyte receptor cluster member 4 (53.4 kD) (MBOAT.0) alternative variant aSep08, mRNA.
<a href="#">MBOAT.0</a>	<a href="#">MBOAT.0.bSep08</a>		9056	3031	3	203	leukocyte receptor cluster member 4 (23.4 kD) (MBOAT.0) alternative variant bSep08, mRNA.
<a href="#">Mboat2</a>	<a href="#">Mboat2.bSep08</a>	<a href="#">313997</a>	109424	600		199	putative protein of eukaryotic origin (Mboat2) alternative variant bSep08, mRNA.
<a href="#">Mboat4</a>	<a href="#">Mboat4.aSep08</a>	<a href="#">306515</a>	8295	1308	3	435	membrane bound O-acyl transferase, MBOAT (Mboat4) alternative variant aSep08, mRNA.
<a href="#">Mbp</a>	<a href="#">Mbp.gSep08</a>	<a href="#">24547</a>	26309	1021	2	146	putative protein (15.8 kD) (Mbp) alternative variant gSep08, mRNA.
<a href="#">Mbp</a>	<a href="#">Mbp.hSep08</a>	<a href="#">24547</a>	30613	798	4	140	myelin basic protein (Mbp) alternative variant hSep08, mRNA.
<a href="#">Mbp</a>	<a href="#">Mbp.iSep08</a>	<a href="#">24547</a>	21053	398	4	120	myelin basic protein (Mbp) alternative variant iSep08, mRNA.
<a href="#">Mbp</a>	<a href="#">Mbp.jSep08</a>	<a href="#">24547</a>	30604	780	3	114	myelin basic protein (12.2 kD) (Mbp) alternative variant jSep08, mRNA.
<a href="#">Mbrl</a>	<a href="#">Mbrl.aSep08</a>	<a href="#">299608</a>	6695	2156	6	588	membralin (Mbrl) alternative variant aSep08, mRNA.
<a href="#">Mbrl</a>	<a href="#">Mbrl.bSep08</a>	<a href="#">299608</a>	2108	1295	6	431	membralin (Mbrl) alternative variant bSep08, mRNA.
<a href="#">Mbrl</a>	<a href="#">Mbrl.cSep08</a>	<a href="#">299608</a>	970	462	2	154	membralin (Mbrl) alternative variant cSep08, mRNA.
<a href="#">Mbrl</a>	<a href="#">Mbrl.dSep08</a>	<a href="#">299608</a>	2652	1988	2	86	membralin (Mbrl) alternative variant dSep08, mRNA.
<a href="#">Mbtps1</a>	<a href="#">Mbtps1.bSep08</a>	<a href="#">89842</a>	12159	887	5	211	membrane-bound transcription factor peptidase, site 1 (Mbtps1) alternative variant bSep08, mRNA.
<a href="#">Mbtps1</a>	<a href="#">Mbtps1.cSep08</a>	<a href="#">89842</a>	2321	737	2	42	membrane-bound transcription factor peptidase, site 1 (Mbtps1) alternative variant cSep08, mRNA.
<a href="#">Mbtps2</a>	<a href="#">Mbtps2.bSep08</a>	<a href="#">302705</a>	7983	1803	1	164	membrane-bound transcription factor peptidase, site 2 (Mbtps2) alternative variant bSep08, mRNA.
<a href="#">Mc2r</a>	<a href="#">Mc2r.aSep08</a>	<a href="#">282839</a>	9784	435	1	62	melanocortin 2 receptor (Mc2r) alternative variant aSep08, mRNA.
<a href="#">Mc2r</a>	<a href="#">Mc2r.bSep08</a>	<a href="#">282839</a>	9797	405	1	70	melanocortin 2 receptor (Mc2r) alternative variant bSep08, mRNA.
<a href="#">Mcam</a>	<a href="#">Mcam.cSep08</a>	<a href="#">78967</a>	519	407	2	97	melanoma cell adhesion molecule (Mcam) alternative variant cSep08, mRNA.



<a href="#">Mcm7</a>	<a href="#">Mcm7.dSep08</a>	<a href="#">288532</a>	1494	491	3	96	putative protein of eukaryotic origin (Mcm7) alternative variant dSep08, mRNA.
<a href="#">Mcm10</a>	<a href="#">Mcm10.bSep08</a>	<a href="#">307126</a>	21664	1482	4	37	putative protein (Mcm10) alternative variant bSep08, mRNA.
<a href="#">Mcoln1</a>	<a href="#">Mcoln1.bSep08</a>	<a href="#">288371</a>	3122	1288	6	265	mucolin 1 (Mcoln1) alternative variant bSep08, mRNA.
<a href="#">Mcoln1</a>	<a href="#">Mcoln1.dSep08</a>	<a href="#">288371</a>	3917	441	3	93	mucolin 1 (Mcoln1) alternative variant dSep08, mRNA.
<a href="#">Mcoln3</a>	<a href="#">Mcoln3.bSep08</a>	<a href="#">308022</a>	6160	2004	3	154	mucolin 3 (Mcoln3) alternative variant bSep08, mRNA.
<a href="#">Mcph1</a>	<a href="#">Mcph1.aSep08</a>	<a href="#">306594</a>	20173	301		100	microcephaly, primary autosomal recessive 1 (Mcph1) mRNA.
<a href="#">Mcpt8l2</a>	<a href="#">Mcpt8l2.aSep08</a>	<a href="#">408240</a>	1609	664		193	mast cell protease 8-like 2 (Mcpt8l2) mRNA.
<a href="#">Mcpt9</a>	<a href="#">Mcpt9.bSep08</a>	<a href="#">54272</a>	2957	1375	1	74	mast cell protease 9 (8.2 kD) (Mcpt9) alternative variant bSep08, mRNA.
<a href="#">Mcrcs1</a>	<a href="#">Mcrcs1.cSep08</a>	<a href="#">300222</a>	3173	487	4	48	microspherule protein 1 (5.3 kD) (Mcrcs1) alternative variant cSep08, mRNA.
<a href="#">Mcrcs1</a>	<a href="#">Mcrcs1.dSep08</a>	<a href="#">300222</a>	1714	412	2	43	microspherule protein 1 (4.7 kD) (Mcrcs1) alternative variant dSep08, mRNA.
<a href="#">Mctp1</a>	<a href="#">Mctp1.aSep08</a>	<a href="#">309928</a>	233924	879		252	multiple C2 domains, transmembrane 1 (29.0 kD) (Mctp1) mRNA.
<a href="#">Mcts1</a>	<a href="#">Mcts1.bSep08</a>	<a href="#">302500</a>	11532	789	6	181	malignant T cell amplified sequence 1 (20.6 kD) (Mcts1) alternative variant bSep08, mRNA.
<a href="#">Mcts1</a>	<a href="#">Mcts1.cSep08</a>	<a href="#">302500</a>	9644	741	5	156	malignant T cell amplified sequence 1 (17.5 kD) (Mcts1) alternative variant cSep08, mRNA.
<a href="#">Mcts1</a>	<a href="#">Mcts1.dSep08</a>	<a href="#">302500</a>	11294	673	6	138	malignant T cell amplified sequence 1 (15.5 kD) (Mcts1) alternative variant dSep08, mRNA.
<a href="#">Mcts1</a>	<a href="#">Mcts1.eSep08</a>	<a href="#">302500</a>	1724	505	2	47	malignant T cell amplified sequence 1 (5.3 kD) (Mcts1) alternative variant eSep08, mRNA.
<a href="#">Mdc1</a>	<a href="#">Mdc1.aSep08</a>	<a href="#">309595</a>	6520	3058		671	mediator of DNA damage checkpoint 1 (71.9 kD) (Mdc1) mRNA.
<a href="#">Mdfic</a>	<a href="#">Mdfic.bSep08</a>	<a href="#">362325</a>	73357	708	3	184	putative nuclear protein of vertebrate origin (19.9 kD) (Mdfic) alternative variant bSep08, mRNA.
<a href="#">Mdfic</a>	<a href="#">Mdfic.cSep08</a>	<a href="#">362325</a>	49931	760	3	134	putative protein of vertebrate origin (Mdfic) alternative variant cSep08, mRNA.
<a href="#">Mdga1</a>	<a href="#">Mdga1.bSep08</a>	<a href="#">309659</a>	8919	982	4	194	glycosylphosphatidylinositol anchor 1 (Mdga1) alternative variant bSep08, mRNA.
<a href="#">Mdga2</a>	<a href="#">Mdga2.bSep08</a>	<a href="#">314180</a>	100806	780	2	259	MAM (Mdga2) alternative variant bSep08, mRNA.
<a href="#">Mdh1</a>	<a href="#">Mdh1.cSep08</a>	<a href="#">24551</a>	1182	570	2	66	malate dehydrogenase 1, NAD (soluble) (7.4 kD) (Mdh1) alternative variant cSep08, mRNA.
<a href="#">Mdh2</a>	<a href="#">Mdh2.bSep08</a>	<a href="#">81829</a>	5849	742	5	247	malate dehydrogenase 2, NAD (mitochondrial) (Mdh2) alternative variant bSep08, mRNA.
<a href="#">Mdh2</a>	<a href="#">Mdh2.cSep08</a>	<a href="#">81829</a>	11872	746	6	215	malate dehydrogenase 2, NAD (mitochondrial) (Mdh2) alternative variant cSep08, mRNA.
<a href="#">Mdk</a>	<a href="#">Mdk.aSep08</a>	<a href="#">81517</a>	1779	658	5	142	midkine (15.7 kD) (Mdk) alternative variant aSep08, mRNA.
<a href="#">Mdk</a>	<a href="#">Mdk.cSep08</a>	<a href="#">81517</a>	1929	746	5	140	midkine (Mdk) alternative variant cSep08, complete mRNA.
<a href="#">Mdk</a>	<a href="#">Mdk.dSep08</a>	<a href="#">81517</a>	1923	732	5	140	midkine (Mdk) alternative variant dSep08, complete mRNA.
<a href="#">Mdk</a>	<a href="#">Mdk.eSep08</a>	<a href="#">81517</a>	851	732	2	117	midkine (Mdk) alternative variant eSep08, mRNA.

<a href="#">Mdm1</a>	<a href="#">Mdm1.bSep08</a>	<a href="#">314859</a>	1782	528	3	89	transformed mouse 3T3 cell double minute 1 (Mdm1) alternative variant bSep08, mRNA.
<a href="#">Mdm1</a>	<a href="#">Mdm1.dSep08</a>	<a href="#">314859</a>	11297	735	6	59	transformed mouse 3T3 cell double minute 1 (6.8 kD) (Mdm1) alternative variant dSep08, mRNA.
<a href="#">Mdm2</a>	<a href="#">Mdm2.bSep08</a>	<a href="#">314856</a>	16003	731	2	243	transformed mouse 3T3 cell double minute 2 (Mdm2) alternative variant bSep08, mRNA.
<a href="#">Mdm2</a>	<a href="#">Mdm2.cSep08</a>	<a href="#">314856</a>	16604	1177	2	211	transformed mouse 3T3 cell double minute 2 (Mdm2) alternative variant cSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.bSep08</a>	<a href="#">304798</a>	36739	2634	7	362	mouse double minute 4 like (41.2 kD) (Mdm4) alternative variant bSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.cSep08</a>	<a href="#">304798</a>	35551	1695	10	212	mouse double minute 4 like (Mdm4) alternative variant cSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.dSep08</a>	<a href="#">304798</a>	10309	1438	4	196	mouse double minute 4 like (22.0 kD) (Mdm4) alternative variant dSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.eSep08</a>	<a href="#">304798</a>	29482	781	8	127	mouse double minute 4 like (14.1 kD) (Mdm4) alternative variant eSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.fSep08</a>	<a href="#">304798</a>	19538	448	5	111	mouse double minute 4 like (Mdm4) alternative variant fSep08, mRNA.
<a href="#">Mdm4</a>	<a href="#">Mdm4.gSep08</a>	<a href="#">304798</a>	1838	430	2	100	mouse double minute 4 like (Mdm4) alternative variant gSep08, mRNA.
<a href="#">Mdp-1</a>	<a href="#">Mdp-1.bSep08</a>	<a href="#">290230</a>	1209	418	1	106	magnesium-dependent phosphatase 1 (12.2 kD) (Mdp-1) alternative variant bSep08, mRNA.
<a href="#">Me1</a>	<a href="#">Me1.bSep08</a>	<a href="#">24552</a>	84821	822	8	182	malic enzyme 1, NADP(+)-dependent, cytosolic (Me1) alternative variant bSep08, mRNA.
<a href="#">Me2</a>	<a href="#">Me2.aSep08</a>	<a href="#">307270</a>	44256	1856		599	malic enzyme 2, NAD(+)-dependent, mitochondrial (Me2) mRNA.
<a href="#">Me3</a>	<a href="#">Me3.aSep08</a>	<a href="#">361602</a>	198042	1363	9	429	malic enzyme 3, NADP(+)-dependent, mitochondrial (Me3) alternative variant aSep08, mRNA.
<a href="#">Mea1</a>	<a href="#">Mea1.aSep08</a>	<a href="#">685131</a>	2126	1597	3	208	male-enhanced antigen 1 (22.4 kD) (Mea1) alternative variant aSep08, mRNA.
<a href="#">Mea1</a>	<a href="#">Mea1.bSep08</a>	<a href="#">685131</a>	1602	667	4	189	male-enhanced antigen 1 (Mea1) alternative variant bSep08, mRNA.
<a href="#">Mea1</a>	<a href="#">Mea1.dSep08</a>	<a href="#">685131</a>	9579	341	2	113	male-enhanced antigen 1 (Mea1) alternative variant dSep08, mRNA.
<a href="#">Mecr</a>	<a href="#">Mecr.bSep08</a>	<a href="#">29470</a>	5789	1119	4	92	mitochondrial trans-2-enoyl-CoA reductase (10.5 kD) (Mecr) alternative variant bSep08, mRNA.
<a href="#">Mecr</a>	<a href="#">Mecr.cSep08</a>	<a href="#">29470</a>	7364	610	5	77	mitochondrial trans-2-enoyl-CoA reductase (Mecr) alternative variant cSep08, mRNA.
<a href="#">Mecr</a>	<a href="#">Mecr.dSep08</a>	<a href="#">29470</a>	12148	557	4	88	mitochondrial trans-2-enoyl-CoA reductase (Mecr) alternative variant dSep08, mRNA.
<a href="#">Med1</a>	<a href="#">Med1.bSep08</a>	<a href="#">497991</a>	28784	2603	10	539	mediator complex subunit 1 (Med1) alternative variant bSep08, mRNA.
<a href="#">Med1</a>	<a href="#">Med1.cSep08</a>	<a href="#">497991</a>	11626	848	5	152	mediator complex subunit 1 (Med1) alternative variant cSep08, mRNA.
<a href="#">Med6</a>	<a href="#">Med6.bSep08</a>	<a href="#">299180</a>	12696	713	7	212	mediator of RNA polymerase II transcription, subunit 6 homolog (yeast) (Med6) alternative variant bSep08, mRNA.

<a href="#">MED7.0</a>	<a href="#">MED7.0.aSep08</a>		4161	1177	3	233	mediator complex (27.2 kD) (MED7.0) alternative variant aSep08, mRNA.
<a href="#">MED7.0</a>	<a href="#">MED7.0.cSep08</a>		3574	770	3	128	mediator complex (MED7.0) alternative variant cSep08, mRNA.
<a href="#">MED7.0</a>	<a href="#">MED7.0.dSep08</a>		14182	699	4	66	putative protein (7.6 kD) (MED7.0) alternative variant dSep08, mRNA.
<a href="#">Med8</a>	<a href="#">Med8.aSep08</a>	<a href="#">362575</a>	6324	1945	7	276	mediator of RNA polymerase II transcription, subunit 8 homolog (yeast) (Med8) alternative variant aSep08, mRNA.
<a href="#">Med10</a>	<a href="#">Med10.aSep08</a>	<a href="#">290939</a>	5660	843		149	mediator of RNA polymerase II transcription, subunit 10 homolog (NUT2, <i>S. cerevisiae</i> ) (Med10) mRNA.
<a href="#">Med11</a>	<a href="#">Med11.bSep08</a>	<a href="#">287456</a>	1809	919	1	137	mediator of RNA polymerase II transcription, subunit 11 homolog ( <i>S. cerevisiae</i> ) (Med11) alternative variant bSep08, mRNA.
<a href="#">Med13</a>	<a href="#">Med13.bSep08</a>	<a href="#">303403</a>	8548	154	1	51	mediator complex subunit 13 (Med13) alternative variant bSep08, mRNA.
<a href="#">Med13l</a>	<a href="#">Med13l.bSep08</a>	<a href="#">360817</a>	13407	1963	7	474	mediator complex subunit 13-like (Med13l) alternative variant bSep08, mRNA.
<a href="#">Med13l</a>	<a href="#">Med13l.cSep08</a>	<a href="#">360817</a>	2415	910	4	215	mediator complex subunit 13-like (Med13l) alternative variant cSep08, mRNA.
<a href="#">Med14</a>	<a href="#">Med14.aSep08</a>	<a href="#">317343</a>	16861	945		315	mediator complex subunit 14 (Med14) mRNA.
<a href="#">Med15</a>	<a href="#">Med15.bSep08</a>	<a href="#">360743</a>	2730	1006	7	311	mediator complex (Med15) alternative variant bSep08, mRNA.
<a href="#">Med15</a>	<a href="#">Med15.cSep08</a>	<a href="#">360743</a>	19323	759	5	252	putative protein (Med15) alternative variant cSep08, mRNA.
<a href="#">Med15</a>	<a href="#">Med15.dSep08</a>	<a href="#">360743</a>	4431	2272	6	223	mediator complex (25.0 kD) (Med15) alternative variant dSep08, mRNA.
<a href="#">Med15</a>	<a href="#">Med15.eSep08</a>	<a href="#">360743</a>	13990	2000	5	211	mediator complex (Med15) alternative variant eSep08, mRNA.
<a href="#">Med15</a>	<a href="#">Med15.fSep08</a>	<a href="#">360743</a>	829	382	2	93	mediator complex (Med15) alternative variant fSep08, mRNA.
<a href="#">Med15</a>	<a href="#">Med15.gSep08</a>	<a href="#">360743</a>	41185	1355	4	84	mediator complex (9.5 kD) (Med15) alternative variant gSep08, complete mRNA.
<a href="#">Med15</a>	<a href="#">Med15.hSep08</a>	<a href="#">360743</a>	708	621	2	68	putative protein (10.1 kD) (Med15) alternative variant hSep08, mRNA.
<a href="#">Med16</a>	<a href="#">Med16.bSep08</a>	<a href="#">299607</a>	4379	822	6	273	mediator complex subunit 16 (Med16) alternative variant bSep08, mRNA.
<a href="#">Med16</a>	<a href="#">Med16.cSep08</a>	<a href="#">299607</a>	2474	880	5	256	mediator complex subunit 16 (Med16) alternative variant cSep08, mRNA.
<a href="#">Med16</a>	<a href="#">Med16.eSep08</a>	<a href="#">299607</a>	4432	739	5	141	mediator complex subunit 16 (Med16) alternative variant eSep08, mRNA.
<a href="#">Med16</a>	<a href="#">Med16.fSep08</a>	<a href="#">299607</a>	1456	397	4	106	mediator complex subunit 16 (Med16) alternative variant fSep08, mRNA.
<a href="#">Med17</a>	<a href="#">Med17.bSep08</a>	<a href="#">300367</a>	9955	1800	5	181	mediator complex subunit 17 (Med17) alternative variant bSep08, mRNA.
<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.bSep08</a>	<a href="#">316209</a>	11545	1582	2	123	mediator complex (Med20andUsp49) alternative variant bSep08, mRNA.

<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.bSep08</a>	<a href="#">316211</a>	11545	1582	2	123	mediator complex (Med20andUsp49) alternative variant bSep08, mRNA.
<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.cSep08</a>	<a href="#">316209</a>	65485	327	2	63	putative protein (Med20andUsp49) alternative variant cSep08, mRNA.
<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.cSep08</a>	<a href="#">316211</a>	65485	327	2	63	putative protein (Med20andUsp49) alternative variant cSep08, mRNA.
<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.dSep08</a>	<a href="#">316209</a>	45961	593	3	48	ubiquitin 49 (Med20andUsp49) alternative variant dSep08, mRNA.
<a href="#">Med20andUsp49</a>	<a href="#">Med20andUsp49.dSep08</a>	<a href="#">316211</a>	45961	593	3	48	ubiquitin 49 (Med20andUsp49) alternative variant dSep08, mRNA.
<a href="#">Med21</a>	<a href="#">Med21.aSep08</a>	<a href="#">312849</a>	6959	771	3	144	mediator complex subunit 21 (15.6 kD) (Med21) alternative variant aSep08, complete mRNA.
<a href="#">Med21</a>	<a href="#">Med21.cSep08</a>	<a href="#">312849</a>	23188	494	4	86	mediator complex subunit 21 (Med21) alternative variant cSep08, mRNA.
<a href="#">Med21</a>	<a href="#">Med21.dSep08</a>	<a href="#">312849</a>	6933	529	1	72	mediator complex subunit 21 (7.9 kD) (Med21) alternative variant dSep08, complete mRNA.
<a href="#">Med22</a>	<a href="#">Med22.bSep08</a>	<a href="#">499762</a>	2570	858	3	140	mediator complex subunit 22 (16.4 kD) (Med22) alternative variant bSep08, mRNA.
<a href="#">Med23</a>	<a href="#">Med23.aSep08</a>	<a href="#">309565</a>	20818	3416	14	960	mediator complex subunit 23 (Med23) alternative variant aSep08, mRNA.
<a href="#">Med23</a>	<a href="#">Med23.bSep08</a>	<a href="#">309565</a>	3413	544	1	181	mediator complex subunit 23 (Med23) alternative variant bSep08, mRNA.
<a href="#">Med24</a>	<a href="#">Med24.bSep08</a>	<a href="#">619436</a>	12530	879	6	206	mediator complex subunit 24 (23.0 kD) (Med24) alternative variant bSep08, mRNA.
<a href="#">Med24</a>	<a href="#">Med24.cSep08</a>	<a href="#">619436</a>	2690	601	3	197	mediator complex subunit 24 (Med24) alternative variant cSep08, mRNA.
<a href="#">Med25</a>	<a href="#">Med25.aSep08</a>	<a href="#">292889</a>	15616	3785	18	802	mediator of RNA polymerase II transcription homolog CRA b (85.3 kD) (Med25) alternative variant aSep08, complete mRNA.
<a href="#">Med25</a>	<a href="#">Med25.bSep08</a>	<a href="#">292889</a>	12822	2180	17	426	mediator of RNA polymerase II transcription homolog CRA b (44.5 kD) (Med25) alternative variant bSep08, mRNA.
<a href="#">Med25</a>	<a href="#">Med25.cSep08</a>	<a href="#">292889</a>	3684	881	6	281	mediator of RNA polymerase II transcription homolog CRA b (Med25) alternative variant cSep08, mRNA.
<a href="#">Med25</a>	<a href="#">Med25.dSep08</a>	<a href="#">292889</a>	1405	847	3	185	mediator of RNA polymerase II transcription homolog CRA b (18.8 kD) (Med25) alternative variant dSep08, mRNA.
<a href="#">Med25</a>	<a href="#">Med25.eSep08</a>	<a href="#">292889</a>	6428	538	4	134	mediator of RNA polymerase II transcription homolog CRA b (Med25) alternative variant eSep08, mRNA.
<a href="#">Med26</a>	<a href="#">Med26.aSep08</a>	<a href="#">306328</a>	48055	506		79	mediator complex subunit 26 (Med26) mRNA.
<a href="#">Med30</a>	<a href="#">Med30.aSep08</a>	<a href="#">299905</a>	21845	998	2	202	mediator complex subunit 30 (Med30) alternative variant aSep08, mRNA.
<a href="#">Med31</a>	<a href="#">Med31.aSep08</a>	<a href="#">287475</a>	3402	646	4	156	mediator of RNA polymerase II transcription, subunit 31 homolog (yeast) (Med31) alternative variant aSep08, mRNA.
<a href="#">Med31</a>	<a href="#">Med31.bSep08</a>	<a href="#">287475</a>	2438	572	3	133	mediator of RNA polymerase II transcription, subunit 31 homolog (yeast) (Med31) alternative variant bSep08, mRNA.



<a href="#">Med31</a>	<a href="#">Med31.cSep08</a>	<a href="#">287475</a>	2030	867	3	100	mediator of RNA polymerase II transcription, subunit 31 homolog (yeast) (Med31) alternative variant cSep08, mRNA.
<a href="#">meeby</a>	<a href="#">meeby.aSep08</a>		168904	1797		526	cyclin-dependent kinase-like 5 (meeby) mRNA.
<a href="#">meechy</a>	<a href="#">meechy.aSep08</a>		867	381		25	putative protein (meechy) mRNA.
<a href="#">meedar</a>	<a href="#">meedar.aSep08</a>		545	415		97	HEAT repeat containing 3 (meedar) mRNA.
<a href="#">meefer</a>	<a href="#">meefer.aSep08</a>		16894	582		118	putative cytoplasmic protein (13.1 kD) (meefer) mRNA.
<a href="#">meeflo</a>	<a href="#">meeflo.aSep08</a>		3621	792		38	putative protein (meeflo) mRNA.
<a href="#">meeflu</a>	<a href="#">meeflu.aSep08</a>		11744	392		38	putative protein (meeflu) mRNA.
<a href="#">meegar</a>	<a href="#">meegar.aSep08</a>		9460	907		300	CRA a (meegar) mRNA.
<a href="#">meejey</a>	<a href="#">meejey.aSep08</a>		1625	656		28	putative protein (3.4 kD) (meejey) mRNA.
<a href="#">meekee</a>	<a href="#">meekee.aSep08</a>		486	348		41	putative protein (4.7 kD) (meekee) mRNA.
<a href="#">meeloy</a>	<a href="#">meeloy.aSep08</a>		12317	1062		76	putative secreted or extracellular protein precursor (8.3 kD) (meeloy) mRNA.
<a href="#">meemee</a>	<a href="#">meemee.aSep08</a>		2568	556	3	53	putative protein (6.0 kD) (meemee) alternative variant aSep08, mRNA.
<a href="#">meemer</a>	<a href="#">meemer.aSep08</a>		5528	730		80	putative protein (8.7 kD) (meemer) mRNA.
<a href="#">meenoy</a>	<a href="#">meenoy.aSep08</a>		19823	390		68	putative protein (meenoy) mRNA.
<a href="#">meepor</a>	<a href="#">meepor.aSep08</a>		694	580		59	putative protein (meepor) mRNA.
<a href="#">meesa</a>	<a href="#">meesa.aSep08</a>		3245	3108		118	putative protein (13.0 kD) (meesa) mRNA.
<a href="#">meeshee</a>	<a href="#">meeshee.aSep08</a>		4317	669		96	ab2-143 like (10.7 kD) (meeshee) mRNA.
<a href="#">meetu</a>	<a href="#">meetu.aSep08</a>		5269	282		93	s1 rna binding domain 1 like (meetu) mRNA.
<a href="#">meevar</a>	<a href="#">meevar.aSep08</a>		1913	331		24	putative protein (meevar) mRNA.
<a href="#">meewey</a>	<a href="#">meewey.aSep08</a>		5652	1412	2	105	putative protein (meewey) mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.bSep08</a>	<a href="#">498607</a>	7113	899	6	171	putative protein of eukaryotic origin (Mef2b) alternative variant bSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.bSep08</a>	<a href="#">688966</a>	7113	899	6	171	putative protein of eukaryotic origin (Mef2b) alternative variant bSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.cSep08</a>	<a href="#">498607</a>	4663	513	5	142	putative protein of eukaryotic origin (Mef2b) alternative variant cSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.cSep08</a>	<a href="#">688966</a>	4663	513	5	142	putative protein of eukaryotic origin (Mef2b) alternative variant cSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.dSep08</a>	<a href="#">498607</a>	5209	427	5	134	MADS box transcription enhancer factor 2 polypeptide B like (Mef2b) alternative variant dSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.dSep08</a>	<a href="#">688966</a>	5209	427	5	134	MADS box transcription enhancer factor 2 polypeptide B like (Mef2b) alternative variant dSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.eSep08</a>	<a href="#">498607</a>	4899	808	4	103	MADS box transcription enhancer factor 2 polypeptide B (Mef2b) alternative variant eSep08, mRNA.
<a href="#">Mef2b</a>	<a href="#">Mef2b.eSep08</a>	<a href="#">688966</a>	4899	808	4	103	MADS box transcription enhancer factor 2 polypeptide B (Mef2b) alternative variant eSep08, mRNA.
<a href="#">Mef2d</a>	<a href="#">Mef2d.bSep08</a>	<a href="#">81518</a>	8905	772	7	257	myocyte enhancer factor 2D (Mef2d) alternative variant bSep08, mRNA.
<a href="#">Mef2d</a>	<a href="#">Mef2d.cSep08</a>	<a href="#">81518</a>	1176	703	2	112	myocyte enhancer factor 2D (13.0 kD) (Mef2d) alternative variant cSep08, mRNA.

<a href="#">Megf10</a>	<a href="#">Megf10.aSep08</a>	<a href="#">291445</a>	1872	852		83	multiple EGF-like domains 10 (Megf10) mRNA.
<a href="#">Megf11</a>	<a href="#">Megf11.aSep08</a>	<a href="#">691517</a>	50874	1350	8	450	multiple EGF-like-domains 11 (Megf11) alternative variant aSep08, mRNA.
<a href="#">Megf11</a>	<a href="#">Megf11.bSep08</a>	<a href="#">691517</a>	1304	980	1	253	multiple EGF-like-domains 11 (Megf11) alternative variant bSep08, mRNA.
<a href="#">Mei1</a>	<a href="#">Mei1.bSep08</a>	<a href="#">315162</a>	20677	1789	1	485	meiosis defective 1 (Mei1) alternative variant bSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.bSep08</a>	<a href="#">311311</a>	195698	2146	8	320	meis1, myeloid ecotropic viral integration site 1 homolog 2 (Meis2) alternative variant bSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.cSep08</a>	<a href="#">311311</a>	6701	826	5	165	meis1, myeloid ecotropic viral integration site 1 homolog 2 (Meis2) alternative variant cSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.dSep08</a>	<a href="#">311311</a>	137452	853	6	126	meis1, myeloid ecotropic viral integration site 1 homolog 2 (Meis2) alternative variant dSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.fSep08</a>	<a href="#">311311</a>	138478	1858	6	116	meis1, myeloid ecotropic viral integration site 1 homolog 2 (Meis2) alternative variant fSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.gSep08</a>	<a href="#">311311</a>	47370	756	2	91	meis1, myeloid ecotropic viral integration site 1 homolog 2 (9.9 kD) (Meis2) alternative variant gSep08, mRNA.
<a href="#">Meis2</a>	<a href="#">Meis2.iSep08</a>	<a href="#">311311</a>	896	391	2	46	meis1, myeloid ecotropic viral integration site 1 homolog 2 (Meis2) alternative variant iSep08, mRNA.
<a href="#">Meis3</a>	<a href="#">Meis3.aSep08</a>	<a href="#">361514</a>	9556	1488	11	490	meis homeobox 3 (Meis3) alternative variant aSep08, mRNA.
<a href="#">Meis3</a>	<a href="#">Meis3.cSep08</a>	<a href="#">361514</a>	7330	1413	9	440	meis homeobox 3 (Meis3) alternative variant cSep08, mRNA.
<a href="#">Mel13</a>	<a href="#">Mel13.bSep08</a>	<a href="#">287661</a>	2977	723	3	137	melanoma nuclear protein 13 (15.6 kD) (Mel13) alternative variant bSep08, mRNA.
<a href="#">Mel13</a>	<a href="#">Mel13.cSep08</a>	<a href="#">287661</a>	2990	1005	2	87	melanoma nuclear protein 13 (9.8 kD) (Mel13) alternative variant cSep08, mRNA.
<a href="#">Melk</a>	<a href="#">Melk.bSep08</a>	<a href="#">362510</a>	34257	1014	1	337	maternal embryonic leucine zipper kinase (Melk) alternative variant bSep08, mRNA.
<a href="#">Memo1</a>	<a href="#">Memo1.bSep08</a>	<a href="#">298787</a>	88730	1811	8	271	mediator of cell motility 1 (Memo1) alternative variant bSep08, mRNA.
<a href="#">Memo1</a>	<a href="#">Memo1.cSep08</a>	<a href="#">298787</a>	11641	1180	3	79	mediator of cell motility 1 (8.8 kD) (Memo1) alternative variant cSep08, mRNA.
<a href="#">Memo1</a>	<a href="#">Memo1.dSep08</a>	<a href="#">298787</a>	3535	568	2	51	mediator of cell motility 1 (Memo1) alternative variant dSep08, mRNA.
<a href="#">Men1</a>	<a href="#">Men1.aSep08</a>	<a href="#">29417</a>	4788	1560	10	519	multiple endocrine neoplasia 1 (Men1) alternative variant aSep08, mRNA.
<a href="#">Men1</a>	<a href="#">Men1.bSep08</a>	<a href="#">29417</a>	2108	1564	3	270	multiple endocrine neoplasia 1 (Men1) alternative variant bSep08, mRNA.
<a href="#">Men1</a>	<a href="#">Men1.dSep08</a>	<a href="#">29417</a>	680	383	2	127	multiple endocrine neoplasia 1 (Men1) alternative variant dSep08, mRNA.
<a href="#">Mepce</a>	<a href="#">Mepce.aSep08</a>	<a href="#">304361</a>	3694	2078		515	methylphosphate capping enzyme (Mepce) mRNA.
<a href="#">merby</a>	<a href="#">merby.aSep08</a>		12687	359		119	cyclin-dependent kinase-like 5 (merby) mRNA.
<a href="#">merchy</a>	<a href="#">merchy.aSep08</a>		4369	619		181	tankyrase protein (merchy) mRNA.
<a href="#">merdar</a>	<a href="#">merdar.aSep08</a>		4382	526	3	104	putative mitochondrial protein (11.6 kD) (merdar) alternative variant aSep08, mRNA.

<a href="#">merdar</a>	<a href="#">merdar.bSep08</a>		1315	401	2	74	putative protein (merdar) alternative variant bSep08, mRNA.
<a href="#">merdar</a>	<a href="#">merdar.cSep08</a>		7856	787	4	42	putative protein (merdar) alternative variant cSep08, mRNA.
<a href="#">merfer</a>	<a href="#">merfer.aSep08</a>		6358	493	2	31	putative protein (merfer) alternative variant aSep08, mRNA.
<a href="#">merfer</a>	<a href="#">merfer.bSep08</a>		2147	262	1	33	putative protein (merfer) alternative variant bSep08, mRNA.
<a href="#">merflo</a>	<a href="#">merflo.aSep08</a>		783	293		63	putative protein (7.0 kD) (merflo) mRNA.
<a href="#">merflu</a>	<a href="#">merflu.aSep08</a>		3536	1157	1	38	putative protein (4.6 kD) (merflu) alternative variant aSep08, mRNA.
<a href="#">merflu</a>	<a href="#">merflu.bSep08</a>		2408	341	1	16	putative protein (merflu) alternative variant bSep08, mRNA.
<a href="#">mergar</a>	<a href="#">mergar.aSep08</a>		11781	717		239	CRA b (mergar) mRNA.
<a href="#">merjey</a>	<a href="#">merjey.aSep08</a>		8505	555		35	putative protein (4.0 kD) (merjey) mRNA.
<a href="#">merkee</a>	<a href="#">merkee.aSep08</a>		26503	973		324	glycoside hydrolase, family 47 (merkee) mRNA.
<a href="#">merloy</a>	<a href="#">merloy.aSep08</a>		1946	941		77	putative protein (merloy) mRNA.
<a href="#">mermee</a>	<a href="#">mermee.aSep08</a>		1243	348		24	putative protein (2.8 kD) (mermee) mRNA.
<a href="#">mermer</a>	<a href="#">mermer.aSep08</a>		894	778		51	putative protein (mermer) mRNA.
<a href="#">merno</a>	<a href="#">merno.aSep08</a>		7732	289		90	putative protein (merno) mRNA.
<a href="#">merpor</a>	<a href="#">merpor.bSep08</a>		1202	357	3	55	putative protein (merpor) alternative variant bSep08, mRNA.
<a href="#">mersa</a>	<a href="#">mersa.aSep08</a>		5888	2303	2	107	putative mitochondrial protein (11.9 kD) (mersa) mRNA.
<a href="#">mershee</a>	<a href="#">mershee.aSep08</a>		97385	818		39	putative protein (4.6 kD) (mershee) mRNA.
<a href="#">Mertk</a>	<a href="#">Mertk.bSep08</a>	<a href="#">65037</a>	25769	681	5	222	c-mer proto-oncogene tyrosine kinase (Mertk) alternative variant bSep08, mRNA.
<a href="#">mertu</a>	<a href="#">mertu.aSep08</a>		4594	775		85	putative nuclear protein (9.0 kD) (mertu) mRNA.
<a href="#">mervar</a>	<a href="#">mervar.aSep08</a>		2610	701		12	putative protein (1.4 kD) (mervar) mRNA.
<a href="#">merwey</a>	<a href="#">merwey.aSep08</a>		1747	664		116	bromodomain PHD finger containing 1 CRA d (merwey) mRNA.
<a href="#">Mesdc2</a>	<a href="#">Mesdc2.bSep08</a>	<a href="#">308796</a>	12572	533	3	153	mesoderm development candidate 2 (Mesdc2) alternative variant bSep08, mRNA.
<a href="#">Metap1</a>	<a href="#">Metap1.bSep08</a>	<a href="#">295500</a>	18802	681	2	226	methionyl aminopeptidase 1 (Metap1) alternative variant bSep08, mRNA.
<a href="#">Metap1</a>	<a href="#">Metap1.cSep08</a>	<a href="#">295500</a>	18979	677	2	225	methionyl aminopeptidase 1 (Metap1) alternative variant cSep08, mRNA.
<a href="#">Metap2</a>	<a href="#">Metap2.bSep08</a>	<a href="#">64370</a>	23418	1046	5	319	methionine aminopeptidase 2 (Metap2) alternative variant bSep08, mRNA.
<a href="#">Metap2</a>	<a href="#">Metap2.cSep08</a>	<a href="#">64370</a>	19368	897	4	263	methionine aminopeptidase 2 (Metap2) alternative variant cSep08, mRNA.
<a href="#">Metap2</a>	<a href="#">Metap2.dSep08</a>	<a href="#">64370</a>	18249	847	3	252	methionine aminopeptidase 2 (Metap2) alternative variant dSep08, mRNA.
<a href="#">Metap2</a>	<a href="#">Metap2.eSep08</a>	<a href="#">64370</a>	27955	2162	7	237	methionine aminopeptidase 2 (26.1 kD) (Metap2) alternative variant eSep08, mRNA.
<a href="#">Metap2</a>	<a href="#">Metap2.fSep08</a>	<a href="#">64370</a>	18243	748	2	213	methionine aminopeptidase 2 (Metap2) alternative variant fSep08, mRNA.

<a href="#">Metap2</a>	<a href="#">Metap2.gSep08</a>	<a href="#">64370</a>	18219	729	2	123	methionine aminopeptidase 2 (13.4 kD) (Metap2) alternative variant gSep08, mRNA.
<a href="#">Metap1</a>	<a href="#">Metap1.bSep08</a>	<a href="#">311748</a>	71373	698		232	methionine aminopeptidase-like 1 (Metap1) alternative variant bSep08, mRNA.
<a href="#">Methyltransf_16.0</a>	<a href="#">Methyltransf_16.0.aSep08</a>		687	409		131	CRA a (Methyltransf_16.0) mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.cSep08</a>	<a href="#">287150</a>	809	595	2	164	meteorin (MetrnandRGD1306126) alternative variant cSep08, mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.cSep08</a>	<a href="#">287151</a>	809	595	2	164	meteorin (MetrnandRGD1306126) alternative variant cSep08, mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.dSep08</a>	<a href="#">287150</a>	1629	716	4	132	putative cytoplasmic protein of ancient origin (14.2 kD) (MetrnandRGD1306126) alternative variant dSep08, mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.dSep08</a>	<a href="#">287151</a>	1629	716	4	132	putative cytoplasmic protein of ancient origin (14.2 kD) (MetrnandRGD1306126) alternative variant dSep08, mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.eSep08</a>	<a href="#">287150</a>	5795	478	4	112	meteorin (MetrnandRGD1306126) alternative variant eSep08, mRNA.
<a href="#">MetrnandRGD1306126</a>	<a href="#">MetrnandRGD1306126.eSep08</a>	<a href="#">287151</a>	5795	478	4	112	meteorin (MetrnandRGD1306126) alternative variant eSep08, mRNA.
<a href="#">Metrnl</a>	<a href="#">Metrnl.bSep08</a>	<a href="#">316842</a>	13472	448	3	146	meteorin, glial cell differentiation regulator-like (Metrnl) alternative variant bSep08, mRNA.
<a href="#">Metrnl</a>	<a href="#">Metrnl.cSep08</a>	<a href="#">316842</a>	5330	432	2	143	meteorin, glial cell differentiation regulator-like (Metrnl) alternative variant cSep08, mRNA.
<a href="#">Mett10d</a>	<a href="#">Mett10d.aSep08</a>	<a href="#">360568</a>	44749	1120	8	339	putative protein of ancient origin (Mett10d) alternative variant aSep08, mRNA.
<a href="#">Mett10d</a>	<a href="#">Mett10d.bSep08</a>	<a href="#">360568</a>	3713	689	1	127	putative protein of ancient origin (Mett10d) alternative variant bSep08, mRNA.
<a href="#">Mett11d1</a>	<a href="#">Mett11d1.aSep08</a>	<a href="#">305845</a>	6618	1521	13	466	methyltransferase type 11 (Mett11d1) alternative variant aSep08, mRNA.
<a href="#">Mett11d1</a>	<a href="#">Mett11d1.bSep08</a>	<a href="#">305845</a>	4563	826	7	77	putative cytoplasmic protein of ancient origin (8.3 kD) (Mett11d1) alternative variant bSep08, mRNA.
<a href="#">Mett11d1</a>	<a href="#">Mett11d1.cSep08</a>	<a href="#">305845</a>	3109	1668	4	80	putative protein of mammalian origin (9.1 kD) (Mett11d1) alternative variant cSep08, mRNA.
<a href="#">Mett11d1</a>	<a href="#">Mett11d1.dSep08</a>	<a href="#">305845</a>	2580	1140	3	80	putative protein of mammalian origin (9.1 kD) (Mett11d1) alternative variant dSep08, mRNA.
<a href="#">Mettl2</a>	<a href="#">Mettl2.bSep08</a>	<a href="#">363687</a>	7679	752	4	117	methyltransferase like 2 (Mettl2) alternative variant bSep08, mRNA.
<a href="#">Mettl3</a>	<a href="#">Mettl3.bSep08</a>	<a href="#">361035</a>	1482	728	3	59	methyltransferase-like 3 (Mettl3) alternative variant bSep08, mRNA.
<a href="#">Mettl4</a>	<a href="#">Mettl4.aSep08</a>	<a href="#">316731</a>	22447	662	4	220	methyltransferase like 4 (Mettl4) alternative variant aSep08, mRNA.
<a href="#">Mettl6</a>	<a href="#">Mettl6.bSep08</a>	<a href="#">290564</a>	2836	884	3	120	methyltransferase like 6 (Mettl6) alternative variant bSep08, mRNA.
<a href="#">meyby</a>	<a href="#">meyby.aSep08</a>		2848	466		31	putative protein (3.6 kD) (meyby) mRNA.
<a href="#">meychy</a>	<a href="#">meychy.aSep08</a>		1407	742		247	tankyrase 1 binding protein like (meychy) mRNA.
<a href="#">meydar</a>	<a href="#">meydar.aSep08</a>		1680	263		19	putative protein (2.2 kD) (meydar) mRNA.

<a href="#">meyfer</a>	<a href="#">meyfer.aSep08</a>		139217	691		230	transcription initiation factor tfiid (meyfer) mRNA.
<a href="#">meyflo</a>	<a href="#">meyflo.aSep08</a>		896	770		38	putative protein (4.1 kD) (meyflo) mRNA.
<a href="#">meyflu</a>	<a href="#">meyflu.aSep08</a>		104640	495		27	putative protein (3.1 kD) (meyflu) mRNA.
<a href="#">meygar</a>	<a href="#">meygar.aSep08</a>		3857	399		108	heavy axonemal (meygar) mRNA.
<a href="#">meyjey</a>	<a href="#">meyjey.aSep08</a>		805	586		103	putative protein (meyjey) mRNA.
<a href="#">meykee</a>	<a href="#">meykee.aSep08</a>		6650	614		189	3 ER degradation mannosidase-like (meykee) mRNA.
<a href="#">meyloy</a>	<a href="#">meyloy.aSep08</a>		3229	333		56	putative protein (meyloy) mRNA.
<a href="#">meymee</a>	<a href="#">meymee.aSep08</a>		28668	452		40	putative protein (meymee) mRNA.
<a href="#">meymer</a>	<a href="#">meymer.aSep08</a>		14216	449		149	tweety 2 (meymer) mRNA.
<a href="#">meynoy</a>	<a href="#">meynoy.aSep08</a>		5673	644	3	169	phosphatidylinositol-3-phosphate phosphatidylinositol type III like (meynoy) alternative variant aSep08, mRNA.
<a href="#">meynoy</a>	<a href="#">meynoy.bSep08</a>		5516	542	3	135	type III (meynoy) alternative variant bSep08, mRNA.
<a href="#">meypor</a>	<a href="#">meypor.aSep08</a>		937	455		85	putative mitochondrial protein (9.2 kD) (meypor) mRNA.
<a href="#">meysa</a>	<a href="#">meysa.aSep08</a>		1191	515		171	rhophilin Rho GTPase binding protein 1 CRA a like (meysa) mRNA.
<a href="#">meyshee</a>	<a href="#">meyshee.aSep08</a>		166285	408		50	putative protein (meyshee) mRNA.
<a href="#">meytu</a>	<a href="#">meytu.aSep08</a>		1103	743		143	putative protein (meytu) mRNA.
<a href="#">meyvar</a>	<a href="#">meyvar.aSep08</a>		5015	697		140	putative protein (meyvar) mRNA.
<a href="#">meywey</a>	<a href="#">meywey.aSep08</a>		2395	1018		111	bromodomain PHD finger containing 1 CRA e (meywey) mRNA.
<a href="#">Mfap1a</a>	<a href="#">Mfap1a.aSep08</a>	<a href="#">499878</a>	18346	3636		439	microfibrillar-associated protein 1A (52.0 kD) (Mfap1a) mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.aSep08</a>	<a href="#">287382</a>	3133	1750	4	281	microfibrillar-associated protein 4 (31.4 kD) (Mfap4) alternative variant aSep08, complete mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.cSep08</a>	<a href="#">287382</a>	1756	742	3	241	microfibrillar-associated protein 4 (26.5 kD) (Mfap4) alternative variant cSep08, mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.dSep08</a>	<a href="#">287382</a>	1860	774	4	217	microfibrillar-associated protein 4 (24.1 kD) (Mfap4) alternative variant dSep08, mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.eSep08</a>	<a href="#">287382</a>	2771	1151	5	210	microfibrillar-associated protein 4 (24.0 kD) (Mfap4) alternative variant eSep08, mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.fSep08</a>	<a href="#">287382</a>	2350	1137	4	135	microfibrillar-associated protein 4 (15.3 kD) (Mfap4) alternative variant fSep08, mRNA.
<a href="#">Mfap4</a>	<a href="#">Mfap4.gSep08</a>	<a href="#">287382</a>	2241	778	5	125	microfibrillar-associated protein 4 (13.9 kD) (Mfap4) alternative variant gSep08, mRNA.
<a href="#">Mfap5</a>	<a href="#">Mfap5.bSep08</a>	<a href="#">362429</a>	3229	669	1	49	microfibrillar associated protein 5 (5.2 kD) (Mfap5) alternative variant bSep08, mRNA.
<a href="#">Mfge8</a>	<a href="#">Mfge8.bSep08</a>	<a href="#">25277</a>	12004	1785	2	473	milk fat globule-EGF factor 8 protein (Mfge8) alternative variant bSep08, mRNA.
<a href="#">Mfhas1</a>	<a href="#">Mfhas1.bSep08</a>	<a href="#">306508</a>	84600	3294	3	990	malignant fibrous histiocytoma amplified sequence 1 (Mfhas1) alternative variant bSep08, mRNA.
<a href="#">Mfhas1</a>	<a href="#">Mfhas1.cSep08</a>	<a href="#">306508</a>	13447	470	3	57	malignant fibrous histiocytoma amplified sequence 1 (Mfhas1) alternative variant cSep08, mRNA.
<a href="#">Mfhas1</a>	<a href="#">Mfhas1.dSep08</a>	<a href="#">306508</a>	8594	457	2	37	malignant fibrous histiocytoma amplified sequence 1 (Mfhas1) alternative variant dSep08, mRNA.

<a href="#">Mfi2</a>	<a href="#">Mfi2.cSep08</a>	<a href="#">288038</a>	1535	736	2	84	antigen p97 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5 (Mfi2) alternative variant cSep08, mRNA.
<a href="#">Mfn1</a>	<a href="#">Mfn1.aSep08</a>	<a href="#">192647</a>	16000	3271	2	416	mitofusin 1 (Mfn1) alternative variant aSep08, mRNA.
<a href="#">Mfn1</a>	<a href="#">Mfn1.bSep08</a>	<a href="#">192647</a>	2536	1677	1	138	mitofusin 1 (Mfn1) alternative variant bSep08, mRNA.
<a href="#">Mfn2</a>	<a href="#">Mfn2.bSep08</a>	<a href="#">64476</a>	4894	2161	3	148	mitofusin 2 (Mfn2) alternative variant bSep08, mRNA.
<a href="#">Mfn2</a>	<a href="#">Mfn2.dSep08</a>	<a href="#">64476</a>	9621	514	4	72	mitofusin 2 (Mfn2) alternative variant dSep08, mRNA.
<a href="#">Mfsd1</a>	<a href="#">Mfsd1.aSep08</a>	<a href="#">361957</a>	19280	1425	16	464	major facilitator superfamily MFS 1 (51.3 kD) (Mfsd1) alternative variant aSep08, mRNA.
<a href="#">Mfsd1</a>	<a href="#">Mfsd1.bSep08</a>	<a href="#">361957</a>	11144	2415	10	258	putative protein, with at least 6 transmembrane domains, of ancient origin (Mfsd1) alternative variant bSep08, mRNA.
<a href="#">Mfsd1</a>	<a href="#">Mfsd1.cSep08</a>	<a href="#">361957</a>	2742	514	3	43	putative protein of metazoan origin (Mfsd1) alternative variant cSep08, mRNA.
<a href="#">Mfsd2</a>	<a href="#">Mfsd2.bSep08</a>	<a href="#">298504</a>	10534	756	3	252	putative protein of ancient origin (Mfsd2) alternative variant bSep08, mRNA.
<a href="#">Mfsd2</a>	<a href="#">Mfsd2.cSep08</a>	<a href="#">298504</a>	5348	622	3	193	putative protein, with at least 2 transmembrane domains, of ancient origin (Mfsd2) alternative variant cSep08, mRNA.
<a href="#">Mfsd3</a>	<a href="#">Mfsd3.bSep08</a>	<a href="#">500899</a>	559	477	2	114	putative protein, with a transmembrane domain, of vertebrate origin (Mfsd3) alternative variant bSep08, mRNA.
<a href="#">Mfsd3</a>	<a href="#">Mfsd3.cSep08</a>	<a href="#">500899</a>	1583	668	4	112	putative protein of mammalian origin (Mfsd3) alternative variant cSep08, mRNA.
<a href="#">Mfsd3</a>	<a href="#">Mfsd3.dSep08</a>	<a href="#">500899</a>	911	815	2	34	putative protein (3.6 kD) (Mfsd3) alternative variant dSep08, mRNA.
<a href="#">Mfsd4</a>	<a href="#">Mfsd4.bSep08</a>	<a href="#">498228</a>	14179	1493	1	159	putative protein, with at least 3 transmembrane domains, of eukaryotic origin (Mfsd4) alternative variant bSep08, mRNA.
<a href="#">Mfsd7</a>	<a href="#">Mfsd7.bSep08</a>	<a href="#">305625</a>	4658	765	1	168	putative protein, with at least 2 transmembrane domains, of ancient origin (Mfsd7) alternative variant bSep08, mRNA.
<a href="#">Mfsd8</a>	<a href="#">Mfsd8.bSep08</a>	<a href="#">361939</a>	9279	377	4	125	putative protein, with a transmembrane domain, of ancient origin (Mfsd8) alternative variant bSep08, mRNA.
<a href="#">Mfsd10</a>	<a href="#">Mfsd10.aSep08</a>	<a href="#">305449</a>	967	423	3	140	putative protein of metazoan origin (Mfsd10) alternative variant aSep08, mRNA.
<a href="#">Mfsd10</a>	<a href="#">Mfsd10.bSep08</a>	<a href="#">305449</a>	1374	738	3	113	putative protein of metazoan origin (12.9 kD) (Mfsd10) alternative variant bSep08, mRNA.
<a href="#">Mga</a>	<a href="#">Mga.aSep08</a>	<a href="#">499874</a>	23794	1800	8	458	MAX gene associated (Mga) alternative variant aSep08, mRNA.
<a href="#">Mga</a>	<a href="#">Mga.bSep08</a>	<a href="#">499874</a>	8587	585	5	194	MAX gene associated (Mga) alternative variant bSep08, mRNA.
<a href="#">Mgat1</a>	<a href="#">Mgat1.bSep08</a>	<a href="#">81519</a>	10586	790	2	199	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (Mgat1) alternative variant bSep08, mRNA.
<a href="#">Mgat1</a>	<a href="#">Mgat1.cSep08</a>	<a href="#">81519</a>	13781	540	2	41	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (4.5 kD) (Mgat1) alternative variant cSep08, mRNA.

<a href="#">Mgat4a</a>	<a href="#">Mgat4a.bSep08</a>	<a href="#">367252</a>	41678	969	3	230	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme A (Mgat4a) alternative variant bSep08, mRNA.
<a href="#">Mgat4a</a>	<a href="#">Mgat4a.cSep08</a>	<a href="#">367252</a>	15507	728	3	166	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme A (Mgat4a) alternative variant cSep08, mRNA.
<a href="#">Mgat4b</a>	<a href="#">Mgat4b.bSep08</a>	<a href="#">303100</a>	2807	1329	10	366	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme B (Mgat4b) alternative variant bSep08, mRNA.
<a href="#">Mgat4b</a>	<a href="#">Mgat4b.cSep08</a>	<a href="#">303100</a>	1749	841	6	237	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme B (Mgat4b) alternative variant cSep08, mRNA.
<a href="#">Mgat4c</a>	<a href="#">Mgat4c.aSep08</a>	<a href="#">299756</a>	205967	385		59	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme C (putative) (Mgat4c) mRNA.
<a href="#">Mgat5</a>	<a href="#">Mgat5.bSep08</a>	<a href="#">65271</a>	16296	895	2	111	mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetylglucosaminyltransferase (Mgat5) alternative variant bSep08, mRNA.
<a href="#">Mgat5</a>	<a href="#">Mgat5.dSep08</a>	<a href="#">65271</a>	20068	590	3	54	mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetylglucosaminyltransferase (6.3 kD) (Mgat5) alternative variant dSep08, mRNA.
<a href="#">MGC72612</a>	<a href="#">MGC72612.bSep08</a>	<a href="#">494340</a>	4336	3220	2	386	similar to expressed sequence AI449175 (45.4 kD) (MGC72612) alternative variant bSep08, mRNA.
<a href="#">MGC72612</a>	<a href="#">MGC72612.cSep08</a>	<a href="#">494340</a>	9359	1719	3	35	similar to expressed sequence AI449175 (MGC72612) alternative variant cSep08, mRNA.
<a href="#">MGC72612</a>	<a href="#">MGC72612.dSep08</a>	<a href="#">494340</a>	12701	732	3	8	similar to expressed sequence AI449175 (0.9 kD) (MGC72612) alternative variant dSep08, mRNA.
<a href="#">MGC72614</a>	<a href="#">MGC72614.bSep08</a>	<a href="#">310540</a>	3723	591	2	147	hypothetical LOC310540 (MGC72614) alternative variant bSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.aSep08</a>	<a href="#">361619</a>	2854	802	3	148	beta-globin (MGC72973) alternative variant aSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.aSep08</a>	<a href="#">689064</a>	2854	802	3	148	beta-globin (MGC72973) alternative variant aSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.bSep08</a>	<a href="#">361619</a>	8086	606	3	147	beta-globin (16.0 kD) (MGC72973) alternative variant bSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.bSep08</a>	<a href="#">689064</a>	8086	606	3	147	beta-globin (16.0 kD) (MGC72973) alternative variant bSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.cSep08</a>	<a href="#">361619</a>	13266	621	3	147	beta-globin (16.0 kD) (MGC72973) alternative variant cSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.cSep08</a>	<a href="#">689064</a>	13266	621	3	147	beta-globin (16.0 kD) (MGC72973) alternative variant cSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.gSep08</a>	<a href="#">361619</a>	894	779	2	133	beta-globin (14.6 kD) (MGC72973) alternative variant gSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.gSep08</a>	<a href="#">689064</a>	894	779	2	133	beta-globin (14.6 kD) (MGC72973) alternative variant gSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.hSep08</a>	<a href="#">361619</a>	8090	689	4	110	beta-globin (12.0 kD) (MGC72973) alternative variant hSep08, mRNA.

<a href="#">MGC72973</a>	<a href="#">MGC72973.hSep08</a>	<a href="#">689064</a>	8090	689	4	110	beta-globin (12.0 kD) (MGC72973) alternative variant hSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.jSep08</a>	<a href="#">361619</a>	7441	384	2	35	beta-globin (3.7 kD) (MGC72973) alternative variant jSep08, mRNA.
<a href="#">MGC72973</a>	<a href="#">MGC72973.jSep08</a>	<a href="#">689064</a>	7441	384	2	35	beta-globin (3.7 kD) (MGC72973) alternative variant jSep08, mRNA.
<a href="#">MGC72974</a>	<a href="#">MGC72974.bSep08</a>	<a href="#">316976</a>	9690	1443	4	255	hypothetical LOC316976 (28.6 kD) (MGC72974) alternative variant bSep08, mRNA.
<a href="#">MGC72974</a>	<a href="#">MGC72974.dSep08</a>	<a href="#">316976</a>	4192	1799	3	53	hypothetical LOC316976 (MGC72974) alternative variant dSep08, mRNA.
<a href="#">MGC72974</a>	<a href="#">MGC72974.eSep08</a>	<a href="#">316976</a>	3611	1349	3	104	hypothetical LOC316976 (12.2 kD) (MGC72974) alternative variant eSep08, complete mRNA.
<a href="#">MGC93975</a>	<a href="#">MGC93975.bSep08</a>	<a href="#">292878</a>	5733	1126	8	264	similar to 2310044H10Rik protein (27.6 kD) (MGC93975) alternative variant bSep08, mRNA.
<a href="#">MGC93975</a>	<a href="#">MGC93975.cSep08</a>	<a href="#">292878</a>	5644	1163	8	236	similar to 2310044H10Rik protein (25.0 kD) (MGC93975) alternative variant cSep08, mRNA.
<a href="#">MGC93975</a>	<a href="#">MGC93975.dSep08</a>	<a href="#">292878</a>	5327	900	7	233	similar to 2310044H10Rik protein (MGC93975) alternative variant dSep08, mRNA.
<a href="#">MGC93975</a>	<a href="#">MGC93975.eSep08</a>	<a href="#">292878</a>	2790	1294	5	154	similar to 2310044H10Rik protein (16.8 kD) (MGC93975) alternative variant eSep08, mRNA.
<a href="#">MGC94190</a>	<a href="#">MGC94190.bSep08</a>	<a href="#">288616</a>	15693	1001	5	245	similar to 0610007L01Rik protein (MGC94190) alternative variant bSep08, mRNA.
<a href="#">MGC94190</a>	<a href="#">MGC94190.cSep08</a>	<a href="#">288616</a>	18141	739	4	197	similar to 0610007L01Rik protein (MGC94190) alternative variant cSep08, mRNA.
<a href="#">MGC94192</a>	<a href="#">MGC94192.bSep08</a>	<a href="#">360550</a>	3705	1177	3	332	similar to PHD zinc finger containing protein JUNE1 (MGC94192) alternative variant bSep08, mRNA.
<a href="#">MGC94192</a>	<a href="#">MGC94192.cSep08</a>	<a href="#">360550</a>	2635	681	3	185	similar to PHD zinc finger containing protein JUNE1 (MGC94192) alternative variant cSep08, mRNA.
<a href="#">MGC94192</a>	<a href="#">MGC94192.dSep08</a>	<a href="#">360550</a>	1268	717	2	163	similar to PHD zinc finger containing protein JUNE1 (MGC94192) alternative variant dSep08, mRNA.
<a href="#">MGC94199</a>	<a href="#">MGC94199.bSep08</a>	<a href="#">362483</a>	13440	865	3	104	similar to RIKEN cDNA 2610301B20; EST A1428449 (MGC94199) alternative variant bSep08, mRNA.
<a href="#">MGC94199</a>	<a href="#">MGC94199.cSep08</a>	<a href="#">362483</a>	13424	986	3	101	similar to RIKEN cDNA 2610301B20; EST A1428449 (MGC94199) alternative variant cSep08, mRNA.
<a href="#">MGC94207</a>	<a href="#">MGC94207.bSep08</a>	<a href="#">362946</a>	550	379	1	119	similar to RIKEN cDNA C030006K11 (MGC94207) alternative variant bSep08, mRNA.
<a href="#">MGC94282</a>	<a href="#">MGC94282.aSep08</a>	<a href="#">297627</a>	5127	1245	3	235	similar to 5930416119Rik protein (27.1 kD) (MGC94282) alternative variant aSep08, mRNA.
<a href="#">MGC94542</a>	<a href="#">MGC94542.bSep08</a>	<a href="#">290631</a>	5977	1765	8	195	similar to RIKEN cDNA 5430437P03 (21.2 kD) (MGC94542) alternative variant bSep08, mRNA.
<a href="#">MGC94542</a>	<a href="#">MGC94542.cSep08</a>	<a href="#">290631</a>	1155	765	2	122	similar to RIKEN cDNA 5430437P03 (13.0 kD) (MGC94542) alternative variant cSep08, mRNA.
<a href="#">MGC94600</a>	<a href="#">MGC94600.aSep08</a>	<a href="#">301013</a>	14276	1156	6	245	scotin (26.4 kD) (MGC94600) alternative variant aSep08, mRNA.
<a href="#">MGC94600</a>	<a href="#">MGC94600.bSep08</a>	<a href="#">301013</a>	14278	1155	6	244	scotin (26.4 kD) (MGC94600) alternative variant bSep08, mRNA.
<a href="#">MGC94600</a>	<a href="#">MGC94600.cSep08</a>	<a href="#">301013</a>	13945	737	6	217	scotin (MGC94600) alternative variant cSep08, mRNA.



<a href="#">MGC94600</a>	<a href="#">MGC94600.dSep08</a>	<a href="#">301013</a>	5846	1025	5	159	scotin (MGC94600) alternative variant dSep08, mRNA.
<a href="#">MGC94600</a>	<a href="#">MGC94600.eSep08</a>	<a href="#">301013</a>	4896	410	3	136	scotin (MGC94600) alternative variant eSep08, mRNA.
<a href="#">MGC94600</a>	<a href="#">MGC94600.gSep08</a>	<a href="#">301013</a>	8879	305	3	101	scotin (MGC94600) alternative variant gSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.bSep08</a>	<a href="#">362712</a>	6204	677	5	225	CRA a like (MGC94915) alternative variant bSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.cSep08</a>	<a href="#">362712</a>	3576	911	4	171	putative protein, with a coiled coil domain, of eukaryotic origin (19.6 kD) (MGC94915) alternative variant cSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.dSep08</a>	<a href="#">362712</a>	5801	582	4	145	putative protein, with a coiled coil domain, of eukaryotic origin (MGC94915) alternative variant dSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.eSep08</a>	<a href="#">362712</a>	2297	405	2	134	CRA a like (MGC94915) alternative variant eSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.fSep08</a>	<a href="#">362712</a>	2127	703	4	82	upf0407 protein c2orf39 homolog like (MGC94915) alternative variant fSep08, mRNA.
<a href="#">MGC94915</a>	<a href="#">MGC94915.gSep08</a>	<a href="#">362712</a>	2670	533	3	74	CRA a like (8.7 kD) (MGC94915) alternative variant gSep08, mRNA.
<a href="#">MGC94941</a>	<a href="#">MGC94941.aSep08</a>	<a href="#">296988</a>	8007	2667	6	376	similar to Mkrn1 protein (MGC94941) alternative variant aSep08, mRNA.
<a href="#">MGC94941</a>	<a href="#">MGC94941.cSep08</a>	<a href="#">296988</a>	13619	736	3	194	similar to Mkrn1 protein (MGC94941) alternative variant cSep08, mRNA.
<a href="#">MGC94941</a>	<a href="#">MGC94941.dSep08</a>	<a href="#">296988</a>	1981	284	2	87	similar to Mkrn1 protein (MGC94941) alternative variant dSep08, mRNA.
<a href="#">MGC95152</a>	<a href="#">MGC95152.bSep08</a>	<a href="#">297109</a>	8550	638	1	120	similar to B230212L03Rik protein (MGC95152) alternative variant bSep08, mRNA.
<a href="#">MGC95208</a>	<a href="#">MGC95208.bSep08</a>	<a href="#">304176</a>	4262	696		195	similar to 4930453N24Rik protein (MGC95208) alternative variant bSep08, mRNA.
<a href="#">MGC95210</a>	<a href="#">MGC95210.bSep08</a>	<a href="#">287798</a>	10030	2019	4	530	DNA segment Chr 11 Wayne State University 47 expressed (58.2 kD) (MGC95210) alternative variant bSep08, mRNA.
<a href="#">MGC95210</a>	<a href="#">MGC95210.cSep08</a>	<a href="#">287798</a>	1259	659	2	219	DNA segment Chr 11 Wayne State University 47 expressed (MGC95210) alternative variant cSep08, mRNA.
<a href="#">MGC95210</a>	<a href="#">MGC95210.dSep08</a>	<a href="#">287798</a>	5883	506	2	131	CRA b (MGC95210) alternative variant dSep08, mRNA.
<a href="#">MGC95210</a>	<a href="#">MGC95210.eSep08</a>	<a href="#">287798</a>	1402	451	1	81	putative mitochondrial protein (8.8 kD) (MGC95210) alternative variant eSep08, mRNA.
<a href="#">MGC105560</a>	<a href="#">MGC105560.bSep08</a>	<a href="#">500941</a>	1920	1255	2	321	similar to Hypothetical protein BC014729 (38.1 kD) (MGC105560) alternative variant bSep08, mRNA.
<a href="#">MGC105649</a>	<a href="#">MGC105649.aSep08</a>	<a href="#">302884</a>	2825	554	2	121	hypothetical LOC302884 (MGC105649) alternative variant aSep08, mRNA.
<a href="#">MGC108747</a>	<a href="#">MGC108747.bSep08</a>	<a href="#">288001</a>	13963	675	7	224	similar to alpha-1 major acute phase protein prepeptide (MGC108747) alternative variant bSep08, mRNA.
<a href="#">MGC108747</a>	<a href="#">MGC108747.cSep08</a>	<a href="#">288001</a>	11216	255	4	80	similar to alpha-1 major acute phase protein prepeptide (MGC108747) alternative variant cSep08, mRNA.
<a href="#">MGC108747</a>	<a href="#">MGC108747.eSep08</a>	<a href="#">288001</a>	1602	413	2	24	similar to alpha-1 major acute phase protein prepeptide (MGC108747) alternative variant eSep08, mRNA.
<a href="#">MGC108785</a>	<a href="#">MGC108785.aSep08</a>	<a href="#">304277</a>	12467	718	3	239	similar to CPSF4 protein (MGC108785) alternative variant aSep08, mRNA.
<a href="#">MGC108785</a>	<a href="#">MGC108785.bSep08</a>	<a href="#">304277</a>	15671	976	4	238	similar to CPSF4 protein (MGC108785) alternative variant bSep08, mRNA.

<a href="#">MGC108785</a>	<a href="#">MGC108785.cSep08</a>	<a href="#">304277</a>	9485	748	2	174	similar to CPSF4 protein (MGC108785) alternative variant cSep08, mRNA.
<a href="#">MGC108896</a>	<a href="#">MGC108896.bSep08</a>	<a href="#">499689</a>	4897	1349	6	114	similar to glutathione transferase GSTM7-7 (13.4 kD) (MGC108896) alternative variant bSep08, mRNA.
<a href="#">MGC108896</a>	<a href="#">MGC108896.cSep08</a>	<a href="#">499689</a>	2684	434	5	78	similar to glutathione transferase GSTM7-7 (MGC108896) alternative variant cSep08, mRNA.
<a href="#">MGC112682</a>	<a href="#">MGC112682.bSep08</a>	<a href="#">497900</a>	7994	910	6	231	similar to RIKEN cDNA 0610039K22 (MGC112682) alternative variant bSep08, mRNA.
<a href="#">MGC112682</a>	<a href="#">MGC112682.cSep08</a>	<a href="#">497900</a>	7780	692	6	177	similar to RIKEN cDNA 0610039K22 (MGC112682) alternative variant cSep08, mRNA.
<a href="#">MGC112682</a>	<a href="#">MGC112682.dSep08</a>	<a href="#">497900</a>	1315	1236	2	66	similar to RIKEN cDNA 0610039K22 (7.5 kD) (MGC112682) alternative variant dSep08, mRNA.
<a href="#">MGC112727</a>	<a href="#">MGC112727.aSep08</a>	<a href="#">360762</a>	1697	859	2	225	similar to RNA polymerase 1-3 (MGC112727) alternative variant aSep08, mRNA.
<a href="#">MGC112727</a>	<a href="#">MGC112727.dSep08</a>	<a href="#">360762</a>	10529	836	2	137	similar to RNA polymerase 1-3 (MGC112727) alternative variant dSep08, mRNA.
<a href="#">MGC112727</a>	<a href="#">MGC112727.eSep08</a>	<a href="#">360762</a>	33220	863	4	71	similar to RNA polymerase 1-3 (MGC112727) alternative variant eSep08, mRNA.
<a href="#">MGC112727</a>	<a href="#">MGC112727.fSep08</a>	<a href="#">360762</a>	467	351	2	48	similar to RNA polymerase 1-3 (MGC112727) alternative variant fSep08, mRNA.
<a href="#">MGC112830</a>	<a href="#">MGC112830.bSep08</a>	<a href="#">361178</a>	38029	1730	3	338	similar to transcription factor (MGC112830) alternative variant bSep08, mRNA.
<a href="#">MGC112830</a>	<a href="#">MGC112830.cSep08</a>	<a href="#">361178</a>	32644	854	4	262	similar to transcription factor (MGC112830) alternative variant cSep08, mRNA.
<a href="#">MGC112830</a>	<a href="#">MGC112830.dSep08</a>	<a href="#">361178</a>	14944	722	1	203	similar to transcription factor (MGC112830) alternative variant dSep08, mRNA.
<a href="#">MGC112830</a>	<a href="#">MGC112830.eSep08</a>	<a href="#">361178</a>	30840	624	3	106	similar to transcription factor (11.2 kD) (MGC112830) alternative variant eSep08, mRNA.
<a href="#">MGC112899</a>	<a href="#">MGC112899.bSep08</a>	<a href="#">500087</a>	8251	491	2	97	similar to RIKEN cDNA 1110001J03 (MGC112899) alternative variant bSep08, mRNA.
<a href="#">MGC112899</a>	<a href="#">MGC112899.cSep08</a>	<a href="#">500087</a>	6814	383	2	89	similar to RIKEN cDNA 1110001J03 (MGC112899) alternative variant cSep08, mRNA.
<a href="#">MGC114440</a>	<a href="#">MGC114440.bSep08</a>	<a href="#">500566</a>	23319	582	1	32	similar to RIKEN cDNA 4930555I21 (3.7 kD) (MGC114440) alternative variant bSep08, mRNA.
<a href="#">MGC114529</a>	<a href="#">MGC114529.bSep08</a>	<a href="#">317590</a>	7618	1045	1	279	similar to melanoma antigen family A, 10 (MGC114529) alternative variant bSep08, mRNA.
<a href="#">MGC116121</a>	<a href="#">MGC116121.bSep08</a>	<a href="#">498830</a>	4825	267	3	88	similar to RIKEN cDNA 2700062C07 (MGC116121) alternative variant bSep08, mRNA.
<a href="#">MGC116121</a>	<a href="#">MGC116121.cSep08</a>	<a href="#">498830</a>	840	765	2	73	similar to RIKEN cDNA 2700062C07 (MGC116121) alternative variant cSep08, mRNA.
<a href="#">MGC116202</a>	<a href="#">MGC116202.bSep08</a>	<a href="#">688736</a>	3984	789	3	230	hypothetical protein LOC688736 (MGC116202) alternative variant bSep08, mRNA.
<a href="#">MGC116202</a>	<a href="#">MGC116202.cSep08</a>	<a href="#">688736</a>	4572	576	4	78	hypothetical protein LOC688736 (MGC116202) alternative variant cSep08, mRNA.
<a href="#">MGC124740</a>	<a href="#">MGC124740.bSep08</a>	<a href="#">688590</a>	31356	610	1	202	similar to Rap2-binding protein 9 (MGC124740) alternative variant bSep08, mRNA.

<a href="#">MGC124740</a>	<a href="#">MGC124740.cSep08</a>	<a href="#">688590</a>	31301	561	1	186	similar to Rap2-binding protein 9 (MGC124740) alternative variant cSep08, mRNA.
<a href="#">MGC124992</a>	<a href="#">MGC124992.bSep08</a>	<a href="#">499697</a>	6061	763	7	246	putative protein of eukaryotic origin (MGC124992) alternative variant bSep08, mRNA.
<a href="#">MGC124992</a>	<a href="#">MGC124992.cSep08</a>	<a href="#">499697</a>	2252	1571	2	236	putative cytoplasmic protein of eukaryotic origin (26.5 kD) (MGC124992) alternative variant cSep08, mRNA.
<a href="#">MGC124992</a>	<a href="#">MGC124992.dSep08</a>	<a href="#">499697</a>	1172	611	3	118	putative protein of vertebrate origin (MGC124992) alternative variant dSep08, mRNA.
<a href="#">MGC125002</a>	<a href="#">MGC125002.bSep08</a>	<a href="#">500501</a>	44147	1628	4	174	similar to RIKEN cDNA 5830433M19 (18.2 kD) (MGC125002) alternative variant bSep08, mRNA.
<a href="#">MGC125002</a>	<a href="#">MGC125002.cSep08</a>	<a href="#">500501</a>	102481	743	7	120	similar to RIKEN cDNA 5830433M19 (13.4 kD) (MGC125002) alternative variant cSep08, mRNA.
<a href="#">MGC125086</a>	<a href="#">MGC125086.bSep08</a>	<a href="#">498695</a>	6718	1041	4	84	similar to RIKEN cDNA 5133401N09 (9.2 kD) (MGC125086) alternative variant bSep08, complete mRNA.
<a href="#">MGC125201</a>	<a href="#">MGC125201.bSep08</a>	<a href="#">497878</a>	58432	773	5	160	HMP19 protein (17.8 kD) (MGC125201) alternative variant bSep08, mRNA.
<a href="#">Mgea5</a>	<a href="#">Mgea5.bSep08</a>	<a href="#">154968</a>	6159	619	4	205	meningioma expressed antigen 5 (hyaluronidase) (Mgea5) alternative variant bSep08, mRNA.
<a href="#">Mgea5</a>	<a href="#">Mgea5.dSep08</a>	<a href="#">154968</a>	13978	3343	6	182	meningioma expressed antigen 5 (hyaluronidase) (20.3 kD) (Mgea5) alternative variant dSep08, mRNA.
<a href="#">Mgea5</a>	<a href="#">Mgea5.eSep08</a>	<a href="#">154968</a>	4511	1356	3	138	meningioma expressed antigen 5 (hyaluronidase) (Mgea5) alternative variant eSep08, mRNA.
<a href="#">Mgea5</a>	<a href="#">Mgea5.fSep08</a>	<a href="#">154968</a>	4981	633	3	136	meningioma expressed antigen 5 (hyaluronidase) (Mgea5) alternative variant fSep08, mRNA.
<a href="#">Mgea5</a>	<a href="#">Mgea5.gSep08</a>	<a href="#">154968</a>	3467	814	2	96	meningioma expressed antigen 5 (hyaluronidase) (10.8 kD) (Mgea5) alternative variant gSep08, mRNA.
<a href="#">Mgl1</a>	<a href="#">Mgl1.bSep08</a>	<a href="#">64195</a>	2647	613	3	113	macrophage galactose N-acetyl-galactosamine specific lectin 1 (Mgl1) alternative variant bSep08, mRNA.
<a href="#">Mgl1</a>	<a href="#">Mgl1.cSep08</a>	<a href="#">64195</a>	2068	563	1	84	macrophage galactose N-acetyl-galactosamine specific lectin 1 (9.2 kD) (Mgl1) alternative variant cSep08, mRNA.
<a href="#">Mgll</a>	<a href="#">Mgll.bSep08</a>	<a href="#">29254</a>	82028	695	5	164	monoglyceride lipase (Mgll) alternative variant bSep08, mRNA.
<a href="#">Mgll</a>	<a href="#">Mgll.cSep08</a>	<a href="#">29254</a>	71331	498	4	123	monoglyceride lipase (13.8 kD) (Mgll) alternative variant cSep08, mRNA.
<a href="#">Mgll</a>	<a href="#">Mgll.dSep08</a>	<a href="#">29254</a>	41720	774	4	112	monoglyceride lipase (12.9 kD) (Mgll) alternative variant dSep08, mRNA.
<a href="#">Mgll</a>	<a href="#">Mgll.eSep08</a>	<a href="#">29254</a>	967	488	2	58	monoglyceride lipase (6.7 kD) (Mgll) alternative variant eSep08, mRNA.
<a href="#">Mgp</a>	<a href="#">Mgp.aSep08</a>	<a href="#">25333</a>	3362	969	3	146	matrix Gla protein (Mgp) alternative variant aSep08, mRNA.
<a href="#">Mgp</a>	<a href="#">Mgp.cSep08</a>	<a href="#">25333</a>	3440	2050	3	78	matrix Gla protein (9.5 kD) (Mgp) alternative variant cSep08, mRNA.
<a href="#">Mgrn1</a>	<a href="#">Mgrn1.bSep08</a>	<a href="#">302938</a>	30198	766	6	209	mahogunin, ring finger 1 (Mgrn1) alternative variant bSep08, mRNA.
<a href="#">Mgrn1</a>	<a href="#">Mgrn1.cSep08</a>	<a href="#">302938</a>	7411	447	5	149	mahogunin, ring finger 1 (Mgrn1) alternative variant cSep08, mRNA.

<a href="#">Mgrn1</a>	<a href="#">Mgrn1.dSep08</a>	<a href="#">302938</a>	6927	1821	4	94	mahogunin, ring finger 1 (Mgrn1) alternative variant dSep08, mRNA.
<a href="#">Mgst2</a>	<a href="#">Mgst2.bSep08</a>	<a href="#">295037</a>	12194	374	2	52	microsomal glutathione S-transferase 2 (Mgst2) alternative variant bSep08, mRNA.
<a href="#">Mgst3</a>	<a href="#">Mgst3.aSep08</a>	<a href="#">289197</a>	20773	628	6	152	microsomal glutathione S-transferase 3 (16.7 kD) (Mgst3) alternative variant aSep08, mRNA.
<a href="#">Mgst3</a>	<a href="#">Mgst3.bSep08</a>	<a href="#">289197</a>	20780	628	6	114	microsomal glutathione S-transferase 3 (12.9 kD) (Mgst3) alternative variant bSep08, mRNA.
<a href="#">MHC_I.0</a>	<a href="#">MHC_I.0.aSep08</a>		891	730	2	243	MHC class I antigen like (MHC_I.0) alternative variant aSep08, mRNA.
<a href="#">MHC_I.1</a>	<a href="#">MHC_I.1.aSep08</a>		1530	1341	2	171	MHC class I antigen like (19.5 kD) (MHC_I.1) alternative variant aSep08, mRNA.
<a href="#">MHC_I.1</a>	<a href="#">MHC_I.1.bSep08</a>		1323	765	2	117	MHC class I (13.4 kD) (MHC_I.1) alternative variant bSep08, mRNA.
<a href="#">MHC_I.1</a>	<a href="#">MHC_I.1.cSep08</a>		2861	1429	7	99	putative protein (10.9 kD) (MHC_I.1) alternative variant cSep08, mRNA.
<a href="#">MHC_I.1</a>	<a href="#">MHC_I.1.dSep08</a>		948	767	2	61	putative protein (6.8 kD) (MHC_I.1) alternative variant dSep08, mRNA.
<a href="#">MIB_HERC2.0</a>	<a href="#">MIB_HERC2.0.aSep08</a>		2513	1014	7	337	mindbomb homolog 2 CRA b (MIB_HERC2.0) alternative variant aSep08, mRNA.
<a href="#">MIB_HERC2.0</a>	<a href="#">MIB_HERC2.0.bSep08</a>		987	865	1	72	mindbomb homolog 2 CRA c (MIB_HERC2.0) alternative variant bSep08, mRNA.
<a href="#">Mical1</a>	<a href="#">Mical1.cSep08</a>	<a href="#">294520</a>	716	433	2	66	putative protein of vertebrate origin (8.1 kD) (Mical1) alternative variant cSep08, mRNA.
<a href="#">Mical2</a>	<a href="#">Mical2.aSep08</a>	<a href="#">365352</a>	91566	1400		399	putative protein of eukaryotic origin (Mical2) mRNA.
<a href="#">Mical3</a>	<a href="#">Mical3.aSep08</a>	<a href="#">362427</a>	46544	1656	6	331	putative protein of eukaryotic origin (37.8 kD) (Mical3) alternative variant aSep08, complete mRNA.
<a href="#">Mical3</a>	<a href="#">Mical3.bSep08</a>	<a href="#">362427</a>	42191	736	2	121	putative protein of eukaryotic origin (Mical3) alternative variant bSep08, mRNA.
<a href="#">Mical3</a>	<a href="#">Mical3.cSep08</a>	<a href="#">362427</a>	42176	725	2	112	putative protein of metazoan origin (Mical3) alternative variant cSep08, mRNA.
<a href="#">Micalcl</a>	<a href="#">Micalcl.bSep08</a>	<a href="#">293180</a>	32630	1779	1	554	MICAL C-terminal like (Micalcl) alternative variant bSep08, mRNA.
<a href="#">Micall2</a>	<a href="#">Micall2.aSep08</a>	<a href="#">288515</a>	6772	1328	9	395	MICAL-like 2 (Micall2) alternative variant aSep08, mRNA.
<a href="#">Micall2</a>	<a href="#">Micall2.bSep08</a>	<a href="#">288515</a>	2679	541	1	33	MICAL-like 2 (3.8 kD) (Micall2) alternative variant bSep08, mRNA.
<a href="#">Mid1</a>	<a href="#">Mid1.bSep08</a>	<a href="#">54252</a>	263870	398	2	72	midline 1 (Mid1) alternative variant bSep08, mRNA.
<a href="#">Mid1ip1</a>	<a href="#">Mid1ip1.bSep08</a>	<a href="#">404280</a>	835	350	2	34	MID1 interacting protein 1 (gastrulation specific G12 homolog (zebrafish)) (Mid1ip1) alternative variant bSep08, mRNA.
<a href="#">Mid2</a>	<a href="#">Mid2.aSep08</a>	<a href="#">363502</a>	13688	440		146	midline 2 (Mid2) mRNA.
<a href="#">Midn</a>	<a href="#">Midn.aSep08</a>	<a href="#">314623</a>	1655	706		108	midnolin (Midn) mRNA.
<a href="#">Mier1</a>	<a href="#">Mier1.bSep08</a>	<a href="#">313418</a>	25923	600	7	143	mesoderm induction early response 1 homolog (Xenopus laevis) (Mier1) alternative variant bSep08, mRNA.
<a href="#">Mier1</a>	<a href="#">Mier1.cSep08</a>	<a href="#">313418</a>	16763	338	6	112	mesoderm induction early response 1 homolog (Xenopus laevis) (Mier1) alternative variant cSep08, mRNA.

<a href="#">Mier1</a>	<a href="#">Mier1.dSep08</a>	<a href="#">313418</a>	26656	717	8	98	mesoderm induction early response 1 homolog ( <i>Xenopus laevis</i> ) ( <i>Mier1</i> ) alternative variant dSep08, mRNA.
<a href="#">Mier2</a>	<a href="#">Mier2.aSep08</a>	<a href="#">362841</a>	9820	1617	12	471	mesoderm induction early response 1 family member 2 CRA d ( <i>Mier2</i> ) alternative variant aSep08, mRNA.
<a href="#">Mier2</a>	<a href="#">Mier2.cSep08</a>	<a href="#">362841</a>	3727	972	4	192	mesoderm induction early response 2 ( <i>Mier2</i> ) alternative variant cSep08, mRNA.
<a href="#">Mier2</a>	<a href="#">Mier2.dSep08</a>	<a href="#">362841</a>	3415	712	6	188	mesoderm induction early response 1 family member 2 CRA c ( <i>Mier2</i> ) alternative variant dSep08, mRNA.
<a href="#">Mier3</a>	<a href="#">Mier3.aSep08</a>	<a href="#">310086</a>	9336	384		127	mesoderm induction early response 1, family member 3 ( <i>Mier3</i> ) mRNA.
<a href="#">Mif</a>	<a href="#">Mif.aSep08</a>	<a href="#">81683</a>	716	594	1	122	macrophage migration inhibitory factor (12.8 kD) ( <i>Mif</i> ) alternative variant aSep08, mRNA.
<a href="#">Mif4gd</a>	<a href="#">Mif4gd.bSep08</a>	<a href="#">360659</a>	4552	1339	5	151	putative cytoplasmic protein of metazoan origin (17.2 kD) ( <i>Mif4gd</i> ) alternative variant bSep08, mRNA.
<a href="#">Mif4gd</a>	<a href="#">Mif4gd.cSep08</a>	<a href="#">360659</a>	3256	908	3	116	putative protein ( <i>Mif4gd</i> ) alternative variant cSep08, mRNA.
<a href="#">Mif4gd</a>	<a href="#">Mif4gd.dSep08</a>	<a href="#">360659</a>	1567	1169	2	103	putative cytoplasmic protein (11.9 kD) ( <i>Mif4gd</i> ) alternative variant dSep08, mRNA.
<a href="#">Mif4gd</a>	<a href="#">Mif4gd.eSep08</a>	<a href="#">360659</a>	3505	398	4	68	putative protein of vertebrate origin ( <i>Mif4gd</i> ) alternative variant eSep08, mRNA.
<a href="#">Mill2</a>	<a href="#">Mill2.bSep08</a>	<a href="#">365212</a>	2151	791	1	154	mill2 gene for MHC class I-like located near the LRC, 2, exon 3, partial cds, strain:LEW.1lm1 ( <i>Mill2</i> ) alternative variant bSep08, mRNA.
<a href="#">Minpp1</a>	<a href="#">Minpp1.bSep08</a>	<a href="#">29688</a>	44977	1091	2	90	multiple inositol polyphosphate histidine phosphatase 1 ( <i>Minpp1</i> ) alternative variant bSep08, mRNA.
<a href="#">Miox</a>	<a href="#">Miox.bSep08</a>	<a href="#">252899</a>	821	550	4	131	myo-inositol oxygenase ( <i>Miox</i> ) alternative variant bSep08, mRNA.
<a href="#">Miox</a>	<a href="#">Miox.cSep08</a>	<a href="#">252899</a>	1124	639	4	114	myo-inositol oxygenase ( <i>Miox</i> ) alternative variant cSep08, mRNA.
<a href="#">Miox</a>	<a href="#">Miox.dSep08</a>	<a href="#">252899</a>	1388	569	5	90	myo-inositol oxygenase ( <i>Miox</i> ) alternative variant dSep08, mRNA.
<a href="#">Miox</a>	<a href="#">Miox.eSep08</a>	<a href="#">252899</a>	916	776	2	73	myo-inositol oxygenase (8.4 kD) ( <i>Miox</i> ) alternative variant eSep08, mRNA.
<a href="#">Mipep</a>	<a href="#">Mipep.bSep08</a>	<a href="#">81684</a>	77191	1141		330	mitochondrial intermediate peptidase ( <i>Mipep</i> ) alternative variant bSep08, mRNA.
<a href="#">Mipep</a>	<a href="#">Mipep.cSep08</a>	<a href="#">81684</a>	25428	419	1	34	mitochondrial intermediate peptidase ( <i>Mipep</i> ) alternative variant cSep08, mRNA.
<a href="#">Miro.1</a>	<a href="#">Miro.1.aSep08</a>		7304	844		265	leucine-rich repeat kinase 2 ( <i>Miro.1</i> ) mRNA.
<a href="#">Mis12</a>	<a href="#">Mis12.aSep08</a>	<a href="#">501706</a>	8967	3153	2	206	MIS12, MIND kinetochore complex component, homolog (yeast) (24.2 kD) ( <i>Mis12</i> ) alternative variant aSep08, mRNA.
<a href="#">Mis12</a>	<a href="#">Mis12.bSep08</a>	<a href="#">501706</a>	6731	501	1	42	MIS12, MIND kinetochore complex component, homolog (yeast) ( <i>Mis12</i> ) alternative variant bSep08, mRNA.
<a href="#">MIT.0</a>	<a href="#">MIT.0.aSep08</a>		8213	565		187	MIT containing protein ( <i>MIT.0</i> ) mRNA.
<a href="#">Mitd1</a>	<a href="#">Mitd1.bSep08</a>	<a href="#">363219</a>	7577	395	3	88	MIT containing protein (9.8 kD) ( <i>Mitd1</i> ) alternative variant bSep08, mRNA.

<a href="#">Mitf</a>	<a href="#">Mitf.aSep08</a>	<a href="#">25094</a>	75047	1010	4	336	microphthalmia-associated transcription factor (Mitf) alternative variant aSep08, mRNA.
<a href="#">Mitf</a>	<a href="#">Mitf.bSep08</a>	<a href="#">25094</a>	10201	923	1	147	microphthalmia-associated transcription factor (Mitf) alternative variant bSep08, mRNA.
<a href="#">Mito_carr.0</a>	<a href="#">Mito_carr.0.aSep08</a>		5743	264		84	mitochondrial carrier protein (Mito_carr.0) mRNA.
<a href="#">Mito_carr.1</a>	<a href="#">Mito_carr.1.bSep08</a>		1095	625	2	70	mitochondrial carrier triple repeat (Mito_carr.1) alternative variant bSep08, mRNA.
<a href="#">Mito_carr.3</a>	<a href="#">Mito_carr.3.aSep08</a>	<a href="#">311122</a>	7260	1177		242	solute carrier family 25 member 12 (Mito_carr.3) mRNA.
<a href="#">Mizf</a>	<a href="#">Mizf.aSep08</a>	<a href="#">300665</a>	2502	1449		322	MBD2-interacting zinc finger (Mizf) mRNA.
<a href="#">Mki67</a>	<a href="#">Mki67.aSep08</a>	<a href="#">291234</a>	6095	2812	3	845	antigen identified by monoclonal antibody Ki-67 (Mki67) alternative variant aSep08, mRNA.
<a href="#">Mki67ip</a>	<a href="#">Mki67ip.aSep08</a>	<a href="#">246042</a>	10056	1720	4	272	mki67 (FHA domain) interacting nucleolar phosphoprotein (31.5 kD) (Mki67ip) alternative variant aSep08, complete mRNA.
<a href="#">Mki67ip</a>	<a href="#">Mki67ip.cSep08</a>	<a href="#">246042</a>	5858	694	2	159	mki67 (FHA domain) interacting nucleolar phosphoprotein (Mki67ip) alternative variant cSep08, mRNA.
<a href="#">Mkl1</a>	<a href="#">Mkl1.aSep08</a>	<a href="#">315151</a>	4459	2220		401	megakaryoblastic leukemia (translocation) 1 (Mkl1) mRNA.
<a href="#">Mknk1</a>	<a href="#">Mknk1.bSep08</a>	<a href="#">500526</a>	16354	554	6	170	MAP kinase-interacting serine/threonine kinase 1 (Mknk1) alternative variant bSep08, mRNA.
<a href="#">Mknk1</a>	<a href="#">Mknk1.cSep08</a>	<a href="#">500526</a>	32845	748	9	168	MAP kinase-interacting serine/threonine kinase 1 (Mknk1) alternative variant cSep08, mRNA.
<a href="#">Mknk1</a>	<a href="#">Mknk1.eSep08</a>	<a href="#">500526</a>	1915	483	2	35	MAP kinase-interacting serine/threonine kinase 1 (Mknk1) alternative variant eSep08, mRNA.
<a href="#">Mknk2</a>	<a href="#">Mknk2.bSep08</a>	<a href="#">299618</a>	6881	737	8	176	MAP kinase-interacting serine/threonine kinase 2 (Mknk2) alternative variant bSep08, mRNA.
<a href="#">Mknk2</a>	<a href="#">Mknk2.cSep08</a>	<a href="#">299618</a>	1185	747	2	98	MAP kinase-interacting serine/threonine kinase 2 (Mknk2) alternative variant cSep08, mRNA.
<a href="#">Mkrn2</a>	<a href="#">Mkrn2.bSep08</a>	<a href="#">297525</a>	999	838	2	92	makorin, ring finger protein, 2 (10.5 kD) (Mkrn2) alternative variant bSep08, mRNA.
<a href="#">Mkx</a>	<a href="#">Mkx.aSep08</a>	<a href="#">291228</a>	1989	295		98	mohawk homeobox (Mkx) mRNA.
<a href="#">Mlc1</a>	<a href="#">Mlc1.bSep08</a>	<a href="#">315215</a>	6147	771	1	93	megalencephalic leukoencephalopathy with subcortical cysts 1 homolog (human) (10.4 kD) (Mlc1) alternative variant bSep08, mRNA.
<a href="#">Mlf1</a>	<a href="#">Mlf1.bSep08</a>	<a href="#">310483</a>	33022	1024		267	myeloid leukemia factor 1 (30.6 kD) (Mlf1) alternative variant bSep08, complete mRNA.
<a href="#">Mlf1ip</a>	<a href="#">Mlf1ip.aSep08</a>	<a href="#">306464</a>	22706	1799	8	542	myeloid leukemia factor 1 interacting protein (Mlf1ip) alternative variant aSep08, mRNA.
<a href="#">Mlf1ip</a>	<a href="#">Mlf1ip.cSep08</a>	<a href="#">306464</a>	15402	889	8	248	myeloid leukemia factor 1 interacting protein (Mlf1ip) alternative variant cSep08, mRNA.
<a href="#">Mlf2</a>	<a href="#">Mlf2.aSep08</a>	<a href="#">312709</a>	4662	1534	9	247	myeloid leukemia factor 2 (28.1 kD) (Mlf2) alternative variant aSep08, complete mRNA.
<a href="#">Mlf2</a>	<a href="#">Mlf2.cSep08</a>	<a href="#">312709</a>	3121	643	7	213	myeloid leukemia factor 2 (Mlf2) alternative variant cSep08, mRNA.
<a href="#">Mlf2</a>	<a href="#">Mlf2.dSep08</a>	<a href="#">312709</a>	2202	995	5	180	myeloid leukemia factor 2 (Mlf2) alternative variant dSep08, mRNA.
<a href="#">Mlh1</a>	<a href="#">Mlh1.bSep08</a>	<a href="#">81685</a>	12055	1100	7	240	mutL homolog 1 (E. coli) (27.7 kD) (Mlh1) alternative variant bSep08, mRNA.

<a href="#">Mlh1</a>	<a href="#">Mlh1.cSep08</a>	<a href="#">81685</a>	683	304	2	83	mutL homolog 1 (E. coli) (Mlh1) alternative variant cSep08, mRNA.
<a href="#">Mlh1</a>	<a href="#">Mlh1.dSep08</a>	<a href="#">81685</a>	5639	509	2	57	mutL homolog 1 (E. coli) (Mlh1) alternative variant dSep08, mRNA.
<a href="#">Mlh1</a>	<a href="#">Mlh1.eSep08</a>	<a href="#">81685</a>	7881	357	4	45	mutL homolog 1 (E. coli) (4.9 kD) (Mlh1) alternative variant eSep08, mRNA.
<a href="#">Mll1</a>	<a href="#">Mll1.cSep08</a>	<a href="#">315606</a>	1827	545	3	115	myeloid/lymphoid or mixed-lineage leukemia 1 (Mll1) alternative variant cSep08, mRNA.
<a href="#">Mll1</a>	<a href="#">Mll1.eSep08</a>	<a href="#">315606</a>	1159	463	2	79	myeloid/lymphoid or mixed-lineage leukemia 1 (Mll1) alternative variant eSep08, mRNA.
<a href="#">Mll5</a>	<a href="#">Mll5.aSep08</a>	<a href="#">311968</a>	14861	1517	8	481	myeloid/lymphoid or mixed-lineage leukemia 5 (trithorax homolog, Drosophila) (Mll5) alternative variant aSep08, mRNA.
<a href="#">Mll5</a>	<a href="#">Mll5.bSep08</a>	<a href="#">311968</a>	4093	698	3	232	myeloid/lymphoid or mixed-lineage leukemia 5 (trithorax homolog, Drosophila) (Mll5) alternative variant bSep08, mRNA.
<a href="#">Mll5</a>	<a href="#">Mll5.cSep08</a>	<a href="#">311968</a>	1933	1235	2	114	myeloid/lymphoid or mixed-lineage leukemia 5 (trithorax homolog, Drosophila) (Mll5) alternative variant cSep08, mRNA.
<a href="#">Mllt1</a>	<a href="#">Mllt1.bSep08</a>	<a href="#">301119</a>	37350	1094	5	316	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 1 (Mllt1) alternative variant bSep08, mRNA.
<a href="#">Mllt3</a>	<a href="#">Mllt3.aSep08</a>	<a href="#">114510</a>	19251	1940		195	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3 (Mllt3) mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.bSep08</a>	<a href="#">26955</a>	21880	3846	8	673	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant bSep08, mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.cSep08</a>	<a href="#">26955</a>	5296	476	3	157	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant cSep08, mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.dSep08</a>	<a href="#">26955</a>	5272	455	3	151	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant dSep08, mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.eSep08</a>	<a href="#">26955</a>	3869	417	3	138	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant eSep08, mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.fSep08</a>	<a href="#">26955</a>	5226	409	3	135	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant fSep08, mRNA.
<a href="#">Mllt4</a>	<a href="#">Mllt4.gSep08</a>	<a href="#">26955</a>	5214	400	3	132	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (Mllt4) alternative variant gSep08, mRNA.
<a href="#">Mllt6</a>	<a href="#">Mllt6.aSep08</a>	<a href="#">303504</a>	7723	2579	10	692	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6 (69.5 kD) (Mllt6) alternative variant aSep08, mRNA.

<a href="#">MlIt6</a>	<a href="#">MlIt6.bSep08</a>	<a href="#">303504</a>	2082	746	3	248	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6 (MlIt6) alternative variant bSep08, mRNA.
<a href="#">MlIt6</a>	<a href="#">MlIt6.cSep08</a>	<a href="#">303504</a>	4197	705	4	235	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6 (MlIt6) alternative variant cSep08, mRNA.
<a href="#">MlIt6</a>	<a href="#">MlIt6.dSep08</a>	<a href="#">303504</a>	6409	4242	3	154	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6 (MlIt6) alternative variant dSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.bSep08</a>	<a href="#">361285</a>	41162	3648	13	684	myeloid lymphoid mixed-lineage leukemia translocated 10 like (MlIt10) alternative variant bSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.cSep08</a>	<a href="#">361285</a>	30036	777	7	233	myeloid lymphoid mixed-lineage leukemia translocated 10 like (MlIt10) alternative variant cSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.dSep08</a>	<a href="#">361285</a>	2083	1584	3	200	myeloid lymphoid mixed-lineage leukemia translocated 10 like (MlIt10) alternative variant dSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.eSep08</a>	<a href="#">361285</a>	51835	580	6	177	myeloid lymphoid mixed-lineage leukemia like (MlIt10) alternative variant eSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.fSep08</a>	<a href="#">361285</a>	31736	526	6	155	myeloid lymphoid mixed-lineage leukemia like (MlIt10) alternative variant fSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.gSep08</a>	<a href="#">361285</a>	52703	392	4	130	myeloid lymphoid mixed-lineage leukemia like (MlIt10) alternative variant gSep08, mRNA.
<a href="#">MlIt10</a>	<a href="#">MlIt10.iSep08</a>	<a href="#">361285</a>	48647	543	6	69	myeloid lymphoid mixed-lineage leukemia like (MlIt10) alternative variant iSep08, mRNA.
<a href="#">MlIt11</a>	<a href="#">MlIt11.aSep08</a>	<a href="#">295264</a>	9655	728	2	108	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 11 (MlIt11) alternative variant aSep08, mRNA.
<a href="#">MlIt11</a>	<a href="#">MlIt11.cSep08</a>	<a href="#">295264</a>	9481	741	3	62	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 11 (MlIt11) alternative variant cSep08, mRNA.
<a href="#">Mlph</a>	<a href="#">Mlph.bSep08</a>	<a href="#">316620</a>	4800	726	1	41	melanophilin (Mlph) alternative variant bSep08, mRNA.
<a href="#">Mlstd2</a>	<a href="#">Mlstd2.bSep08</a>	<a href="#">293173</a>	23914	4312	9	341	fatty acyl CoA reductase 1 (Mlstd2) alternative variant bSep08, mRNA.
<a href="#">Mlx</a>	<a href="#">Mlx.bSep08</a>	<a href="#">360631</a>	4757	2823	7	264	MAX-like protein X (Mlx) alternative variant bSep08, mRNA.
<a href="#">Mlx</a>	<a href="#">Mlx.cSep08</a>	<a href="#">360631</a>	4403	1457	8	167	MAX-like protein X (19.0 kD) (Mlx) alternative variant cSep08, mRNA.
<a href="#">Mlx</a>	<a href="#">Mlx.dSep08</a>	<a href="#">360631</a>	2441	417	6	138	MAX-like protein X (Mlx) alternative variant dSep08, mRNA.
<a href="#">Mlxip</a>	<a href="#">Mlxip.aSep08</a>	<a href="#">304479</a>	1680	770		256	MLX interacting protein (Mlxip) mRNA.
<a href="#">Mlycd</a>	<a href="#">Mlycd.bSep08</a>	<a href="#">85239</a>	2956	816	1	225	malonyl-CoA decarboxylase (25.1 kD) (Mlycd) alternative variant bSep08, mRNA.
<a href="#">Mlze</a>	<a href="#">Mlze.bSep08</a>	<a href="#">299908</a>	1218	745	2	102	melanoma-derived leucine zipper, extra-nuclear factor (11.3 kD) (Mlze) alternative variant bSep08, mRNA.
<a href="#">Mmaa</a>	<a href="#">Mmaa.bSep08</a>	<a href="#">291939</a>	16865	733	4	158	methylmalonic aciduria cblA type (18.1 kD) (Mmaa) alternative variant bSep08, mRNA.
<a href="#">Mmaa</a>	<a href="#">Mmaa.cSep08</a>	<a href="#">291939</a>	1792	941	2	137	methylmalonic aciduria cblA type (15.9 kD) (Mmaa) alternative variant cSep08, mRNA.



<a href="#">Mmachc</a>	<a href="#">Mmachc.bSep08</a>	<a href="#">313520</a>	6371	1836	2	144	methylmalonic aciduria (cobalamin deficiency) cbIC type, with homocystinuria (Mmachc) alternative variant bSep08, mRNA.
<a href="#">Mmachc</a>	<a href="#">Mmachc.cSep08</a>	<a href="#">313520</a>	3661	751	1	134	methylmalonic aciduria (cobalamin deficiency) cbIC type, with homocystinuria (Mmachc) alternative variant cSep08, mRNA.
<a href="#">Mmd</a>	<a href="#">Mmd.bSep08</a>	<a href="#">303439</a>	9813	390	1	77	monocyte to macrophage differentiation-associated (Mmd) alternative variant bSep08, mRNA.
<a href="#">Mme</a>	<a href="#">Mme.bSep08</a>	<a href="#">24590</a>	29294	551	1	174	membrane metallo endopeptidase (Mme) alternative variant bSep08, mRNA.
<a href="#">Mme</a>	<a href="#">Mme.cSep08</a>	<a href="#">24590</a>	29158	576	1	150	membrane metallo endopeptidase (Mme) alternative variant cSep08, mRNA.
<a href="#">Mmel1</a>	<a href="#">Mmel1.bSep08</a>	<a href="#">313755</a>	30370	2717	24	521	membrane metallo-endopeptidase-like 1 (60.5 kD) (Mmel1) alternative variant bSep08, mRNA.
<a href="#">Mmel1</a>	<a href="#">Mmel1.cSep08</a>	<a href="#">313755</a>	22958	3116	9	313	membrane metallo-endopeptidase-like 1 (35.4 kD) (Mmel1) alternative variant cSep08, mRNA.
<a href="#">Mmel1</a>	<a href="#">Mmel1.dSep08</a>	<a href="#">313755</a>	17845	572	5	190	metallo-endopeptidase-like 1 (Mmel1) alternative variant dSep08, mRNA.
<a href="#">Mmel1</a>	<a href="#">Mmel1.fSep08</a>	<a href="#">313755</a>	846	494	3	62	metallo-endopeptidase-like 1 (Mmel1) alternative variant fSep08, mRNA.
<a href="#">Mmp1b</a>	<a href="#">Mmp1b.aSep08</a>	<a href="#">300338</a>	52415	675		160	matrix metalloproteinase 1b (interstitial collagenase) (Mmp1b) mRNA.
<a href="#">Mmp2</a>	<a href="#">Mmp2.bSep08</a>	<a href="#">81686</a>	25562	2947	2	586	matrix metalloproteinase 2 (Mmp2) alternative variant bSep08, mRNA.
<a href="#">Mmp2</a>	<a href="#">Mmp2.cSep08</a>	<a href="#">81686</a>	21474	2042	3	527	matrix metalloproteinase 2 (Mmp2) alternative variant cSep08, mRNA.
<a href="#">Mmp2</a>	<a href="#">Mmp2.dSep08</a>	<a href="#">81686</a>	3634	742	1	121	matrix metalloproteinase 2 (Mmp2) alternative variant dSep08, mRNA.
<a href="#">Mmp11</a>	<a href="#">Mmp11.bSep08</a>	<a href="#">25481</a>	1717	1057	1	135	matrix metalloproteinase 11 (Mmp11) alternative variant bSep08, mRNA.
<a href="#">Mmp13</a>	<a href="#">Mmp13.aSep08</a>	<a href="#">171052</a>	2881	1519		114	matrix metalloproteinase 13 (Mmp13) mRNA.
<a href="#">Mmp15</a>	<a href="#">Mmp15.bSep08</a>	<a href="#">291848</a>	1067	764	1	108	matrix metalloproteinase 15 (Mmp15) alternative variant bSep08, mRNA.
<a href="#">Mmp16</a>	<a href="#">Mmp16.bSep08</a>	<a href="#">65205</a>	6886	1827	3	183	matrix metalloproteinase 16 (Mmp16) alternative variant bSep08, mRNA.
<a href="#">Mmp16</a>	<a href="#">Mmp16.cSep08</a>	<a href="#">65205</a>	646	550	1	68	matrix metalloproteinase 16 (Mmp16) alternative variant cSep08, mRNA.
<a href="#">Mmp17</a>	<a href="#">Mmp17.aSep08</a>	<a href="#">288626</a>	4325	1785	1	592	matrix metalloproteinase 17 (Mmp17) alternative variant aSep08, mRNA.
<a href="#">Mmp19</a>	<a href="#">Mmp19.bSep08</a>	<a href="#">304608</a>	4123	384	1	128	matrix metalloproteinase 19 (Mmp19) alternative variant bSep08, mRNA.
<a href="#">Mmp23</a>	<a href="#">Mmp23.aSep08</a>	<a href="#">94339</a>	2577	1259	7	262	metalloproteinase (Mmp23) alternative variant aSep08, mRNA.
<a href="#">Mmp23</a>	<a href="#">Mmp23.bSep08</a>	<a href="#">94339</a>	1463	729	5	224	metalloproteinase (Mmp23) alternative variant bSep08, mRNA.
<a href="#">Mmp23</a>	<a href="#">Mmp23.cSep08</a>	<a href="#">94339</a>	1031	644	4	214	metalloproteinase (Mmp23) alternative variant cSep08, mRNA.

<a href="#">Mmp23</a>	<a href="#">Mmp23.dSep08</a>	<a href="#">94339</a>	1097	790	4	171	metalloprotease (20.1 kD) (Mmp23) alternative variant dSep08, mRNA.
<a href="#">Mmp23</a>	<a href="#">Mmp23.eSep08</a>	<a href="#">94339</a>	1350	733	3	161	matrix metalloproteinase 23 (Mmp23) alternative variant eSep08, mRNA.
<a href="#">Mmp28</a>	<a href="#">Mmp28.bSep08</a>	<a href="#">303384</a>	16515	1783	4	546	matrix metalloproteinase 28 (epilysin) (Mmp28) alternative variant bSep08, mRNA.
<a href="#">Mmrn1</a>	<a href="#">Mmrn1.aSep08</a>	<a href="#">500152</a>	21618	1461		236	multimerin 1 (Mmrn1) mRNA.
<a href="#">Mnd1</a>	<a href="#">Mnd1.aSep08</a>	<a href="#">295160</a>	48039	647	2	162	meiotic nuclear divisions 1 homolog (S. cerevisiae) (Mnd1) alternative variant aSep08, mRNA.
<a href="#">Mnd1</a>	<a href="#">Mnd1.bSep08</a>	<a href="#">295160</a>	12740	424	1	25	meiotic nuclear divisions 1 homolog (S. cerevisiae) (Mnd1) alternative variant bSep08, mRNA.
<a href="#">Moap1</a>	<a href="#">Moap1.bSep08</a>	<a href="#">299261</a>	1238	1152	2	285	putative protein (Moap1) alternative variant bSep08, mRNA.
<a href="#">Moap1</a>	<a href="#">Moap1.dSep08</a>	<a href="#">299261</a>	4527	920	2	97	CRA b like (10.2 kD) (Moap1) alternative variant dSep08, complete mRNA.
<a href="#">Moap1</a>	<a href="#">Moap1.eSep08</a>	<a href="#">299261</a>	16127	1058	2	75	CRA b like (7.9 kD) (Moap1) alternative variant eSep08, mRNA.
<a href="#">Moap1</a>	<a href="#">Moap1.fSep08</a>	<a href="#">299261</a>	5688	339	2	69	paraneoplastic antigen MA1 like (7.8 kD) (Moap1) alternative variant fSep08, mRNA.
<a href="#">Mobkl2a</a>	<a href="#">Mobkl2a.aSep08</a>	<a href="#">362833</a>	14464	1422	2	246	MOB1, Mps One Binder kinase activator-like 2A (yeast) (28.6 kD) (Mobkl2a) alternative variant aSep08, complete mRNA.
<a href="#">Mobkl2b</a>	<a href="#">Mobkl2b.aSep08</a>	<a href="#">366352</a>	111904	651		191	MOB1, Mps One Binder kinase activator-like 2B (yeast) (Mobkl2b) mRNA.
<a href="#">Mobkl2c</a>	<a href="#">Mobkl2c.bSep08</a>	<a href="#">313511</a>	3100	766	2	216	MOB1, Mps One Binder kinase activator-like 2C (yeast) (25.6 kD) (Mobkl2c) alternative variant bSep08, mRNA.
<a href="#">Mobkl3</a>	<a href="#">Mobkl3.bSep08</a>	<a href="#">171050</a>	22404	905	6	126	MOB1, Mps One Binder kinase activator-like 3 (yeast) (14.6 kD) (Mobkl3) alternative variant bSep08, complete mRNA.
<a href="#">Mobkl3</a>	<a href="#">Mobkl3.cSep08</a>	<a href="#">171050</a>	3334	685	2	97	MOB1, Mps One Binder kinase activator-like 3 (yeast) (11.3 kD) (Mobkl3) alternative variant cSep08, mRNA.
<a href="#">Mobp</a>	<a href="#">Mobp.aSep08</a>	<a href="#">25037</a>	29681	2410	4	81	myelin-associated oligodendrocytic basic protein (9.7 kD) (Mobp) alternative variant aSep08, complete mRNA.
<a href="#">Mobp</a>	<a href="#">Mobp.bSep08</a>	<a href="#">25037</a>	28069	875	5	81	myelin-associated oligodendrocytic basic protein (9.7 kD) (Mobp) alternative variant bSep08, complete mRNA.
<a href="#">Mobp</a>	<a href="#">Mobp.cSep08</a>	<a href="#">25037</a>	17862	792	4	170	myelin-associated oligodendrocytic basic protein (19.1 kD) (Mobp) alternative variant cSep08, complete mRNA.
<a href="#">Mobp</a>	<a href="#">Mobp.dSep08</a>	<a href="#">25037</a>	17515	409	3	99	myelin-associated oligodendrocytic basic protein (Mobp) alternative variant dSep08, mRNA.
<a href="#">Mobp</a>	<a href="#">Mobp.eSep08</a>	<a href="#">25037</a>	22153	344	2		
<a href="#">moby</a>	<a href="#">moby.aSep08</a>		2695	1279		51	putative protein (moby) mRNA.
<a href="#">mochy</a>	<a href="#">mochy.aSep08</a>		11695	803	3	247	CRA b (mochy) alternative variant aSep08, mRNA.
<a href="#">mochy</a>	<a href="#">mochy.bSep08</a>		6992	520	1	136	putative protein of vertebrate origin (mochy) alternative variant bSep08, mRNA.
<a href="#">Mocos</a>	<a href="#">Mocos.bSep08</a>	<a href="#">361300</a>	18953	549	5	183	molybdenum cofactor sulfurase (Mocos) alternative variant bSep08, mRNA.

<a href="#">Mocos</a>	<a href="#">Mocos.cSep08</a>	<a href="#">361300</a>	3357	760	2	154	molybdenum cofactor sulfurase (Mocos) alternative variant cSep08, mRNA.
<a href="#">Mocos</a>	<a href="#">Mocos.dSep08</a>	<a href="#">361300</a>	16193	496	4	103	molybdenum cofactor sulfurase (Mocos) alternative variant dSep08, mRNA.
<a href="#">Mocos</a>	<a href="#">Mocos.eSep08</a>	<a href="#">361300</a>	4665	582	3	62	molybdenum cofactor sulfurase (6.8 kD) (Mocos) alternative variant eSep08, mRNA.
<a href="#">Mocos</a>	<a href="#">Mocos.fSep08</a>	<a href="#">361300</a>	4874	424	4	54	molybdenum cofactor sulfurase (Mocos) alternative variant fSep08, mRNA.
<a href="#">Mocs1</a>	<a href="#">Mocs1.bSep08</a>	<a href="#">301221</a>	13099	754	6	230	molybdenum cofactor synthesis 1 (Mocs1) alternative variant bSep08, mRNA.
<a href="#">Mocs1</a>	<a href="#">Mocs1.cSep08</a>	<a href="#">301221</a>	10401	442	3	103	molybdenum cofactor synthesis 1 (Mocs1) alternative variant cSep08, mRNA.
<a href="#">Mocs1</a>	<a href="#">Mocs1.dSep08</a>	<a href="#">301221</a>	6485	1583	4	76	molybdenum cofactor synthesis 1 (Mocs1) alternative variant dSep08, mRNA.
<a href="#">Mocs2</a>	<a href="#">Mocs2.bSep08</a>	<a href="#">294753</a>	11011	830	7	189	molybdenum cofactor synthesis 2 CRA a (21.0 kD) (Mocs2) alternative variant bSep08, complete mRNA.
<a href="#">Mocs2</a>	<a href="#">Mocs2.cSep08</a>	<a href="#">294753</a>	7397	638	6	110	molybdenum cofactor synthesis 2 CRA a (Mocs2) alternative variant cSep08, mRNA.
<a href="#">Mocs2</a>	<a href="#">Mocs2.dSep08</a>	<a href="#">294753</a>	7420	612	5	88	molybdopterin synthase Mocs2A (9.7 kD) (Mocs2) alternative variant dSep08, mRNA.
<a href="#">modar</a>	<a href="#">modar.aSep08</a>		4650	344		109	ubiquitin carboxyl-terminal hydrolase cyld (modar) mRNA.
<a href="#">mofer</a>	<a href="#">mofer.aSep08</a>		16756	804	4	100	putative protein (mofer) alternative variant aSep08, mRNA.
<a href="#">mofer</a>	<a href="#">mofer.bSep08</a>		8619	635	5	95	putative protein (10.6 kD) (mofer) alternative variant bSep08, mRNA.
<a href="#">mofer</a>	<a href="#">mofer.cSep08</a>		3465	1137	2	45	putative protein (5.0 kD) (mofer) alternative variant cSep08, mRNA.
<a href="#">mofer</a>	<a href="#">mofer.dSep08</a>		43600	717	5	70	putative protein (8.0 kD) (mofer) alternative variant dSep08, mRNA.
<a href="#">moflo</a>	<a href="#">moflo.aSep08</a>		1092	309		60	putative protein (moflo) mRNA.
<a href="#">moflu</a>	<a href="#">moflu.aSep08</a>		129328	568		81	putative protein (moflu) mRNA.
<a href="#">mogar</a>	<a href="#">mogar.aSep08</a>		746	235		39	putative protein (mogar) mRNA.
<a href="#">Mogat1</a>	<a href="#">Mogat1.bSep08</a>	<a href="#">363261</a>	9268	897	2	163	monoacylglycerol O-acyltransferase 1 (18.4 kD) (Mogat1) alternative variant bSep08, mRNA.
<a href="#">Mogat2</a>	<a href="#">Mogat2.aSep08</a>	<a href="#">681211</a>	24369	1748	6	334	monoacylglycerol O-acyltransferase 2 (Mogat2) alternative variant aSep08, complete mRNA.
<a href="#">Mogat2</a>	<a href="#">Mogat2.cSep08</a>	<a href="#">681211</a>	2779	702	2	46	monoacylglycerol O-acyltransferase 2 (Mogat2) alternative variant cSep08, mRNA.
<a href="#">mojey</a>	<a href="#">mojey.aSep08</a>		2008	376		125	protein Tyrosine phosphatase non-receptor type 13 (mojey) mRNA.
<a href="#">mokee</a>	<a href="#">mokee.aSep08</a>		7119	403		134	protein odr-4 homolog (mokee) mRNA.
<a href="#">moloy</a>	<a href="#">moloy.aSep08</a>		4188	616		126	putative protein (13.9 kD) (moloy) alternative variant aSep08, mRNA.
<a href="#">moloy</a>	<a href="#">moloy.bSep08</a>		4233	808	1	91	putative protein (moloy) alternative variant bSep08, mRNA.
<a href="#">momee</a>	<a href="#">momee.aSep08</a>		962	362	3	46	wd repeat-containing protein 90 like (momee) alternative variant aSep08, mRNA.

<a href="#">momer</a>	<a href="#">momer.bSep08</a>		1204	720	2	99	putative protein (momer) alternative variant bSep08, mRNA.
<a href="#">Mon1a</a>	<a href="#">Mon1a.aSep08</a>	<a href="#">315999</a>	20936	2044		555	MON1 homolog A (yeast) (62.2 kD) (Mon1a) mRNA.
<a href="#">Mon1b</a>	<a href="#">Mon1b.bSep08</a>	<a href="#">307868</a>	2926	939	3	228	MON1 homolog b (yeast) (Mon1b) alternative variant bSep08, mRNA.
<a href="#">Mon2</a>	<a href="#">Mon2.aSep08</a>	<a href="#">314894</a>	26061	4065	13	771	mon2 homolog CRA c (Mon2) alternative variant aSep08, mRNA.
<a href="#">Mon2</a>	<a href="#">Mon2.bSep08</a>	<a href="#">314894</a>	57929	2593	13	674	mon2 homolog CRA b (Mon2) alternative variant bSep08, mRNA.
<a href="#">Mon2</a>	<a href="#">Mon2.cSep08</a>	<a href="#">314894</a>	5300	628	4	208	mon2 homolog CRA d (Mon2) alternative variant cSep08, mRNA.
<a href="#">Mon2</a>	<a href="#">Mon2.dSep08</a>	<a href="#">314894</a>	18920	672	4	152	putative protein (Mon2) alternative variant dSep08, mRNA.
<a href="#">monoy</a>	<a href="#">monoy.aSep08</a>		18837	732		62	putative protein (monoy) mRNA.
<a href="#">mopor</a>	<a href="#">mopor.aSep08</a>		10346	733		243	D-glucuronyl (mopor) mRNA.
<a href="#">morby</a>	<a href="#">morby.aSep08</a>		251605	437		27	putative protein (morby) mRNA.
<a href="#">Morc1</a>	<a href="#">Morc1.aSep08</a>	<a href="#">360712</a>	32859	1784		435	microrchidia 1 (Morc1) alternative variant aSep08, mRNA.
<a href="#">Morc1</a>	<a href="#">Morc1.bSep08</a>	<a href="#">360712</a>	32346	765		180	microrchidia 1 (Morc1) alternative variant bSep08, mRNA.
<a href="#">Morc3</a>	<a href="#">Morc3.bSep08</a>	<a href="#">304074</a>	8225	965	5	321	microrchidia 3 (Morc3) alternative variant bSep08, mRNA.
<a href="#">Morc3</a>	<a href="#">Morc3.cSep08</a>	<a href="#">304074</a>	19739	1048	9	306	zinc finger, CW-type (Morc3) alternative variant cSep08, mRNA.
<a href="#">Morc3</a>	<a href="#">Morc3.dSep08</a>	<a href="#">304074</a>	1229	403	2	117	putative protein of eukaryotic origin (Morc3) alternative variant dSep08, mRNA.
<a href="#">Morc3</a>	<a href="#">Morc3.fSep08</a>	<a href="#">304074</a>	2603	635	3	38	putative protein of vertebrate origin (Morc3) alternative variant fSep08, mRNA.
<a href="#">Morc4</a>	<a href="#">Morc4.aSep08</a>	<a href="#">315914</a>	33317	3057	11	617	microrchidia 4 (Morc4) alternative variant aSep08, mRNA.
<a href="#">Morc4</a>	<a href="#">Morc4.bSep08</a>	<a href="#">315914</a>	1347	1045	2	114	microrchidia 4 (Morc4) alternative variant bSep08, mRNA.
<a href="#">morchy</a>	<a href="#">morchy.aSep08</a>		18516	782		105	putative protein (12.0 kD) (morchy) mRNA.
<a href="#">mordar</a>	<a href="#">mordar.aSep08</a>		3356	661	4	74	putative protein (mordar) alternative variant aSep08, mRNA.
<a href="#">Morf4I1.1</a>	<a href="#">Morf4I1.1.bSep08</a>	<a href="#">300891</a>	18646	1041	11	296	mortality factor 4 like 1 (Morf4I1.1) alternative variant bSep08, mRNA.
<a href="#">Morf4I1.1</a>	<a href="#">Morf4I1.1.cSep08</a>	<a href="#">300891</a>	2809	414	4	138	mortality factor 4 like 1 (Morf4I1.1) alternative variant cSep08, mRNA.
<a href="#">Morf4I1.1</a>	<a href="#">Morf4I1.1.dSep08</a>	<a href="#">300891</a>	3207	1464	2	43	mortality factor 4 like 1 (4.8 kD) (Morf4I1.1) alternative variant dSep08, mRNA.
<a href="#">Morf4I2</a>	<a href="#">Morf4I2.aSep08</a>	<a href="#">317413</a>	7787	932	3	251	mortality factor 4 like 2 (Morf4I2) alternative variant aSep08, mRNA.
<a href="#">Morf4I2</a>	<a href="#">Morf4I2.bSep08</a>	<a href="#">317413</a>	8601	1847	4	288	mortality factor 4 like 2 (32.2 kD) (Morf4I2) alternative variant bSep08, mRNA.
<a href="#">Morf4I2</a>	<a href="#">Morf4I2.dSep08</a>	<a href="#">317413</a>	6726	1281	3	78	mortality factor 4 like 2 (Morf4I2) alternative variant dSep08, mRNA.
<a href="#">Morf4I2</a>	<a href="#">Morf4I2.fSep08</a>	<a href="#">317413</a>	9154	718	4	136	mortality factor 4 like 2 (Morf4I2) alternative variant fSep08, mRNA.
<a href="#">Morf4I2</a>	<a href="#">Morf4I2.gSep08</a>	<a href="#">317413</a>	3358	367	2	24	mortality factor 4 like 2 (2.8 kD) (Morf4I2) alternative variant gSep08, mRNA.

<a href="#">morfer</a>	<a href="#">morfer.aSep08</a>		5998	1777		54	putative protein (morfer) alternative variant aSep08, mRNA.
<a href="#">morfer</a>	<a href="#">morfer.bSep08</a>		5192	411		54	putative protein (morfer) alternative variant bSep08, mRNA.
<a href="#">morflo</a>	<a href="#">morflo.aSep08</a>		2153	2026	2	233	sema 4G (morflo) alternative variant aSep08, mRNA.
<a href="#">morflu</a>	<a href="#">morflu.aSep08</a>		37375	465		62	putative protein (7.1 kD) (morflu) mRNA.
<a href="#">Morg1</a>	<a href="#">Morg1.bSep08</a>	<a href="#">288924</a>	5032	747	3	173	mitogen-activated protein kinase organizer 1 (Morg1) alternative variant bSep08, mRNA.
<a href="#">Morg1</a>	<a href="#">Morg1.cSep08</a>	<a href="#">288924</a>	3915	567	1	127	mitogen-activated protein kinase organizer 1 (Morg1) alternative variant cSep08, mRNA.
<a href="#">morgar</a>	<a href="#">morgar.aSep08</a>		21820	801	8	266	adaptor protein phosphotyrosine interaction PH domain leucine zipper containing 1 (morgar) alternative variant aSep08, mRNA.
<a href="#">morgar</a>	<a href="#">morgar.bSep08</a>		1741	348	1	35	putative protein (4.1 kD) (morgar) alternative variant bSep08, mRNA.
<a href="#">morja</a>	<a href="#">morja.aSep08</a>		34652	680		167	putative protein (morja) mRNA.
<a href="#">morjey</a>	<a href="#">morjey.aSep08</a>		6672	412		136	putative protein of vertebrate origin (morjey) mRNA.
<a href="#">morkee</a>	<a href="#">morkee.aSep08</a>		661	398		94	putative protein (morkee) mRNA.
<a href="#">morloy</a>	<a href="#">morloy.aSep08</a>		4146	768		79	protease (morloy) mRNA.
<a href="#">mormee</a>	<a href="#">mormee.aSep08</a>		17058	756		66	putative protein (7.5 kD) (mormee) mRNA.
<a href="#">mormer</a>	<a href="#">mormer.aSep08</a>		1253	572		154	kinesin family member 19A (mormer) mRNA.
<a href="#">Morn1</a>	<a href="#">Morn1.bSep08</a>	<a href="#">298676</a>	42861	1081	2	233	MORN repeat containing 1 (Morn1) alternative variant bSep08, mRNA.
<a href="#">Morn2</a>	<a href="#">Morn2.aSep08</a>	<a href="#">500606</a>	6727	615	1	165	MORN repeat containing 2 (Morn2) alternative variant aSep08, mRNA.
<a href="#">mornoy</a>	<a href="#">mornoy.aSep08</a>		12217	402		134	type III (mornoy) mRNA.
<a href="#">morpor</a>	<a href="#">morpor.aSep08</a>		5641	597		198	zwilch (morpor) mRNA.
<a href="#">morsa</a>	<a href="#">morsa.aSep08</a>		4409	597		97	putative protein (morsa) mRNA.
<a href="#">morshee</a>	<a href="#">morshee.aSep08</a>		1874	576		82	putative mitochondrial protein (9.0 kD) (morshee) mRNA.
<a href="#">mortu</a>	<a href="#">mortu.aSep08</a>		30802	301		92	uncharacterized protein c2orf61 homolog like (mortu) mRNA.
<a href="#">morvar</a>	<a href="#">morvar.aSep08</a>		1557	299		94	putative protein (morvar) mRNA.
<a href="#">morwey</a>	<a href="#">morwey.aSep08</a>		4985	543		66	putative protein (morwey) mRNA.
<a href="#">mosa</a>	<a href="#">mosa.aSep08</a>		6244	685		76	putative protein (mosa) mRNA.
<a href="#">Mosc2</a>	<a href="#">Mosc2.bSep08</a>	<a href="#">171451</a>	39830	596	3	177	MOSC, N-terminal beta barrel (Mosc2) alternative variant bSep08, mRNA.
<a href="#">Mosc2</a>	<a href="#">Mosc2.cSep08</a>	<a href="#">171451</a>	7448	690	4	74	putative mitochondrial protein of ancient origin (8.4 kD) (Mosc2) alternative variant cSep08, mRNA.
<a href="#">Mosc2</a>	<a href="#">Mosc2.dSep08</a>	<a href="#">171451</a>	3468	689	2	60	putative protein of metazoan origin (Mosc2) alternative variant dSep08, mRNA.
<a href="#">moshee</a>	<a href="#">moshee.aSep08</a>		2070	558		83	putative protein of metazoan origin (moshee) mRNA.
<a href="#">Mospd1</a>	<a href="#">Mospd1.bSep08</a>	<a href="#">317312</a>	24525	768	5	191	major sperm protein (Mospd1) alternative variant bSep08, mRNA.
<a href="#">Mospd3</a>	<a href="#">Mospd3.bSep08</a>	<a href="#">288557</a>	5045	935	5	173	major sperm protein (18.8 kD) (Mospd3) alternative variant bSep08, mRNA.

<a href="#">Mospd3</a>	<a href="#">Mospd3.cSep08</a>	<a href="#">288557</a>	3462	970	3	129	putative protein of mammalian origin (Mospd3) alternative variant cSep08, mRNA.
<a href="#">motu</a>	<a href="#">motu.aSep08</a>		2775	656		66	putative protein (motu) mRNA.
<a href="#">Mov10l1</a>	<a href="#">Mov10l1.bSep08</a>	<a href="#">300141</a>	10483	603	1	185	moloney leukemia virus 10-like 1 (Mov10l1) alternative variant bSep08, mRNA.
<a href="#">movar</a>	<a href="#">movar.aSep08</a>		598	525		71	putative protein (movar) mRNA.
<a href="#">mowey</a>	<a href="#">mowey.aSep08</a>		1101	461		40	putative protein (4.6 kD) (mowey) mRNA.
<a href="#">Moxd1</a>	<a href="#">Moxd1.aSep08</a>	<a href="#">294119</a>	59666	2486		501	monooxygenase, DBH-like 1 (Moxd1) mRNA.
<a href="#">moyby</a>	<a href="#">moyby.aSep08</a>		2620	287		31	putative protein (3.7 kD) (moyby) mRNA.
<a href="#">moychy</a>	<a href="#">moychy.aSep08</a>		700	400		52	putative protein (moychy) mRNA.
<a href="#">moydar</a>	<a href="#">moydar.aSep08</a>		39774	385		127	putative protein (moydar) mRNA.
<a href="#">moyfer</a>	<a href="#">moyfer.aSep08</a>		4931	737		69	putative protein (moyfer) mRNA.
<a href="#">moyflo</a>	<a href="#">moyflo.aSep08</a>		559	383		95	putative protein of mammalian origin (moyflo) mRNA.
<a href="#">moyflu</a>	<a href="#">moyflu.aSep08</a>		21815	603		39	putative protein (moyflu) mRNA.
<a href="#">moygar</a>	<a href="#">moygar.bSep08</a>		1758	428	3	49	putative protein (moygar) alternative variant bSep08, mRNA.
<a href="#">moyja</a>	<a href="#">moyja.aSep08</a>		10592	565		21	putative protein (2.4 kD) (moyja) mRNA.
<a href="#">moyjey</a>	<a href="#">moyjey.aSep08</a>		4532	545		181	putative protein of metazoan origin (moyjey) mRNA.
<a href="#">moykee</a>	<a href="#">moykee.aSep08</a>		97831	776		43	putative protein (4.8 kD) (moykee) mRNA.
<a href="#">moyloy</a>	<a href="#">moyloy.aSep08</a>		62478	1336		445	vacuolar protein sorting 8 homolog (moyloy) mRNA.
<a href="#">moymee</a>	<a href="#">moymee.aSep08</a>		1146	759		69	putative protein (moymee) mRNA.
<a href="#">moymmer</a>	<a href="#">moymmer.aSep08</a>		1435	995		169	kinesin family member 19A (moymmer) mRNA.
<a href="#">moynoy</a>	<a href="#">moynoy.aSep08</a>		4204	470		111	type III (moynoy) mRNA.
<a href="#">moypor</a>	<a href="#">moypor.aSep08</a>		6476	561		81	11-like domains precursor (8.8 kD) (moypor) mRNA.
<a href="#">moysa</a>	<a href="#">moysa.aSep08</a>		1215	765		30	putative protein (3.2 kD) (moysa) mRNA.
<a href="#">moyshee</a>	<a href="#">moyshee.bSep08</a>		75245	341	3	45	putative protein (moyshee) alternative variant bSep08, mRNA.
<a href="#">moytu</a>	<a href="#">moytu.aSep08</a>		8525	755		109	putative protein (moytu) mRNA.
<a href="#">moyvar</a>	<a href="#">moyvar.aSep08</a>		515	425		47	putative protein (moyvar) mRNA.
<a href="#">moywey</a>	<a href="#">moywey.aSep08</a>		3346	465		50	putative protein (moywey) mRNA.
<a href="#">Mpdu1</a>	<a href="#">Mpdu1.bSep08</a>	<a href="#">303244</a>	4992	626	4	157	mannose-P-dolichol utilization defect 1 (Mpdu1) alternative variant bSep08, mRNA.
<a href="#">Mpdu1</a>	<a href="#">Mpdu1.cSep08</a>	<a href="#">303244</a>	5125	694	5	121	mannose-P-dolichol utilization defect 1 (13.4 kD) (Mpdu1) alternative variant cSep08, mRNA.
<a href="#">Mpdu1</a>	<a href="#">Mpdu1.dSep08</a>	<a href="#">303244</a>	873	722	2	65	mannose-P-dolichol utilization defect 1 (6.7 kD) (Mpdu1) alternative variant dSep08, mRNA.
<a href="#">Mpeg1</a>	<a href="#">Mpeg1.aSep08</a>	<a href="#">64552</a>	880	598		156	putative protein of metazoan origin (Mpeg1) mRNA.
<a href="#">Mpg</a>	<a href="#">Mpg.bSep08</a>	<a href="#">24561</a>	3643	1452	2	162	N-methylpurine-DNA glycosylase (18.0 kD) (Mpg) alternative variant bSep08, mRNA.
<a href="#">Mphosph1</a>	<a href="#">Mphosph1.bSep08</a>	<a href="#">309523</a>	30894	3018	8	1006	M-phase phosphoprotein 1 (Mphosph1) alternative variant bSep08, mRNA.
<a href="#">Mphosph1</a>	<a href="#">Mphosph1.cSep08</a>	<a href="#">309523</a>	15061	1024	2	341	M-phase phosphoprotein 1 (Mphosph1) alternative variant cSep08, mRNA.

<a href="#">Mphosph8</a>	<a href="#">Mphosph8.bSep08</a>	<a href="#">290270</a>	4184	894	4	92	M-phase phosphoprotein 8 (10.2 kD) (Mphosph8) alternative variant bSep08, mRNA.
<a href="#">Mphosph8</a>	<a href="#">Mphosph8.cSep08</a>	<a href="#">290270</a>	3828	1267	2	81	M-phase phosphoprotein 8 (Mphosph8) alternative variant cSep08, mRNA.
<a href="#">Mphosph9</a>	<a href="#">Mphosph9.aSep08</a>	<a href="#">288654</a>	17125	583		193	M-phase phosphoprotein 9 (Mphosph9) mRNA.
<a href="#">Mphosph10</a>	<a href="#">Mphosph10.bSep08</a>	<a href="#">293828</a>	8555	1776	1	482	M-phase phosphoprotein 10 (U3 small nucleolar ribonucleoprotein) (Mphosph10) alternative variant bSep08, mRNA.
<a href="#">Mpi</a>	<a href="#">Mpi.bSep08</a>	<a href="#">300741</a>	8133	1891	6	315	mannose phosphate isomerase (mapped) (34.6 kD) (Mpi) alternative variant bSep08, mRNA.
<a href="#">Mpi</a>	<a href="#">Mpi.cSep08</a>	<a href="#">300741</a>	7313	1075	7	233	mannose phosphate isomerase (mapped) (Mpi) alternative variant cSep08, mRNA.
<a href="#">Mpi</a>	<a href="#">Mpi.eSep08</a>	<a href="#">300741</a>	6058	445	4	55	mannose phosphate isomerase (mapped) (6.0 kD) (Mpi) alternative variant eSep08, mRNA.
<a href="#">Mpo</a>	<a href="#">Mpo.bSep08</a>	<a href="#">303413</a>	6577	1176	3	391	myeloperoxidase (Mpo) alternative variant bSep08, mRNA.
<a href="#">Mpp2</a>	<a href="#">Mpp2.aSep08</a>	<a href="#">85275</a>	25411	395	3	100	membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) (Mpp2) alternative variant aSep08, mRNA.
<a href="#">Mpp2</a>	<a href="#">Mpp2.bSep08</a>	<a href="#">85275</a>	25153	415	3	83	membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) (Mpp2) alternative variant bSep08, mRNA.
<a href="#">Mpp2</a>	<a href="#">Mpp2.cSep08</a>	<a href="#">85275</a>	4238	634	2	38	membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) (Mpp2) alternative variant cSep08, mRNA.
<a href="#">Mpp3</a>	<a href="#">Mpp3.aSep08</a>	<a href="#">114202</a>	28823	2197	10	568	membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3) (Mpp3) alternative variant aSep08, mRNA.
<a href="#">Mpp3</a>	<a href="#">Mpp3.bSep08</a>	<a href="#">114202</a>	20804	1120	12	372	membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3) (Mpp3) alternative variant bSep08, mRNA.
<a href="#">Mpp3</a>	<a href="#">Mpp3.cSep08</a>	<a href="#">114202</a>	11265	1829	8	274	membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3) (31.2 kD) (Mpp3) alternative variant cSep08, mRNA.
<a href="#">Mpp4</a>	<a href="#">Mpp4.bSep08</a>	<a href="#">58808</a>	26593	847	7	101	membrane protein palmitoylated 4 (Mpp4) alternative variant bSep08, mRNA.
<a href="#">Mpp4</a>	<a href="#">Mpp4.cSep08</a>	<a href="#">58808</a>	9853	418	4	65	putative protein (6.8 kD) (Mpp4) alternative variant cSep08, mRNA.
<a href="#">Mpp4</a>	<a href="#">Mpp4.dSep08</a>	<a href="#">58808</a>	26464	718	7	58	membrane protein palmitoylated 4 (Mpp4) alternative variant dSep08, mRNA.
<a href="#">Mpp6</a>	<a href="#">Mpp6.aSep08</a>	<a href="#">362359</a>	80010	2634	7	504	membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6) (Mpp6) alternative variant aSep08, mRNA.
<a href="#">Mpp6</a>	<a href="#">Mpp6.bSep08</a>	<a href="#">362359</a>	12749	467	1	155	membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6) (Mpp6) alternative variant bSep08, mRNA.
<a href="#">Mpp6</a>	<a href="#">Mpp6.cSep08</a>	<a href="#">362359</a>	12732	501	3	58	membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6) (6.6 kD) (Mpp6) alternative variant cSep08, mRNA.
<a href="#">Mpp7</a>	<a href="#">Mpp7.aSep08</a>	<a href="#">307035</a>	215144	734	1	179	membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7) (Mpp7) alternative variant aSep08, mRNA.
<a href="#">Mpp7</a>	<a href="#">Mpp7.bSep08</a>	<a href="#">307035</a>	152314	608	1	157	membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7) (Mpp7) alternative variant bSep08, mRNA.
<a href="#">Mppe1</a>	<a href="#">Mppe1.bSep08</a>	<a href="#">361344</a>	3110	299	2	84	metallophosphoesterase 1 (Mppe1) alternative variant bSep08, mRNA.

<a href="#">Mpped2</a>	<a href="#">Mpped2.aSep08</a>	<a href="#">362185</a>	176795	2674	5	294	metallophosphoesterase (33.3 kD) (Mpped2) alternative variant aSep08, mRNA.
<a href="#">Mpped2</a>	<a href="#">Mpped2.bSep08</a>	<a href="#">362185</a>	3343	774	1	93	putative protein (Mpped2) alternative variant bSep08, mRNA.
<a href="#">Mpst</a>	<a href="#">Mpst.bSep08</a>	<a href="#">192172</a>	5155	1525	2	189	mercaptopyruvate sulfurtransferase (21.1 kD) (Mpst) alternative variant bSep08, mRNA.
<a href="#">Mpv17l</a>	<a href="#">Mpv17l.aSep08</a>	<a href="#">360463</a>	13427	450		111	mpv17 transgene, kidney disease mutant-like (Mpv17l) mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.aSep08</a>	<a href="#">360463</a>	9727	1038	7	154	mpv17 transgene, kidney disease mutant-like (17.1 kD) (Mpv17l.1) alternative variant aSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.cSep08</a>	<a href="#">360463</a>	9242	801	5	150	mpv17 transgene, kidney disease mutant-like (Mpv17l.1) alternative variant cSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.dSep08</a>	<a href="#">360463</a>	9145	651	5	149	mpv17 transgene, kidney disease mutant-like (16.2 kD) (Mpv17l.1) alternative variant dSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.eSep08</a>	<a href="#">360463</a>	9335	613	6	143	mpv17 transgene, kidney disease mutant-like (15.8 kD) (Mpv17l.1) alternative variant eSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.fSep08</a>	<a href="#">360463</a>	9875	1118	6	136	mpv17 transgene, kidney disease mutant-like (Mpv17l.1) alternative variant fSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.gSep08</a>	<a href="#">360463</a>	9455	698	6	133	mpv17 transgene, kidney disease mutant-like (14.4 kD) (Mpv17l.1) alternative variant gSep08, mRNA.
<a href="#">Mpv17l.1</a>	<a href="#">Mpv17l.1.hSep08</a>	<a href="#">360463</a>	1253	392	3	67	mpv17 transgene, kidney disease mutant-like (7.4 kD) (Mpv17l.1) alternative variant hSep08, complete mRNA.
<a href="#">Mpz</a>	<a href="#">Mpz.bSep08</a>	<a href="#">24564</a>	3626	577	3	170	myelin protein zero (18.8 kD) (Mpz) alternative variant bSep08, mRNA.
<a href="#">Mpzl1</a>	<a href="#">Mpzl1.bSep08</a>	<a href="#">360871</a>	35886	1273		250	myelin protein zero-like 1 (Mpzl1) alternative variant bSep08, mRNA.
<a href="#">Mpzl3</a>	<a href="#">Mpzl3.bSep08</a>	<a href="#">363054</a>	6446	375	1	124	myelin protein zero-like 3 (Mpzl3) alternative variant bSep08, mRNA.
<a href="#">Mrap</a>	<a href="#">Mrap.aSep08</a>	<a href="#">288271</a>	10494	837	1	153	melanocortin 2 receptor accessory protein (Mrap) alternative variant aSep08, mRNA.
<a href="#">Mrap</a>	<a href="#">Mrap.bSep08</a>	<a href="#">288271</a>	10173	568	1	92	melanocortin 2 receptor accessory protein (Mrap) alternative variant bSep08, mRNA.
<a href="#">Mras</a>	<a href="#">Mras.bSep08</a>	<a href="#">25482</a>	3311	407	1	124	putative protein (Mras) alternative variant bSep08, mRNA.
<a href="#">Mrc2</a>	<a href="#">Mrc2.bSep08</a>	<a href="#">498011</a>	11618	3158	13	615	mannose receptor, C type 2 (Mrc2) alternative variant bSep08, mRNA.
<a href="#">Mrc2</a>	<a href="#">Mrc2.cSep08</a>	<a href="#">498011</a>	612	390	2	129	mannose receptor, C type 2 (Mrc2) alternative variant cSep08, mRNA.
<a href="#">Mreg</a>	<a href="#">Mreg.aSep08</a>	<a href="#">501162</a>	34308	602		160	melanoregulin (18.9 kD) (Mreg) mRNA.
<a href="#">Mrlcb</a>	<a href="#">Mrlcb.aSep08</a>	<a href="#">50685</a>	14798	1167	4	172	myosin light chain, regulatory B (19.8 kD) (Mrlcb) alternative variant aSep08, complete mRNA.
<a href="#">Mrm1</a>	<a href="#">Mrm1.aSep08</a>	<a href="#">363661</a>	1447	1157	2	221	mitochondrial rRNA methyltransferase 1 homolog (S. cerevisiae) (24.3 kD) (Mrm1) alternative variant aSep08, mRNA.
<a href="#">Mrm1</a>	<a href="#">Mrm1.bSep08</a>	<a href="#">363661</a>	3351	542	3	141	mitochondrial rRNA methyltransferase 1 homolog (S. cerevisiae) (Mrm1) alternative variant bSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.aSep08</a>	<a href="#">361348</a>	18456	716	5	238	maestro (Mro) alternative variant aSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.aSep08</a>	<a href="#">680524</a>	18456	716	5	238	maestro (Mro) alternative variant aSep08, mRNA.



<a href="#">Mro</a>	<a href="#">Mro.bSep08</a>	<a href="#">361348</a>	18542	961	5	169	putative protein (Mro) alternative variant bSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.bSep08</a>	<a href="#">680524</a>	18542	961	5	169	putative protein (Mro) alternative variant bSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.cSep08</a>	<a href="#">361348</a>	10282	474	4	132	putative protein (Mro) alternative variant cSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.cSep08</a>	<a href="#">680524</a>	10282	474	4	132	putative protein (Mro) alternative variant cSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.dSep08</a>	<a href="#">361348</a>	7988	523	3	116	putative protein (Mro) alternative variant dSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.dSep08</a>	<a href="#">680524</a>	7988	523	3	116	putative protein (Mro) alternative variant dSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.eSep08</a>	<a href="#">361348</a>	1971	1093	3	105	maestro (Mro) alternative variant eSep08, mRNA.
<a href="#">Mro</a>	<a href="#">Mro.eSep08</a>	<a href="#">680524</a>	1971	1093	3	105	maestro (Mro) alternative variant eSep08, mRNA.
<a href="#">Mrpl1</a>	<a href="#">Mrpl1.bSep08</a>	<a href="#">289491</a>	6401	334	1	51	mitochondrial ribosomal protein L1 (Mrpl1) alternative variant bSep08, mRNA.
<a href="#">Mrpl2</a>	<a href="#">Mrpl2.bSep08</a>	<a href="#">301240</a>	2542	733	1	140	mitochondrial ribosomal protein L2 (15.5 kD) (Mrpl2) alternative variant bSep08, mRNA.
<a href="#">Mrpl3</a>	<a href="#">Mrpl3.bSep08</a>	<a href="#">300974</a>	3304	580	5	193	mitochondrial ribosomal protein L3 (Mrpl3) alternative variant bSep08, mRNA.
<a href="#">Mrpl3</a>	<a href="#">Mrpl3.cSep08</a>	<a href="#">300974</a>	3392	583	4	145	mitochondrial ribosomal protein L3 (Mrpl3) alternative variant cSep08, mRNA.
<a href="#">Mrpl3</a>	<a href="#">Mrpl3.dSep08</a>	<a href="#">300974</a>	5756	792	4	130	mitochondrial ribosomal protein L3 (14.7 kD) (Mrpl3) alternative variant dSep08, mRNA.
<a href="#">Mrpl3</a>	<a href="#">Mrpl3.eSep08</a>	<a href="#">300974</a>	3783	358	3	106	mitochondrial ribosomal protein L3 (Mrpl3) alternative variant eSep08, mRNA.
<a href="#">Mrpl3</a>	<a href="#">Mrpl3.gSep08</a>	<a href="#">300974</a>	1333	337	2	44	mitochondrial ribosomal protein L3 (Mrpl3) alternative variant gSep08, mRNA.
<a href="#">Mrpl4</a>	<a href="#">Mrpl4.bSep08</a>	<a href="#">363023</a>	4620	800	3	159	mitochondrial ribosomal protein L4 (Mrpl4) alternative variant bSep08, mRNA.
<a href="#">Mrpl4</a>	<a href="#">Mrpl4.cSep08</a>	<a href="#">363023</a>	1417	877	2	129	mitochondrial ribosomal protein L4 (Mrpl4) alternative variant cSep08, mRNA.
<a href="#">mrpl9</a>	<a href="#">mrpl9.bSep08</a>	<a href="#">310653</a>	3271	783	5	178	mitochondrial ribosomal protein L9 (20.4 kD) (mrpl9) alternative variant bSep08, mRNA.
<a href="#">mrpl9</a>	<a href="#">mrpl9.cSep08</a>	<a href="#">310653</a>	2923	416	4	131	mitochondrial ribosomal protein L9 (mrpl9) alternative variant cSep08, mRNA.
<a href="#">mrpl11</a>	<a href="#">mrpl11.bSep08</a>	<a href="#">293666</a>	3357	1211	2	160	mitochondrial ribosomal protein L11 (17.0 kD) (mrpl11) alternative variant bSep08, complete mRNA.
<a href="#">mrpl11</a>	<a href="#">mrpl11.cSep08</a>	<a href="#">293666</a>	2840	651	3	152	mitochondrial ribosomal protein L11 (16.6 kD) (mrpl11) alternative variant cSep08, complete mRNA.
<a href="#">Mrpl13</a>	<a href="#">Mrpl13.aSep08</a>	<a href="#">299938</a>	21902	847	6	237	mitochondrial ribosomal protein L13 (Mrpl13) alternative variant aSep08, mRNA.
<a href="#">Mrpl13</a>	<a href="#">Mrpl13.cSep08</a>	<a href="#">299938</a>	21608	969	7	162	mitochondrial ribosomal protein L13 (18.7 kD) (Mrpl13) alternative variant cSep08, complete mRNA.
<a href="#">Mrpl14</a>	<a href="#">Mrpl14.aSep08</a>	<a href="#">301250</a>	10801	621	1	175	mitochondrial ribosomal protein L14 (Mrpl14) alternative variant aSep08, mRNA.
<a href="#">Mrpl15</a>	<a href="#">Mrpl15.aSep08</a>	<a href="#">297799</a>	9495	2191	5	317	mitochondrial ribosomal protein L15 (Mrpl15) alternative variant aSep08, mRNA.
<a href="#">Mrpl16</a>	<a href="#">Mrpl16.bSep08</a>	<a href="#">293754</a>	886	611	2	65	mitochondrial ribosomal protein L16 (7.4 kD) (Mrpl16) alternative variant bSep08, mRNA.
<a href="#">Mrpl17</a>	<a href="#">Mrpl17.aSep08</a>	<a href="#">171061</a>	1666	974	2	197	mitochondrial ribosomal protein L17 (Mrpl17) alternative variant aSep08, mRNA.

<a href="#">Mrpl17</a>	<a href="#">Mrpl17.cSep08</a>	<a href="#">171061</a>	1584	635	2	118	mitochondrial ribosomal protein L17 (Mrpl17) alternative variant cSep08, mRNA.
<a href="#">Mrpl18</a>	<a href="#">Mrpl18.bSep08</a>	<a href="#">292244</a>	4087	678	1	163	mitochondrial ribosomal protein L18 (18.4 kD) (Mrpl18) alternative variant bSep08, complete mRNA.
<a href="#">Mrpl22</a>	<a href="#">Mrpl22.bSep08</a>	<a href="#">287302</a>	10295	578	4	126	mitochondrial ribosomal protein L22 (14.6 kD) (Mrpl22) alternative variant bSep08, complete mRNA.
<a href="#">Mrpl22</a>	<a href="#">Mrpl22.cSep08</a>	<a href="#">287302</a>	3834	705	1	116	mitochondrial ribosomal protein L22 (13.4 kD) (Mrpl22) alternative variant cSep08, mRNA.
<a href="#">Mrpl23</a>	<a href="#">Mrpl23.aSep08</a>	<a href="#">64360</a>	7438	562	1	158	mitochondrial ribosomal protein L23 (Mrpl23) alternative variant aSep08, mRNA.
<a href="#">Mrpl23</a>	<a href="#">Mrpl23.cSep08</a>	<a href="#">64360</a>	7734	1343	1	101	mitochondrial ribosomal protein L23 (11.8 kD) (Mrpl23) alternative variant cSep08, mRNA.
<a href="#">mrpl24</a>	<a href="#">mrpl24.aSep08</a>	<a href="#">295224</a>	5127	821	6	216	mitochondrial ribosomal protein L24 (25.0 kD) (mrpl24) alternative variant aSep08, complete mRNA.
<a href="#">mrpl24</a>	<a href="#">mrpl24.bSep08</a>	<a href="#">295224</a>	5245	859	6	216	mitochondrial ribosomal protein L24 (25.0 kD) (mrpl24) alternative variant bSep08, complete mRNA.
<a href="#">mrpl24</a>	<a href="#">mrpl24.dSep08</a>	<a href="#">295224</a>	1447	851	4	189	mitochondrial ribosomal protein L24 (21.9 kD) (mrpl24) alternative variant dSep08, mRNA.
<a href="#">mrpl24</a>	<a href="#">mrpl24.eSep08</a>	<a href="#">295224</a>	4746	604	5	146	mitochondrial ribosomal protein L24 (mrpl24) alternative variant eSep08, mRNA.
<a href="#">Mrpl28</a>	<a href="#">Mrpl28.aSep08</a>	<a href="#">497876</a>	2922	1084	5	283	mitochondrial ribosomal protein L28 (Mrpl28) alternative variant aSep08, mRNA.
<a href="#">Mrpl28</a>	<a href="#">Mrpl28.bSep08</a>	<a href="#">497876</a>	2028	636	4	152	mitochondrial ribosomal protein L28 (18.0 kD) (Mrpl28) alternative variant bSep08, mRNA.
<a href="#">Mrpl28</a>	<a href="#">Mrpl28.dSep08</a>	<a href="#">497876</a>	1210	473	2	88	mitochondrial ribosomal protein L28 (Mrpl28) alternative variant dSep08, mRNA.
<a href="#">Mrpl30</a>	<a href="#">Mrpl30.bSep08</a>	<a href="#">301352</a>	11806	2076	1	160	mitochondrial ribosomal protein L30 (18.4 kD) (Mrpl30) alternative variant bSep08, complete mRNA.
<a href="#">Mrpl35</a>	<a href="#">Mrpl35.bSep08</a>	<a href="#">297334</a>	2720	563	2	132	mitochondrial ribosomal protein L35 (Mrpl35) alternative variant bSep08, mRNA.
<a href="#">Mrpl35</a>	<a href="#">Mrpl35.cSep08</a>	<a href="#">297334</a>	8655	781	2	78	mitochondrial ribosomal protein L35 (8.6 kD) (Mrpl35) alternative variant cSep08, mRNA.
<a href="#">Mrpl35</a>	<a href="#">Mrpl35.dSep08</a>	<a href="#">297334</a>	6647	705	3	88	mitochondrial ribosomal protein L35 (Mrpl35) alternative variant dSep08, mRNA.
<a href="#">Mrpl36</a>	<a href="#">Mrpl36.aSep08</a>	<a href="#">364656</a>	1096	857	1	133	mitochondrial ribosomal protein L36 (Mrpl36) alternative variant aSep08, mRNA.
<a href="#">Mrpl37</a>	<a href="#">Mrpl37.bSep08</a>	<a href="#">56281</a>	9118	1026	4	286	mitochondrial ribosomal protein L37 (Mrpl37) alternative variant bSep08, mRNA.
<a href="#">Mrpl37</a>	<a href="#">Mrpl37.cSep08</a>	<a href="#">56281</a>	12800	2001	4	180	mitochondrial ribosomal protein L37 (20.6 kD) (Mrpl37) alternative variant cSep08, mRNA.
<a href="#">Mrpl37</a>	<a href="#">Mrpl37.dSep08</a>	<a href="#">56281</a>	5831	748	5	156	mitochondrial ribosomal protein L37 (Mrpl37) alternative variant dSep08, mRNA.
<a href="#">Mrpl38</a>	<a href="#">Mrpl38.bSep08</a>	<a href="#">303685</a>	3319	405	3	133	mitochondrial ribosomal protein l38 CRA a (Mrpl38) alternative variant bSep08, mRNA.
<a href="#">Mrpl38</a>	<a href="#">Mrpl38.cSep08</a>	<a href="#">303685</a>	2679	553	4	126	PEBP (Mrpl38) alternative variant cSep08, mRNA.
<a href="#">Mrpl38</a>	<a href="#">Mrpl38.dSep08</a>	<a href="#">303685</a>	3480	705	4	99	mitochondrial ribosomal protein L38 CRA a (11.5 kD) (Mrpl38) alternative variant dSep08, mRNA.

<a href="#">Mrpl38</a>	<a href="#">Mrpl38.eSep08</a>	<a href="#">303685</a>	1266	1066	2	95	putative protein of eukaryotic origin (Mrpl38) alternative variant eSep08, mRNA.
<a href="#">Mrpl42</a>	<a href="#">Mrpl42.aSep08</a>	<a href="#">299743</a>	22293	568	1	147	mitochondrial ribosomal protein L42 (Mrpl42) alternative variant aSep08, mRNA.
<a href="#">Mrpl42</a>	<a href="#">Mrpl42.cSep08</a>	<a href="#">299743</a>	23733	663	3	141	mitochondrial ribosomal protein L42 (16.6 kD) (Mrpl42) alternative variant cSep08, complete mRNA.
<a href="#">Mrpl42</a>	<a href="#">Mrpl42.dSep08</a>	<a href="#">299743</a>	23736	545	2	114	mitochondrial ribosomal protein L42 (13.4 kD) (Mrpl42) alternative variant dSep08, complete mRNA.
<a href="#">Mrpl43</a>	<a href="#">Mrpl43.aSep08</a>	<a href="#">309440</a>	869	664	2	174	mitochondrial ribosomal protein L43 (Mrpl43) alternative variant aSep08, mRNA.
<a href="#">Mrpl45</a>	<a href="#">Mrpl45.bSep08</a>	<a href="#">287656</a>	8719	961	1	171	mitochondrial ribosomal protein L45 (Mrpl45) alternative variant bSep08, mRNA.
<a href="#">Mrpl47</a>	<a href="#">Mrpl47.bSep08</a>	<a href="#">294963</a>	10313	828	7	213	mitochondrial ribosomal protein L47 (25.7 kD) (Mrpl47) alternative variant bSep08, mRNA.
<a href="#">Mrpl47</a>	<a href="#">Mrpl47.dSep08</a>	<a href="#">294963</a>	6281	336	4	53	mitochondrial ribosomal protein L47 (Mrpl47) alternative variant dSep08, mRNA.
<a href="#">Mrpl48</a>	<a href="#">Mrpl48.aSep08</a>	<a href="#">293149</a>	42230	930	8	246	mitochondrial ribosomal protein L48 (Mrpl48) alternative variant aSep08, mRNA.
<a href="#">Mrpl48</a>	<a href="#">Mrpl48.bSep08</a>	<a href="#">293149</a>	15956	776	4	53	mitochondrial ribosomal protein L48 (5.7 kD) (Mrpl48) alternative variant bSep08, complete mRNA.
<a href="#">Mrpl48</a>	<a href="#">Mrpl48.cSep08</a>	<a href="#">293149</a>	4237	633	4	26	mitochondrial ribosomal protein L48 (Mrpl48) alternative variant cSep08, mRNA.
<a href="#">Mrpl48</a>	<a href="#">Mrpl48.dSep08</a>	<a href="#">293149</a>	1665	515	2	45	mitochondrial ribosomal protein L48 (Mrpl48) alternative variant dSep08, mRNA.
<a href="#">Mrpl50</a>	<a href="#">Mrpl50.cSep08</a>	<a href="#">362517</a>	448	239	2	58	mitochondrial ribosomal protein L50 (Mrpl50) alternative variant cSep08, mRNA.
<a href="#">Mrpl51</a>	<a href="#">Mrpl51.bSep08</a>	<a href="#">297601</a>	1079	740	2	51	mitochondrial ribosomal protein L51 (5.9 kD) (Mrpl51) alternative variant bSep08, mRNA.
<a href="#">Mrpl52</a>	<a href="#">Mrpl52.aSep08</a>	<a href="#">361037</a>	3813	1584	5	122	mitochondrial ribosomal protein L52 CRA c (13.7 kD) (Mrpl52) alternative variant aSep08, complete mRNA.
<a href="#">Mrpl52</a>	<a href="#">Mrpl52.cSep08</a>	<a href="#">361037</a>	2619	1750	3	63	mitochondrial ribosomal protein L52 CRA a (7.5 kD) (Mrpl52) alternative variant cSep08, complete mRNA.
<a href="#">Mrpl52</a>	<a href="#">Mrpl52.dSep08</a>	<a href="#">361037</a>	2631	2486	3	55	mitochondrial ribosomal protein L52 CRA e (5.8 kD) (Mrpl52) alternative variant dSep08, complete mRNA.
<a href="#">Mrpl52</a>	<a href="#">Mrpl52.eSep08</a>	<a href="#">361037</a>	2607	447	4	63	mitochondrial ribosomal protein L52 CRA a (7.5 kD) (Mrpl52) alternative variant eSep08, complete mRNA.
<a href="#">Mrpl55</a>	<a href="#">Mrpl55.aSep08</a>	<a href="#">287356</a>	2798	816	4	138	mitochondrial ribosomal protein L55 (15.8 kD) (Mrpl55) alternative variant aSep08, mRNA.
<a href="#">Mrpl55</a>	<a href="#">Mrpl55.bSep08</a>	<a href="#">287356</a>	2724	536	3	127	mitochondrial ribosomal protein L55 (15.1 kD) (Mrpl55) alternative variant bSep08, mRNA.
<a href="#">Mrpl55</a>	<a href="#">Mrpl55.cSep08</a>	<a href="#">287356</a>	2803	602	3	127	mitochondrial ribosomal protein L55 (15.1 kD) (Mrpl55) alternative variant cSep08, mRNA.
<a href="#">Mrpl55</a>	<a href="#">Mrpl55.dSep08</a>	<a href="#">287356</a>	2841	620	3	127	mitochondrial ribosomal protein L55 (15.1 kD) (Mrpl55) alternative variant dSep08, mRNA.
<a href="#">Mrpl55</a>	<a href="#">Mrpl55.fSep08</a>	<a href="#">287356</a>	2785	594	3	125	mitochondrial ribosomal protein L55 (Mrpl55) alternative variant fSep08, mRNA.

<a href="#">Mrpl55</a>	<a href="#">Mrpl55.gSep08</a>	<a href="#">287356</a>	1052	768	1	54	mitochondrial ribosomal protein L55 (Mrpl55) alternative variant gSep08, mRNA.
<a href="#">Mrps2</a>	<a href="#">Mrps2.aSep08</a>	<a href="#">362094</a>	3178	2263	3	300	mitochondrial ribosomal protein S2 (33.5 kD) (Mrps2) alternative variant aSep08, complete mRNA.
<a href="#">Mrps2</a>	<a href="#">Mrps2.bSep08</a>	<a href="#">362094</a>	1826	784	3	260	mitochondrial ribosomal protein S2 (Mrps2) alternative variant bSep08, mRNA.
<a href="#">Mrps2</a>	<a href="#">Mrps2.cSep08</a>	<a href="#">362094</a>	1017	544	2	62	mitochondrial ribosomal protein S2 (Mrps2) alternative variant cSep08, mRNA.
<a href="#">Mrps7</a>	<a href="#">Mrps7.aSep08</a>	<a href="#">113958</a>	3835	1684	5	242	mitochondrial ribosomal protein S7 (28.2 kD) (Mrps7) alternative variant aSep08, mRNA.
<a href="#">Mrps9</a>	<a href="#">Mrps9.bSep08</a>	<a href="#">301371</a>	25978	685	7	185	mitochondrial ribosomal protein S9 (Mrps9) alternative variant bSep08, mRNA.
<a href="#">Mrps11</a>	<a href="#">Mrps11.bSep08</a>	<a href="#">499185</a>	1323	694	2	87	mitochondrial ribosomal protein S11 (9.8 kD) (Mrps11) alternative variant bSep08, mRNA.
<a href="#">Mrps12</a>	<a href="#">Mrps12.bSep08</a>	<a href="#">292758</a>	2895	723	3	139	mitochondrial ribosomal protein S12 (15.4 kD) (Mrps12) alternative variant bSep08, complete mRNA.
<a href="#">Mrps14</a>	<a href="#">Mrps14.aSep08</a>	<a href="#">289143</a>	5105	978	3	136	mitochondrial ribosomal protein S14 (Mrps14) alternative variant aSep08, mRNA.
<a href="#">Mrps14</a>	<a href="#">Mrps14.cSep08</a>	<a href="#">289143</a>	2154	401	2	97	mitochondrial ribosomal protein S14 (11.2 kD) (Mrps14) alternative variant cSep08, mRNA.
<a href="#">Mrps16</a>	<a href="#">Mrps16.aSep08</a>	<a href="#">688912</a>	2781	999	2	147	mitochondrial ribosomal protein S16 (Mrps16) alternative variant aSep08, mRNA.
<a href="#">Mrps17</a>	<a href="#">Mrps17.aSep08</a>	<a href="#">288621</a>	2936	692	2	127	mitochondrial ribosomal protein S17 (14.0 kD) (Mrps17) alternative variant aSep08, mRNA.
<a href="#">Mrps18a</a>	<a href="#">Mrps18a.bSep08</a>	<a href="#">301249</a>	13137	413	2	137	mitochondrial ribosomal protein S18A (Mrps18a) alternative variant bSep08, mRNA.
<a href="#">Mrps18a</a>	<a href="#">Mrps18a.cSep08</a>	<a href="#">301249</a>	16219	519	3	98	mitochondrial ribosomal protein S18A (11.5 kD) (Mrps18a) alternative variant cSep08, mRNA.
<a href="#">Mrps18b</a>	<a href="#">Mrps18b.bSep08</a>	<a href="#">294230</a>	5519	454	1	150	mitochondrial ribosomal protein S18B (Mrps18b) alternative variant bSep08, mRNA.
<a href="#">Mrps18b</a>	<a href="#">Mrps18b.cSep08</a>	<a href="#">294230</a>	6017	885		87	mitochondrial ribosomal protein S18B (Mrps18b) alternative variant cSep08, mRNA.
<a href="#">Mrps18c</a>	<a href="#">Mrps18c.bSep08</a>	<a href="#">289469</a>	5863	395	2	87	mitochondrial ribosomal protein S18C (Mrps18c) alternative variant bSep08, mRNA.
<a href="#">Mrps18c</a>	<a href="#">Mrps18c.cSep08</a>	<a href="#">289469</a>	24183	382	1	57	mitochondrial ribosomal protein S18C (Mrps18c) alternative variant cSep08, mRNA.
<a href="#">Mrps21</a>	<a href="#">Mrps21.bSep08</a>	<a href="#">689432</a>	7739	469	3	87	mitochondrial ribosomal protein S21 (10.6 kD) (Mrps21) alternative variant bSep08, mRNA.
<a href="#">Mrps23</a>	<a href="#">Mrps23.bSep08</a>	<a href="#">360594</a>	7209	726	5	172	mitochondrial ribosomal protein S23 (Mrps23) alternative variant bSep08, mRNA.
<a href="#">Mrps23</a>	<a href="#">Mrps23.cSep08</a>	<a href="#">360594</a>	8184	551	4	103	mitochondrial ribosomal protein S23 (Mrps23) alternative variant cSep08, mRNA.
<a href="#">Mrps26</a>	<a href="#">Mrps26.bSep08</a>	<a href="#">362216</a>	1503	660		108	mitochondrial ribosomal protein S26 (12.7 kD) (Mrps26) alternative variant bSep08, mRNA.
<a href="#">Mrps27</a>	<a href="#">Mrps27.aSep08</a>	<a href="#">361883</a>	68605	1904	11	425	mitochondrial ribosomal protein S27 CRA a (Mrps27) alternative variant aSep08, mRNA.

<a href="#">Mrps27</a>	<a href="#">Mrps27.cSep08</a>	<a href="#">361883</a>	2363	375	2	94	mitochondrial ribosomal protein S27 CRA g (Mrps27) alternative variant cSep08, mRNA.
<a href="#">Mrps27</a>	<a href="#">Mrps27.dSep08</a>	<a href="#">361883</a>	37456	691	4	75	mitochondrial ribosomal protein S27 CRA c (Mrps27) alternative variant dSep08, mRNA.
<a href="#">Mrps27</a>	<a href="#">Mrps27.eSep08</a>	<a href="#">361883</a>	30168	597	2	56	putative protein (6.5 kD) (Mrps27) alternative variant eSep08, mRNA.
<a href="#">Mrps27</a>	<a href="#">Mrps27.gSep08</a>	<a href="#">361883</a>	30152	379	4	50	mitochondrial ribosomal protein S27 CRA g (Mrps27) alternative variant gSep08, mRNA.
<a href="#">Mrps28</a>	<a href="#">Mrps28.bSep08</a>	<a href="#">689025</a>	35675	407	1	135	mitochondrial ribosomal protein S28 (Mrps28) alternative variant bSep08, mRNA.
<a href="#">Mrps28</a>	<a href="#">Mrps28.cSep08</a>	<a href="#">689025</a>	37621	376	2	71	mitochondrial ribosomal protein S28 (Mrps28) alternative variant cSep08, mRNA.
<a href="#">Mrps30</a>	<a href="#">Mrps30.bSep08</a>	<a href="#">294767</a>	6488	1148	6	351	mitochondrial ribosomal protein S30 (Mrps30) alternative variant bSep08, mRNA.
<a href="#">Mrps30</a>	<a href="#">Mrps30.cSep08</a>	<a href="#">294767</a>	2636	729	2	209	mitochondrial ribosomal protein S30 (22.9 kD) (Mrps30) alternative variant cSep08, mRNA.
<a href="#">Mrps30</a>	<a href="#">Mrps30.eSep08</a>	<a href="#">294767</a>	3064	1042	2	108	mitochondrial ribosomal protein S30 CRA b (12.5 kD) (Mrps30) alternative variant eSep08, mRNA.
<a href="#">Mrps31</a>	<a href="#">Mrps31.bSep08</a>	<a href="#">290850</a>	1828	980	2	96	mitochondrial ribosomal protein S31 (11.5 kD) (Mrps31) alternative variant bSep08, mRNA.
<a href="#">Mrps31</a>	<a href="#">Mrps31.cSep08</a>	<a href="#">290850</a>	3550	526	2	75	putative protein (Mrps31) alternative variant cSep08, mRNA.
<a href="#">Mrps31</a>	<a href="#">Mrps31.dSep08</a>	<a href="#">290850</a>	3078	571	2	74	mitochondrial ribosomal protein S31 like (Mrps31) alternative variant dSep08, mRNA.
<a href="#">Mrps34andNme3</a>	<a href="#">Mrps34andNme3.cSep08</a>	<a href="#">85269</a>	1011	772	4	162	non-metastatic cells like (18.2 kD) (Mrps34andNme3) alternative variant cSep08, mRNA.
<a href="#">Mrps34andNme3</a>	<a href="#">Mrps34andNme3.cSep08</a>	<a href="#">287126</a>	1011	772	4	162	non-metastatic cells like (18.2 kD) (Mrps34andNme3) alternative variant cSep08, mRNA.
<a href="#">Mrps34andNme3</a>	<a href="#">Mrps34andNme3.eSep08</a>	<a href="#">85269</a>	962	537	4	91	nucleoside diphosphate (Mrps34andNme3) alternative variant eSep08, mRNA.
<a href="#">Mrps34andNme3</a>	<a href="#">Mrps34andNme3.eSep08</a>	<a href="#">287126</a>	962	537	4	91	nucleoside diphosphate (Mrps34andNme3) alternative variant eSep08, mRNA.
<a href="#">Mrps35</a>	<a href="#">Mrps35.bSep08</a>	<a href="#">297727</a>	29425	813	5	266	mitochondrial ribosomal protein S35 (Mrps35) alternative variant bSep08, mRNA.
<a href="#">Mrps35</a>	<a href="#">Mrps35.cSep08</a>	<a href="#">297727</a>	15281	464	2	97	mitochondrial ribosomal protein S35 (Mrps35) alternative variant cSep08, mRNA.
<a href="#">Mrps36</a>	<a href="#">Mrps36.aSep08</a>	<a href="#">294696</a>	7755	892	2	117	mitochondrial ribosomal protein S36 (Mrps36) alternative variant aSep08, mRNA.
<a href="#">Mrps36</a>	<a href="#">Mrps36.bSep08</a>	<a href="#">294696</a>	7480	429	1	58	mitochondrial ribosomal protein S36 (6.2 kD) (Mrps36) alternative variant bSep08, complete mRNA.
<a href="#">Mrpf</a>	<a href="#">Mrpf.aSep08</a>	<a href="#">311903</a>	16994	1798	5	535	mitochondrial ribosome recycling factor (Mrpf) alternative variant aSep08, complete mRNA.
<a href="#">Mrs2</a>	<a href="#">Mrs2.bSep08</a>	<a href="#">79032</a>	17829	2312	2	450	MRS2 magnesium homeostasis factor homolog (S. cerevisiae) (51.1 kD) (Mrs2) alternative variant bSep08, complete mRNA.

<a href="#">Mrs2</a>	<a href="#">Mrs2.cSep08</a>	<a href="#">79032</a>	19207	3246	4	200	MRS2 magnesium homeostasis factor homolog (S. cerevisiae) (22.8 kD) (Mrs2) alternative variant cSep08, complete mRNA.
<a href="#">Mrto4</a>	<a href="#">Mrto4.aSep08</a>	<a href="#">298586</a>	6819	2518	2	240	MRT4, mRNA turnover 4, homolog (S. cerevisiae) (27.6 kD) (Mrto4) alternative variant aSep08, mRNA.
<a href="#">Ms4a1</a>	<a href="#">Ms4a1.bSep08</a>	<a href="#">309217</a>	9390	1084	5	251	membrane-spanning 4-domains, subfamily A, member 1 (Ms4a1) alternative variant bSep08, mRNA.
<a href="#">Ms4a1</a>	<a href="#">Ms4a1.cSep08</a>	<a href="#">309217</a>	5095	1159	4	199	membrane-spanning 4-domains, subfamily A, member 1 (Ms4a1) alternative variant cSep08, mRNA.
<a href="#">Ms4a3</a>	<a href="#">Ms4a3.aSep08</a>	<a href="#">293753</a>	15952	729		185	membrane-spanning 4-domains, subfamily A, member 3 (Ms4a3) mRNA.
<a href="#">Ms4a4a</a>	<a href="#">Ms4a4a.aSep08</a>	<a href="#">361734</a>	21951	1098		236	membrane-spanning 4-domains, subfamily A, member 4A (25.3 kD) (Ms4a4a) alternative variant aSep08, mRNA.
<a href="#">Ms4a6a</a>	<a href="#">Ms4a6a.aSep08</a>	<a href="#">293750</a>	3089	534		60	membrane-spanning 4-domains, subfamily A, member 6A (6.6 kD) (Ms4a6a) mRNA.
<a href="#">Ms4a7</a>	<a href="#">Ms4a7.bSep08</a>	<a href="#">293744</a>	16188	978	5	225	membrane-spanning 4-domains, subfamily A, member 7 (24.5 kD) (Ms4a7) alternative variant bSep08, mRNA.
<a href="#">Ms4a7</a>	<a href="#">Ms4a7.cSep08</a>	<a href="#">293744</a>	3064	514	2	156	membrane-spanning 4-domains, subfamily A, member 7 (Ms4a7) alternative variant cSep08, mRNA.
<a href="#">Ms4a8a</a>	<a href="#">Ms4a8a.bSep08</a>	<a href="#">361733</a>	12124	823	4	148	membrane-spanning 4-domains, subfamily A, member 8A (Ms4a8a) alternative variant bSep08, mRNA.
<a href="#">Ms4a10</a>	<a href="#">Ms4a10.aSep08</a>	<a href="#">293739</a>	13130	886	8	259	membrane-spanning 4-domains, subfamily A, member 10 (Ms4a10) alternative variant aSep08, mRNA.
<a href="#">Ms4a11</a>	<a href="#">Ms4a11.aSep08</a>	<a href="#">361735</a>	12225	1374	5	245	membrane-spanning 4-domains, subfamily A, member 11 (26.7 kD) (Ms4a11) alternative variant aSep08, mRNA.
<a href="#">Ms4a11</a>	<a href="#">Ms4a11.bSep08</a>	<a href="#">361735</a>	3152	607	1	113	membrane-spanning 4-domains, subfamily A, member 11 (12.3 kD) (Ms4a11) alternative variant bSep08, mRNA.
<a href="#">Msc</a>	<a href="#">Msc.bSep08</a>	<a href="#">312897</a>	973	290	1	32	musculin (Msc) alternative variant bSep08, mRNA.
<a href="#">Msc</a>	<a href="#">Msc.dSep08</a>	<a href="#">312897</a>	1458	483	2	79	musculin (Msc) alternative variant dSep08, mRNA.
<a href="#">Msh2</a>	<a href="#">Msh2.bSep08</a>	<a href="#">81709</a>	4347	944	1	39	mutS homolog 2 (E. coli) (4.7 kD) (Msh2) alternative variant bSep08, mRNA.
<a href="#">Msh3</a>	<a href="#">Msh3.aSep08</a>	<a href="#">499505</a>	44703	1414	6	316	mutS homolog 3 (E. coli) (Msh3) alternative variant aSep08, mRNA.
<a href="#">Msh3</a>	<a href="#">Msh3.bSep08</a>	<a href="#">499505</a>	21730	784	4	209	mutS homolog 3 (E. coli) (Msh3) alternative variant bSep08, mRNA.
<a href="#">Msh3</a>	<a href="#">Msh3.cSep08</a>	<a href="#">499505</a>	16094	708	2	194	mutS homolog 3 (E. coli) (Msh3) alternative variant cSep08, mRNA.
<a href="#">Msh5</a>	<a href="#">Msh5.bSep08</a>	<a href="#">294252</a>	3614	1264	8	244	mutS homolog 5 (E. coli) (Msh5) alternative variant bSep08, mRNA.
<a href="#">Msh5</a>	<a href="#">Msh5.cSep08</a>	<a href="#">294252</a>	9662	688	9	229	mutS homolog 5 (E. coli) (Msh5) alternative variant cSep08, mRNA.
<a href="#">Msh5</a>	<a href="#">Msh5.dSep08</a>	<a href="#">294252</a>	2097	1197	4	95	mutS homolog 5 (E. coli) (Msh5) alternative variant dSep08, mRNA.
<a href="#">Msi1</a>	<a href="#">Msi1.bSep08</a>	<a href="#">259272</a>	22353	2656	2	264	musashi homolog 1 (Drosophila) (Msi1) alternative variant bSep08, mRNA.
<a href="#">Msi2</a>	<a href="#">Msi2.aSep08</a>	<a href="#">360596</a>	72963	707	5	178	musashi homolog 2 (Drosophila) (Msi2) alternative variant aSep08, mRNA.

<a href="#">Msi2</a>	<a href="#">Msi2.bSep08</a>	<a href="#">360596</a>	72961	751	6	177	musashi homolog 2 (Drosophila) (Msi2) alternative variant bSep08, mRNA.
<a href="#">Msi2</a>	<a href="#">Msi2.cSep08</a>	<a href="#">360596</a>	39282	3280	5	129	musashi homolog 2 (Drosophila) (Msi2) alternative variant cSep08, mRNA.
<a href="#">Msi2</a>	<a href="#">Msi2.dSep08</a>	<a href="#">360596</a>	36532	248	3	66	musashi homolog 2 (Drosophila) (Msi2) alternative variant dSep08, mRNA.
<a href="#">Msi2</a>	<a href="#">Msi2.eSep08</a>	<a href="#">360596</a>	50287	380	3	33	musashi homolog 2 (Drosophila) (Msi2) alternative variant eSep08, mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.bSep08</a>	<a href="#">317464</a>	17682	1196	9	291	male-specific lethal-3 homolog 1 (Drosophila) (33.6 kD) (Msl31) alternative variant bSep08, complete mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.cSep08</a>	<a href="#">317464</a>	3608	809	6	209	male-specific lethal-3 homolog 1 (Drosophila) (Msl31) alternative variant cSep08, mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.dSep08</a>	<a href="#">317464</a>	3382	449	5	149	male-specific lethal-3 homolog 1 (Drosophila) (Msl31) alternative variant dSep08, mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.eSep08</a>	<a href="#">317464</a>	1670	350	4	116	male-specific lethal-3 homolog 1 (Drosophila) (Msl31) alternative variant eSep08, mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.fSep08</a>	<a href="#">317464</a>	7902	743	6	114	male-specific lethal-3 homolog 1 (Drosophila) (13.1 kD) (Msl31) alternative variant fSep08, mRNA.
<a href="#">Msl31</a>	<a href="#">Msl31.hSep08</a>	<a href="#">317464</a>	2116	327	3	108	male-specific lethal-3 homolog 1 (Drosophila) (Msl31) alternative variant hSep08, mRNA.
<a href="#">Msln</a>	<a href="#">Msln.bSep08</a>	<a href="#">60333</a>	5695	405	2	117	mesothelin (Msln) alternative variant bSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.cSep08</a>	<a href="#">29311</a>	38503	771	7	238	nuclear receptor coactivator 4 (MsmbandNcoa4) alternative variant cSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.cSep08</a>	<a href="#">619385</a>	38503	771	7	238	nuclear receptor coactivator 4 (MsmbandNcoa4) alternative variant cSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.dSep08</a>	<a href="#">29311</a>	36908	673	7	223	nuclear receptor coactivator 4 (MsmbandNcoa4) alternative variant dSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.dSep08</a>	<a href="#">619385</a>	36908	673	7	223	nuclear receptor coactivator 4 (MsmbandNcoa4) alternative variant dSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.eSep08</a>	<a href="#">29311</a>	7073	793	4	132	nuclear receptor coactivator 4 (15.3 kD) (MsmbandNcoa4) alternative variant eSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.eSep08</a>	<a href="#">619385</a>	7073	793	4	132	nuclear receptor coactivator 4 (15.3 kD) (MsmbandNcoa4) alternative variant eSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.gSep08</a>	<a href="#">29311</a>	8355	742	3	84	beta-microseminoprotein (MsmbandNcoa4) alternative variant gSep08, mRNA.
<a href="#">MsmbandNcoa4</a>	<a href="#">MsmbandNcoa4.gSep08</a>	<a href="#">619385</a>	8355	742	3	84	beta-microseminoprotein (MsmbandNcoa4) alternative variant gSep08, mRNA.
<a href="#">Msn</a>	<a href="#">Msn.bSep08</a>	<a href="#">81521</a>	7412	672	4	158	moesin (Msn) alternative variant bSep08, mRNA.
<a href="#">Msr1</a>	<a href="#">Msr1.aSep08</a>	<a href="#">498638</a>	10424	799		77	macrophage scavenger receptor 1 (Msr1) mRNA.
<a href="#">Msra</a>	<a href="#">Msra.bSep08</a>	<a href="#">29447</a>	227945	581	2	193	methionine sulfoxide reductase A (Msra) alternative variant bSep08, mRNA.
<a href="#">Msra</a>	<a href="#">Msra.cSep08</a>	<a href="#">29447</a>	149822	432	1	144	methionine sulfoxide reductase A (Msra) alternative variant cSep08, mRNA.
<a href="#">Msrb2</a>	<a href="#">Msrb2.bSep08</a>	<a href="#">361286</a>	25649	856		146	methionine sulfoxide reductase B2 (Msrb2) alternative variant bSep08, mRNA.
<a href="#">Mst1</a>	<a href="#">Mst1.bSep08</a>	<a href="#">24566</a>	1908	736	4	135	macrophage stimulating 1 (hepatocyte growth factor-like) (Mst1) alternative variant bSep08, mRNA.

<a href="#">Mst4</a>	<a href="#">Mst4.aSep08</a>	<a href="#">317589</a>	44402	1002		334	serine/threonine protein kinase MST4 (Mst4) mRNA.
<a href="#">Msto1</a>	<a href="#">Msto1.bSep08</a>	<a href="#">295237</a>	730	337	3	112	misato homolog 1 (Drosophila) (Msto1) alternative variant bSep08, mRNA.
<a href="#">Mt1a</a>	<a href="#">Mt1a.bSep08</a>	<a href="#">24567</a>	1148	982	2	85	metallothionein 1a (8.5 kD) (Mt1a) alternative variant bSep08, mRNA.
<a href="#">Mt1a</a>	<a href="#">Mt1a.cSep08</a>	<a href="#">24567</a>	1006	544	2	67	metallothionein 1a (Mt1a) alternative variant cSep08, mRNA.
<a href="#">Mt2A</a>	<a href="#">Mt2A.aSep08</a>	<a href="#">689415</a>	774	382	3	61	metallothionein 2A (Mt2A) alternative variant aSep08, complete mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.bSep08</a>	<a href="#">64520</a>	2989	1794	3	536	metastasis associated 1 (Mta1) alternative variant bSep08, mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.cSep08</a>	<a href="#">64520</a>	17373	1783	14	361	metastasis associated 1 (Mta1) alternative variant cSep08, mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.dSep08</a>	<a href="#">64520</a>	4391	1158	8	280	metastasis associated 1 (Mta1) alternative variant dSep08, mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.eSep08</a>	<a href="#">64520</a>	11478	551	6	183	metastasis associated 1 (Mta1) alternative variant eSep08, mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.gSep08</a>	<a href="#">64520</a>	3765	944	5	134	metastasis associated 1 (Mta1) alternative variant gSep08, mRNA.
<a href="#">Mta1</a>	<a href="#">Mta1.hSep08</a>	<a href="#">64520</a>	1296	894	3	119	metastasis associated 1 (Mta1) alternative variant hSep08, mRNA.
<a href="#">Mta3</a>	<a href="#">Mta3.aSep08</a>	<a href="#">298763</a>	102173	1582	3	293	metastasis associated 1 family, member 3 (Mta3) alternative variant aSep08, mRNA.
<a href="#">Mta3</a>	<a href="#">Mta3.bSep08</a>	<a href="#">298763</a>	45950	1195	2	276	metastasis associated 1 family, member 3 (31.4 kD) (Mta3) alternative variant bSep08, mRNA.
<a href="#">Mtap</a>	<a href="#">Mtap.bSep08</a>	<a href="#">298227</a>	7797	1236	1	167	methylthioadenosine phosphorylase (Mtap) alternative variant bSep08, mRNA.
<a href="#">Mtch1</a>	<a href="#">Mtch1.bSep08</a>	<a href="#">294313</a>	22866	1444	12	260	mitochondrial carrier homolog 1 (29.0 kD) (Mtch1) alternative variant bSep08, mRNA.
<a href="#">Mtch1</a>	<a href="#">Mtch1.cSep08</a>	<a href="#">294313</a>	15004	755	7	164	mitochondrial carrier homolog 1 (Mtch1) alternative variant cSep08, mRNA.
<a href="#">Mtch1</a>	<a href="#">Mtch1.dSep08</a>	<a href="#">294313</a>	14980	646	7	129	mitochondrial carrier homolog 1 (Mtch1) alternative variant dSep08, mRNA.
<a href="#">Mtch2</a>	<a href="#">Mtch2.bSep08</a>	<a href="#">295922</a>	15293	694	2	227	mitochondrial carrier homolog 2 (Mtch2) alternative variant bSep08, mRNA.
<a href="#">Mtch2</a>	<a href="#">Mtch2.cSep08</a>	<a href="#">295922</a>	17187	738	2	225	mitochondrial carrier homolog 2 (Mtch2) alternative variant cSep08, mRNA.
<a href="#">Mtcp1</a>	<a href="#">Mtcp1.aSep08</a>	<a href="#">498814</a>	3931	898	5	107	mature T-cell proliferation 1 and similar to P8 MTCP-1 protein (Mature T-cell proliferation-1 type A) (MTCP-1 type A) (P8MTCP1) (12.9 kD) (Mtcp1) alternative variant aSep08, mRNA.
<a href="#">Mtcp1</a>	<a href="#">Mtcp1.aSep08</a>	<a href="#">679930</a>	3931	898	5	107	mature T-cell proliferation 1 and similar to P8 MTCP-1 protein (Mature T-cell proliferation-1 type A) (MTCP-1 type A) (P8MTCP1) (12.9 kD) (Mtcp1) alternative variant aSep08, mRNA.



<a href="#">Mtcp1</a>	<a href="#">Mtcp1.bSep08</a>	<a href="#">498814</a>	632	426	2	69	mature T-cell proliferation 1 and similar to P8 MTCP-1 protein (Mature T-cell proliferation-1 type A) (MTCP-1 type A) (P8MTCP1) (Mtcp1) alternative variant bSep08, mRNA.
<a href="#">Mtcp1</a>	<a href="#">Mtcp1.bSep08</a>	<a href="#">679930</a>	632	426	2	69	mature T-cell proliferation 1 and similar to P8 MTCP-1 protein (Mature T-cell proliferation-1 type A) (MTCP-1 type A) (P8MTCP1) (Mtcp1) alternative variant bSep08, mRNA.
<a href="#">Mtdh</a>	<a href="#">Mtdh.bSep08</a>	<a href="#">170910</a>	30695	4048	9	393	metadherin (Mtdh) alternative variant bSep08, mRNA.
<a href="#">Mtdh</a>	<a href="#">Mtdh.cSep08</a>	<a href="#">170910</a>	20820	957	8	319	metadherin (Mtdh) alternative variant cSep08, mRNA.
<a href="#">Mtdh</a>	<a href="#">Mtdh.dSep08</a>	<a href="#">170910</a>	48403	950	6	199	metadherin (Mtdh) alternative variant dSep08, mRNA.
<a href="#">Mtdh</a>	<a href="#">Mtdh.eSep08</a>	<a href="#">170910</a>	16780	684	5	120	metadherin (Mtdh) alternative variant eSep08, mRNA.
<a href="#">Mterf</a>	<a href="#">Mterf.bSep08</a>	<a href="#">85261</a>	6468	844	2	240	mitochondrial transcription termination factor (Mterf) alternative variant bSep08, mRNA.
<a href="#">Mterf</a>	<a href="#">Mterf.cSep08</a>	<a href="#">85261</a>	6347	772	3	39	putative protein (Mterf) alternative variant cSep08, mRNA.
<a href="#">Mterf</a>	<a href="#">Mterf.dSep08</a>	<a href="#">85261</a>	6323	745	3	194	mitochondrial transcription termination factor (Mterf) alternative variant dSep08, mRNA.
<a href="#">Mterfd1</a>	<a href="#">Mterfd1.bSep08</a>	<a href="#">299514</a>	11031	760	5	161	putative protein of mammalian origin (17.7 kD) (Mterfd1) alternative variant bSep08, mRNA.
<a href="#">Mterfd1</a>	<a href="#">Mterfd1.cSep08</a>	<a href="#">299514</a>	18255	1086	6	155	putative protein of mammalian origin (17.1 kD) (Mterfd1) alternative variant cSep08, complete mRNA.
<a href="#">Mterfd1</a>	<a href="#">Mterfd1.dSep08</a>	<a href="#">299514</a>	2971	856	2	103	putative cytoplasmic protein of metazoan origin (12.1 kD) (Mterfd1) alternative variant dSep08, mRNA.
<a href="#">Mterfd1</a>	<a href="#">Mterfd1.eSep08</a>	<a href="#">299514</a>	2691	734	2	99	putative protein of mammalian origin (Mterfd1) alternative variant eSep08, mRNA.
<a href="#">Mterfd1</a>	<a href="#">Mterfd1.gSep08</a>	<a href="#">299514</a>	13516	679	6	87	putative protein of mammalian origin (9.5 kD) (Mterfd1) alternative variant gSep08, complete mRNA.
<a href="#">Mterfd3</a>	<a href="#">Mterfd3.cSep08</a>	<a href="#">366856</a>	5930	993	4	34	putative protein (Mterfd3) alternative variant cSep08, mRNA.
<a href="#">Mtf2</a>	<a href="#">Mtf2.aSep08</a>	<a href="#">360905</a>	43300	2784	15	593	metal response element binding transcription factor 2 (66.9 kD) (Mtf2) alternative variant aSep08, complete mRNA.
<a href="#">Mtf2</a>	<a href="#">Mtf2.bSep08</a>	<a href="#">360905</a>	16126	968	9	280	metal response element binding transcription factor 2 (Mtf2) alternative variant bSep08, mRNA.
<a href="#">Mtf2</a>	<a href="#">Mtf2.cSep08</a>	<a href="#">360905</a>	21066	863	6	129	metal response element binding transcription factor 2 (Mtf2) alternative variant cSep08, mRNA.
<a href="#">Mtf2</a>	<a href="#">Mtf2.dSep08</a>	<a href="#">360905</a>	3794	1033	2	30	metal response element binding transcription factor 2 (Mtf2) alternative variant dSep08, mRNA.
<a href="#">Mtfmt</a>	<a href="#">Mtfmt.bSep08</a>	<a href="#">315763</a>	3941	624	2	56	mitochondrial methionyl-tRNA formyltransferase (6.5 kD) (Mtfmt) alternative variant bSep08, mRNA.
<a href="#">Mtfr1</a>	<a href="#">Mtfr1.bSep08</a>	<a href="#">311403</a>	11759	726	5	227	mitochondrial fission regulator 1 (Mtfr1) alternative variant bSep08, mRNA.
<a href="#">Mthfd1l</a>	<a href="#">Mthfd1l.aSep08</a>	<a href="#">361472</a>	193421	3568	23	1021	methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1-like (Mthfd1l) alternative variant aSep08, mRNA.
<a href="#">Mthfd2l</a>	<a href="#">Mthfd2l.bSep08</a>	<a href="#">305248</a>	53671	457	1	90	methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2-like (Mthfd2l) alternative variant bSep08, mRNA.
<a href="#">Mthfr</a>	<a href="#">Mthfr.aSep08</a>	<a href="#">362657</a>	10815	1594	7	530	5,10-methylenetetrahydrofolate reductase (NADPH) (Mthfr) alternative variant aSep08, mRNA.

<a href="#">Mthfr</a>	<a href="#">Mthfr.bSep08</a>	<a href="#">362657</a>	5172	1784	5	434	5,10-methylenetetrahydrofolate reductase (NADPH) (Mthfr) alternative variant bSep08, mRNA.
<a href="#">Mthfr</a>	<a href="#">Mthfr.cSep08</a>	<a href="#">362657</a>	1469	384	1	66	5,10-methylenetetrahydrofolate reductase (NADPH) (Mthfr) alternative variant cSep08, mRNA.
<a href="#">Mtif2</a>	<a href="#">Mtif2.bSep08</a>	<a href="#">305606</a>	12951	1268	9	341	mitochondrial translational initiation factor 2 (37.8 kD) (Mtif2) alternative variant bSep08, mRNA.
<a href="#">Mtl5</a>	<a href="#">Mtl5.bSep08</a>	<a href="#">309142</a>	3244	298	1	87	metallothionein-like 5, testis-specific (tesmin) (Mtl5) alternative variant bSep08, mRNA.
<a href="#">Mtm1</a>	<a href="#">Mtm1.bSep08</a>	<a href="#">288762</a>	3971	1655	5	243	X-linked myotubular myopathy gene 1 (Mtm1) alternative variant bSep08, mRNA.
<a href="#">Mtm1</a>	<a href="#">Mtm1.cSep08</a>	<a href="#">288762</a>	3093	452	2	137	X-linked myotubular myopathy gene 1 (Mtm1) alternative variant cSep08, mRNA.
<a href="#">Mtmr1</a>	<a href="#">Mtmr1.aSep08</a>	<a href="#">317296</a>	18912	1792	5	565	myotubularin related protein 1 (Mtmr1) alternative variant aSep08, mRNA.
<a href="#">Mtmr1</a>	<a href="#">Mtmr1.bSep08</a>	<a href="#">317296</a>	17594	2943	11	515	myotubularin related protein 1 (Mtmr1) alternative variant bSep08, mRNA.
<a href="#">Mtmr1</a>	<a href="#">Mtmr1.cSep08</a>	<a href="#">317296</a>	3356	914	2	87	myotubularin related protein 1 (8.9 kD) (Mtmr1) alternative variant cSep08, complete mRNA.
<a href="#">Mtmr1</a>	<a href="#">Mtmr1.dSep08</a>	<a href="#">317296</a>	2998	579	2	80	myotubularin related protein 1 (8.1 kD) (Mtmr1) alternative variant dSep08, complete mRNA.
<a href="#">Mtmr2</a>	<a href="#">Mtmr2.bSep08</a>	<a href="#">315422</a>	4188	659	4	112	myotubularin related protein 2 (Mtmr2) alternative variant bSep08, mRNA.
<a href="#">Mtmr2</a>	<a href="#">Mtmr2.dSep08</a>	<a href="#">315422</a>	3533	396	2	60	myotubularin related protein 2 (Mtmr2) alternative variant dSep08, mRNA.
<a href="#">Mtmr2</a>	<a href="#">Mtmr2.eSep08</a>	<a href="#">315422</a>	2797	1284	2	57	myotubularin related protein 2 (6.4 kD) (Mtmr2) alternative variant eSep08, mRNA.
<a href="#">Mtmr2</a>	<a href="#">Mtmr2.fSep08</a>	<a href="#">315422</a>	27659	371	4	47	myotubularin related protein 2 (Mtmr2) alternative variant fSep08, mRNA.
<a href="#">Mtmr3</a>	<a href="#">Mtmr3.bSep08</a>	<a href="#">305482</a>	6749	2688	4	261	myotubularin related protein 3 (Mtmr3) alternative variant bSep08, mRNA.
<a href="#">Mtmr3</a>	<a href="#">Mtmr3.dSep08</a>	<a href="#">305482</a>	23719	477	6	97	myotubularin-related protein 3 (Mtmr3) alternative variant dSep08, mRNA.
<a href="#">Mtmr3</a>	<a href="#">Mtmr3.eSep08</a>	<a href="#">305482</a>	49352	1536	3	84	putative protein (Mtmr3) alternative variant eSep08, mRNA.
<a href="#">Mtmr3</a>	<a href="#">Mtmr3.gSep08</a>	<a href="#">305482</a>	6292	1358	2	72	ab2-143 like (8.0 kD) (Mtmr3) alternative variant gSep08, mRNA.
<a href="#">Mtmr4</a>	<a href="#">Mtmr4.bSep08</a>	<a href="#">287607</a>	2881	451	4	149	myotubularin related protein 4 and hypothetical protein LOC689785 (Mtmr4) alternative variant bSep08, mRNA.
<a href="#">Mtmr4</a>	<a href="#">Mtmr4.bSep08</a>	<a href="#">689785</a>	2881	451	4	149	myotubularin related protein 4 and hypothetical protein LOC689785 (Mtmr4) alternative variant bSep08, mRNA.
<a href="#">Mtmr6</a>	<a href="#">Mtmr6.bSep08</a>	<a href="#">305935</a>	9549	971	1	323	myotubularin related protein 6 (Mtmr6) alternative variant bSep08, mRNA.
<a href="#">Mtmr7</a>	<a href="#">Mtmr7.bSep08</a>	<a href="#">306490</a>	12522	932	7	310	myotubularin related protein 7 (Mtmr7) alternative variant bSep08, mRNA.
<a href="#">Mtmr7</a>	<a href="#">Mtmr7.cSep08</a>	<a href="#">306490</a>	49626	571	5	169	myotubularin related protein 7 (Mtmr7) alternative variant cSep08, mRNA.
<a href="#">Mtmr7</a>	<a href="#">Mtmr7.dSep08</a>	<a href="#">306490</a>	3573	777	3	97	myotubularin related protein 7 (Mtmr7) alternative variant dSep08, mRNA.

<a href="#">Mtmr10</a>	<a href="#">Mtmr10.aSep08</a>	<a href="#">309255</a>	51444	4649		776	myotubularin related protein 10 (Mtmr10) mRNA.
<a href="#">Mtmr11</a>	<a href="#">Mtmr11.aSep08</a>	<a href="#">689613</a>	5472	1268	12	387	myotubularin related protein 11 (Mtmr11) alternative variant aSep08, mRNA.
<a href="#">Mtmr11</a>	<a href="#">Mtmr11.bSep08</a>	<a href="#">689613</a>	3417	1922	4	270	myotubularin related protein 11 CRA f (30.7 kD) (Mtmr11) alternative variant bSep08, mRNA.
<a href="#">Mtmr11</a>	<a href="#">Mtmr11.cSep08</a>	<a href="#">689613</a>	4914	1076	10	207	myotubularin related protein 11 (Mtmr11) alternative variant cSep08, mRNA.
<a href="#">Mtmr11</a>	<a href="#">Mtmr11.dSep08</a>	<a href="#">689613</a>	5682	1500	8	119	myotubularin related protein 11 CRA f (Mtmr11) alternative variant dSep08, mRNA.
<a href="#">Mtmr12</a>	<a href="#">Mtmr12.bSep08</a>	<a href="#">310155</a>	57580	2634	11	223	myotubularin related protein 12 (25.8 kD) (Mtmr12) alternative variant bSep08, mRNA.
<a href="#">Mtmr14</a>	<a href="#">Mtmr14.bSep08</a>	<a href="#">312634</a>	11120	575	7	128	myotubularin related protein 14 (Mtmr14) alternative variant bSep08, mRNA.
<a href="#">Mtmr14</a>	<a href="#">Mtmr14.dSep08</a>	<a href="#">312634</a>	1097	463	2	71	myotubularin related protein 14 (Mtmr14) alternative variant dSep08, mRNA.
<a href="#">Mtmr15</a>	<a href="#">Mtmr15.aSep08</a>	<a href="#">309256</a>	4132	1179		155	myotubularin related protein 15 (Mtmr15) mRNA.
<a href="#">Mtp18</a>	<a href="#">Mtp18.bSep08</a>	<a href="#">289745</a>	3233	789	2	157	mitochondrial protein 18 kDa (17.0 kD) (Mtp18) alternative variant bSep08, mRNA.
<a href="#">Mtrf1l</a>	<a href="#">Mtrf1l.bSep08</a>	<a href="#">361473</a>	9441	894	6	143	mitochondrial translational release factor 1-like (15.9 kD) (Mtrf1l) alternative variant bSep08, mRNA.
<a href="#">Mtrf1l</a>	<a href="#">Mtrf1l.cSep08</a>	<a href="#">361473</a>	3327	421	2	135	mitochondrial translational release factor 1-like (Mtrf1l) alternative variant cSep08, mRNA.
<a href="#">Mtrf1l</a>	<a href="#">Mtrf1l.eSep08</a>	<a href="#">361473</a>	1146	544	2	93	mitochondrial translational release factor 1-like (11.1 kD) (Mtrf1l) alternative variant eSep08, mRNA.
<a href="#">Mtrf1l</a>	<a href="#">Mtrf1l.fSep08</a>	<a href="#">361473</a>	1841	395	2	87	mitochondrial translational release factor 1-like (Mtrf1l) alternative variant fSep08, mRNA.
<a href="#">Mtrr</a>	<a href="#">Mtrr.bSep08</a>	<a href="#">290947</a>	9815	971	7	323	5-methyltetrahydrofolate-homocysteine methyltransferase reductase (Mtrr) alternative variant bSep08, mRNA.
<a href="#">Mtrr</a>	<a href="#">Mtrr.cSep08</a>	<a href="#">290947</a>	17139	660	5	219	5-methyltetrahydrofolate-homocysteine methyltransferase reductase (Mtrr) alternative variant cSep08, mRNA.
<a href="#">Mtrr</a>	<a href="#">Mtrr.dSep08</a>	<a href="#">290947</a>	8595	837	5	95	5-methyltetrahydrofolate-homocysteine methyltransferase reductase (10.6 kD) (Mtrr) alternative variant dSep08, mRNA.
<a href="#">Mtss1</a>	<a href="#">Mtss1.bSep08</a>	<a href="#">362918</a>	2005	867	2	263	metastasis suppressor 1 (Mtss1) alternative variant bSep08, mRNA.
<a href="#">Mtus1</a>	<a href="#">Mtus1.aSep08</a>	<a href="#">306487</a>	142806	4317	14	1210	mitochondrial tumor suppressor 1 (134.2 kD) (Mtus1) alternative variant aSep08, mRNA.
<a href="#">Mtus1</a>	<a href="#">Mtus1.cSep08</a>	<a href="#">306487</a>	17315	751	6	182	mitochondrial tumor suppressor 1 (Mtus1) alternative variant cSep08, mRNA.
<a href="#">Mtvr2</a>	<a href="#">Mtvr2.cSep08</a>	<a href="#">309170</a>	1608	1253	2	118	mammary tumor virus receptor 2 (11.8 kD) (Mtvr2) alternative variant cSep08, mRNA.
<a href="#">Mtx2</a>	<a href="#">Mtx2.bSep08</a>	<a href="#">288150</a>	32226	967	7	177	metaxin 2 (Mtx2) alternative variant bSep08, mRNA.
<a href="#">muby</a>	<a href="#">muby.aSep08</a>		1152	423		36	putative protein (4.4 kD) (muby) mRNA.
<a href="#">Muc4</a>	<a href="#">Muc4.aSep08</a>	<a href="#">303887</a>	2322	1007	4	215	mucin 4, cell surface associated (Muc4) alternative variant aSep08, mRNA.
<a href="#">Muc13</a>	<a href="#">Muc13.aSep08</a>	<a href="#">207126</a>	22252	2223	1	380	mucin 13, cell surface associated (Muc13) alternative variant aSep08, mRNA.

<a href="#">Muc13</a>	<a href="#">Muc13.bSep08</a>	<a href="#">207126</a>	2618	401	2	48	mucin 13, cell surface associated (Muc13) alternative variant bSep08, mRNA.
<a href="#">Muc15</a>	<a href="#">Muc15.bSep08</a>	<a href="#">690914</a>	6770	724	3	75	mucin 15, cell surface associated (8.9 kD) (Muc15) alternative variant bSep08, mRNA.
<a href="#">Muc16</a>	<a href="#">Muc16.aSep08</a>	<a href="#">315451</a>	27077	1360		431	mucin 16, cell surface associated (Muc16) mRNA.
<a href="#">Muc19</a>	<a href="#">Muc19.aSep08</a>	<a href="#">315239</a>	6422	709		181	mucin 19 (Muc19) mRNA.
<a href="#">Mucdhl</a>	<a href="#">Mucdhl.bSep08</a>	<a href="#">171554</a>	3654	981	5	304	mucin and cadherin like (Mucdhl) alternative variant bSep08, mRNA.
<a href="#">Mucdhl</a>	<a href="#">Mucdhl.cSep08</a>	<a href="#">171554</a>	5022	300	3	99	mucin and cadherin like (Mucdhl) alternative variant cSep08, mRNA.
<a href="#">muchy</a>	<a href="#">muchy.aSep08</a>		1093	303		96	putative protein (muchy) mRNA.
<a href="#">mudar</a>	<a href="#">mudar.aSep08</a>		7173	676		77	putative nuclear protein (8.3 kD) (mudar) mRNA.
<a href="#">mufer</a>	<a href="#">mufer.aSep08</a>		3031	270		45	putative protein (mufer) mRNA.
<a href="#">muflo</a>	<a href="#">muflo.aSep08</a>		4674	928		155	hermansky-Pudlak syndrome protein like (muflo) mRNA.
<a href="#">muflu</a>	<a href="#">muflu.aSep08</a>		1358	1213		230	ATPase class V type 10A (24.7 kD) (muflu) mRNA.
<a href="#">Mug2</a>	<a href="#">Mug2.aSep08</a>	<a href="#">408236</a>	14417	1169		389	murinoglobulin 2 (Mug2) mRNA.
<a href="#">mugar</a>	<a href="#">mugar.aSep08</a>		2660	216		52	putative protein (5.8 kD) (mugar) mRNA.
<a href="#">mujej</a>	<a href="#">mujej.aSep08</a>		5087	393	2	131	protein Tyrosine phosphatase non-receptor type 13 (mujej) alternative variant aSep08, mRNA.
<a href="#">mukee</a>	<a href="#">mukee.aSep08</a>		3756	786		104	CRA a like (mukee) mRNA.
<a href="#">muloy</a>	<a href="#">muloy.aSep08</a>		59626	487		79	putative protein (muloy) mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.aSep08</a>	<a href="#">362838</a>	16896	2572	14	698	melanoma associated antigen (mutated) 1 (78.0 kD) (Mum1) alternative variant aSep08, complete mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.cSep08</a>	<a href="#">362838</a>	5500	700	5	201	melanoma associated antigen (mutated) 1 (Mum1) alternative variant cSep08, mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.dSep08</a>	<a href="#">362838</a>	4534	795	5	192	melanoma associated antigen (mutated) 1 (Mum1) alternative variant dSep08, mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.eSep08</a>	<a href="#">362838</a>	959	604	2	118	melanoma associated antigen (mutated) 1 (Mum1) alternative variant eSep08, mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.gSep08</a>	<a href="#">362838</a>	5194	731	3	96	melanoma associated antigen (mutated) 1 (Mum1) alternative variant gSep08, mRNA.
<a href="#">Mum1</a>	<a href="#">Mum1.hSep08</a>	<a href="#">362838</a>	3030	806	3	57	melanoma associated antigen (mutated) 1 (6.8 kD) (Mum1) alternative variant hSep08, mRNA.
<a href="#">Mum111</a>	<a href="#">Mum111.aSep08</a>	<a href="#">501630</a>	4227	3602	2	679	melanoma associated antigen (mutated) 1-like 1 (77.2 kD) (Mum111) alternative variant aSep08, mRNA.
<a href="#">Mum111</a>	<a href="#">Mum111.cSep08</a>	<a href="#">501630</a>	29092	554	4	74	melanoma associated antigen (mutated) 1-like 1 (Mum111) alternative variant cSep08, mRNA.
<a href="#">mumee</a>	<a href="#">mumee.aSep08</a>		20601	1689	4	226	lipase maturation factor 1 (mumee) alternative variant aSep08, mRNA.
<a href="#">mumee</a>	<a href="#">mumee.bSep08</a>		6272	795	3	149	lipase maturation factor 1 (mumee) alternative variant bSep08, mRNA.
<a href="#">mumer</a>	<a href="#">mumer.aSep08</a>		3482	728		36	putative protein (4.3 kD) (mumer) mRNA.
<a href="#">munoy</a>	<a href="#">munoy.aSep08</a>		20189	632		78	putative protein (munoy) mRNA.
<a href="#">mupor</a>	<a href="#">mupor.aSep08</a>		11964	601	2	37	CRA b like (4.0 kD) (mupor) alternative variant aSep08, mRNA.

<a href="#">mupor</a>	<a href="#">mupor.bSep08</a>		10241	575	1	28	putative protein (3.0 kD) (mupor) alternative variant bSep08, mRNA.
<a href="#">Mus81</a>	<a href="#">Mus81.bSep08</a>	<a href="#">293678</a>	1491	599	6	140	mus81 endonuclease (Mus81) alternative variant bSep08, mRNA.
<a href="#">Mus81</a>	<a href="#">Mus81.cSep08</a>	<a href="#">293678</a>	1003	516	3	72	mus81 endonuclease (7.8 kD) (Mus81) alternative variant cSep08, mRNA.
<a href="#">Mus81</a>	<a href="#">Mus81.dSep08</a>	<a href="#">293678</a>	1895	1096	6	83	mus81 endonuclease (Mus81) alternative variant dSep08, mRNA.
<a href="#">musa</a>	<a href="#">musa.aSep08</a>		9398	918		305	putative protein of eukaryotic origin (musa) mRNA.
<a href="#">mushee</a>	<a href="#">mushee.aSep08</a>		10453	1025		144	putative protein of metazoan origin (mushee) mRNA.
<a href="#">Mut</a>	<a href="#">Mut.aSep08</a>	<a href="#">688517</a>	13889	2187		288	methylmalonyl-Coenzyme A mutase (Mut) mRNA.
<a href="#">MutS_II.0</a>	<a href="#">MutS_II.0.aSep08</a>		81400	1362		451	mutS homolog 3 (MutS_II.0) mRNA.
<a href="#">MutS_V.0</a>	<a href="#">MutS_V.0.aSep08</a>		4786	2188	7	603	mutS homolog (MutS_V.0) alternative variant aSep08, mRNA.
<a href="#">mutu</a>	<a href="#">mutu.aSep08</a>		4024	733		53	putative protein (mutu) mRNA.
<a href="#">Mutyh</a>	<a href="#">Mutyh.bSep08</a>	<a href="#">170841</a>	9560	732	10	143	mutY homolog (Mutyh) alternative variant bSep08, mRNA.
<a href="#">Mutyh</a>	<a href="#">Mutyh.cSep08</a>	<a href="#">170841</a>	2444	853	6	119	mutY homolog (Mutyh) alternative variant cSep08, mRNA.
<a href="#">Mutyh</a>	<a href="#">Mutyh.dSep08</a>	<a href="#">170841</a>	9106	1043	3	111	mutY homolog (12.9 kD) (Mutyh) alternative variant dSep08, complete mRNA.
<a href="#">Mutyh</a>	<a href="#">Mutyh.eSep08</a>	<a href="#">170841</a>	1620	649	3	92	putative mitochondrial protein (9.3 kD) (Mutyh) alternative variant eSep08, mRNA.
<a href="#">muvar</a>	<a href="#">muvar.aSep08</a>		1018	493		65	putative protein (6.8 kD) (muvar) mRNA.
<a href="#">muwey</a>	<a href="#">muwey.aSep08</a>		7993	1061		185	ER degradation enhancer mannosidase alpha-like 1 (muwey) mRNA.
<a href="#">Mvd</a>	<a href="#">Mvd.bSep08</a>	<a href="#">81726</a>	491	381	2	44	mevalonate (diphospho) decarboxylase (4.7 kD) (Mvd) alternative variant bSep08, mRNA.
<a href="#">Mvp</a>	<a href="#">Mvp.bSep08</a>	<a href="#">64681</a>	1643	658	1	148	major vault protein (15.7 kD) (Mvp) alternative variant bSep08, mRNA.
<a href="#">Mx1</a>	<a href="#">Mx1.bSep08</a>	<a href="#">24575</a>	9191	386	3	69	myxovirus (influenza virus) resistance 1 (Mx1) alternative variant bSep08, mRNA.
<a href="#">Mx1</a>	<a href="#">Mx1.cSep08</a>	<a href="#">24575</a>	2178	288	2	26	myxovirus (influenza virus) resistance 1 (Mx1) alternative variant cSep08, mRNA.
<a href="#">Mxd1</a>	<a href="#">Mxd1.aSep08</a>	<a href="#">362391</a>	18410	946	5	231	max dimerization protein 1 (Mxd1) alternative variant aSep08, mRNA.
<a href="#">Mxd1</a>	<a href="#">Mxd1.bSep08</a>	<a href="#">362391</a>	17914	494	5	164	max dimerization protein 1 (Mxd1) alternative variant bSep08, mRNA.
<a href="#">Mxd4</a>	<a href="#">Mxd4.aSep08</a>	<a href="#">360961</a>	14051	3930	6	273	max dimerization protein 4 (Mxd4) alternative variant aSep08, mRNA.
<a href="#">Mxd4</a>	<a href="#">Mxd4.bSep08</a>	<a href="#">360961</a>	9841	404	5	134	max dimerization protein 4 (Mxd4) alternative variant bSep08, mRNA.
<a href="#">Mxd4</a>	<a href="#">Mxd4.eSep08</a>	<a href="#">360961</a>	1529	475	2	32	max dimerization protein 4 (3.7 kD) (Mxd4) alternative variant eSep08, mRNA.
<a href="#">Mxi1</a>	<a href="#">Mxi1.bSep08</a>	<a href="#">25701</a>	27516	756	5	154	max interacting protein 1 (17.6 kD) (Mxi1) alternative variant bSep08, mRNA.
<a href="#">Mxi1</a>	<a href="#">Mxi1.cSep08</a>	<a href="#">25701</a>	55103	380	4	126	max interacting protein 1 (Mxi1) alternative variant cSep08, mRNA.

<a href="#">Mxra7</a>	<a href="#">Mxra7.aSep08</a>	<a href="#">690599</a>	2773	463	2	76	matrix-remodelling associated 7 (Mxra7) alternative variant aSep08, mRNA.
<a href="#">Mxra7</a>	<a href="#">Mxra7.bSep08</a>	<a href="#">690599</a>	23007	1782	4		
<a href="#">Mxra7</a>	<a href="#">Mxra7.dSep08</a>	<a href="#">690599</a>	2603	373	3	61	matrix-remodelling associated 7 (6.5 kD) (Mxra7) alternative variant dSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.aSep08</a>	<a href="#">83721</a>	11340	2706	14	785	dishevelled 1 (Mxra8andDvl1) alternative variant aSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.aSep08</a>	<a href="#">313770</a>	11340	2706	14	785	dishevelled 1 (Mxra8andDvl1) alternative variant aSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.bSep08</a>	<a href="#">83721</a>	11902	2984	14	563	dishevelled 1 (61.1 kD) (Mxra8andDvl1) alternative variant bSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.bSep08</a>	<a href="#">313770</a>	11902	2984	14	563	dishevelled 1 (61.1 kD) (Mxra8andDvl1) alternative variant bSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.cSep08</a>	<a href="#">83721</a>	14610	2533	15	447	dishevelled 1 (49.2 kD) (Mxra8andDvl1) alternative variant cSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.cSep08</a>	<a href="#">313770</a>	14610	2533	15	447	dishevelled 1 (49.2 kD) (Mxra8andDvl1) alternative variant cSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.dSep08</a>	<a href="#">83721</a>	4356	2121	10	439	matrix-remodelling associated 8 (49.5 kD) (Mxra8andDvl1) alternative variant dSep08, complete mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.dSep08</a>	<a href="#">313770</a>	4356	2121	10	439	matrix-remodelling associated 8 (49.5 kD) (Mxra8andDvl1) alternative variant dSep08, complete mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.fSep08</a>	<a href="#">83721</a>	1691	717	7	238	dishevelled 1 (Mxra8andDvl1) alternative variant fSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.fSep08</a>	<a href="#">313770</a>	1691	717	7	238	dishevelled 1 (Mxra8andDvl1) alternative variant fSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.gSep08</a>	<a href="#">83721</a>	1941	703	5	190	matrix-remodelling associated 8 (Mxra8andDvl1) alternative variant gSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.gSep08</a>	<a href="#">313770</a>	1941	703	5	190	matrix-remodelling associated 8 (Mxra8andDvl1) alternative variant gSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.hSep08</a>	<a href="#">83721</a>	1657	468	2	145	dishevelled dsh homolog 1 CRA a like (Mxra8andDvl1) alternative variant hSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.hSep08</a>	<a href="#">313770</a>	1657	468	2	145	dishevelled dsh homolog 1 CRA a like (Mxra8andDvl1) alternative variant hSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.jSep08</a>	<a href="#">83721</a>	1203	763	2	129	dishevelled dsh homolog 1 CRA b (13.3 kD) (Mxra8andDvl1) alternative variant jSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.jSep08</a>	<a href="#">313770</a>	1203	763	2	129	dishevelled dsh homolog 1 CRA b (13.3 kD) (Mxra8andDvl1) alternative variant jSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.kSep08</a>	<a href="#">83721</a>	2009	859	3	129	dishevelled dsh homolog 1 CRA b (13.3 kD) (Mxra8andDvl1) alternative variant kSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.kSep08</a>	<a href="#">313770</a>	2009	859	3	129	dishevelled dsh homolog 1 CRA b (13.3 kD) (Mxra8andDvl1) alternative variant kSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.lSep08</a>	<a href="#">83721</a>	1410	522	2	77	putative protein (Mxra8andDvl1) alternative variant lSep08, mRNA.
<a href="#">Mxra8andDvl1</a>	<a href="#">Mxra8andDvl1.lSep08</a>	<a href="#">313770</a>	1410	522	2	77	putative protein (Mxra8andDvl1) alternative variant lSep08, mRNA.
<a href="#">Myadm</a>	<a href="#">Myadm.bSep08</a>	<a href="#">369016</a>	3793	761	2	187	myeloid-associated differentiation marker (Myadm) alternative variant bSep08, mRNA.

<a href="#">Myadm</a>	<a href="#">Myadm.cSep08</a>	<a href="#">369016</a>	9820	1210	4	86	myeloid-associated differentiation marker (9.3 kD) (Myadm) alternative variant cSep08, mRNA.
<a href="#">Mybbp1a</a>	<a href="#">Mybbp1a.bSep08</a>	<a href="#">60571</a>	2242	797	6	265	MYB binding protein (P160) 1a (Mybbp1a) alternative variant bSep08, mRNA.
<a href="#">Mybbp1a</a>	<a href="#">Mybbp1a.cSep08</a>	<a href="#">60571</a>	1171	646	5	214	MYB binding protein (P160) 1a (Mybbp1a) alternative variant cSep08, mRNA.
<a href="#">Mybpc1</a>	<a href="#">Mybpc1.aSep08</a>	<a href="#">362867</a>	37256	1896	10	493	myosin binding protein C, slow type (Mybpc1) alternative variant aSep08, mRNA.
<a href="#">Mybpc1</a>	<a href="#">Mybpc1.bSep08</a>	<a href="#">362867</a>	16896	1325	8	381	myosin binding protein C, slow type (Mybpc1) alternative variant bSep08, mRNA.
<a href="#">Mybpc1</a>	<a href="#">Mybpc1.cSep08</a>	<a href="#">362867</a>	13184	1038	7	242	myosin binding protein C, slow type (Mybpc1) alternative variant cSep08, mRNA.
<a href="#">Mybpc1</a>	<a href="#">Mybpc1.dSep08</a>	<a href="#">362867</a>	8239	470	4	156	myosin binding protein C, slow type (Mybpc1) alternative variant dSep08, mRNA.
<a href="#">Mybpc1</a>	<a href="#">Mybpc1.eSep08</a>	<a href="#">362867</a>	5251	439	4	42	myosin binding protein C, slow type (Mybpc1) alternative variant eSep08, mRNA.
<a href="#">Mybpc2</a>	<a href="#">Mybpc2.bSep08</a>	<a href="#">292879</a>	8189	756	8	231	myosin binding protein C, fast-type (Mybpc2) alternative variant bSep08, mRNA.
<a href="#">Mybpc2</a>	<a href="#">Mybpc2.cSep08</a>	<a href="#">292879</a>	1835	691	3	143	myosin binding protein C, fast-type (Mybpc2) alternative variant cSep08, mRNA.
<a href="#">Mybpc2</a>	<a href="#">Mybpc2.dSep08</a>	<a href="#">292879</a>	1044	453	3	96	myosin binding protein C, fast-type (Mybpc2) alternative variant dSep08, mRNA.
<a href="#">Mybpc2</a>	<a href="#">Mybpc2.eSep08</a>	<a href="#">292879</a>	465	288	2	92	myosin binding protein C, fast-type (Mybpc2) alternative variant eSep08, mRNA.
<a href="#">Mybpc3</a>	<a href="#">Mybpc3.bSep08</a>	<a href="#">295929</a>	2816	741	9	216	myosin binding protein C, cardiac (Mybpc3) alternative variant bSep08, mRNA.
<a href="#">Mybpc3</a>	<a href="#">Mybpc3.cSep08</a>	<a href="#">295929</a>	1704	891	6	190	myosin binding protein C, cardiac (Mybpc3) alternative variant cSep08, mRNA.
<a href="#">Mybph</a>	<a href="#">Mybph.cSep08</a>	<a href="#">83708</a>	825	325	2	96	myosin binding protein H (Mybph) alternative variant cSep08, mRNA.
<a href="#">Mybphl</a>	<a href="#">Mybphl.aSep08</a>	<a href="#">310782</a>	2169	1785	3	486	myosin binding protein H-like (Mybphl) alternative variant aSep08, mRNA.
<a href="#">myby</a>	<a href="#">myby.aSep08</a>		159739	481		37	putative protein (myby) mRNA.
<a href="#">Myb_DNA-binding.0</a>	<a href="#">Myb_DNA-binding.0.aSep08</a>		4806	395		131	myb-like SWIRM MPN domains 1 (Myb_DNA-binding.0) mRNA.
<a href="#">Myb_DNA-binding.1</a>	<a href="#">Myb_DNA-binding.1.aSep08</a>		2415	424		140	myeloblastosis oncogene CRA b (Myb_DNA-binding.1) mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.cSep08</a>	<a href="#">290447</a>	15549	846	7	282	MYC binding protein 2 (Mycbp2) alternative variant cSep08, mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.dSep08</a>	<a href="#">290447</a>	29387	732	7	243	MYC binding protein 2 (Mycbp2) alternative variant dSep08, mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.eSep08</a>	<a href="#">290447</a>	5946	460	5	153	MYC binding protein 2 (Mycbp2) alternative variant eSep08, mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.fSep08</a>	<a href="#">290447</a>	5934	738	5	152	MYC binding protein 2 (Mycbp2) alternative variant fSep08, mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.gSep08</a>	<a href="#">290447</a>	2878	454	4	151	MYC binding protein 2 (Mycbp2) alternative variant gSep08, mRNA.

<a href="#">Mycbp2</a>	<a href="#">Mycbp2.hSep08</a>	<a href="#">290447</a>	1406	314	2	104	MYC binding protein 2 (Mycbp2) alternative variant hSep08, mRNA.
<a href="#">Mycbp2</a>	<a href="#">Mycbp2.iSep08</a>	<a href="#">290447</a>	15288	974	3	86	MYC binding protein 2 (Mycbp2) alternative variant iSep08, mRNA.
<a href="#">mychy</a>	<a href="#">mychy.aSep08</a>		39877	846		181	CRA a (20.6 kD) (mychy) mRNA.
<a href="#">mydar</a>	<a href="#">mydar.aSep08</a>		4115	596		105	putative mitochondrial protein (11.4 kD) (mydar) mRNA.
<a href="#">Myef2</a>	<a href="#">Myef2.bSep08</a>	<a href="#">679712</a>	18530	1008	10	336	myelin basic protein expression factor 2 repressor CRA c (Myef2) alternative variant bSep08, mRNA.
<a href="#">Myef2</a>	<a href="#">Myef2.cSep08</a>	<a href="#">679712</a>	7971	1684	3	183	myelin basic protein expression factor 2 repressor (20.1 kD) (Myef2) alternative variant cSep08, mRNA.
<a href="#">Myef2</a>	<a href="#">Myef2.dSep08</a>	<a href="#">679712</a>	10991	1684	4	128	myelin basic protein expression factor 2 repressor CRA c (Myef2) alternative variant dSep08, mRNA.
<a href="#">Myef2</a>	<a href="#">Myef2.eSep08</a>	<a href="#">679712</a>	1105	529	3	58	myelin expression factor 2 (Myef2) alternative variant eSep08, mRNA.
<a href="#">Myeov2</a>	<a href="#">Myeov2.aSep08</a>	<a href="#">681389</a>	4619	258	1	77	myeloma overexpressed 2 (Myeov2) alternative variant aSep08, mRNA.
<a href="#">myfer</a>	<a href="#">myfer.aSep08</a>		6108	1675		181	GTP binding protein 4 like (myfer) mRNA.
<a href="#">myflo</a>	<a href="#">myflo.aSep08</a>		12061	404		71	putative protein (8.2 kD) (myflo) mRNA.
<a href="#">myflu</a>	<a href="#">myflu.aSep08</a>		6650	751		250	e1-E2 ATPase (myflu) mRNA.
<a href="#">mygar</a>	<a href="#">mygar.aSep08</a>		2823	1784		18	putative protein (1.7 kD) (mygar) alternative variant aSep08, mRNA.
<a href="#">Myh4</a>	<a href="#">Myh4.bSep08</a>	<a href="#">360543</a>	582	487	1	132	myosin, heavy polypeptide 4, skeletal muscle (14.7 kD) (Myh4) alternative variant bSep08, mRNA.
<a href="#">Myh6</a>	<a href="#">Myh6.bSep08</a>	<a href="#">29556</a>	14236	1457	9	349	myosin, heavy polypeptide 6, cardiac muscle, alpha (Myh6) alternative variant bSep08, mRNA.
<a href="#">Myh6</a>	<a href="#">Myh6.cSep08</a>	<a href="#">29556</a>	5354	964	7	298	myosin, heavy polypeptide 6, cardiac muscle, alpha (Myh6) alternative variant cSep08, mRNA.
<a href="#">Myh6</a>	<a href="#">Myh6.dSep08</a>	<a href="#">29556</a>	1688	755	3	230	myosin, heavy polypeptide 6, cardiac muscle, alpha (Myh6) alternative variant dSep08, mRNA.
<a href="#">Myh6</a>	<a href="#">Myh6.eSep08</a>	<a href="#">29556</a>	1691	1374	2	100	myosin, heavy polypeptide 6, cardiac muscle, alpha (Myh6) alternative variant eSep08, mRNA.
<a href="#">Myh6</a>	<a href="#">Myh6.fSep08</a>	<a href="#">29556</a>	1452	1313	2	88	myosin, heavy polypeptide 6, cardiac muscle, alpha (Myh6) alternative variant fSep08, mRNA.
<a href="#">Myh8</a>	<a href="#">Myh8.aSep08</a>	<a href="#">252942</a>	2916	686		179	myosin, heavy polypeptide 8, skeletal muscle, perinatal (Myh8) mRNA.
<a href="#">Myh9</a>	<a href="#">Myh9.bSep08</a>	<a href="#">25745</a>	2343	773	4	257	myosin, heavy polypeptide 9, non-muscle (Myh9) alternative variant bSep08, mRNA.
<a href="#">Myh9</a>	<a href="#">Myh9.cSep08</a>	<a href="#">25745</a>	33258	695	3	159	myosin, heavy polypeptide 9, non-muscle (Myh9) alternative variant cSep08, mRNA.
<a href="#">Myh10</a>	<a href="#">Myh10.bSep08</a>	<a href="#">79433</a>	1072	478	2	105	myosin heavy chain (11.8 kD) (Myh10) alternative variant bSep08, mRNA.
<a href="#">Myh10</a>	<a href="#">Myh10.cSep08</a>	<a href="#">79433</a>	1379	400	2	101	myosin heavy chain non-muscle (Myh10) alternative variant cSep08, mRNA.
<a href="#">Myh11</a>	<a href="#">Myh11.aSep08</a>	<a href="#">24582</a>	70548	2982	23	960	myosin, heavy polypeptide 11, smooth muscle (Myh11) alternative variant aSep08, mRNA.



<a href="#">Myh11</a>	<a href="#">Myh11.bSep08</a>	<a href="#">24582</a>	5543	605	3	201	myosin, heavy polypeptide 11, smooth muscle (Myh11) alternative variant bSep08, mRNA.
<a href="#">Myh11</a>	<a href="#">Myh11.cSep08</a>	<a href="#">24582</a>	12422	395	3	62	myosin, heavy polypeptide 11, smooth muscle (Myh11) alternative variant cSep08, mRNA.
<a href="#">Myh13</a>	<a href="#">Myh13.aSep08</a>	<a href="#">29605</a>	2829	510		170	myosin, heavy polypeptide 13, skeletal muscle (Myh13) mRNA.
<a href="#">Myh14</a>	<a href="#">Myh14.bSep08</a>	<a href="#">308572</a>	2969	707	4	160	myosin, heavy polypeptide 14 (Myh14) alternative variant bSep08, mRNA.
<a href="#">Myh14</a>	<a href="#">Myh14.cSep08</a>	<a href="#">308572</a>	6979	307	3	102	myosin, heavy polypeptide 14 (Myh14) alternative variant cSep08, mRNA.
<a href="#">myjey</a>	<a href="#">myjey.aSep08</a>		2984	669		69	putative protein (7.5 kD) (myjey) mRNA.
<a href="#">mykee</a>	<a href="#">mykee.aSep08</a>		1238	543		36	putative protein (4.0 kD) (mykee) mRNA.
<a href="#">Myl1</a>	<a href="#">Myl1.aSep08</a>	<a href="#">56781</a>	10154	1113	6	197	myosin, light polypeptide 1 (21.9 kD) (Myl1) alternative variant aSep08, mRNA.
<a href="#">Myl1</a>	<a href="#">Myl1.cSep08</a>	<a href="#">56781</a>	10348	792	6	162	myosin, light polypeptide 1 (17.9 kD) (Myl1) alternative variant cSep08, mRNA.
<a href="#">Myl1</a>	<a href="#">Myl1.eSep08</a>	<a href="#">56781</a>	7827	742	3	64	myosin, light polypeptide 1 (7.1 kD) (Myl1) alternative variant eSep08, mRNA.
<a href="#">Myl1</a>	<a href="#">Myl1.fSep08</a>	<a href="#">56781</a>	10509	978	7	92	myosin, light polypeptide 1 (10.3 kD) (Myl1) alternative variant fSep08, mRNA.
<a href="#">Myl3</a>	<a href="#">Myl3.bSep08</a>	<a href="#">24585</a>	28168	1394	3	200	myosin, light polypeptide 3 (22.2 kD) (Myl3) alternative variant bSep08, mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.aSep08</a>	<a href="#">685867</a>	2996	675	7	151	myosin, light chain 6, alkali, smooth muscle and non-muscle (16.9 kD) (Myl6) alternative variant aSep08, complete mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.cSep08</a>	<a href="#">685867</a>	2982	1668	6	117	myosin, light chain 6, alkali, smooth muscle and non-muscle (13.0 kD) (Myl6) alternative variant cSep08, complete mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.dSep08</a>	<a href="#">685867</a>	2682	1007	7	116	myosin, light chain 6, alkali, smooth muscle and non-muscle (12.9 kD) (Myl6) alternative variant dSep08, mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.eSep08</a>	<a href="#">685867</a>	1728	488	4	99	myosin, light chain 6, alkali, smooth muscle and non-muscle (11.1 kD) (Myl6) alternative variant eSep08, mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.fSep08</a>	<a href="#">685867</a>	1621	1208	3	81	myosin, light chain 6, alkali, smooth muscle and non-muscle (Myl6) alternative variant fSep08, mRNA.
<a href="#">Myl6</a>	<a href="#">Myl6.gSep08</a>	<a href="#">685867</a>	1066	822	2	87	myosin, light chain 6, alkali, smooth muscle and non-muscle (9.7 kD) (Myl6) alternative variant gSep08, mRNA.
<a href="#">Myl6b</a>	<a href="#">Myl6b.aSep08</a>	<a href="#">685883</a>	3071	907	6	215	myosin, light chain 6B, alkali, smooth muscle and non-muscle (Myl6b) alternative variant aSep08, mRNA.
<a href="#">Myl6b</a>	<a href="#">Myl6b.bSep08</a>	<a href="#">685883</a>	1559	387	3	56	myosin, light chain 6B, alkali, smooth muscle and non-muscle (Myl6b) alternative variant bSep08, mRNA.
<a href="#">Myl7</a>	<a href="#">Myl7.aSep08</a>	<a href="#">289759</a>	3116	1487	7	177	myosin, light polypeptide 7, regulatory (Myl7) alternative variant aSep08, mRNA.
<a href="#">Myl7</a>	<a href="#">Myl7.cSep08</a>	<a href="#">289759</a>	1964	455	5	136	myosin, light polypeptide 7, regulatory (Myl7) alternative variant cSep08, mRNA.
<a href="#">Myl9</a>	<a href="#">Myl9.bSep08</a>	<a href="#">296313</a>	5949	716	1	204	myosin, light chain 9, regulatory (Myl9) alternative variant bSep08, mRNA.

<a href="#">Mylip</a>	<a href="#">Mylip.bSep08</a>	<a href="#">306825</a>	6392	1310	4	257	myosin regulatory light chain interacting protein (Mylip) alternative variant bSep08, mRNA.
<a href="#">Mylip</a>	<a href="#">Mylip.cSep08</a>	<a href="#">306825</a>	1491	384	2	79	myosin regulatory light chain interacting protein (Mylip) alternative variant cSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.bSep08</a>	<a href="#">288057</a>	5376	698	3	152	myosin, light polypeptide kinase (Mylk) alternative variant bSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.cSep08</a>	<a href="#">288057</a>	5382	701	3	150	myosin, light polypeptide kinase (Mylk) alternative variant cSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.dSep08</a>	<a href="#">288057</a>	16005	420	3	140	myosin, light polypeptide kinase (Mylk) alternative variant dSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.eSep08</a>	<a href="#">288057</a>	18568	464	5	127	myosin, light polypeptide kinase (Mylk) alternative variant eSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.fSep08</a>	<a href="#">288057</a>	26802	411	5	101	myosin, light polypeptide kinase (Mylk) alternative variant fSep08, mRNA.
<a href="#">Mylk</a>	<a href="#">Mylk.gSep08</a>	<a href="#">288057</a>	52369	312	3	77	myosin, light polypeptide kinase (Mylk) alternative variant gSep08, mRNA.
<a href="#">Mylk3</a>	<a href="#">Mylk3.bSep08</a>	<a href="#">291926</a>	9266	744	3	164	myosin light chain kinase 3 (Mylk3) alternative variant bSep08, mRNA.
<a href="#">myloy</a>	<a href="#">myloy.aSep08</a>		5103	896		152	putative protein (myloy) mRNA.
<a href="#">mymee</a>	<a href="#">mymee.aSep08</a>		862	718		88	putative protein (mymee) mRNA.
<a href="#">mymer</a>	<a href="#">mymer.aSep08</a>		1495	907		84	putative protein (mymer) mRNA.
<a href="#">Mynn</a>	<a href="#">Mynn.aSep08</a>	<a href="#">361924</a>	16754	3661	4	610	myoneurin (68.7 kD) (Mynn) alternative variant aSep08, complete mRNA.
<a href="#">Mynn</a>	<a href="#">Mynn.cSep08</a>	<a href="#">361924</a>	10105	1329	2	153	myoneurin (17.0 kD) (Mynn) alternative variant cSep08, mRNA.
<a href="#">mynoy</a>	<a href="#">mynoy.aSep08</a>		661	399		29	putative protein (3.1 kD) (mynoy) mRNA.
<a href="#">Myo1a</a>	<a href="#">Myo1a.aSep08</a>	<a href="#">299509</a>	1668	836		212	myosin IA (Myo1a) mRNA.
<a href="#">Myo1b</a>	<a href="#">Myo1b.aSep08</a>	<a href="#">117057</a>	66983	511		117	myosin Ib (Myo1b) mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.aSep08</a>	<a href="#">65261</a>	21378	4479	32	954	myosin-1c (109.4 kD) (Myo1c) alternative variant aSep08, mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.bSep08</a>	<a href="#">65261</a>	1605	575	5	148	myosin 1C (Myo1c) alternative variant bSep08, mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.cSep08</a>	<a href="#">65261</a>	4244	1134	6	139	myosin 1C (16.3 kD) (Myo1c) alternative variant cSep08, mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.dSep08</a>	<a href="#">65261</a>	7450	376	4	120	myosin 1C (Myo1c) alternative variant dSep08, mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.eSep08</a>	<a href="#">65261</a>	634	531	2	107	putative protein (Myo1c) alternative variant eSep08, mRNA.
<a href="#">Myo1c</a>	<a href="#">Myo1c.fSep08</a>	<a href="#">65261</a>	2129	436	4	84	myosin 1C (Myo1c) alternative variant fSep08, mRNA.
<a href="#">Myo1d</a>	<a href="#">Myo1d.aSep08</a>	<a href="#">25485</a>	96780	1150		381	myosin ID (Myo1d) mRNA.
<a href="#">Myo1e</a>	<a href="#">Myo1e.bSep08</a>	<a href="#">25484</a>	28556	1034	9	344	myosin IE (Myo1e) alternative variant bSep08, mRNA.
<a href="#">Myo1e</a>	<a href="#">Myo1e.cSep08</a>	<a href="#">25484</a>	18328	1310	3	111	myosin IE (Myo1e) alternative variant cSep08, mRNA.
<a href="#">Myo1f</a>	<a href="#">Myo1f.bSep08</a>	<a href="#">314654</a>	8582	1010	7	336	myosin IF (Myo1f) alternative variant bSep08, mRNA.
<a href="#">Myo1f</a>	<a href="#">Myo1f.cSep08</a>	<a href="#">314654</a>	510	407	2	77	myosin IF (Myo1f) alternative variant cSep08, mRNA.
<a href="#">Myo1g</a>	<a href="#">Myo1g.bSep08</a>	<a href="#">289785</a>	2609	641	2	213	myosin IG (Myo1g) alternative variant bSep08, mRNA.
<a href="#">Myo1g</a>	<a href="#">Myo1g.cSep08</a>	<a href="#">289785</a>	4240	692	2	180	myosin IG (Myo1g) alternative variant cSep08, mRNA.

<a href="#">Myo5a</a>	<a href="#">Myo5a.aSep08</a>	<a href="#">25017</a>	33544	3892	9	440	myosin Va (50.4 kD) (Myo5a) alternative variant aSep08, mRNA.
<a href="#">Myo5a</a>	<a href="#">Myo5a.bSep08</a>	<a href="#">25017</a>	2924	643	2	89	myosin Va (Myo5a) alternative variant bSep08, mRNA.
<a href="#">Myo5b</a>	<a href="#">Myo5b.aSep08</a>	<a href="#">25132</a>	31660	1993		664	myosin Vb (Myo5b) mRNA.
<a href="#">Myo5c</a>	<a href="#">Myo5c.bSep08</a>	<a href="#">315820</a>	4242	735	2	128	myosin VC (Myo5c) alternative variant bSep08, mRNA.
<a href="#">Myo6</a>	<a href="#">Myo6.aSep08</a>	<a href="#">315840</a>	92409	835		213	myosin VI (Myo6) mRNA.
<a href="#">Myo7a</a>	<a href="#">Myo7a.bSep08</a>	<a href="#">266714</a>	2042	745	3	65	myosin VIIA (7.1 kD) (Myo7a) alternative variant bSep08, mRNA.
<a href="#">Myo7b</a>	<a href="#">Myo7b.aSep08</a>	<a href="#">498834</a>	8362	1038		197	myosin VIIb (Myo7b) mRNA.
<a href="#">Myo9b</a>	<a href="#">Myo9b.bSep08</a>	<a href="#">25486</a>	4042	883	7	294	myosin IXb (Myo9b) alternative variant bSep08, mRNA.
<a href="#">Myo9b</a>	<a href="#">Myo9b.cSep08</a>	<a href="#">25486</a>	2348	684	7	227	myosin IXb (Myo9b) alternative variant cSep08, mRNA.
<a href="#">Myo9b</a>	<a href="#">Myo9b.dSep08</a>	<a href="#">25486</a>	1772	1343	3	88	myosin IXb (Myo9b) alternative variant dSep08, mRNA.
<a href="#">Myo9b</a>	<a href="#">Myo9b.fSep08</a>	<a href="#">25486</a>	6083	1087	3	60	myosin IXb (6.6 kD) (Myo9b) alternative variant fSep08, mRNA.
<a href="#">Myo15</a>	<a href="#">Myo15.aSep08</a>	<a href="#">501699</a>	2051	409		136	myosin XV (Myo15) mRNA.
<a href="#">Myo16</a>	<a href="#">Myo16.bSep08</a>	<a href="#">192253</a>	27082	834	3	73	myosin XVI (Myo16) alternative variant bSep08, mRNA.
<a href="#">Myocd</a>	<a href="#">Myocd.bSep08</a>	<a href="#">246297</a>	47895	384	1	93	transcription factor myocardin (Myocd) alternative variant bSep08, mRNA.
<a href="#">Myom1</a>	<a href="#">Myom1.aSep08</a>	<a href="#">316740</a>	63552	3280	24	932	myomesin 1 (Myom1) alternative variant aSep08, mRNA.
<a href="#">Myom1</a>	<a href="#">Myom1.bSep08</a>	<a href="#">316740</a>	582	280	2	55	myomesin 1 (Myom1) alternative variant bSep08, mRNA.
<a href="#">Myom2</a>	<a href="#">Myom2.aSep08</a>	<a href="#">306616</a>	29212	2288		762	myomesin 2 (Myom2) mRNA.
<a href="#">Myosin_head.0</a>	<a href="#">Myosin_head.0.aSep08</a>		25413	1802		519	myosin VC CRA d (Myosin_head.0) alternative variant aSep08, mRNA.
<a href="#">Myosin_head.0</a>	<a href="#">Myosin_head.0.bSep08</a>		18029	969		322	myosin VC CRA b (Myosin_head.0) alternative variant bSep08, mRNA.
<a href="#">Myosin_head.0</a>	<a href="#">Myosin_head.0.cSep08</a>		5666	755		101	myosin VC CRA b (Myosin_head.0) alternative variant cSep08, mRNA.
<a href="#">Myosin_N.0</a>	<a href="#">Myosin_N.0.aSep08</a>		2255	333		89	myosin heavy chain (Myosin_N.0) mRNA.
<a href="#">Myosin_N.1</a>	<a href="#">Myosin_N.1.aSep08</a>		1814	330		85	myosin heavy chain (Myosin_N.1) mRNA.
<a href="#">Myosin_tail_1.0</a>	<a href="#">Myosin_tail_1.0.aSep08</a>		21860	3286	17	932	myosin tail (Myosin_tail_1.0) alternative variant aSep08, mRNA.
<a href="#">Myosin_tail_1.0</a>	<a href="#">Myosin_tail_1.0.bSep08</a>		3844	720	4	111	myosin heavy chain (Myosin_tail_1.0) alternative variant bSep08, mRNA.
<a href="#">Myosin_tail_1.0</a>	<a href="#">Myosin_tail_1.0.cSep08</a>		6202	739	3	78	putative protein human specific (8.9 kD) (Myosin_tail_1.0) alternative variant cSep08, mRNA.
<a href="#">Myosin_tail_1.1</a>	<a href="#">Myosin_tail_1.1.aSep08</a>		9405	2820		902	myosin heavy chain skeletal muscle adult (Myosin_tail_1.1) mRNA.
<a href="#">Myot</a>	<a href="#">Myot.bSep08</a>	<a href="#">291605</a>	7561	671	1	175	myotilin (Myot) alternative variant bSep08, mRNA.
<a href="#">Myotub-related.0</a>	<a href="#">Myotub-related.0.aSep08</a>		6785	1766	11	588	CRA b (Myotub-related.0) alternative variant aSep08, mRNA.
<a href="#">Myotub-related.0</a>	<a href="#">Myotub-related.0.bSep08</a>		4128	605	4	201	CRA b (Myotub-related.0) alternative variant bSep08, mRNA.
<a href="#">Myoz2</a>	<a href="#">Myoz2.aSep08</a>	<a href="#">295426</a>	26705	1119	3	264	myozenin 2 (29.8 kD) (Myoz2) alternative variant aSep08, mRNA.
<a href="#">Myoz2</a>	<a href="#">Myoz2.cSep08</a>	<a href="#">295426</a>	18274	840	2	144	myozenin 2 (Myoz2) alternative variant cSep08, mRNA.

<a href="#">mypor</a>	<a href="#">mypor.aSep08</a>		9754	795		54	putative protein (5.8 kD) (mypor) mRNA.
<a href="#">Myrip</a>	<a href="#">Myrip.bSep08</a>	<a href="#">360034</a>	69397	1776	3	447	myosin VIIA and Rab interacting protein (Myrip) alternative variant bSep08, mRNA.
<a href="#">Myrip</a>	<a href="#">Myrip.cSep08</a>	<a href="#">360034</a>	28720	766	4	233	myosin VIIA and Rab interacting protein (Myrip) alternative variant cSep08, mRNA.
<a href="#">mysa</a>	<a href="#">mysa.aSep08</a>		3655	328		30	putative protein (mysa) mRNA.
<a href="#">myshee</a>	<a href="#">myshee.aSep08</a>		2333	761		110	repeat domain 50 (myshee) mRNA.
<a href="#">Mysm1</a>	<a href="#">Mysm1.aSep08</a>	<a href="#">298247</a>	6748	916		264	myb-like, SWIRM and MPN domains 1 (Mysm1) mRNA.
<a href="#">Myst1</a>	<a href="#">Myst1.bSep08</a>	<a href="#">310194</a>	5082	700	4	172	myst histone acetyltransferase (Myst1) alternative variant bSep08, mRNA.
<a href="#">Myst1</a>	<a href="#">Myst1.cSep08</a>	<a href="#">310194</a>	1329	731	4	153	myst histone acetyltransferase (17.5 kD) (Myst1) alternative variant cSep08, mRNA.
<a href="#">Myst1</a>	<a href="#">Myst1.dSep08</a>	<a href="#">310194</a>	4942	2985	3	126	histone acetyltransferase (Myst1) alternative variant dSep08, mRNA.
<a href="#">Myst2</a>	<a href="#">Myst2.bSep08</a>	<a href="#">303470</a>	34593	2447	12	516	MYST histone acetyltransferase 2 (59.4 kD) (Myst2) alternative variant bSep08, mRNA.
<a href="#">Myst2</a>	<a href="#">Myst2.cSep08</a>	<a href="#">303470</a>	2130	790	2	37	MYST histone acetyltransferase 2 (Myst2) alternative variant cSep08, mRNA.
<a href="#">Myst3</a>	<a href="#">Myst3.aSep08</a>	<a href="#">306571</a>	20052	1255	1	418	MYST histone acetyltransferase (monocytic leukemia) 3 (Myst3) alternative variant aSep08, mRNA.
<a href="#">Myst3</a>	<a href="#">Myst3.bSep08</a>	<a href="#">306571</a>	9778	419	1	139	MYST histone acetyltransferase (monocytic leukemia) 3 (Myst3) alternative variant bSep08, mRNA.
<a href="#">Myt11</a>	<a href="#">Myt11.bSep08</a>	<a href="#">116668</a>	67935	1785	7	542	myelin transcription factor 1-like (Myt11) alternative variant bSep08, mRNA.
<a href="#">Myt11</a>	<a href="#">Myt11.cSep08</a>	<a href="#">116668</a>	34628	1699	7	300	myelin transcription factor 1-like (Myt11) alternative variant cSep08, mRNA.
<a href="#">Myt11</a>	<a href="#">Myt11.dSep08</a>	<a href="#">116668</a>	208597	473	4	32	myelin transcription factor 1-like (3.2 kD) (Myt11) alternative variant dSep08, mRNA.
<a href="#">mytu</a>	<a href="#">mytu.aSep08</a>		8096	409		135	putative protein of eukaryotic origin (mytu) mRNA.
<a href="#">myvar</a>	<a href="#">myvar.aSep08</a>		3571	437		145	zinc finger protein 362 (myvar) mRNA.
<a href="#">mywey</a>	<a href="#">mywey.aSep08</a>		5596	262		87	ER degradation enhancer mannosidase alpha-like 1 (mywey) mRNA.
<a href="#">Mzf1</a>	<a href="#">Mzf1.cSep08</a>	<a href="#">361508</a>	2100	741	2	204	zinc finger (Mzf1) alternative variant cSep08, mRNA.
<a href="#">Mzf1</a>	<a href="#">Mzf1.dSep08</a>	<a href="#">361508</a>	11916	2263	3	136	zinc finger (15.7 kD) (Mzf1) alternative variant dSep08, mRNA.
<a href="#">Mzf1</a>	<a href="#">Mzf1.eSep08</a>	<a href="#">361508</a>	7096	578	4	134	CRA d (14.6 kD) (Mzf1) alternative variant eSep08, mRNA.
<a href="#">N4bp1</a>	<a href="#">N4bp1.aSep08</a>	<a href="#">291921</a>	11721	736		244	nedd4 binding protein 1 (N4bp1) mRNA.
<a href="#">N4bp2</a>	<a href="#">N4bp2.aSep08</a>	<a href="#">305342</a>	8390	293		97	NEDD4 binding protein 2 (N4bp2) mRNA.
<a href="#">N4bp2l2</a>	<a href="#">N4bp2l2.cSep08</a>	<a href="#">288416</a>	2333	444	2	148	NEDD4 binding protein 2-like 2 (N4bp2l2) alternative variant cSep08, mRNA.
<a href="#">N4bp2l2</a>	<a href="#">N4bp2l2.dSep08</a>	<a href="#">288416</a>	7307	292	3	37	NEDD4 binding protein 2-like 2 (N4bp2l2) alternative variant dSep08, mRNA.
<a href="#">N6amt1</a>	<a href="#">N6amt1.aSep08</a>	<a href="#">288309</a>	12476	949	3	214	N-6 adenine-specific DNA methyltransferase 1 (putative) (22.9 kD) (N6amt1) alternative variant aSep08, mRNA.
<a href="#">N6amt1</a>	<a href="#">N6amt1.bSep08</a>	<a href="#">288309</a>	7934	711	1	150	N-6 adenine-specific DNA methyltransferase 1 (putative) (N6amt1) alternative variant bSep08, complete mRNA.

<a href="#">N6amt2</a>	<a href="#">N6amt2.aSep08</a>	<a href="#">290279</a>	17043	883	4	246	N-6 adenine-specific DNA methyltransferase 2 (N6amt2) alternative variant aSep08, mRNA.
<a href="#">N6amt2</a>	<a href="#">N6amt2.bSep08</a>	<a href="#">290279</a>	13396	1277	4	244	N-6 adenine-specific DNA methyltransferase 2 (27.8 kD) (N6amt2) alternative variant bSep08, mRNA.
<a href="#">N6amt2</a>	<a href="#">N6amt2.cSep08</a>	<a href="#">290279</a>	11880	717	4	145	N-6 adenine-specific DNA methyltransferase 2 (N6amt2) alternative variant cSep08, mRNA.
<a href="#">N6amt2</a>	<a href="#">N6amt2.dSep08</a>	<a href="#">290279</a>	12098	727	2	130	N-6 adenine-specific DNA methyltransferase 2 (15.2 kD) (N6amt2) alternative variant dSep08, mRNA.
<a href="#">N6amt2</a>	<a href="#">N6amt2.eSep08</a>	<a href="#">290279</a>	2422	230	1	64	CRA a like (N6amt2) alternative variant eSep08, mRNA.
<a href="#">Naalad2</a>	<a href="#">Naalad2.bSep08</a>	<a href="#">300384</a>	73981	2998	15	346	N-acetylated alpha-linked acidic dipeptidase 2 (39.3 kD) (Naalad2) alternative variant bSep08, mRNA.
<a href="#">Naalad2</a>	<a href="#">Naalad2.cSep08</a>	<a href="#">300384</a>	22662	692	1	186	N-acetylated alpha-linked acidic dipeptidase 2 (Naalad2) alternative variant cSep08, mRNA.
<a href="#">Naalad1</a>	<a href="#">Naalad1.bSep08</a>	<a href="#">83568</a>	1365	1023	4	152	N-acetylated alpha-linked acidic dipeptidase-like 1 (Naalad1) alternative variant bSep08, mRNA.
<a href="#">Nab2</a>	<a href="#">Nab2.bSep08</a>	<a href="#">314910</a>	3406	1243	5	361	ngfi-A binding protein 2 (Nab2) alternative variant bSep08, mRNA.
<a href="#">naby</a>	<a href="#">naby.aSep08</a>		16245	516		100	ab2-143 like (naby) mRNA.
<a href="#">Naca</a>	<a href="#">Naca.aSep08</a>	<a href="#">288770</a>	12278	872	1	245	nascent-polypeptide-associated complex alpha polypeptide (Naca) alternative variant aSep08, mRNA.
<a href="#">Nacad</a>	<a href="#">Nacad.aSep08</a>	<a href="#">289786</a>	3406	2656	5	696	nacad protein (Nacad) alternative variant aSep08, mRNA.
<a href="#">Nacad</a>	<a href="#">Nacad.bSep08</a>	<a href="#">289786</a>	804	469	1	156	nascent polypeptide-associated complex NAC (Nacad) alternative variant bSep08, mRNA.
<a href="#">nachy</a>	<a href="#">nachy.aSep08</a>		5092	408		136	protein tyrosine phosphatase receptor type J (nachy) mRNA.
<a href="#">nadar</a>	<a href="#">nadar.aSep08</a>		3875	881		167	nedd4 binding protein 1 like (nadar) mRNA.
<a href="#">NADH_oxidored.0</a>	<a href="#">NADH_oxidored.0.aSep08</a>		8559	549	3	57	nadh dehydrogenase 1 beta subcomplex (7.0 kD) (NADH_oxidored.0) alternative variant aSep08, mRNA.
<a href="#">NADH_oxidored.0</a>	<a href="#">NADH_oxidored.0.bSep08</a>		8455	448	3	57	nadh dehydrogenase 1 beta subcomplex (7.0 kD) (NADH_oxidored.0) alternative variant bSep08, mRNA.
<a href="#">NADH_ub_rd_NUML.0</a>	<a href="#">NADH_ub_rd_NUML.0.aSep08</a>		2123	938	3	86	nadh dehydrogenase 1 alpha subcomplex 4-like 2 (NADH_ub_rd_NUML.0) alternative variant aSep08, mRNA.
<a href="#">NADH_ub_rd_NUML.0</a>	<a href="#">NADH_ub_rd_NUML.0.bSep08</a>		1126	772	1	39	NADH dehydrogenase 1 alpha subcomplex 4-like 2 (NADH_ub_rd_NUML.0) alternative variant bSep08, mRNA.
<a href="#">Nadk</a>	<a href="#">Nadk.aSep08</a>	<a href="#">100125370</a>	28517	1931	12	455	NAD kinase (50.4 kD) (Nadk) alternative variant aSep08, mRNA.
<a href="#">Nadk</a>	<a href="#">Nadk.cSep08</a>	<a href="#">100125370</a>	713	515	2	43	NAD kinase (4.9 kD) (Nadk) alternative variant cSep08, mRNA.
<a href="#">Nadsyn1</a>	<a href="#">Nadsyn1.bSep08</a>	<a href="#">353255</a>	2148	446	2	111	NAD synthetase 1 (Nadsyn1) alternative variant bSep08, mRNA.
<a href="#">Nadsyn1</a>	<a href="#">Nadsyn1.dSep08</a>	<a href="#">353255</a>	2191	639	4	86	NAD synthetase 1 (9.2 kD) (Nadsyn1) alternative variant dSep08, mRNA.
<a href="#">NAD_Gly3P_dh_C.0</a>	<a href="#">NAD_Gly3P_dh_C.0.aSep08</a>		1321	348		110	dehydrogenase 1-like CRA d (NAD_Gly3P_dh_C.0) mRNA.
<a href="#">Nae1</a>	<a href="#">Nae1.aSep08</a>	<a href="#">84019</a>	15263	1311		392	NEDD8 activating enzyme E1 subunit 1 (Nae1) mRNA.

<a href="#">Naf1</a>	<a href="#">Naf1.bSep08</a>	<a href="#">306387</a>	20223	2324	8	421	nuclear assembly factor 1 homolog ( <i>S. cerevisiae</i> ) (Naf1) alternative variant bSep08, mRNA.
<a href="#">nafer</a>	<a href="#">nafer.aSep08</a>		138869	739		72	putative protein (nafer) mRNA.
<a href="#">naflo</a>	<a href="#">naflo.aSep08</a>		68717	1219		60	putative protein (7.1 kD) (naflo) mRNA.
<a href="#">naflu</a>	<a href="#">naflu.aSep08</a>		14609	753		84	putative mitochondrial protein (9.6 kD) (naflu) mRNA.
<a href="#">Naga</a>	<a href="#">Naga.bSep08</a>	<a href="#">315165</a>	4117	850	7	283	N-acetyl galactosaminidase, alpha (Naga) alternative variant bSep08, mRNA.
<a href="#">Naga</a>	<a href="#">Naga.cSep08</a>	<a href="#">315165</a>	2917	486	5	99	N-acetyl galactosaminidase, alpha (Naga) alternative variant cSep08, mRNA.
<a href="#">nagar</a>	<a href="#">nagar.aSep08</a>		2917	717		69	putative protein (nagar) mRNA.
<a href="#">Nagk</a>	<a href="#">Nagk.bSep08</a>	<a href="#">297393</a>	6316	784	8	247	N-acetylglucosamine kinase (Nagk) alternative variant bSep08, mRNA.
<a href="#">Naglt1</a>	<a href="#">Naglt1.bSep08</a>	<a href="#">337920</a>	8062	760	1	253	na <sup>+</sup> dependent glucose transporter 1 (Naglt1) alternative variant bSep08, mRNA.
<a href="#">Nagpa</a>	<a href="#">Nagpa.bSep08</a>	<a href="#">360476</a>	2299	725	3	234	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (Nagpa) alternative variant bSep08, mRNA.
<a href="#">Nagpa</a>	<a href="#">Nagpa.cSep08</a>	<a href="#">360476</a>	4438	722	5	139	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (Nagpa) alternative variant cSep08, mRNA.
<a href="#">Nagpa</a>	<a href="#">Nagpa.dSep08</a>	<a href="#">360476</a>	3948	374	4	124	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (Nagpa) alternative variant dSep08, mRNA.
<a href="#">Nagpa</a>	<a href="#">Nagpa.eSep08</a>	<a href="#">360476</a>	2063	586	2	42	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (Nagpa) alternative variant eSep08, mRNA.
<a href="#">Nagpa</a>	<a href="#">Nagpa.fSep08</a>	<a href="#">360476</a>	3660	721	5	103	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (Nagpa) alternative variant fSep08, mRNA.
<a href="#">Naip2</a>	<a href="#">Naip2.aSep08</a>	<a href="#">191568</a>	9718	726		241	NLR family, apoptosis inhibitory protein 2 (Naip2) mRNA.
<a href="#">naja</a>	<a href="#">naja.aSep08</a>		5084	745		92	putative secreted or extracellular protein precursor (9.4 kD) (naja) mRNA.
<a href="#">najey</a>	<a href="#">najey.aSep08</a>		4552	345		114	beige/BEACH (najey) mRNA.
<a href="#">nakee</a>	<a href="#">nakee.aSep08</a>		12245	402		134	putative protein of ancient origin (nakee) mRNA.
<a href="#">Nalcn</a>	<a href="#">Nalcn.aSep08</a>	<a href="#">266760</a>	242410	4501	36	1499	sodium leak channel, non-selective (Nalcn) alternative variant aSep08, mRNA.
<a href="#">Nalcn</a>	<a href="#">Nalcn.cSep08</a>	<a href="#">266760</a>	7115	1934	3	132	sodium leak channel, non-selective (Nalcn) alternative variant cSep08, mRNA.
<a href="#">Nalcn</a>	<a href="#">Nalcn.dSep08</a>	<a href="#">266760</a>	4343	333	3	111	sodium leak channel, non-selective (Nalcn) alternative variant dSep08, mRNA.
<a href="#">naloy</a>	<a href="#">naloy.aSep08</a>		5849	468	3	66	vacuolar protein sorting 8 homolog (naloy) alternative variant aSep08, mRNA.
<a href="#">namee</a>	<a href="#">namee.aSep08</a>		7744	423		62	putative protein (6.8 kD) (namee) mRNA.
<a href="#">namer</a>	<a href="#">namer.aSep08</a>		2393	596		71	polymeric immunoglobulin receptor like (namer) mRNA.
<a href="#">Nampt</a>	<a href="#">Nampt.bSep08</a>	<a href="#">297508</a>	3112	263	2	44	nicotinamide phosphoribosyltransferase (5.0 kD) (Nampt) alternative variant bSep08, mRNA.

<a href="#">Nanog</a>	<a href="#">Nanog.bSep08</a>	<a href="#">414065</a>	5956	936	1	311	nanog homeobox (Nanog) alternative variant bSep08, mRNA.
<a href="#">nanoy</a>	<a href="#">nanoy.aSep08</a>		1754	860		117	type III (13.4 kD) (nanoy) mRNA.
<a href="#">Nans</a>	<a href="#">Nans.bSep08</a>	<a href="#">298071</a>	2456	717	3	172	N-acetylneuraminic acid synthase (sialic acid synthase) (Nans) alternative variant bSep08, mRNA.
<a href="#">Nans</a>	<a href="#">Nans.cSep08</a>	<a href="#">298071</a>	17512	1127	3	147	N-acetylneuraminic acid synthase (sialic acid synthase) (16.7 kD) (Nans) alternative variant cSep08, complete mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.bSep08</a>	<a href="#">89825</a>	23282	1293	14	328	nucleosome assembly protein 1-like 1 (38.2 kD) (Nap111) alternative variant bSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.cSep08</a>	<a href="#">89825</a>	4970	1170	9	290	nucleosome assembly protein 1-like 1 (Nap111) alternative variant cSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.dSep08</a>	<a href="#">89825</a>	5065	1590	8	231	nucleosome assembly protein 1-like 1 (Nap111) alternative variant dSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.eSep08</a>	<a href="#">89825</a>	18826	767	6	154	nucleosome assembly protein 1-like 1 CRA e (18.2 kD) (Nap111) alternative variant eSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.fSep08</a>	<a href="#">89825</a>	1746	337	5	101	nucleosome assembly protein 1-like 1 (Nap111) alternative variant fSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.gSep08</a>	<a href="#">89825</a>	4654	2301	6	93	nucleosome assembly protein 1-like 1 (Nap111) alternative variant gSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.hSep08</a>	<a href="#">89825</a>	4191	2552	4	44	putative protein (5.2 kD) (Nap111) alternative variant hSep08, mRNA.
<a href="#">Nap111</a>	<a href="#">Nap111.iSep08</a>	<a href="#">89825</a>	5997	387	5	50	putative protein (6.1 kD) (Nap111) alternative variant iSep08, mRNA.
<a href="#">Nap114</a>	<a href="#">Nap114.bSep08</a>	<a href="#">361684</a>	20887	816	9	247	nucleosome assembly protein 1-like 4 (Nap114) alternative variant bSep08, mRNA.
<a href="#">Nap114</a>	<a href="#">Nap114.cSep08</a>	<a href="#">361684</a>	13310	1455	7	126	nucleosome assembly protein 1-like 4 (Nap114) alternative variant cSep08, mRNA.
<a href="#">Nap115</a>	<a href="#">Nap115.bSep08</a>	<a href="#">688843</a>	1850	1445	2	189	nucleosome assembly protein 1-like 5 (Nap115) alternative variant bSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.bSep08</a>	<a href="#">140673</a>	18014	906	11	265	N-ethylmaleimide sensitive fusion protein attachment protein alpha (29.6 kD) (Napa) alternative variant bSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.cSep08</a>	<a href="#">140673</a>	18000	789	10	247	N-ethylmaleimide sensitive fusion protein attachment protein alpha (Napa) alternative variant cSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.dSep08</a>	<a href="#">140673</a>	15773	777	7	192	N-ethylmaleimide sensitive fusion protein attachment protein alpha (Napa) alternative variant dSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.eSep08</a>	<a href="#">140673</a>	14538	391	4	109	N-ethylmaleimide sensitive fusion protein attachment protein alpha (Napa) alternative variant eSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.fSep08</a>	<a href="#">140673</a>	5166	389	6	97	N-ethylmaleimide sensitive fusion protein attachment protein alpha (Napa) alternative variant fSep08, mRNA.
<a href="#">Napa</a>	<a href="#">Napa.gSep08</a>	<a href="#">140673</a>	1407	775	3	60	N-ethylmaleimide sensitive fusion protein attachment protein alpha (Napa) alternative variant gSep08, mRNA.
<a href="#">Napg</a>	<a href="#">Napg.aSep08</a>	<a href="#">307382</a>	14221	862	11	287	N-ethylmaleimide sensitive fusion protein attachment protein gamma (Napg) alternative variant aSep08, mRNA.

<a href="#">Napg</a>	<a href="#">Napg.cSep08</a>	<a href="#">307382</a>	2505	678	2	51	N-ethylmaleimide sensitive fusion protein attachment protein gamma (5.1 kD) (Napg) alternative variant cSep08, mRNA.
<a href="#">napor</a>	<a href="#">napor.aSep08</a>		4414	1125	2	82	ac1576 like (9.4 kD) (napor) alternative variant aSep08, mRNA.
<a href="#">napor</a>	<a href="#">napor.bSep08</a>		3225	765	1	81	putative protein (napor) alternative variant bSep08, mRNA.
<a href="#">Naprt1</a>	<a href="#">Naprt1.bSep08</a>	<a href="#">315085</a>	2030	1036	7	345	putative protein of ancient origin (Naprt1) alternative variant bSep08, mRNA.
<a href="#">Naprt1</a>	<a href="#">Naprt1.cSep08</a>	<a href="#">315085</a>	1921	747	6	172	putative protein of ancient origin (Naprt1) alternative variant cSep08, mRNA.
<a href="#">Naprt1</a>	<a href="#">Naprt1.dSep08</a>	<a href="#">315085</a>	839	663	3	168	putative protein of ancient origin (Naprt1) alternative variant dSep08, mRNA.
<a href="#">narby</a>	<a href="#">narby.aSep08</a>		6735	355	3	68	putative protein (7.6 kD) (narby) alternative variant aSep08, mRNA.
<a href="#">narby</a>	<a href="#">narby.bSep08</a>		13952	1456	2	74	putative protein (8.4 kD) (narby) alternative variant bSep08, mRNA.
<a href="#">narchy</a>	<a href="#">narchy.aSep08</a>		5732	693		207	ATP GTP binding protein-like 2 (narchy) mRNA.
<a href="#">nardar</a>	<a href="#">nardar.aSep08</a>		1175	589	1	110	putative protein (nardar) alternative variant aSep08, mRNA.
<a href="#">nardar</a>	<a href="#">nardar.bSep08</a>		1003	422	1	80	putative protein (nardar) alternative variant bSep08, mRNA.
<a href="#">narfer</a>	<a href="#">narfer.aSep08</a>		1202	791		83	putative protein (9.4 kD) (narfer) mRNA.
<a href="#">Narfl</a>	<a href="#">Narfl.bSep08</a>	<a href="#">360496</a>	3494	406	4	73	nuclear prelamin A recognition factor-like (8.0 kD) (Narfl) alternative variant bSep08, mRNA.
<a href="#">Narfl</a>	<a href="#">Narfl.dSep08</a>	<a href="#">360496</a>	1560	702	3	54	nuclear prelamin A recognition factor-like (Narfl) alternative variant dSep08, mRNA.
<a href="#">narflo</a>	<a href="#">narflo.aSep08</a>		733	651	1	46	putative protein (5.4 kD) (narflo) alternative variant aSep08, mRNA.
<a href="#">narflo</a>	<a href="#">narflo.bSep08</a>		1083	650	2	46	putative protein (5.4 kD) (narflo) alternative variant bSep08, mRNA.
<a href="#">narflu</a>	<a href="#">narflu.aSep08</a>		438	257		78	putative protein (narflu) mRNA.
<a href="#">Narg1</a>	<a href="#">Narg1.bSep08</a>	<a href="#">310399</a>	5416	820	5	215	NMDA receptor-regulated gene 1 (24.4 kD) (Narg1) alternative variant bSep08, mRNA.
<a href="#">Narg1</a>	<a href="#">Narg1.cSep08</a>	<a href="#">310399</a>	3539	381	3	126	NMDA receptor-regulated gene 1 (Narg1) alternative variant cSep08, mRNA.
<a href="#">Narg2</a>	<a href="#">Narg2.aSep08</a>	<a href="#">691379</a>	3990	362		112	NMDA receptor-regulated gene 2 (Narg2) mRNA.
<a href="#">nargar</a>	<a href="#">nargar.aSep08</a>		9241	249		50	putative protein (nargar) mRNA.
<a href="#">narja</a>	<a href="#">narja.aSep08</a>		666	375		65	putative protein (narja) mRNA.
<a href="#">narjey</a>	<a href="#">narjey.aSep08</a>		2928	1723		309	CRA b (34.3 kD) (narjey) mRNA.
<a href="#">narkee</a>	<a href="#">narkee.aSep08</a>		665	260		28	putative protein (narkee) mRNA.
<a href="#">narkler</a>	<a href="#">narkler.aSep08</a>		2594	731	2	80	endonuclease reverse transcriptase (9.3 kD) (narkler) alternative variant aSep08, mRNA.
<a href="#">narkler</a>	<a href="#">narkler.bSep08</a>		3584	687	2	35	putative protein (narkler) alternative variant bSep08, mRNA.
<a href="#">narloy</a>	<a href="#">narloy.aSep08</a>		515	294		98	chordin (narloy) mRNA.
<a href="#">narmee</a>	<a href="#">narmee.aSep08</a>		928	474		30	putative protein (narmee) mRNA.



<a href="#">narmer</a>	<a href="#">narmer.bSep08</a>		1109	1000			
<a href="#">naroy</a>	<a href="#">naroy.aSep08</a>		31289	563		37	putative protein (naroy) mRNA.
<a href="#">narpor</a>	<a href="#">narpor.aSep08</a>		3654	246		32	putative protein (3.5 kD) (narpor) mRNA.
<a href="#">Nars</a>	<a href="#">Nars.aSep08</a>	<a href="#">291556</a>	14962	1913	13	475	asparaginyl-tRNA synthetase (54.3 kD) (Nars) alternative variant aSep08, mRNA.
<a href="#">Nars</a>	<a href="#">Nars.cSep08</a>	<a href="#">291556</a>	12005	1706	9	273	asparaginyl-tRNA synthetase (Nars) alternative variant cSep08, mRNA.
<a href="#">Nars2</a>	<a href="#">Nars2.bSep08</a>	<a href="#">293128</a>	68175	2165	9	277	asparaginyl-tRNA synthetase 2 (mitochondrial)(putative) (Nars2) alternative variant bSep08, mRNA.
<a href="#">Nars2</a>	<a href="#">Nars2.dSep08</a>	<a href="#">293128</a>	24720	434	2	45	asparaginyl-tRNA synthetase 2 (mitochondrial)(putative) (Nars2) alternative variant dSep08, mRNA.
<a href="#">narsa</a>	<a href="#">narsa.aSep08</a>		18486	1571	13	523	CRA b (narsa) alternative variant aSep08, mRNA.
<a href="#">narsa</a>	<a href="#">narsa.bSep08</a>		2202	828	5	240	CRA b like (narsa) alternative variant bSep08, mRNA.
<a href="#">narsa</a>	<a href="#">narsa.cSep08</a>		3527	470	1	109	CRA b (narsa) alternative variant cSep08, mRNA.
<a href="#">narshee</a>	<a href="#">narshee.aSep08</a>		1511	363		37	putative protein (narshee) mRNA.
<a href="#">nartu</a>	<a href="#">nartu.aSep08</a>		1748	744		67	putative protein (7.8 kD) (nartu) mRNA.
<a href="#">narvar</a>	<a href="#">narvar.aSep08</a>		4707	753		34	putative protein (narvar) mRNA.
<a href="#">narwey</a>	<a href="#">narwey.aSep08</a>		6663	867		289	cullin-associated 2 (narwey) mRNA.
<a href="#">nasa</a>	<a href="#">nasa.aSep08</a>		3503	699		233	putative protein of mammalian origin (nasa) mRNA.
<a href="#">nashee</a>	<a href="#">nashee.aSep08</a>		1729	1109		332	forkhead box O1A (nashee) mRNA.
<a href="#">Nasp</a>	<a href="#">Nasp.bSep08</a>	<a href="#">298441</a>	25634	2132	7	492	nuclear autoantigenic sperm protein (histone-binding) (Nasp) alternative variant bSep08, mRNA.
<a href="#">Nasp</a>	<a href="#">Nasp.cSep08</a>	<a href="#">298441</a>	25636	2126	14	482	nuclear autoantigenic sperm protein (histone-binding) (Nasp) alternative variant cSep08, mRNA.
<a href="#">Nasp</a>	<a href="#">Nasp.dSep08</a>	<a href="#">298441</a>	8431	862	3	73	nuclear autoantigenic sperm protein (histone-binding) (7.4 kD) (Nasp) alternative variant dSep08, mRNA.
<a href="#">Nat5</a>	<a href="#">Nat5.bSep08</a>	<a href="#">362228</a>	16834	2690	4	178	N-acetyltransferase 5 (ARD1 homolog, <i>S. cerevisiae</i> ) (20.4 kD) (Nat5) alternative variant bSep08, complete mRNA.
<a href="#">Nat5</a>	<a href="#">Nat5.cSep08</a>	<a href="#">362228</a>	14353	462	3	143	N-acetyltransferase 5 (ARD1 homolog, <i>S. cerevisiae</i> ) (Nat5) alternative variant cSep08, mRNA.
<a href="#">Nat5</a>	<a href="#">Nat5.dSep08</a>	<a href="#">362228</a>	14448	585	3	106	N-acetyltransferase 5 (ARD1 homolog, <i>S. cerevisiae</i> ) (12.2 kD) (Nat5) alternative variant dSep08, complete mRNA.
<a href="#">Nat8l</a>	<a href="#">Nat8l.aSep08</a>	<a href="#">289727</a>	6370	3013		241	N-acetyltransferase 8-like (Nat8l) mRNA.
<a href="#">Nat9</a>	<a href="#">Nat9.aSep08</a>	<a href="#">303669</a>	4836	1163	6	329	NAT9 (Nat9) alternative variant aSep08, mRNA.
<a href="#">Nat9</a>	<a href="#">Nat9.cSep08</a>	<a href="#">303669</a>	1117	653	1	88	NAT9 (10.2 kD) (Nat9) alternative variant cSep08, mRNA.
<a href="#">Nat11</a>	<a href="#">Nat11.bSep08</a>	<a href="#">361718</a>	15978	430		141	N-acetyltransferase 11 (Nat11) alternative variant bSep08, mRNA.
<a href="#">Nat13</a>	<a href="#">Nat13.aSep08</a>	<a href="#">288108</a>	19642	1833	2	169	N-acetyltransferase 13 (19.4 kD) (Nat13) alternative variant aSep08, mRNA.
<a href="#">Nat14</a>	<a href="#">Nat14.bSep08</a>	<a href="#">361500</a>	1253	539	3	115	N-acetyltransferase 14 (Nat14) alternative variant bSep08, mRNA.
<a href="#">natu</a>	<a href="#">natu.bSep08</a>		14841	1377		124	putative protein, with a transmembrane domain (13.9 kD) (natu) alternative variant bSep08, mRNA.
<a href="#">Nav2</a>	<a href="#">Nav2.aSep08</a>	<a href="#">171563</a>	8809	837		175	neuron navigator 2 (Nav2) mRNA.
<a href="#">Nav3</a>	<a href="#">Nav3.aSep08</a>	<a href="#">314814</a>	4933	569		46	neuron navigator 3 (Nav3) mRNA.

<a href="#">navar</a>	<a href="#">navar.aSep08</a>		1717	713		75	putative secreted or extracellular protein precursor (8.5 kD) (navar) mRNA.
<a href="#">nawby</a>	<a href="#">nawby.aSep08</a>		1172	423	3	74	arrestin 3 retinal (nawby) alternative variant aSep08, mRNA.
<a href="#">nawchy</a>	<a href="#">nawchy.aSep08</a>		3322	305		69	ATP GTP binding protein-like 2 (nawchy) mRNA.
<a href="#">nawdar</a>	<a href="#">nawdar.aSep08</a>		861	686		110	hook homolog 2 (nawdar) mRNA.
<a href="#">nawey</a>	<a href="#">nawey.aSep08</a>		2926	525		102	putative protein (nawey) mRNA.
<a href="#">nawfer</a>	<a href="#">nawfer.bSep08</a>		2742	438		43	putative protein (4.9 kD) (nawfer) alternative variant bSep08, mRNA.
<a href="#">nawflo</a>	<a href="#">nawflo.aSep08</a>		818	467		67	putative protein (nawflo) mRNA.
<a href="#">nawflu</a>	<a href="#">nawflu.aSep08</a>		6738	747		66	putative protein (7.8 kD) (nawflu) mRNA.
<a href="#">nawgar</a>	<a href="#">nawgar.aSep08</a>		4916	339		113	leucine-rich repeats transmembrane domains 1 (nawgar) mRNA.
<a href="#">nawja</a>	<a href="#">nawja.aSep08</a>		1136	699		43	putative protein (nawja) mRNA.
<a href="#">nawjey</a>	<a href="#">nawjey.aSep08</a>		5391	586	4	87	putative protein (10.0 kD) (nawjey) alternative variant aSep08, mRNA.
<a href="#">nawjey</a>	<a href="#">nawjey.bSep08</a>		14071	672	3	40	putative protein (nawjey) alternative variant bSep08, mRNA.
<a href="#">nawkee</a>	<a href="#">nawkee.aSep08</a>		2227	443		48	putative protein (nawkee) mRNA.
<a href="#">nawkler</a>	<a href="#">nawkler.aSep08</a>		639	140		34	putative protein (nawkler) mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.aSep08</a>		8629	2628	6	368	CRA c (nawloy) alternative variant aSep08, mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.bSep08</a>		5728	757	5	252	CRA c (nawloy) alternative variant bSep08, mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.cSep08</a>		5970	710	5	189	CRA c (nawloy) alternative variant cSep08, mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.dSep08</a>		7241	1057	6	150	CRA c (16.1 kD) (nawloy) alternative variant dSep08, complete mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.eSep08</a>		6292	621	5	121	putative protein (nawloy) alternative variant eSep08, mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.fSep08</a>		1124	639	2	95	CRA a like (nawloy) alternative variant fSep08, mRNA.
<a href="#">nawloy</a>	<a href="#">nawloy.hSep08</a>		2670	569	3	76	putative protein (nawloy) alternative variant hSep08, mRNA.
<a href="#">nawmee</a>	<a href="#">nawmee.aSep08</a>		2132	603	3	29	putative protein (nawmee) alternative variant aSep08, mRNA.
<a href="#">nawmee</a>	<a href="#">nawmee.bSep08</a>		606	420	2	29	putative protein (nawmee) alternative variant bSep08, mRNA.
<a href="#">nawmer</a>	<a href="#">nawmer.aSep08</a>		1757	581		193	CRA b (nawmer) mRNA.
<a href="#">nawnoy</a>	<a href="#">nawnoy.bSep08</a>		33053	417	3	38	putative protein (4.2 kD) (nawnoy) alternative variant bSep08, mRNA.
<a href="#">nawpor</a>	<a href="#">nawpor.aSep08</a>		3114	1259		71	putative protein of bilateral origin (nawpor) mRNA.
<a href="#">nawsa</a>	<a href="#">nawsa.aSep08</a>		9846	1839	4	318	cysteine histidine rich 1 (33.8 kD) (nawsa) alternative variant aSep08, mRNA.
<a href="#">nawsa</a>	<a href="#">nawsa.bSep08</a>		9705	1624	4	287	cysteine histidine rich 1 (nawsa) alternative variant bSep08, mRNA.
<a href="#">nawsa</a>	<a href="#">nawsa.cSep08</a>		5502	3356	3	264	histidine-rich 1 (nawsa) alternative variant cSep08, mRNA.
<a href="#">nawsa</a>	<a href="#">nawsa.dSep08</a>		1806	1178	2	171	cysteine histidine rich 1 (19.4 kD) (nawsa) alternative variant dSep08, mRNA.

<a href="#">nawshee</a>	<a href="#">nawshee.aSep08</a>		17116	221		28	putative protein (nawshee) mRNA.
<a href="#">nawtu</a>	<a href="#">nawtu.aSep08</a>		3259	199		63	gag protein like (nawtu) mRNA.
<a href="#">nawvar</a>	<a href="#">nawvar.aSep08</a>		3154	513		104	putative protein (nawvar) mRNA.
<a href="#">nawwey</a>	<a href="#">nawwey.aSep08</a>		535	376		53	putative protein (6.2 kD) (nawwey) complete mRNA.
<a href="#">Nbeal1</a>	<a href="#">Nbeal1.aSep08</a>	<a href="#">689253</a>	2582	627		52	neurobeachin like 1 (Nbeal1) mRNA.
<a href="#">Nbeal2</a>	<a href="#">Nbeal2.aSep08</a>	<a href="#">316014</a>	3540	1558	10	459	neurobeachin-like 2 (Nbeal2) alternative variant aSep08, mRNA.
<a href="#">Nbeal2</a>	<a href="#">Nbeal2.bSep08</a>	<a href="#">316014</a>	942	730	3	165	neurobeachin-like 2 (Nbeal2) alternative variant bSep08, mRNA.
<a href="#">Nbeal2</a>	<a href="#">Nbeal2.cSep08</a>	<a href="#">316014</a>	750	669	2	158	neurobeachin-like 2 (Nbeal2) alternative variant cSep08, mRNA.
<a href="#">Nbl1</a>	<a href="#">Nbl1.bSep08</a>	<a href="#">50594</a>	10192	659		108	neuroblastoma, suppression of tumorigenicity 1 (Nbl1) alternative variant bSep08, mRNA.
<a href="#">Nbr1</a>	<a href="#">Nbr1.aSep08</a>	<a href="#">303554</a>	10406	2513	5	436	neighbor of Brca1 gene 1 (Nbr1) alternative variant aSep08, mRNA.
<a href="#">Nbr1</a>	<a href="#">Nbr1.bSep08</a>	<a href="#">303554</a>	5995	818	4	184	neighbor of Brca1 gene 1 (Nbr1) alternative variant bSep08, mRNA.
<a href="#">Ncald</a>	<a href="#">Ncald.aSep08</a>	<a href="#">553106</a>	39719	806		156	neurocalcin delta (16.5 kD) (Ncald) mRNA.
<a href="#">Ncam1</a>	<a href="#">Ncam1.bSep08</a>	<a href="#">24586</a>	18012	747	5	248	neural cell adhesion molecule 1 (Ncam1) alternative variant bSep08, mRNA.
<a href="#">Ncam1</a>	<a href="#">Ncam1.cSep08</a>	<a href="#">24586</a>	25948	524	5	145	neural cell adhesion molecule 1 (Ncam1) alternative variant cSep08, mRNA.
<a href="#">Ncapd2</a>	<a href="#">Ncapd2.aSep08</a>	<a href="#">362438</a>	17668	4148	25	1305	non-SMC condensin I complex, subunit D2 (Ncapd2) alternative variant aSep08, mRNA.
<a href="#">Ncapd2</a>	<a href="#">Ncapd2.bSep08</a>	<a href="#">362438</a>	2287	766	5	254	non-SMC condensin I complex, subunit D2 (Ncapd2) alternative variant bSep08, mRNA.
<a href="#">Ncapd2</a>	<a href="#">Ncapd2.cSep08</a>	<a href="#">362438</a>	2588	1235	2	223	non-SMC condensin I complex, subunit D2 (25.3 kD) (Ncapd2) alternative variant cSep08, mRNA.
<a href="#">Ncapd3</a>	<a href="#">Ncapd3.bSep08</a>	<a href="#">315508</a>	4740	512	5	91	non-SMC condensin II complex, subunit D3 (Ncapd3) alternative variant bSep08, mRNA.
<a href="#">Ncapd3</a>	<a href="#">Ncapd3.dSep08</a>	<a href="#">315508</a>	863	776	2	36	non-SMC condensin II complex, subunit D3 (Ncapd3) alternative variant dSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.aSep08</a>	<a href="#">680002</a>	22717	2956	2	654	condensin complex (Ncaph) alternative variant aSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.aSep08</a>	<a href="#">680089</a>	22717	2956	2	654	condensin complex (Ncaph) alternative variant aSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.bSep08</a>	<a href="#">680002</a>	19759	1787		560	condensin complex (Ncaph) alternative variant bSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.bSep08</a>	<a href="#">680089</a>	19759	1787		560	condensin complex (Ncaph) alternative variant bSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.cSep08</a>	<a href="#">680002</a>	8161	951	2	178	condensin complex (Ncaph) alternative variant cSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.cSep08</a>	<a href="#">680089</a>	8161	951	2	178	condensin complex (Ncaph) alternative variant cSep08, mRNA.
<a href="#">Ncaph</a>	<a href="#">Ncaph.dSep08</a>	<a href="#">680002</a>	4216	773	1	124	condensin complex (Ncaph) alternative variant dSep08, mRNA.

<a href="#">Ncaph</a>	<a href="#">Ncaph.dSep08</a>	<a href="#">680089</a>	4216	773	1	124	condensin complex (Ncaph) alternative variant dSep08, mRNA.
<a href="#">Ncaph2</a>	<a href="#">Ncaph2.bSep08</a>	<a href="#">300149</a>	8627	784	8	241	non-SMC condensin II complex, subunit H2 (Ncaph2) alternative variant bSep08, mRNA.
<a href="#">Ncaph2</a>	<a href="#">Ncaph2.cSep08</a>	<a href="#">300149</a>	8074	670	7	206	non-SMC condensin II complex, subunit H2 (Ncaph2) alternative variant cSep08, mRNA.
<a href="#">Ncbp1</a>	<a href="#">Ncbp1.bSep08</a>	<a href="#">298075</a>	8471	1911	9	280	nuclear cap binding protein subunit 1, 80kDa (32.9 kD) (Ncbp1) alternative variant bSep08, mRNA.
<a href="#">Ncbp1</a>	<a href="#">Ncbp1.cSep08</a>	<a href="#">298075</a>	1808	728	2	72	nuclear cap binding protein subunit 1, 80kDa (8.5 kD) (Ncbp1) alternative variant cSep08, mRNA.
<a href="#">Ncdn</a>	<a href="#">Ncdn.bSep08</a>	<a href="#">89791</a>	3032	688	3	228	neurochondrin (Ncdn) alternative variant bSep08, mRNA.
<a href="#">Ncdn</a>	<a href="#">Ncdn.cSep08</a>	<a href="#">89791</a>	2112	572	3	129	neurochondrin (Ncdn) alternative variant cSep08, mRNA.
<a href="#">Nckap1</a>	<a href="#">Nckap1.bSep08</a>	<a href="#">58823</a>	29115	855	8	285	NCK-associated protein 1 (Nckap1) alternative variant bSep08, mRNA.
<a href="#">Nckap1</a>	<a href="#">Nckap1.cSep08</a>	<a href="#">58823</a>	1249	757	2	42	NCK-associated protein 1 (Nckap1) alternative variant cSep08, mRNA.
<a href="#">Nckap1l</a>	<a href="#">Nckap1l.bSep08</a>	<a href="#">315348</a>	34898	1782	14	514	NCK associated protein 1 like (Nckap1l) alternative variant bSep08, mRNA.
<a href="#">Nckap1l</a>	<a href="#">Nckap1l.cSep08</a>	<a href="#">315348</a>	9742	778	6	207	NCK associated protein 1 like (Nckap1l) alternative variant cSep08, mRNA.
<a href="#">Nckap1l</a>	<a href="#">Nckap1l.dSep08</a>	<a href="#">315348</a>	14922	282	3	71	NCK associated protein 1 like (Nckap1l) alternative variant dSep08, mRNA.
<a href="#">Nckipsd</a>	<a href="#">Nckipsd.bSep08</a>	<a href="#">301009</a>	1526	753	7	251	NCK interacting protein with SH3 domain (Nckipsd) alternative variant bSep08, mRNA.
<a href="#">Nckipsd</a>	<a href="#">Nckipsd.cSep08</a>	<a href="#">301009</a>	2770	614	3	169	NCK interacting protein with SH3 domain (Nckipsd) alternative variant cSep08, mRNA.
<a href="#">Ncl</a>	<a href="#">Ncl.bSep08</a>	<a href="#">25135</a>	2849	821	6	273	nucleolin (Ncl) alternative variant bSep08, mRNA.
<a href="#">Ncln</a>	<a href="#">Ncln.bSep08</a>	<a href="#">314648</a>	2671	1231	4	262	nicalin (Ncln) alternative variant bSep08, mRNA.
<a href="#">Ncln</a>	<a href="#">Ncln.cSep08</a>	<a href="#">314648</a>	2032	522	5	174	nicalin (Ncln) alternative variant cSep08, mRNA.
<a href="#">Ncln</a>	<a href="#">Ncln.dSep08</a>	<a href="#">314648</a>	1693	228	3	65	nicalin (Ncln) alternative variant dSep08, mRNA.
<a href="#">Ncln</a>	<a href="#">Ncln.eSep08</a>	<a href="#">314648</a>	1547	1096	4	65	nicalin (Ncln) alternative variant eSep08, mRNA.
<a href="#">Ncln</a>	<a href="#">Ncln.fSep08</a>	<a href="#">314648</a>	1842	1558	2	45	nicalin (Ncln) alternative variant fSep08, mRNA.
<a href="#">Ncoa1</a>	<a href="#">Ncoa1.bSep08</a>	<a href="#">313929</a>	54029	4550	11	803	nuclear receptor coactivator 1 (Ncoa1) alternative variant bSep08, mRNA.
<a href="#">Ncoa1</a>	<a href="#">Ncoa1.cSep08</a>	<a href="#">313929</a>	19784	1207	6	293	nuclear receptor coactivator 1 (Ncoa1) alternative variant cSep08, mRNA.
<a href="#">Ncoa1</a>	<a href="#">Ncoa1.dSep08</a>	<a href="#">313929</a>	13193	692	5	122	nuclear receptor coactivator 1 (Ncoa1) alternative variant dSep08, mRNA.
<a href="#">Ncoa1</a>	<a href="#">Ncoa1.fSep08</a>	<a href="#">313929</a>	155904	645	5	67	nuclear receptor coactivator 1 (Ncoa1) alternative variant fSep08, mRNA.
<a href="#">Ncoa2</a>	<a href="#">Ncoa2.aSep08</a>	<a href="#">83724</a>	194058	824	7	222	nuclear receptor coactivator 2 (Ncoa2) alternative variant aSep08, mRNA.
<a href="#">Ncoa2</a>	<a href="#">Ncoa2.aSep08</a>	<a href="#">680234</a>	194058	824	7	222	nuclear receptor coactivator 2 (Ncoa2) alternative variant aSep08, mRNA.
<a href="#">Ncoa2</a>	<a href="#">Ncoa2.bSep08</a>	<a href="#">83724</a>	154868	1002	3	44	putative protein (Ncoa2) alternative variant bSep08, mRNA.

<a href="#">Ncoa2</a>	<a href="#">Ncoa2.bSep08</a>	<a href="#">680234</a>	154868	1002	3	44	putative protein (Ncoa2) alternative variant bSep08, mRNA.
<a href="#">Ncoa2</a>	<a href="#">Ncoa2.dSep08</a>	<a href="#">83724</a>	89122	526	2	43	putative protein (Ncoa2) alternative variant dSep08, mRNA.
<a href="#">Ncoa2</a>	<a href="#">Ncoa2.dSep08</a>	<a href="#">680234</a>	89122	526	2	43	putative protein (Ncoa2) alternative variant dSep08, mRNA.
<a href="#">Ncoa3</a>	<a href="#">Ncoa3.aSep08</a>	<a href="#">84584</a>	18270	4790		582	nuclear receptor coactivator 3 (Ncoa3) mRNA.
<a href="#">Ncoa5</a>	<a href="#">Ncoa5.bSep08</a>	<a href="#">296372</a>	3520	783	3	174	nuclear receptor coactivator 5 (Ncoa5) alternative variant bSep08, mRNA.
<a href="#">Ncoa5</a>	<a href="#">Ncoa5.cSep08</a>	<a href="#">296372</a>	2206	1136	2	148	nuclear receptor coactivator 5 (Ncoa5) alternative variant cSep08, mRNA.
<a href="#">Ncoa5</a>	<a href="#">Ncoa5.dSep08</a>	<a href="#">296372</a>	12095	1013	3	85	nuclear receptor coactivator 5 (9.5 kD) (Ncoa5) alternative variant dSep08, mRNA.
<a href="#">Ncoa6</a>	<a href="#">Ncoa6.aSep08</a>	<a href="#">116464</a>	16534	3494	5	989	nuclear receptor coactivator 6 (Ncoa6) alternative variant aSep08, mRNA.
<a href="#">Ncoa6</a>	<a href="#">Ncoa6.bSep08</a>	<a href="#">116464</a>	8257	1018	3	104	nuclear receptor coactivator 6 (11.4 kD) (Ncoa6) alternative variant bSep08, mRNA.
<a href="#">Ncoa6</a>	<a href="#">Ncoa6.cSep08</a>	<a href="#">116464</a>	7970	728	3	70	nuclear receptor coactivator 6 (Ncoa6) alternative variant cSep08, mRNA.
<a href="#">Ncoa7</a>	<a href="#">Ncoa7.aSep08</a>	<a href="#">498995</a>	38144	397		132	nuclear receptor coactivator 7 (Ncoa7) mRNA.
<a href="#">Ncor1</a>	<a href="#">Ncor1.aSep08</a>	<a href="#">54299</a>	45305	1198	11	398	nuclear receptor co-repressor 1 (Ncor1) alternative variant aSep08, mRNA.
<a href="#">Ncor1</a>	<a href="#">Ncor1.bSep08</a>	<a href="#">54299</a>	18778	378	6	126	nuclear receptor co-repressor 1 (Ncor1) alternative variant bSep08, mRNA.
<a href="#">Ncor1</a>	<a href="#">Ncor1.cSep08</a>	<a href="#">54299</a>	27682	1577	4	108	nuclear receptor co-repressor 1 (11.9 kD) (Ncor1) alternative variant cSep08, mRNA.
<a href="#">Ncor2</a>	<a href="#">Ncor2.bSep08</a>	<a href="#">360801</a>	9051	2027	10	562	nuclear receptor co-repressor 2 (Ncor2) alternative variant bSep08, mRNA.
<a href="#">Ncor2</a>	<a href="#">Ncor2.cSep08</a>	<a href="#">360801</a>	9194	1816	8	484	nuclear receptor co-repressor 2 (Ncor2) alternative variant cSep08, mRNA.
<a href="#">Ncor2</a>	<a href="#">Ncor2.dSep08</a>	<a href="#">360801</a>	4836	707	6	224	nuclear receptor co-repressor 2 (Ncor2) alternative variant dSep08, mRNA.
<a href="#">Ncor2</a>	<a href="#">Ncor2.eSep08</a>	<a href="#">360801</a>	2007	418	3	139	nuclear receptor co-repressor 2 (Ncor2) alternative variant eSep08, mRNA.
<a href="#">Ncor2</a>	<a href="#">Ncor2.iSep08</a>	<a href="#">360801</a>	54029	572	3	61	nuclear receptor co-repressor 2 (6.3 kD) (Ncor2) alternative variant iSep08, mRNA.
<a href="#">Ncstn</a>	<a href="#">Ncstn.bSep08</a>	<a href="#">289231</a>	15991	2748	16	602	nicastrin (67.2 kD) (Ncstn) alternative variant bSep08, mRNA.
<a href="#">Ncstn</a>	<a href="#">Ncstn.cSep08</a>	<a href="#">289231</a>	8698	936	7	150	nicastrin (Ncstn) alternative variant cSep08, mRNA.
<a href="#">Ncstn</a>	<a href="#">Ncstn.dSep08</a>	<a href="#">289231</a>	1549	703	3	140	nicastrin (Ncstn) alternative variant dSep08, mRNA.
<a href="#">Ncstn</a>	<a href="#">Ncstn.eSep08</a>	<a href="#">289231</a>	2188	989	5	83	nicastrin (Ncstn) alternative variant eSep08, mRNA.
<a href="#">Ncstn</a>	<a href="#">Ncstn.fSep08</a>	<a href="#">289231</a>	483	327	2	69	nicastrin (Ncstn) alternative variant fSep08, mRNA.
<a href="#">Ndc80</a>	<a href="#">Ndc80.bSep08</a>	<a href="#">301701</a>	16744	1783	1	384	NDC80 homolog, kinetochore complex component (S. cerevisiae) (Ndc80) alternative variant bSep08, mRNA.
<a href="#">Ndel1</a>	<a href="#">Ndel1.aSep08</a>	<a href="#">170845</a>	33779	2357	10	378	nuclear distribution gene E-like homolog 1 (A. nidulans) (Ndel1) alternative variant aSep08, mRNA.

<a href="#">Ndel1</a>	<a href="#">Ndel1.cSep08</a>	<a href="#">170845</a>	53024	779	5	162	nuclear distribution gene E-like homolog 1 ( <i>A. nidulans</i> ) (Ndel1) alternative variant cSep08, mRNA.
<a href="#">Ndel1</a>	<a href="#">Ndel1.dSep08</a>	<a href="#">170845</a>	16701	733	5	147	nuclear distribution gene E-like homolog 1 ( <i>A. nidulans</i> ) (Ndel1) alternative variant dSep08, mRNA.
<a href="#">Ndfip1</a>	<a href="#">Ndfip1.bSep08</a>	<a href="#">291609</a>	50210	1756	8	263	nedd4 family interacting protein 1 (Ndfip1) alternative variant bSep08, mRNA.
<a href="#">Ndfip1</a>	<a href="#">Ndfip1.cSep08</a>	<a href="#">291609</a>	31102	412	3	137	nedd4 family interacting protein 1 (Ndfip1) alternative variant cSep08, mRNA.
<a href="#">Ndfip1</a>	<a href="#">Ndfip1.dSep08</a>	<a href="#">291609</a>	5421	1138	2	66	nedd4 family interacting protein 1 (7.5 kD) (Ndfip1) alternative variant dSep08, mRNA.
<a href="#">Ndfip1</a>	<a href="#">Ndfip1.eSep08</a>	<a href="#">291609</a>	7942	294	3	33	nedd4 family interacting protein 1 (3.9 kD) (Ndfip1) alternative variant eSep08, mRNA.
<a href="#">Ndfip1</a>	<a href="#">Ndfip1.fSep08</a>	<a href="#">291609</a>	6178	194	2	34	nedd4 family interacting protein 1 (3.6 kD) (Ndfip1) alternative variant fSep08, mRNA.
<a href="#">Ndfip2</a>	<a href="#">Ndfip2.aSep08</a>	<a href="#">361089</a>	59154	2108	1	337	nedd4 family interacting protein 2 (36.3 kD) (Ndfip2) alternative variant aSep08, mRNA.
<a href="#">Ndg2</a>	<a href="#">Ndg2.bSep08</a>	<a href="#">361824</a>	2321	1327	2	92	nur77 downstream gene 2 (Ndg2) alternative variant bSep08, mRNA.
<a href="#">Ndor1</a>	<a href="#">Ndor1.bSep08</a>	<a href="#">311799</a>	954	744	3	159	NADPH dependent diflavin oxidoreductase 1 (Ndor1) alternative variant bSep08, mRNA.
<a href="#">Ndor1</a>	<a href="#">Ndor1.cSep08</a>	<a href="#">311799</a>	5133	700	3	58	NADPH dependent diflavin oxidoreductase 1 (6.8 kD) (Ndor1) alternative variant cSep08, mRNA.
<a href="#">Ndph</a>	<a href="#">Ndph.bSep08</a>	<a href="#">363443</a>	24450	1547		89	norrie disease homolog (9.5 kD) (Ndph) alternative variant bSep08, mRNA.
<a href="#">Ndrg1</a>	<a href="#">Ndrg1.aSep08</a>	<a href="#">299923</a>	35180	2882	1	441	N-myc downstream regulated gene 1 (Ndrg1) alternative variant aSep08, mRNA.
<a href="#">Ndrg1</a>	<a href="#">Ndrg1.bSep08</a>	<a href="#">299923</a>	32902	1028	1	232	N-myc downstream regulated gene 1 (Ndrg1) alternative variant bSep08, mRNA.
<a href="#">Ndrg1</a>	<a href="#">Ndrg1.cSep08</a>	<a href="#">299923</a>	29232	704	1	198	N-myc downstream regulated gene 1 (Ndrg1) alternative variant cSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.bSep08</a>	<a href="#">171114</a>	8367	1718	15	359	ndrg family member 2 (39.5 kD) (Ndrg2) alternative variant bSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.cSep08</a>	<a href="#">171114</a>	6890	2092	14	341	ndrg family member 2 (37.5 kD) (Ndrg2) alternative variant cSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.dSep08</a>	<a href="#">171114</a>	6076	849	12	257	ndrg 2 (Ndrg2) alternative variant dSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.eSep08</a>	<a href="#">171114</a>	6094	825	11	257	ndrg family member 2 (Ndrg2) alternative variant eSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.fSep08</a>	<a href="#">171114</a>	5020	599	8	184	ndrg family member 2 (Ndrg2) alternative variant fSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.gSep08</a>	<a href="#">171114</a>	4307	682	9	151	ndrg family member 2 CRA c (Ndrg2) alternative variant gSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.hSep08</a>	<a href="#">171114</a>	2721	403	2	105	putative protein (Ndrg2) alternative variant hSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.iSep08</a>	<a href="#">171114</a>	2597	429	4	100	ndrg family member 2 (Ndrg2) alternative variant iSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.jSep08</a>	<a href="#">171114</a>	834	703	2	49	ndrg family member 2 (Ndrg2) alternative variant jSep08, mRNA.
<a href="#">Ndrg2</a>	<a href="#">Ndrg2.lSep08</a>	<a href="#">171114</a>	1141	749	2	51	putative protein (Ndrg2) alternative variant lSep08, mRNA.

<a href="#">Ndrg2</a>	<a href="#">Ndrg2.mSep08</a>	<a href="#">171114</a>	804	414	3	53	putative protein (6.6 kD) (Ndrg2) alternative variant mSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.aSep08</a>	<a href="#">296318</a>	28593	1453	13	324	N-myc downstream regulated gene 3 (Ndrg3) alternative variant aSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.bSep08</a>	<a href="#">296318</a>	20178	2138	8	190	N-myc downstream regulated gene 3 (21.0 kD) (Ndrg3) alternative variant bSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.cSep08</a>	<a href="#">296318</a>	18965	925	8	185	N-myc downstream regulated gene 3 (Ndrg3) alternative variant cSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.dSep08</a>	<a href="#">296318</a>	13574	471	8	156	N-myc downstream regulated gene 3 (Ndrg3) alternative variant dSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.eSep08</a>	<a href="#">296318</a>	15880	716	6	153	N-myc downstream regulated gene 3 (16.7 kD) (Ndrg3) alternative variant eSep08, mRNA.
<a href="#">Ndrg3</a>	<a href="#">Ndrg3.fSep08</a>	<a href="#">296318</a>	5477	729	3	85	N-myc downstream regulated gene 3 (Ndrg3) alternative variant fSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.aSep08</a>	<a href="#">64457</a>	11048	2810	15	352	N-myc downstream regulated gene 4 (38.5 kD) (Ndrg4) alternative variant aSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.bSep08</a>	<a href="#">64457</a>	17253	691	9	229	brain heart protein (Ndrg4) alternative variant bSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.cSep08</a>	<a href="#">64457</a>	4308	795	9	206	brain heart protein (Ndrg4) alternative variant cSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.eSep08</a>	<a href="#">64457</a>	2235	739	8	184	brain heart protein (Ndrg4) alternative variant eSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.fSep08</a>	<a href="#">64457</a>	1519	772	2	118	N-myc downstream regulated gene 4 (12.5 kD) (Ndrg4) alternative variant fSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.gSep08</a>	<a href="#">64457</a>	2754	454	4	107	N-myc downstream regulated gene 4 (Ndrg4) alternative variant gSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.iSep08</a>	<a href="#">64457</a>	28341	367	5	88	brain heart protein (Ndrg4) alternative variant iSep08, mRNA.
<a href="#">Ndrg4</a>	<a href="#">Ndrg4.jSep08</a>	<a href="#">64457</a>	1522	437	4	47	N-myc downstream regulated gene 4 (Ndrg4) alternative variant jSep08, mRNA.
<a href="#">Ndst2</a>	<a href="#">Ndst2.bSep08</a>	<a href="#">114002</a>	2806	1467	5	165	N-deacetylase N-sulfotransferase 2 (Ndst2) alternative variant bSep08, mRNA.
<a href="#">Ndst2</a>	<a href="#">Ndst2.cSep08</a>	<a href="#">114002</a>	914	789	2	71	putative protein (7.6 kD) (Ndst2) alternative variant cSep08, mRNA.
<a href="#">Ndst4</a>	<a href="#">Ndst4.aSep08</a>	<a href="#">362035</a>	11856	632		145	N-deacetylase/N-sulfotransferase (heparin glucosaminy) 4 (Ndst4) mRNA.
<a href="#">NDT80_Phog.0</a>	<a href="#">NDT80_Phog.0.aSep08</a>		10386	735	2	206	NDT80/Phog like DNA-binding (NDT80_Phog.0) alternative variant aSep08, mRNA.
<a href="#">NDT80_Phog.0</a>	<a href="#">NDT80_Phog.0.bSep08</a>		9784	430	1	142	CRA b (NDT80_Phog.0) alternative variant bSep08, mRNA.
<a href="#">Ndufa1</a>	<a href="#">Ndufa1.bSep08</a>	<a href="#">363441</a>	3612	2541	1	66	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.7 kD) (Ndufa1) alternative variant bSep08, complete mRNA.
<a href="#">Ndufa3</a>	<a href="#">Ndufa3.bSep08</a>	<a href="#">691001</a>	2643	359	4	84	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9.4 kD) (Ndufa3) alternative variant bSep08, complete mRNA.

<a href="#">Ndufa3</a>	<a href="#">Ndufa3.dSep08</a>	<a href="#">691001</a>	2602	527	3	53	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (6.0 kD) (Ndufa3) alternative variant dSep08, complete mRNA.
<a href="#">Ndufa5</a>	<a href="#">Ndufa5.bSep08</a>	<a href="#">25488</a>	1875	1544	2	78	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 5 (8.9 kD) (Ndufa5) alternative variant bSep08, mRNA.
<a href="#">Ndufa7</a>	<a href="#">Ndufa7.aSep08</a>	<a href="#">299643</a>	13311	1007	2	120	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a) (Ndufa7) alternative variant aSep08, mRNA.
<a href="#">Ndufa8</a>	<a href="#">Ndufa8.bSep08</a>	<a href="#">296658</a>	16452	1248	4	172	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8 (20.0 kD) (Ndufa8) alternative variant bSep08, mRNA.
<a href="#">Ndufa8</a>	<a href="#">Ndufa8.cSep08</a>	<a href="#">296658</a>	15901	693	4	117	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8 (Ndufa8) alternative variant cSep08, mRNA.
<a href="#">Ndufa10</a>	<a href="#">Ndufa10.cSep08</a>	<a href="#">678759</a>	1758	318	2	46	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 10 (Ndufa10) alternative variant cSep08, mRNA.
<a href="#">Ndufa10I1</a>	<a href="#">Ndufa10I1.bSep08</a>	<a href="#">314071</a>	1937	1803	2	273	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 10-like 1 (31.8 kD) (Ndufa10I1) alternative variant bSep08, complete mRNA.
<a href="#">Ndufa10I1</a>	<a href="#">Ndufa10I1.bSep08</a>	<a href="#">316632</a>	1937	1803	2	273	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 10-like 1 (31.8 kD) (Ndufa10I1) alternative variant bSep08, complete mRNA.
<a href="#">Ndufa13-ps</a>	<a href="#">Ndufa13-ps.aSep08</a>	<a href="#">290671</a>	6977	513	4	144	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 13 pseudogene (16.8 kD) (Ndufa13-ps) alternative variant aSep08, complete mRNA.
<a href="#">Ndufa13-ps</a>	<a href="#">Ndufa13-ps.bSep08</a>	<a href="#">290671</a>	1293	547	3	110	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 13 pseudogene (13.1 kD) (Ndufa13-ps) alternative variant bSep08, mRNA.
<a href="#">Ndufab1</a>	<a href="#">Ndufab1.aSep08</a>	<a href="#">293453</a>	9733	642	5	165	NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (Ndufab1) alternative variant aSep08, mRNA.
<a href="#">Ndufab1</a>	<a href="#">Ndufab1.cSep08</a>	<a href="#">293453</a>	9283	849	4	132	NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (Ndufab1) alternative variant cSep08, mRNA.
<a href="#">Ndufaf2</a>	<a href="#">Ndufaf2.aSep08</a>	<a href="#">361894</a>	113420	602		166	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, assembly factor 2 (19.6 kD) (Ndufaf2) mRNA.
<a href="#">Ndufb2</a>	<a href="#">Ndufb2.aSep08</a>	<a href="#">362344</a>	7079	456	4	105	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (Ndufb2) alternative variant aSep08, complete mRNA.
<a href="#">Ndufb3</a>	<a href="#">Ndufb3.bSep08</a>	<a href="#">301427</a>	10177	533		58	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 3 (Ndufb3) alternative variant bSep08, mRNA.
<a href="#">Ndufb8</a>	<a href="#">Ndufb8.bSep08</a>	<a href="#">293991</a>	4669	435	5	144	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 8 (Ndufb8) alternative variant bSep08, mRNA.
<a href="#">Ndufb8</a>	<a href="#">Ndufb8.cSep08</a>	<a href="#">293991</a>	4384	1493	4	110	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 8 (13.3 kD) (Ndufb8) alternative variant cSep08, mRNA.
<a href="#">Ndufb8</a>	<a href="#">Ndufb8.dSep08</a>	<a href="#">293991</a>	3730	840	3	110	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 8 (13.3 kD) (Ndufb8) alternative variant dSep08, mRNA.
<a href="#">Ndufb9</a>	<a href="#">Ndufb9.bSep08</a>	<a href="#">299954</a>	6370	649	4	175	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9 (21.4 kD) (Ndufb9) alternative variant bSep08, complete mRNA.



<a href="#">Ndufb9</a>	<a href="#">Ndufb9.cSep08</a>	<a href="#">299954</a>	424	278	2	46	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9 (5.6 kD) (Ndufb9) alternative variant cSep08, mRNA.
<a href="#">Ndufb11</a>	<a href="#">Ndufb11.bSep08</a>	<a href="#">299310</a>	2258	1016	2	125	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11 (14.6 kD) (Ndufb11) alternative variant bSep08, complete mRNA.
<a href="#">Ndufc1</a>	<a href="#">Ndufc1.aSep08</a>	<a href="#">689938</a>	3787	528	4	82	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (Ndufc1) alternative variant aSep08, mRNA.
<a href="#">Ndufc1</a>	<a href="#">Ndufc1.bSep08</a>	<a href="#">689938</a>	3788	569	5	80	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (Ndufc1) alternative variant bSep08, mRNA.
<a href="#">Ndufc2</a>	<a href="#">Ndufc2.bSep08</a>	<a href="#">293130</a>	6230	623	3	94	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (10.3 kD) (Ndufc2) alternative variant bSep08, complete mRNA.
<a href="#">Ndufs1</a>	<a href="#">Ndufs1.bSep08</a>	<a href="#">301458</a>	34022	1788	12	431	NADH dehydrogenase (ubiquinone) Fe-S protein 1 (Ndufs1) alternative variant bSep08, complete mRNA.
<a href="#">Ndufs1</a>	<a href="#">Ndufs1.cSep08</a>	<a href="#">301458</a>	5178	524	4	152	NADH dehydrogenase (ubiquinone) Fe-S protein 1 (Ndufs1) alternative variant cSep08, mRNA.
<a href="#">Ndufs2</a>	<a href="#">Ndufs2.bSep08</a>	<a href="#">289218</a>	2804	753	5	185	NADH dehydrogenase (ubiquinone) Fe-S protein 2 (Ndufs2) alternative variant bSep08, mRNA.
<a href="#">Ndufs2</a>	<a href="#">Ndufs2.cSep08</a>	<a href="#">289218</a>	8610	531	4	109	NADH dehydrogenase (ubiquinone) Fe-S protein 2 (Ndufs2) alternative variant cSep08, mRNA.
<a href="#">Ndufs2</a>	<a href="#">Ndufs2.dSep08</a>	<a href="#">289218</a>	8644	578	4	46	NADH dehydrogenase (ubiquinone) Fe-S protein 2 (Ndufs2) alternative variant dSep08, mRNA.
<a href="#">Ndufs3</a>	<a href="#">Ndufs3.bSep08</a>	<a href="#">295923</a>	4639	574	6	191	NADH dehydrogenase (ubiquinone) Fe-S protein 3 (Ndufs3) alternative variant bSep08, mRNA.
<a href="#">Ndufs3</a>	<a href="#">Ndufs3.cSep08</a>	<a href="#">295923</a>	5538	1548	4	92	NADH dehydrogenase (ubiquinone) Fe-S protein 3 (10.9 kD) (Ndufs3) alternative variant cSep08, mRNA.
<a href="#">Ndufs4</a>	<a href="#">Ndufs4.bSep08</a>	<a href="#">499529</a>	8018	740	1	64	NADH dehydrogenase (ubiquinone) Fe-S protein 4 (Ndufs4) alternative variant bSep08, mRNA.
<a href="#">Ndufs5</a>	<a href="#">Ndufs5.aSep08</a>	<a href="#">362588</a>	5868	436	3	106	NADH dehydrogenase (ubiquinone) Fe-S protein 5 (12.7 kD) (Ndufs5) alternative variant aSep08, mRNA.
<a href="#">Ndufs7</a>	<a href="#">Ndufs7.bSep08</a>	<a href="#">362837</a>	9749	3723	8	228	NADH dehydrogenase (ubiquinone) Fe-S protein 7 (25.1 kD) (Ndufs7) alternative variant bSep08, mRNA.
<a href="#">Ndufs7</a>	<a href="#">Ndufs7.cSep08</a>	<a href="#">362837</a>	2217	707	3	142	NADH dehydrogenase (ubiquinone) Fe-S protein 7 (Ndufs7) alternative variant cSep08, mRNA.
<a href="#">Ndufs8</a>	<a href="#">Ndufs8.aSep08</a>	<a href="#">293652</a>	3771	880	3	212	NADH dehydrogenase (ubiquinone) Fe-S protein 8 (24.0 kD) (Ndufs8) alternative variant aSep08, mRNA.
<a href="#">Ndufs8</a>	<a href="#">Ndufs8.cSep08</a>	<a href="#">293652</a>	1848	556	1	119	NADH dehydrogenase (ubiquinone) Fe-S protein 8 (Ndufs8) alternative variant cSep08, mRNA.
<a href="#">Ndufv1</a>	<a href="#">Ndufv1.bSep08</a>	<a href="#">293655</a>	3885	1121	5	232	nadh dehydrogenase flavoprotein 1 (25.6 kD) (Ndufv1) alternative variant bSep08, mRNA.
<a href="#">Ndufv1</a>	<a href="#">Ndufv1.cSep08</a>	<a href="#">293655</a>	871	655	3	73	nadh dehydrogenase flavoprotein 1 CRA a (8.3 kD) (Ndufv1) alternative variant cSep08, mRNA.
<a href="#">Ndufv2</a>	<a href="#">Ndufv2.bSep08</a>	<a href="#">81728</a>	14820	778	5	184	NADH dehydrogenase (ubiquinone) flavoprotein 2 (20.7 kD) (Ndufv2) alternative variant bSep08, complete mRNA.
<a href="#">Ndufv2</a>	<a href="#">Ndufv2.cSep08</a>	<a href="#">81728</a>	1811	432	2	31	NADH dehydrogenase (ubiquinone) flavoprotein 2 (Ndufv2) alternative variant cSep08, mRNA.

<a href="#">Ndufv3l</a>	<a href="#">Ndufv3l.cSep08</a>	<a href="#">64539</a>	2103	725	1	73	NADH dehydrogenase (ubiquinone) flavoprotein 3-like (7.8 kD) (Ndufv3l) alternative variant cSep08, mRNA.
<a href="#">Neb</a>	<a href="#">Neb.aSep08</a>	<a href="#">311029</a>	9110	1075	8	332	nebulin (Neb) alternative variant aSep08, mRNA.
<a href="#">Neb</a>	<a href="#">Neb.bSep08</a>	<a href="#">311029</a>	8668	1036	8	319	nebulin (Neb) alternative variant bSep08, mRNA.
<a href="#">Neb</a>	<a href="#">Neb.cSep08</a>	<a href="#">311029</a>	20318	1461	8	285	nebulin (Neb) alternative variant cSep08, mRNA.
<a href="#">Neb</a>	<a href="#">Neb.dSep08</a>	<a href="#">311029</a>	13020	749	8	249	nebulin (Neb) alternative variant dSep08, mRNA.
<a href="#">Neb</a>	<a href="#">Neb.eSep08</a>	<a href="#">311029</a>	784	454	2	56	nebulin (Neb) alternative variant eSep08, mRNA.
<a href="#">Neb1</a>	<a href="#">Neb1.aSep08</a>	<a href="#">307189</a>	74373	557	4	85	nebulette (Neb1) alternative variant aSep08, mRNA.
<a href="#">Nebulin.0</a>	<a href="#">Nebulin.0.aSep08</a>		10892	396		131	nebulette (Nebulin.0) mRNA.
<a href="#">Nebulin.1</a>	<a href="#">Nebulin.1.aSep08</a>		9857	1779		592	nebulin (Nebulin.1) alternative variant aSep08, mRNA.
<a href="#">Nebulin.1</a>	<a href="#">Nebulin.1.bSep08</a>		4409	603		201	nebulin (Nebulin.1) alternative variant bSep08, mRNA.
<a href="#">Nebulin.1</a>	<a href="#">Nebulin.1.cSep08</a>		898	200		66	putative protein of vertebrate origin (Nebulin.1) alternative variant cSep08, mRNA.
<a href="#">Nebulin.2</a>	<a href="#">Nebulin.2.aSep08</a>		1298	396		131	nebulin (Nebulin.2) mRNA.
<a href="#">Nebulin.3</a>	<a href="#">Nebulin.3.aSep08</a>		2686	374		124	nebulin (Nebulin.3) mRNA.
<a href="#">Nebulin.4</a>	<a href="#">Nebulin.4.aSep08</a>		1003	397		132	nebulin (Nebulin.4) mRNA.
<a href="#">Nebulin.5</a>	<a href="#">Nebulin.5.aSep08</a>		9906	461		153	nebulin (Nebulin.5) mRNA.
<a href="#">Necab1</a>	<a href="#">Necab1.bSep08</a>	<a href="#">64169</a>	3855	411	1	85	N-terminal EF-hand calcium binding protein 1 (Necab1) alternative variant bSep08, mRNA.
<a href="#">Necab2</a>	<a href="#">Necab2.aSep08</a>	<a href="#">170928</a>	6995	694	8	198	N-terminal EF-hand calcium binding protein 2 (22.4 kD) (Necab2) alternative variant aSep08, mRNA.
<a href="#">Necab2</a>	<a href="#">Necab2.bSep08</a>	<a href="#">170928</a>	5559	628	6	150	N-terminal EF-hand calcium binding protein 2 (16.8 kD) (Necab2) alternative variant bSep08, mRNA.
<a href="#">Necab3</a>	<a href="#">Necab3.bSep08</a>	<a href="#">311562</a>	3705	739	8	228	antibiotic biosynthesis monooxygenase (Necab3) alternative variant bSep08, mRNA.
<a href="#">Necab3</a>	<a href="#">Necab3.cSep08</a>	<a href="#">311562</a>	2822	787	4	102	putative protein of metazoan origin (Necab3) alternative variant cSep08, mRNA.
<a href="#">Necab3</a>	<a href="#">Necab3.dSep08</a>	<a href="#">311562</a>	11860	654	7	85	calcium-binding EF-hand containing protein (9.3 kD) (Necab3) alternative variant dSep08, mRNA.
<a href="#">Necab3</a>	<a href="#">Necab3.eSep08</a>	<a href="#">311562</a>	1159	711	3	65	putative protein of vertebrate origin (Necab3) alternative variant eSep08, mRNA.
<a href="#">Necap1</a>	<a href="#">Necap1.bSep08</a>	<a href="#">312694</a>	14968	1193	3	218	NECAP endocytosis associated 1 (23.0 kD) (Necap1) alternative variant bSep08, mRNA.
<a href="#">Necap2</a>	<a href="#">Necap2.aSep08</a>	<a href="#">298598</a>	11287	1180	7	286	NECAP endocytosis associated 2 (Necap2) alternative variant aSep08, mRNA.
<a href="#">Necap2</a>	<a href="#">Necap2.bSep08</a>	<a href="#">298598</a>	12386	1911	8	263	NECAP endocytosis associated 2 (Necap2) alternative variant bSep08, complete mRNA.
<a href="#">Nedd1</a>	<a href="#">Nedd1.bSep08</a>	<a href="#">299730</a>	8625	505	1	168	neural precursor cell expressed, developmentally down-regulated gene 1 (Nedd1) alternative variant bSep08, mRNA.
<a href="#">Nedd4</a>	<a href="#">Nedd4.bSep08</a>	<a href="#">25489</a>	4111	864	3	111	neural precursor cell expressed, developmentally down-regulated gene 4 (Nedd4) alternative variant bSep08, mRNA.

<a href="#">Nedd4</a>	<a href="#">Nedd4.cSep08</a>	<a href="#">25489</a>	1324	646	2	71	neural precursor cell expressed, developmentally down-regulated gene 4 (8.2 kD) (Nedd4) alternative variant cSep08, mRNA.
<a href="#">Nedd4</a>	<a href="#">Nedd4.dSep08</a>	<a href="#">25489</a>	9243	425	7	51	neural precursor cell expressed, developmentally down-regulated gene 4 (Nedd4) alternative variant dSep08, mRNA.
<a href="#">Nedd4l</a>	<a href="#">Nedd4l.bSep08</a>	<a href="#">291553</a>	157538	1129	13	355	neural precursor cell expressed, developmentally down-regulated gene 4-like (Nedd4l) alternative variant bSep08, mRNA.
<a href="#">Nedd4l</a>	<a href="#">Nedd4l.cSep08</a>	<a href="#">291553</a>	14491	389	4	129	neural precursor cell expressed, developmentally down-regulated gene 4-like (Nedd4l) alternative variant cSep08, mRNA.
<a href="#">Nedd4l</a>	<a href="#">Nedd4l.dSep08</a>	<a href="#">291553</a>	9871	375	5	125	neural precursor cell expressed, developmentally down-regulated gene 4-like (Nedd4l) alternative variant dSep08, mRNA.
<a href="#">Nedd9</a>	<a href="#">Nedd9.bSep08</a>	<a href="#">291044</a>	147526	695	3	177	neural precursor cell expressed, developmentally down-regulated gene 9 (Nedd9) alternative variant bSep08, mRNA.
<a href="#">neeby</a>	<a href="#">neeby.aSep08</a>		494	377		62	KIF4B (neeby) mRNA.
<a href="#">neechy</a>	<a href="#">neechy.aSep08</a>		3640	870		83	cytosolic carboxypeptidase 2 (neechy) mRNA.
<a href="#">needar</a>	<a href="#">needar.aSep08</a>		3033	426		141	hook homolog 2 CRA a (needar) mRNA.
<a href="#">neefer</a>	<a href="#">neefer.aSep08</a>		1065	720		70	polyprotein (8.3 kD) (neefer) mRNA.
<a href="#">neeflo</a>	<a href="#">neeflo.aSep08</a>		378	239		41	putative protein (neeflo) mRNA.
<a href="#">neeflu</a>	<a href="#">neeflu.aSep08</a>		340	97		24	putative protein (neeflu) mRNA.
<a href="#">neegar</a>	<a href="#">neegar.aSep08</a>		16410	585		41	putative protein (4.5 kD) (neegar) mRNA.
<a href="#">neeja</a>	<a href="#">neeja.aSep08</a>		16750	1634		97	putative protein (neeja) mRNA.
<a href="#">neeje</a>	<a href="#">neeje.aSep08</a>		8427	1141	4	50	putative protein (neeje) alternative variant aSep08, mRNA.
<a href="#">neeje</a>	<a href="#">neeje.bSep08</a>		7180	882	4	66	putative protein (neeje) alternative variant bSep08, mRNA.
<a href="#">neeje</a>	<a href="#">neeje.cSep08</a>		6903	781	2	39	putative protein (4.2 kD) (neeje) alternative variant cSep08, mRNA.
<a href="#">neekee</a>	<a href="#">neekee.aSep08</a>		12382	254		84	putative protein of mammalian origin (neekee) mRNA.
<a href="#">neekler</a>	<a href="#">neekler.aSep08</a>		1051	266		38	putative protein (4.1 kD) (neekler) mRNA.
<a href="#">neeloy</a>	<a href="#">neeloy.aSep08</a>		1851	1188		159	endothelin converting enzyme 2 (17.8 kD) (neeloy) mRNA.
<a href="#">neemee</a>	<a href="#">neemee.aSep08</a>		4125	961	4	83	putative mitochondrial protein (9.3 kD) (neemee) alternative variant aSep08, mRNA.
<a href="#">neemer</a>	<a href="#">neemer.aSep08</a>		1270	595		197	CRA b (neemer) mRNA.
<a href="#">neenoy</a>	<a href="#">neenoy.aSep08</a>		3229	2278		112	polyprotein (12.4 kD) (neenoy) mRNA.
<a href="#">neepor</a>	<a href="#">neepor.aSep08</a>		3492	247		57	putative protein (neepor) mRNA.
<a href="#">neesa</a>	<a href="#">neesa.aSep08</a>		720	573		65	putative protein of mammalian origin (6.7 kD) (neesa) mRNA.
<a href="#">neeshee</a>	<a href="#">neeshee.aSep08</a>		913	613		103	putative protein (neeshee) mRNA.
<a href="#">neetu</a>	<a href="#">neetu.bSep08</a>		674	622	2	11	putative protein (neetu) alternative variant bSep08, mRNA.
<a href="#">neever</a>	<a href="#">neever.aSep08</a>		925	307		66	putative protein (neever) mRNA.

<a href="#">neewey</a>	<a href="#">neewey.aSep08</a>		9482	1597	7	389	intraflagellar transport 122 homolog (43.9 kD) (neewey) alternative variant aSep08, mRNA.
<a href="#">neewey</a>	<a href="#">neewey.bSep08</a>		3523	1080	5	236	intraflagellar transport 122 homolog CRA d (neewey) alternative variant bSep08, mRNA.
<a href="#">neewey</a>	<a href="#">neewey.cSep08</a>		3406	798	4	143	intraflagellar transport 122 homolog CRA c (neewey) alternative variant cSep08, mRNA.
<a href="#">neewey</a>	<a href="#">neewey.dSep08</a>		935	495	1	94	intraflagellar transport 122 homolog CRA c (neewey) alternative variant dSep08, mRNA.
<a href="#">Nefl</a>	<a href="#">Nefl.cSep08</a>	<a href="#">83613</a>	696	313	2	71	neurofilament, light polypeptide (Nefl) alternative variant cSep08, mRNA.
<a href="#">Nefm</a>	<a href="#">Nefm.bSep08</a>	<a href="#">24588</a>	1997	579	1	151	neurofilament, medium polypeptide (Nefm) alternative variant bSep08, mRNA.
<a href="#">Neil1</a>	<a href="#">Neil1.bSep08</a>	<a href="#">367090</a>	2433	744	3	206	nei endonuclease VIII-like 1 (E. coli) (Neil1) alternative variant bSep08, mRNA.
<a href="#">Neil1</a>	<a href="#">Neil1.cSep08</a>	<a href="#">367090</a>	1566	741	4	94	nei endonuclease VIII-like 1 (E. coli) (Neil1) alternative variant cSep08, mRNA.
<a href="#">Neil1</a>	<a href="#">Neil1.dSep08</a>	<a href="#">367090</a>	1591	373	3	58	nei endonuclease VIII-like 1 (E. coli) (Neil1) alternative variant dSep08, mRNA.
<a href="#">Neil1</a>	<a href="#">Neil1.eSep08</a>	<a href="#">367090</a>	902	577	3	26	nei endonuclease VIII-like 1 (E. coli) (2.9 kD) (Neil1) alternative variant eSep08, mRNA.
<a href="#">Neil3</a>	<a href="#">Neil3.aSep08</a>	<a href="#">290729</a>	12624	1396		302	nei like 3 (E. coli) (33.6 kD) (Neil3) mRNA.
<a href="#">Nek1</a>	<a href="#">Nek1.bSep08</a>	<a href="#">290705</a>	18816	501	4	166	NIMA (never in mitosis gene a)-related expressed kinase 1 (Nek1) alternative variant bSep08, mRNA.
<a href="#">Nek1</a>	<a href="#">Nek1.cSep08</a>	<a href="#">290705</a>	78967	531	6	153	NIMA (never in mitosis gene a)-related expressed kinase 1 (Nek1) alternative variant cSep08, mRNA.
<a href="#">Nek3</a>	<a href="#">Nek3.aSep08</a>	<a href="#">306576</a>	20740	1592	11	399	NIMA (never in mitosis gene a)-related expressed kinase 3 (Nek3) alternative variant aSep08, mRNA.
<a href="#">Nek4</a>	<a href="#">Nek4.bSep08</a>	<a href="#">306252</a>	10246	977	6	325	NIMA (never in mitosis gene a)-related expressed kinase 4 (Nek4) alternative variant bSep08, mRNA.
<a href="#">Nek4</a>	<a href="#">Nek4.cSep08</a>	<a href="#">306252</a>	14509	1183	8	226	NIMA (never in mitosis gene a)-related expressed kinase 4 (25.7 kD) (Nek4) alternative variant cSep08, mRNA.
<a href="#">Nek6</a>	<a href="#">Nek6.aSep08</a>	<a href="#">360161</a>	71775	2287	8	345	NIMA (never in mitosis gene a)-related expressed kinase 6 (Nek6) alternative variant aSep08, mRNA.
<a href="#">Nek6</a>	<a href="#">Nek6.bSep08</a>	<a href="#">360161</a>	47004	809	2	147	NIMA (never in mitosis gene a)-related expressed kinase 6 (Nek6) alternative variant bSep08, mRNA.
<a href="#">Nek6</a>	<a href="#">Nek6.cSep08</a>	<a href="#">360161</a>	36138	493	3	105	NIMA (never in mitosis gene a)-related expressed kinase 6 (Nek6) alternative variant cSep08, mRNA.
<a href="#">Nek7</a>	<a href="#">Nek7.bSep08</a>	<a href="#">360850</a>	51119	646	7	215	NIMA (never in mitosis gene a)-related expressed kinase 7 (Nek7) alternative variant bSep08, mRNA.
<a href="#">Nek7</a>	<a href="#">Nek7.cSep08</a>	<a href="#">360850</a>	14876	601	2	89	NIMA (never in mitosis gene a)-related expressed kinase 7 (Nek7) alternative variant cSep08, mRNA.
<a href="#">Nek9</a>	<a href="#">Nek9.bSep08</a>	<a href="#">299204</a>	15873	1848	7	448	NIMA (never in mitosis gene a)-related kinase 9 (Nek9) alternative variant bSep08, mRNA.
<a href="#">Nek11</a>	<a href="#">Nek11.bSep08</a>	<a href="#">315978</a>	79318	772		187	NIMA (never in mitosis gene a)-related expressed kinase 11 (Nek11) alternative variant bSep08, mRNA.
<a href="#">Nelf</a>	<a href="#">Nelf.aSep08</a>	<a href="#">117536</a>	5702	1093	7	310	nasal embryonic LHRH factor (Nelf) alternative variant aSep08, mRNA.

<a href="#">Nelf</a>	<a href="#">Nelf.bSep08</a>	<a href="#">117536</a>	4973	1974	11	292	nasal embryonic LHRH factor (Nelf) alternative variant bSep08, mRNA.
<a href="#">Nelf</a>	<a href="#">Nelf.cSep08</a>	<a href="#">117536</a>	5103	740	5	246	nasal embryonic LHRH factor (Nelf) alternative variant cSep08, mRNA.
<a href="#">Nell2</a>	<a href="#">Nell2.bSep08</a>	<a href="#">81734</a>	44436	750	6	194	NEL-like 2 (chicken) (Nell2) alternative variant bSep08, mRNA.
<a href="#">Nell2</a>	<a href="#">Nell2.cSep08</a>	<a href="#">81734</a>	101063	598	5	183	NEL-like 2 (chicken) (Nell2) alternative variant cSep08, mRNA.
<a href="#">Nenf</a>	<a href="#">Nenf.bSep08</a>	<a href="#">289380</a>	8903	546	4	136	neuron derived neurotrophic factor (Nenf) alternative variant bSep08, mRNA.
<a href="#">Neo1</a>	<a href="#">Neo1.aSep08</a>	<a href="#">81735</a>	19337	3370		332	neogenin (Neo1) mRNA.
<a href="#">nerby</a>	<a href="#">nerby.aSep08</a>		2023	380		126	KIF4B (nerby) mRNA.
<a href="#">nerchy</a>	<a href="#">nerchy.aSep08</a>		27878	681	1	104	putative protein (nerchy) alternative variant aSep08, mRNA.
<a href="#">nerchy</a>	<a href="#">nerchy.bSep08</a>		27863	607		80	putative protein (nerchy) alternative variant bSep08, mRNA.
<a href="#">nerdar</a>	<a href="#">nerdar.aSep08</a>		4096	688		112	putative protein (12.5 kD) (nerdar) mRNA.
<a href="#">nerfer</a>	<a href="#">nerfer.aSep08</a>		76195	626		208	putative protein of vertebrate origin (nerfer) mRNA.
<a href="#">nerflo</a>	<a href="#">nerflo.aSep08</a>		8738	465		127	putative nuclear protein (14.2 kD) (nerflo) mRNA.
<a href="#">nerflu</a>	<a href="#">nerflu.aSep08</a>		3970	599		90	putative nuclear protein (9.7 kD) (nerflu) mRNA.
<a href="#">nergar</a>	<a href="#">nergar.aSep08</a>		934	726		59	putative protein (7.0 kD) (nergar) mRNA.
<a href="#">nerja</a>	<a href="#">nerja.aSep08</a>		4890	959		80	ab2-143 like (nerja) mRNA.
<a href="#">nerjey</a>	<a href="#">nerjey.aSep08</a>		1095	247		50	putative protein (nerjey) mRNA.
<a href="#">nerkee</a>	<a href="#">nerkee.aSep08</a>		4052	492		41	putative protein (5.0 kD) (nerkee) mRNA.
<a href="#">nerkler</a>	<a href="#">nerkler.aSep08</a>		622	238		79	putative protein (nerkler) mRNA.
<a href="#">nerloy</a>	<a href="#">nerloy.aSep08</a>		465	286		34	putative protein (nerloy) mRNA.
<a href="#">nermee</a>	<a href="#">nermee.aSep08</a>		14016	733		64	putative protein (7.4 kD) (nermee) mRNA.
<a href="#">nermer</a>	<a href="#">nermer.aSep08</a>		1181	373		124	CRA b (nermer) mRNA.
<a href="#">nernoy</a>	<a href="#">nernoy.aSep08</a>		9643	183		29	putative protein (nernoy) mRNA.
<a href="#">nerpor</a>	<a href="#">nerpor.aSep08</a>		12812	408	1	135	casein kinase 1 gamma CRA a (nerpor) alternative variant aSep08, mRNA.
<a href="#">nerpor</a>	<a href="#">nerpor.bSep08</a>		17381	576	1	87	casein kinase 1 gamma CRA a (nerpor) alternative variant bSep08, mRNA.
<a href="#">nersa</a>	<a href="#">nersa.aSep08</a>		5127	308		28	putative protein (nersa) mRNA.
<a href="#">nershee</a>	<a href="#">nershee.aSep08</a>		8549	970		51	putative protein (nershee) mRNA.
<a href="#">nertu</a>	<a href="#">nertu.aSep08</a>		626	249		52	putative protein (nertu) mRNA.
<a href="#">nervar</a>	<a href="#">nervar.aSep08</a>		743	443		107	putative protein (nervar) mRNA.
<a href="#">nerwey</a>	<a href="#">nerwey.aSep08</a>		27583	708	1	157	transmembrane coiled-coil domain family 1 CRA a (nerwey) alternative variant aSep08, mRNA.
<a href="#">nerwey</a>	<a href="#">nerwey.bSep08</a>		26484	336	1	111	transmembrane coiled-coil domain family 1 CRA a (nerwey) alternative variant bSep08, mRNA.
<a href="#">Net1</a>	<a href="#">Net1.bSep08</a>	<a href="#">307098</a>	24453	812	3	270	neuroepithelial cell transforming gene 1 (Net1) alternative variant bSep08, mRNA.

<a href="#">Neto1</a>	<a href="#">Neto1.bSep08</a>	<a href="#">307206</a>	7243	537	1	74	neuropilin (NRP) and tolloid (TLL)-like 1 (8.4 kD) (Neto1) alternative variant bSep08, mRNA.
<a href="#">Neto2</a>	<a href="#">Neto2.bSep08</a>	<a href="#">307757</a>	31099	932	3	310	neuropilin (NRP) and tolloid (TLL)-like 2 (Neto2) alternative variant bSep08, mRNA.
<a href="#">Neu1</a>	<a href="#">Neu1.bSep08</a>	<a href="#">24591</a>	2328	2232	2	320	neuraminidase 1 (Neu1) alternative variant bSep08, mRNA.
<a href="#">Neu1</a>	<a href="#">Neu1.cSep08</a>	<a href="#">24591</a>	2204	643	3	206	neuraminidase 1 (Neu1) alternative variant cSep08, mRNA.
<a href="#">Neu2</a>	<a href="#">Neu2.bSep08</a>	<a href="#">29204</a>	22025	785	1	222	neuraminidase 2 (Neu2) alternative variant bSep08, mRNA.
<a href="#">Neu3</a>	<a href="#">Neu3.aSep08</a>	<a href="#">117185</a>	11528	2773	2	489	neuraminidase 3 (Neu3) alternative variant aSep08, mRNA.
<a href="#">Neuralized.0</a>	<a href="#">Neuralized.0.bSep08</a>		69226	489	2	162	neuralized-like (Neuralized.0) alternative variant bSep08, mRNA.
<a href="#">Neurl</a>	<a href="#">Neurl.aSep08</a>	<a href="#">309459</a>	6717	2403		218	neuralized-like homolog (Drosophila) (Neurl) mRNA.
<a href="#">Nexn</a>	<a href="#">Nexn.cSep08</a>	<a href="#">246172</a>	31653	2023	10	622	nexilin (Nexn) alternative variant cSep08, mRNA.
<a href="#">Nexn</a>	<a href="#">Nexn.eSep08</a>	<a href="#">246172</a>	1257	1171	2	91	nexilin (10.4 kD) (Nexn) alternative variant eSep08, mRNA.
<a href="#">Nexn</a>	<a href="#">Nexn.fSep08</a>	<a href="#">246172</a>	12354	252	4	32	nexilin CRA c (Nexn) alternative variant fSep08, mRNA.
<a href="#">Nexn</a>	<a href="#">Nexn.gSep08</a>	<a href="#">246172</a>	23100	395	5		
<a href="#">Nexn</a>	<a href="#">Nexn.hSep08</a>	<a href="#">246172</a>	437	361	2	37	putative protein (Nexn) alternative variant hSep08, mRNA.
<a href="#">neyby</a>	<a href="#">neyby.aSep08</a>		15395	499		45	putative protein (5.0 kD) (neyby) mRNA.
<a href="#">neychy</a>	<a href="#">neychy.aSep08</a>		14112	373		107	damage-specific DNA binding protein 2 like (neychy) mRNA.
<a href="#">neydar</a>	<a href="#">neydar.aSep08</a>		5719	1783	4	513	putative protein (neydar) alternative variant aSep08, mRNA.
<a href="#">neydar</a>	<a href="#">neydar.bSep08</a>		4093	550	2	92	putative protein (neydar) alternative variant bSep08, mRNA.
<a href="#">neydar</a>	<a href="#">neydar.cSep08</a>		2656	840	4	60	putative protein (6.9 kD) (neydar) alternative variant cSep08, mRNA.
<a href="#">neyfer</a>	<a href="#">neyfer.aSep08</a>		13069	741		45	putative protein (neyfer) mRNA.
<a href="#">neyflo</a>	<a href="#">neyflo.aSep08</a>		14048	1153	1	53	putative protein (5.8 kD) (neyflo) alternative variant aSep08, mRNA.
<a href="#">neyflo</a>	<a href="#">neyflo.bSep08</a>		12455	579	1	67	putative protein (neyflo) alternative variant bSep08, mRNA.
<a href="#">neyflu</a>	<a href="#">neyflu.aSep08</a>		15583	441		146	putative protein of metazoan origin (neyflu) mRNA.
<a href="#">neygar</a>	<a href="#">neygar.aSep08</a>		5127	675	2	64	putative protein (6.8 kD) (neygar) alternative variant aSep08, mRNA.
<a href="#">neygar</a>	<a href="#">neygar.bSep08</a>		4350	717	2	51	putative protein (5.7 kD) (neygar) alternative variant bSep08, mRNA.
<a href="#">neyja</a>	<a href="#">neyja.aSep08</a>		745	303		93	putative protein, with a coiled coil domain, of vertebrate origin (neyja) mRNA.
<a href="#">neyjey</a>	<a href="#">neyjey.aSep08</a>		621	334		70	putative protein (neyjey) mRNA.
<a href="#">neykee</a>	<a href="#">neykee.aSep08</a>		96616	604		123	putative protein (neykee) mRNA.
<a href="#">neykler</a>	<a href="#">neykler.aSep08</a>		11759	658		100	putative protein (11.1 kD) (neykler) mRNA.
<a href="#">neyloy</a>	<a href="#">neyloy.aSep08</a>		6495	541	1	85	putative protein (neyloy) alternative variant aSep08, mRNA.

<a href="#">neyloy</a>	<a href="#">neyloy.bSep08</a>		6548	588	1	73	putative protein (neyloy) alternative variant bSep08, mRNA.
<a href="#">neyloy</a>	<a href="#">neyloy.cSep08</a>		6824	764	2	50	putative protein (5.4 kD) (neyloy) alternative variant cSep08, mRNA.
<a href="#">neyloy</a>	<a href="#">neyloy.dSep08</a>		6800	734	2	48	putative protein (5.2 kD) (neyloy) alternative variant dSep08, mRNA.
<a href="#">neymee</a>	<a href="#">neymee.aSep08</a>		999	700		46	putative protein (5.2 kD) (neymee) mRNA.
<a href="#">neymer</a>	<a href="#">neymer.aSep08</a>		1877	463		73	myosin-xvb like (neymer) mRNA.
<a href="#">neynoy</a>	<a href="#">neynoy.aSep08</a>		8704	271		14	putative protein (1.5 kD) (neynoy) mRNA.
<a href="#">neypor</a>	<a href="#">neypor.aSep08</a>		867	419		42	putative protein (neypor) mRNA.
<a href="#">neysa</a>	<a href="#">neysa.aSep08</a>		613	489		45	putative protein (neysa) mRNA.
<a href="#">neyshee</a>	<a href="#">neyshee.aSep08</a>		4984	240		13	putative protein (neyshee) mRNA.
<a href="#">neytu</a>	<a href="#">neytu.aSep08</a>		10171	664	1	126	SALF (neytu) alternative variant aSep08, mRNA.
<a href="#">neytu</a>	<a href="#">neytu.bSep08</a>		15687	384	1	65	CRA b like (neytu) alternative variant bSep08, mRNA.
<a href="#">neyvar</a>	<a href="#">neyvar.aSep08</a>		680	510		66	putative protein (neyvar) mRNA.
<a href="#">neywey</a>	<a href="#">neywey.aSep08</a>		1939	988		127	putative protein (15.2 kD) (neywey) mRNA.
<a href="#">Nf1</a>	<a href="#">Nf1.aSep08</a>	<a href="#">24592</a>	18108	1456		485	neurofibromatosis 1 (Nf1) mRNA.
<a href="#">Nf2</a>	<a href="#">Nf2.bSep08</a>	<a href="#">25744</a>	14102	1784	4	465	neurofibromatosis 2 (Nf2) alternative variant bSep08, mRNA.
<a href="#">Nf2</a>	<a href="#">Nf2.cSep08</a>	<a href="#">25744</a>	14311	678	2	81	neurofibromatosis 2 (8.5 kD) (Nf2) alternative variant cSep08, mRNA.
<a href="#">Nfam1</a>	<a href="#">Nfam1.aSep08</a>	<a href="#">362966</a>	18110	735	5	168	nfat activating molecule with ITAM motif 1 (Nfam1) alternative variant aSep08, mRNA.
<a href="#">Nfam1</a>	<a href="#">Nfam1.bSep08</a>	<a href="#">362966</a>	2408	758	3	88	nfat activating molecule with ITAM motif 1 (Nfam1) alternative variant bSep08, mRNA.
<a href="#">Nfam1</a>	<a href="#">Nfam1.cSep08</a>	<a href="#">362966</a>	5381	923	2	57	nfat activating molecule with ITAM motif 1 (Nfam1) alternative variant cSep08, mRNA.
<a href="#">Nfasc</a>	<a href="#">Nfasc.bSep08</a>	<a href="#">116690</a>	5064	569	5	189	neurofascin (Nfasc) alternative variant bSep08, mRNA.
<a href="#">Nfatc2ip</a>	<a href="#">Nfatc2ip.aSep08</a>	<a href="#">308983</a>	16534	1800	3	538	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2 interacting protein (Nfatc2ip) alternative variant aSep08, mRNA.
<a href="#">Nfatc3</a>	<a href="#">Nfatc3.bSep08</a>	<a href="#">361400</a>	22568	1549	3	469	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 (50.6 kD) (Nfatc3) alternative variant bSep08, mRNA.
<a href="#">Nfatc3</a>	<a href="#">Nfatc3.cSep08</a>	<a href="#">361400</a>	21076	1105	2	354	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 (Nfatc3) alternative variant cSep08, mRNA.
<a href="#">Nfe2</a>	<a href="#">Nfe2.bSep08</a>	<a href="#">366998</a>	5227	706	1	185	nuclear factor, erythroid derived 2 (Nfe2) alternative variant bSep08, mRNA.
<a href="#">Nfe211</a>	<a href="#">Nfe211.bSep08</a>	<a href="#">360610</a>	6121	3496	2	577	nuclear factor, erythroid derived 2,-like 1 (Nfe211) alternative variant bSep08, mRNA.
<a href="#">Nfe211</a>	<a href="#">Nfe211.cSep08</a>	<a href="#">360610</a>	6452	408	2	64	nuclear factor, erythroid derived 2,-like 1 (Nfe211) alternative variant cSep08, mRNA.
<a href="#">Nfe212</a>	<a href="#">Nfe212.bSep08</a>	<a href="#">83619</a>	26470	1372	5	428	nuclear factor, erythroid derived 2, like 2 (Nfe212) alternative variant bSep08, mRNA.

<a href="#">Nfe2l2</a>	<a href="#">Nfe2l2.dSep08</a>	<a href="#">83619</a>	24073	689	4	111	nuclear factor, erythroid derived 2, like 2 (Nfe2l2) alternative variant dSep08, mRNA.
<a href="#">Nfe2l3</a>	<a href="#">Nfe2l3.aSep08</a>	<a href="#">312331</a>	26456	1010		336	nuclear factor, erythroid derived 2, like 3 (Nfe2l3) mRNA.
<a href="#">Nfia</a>	<a href="#">Nfia.bSep08</a>	<a href="#">25492</a>	341814	1952	5	478	nuclear factor I/A (Nfia) alternative variant bSep08, mRNA.
<a href="#">Nfia</a>	<a href="#">Nfia.cSep08</a>	<a href="#">25492</a>	341466	1283	2	371	nuclear factor I/A (Nfia) alternative variant cSep08, mRNA.
<a href="#">Nfia</a>	<a href="#">Nfia.dSep08</a>	<a href="#">25492</a>	343738	3850	5	357	nuclear factor I/A (Nfia) alternative variant dSep08, mRNA.
<a href="#">Nfib</a>	<a href="#">Nfib.bSep08</a>	<a href="#">29227</a>	58054	1170	5	183	nuclear factor I/B (Nfib) alternative variant bSep08, mRNA.
<a href="#">Nfib</a>	<a href="#">Nfib.cSep08</a>	<a href="#">29227</a>	31010	2396	4	121	nuclear factor I/B (13.2 kD) (Nfib) alternative variant cSep08, mRNA.
<a href="#">Nfil3</a>	<a href="#">Nfil3.bSep08</a>	<a href="#">114519</a>	12107	697	2	126	nuclear factor, interleukin 3 regulated (Nfil3) alternative variant bSep08, mRNA.
<a href="#">Nfix</a>	<a href="#">Nfix.bSep08</a>	<a href="#">81524</a>	66129	1425	9	392	nuclear factor I/X (43.7 kD) (Nfix) alternative variant bSep08, mRNA.
<a href="#">Nfix</a>	<a href="#">Nfix.cSep08</a>	<a href="#">81524</a>	19292	1004	6	156	nuclear factor I/X (16.3 kD) (Nfix) alternative variant cSep08, mRNA.
<a href="#">Nfkb1</a>	<a href="#">Nfkb1.aSep08</a>	<a href="#">81736</a>	44673	2378	18	792	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1, p105 (Nfkb1) alternative variant aSep08, mRNA.
<a href="#">Nfkb1</a>	<a href="#">Nfkb1.bSep08</a>	<a href="#">81736</a>	9178	1348	6	203	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1, p105 (Nfkb1) alternative variant bSep08, mRNA.
<a href="#">Nfkb1</a>	<a href="#">Nfkb1.cSep08</a>	<a href="#">81736</a>	8545	1185	6	198	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1, p105 (21.7 kD) (Nfkb1) alternative variant cSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.bSep08</a>	<a href="#">309452</a>	7252	1783	7	552	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant bSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.bSep08</a>	<a href="#">689848</a>	7252	1783	7	552	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant bSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.cSep08</a>	<a href="#">309452</a>	1647	749	6	181	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant cSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.cSep08</a>	<a href="#">689848</a>	1647	749	6	181	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant cSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.dSep08</a>	<a href="#">309452</a>	700	498	3	166	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant dSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.dSep08</a>	<a href="#">689848</a>	700	498	3	166	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant dSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.eSep08</a>	<a href="#">309452</a>	2662	625	7	155	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant eSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.eSep08</a>	<a href="#">689848</a>	2662	625	7	155	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant eSep08, mRNA.



<a href="#">Nfkb2</a>	<a href="#">Nfkb2.fSep08</a>	<a href="#">309452</a>	482	397	2	89	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant fSep08, mRNA.
<a href="#">Nfkb2</a>	<a href="#">Nfkb2.fSep08</a>	<a href="#">689848</a>	482	397	2	89	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100 and hypothetical protein LOC689848 (Nfkb2) alternative variant fSep08, mRNA.
<a href="#">Nfkbia</a>	<a href="#">Nfkbia.aSep08</a>	<a href="#">25493</a>	3111	1963	5	327	kappa alpha (Nfkbia) alternative variant aSep08, complete mRNA.
<a href="#">Nfkbia</a>	<a href="#">Nfkbia.cSep08</a>	<a href="#">25493</a>	1905	1120	3	211	nuclear factor of kappa light gene enhancer in B-cells inhibitor alpha (23.4 kD) (Nfkbia) alternative variant cSep08, mRNA.
<a href="#">Nfkbia</a>	<a href="#">Nfkbia.dSep08</a>	<a href="#">25493</a>	2118	1755	3	129	kappa alpha (14.4 kD) (Nfkbia) alternative variant dSep08, mRNA.
<a href="#">Nfkbid</a>	<a href="#">Nfkbid.aSep08</a>	<a href="#">308496</a>	2550	788		148	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, delta (Nfkbid) mRNA.
<a href="#">Nfkbie</a>	<a href="#">Nfkbie.bSep08</a>	<a href="#">316241</a>	7636	2039	3	308	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon (Nfkbie) alternative variant bSep08, mRNA.
<a href="#">Nfkbil1</a>	<a href="#">Nfkbil1.bSep08</a>	<a href="#">361794</a>	15795	996	5	270	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1 (Nfkbil1) alternative variant bSep08, mRNA.
<a href="#">Nfkbil1</a>	<a href="#">Nfkbil1.dSep08</a>	<a href="#">361794</a>	14751	553	4	164	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1 (Nfkbil1) alternative variant dSep08, mRNA.
<a href="#">Nfrkb</a>	<a href="#">Nfrkb.bSep08</a>	<a href="#">315523</a>	3110	694	5	169	nuclear factor related to kappa B binding protein (Nfrkb) alternative variant bSep08, mRNA.
<a href="#">Nfrkb</a>	<a href="#">Nfrkb.dSep08</a>	<a href="#">315523</a>	1454	545	3	149	nuclear factor related to kappa B binding protein (Nfrkb) alternative variant dSep08, mRNA.
<a href="#">Nfrkb</a>	<a href="#">Nfrkb.eSep08</a>	<a href="#">315523</a>	2506	404	5	134	nuclear factor related to kappa B binding protein (Nfrkb) alternative variant eSep08, mRNA.
<a href="#">Nfrkb</a>	<a href="#">Nfrkb.fSep08</a>	<a href="#">315523</a>	1544	1447	2	98	nuclear factor related to kappa B binding protein (Nfrkb) alternative variant fSep08, mRNA.
<a href="#">Nfrkb</a>	<a href="#">Nfrkb.gSep08</a>	<a href="#">315523</a>	945	221	2	35	nuclear factor related to kappa B binding protein (3.9 kD) (Nfrkb) alternative variant gSep08, mRNA.
<a href="#">Nfs1</a>	<a href="#">Nfs1.bSep08</a>	<a href="#">84594</a>	15567	1358	7	292	nitrogen fixation 1 (32.1 kD) (Nfs1) alternative variant bSep08, mRNA.
<a href="#">Nfs1</a>	<a href="#">Nfs1.cSep08</a>	<a href="#">84594</a>	3974	1646	4	102	nfs1 nitrogen fixation 1 (Nfs1) alternative variant cSep08, mRNA.
<a href="#">Nfs1</a>	<a href="#">Nfs1.dSep08</a>	<a href="#">84594</a>	1663	798	2	44	putative protein (Nfs1) alternative variant dSep08, mRNA.
<a href="#">Nfu1</a>	<a href="#">Nfu1.aSep08</a>	<a href="#">297416</a>	20782	909	8	254	NFU1 iron-sulfur cluster scaffold homolog ( <i>S. cerevisiae</i> ) (Nfu1) alternative variant aSep08, mRNA.
<a href="#">Nfu1</a>	<a href="#">Nfu1.bSep08</a>	<a href="#">297416</a>	21547	1538	7	114	NFU1 iron-sulfur cluster scaffold homolog ( <i>S. cerevisiae</i> ) (12.5 kD) (Nfu1) alternative variant bSep08, mRNA.
<a href="#">Nfu1</a>	<a href="#">Nfu1.dSep08</a>	<a href="#">297416</a>	4673	1788	3	13	NFU1 iron-sulfur cluster scaffold homolog ( <i>S. cerevisiae</i> ) (Nfu1) alternative variant dSep08, mRNA.
<a href="#">Nfx1</a>	<a href="#">Nfx1.bSep08</a>	<a href="#">313166</a>	20733	1356	8	281	nuclear transcription factor X-box binding 1 like (Nfx1) alternative variant bSep08, mRNA.

<a href="#">Nfx1</a>	<a href="#">Nfx1.cSep08</a>	<a href="#">313166</a>	13526	744	7	247	nuclear transcription factor X-box binding 1 like (Nfx1) alternative variant cSep08, mRNA.
<a href="#">Nfx1</a>	<a href="#">Nfx1.dSep08</a>	<a href="#">313166</a>	2643	807	2	93	nuclear transcription factor X-box binding 1 like (10.2 kD) (Nfx1) alternative variant dSep08, mRNA.
<a href="#">Nfx1</a>	<a href="#">Nfx1.eSep08</a>	<a href="#">313166</a>	4837	431	2	84	nuclear transcription factor X-box binding 1 like (Nfx1) alternative variant eSep08, mRNA.
<a href="#">Nfya</a>	<a href="#">Nfya.aSep08</a>	<a href="#">29508</a>	15550	1777	5	502	nuclear transcription factor-Y alpha (Nfya) alternative variant aSep08, mRNA.
<a href="#">Nfya</a>	<a href="#">Nfya.dSep08</a>	<a href="#">29508</a>	13762	364	4	69	nuclear transcription factor-Y alpha (Nfya) alternative variant dSep08, mRNA.
<a href="#">Nfyb</a>	<a href="#">Nfyb.bSep08</a>	<a href="#">25336</a>	3793	396	2	113	nuclear transcription factor-Y beta (Nfyb) alternative variant bSep08, mRNA.
<a href="#">Nfyb</a>	<a href="#">Nfyb.cSep08</a>	<a href="#">25336</a>	13452	1392	4	110	nuclear transcription factor-Y beta (12.0 kD) (Nfyb) alternative variant cSep08, mRNA.
<a href="#">Nfyc</a>	<a href="#">Nfyc.bSep08</a>	<a href="#">25337</a>	57883	767	7	244	nuclear transcription factor Y (Nfyc) alternative variant bSep08, mRNA.
<a href="#">Nfyc</a>	<a href="#">Nfyc.cSep08</a>	<a href="#">25337</a>	8825	1823	4	177	putative mitochondrial protein human specific (19.3 kD) (Nfyc) alternative variant cSep08, mRNA.
<a href="#">Nfyc</a>	<a href="#">Nfyc.dSep08</a>	<a href="#">25337</a>	3796	533	3	98	putative secreted or extracellular protein precursor human specific (10.5 kD) (Nfyc) alternative variant dSep08, mRNA.
<a href="#">Nfyc</a>	<a href="#">Nfyc.eSep08</a>	<a href="#">25337</a>	2015	1022	2	65	nuclear transcription factor-Y gamma CRA a like (7.0 kD) (Nfyc) alternative variant eSep08, mRNA.
<a href="#">Ng23</a>	<a href="#">Ng23.bSep08</a>	<a href="#">406170</a>	1795	863	1	142	ng23 protein (Ng23) alternative variant bSep08, mRNA.
<a href="#">Ngb</a>	<a href="#">Ngb.aSep08</a>	<a href="#">85382</a>	5277	1763		176	neuroglobin (19.8 kD) (Ngb) mRNA.
<a href="#">Ngdn</a>	<a href="#">Ngdn.bSep08</a>	<a href="#">305887</a>	2095	726	6	217	neuroguidin, EIF4E binding protein (Ngdn) alternative variant bSep08, mRNA.
<a href="#">Ngdn</a>	<a href="#">Ngdn.cSep08</a>	<a href="#">305887</a>	813	414	4	81	neuroguidin, EIF4E binding protein (Ngdn) alternative variant cSep08, mRNA.
<a href="#">Ngdn</a>	<a href="#">Ngdn.eSep08</a>	<a href="#">305887</a>	4249	428	4	48	neuroguidin, EIF4E binding protein (Ngdn) alternative variant eSep08, mRNA.
<a href="#">Ngef</a>	<a href="#">Ngef.aSep08</a>	<a href="#">246217</a>	10521	1816	8	361	neuronal guanine nucleotide exchange factor (Ngef) alternative variant aSep08, mRNA.
<a href="#">Ngef</a>	<a href="#">Ngef.bSep08</a>	<a href="#">246217</a>	7392	1670	6	313	neuronal guanine nucleotide exchange factor (Ngef) alternative variant bSep08, mRNA.
<a href="#">Ngef</a>	<a href="#">Ngef.cSep08</a>	<a href="#">246217</a>	882	502	1	95	neuronal guanine nucleotide exchange factor (Ngef) alternative variant cSep08, mRNA.
<a href="#">Ngfrap1</a>	<a href="#">Ngfrap1.aSep08</a>	<a href="#">117089</a>	1547	835	2	130	nerve growth factor receptor (TNFRSF16) associated protein 1 (15.3 kD) (Ngfrap1) alternative variant aSep08, complete mRNA.
<a href="#">Ngly1</a>	<a href="#">Ngly1.bSep08</a>	<a href="#">361014</a>	17791	1268	5	260	N-glycanase 1 (Ngly1) alternative variant bSep08, mRNA.
<a href="#">Ngly1</a>	<a href="#">Ngly1.cSep08</a>	<a href="#">361014</a>	9165	689	5	208	N-glycanase 1 (Ngly1) alternative variant cSep08, mRNA.
<a href="#">Ngly1</a>	<a href="#">Ngly1.dSep08</a>	<a href="#">361014</a>	15458	1329	4	203	N-glycanase 1 (Ngly1) alternative variant dSep08, mRNA.
<a href="#">Ngly1</a>	<a href="#">Ngly1.eSep08</a>	<a href="#">361014</a>	16913	545	4	65	N-glycanase 1 (Ngly1) alternative variant eSep08, mRNA.
<a href="#">Nhedc2</a>	<a href="#">Nhedc2.aSep08</a>	<a href="#">365946</a>	26075	1250	8	356	sodium/hydrogen exchanger (Nhedc2) mRNA.

<a href="#">Nhej1</a>	<a href="#">Nhej1.bSep08</a>	<a href="#">363251</a>	5785	376	3	79	nonhomologous end-joining factor 1 (Nhej1) alternative variant bSep08, mRNA.
<a href="#">Nhej1</a>	<a href="#">Nhej1.cSep08</a>	<a href="#">363251</a>	4936	668	3	60	nonhomologous end-joining factor 1 (Nhej1) alternative variant cSep08, mRNA.
<a href="#">Nhlh1</a>	<a href="#">Nhlh1.bSep08</a>	<a href="#">289230</a>	3806	1159	1	133	nescient helix loop helix 1 (14.8 kD) (Nhlh1) alternative variant bSep08, mRNA.
<a href="#">Nhlh2</a>	<a href="#">Nhlh2.bSep08</a>	<a href="#">295327</a>	3435	1440	2	117	nescient helix loop helix 2 (12.1 kD) (Nhlh2) alternative variant bSep08, mRNA.
<a href="#">Nhirc2</a>	<a href="#">Nhirc2.bSep08</a>	<a href="#">307986</a>	4816	1674	3	189	NHL repeat containing 2 (Nhirc2) alternative variant bSep08, mRNA.
<a href="#">Nhirc2</a>	<a href="#">Nhirc2.cSep08</a>	<a href="#">307986</a>	932	531	2	105	NHL repeat containing 2 (Nhirc2) alternative variant cSep08, mRNA.
<a href="#">Nibp</a>	<a href="#">Nibp.aSep08</a>	<a href="#">315059</a>	346208	944	7	314	trafficking protein particle complex 9 (Nibp) alternative variant aSep08, mRNA.
<a href="#">Nibp</a>	<a href="#">Nibp.bSep08</a>	<a href="#">315059</a>	73820	515	6	171	trafficking protein particle complex 9 (Nibp) alternative variant bSep08, mRNA.
<a href="#">Nibp</a>	<a href="#">Nibp.cSep08</a>	<a href="#">315059</a>	96688	1268	3	150	trafficking protein particle complex 9 (16.7 kD) (Nibp) alternative variant cSep08, mRNA.
<a href="#">Nid1</a>	<a href="#">Nid1.aSep08</a>	<a href="#">25494</a>	10594	2929		272	nidogen 1 (Nid1) mRNA.
<a href="#">Nid2</a>	<a href="#">Nid2.bSep08</a>	<a href="#">302248</a>	18020	1782	8	415	nidogen 2 CRA b (Nid2) alternative variant bSep08, mRNA.
<a href="#">Nid2</a>	<a href="#">Nid2.cSep08</a>	<a href="#">302248</a>	12255	1376	6	358	nidogen 2 CRA a (38.8 kD) (Nid2) alternative variant cSep08, mRNA.
<a href="#">Nid2</a>	<a href="#">Nid2.dSep08</a>	<a href="#">302248</a>	7152	1353	6	262	nidogen 2 (Nid2) alternative variant dSep08, mRNA.
<a href="#">Nid2</a>	<a href="#">Nid2.eSep08</a>	<a href="#">302248</a>	4281	871	4	163	nidogen 2 (Nid2) alternative variant eSep08, mRNA.
<a href="#">Nid67</a>	<a href="#">Nid67.aSep08</a>	<a href="#">286910</a>	26844	897	1	148	putative small membrane protein NID67 (16.2 kD) (Nid67) alternative variant aSep08, mRNA.
<a href="#">NIDO.0</a>	<a href="#">NIDO.0.aSep08</a>		3369	357		118	nidogen (NIDO.0) mRNA.
<a href="#">Nif311</a>	<a href="#">Nif311.aSep08</a>	<a href="#">301431</a>	19925	1704	8	376	ngg1 interacting factor 3-like 1 (S. pombe) (41.5 kD) (Nif311) alternative variant aSep08, mRNA.
<a href="#">Nif311</a>	<a href="#">Nif311.dSep08</a>	<a href="#">301431</a>	2745	738	3	161	ngg1 interacting factor 3-like 1 (S. pombe) (18.1 kD) (Nif311) alternative variant dSep08, mRNA.
<a href="#">Nin</a>	<a href="#">Nin.bSep08</a>	<a href="#">299117</a>	10775	1179	1	243	ninein (Nin) alternative variant bSep08, mRNA.
<a href="#">Ninj1</a>	<a href="#">Ninj1.cSep08</a>	<a href="#">25338</a>	2501	530	2	71	ninjurin 1 (Ninj1) alternative variant cSep08, mRNA.
<a href="#">Ninj2</a>	<a href="#">Ninj2.bSep08</a>	<a href="#">59115</a>	1788	669	1	113	ninjurin 2 (Ninj2) alternative variant bSep08, mRNA.
<a href="#">Nip30</a>	<a href="#">Nip30.bSep08</a>	<a href="#">307652</a>	21434	755	2	167	NEFA-interacting nuclear protein NIP30 (20.1 kD) (Nip30) alternative variant bSep08, mRNA.
<a href="#">Nip30</a>	<a href="#">Nip30.cSep08</a>	<a href="#">307652</a>	21431	748	2	158	NEFA-interacting nuclear protein NIP30 (Nip30) alternative variant cSep08, mRNA.
<a href="#">Nip30</a>	<a href="#">Nip30.dSep08</a>	<a href="#">307652</a>	11773	754	2	89	NEFA-interacting nuclear protein NIP30 (Nip30) alternative variant dSep08, mRNA.
<a href="#">Nipa2</a>	<a href="#">Nipa2.aSep08</a>	<a href="#">308667</a>	24129	2382	9	401	non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human) (Nipa2) alternative variant aSep08, mRNA.
<a href="#">Nipa2</a>	<a href="#">Nipa2.dSep08</a>	<a href="#">308667</a>	2433	830	2	40	non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human) (4.8 kD) (Nipa2) alternative variant dSep08, mRNA.

<a href="#">Nipa2</a>	<a href="#">Nipa2.eSep08</a>	<a href="#">308667</a>	12257	715	5	65	non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human) (Nipa2) alternative variant eSep08, mRNA.
<a href="#">NIPBL</a>	<a href="#">NIPBL.bSep08</a>	<a href="#">294787</a>	6568	1776	4	276	delangin (NIPBL) alternative variant bSep08, mRNA.
<a href="#">NIPBL</a>	<a href="#">NIPBL.cSep08</a>	<a href="#">294787</a>	7957	750	7	250	delangin (NIPBL) alternative variant cSep08, mRNA.
<a href="#">NIPBL</a>	<a href="#">NIPBL.dSep08</a>	<a href="#">294787</a>	11221	1615	6	227	delangin (25.5 kD) (NIPBL) alternative variant dSep08, mRNA.
<a href="#">NIPBL</a>	<a href="#">NIPBL.eSep08</a>	<a href="#">294787</a>	81763	710	6	150	delangin (NIPBL) alternative variant eSep08, mRNA.
<a href="#">NIPBL</a>	<a href="#">NIPBL.fSep08</a>	<a href="#">294787</a>	79590	562	4	105	delangin (NIPBL) alternative variant fSep08, mRNA.
<a href="#">Nipsnap1</a>	<a href="#">Nipsnap1.aSep08</a>	<a href="#">360971</a>	24116	1880	10	284	4-nitrophenylphosphatase domain non-neuronal protein homolog 1 CRA b (33.3 kD) (Nipsnap1) alternative variant aSep08, complete mRNA.
<a href="#">Nipsnap1</a>	<a href="#">Nipsnap1.bSep08</a>	<a href="#">360971</a>	3838	474	4	88	4-nitrophenylphosphatase domain non-neuronal protein homolog 1 CRA c (10.6 kD) (Nipsnap1) alternative variant bSep08, mRNA.
<a href="#">Nipsnap1</a>	<a href="#">Nipsnap1.eSep08</a>	<a href="#">360971</a>	1761	350	2	37	putative protein (Nipsnap1) alternative variant eSep08, mRNA.
<a href="#">Nipsnap3a</a>	<a href="#">Nipsnap3a.bSep08</a>	<a href="#">313211</a>	4969	695		120	nipsnap homolog 3B (Nipsnap3a) alternative variant bSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.aSep08</a>	<a href="#">306255</a>	36212	5564	21	1653	nischarin (Nisch) alternative variant aSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.bSep08</a>	<a href="#">306255</a>	3174	2357	3	236	nischarin (26.5 kD) (Nisch) alternative variant bSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.cSep08</a>	<a href="#">306255</a>	4851	1982	6	207	nischarin (Nisch) alternative variant cSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.dSep08</a>	<a href="#">306255</a>	3630	619	3	172	nischarin (Nisch) alternative variant dSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.eSep08</a>	<a href="#">306255</a>	2485	727	3	129	nischarin (Nisch) alternative variant eSep08, mRNA.
<a href="#">Nisch</a>	<a href="#">Nisch.fSep08</a>	<a href="#">306255</a>	6026	907	4	124	nischarin (12.9 kD) (Nisch) alternative variant fSep08, mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.cSep08</a>	<a href="#">289222</a>	3373	1271	7	321	nitrilase 1 (35.4 kD) (Nit1) alternative variant cSep08, complete mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.dSep08</a>	<a href="#">289222</a>	2085	964	5	315	nitrilase 1 (Nit1) alternative variant dSep08, mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.eSep08</a>	<a href="#">289222</a>	2136	911	4	230	nitrilase 1 (Nit1) alternative variant eSep08, mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.fSep08</a>	<a href="#">289222</a>	2529	1500	6	226	nitrilase 1 (24.8 kD) (Nit1) alternative variant fSep08, mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.gSep08</a>	<a href="#">289222</a>	2919	1706	6	224	nitrilase 1 (Nit1) alternative variant gSep08, mRNA.
<a href="#">Nit1</a>	<a href="#">Nit1.hSep08</a>	<a href="#">289222</a>	2995	1821	4	161	nitrilase 1 (18.0 kD) (Nit1) alternative variant hSep08, mRNA.
<a href="#">Nit2</a>	<a href="#">Nit2.aSep08</a>	<a href="#">288174</a>	11027	1347	8	312	nitrilase family, member 2 (Nit2) alternative variant aSep08, mRNA.
<a href="#">Nit2</a>	<a href="#">Nit2.cSep08</a>	<a href="#">288174</a>	10758	2048	7	191	nitrilase family, member 2 (21.0 kD) (Nit2) alternative variant cSep08, complete mRNA.
<a href="#">Nkain4</a>	<a href="#">Nkain4.aSep08</a>	<a href="#">296469</a>	20503	838	7	194	na <sup>+</sup> /K <sup>+</sup> transporting ATPase interacting 4 (Nkain4) alternative variant aSep08, mRNA.
<a href="#">Nkd1</a>	<a href="#">Nkd1.bSep08</a>	<a href="#">364952</a>	64810	645	4	214	naked cuticle 1 homolog (Drosophila) (Nkd1) alternative variant bSep08, mRNA.
<a href="#">Nkiras1</a>	<a href="#">Nkiras1.bSep08</a>	<a href="#">305751</a>	8001	910	1	225	NFKB inhibitor interacting Ras-like protein 1 (Nkiras1) alternative variant bSep08, mRNA.

<a href="#">Nkiras2</a>	<a href="#">Nkiras2.bSep08</a>	<a href="#">287707</a>	2931	869	3	191	NFKB inhibitor interacting Ras-like protein 2 (21.5 kD) (Nkiras2) alternative variant bSep08, mRNA.
<a href="#">Nkiras2</a>	<a href="#">Nkiras2.cSep08</a>	<a href="#">287707</a>	4112	1882	3	135	NFKB inhibitor interacting Ras-like protein 2 (14.9 kD) (Nkiras2) alternative variant cSep08, mRNA.
<a href="#">Nkrf</a>	<a href="#">Nkrf.aSep08</a>	<a href="#">298316</a>	13011	531		176	NF-kappaB repressing factor (Nkrf) mRNA.
<a href="#">Nlgn2</a>	<a href="#">Nlgn2.bSep08</a>	<a href="#">117096</a>	414	305	2	88	neuroligin 2 (Nlgn2) alternative variant bSep08, mRNA.
<a href="#">Nlgn2</a>	<a href="#">Nlgn2.cSep08</a>	<a href="#">117096</a>	4055	628	2	63	neuroligin 2 (Nlgn2) alternative variant cSep08, mRNA.
<a href="#">Nlk</a>	<a href="#">Nlk.aSep08</a>	<a href="#">497961</a>	15154	510		170	nemo like kinase (Nlk) mRNA.
<a href="#">Nln</a>	<a href="#">Nln.bSep08</a>	<a href="#">117041</a>	81071	1783	7	386	neurolysin (metallopeptidase M3 family) (Nln) alternative variant bSep08, mRNA.
<a href="#">Nln</a>	<a href="#">Nln.cSep08</a>	<a href="#">117041</a>	12267	537	4	128	neurolysin (metallopeptidase M3 family) (Nln) alternative variant cSep08, mRNA.
<a href="#">Nln</a>	<a href="#">Nln.dSep08</a>	<a href="#">117041</a>	1340	438	2	91	neurolysin (metallopeptidase M3 family) (Nln) alternative variant dSep08, mRNA.
<a href="#">Nln</a>	<a href="#">Nln.eSep08</a>	<a href="#">117041</a>	5519	2068	2	42	neurolysin (metallopeptidase M3 family) (4.8 kD) (Nln) alternative variant eSep08, mRNA.
<a href="#">Nlp</a>	<a href="#">Nlp.aSep08</a>	<a href="#">311529</a>	30516	659		181	ninein-like (Nlp) mRNA.
<a href="#">Nlrp1a</a>	<a href="#">Nlrp1a.aSep08</a>	<a href="#">360557</a>	17506	354		117	NACHT- LRR- PYD-containing protein 1 paralog b (Nlrp1a) mRNA.
<a href="#">Nlrp3</a>	<a href="#">Nlrp3.aSep08</a>	<a href="#">287362</a>	16063	844	6	281	leucine-rich repeat containing protein (Nlrp3) alternative variant aSep08, mRNA.
<a href="#">Nlrp3</a>	<a href="#">Nlrp3.bSep08</a>	<a href="#">287362</a>	15066	928	6	158	leucine-rich repeat containing protein (17.6 kD) (Nlrp3) alternative variant bSep08, mRNA.
<a href="#">Nlrp3</a>	<a href="#">Nlrp3.cSep08</a>	<a href="#">287362</a>	1178	1033	2	109	putative protein of vertebrate origin (Nlrp3) alternative variant cSep08, mRNA.
<a href="#">Nlrp4e</a>	<a href="#">Nlrp4e.aSep08</a>	<a href="#">499069</a>	5917	671		117	containing NLR family pyrin domain 4 (Nlrp4e) mRNA.
<a href="#">Nlrp6</a>	<a href="#">Nlrp6.aSep08</a>	<a href="#">171390</a>	3883	1613		291	vasopressin receptor (Nlrp6) mRNA.
<a href="#">Nmd3</a>	<a href="#">Nmd3.bSep08</a>	<a href="#">310512</a>	25467	1848	14	383	NMD3 homolog (S. cerevisiae) (43.6 kD) (Nmd3) alternative variant bSep08, mRNA.
<a href="#">Nmd3</a>	<a href="#">Nmd3.cSep08</a>	<a href="#">310512</a>	3004	372	1	85	NMD3 homolog (S. cerevisiae) (10.5 kD) (Nmd3) alternative variant cSep08, mRNA.
<a href="#">Nme1</a>	<a href="#">Nme1.aSep08</a>	<a href="#">191575</a>	9696	1094	4	152	non-metastatic cells 1, protein (NM23A) expressed in (17.2 kD) (Nme1) alternative variant aSep08, complete mRNA.
<a href="#">Nme1</a>	<a href="#">Nme1.cSep08</a>	<a href="#">191575</a>	2637	500	1	80	non-metastatic cells 1, protein (NM23A) expressed in (Nme1) alternative variant cSep08, mRNA.
<a href="#">Nme2</a>	<a href="#">Nme2.aSep08</a>	<a href="#">83782</a>	5167	610	4	164	non-metastatic cells 2, protein (NM23B) expressed in (Nme2) alternative variant aSep08, mRNA.
<a href="#">Nme2</a>	<a href="#">Nme2.cSep08</a>	<a href="#">83782</a>	5384	524	4	118	non-metastatic cells 2, protein (NM23B) expressed in (13.4 kD) (Nme2) alternative variant cSep08, mRNA.
<a href="#">Nme6</a>	<a href="#">Nme6.aSep08</a>	<a href="#">58964</a>	6707	1407	3	141	non-metastatic cells 6, protein expressed in (nucleoside-diphosphate kinase) (16.3 kD) (Nme6) alternative variant aSep08, mRNA.
<a href="#">Nme6</a>	<a href="#">Nme6.bSep08</a>	<a href="#">58964</a>	6308	1018	2	129	non-metastatic cells 6, protein expressed in (nucleoside-diphosphate kinase) (Nme6) alternative variant bSep08, mRNA.

<a href="#">Nme7</a>	<a href="#">Nme7.bSep08</a>	<a href="#">171566</a>	24171	377	3	100	non-metastatic cells 7, protein expressed in (nucleoside-diphosphate kinase) (Nme7) alternative variant bSep08, mRNA.
<a href="#">Nme7</a>	<a href="#">Nme7.cSep08</a>	<a href="#">171566</a>	3716	421	3	35	non-metastatic cells 7, protein expressed in (nucleoside-diphosphate kinase) (Nme7) alternative variant cSep08, mRNA.
<a href="#">Nmi</a>	<a href="#">Nmi.aSep08</a>	<a href="#">311021</a>	22719	801	5	242	N-myc (and STAT) interactor (Nmi) alternative variant aSep08, mRNA.
<a href="#">Nmi</a>	<a href="#">Nmi.cSep08</a>	<a href="#">311021</a>	10157	381	2	41	N-myc (and STAT) interactor (Nmi) alternative variant cSep08, mRNA.
<a href="#">Nmnat1</a>	<a href="#">Nmnat1.bSep08</a>	<a href="#">298653</a>	2374	2120	2	75	nicotinamide nucleotide adenyllyltransferase 1 (Nmnat1) alternative variant bSep08, mRNA.
<a href="#">Nmnat2</a>	<a href="#">Nmnat2.bSep08</a>	<a href="#">289095</a>	14643	307	1	97	nicotinamide nucleotide adenyllyltransferase 2 (Nmnat2) alternative variant bSep08, mRNA.
<a href="#">Nmnat3</a>	<a href="#">Nmnat3.aSep08</a>	<a href="#">363118</a>	111832	1855	5	245	nicotinamide nucleotide adenyllyltransferase 3 (27.6 kD) (Nmnat3) alternative variant aSep08, mRNA.
<a href="#">Nmnat3</a>	<a href="#">Nmnat3.bSep08</a>	<a href="#">363118</a>	7867	761	2	153	nicotinamide nucleotide adenyllyltransferase 3 (17.0 kD) (Nmnat3) alternative variant bSep08, mRNA.
<a href="#">Nmnat3</a>	<a href="#">Nmnat3.dSep08</a>	<a href="#">363118</a>	103617	743	4	117	nicotinamide nucleotide adenyllyltransferase 3 (Nmnat3) alternative variant dSep08, mRNA.
<a href="#">Nmral1</a>	<a href="#">Nmral1.aSep08</a>	<a href="#">287063</a>	8342	1150	2	299	NmrA-like and TrkA-N (33.1 kD) (Nmral1) alternative variant aSep08, mRNA.
<a href="#">Nmral1</a>	<a href="#">Nmral1.bSep08</a>	<a href="#">287063</a>	8378	1754	1	197	NmrA-like and TrkA-N (21.6 kD) (Nmral1) alternative variant bSep08, mRNA.
<a href="#">Nmral1</a>	<a href="#">Nmral1.cSep08</a>	<a href="#">287063</a>	8488	1105	1	179	NmrA-like and TrkA-N (19.7 kD) (Nmral1) alternative variant cSep08, mRNA.
<a href="#">Nmral1</a>	<a href="#">Nmral1.dSep08</a>	<a href="#">287063</a>	5604	714	2	149	NmrA-like (Nmral1) alternative variant dSep08, mRNA.
<a href="#">Nmral1</a>	<a href="#">Nmral1.eSep08</a>	<a href="#">287063</a>	8353	1776	1	27	putative protein (2.8 kD) (Nmral1) alternative variant eSep08, mRNA.
<a href="#">Nmt1</a>	<a href="#">Nmt1.bSep08</a>	<a href="#">259274</a>	12954	1684	10	395	N-myristoyltransferase 1 (45.9 kD) (Nmt1) alternative variant bSep08, mRNA.
<a href="#">Nmt1</a>	<a href="#">Nmt1.cSep08</a>	<a href="#">259274</a>	11897	1792	7	216	N-myristoyltransferase 1 (Nmt1) alternative variant cSep08, mRNA.
<a href="#">Nmu</a>	<a href="#">Nmu.bSep08</a>	<a href="#">63887</a>	28164	746	3	137	neuromedin U (Nmu) alternative variant bSep08, mRNA.
<a href="#">Nnat</a>	<a href="#">Nnat.bSep08</a>	<a href="#">94270</a>	1955	764	3	129	neuronatin (Nnat) alternative variant bSep08, mRNA.
<a href="#">Nnat</a>	<a href="#">Nnat.eSep08</a>	<a href="#">94270</a>	1825	715	4	79	neuronatin (Nnat) alternative variant eSep08, mRNA.
<a href="#">Nnt</a>	<a href="#">Nnt.bSep08</a>	<a href="#">310378</a>	14667	790	4	262	nicotinamide nucleotide transhydrogenase (Nnt) alternative variant bSep08, mRNA.
<a href="#">Nnt</a>	<a href="#">Nnt.cSep08</a>	<a href="#">310378</a>	13781	703	4	198	nicotinamide nucleotide transhydrogenase (Nnt) alternative variant cSep08, mRNA.
<a href="#">Nnt</a>	<a href="#">Nnt.dSep08</a>	<a href="#">310378</a>	497	355	2	46	nicotinamide nucleotide transhydrogenase (Nnt) alternative variant dSep08, mRNA.
<a href="#">Nob1</a>	<a href="#">Nob1.bSep08</a>	<a href="#">291996</a>	7972	906	7	295	NIN one binding protein like (Nob1) alternative variant bSep08, mRNA.
<a href="#">Nob1</a>	<a href="#">Nob1.cSep08</a>	<a href="#">291996</a>	8823	1075	7	273	NIN one binding protein like (31.0 kD) (Nob1) alternative variant cSep08, mRNA.

<a href="#">Nob1</a>	<a href="#">Nob1.dSep08</a>	<a href="#">291996</a>	7267	810	7	234	NIN one binding protein like (Nob1) alternative variant dSep08, mRNA.
<a href="#">Nob1</a>	<a href="#">Nob1.eSep08</a>	<a href="#">291996</a>	3865	810	2	59	binding protein 1 like (Nob1) alternative variant eSep08, mRNA.
<a href="#">noby</a>	<a href="#">noby.aSep08</a>		9401	652		33	putative protein (noby) mRNA.
<a href="#">Noc4l</a>	<a href="#">Noc4l.aSep08</a>	<a href="#">360828</a>	1163	659	5	164	nucleolar complex associated 4 homolog ( <i>S. cerevisiae</i> ) (Noc4l) alternative variant aSep08, mRNA.
<a href="#">Noc4l</a>	<a href="#">Noc4l.bSep08</a>	<a href="#">360828</a>	1806	732	6	132	nucleolar complex associated 4 homolog ( <i>S. cerevisiae</i> ) (15.0 kD) (Noc4l) alternative variant bSep08, mRNA.
<a href="#">Noc4l</a>	<a href="#">Noc4l.cSep08</a>	<a href="#">360828</a>	1600	892	5	125	nucleolar complex associated 4 homolog ( <i>S. cerevisiae</i> ) (13.7 kD) (Noc4l) alternative variant cSep08, mRNA.
<a href="#">nochy</a>	<a href="#">nochy.aSep08</a>		3541	718		239	ATP GTP binding protein-like 2 (nochy) mRNA.
<a href="#">Nod1</a>	<a href="#">Nod1.bSep08</a>	<a href="#">500133</a>	5735	1455	3	63	putative protein of vertebrate origin (7.3 kD) (Nod1) alternative variant bSep08, mRNA.
<a href="#">nodar</a>	<a href="#">nodar.aSep08</a>		2723	1042		80	putative protein (9.1 kD) (nodar) mRNA.
<a href="#">NODP.0</a>	<a href="#">NODP.0.aSep08</a>		4002	391		130	notch homolog 3 (NODP.0) mRNA.
<a href="#">nofer</a>	<a href="#">nofer.bSep08</a>		724	532	2	72	putative protein (8.0 kD) (nofer) alternative variant bSep08, mRNA.
<a href="#">nofer</a>	<a href="#">nofer.cSep08</a>		689	488	3	59	putative protein (nofer) alternative variant cSep08, mRNA.
<a href="#">noflo</a>	<a href="#">noflo.aSep08</a>		4125	1106		63	gag protein like (noflo) mRNA.
<a href="#">noflu</a>	<a href="#">noflu.aSep08</a>		72407	488		43	putative protein (5.2 kD) (noflu) mRNA.
<a href="#">nogar</a>	<a href="#">nogar.aSep08</a>		7558	495	3	164	putative protein of vertebrate origin (nogar) alternative variant aSep08, mRNA.
<a href="#">nogar</a>	<a href="#">nogar.bSep08</a>		2062	398	2	92	putative protein of vertebrate origin (nogar) alternative variant bSep08, mRNA.
<a href="#">nogar</a>	<a href="#">nogar.cSep08</a>		8990	474	5	58	putative protein of vertebrate origin (nogar) alternative variant cSep08, mRNA.
<a href="#">nogar</a>	<a href="#">nogar.dSep08</a>		9024	355	4	49	putative protein of vertebrate origin (nogar) alternative variant dSep08, mRNA.
<a href="#">noja</a>	<a href="#">noja.aSep08</a>		4927	666		222	repeat 18 (noja) mRNA.
<a href="#">nojey</a>	<a href="#">nojey.aSep08</a>		790	609		81	putative protein (nojey) mRNA.
<a href="#">nokee</a>	<a href="#">nokee.aSep08</a>		893	342		113	smg-7 homolog (nokee) mRNA.
<a href="#">nokler</a>	<a href="#">nokler.aSep08</a>		2584	249		43	putative protein (4.9 kD) (nokler) mRNA.
<a href="#">Nol1</a>	<a href="#">Nol1.aSep08</a>	<a href="#">314969</a>	4452	1345		410	nucleolar protein 1 (Nol1) mRNA.
<a href="#">Nol3</a>	<a href="#">Nol3.bSep08</a>	<a href="#">85383</a>	4161	1536	4	218	nucleolar protein 3 (apoptosis repressor with CARD domain) (24.2 kD) (Nol3) alternative variant bSep08, mRNA.
<a href="#">Nol5</a>	<a href="#">Nol5.bSep08</a>	<a href="#">60373</a>	8878	1908	6	187	nucleolar protein 5 CRA a (Nol5) alternative variant bSep08, mRNA.
<a href="#">Nol5</a>	<a href="#">Nol5.dSep08</a>	<a href="#">60373</a>	5159	379	5	95	nucleolar protein 5 CRA a (Nol5) alternative variant dSep08, mRNA.
<a href="#">Nol5</a>	<a href="#">Nol5.eSep08</a>	<a href="#">60373</a>	5688	350	3	49	nucleolar protein (Nol5) alternative variant eSep08, mRNA.
<a href="#">Nol5a</a>	<a href="#">Nol5a.bSep08</a>	<a href="#">362214</a>	3112	1991	7	252	nucleolar protein 5A CRA d (27.9 kD) (Nol5a) alternative variant bSep08, mRNA.

<a href="#">Nol5a</a>	<a href="#">Nol5a.cSep08</a>	<a href="#">362214</a>	2654	1633	5	197	nucleolar protein 5A (Nol5a) alternative variant cSep08, mRNA.
<a href="#">Nol5a</a>	<a href="#">Nol5a.dSep08</a>	<a href="#">362214</a>	1721	1107	2	188	CRA d like (21.1 kD) (Nol5a) alternative variant dSep08, mRNA.
<a href="#">Nol5a</a>	<a href="#">Nol5a.eSep08</a>	<a href="#">362214</a>	2179	638	6	160	nucleolar protein 5A (Nol5a) alternative variant eSep08, mRNA.
<a href="#">Nol6</a>	<a href="#">Nol6.aSep08</a>	<a href="#">313167</a>	6474	3738	15	622	nucleolar protein family 6 (RNA-associated) (Nol6) alternative variant aSep08, mRNA.
<a href="#">Nol6</a>	<a href="#">Nol6.bSep08</a>	<a href="#">313167</a>	9566	1749	8	299	nucleolar protein family 6 (RNA-associated) (34.0 kD) (Nol6) alternative variant bSep08, mRNA.
<a href="#">Nol6</a>	<a href="#">Nol6.cSep08</a>	<a href="#">313167</a>	2137	817	6	272	nucleolar protein family 6 (RNA-associated) (Nol6) alternative variant cSep08, mRNA.
<a href="#">Nol6</a>	<a href="#">Nol6.dSep08</a>	<a href="#">313167</a>	1924	678	5	225	nucleolar protein family 6 (RNA-associated) (Nol6) alternative variant dSep08, mRNA.
<a href="#">Nol7</a>	<a href="#">Nol7.aSep08</a>	<a href="#">498727</a>	3202	1149	6	229	nucleolar protein 7 (Nol7) alternative variant aSep08, mRNA.
<a href="#">Nol7</a>	<a href="#">Nol7.bSep08</a>	<a href="#">498727</a>	799	723	2	130	nucleolar protein 7 (14.4 kD) (Nol7) alternative variant bSep08, mRNA.
<a href="#">Nol7</a>	<a href="#">Nol7.cSep08</a>	<a href="#">498727</a>	2707	730	5	126	nucleolar protein 7 (Nol7) alternative variant cSep08, mRNA.
<a href="#">Nol8</a>	<a href="#">Nol8.bSep08</a>	<a href="#">361221</a>	7275	1273	6	423	nucleolar protein 8 (Nol8) alternative variant bSep08, mRNA.
<a href="#">Nol8</a>	<a href="#">Nol8.cSep08</a>	<a href="#">361221</a>	2614	554	4	144	nucleolar protein 8 CRA b (Nol8) alternative variant cSep08, mRNA.
<a href="#">Nol8</a>	<a href="#">Nol8.dSep08</a>	<a href="#">361221</a>	3730	930	4	124	nucleolar protein 8 (14.4 kD) (Nol8) alternative variant dSep08, mRNA.
<a href="#">Nol9</a>	<a href="#">Nol9.aSep08</a>	<a href="#">313744</a>	8294	2419		308	nucleolar protein 9 (Nol9) mRNA.
<a href="#">Nol10</a>	<a href="#">Nol10.bSep08</a>	<a href="#">313981</a>	5134	366	1	28	nucleolar protein 10 (Nol10) alternative variant bSep08, mRNA.
<a href="#">Nol11</a>	<a href="#">Nol11.aSep08</a>	<a href="#">688885</a>	12786	1408	11	468	nucleolar protein 11 (Nol11) alternative variant aSep08, mRNA.
<a href="#">Nol11</a>	<a href="#">Nol11.bSep08</a>	<a href="#">688885</a>	2701	522	2	108	nucleolar protein 11 (Nol11) alternative variant bSep08, mRNA.
<a href="#">Nol14</a>	<a href="#">Nol14.bSep08</a>	<a href="#">289724</a>	5162	816	3	264	nucleolar protein 14 (Nol14) alternative variant bSep08, mRNA.
<a href="#">Nol14</a>	<a href="#">Nol14.cSep08</a>	<a href="#">289724</a>	2625	665	1	191	nucleolar protein 14 precursor (22.3 kD) (Nol14) alternative variant cSep08, mRNA.
<a href="#">Nol14</a>	<a href="#">Nol14.dSep08</a>	<a href="#">289724</a>	5884	891	4	156	nucleolar protein 14 (18.4 kD) (Nol14) alternative variant dSep08, mRNA.
<a href="#">Nola1</a>	<a href="#">Nola1.bSep08</a>	<a href="#">499709</a>	6890	1188	6	222	nucleolar protein family A, member 1 (H/ACA small nucleolar RNPs) (22.7 kD) (Nola1) alternative variant bSep08, mRNA.
<a href="#">Nola2</a>	<a href="#">Nola2.aSep08</a>	<a href="#">287273</a>	3359	746	1	153	nucleolar protein family A, member 2 (17.3 kD) (Nola2) alternative variant aSep08, mRNA.
<a href="#">Nolc1</a>	<a href="#">Nolc1.bSep08</a>	<a href="#">64896</a>	7397	983	8	322	nucleolar and coiled-body phosphoprotein 1 (Nolc1) alternative variant bSep08, mRNA.



<a href="#">Nolc1</a>	<a href="#">Nolc1.cSep08</a>	<a href="#">64896</a>	2997	309	2	73	nucleolar and coiled-body phosphoprotein 1 (Nolc1) alternative variant cSep08, mRNA.
<a href="#">Nolc1</a>	<a href="#">Nolc1.dSep08</a>	<a href="#">64896</a>	1405	415	2	37	nucleolar and coiled-body phosphoprotein 1 (4.5 kD) (Nolc1) alternative variant dSep08, mRNA.
<a href="#">noloy</a>	<a href="#">noloy.aSep08</a>		1450	415		137	chordin (noloy) mRNA.
<a href="#">nomee</a>	<a href="#">nomee.aSep08</a>		1854	161		53	putative protein of vertebrate origin (nomee) mRNA.
<a href="#">nomer</a>	<a href="#">nomer.aSep08</a>		5790	671		101	putative protein (nomer) mRNA.
<a href="#">Nomo1</a>	<a href="#">Nomo1.bSep08</a>	<a href="#">361578</a>	11558	838	8	279	nodal modulator 1 (Nomo1) alternative variant bSep08, mRNA.
<a href="#">Nomo1</a>	<a href="#">Nomo1.cSep08</a>	<a href="#">361578</a>	6874	636	6	212	nodal modulator 1 (Nomo1) alternative variant cSep08, mRNA.
<a href="#">Nomo1</a>	<a href="#">Nomo1.dSep08</a>	<a href="#">361578</a>	5947	418	4	86	nodal modulator 1 (Nomo1) alternative variant dSep08, mRNA.
<a href="#">Nono</a>	<a href="#">Nono.bSep08</a>	<a href="#">317259</a>	5051	1215	7	172	non-POU-domain-containing, octamer-binding protein (21.3 kD) (Nono) alternative variant bSep08, mRNA.
<a href="#">Nono</a>	<a href="#">Nono.cSep08</a>	<a href="#">317259</a>	1476	398	3	102	non-POU-domain-containing, octamer-binding protein (Nono) alternative variant cSep08, mRNA.
<a href="#">nonoy</a>	<a href="#">nonoy.aSep08</a>		100372	693	2	12	putative protein (1.4 kD) (nonoy) alternative variant aSep08, mRNA.
<a href="#">nonoy</a>	<a href="#">nonoy.bSep08</a>		100190	402	1	64	putative protein (nonoy) alternative variant bSep08, mRNA.
<a href="#">Nope</a>	<a href="#">Nope.bSep08</a>	<a href="#">363081</a>	8088	1786	2	486	neighbor of Punc E11 (Nope) alternative variant bSep08, mRNA.
<a href="#">nopor</a>	<a href="#">nopor.aSep08</a>		2387	669		144	solute carrier family 24 member 1 (nopor) mRNA.
<a href="#">norby</a>	<a href="#">norby.aSep08</a>		917	369		109	interleukin 2 receptor (norby) mRNA.
<a href="#">norchy</a>	<a href="#">norchy.aSep08</a>		1286	264		15	putative protein (norchy) mRNA.
<a href="#">nordar</a>	<a href="#">nordar.aSep08</a>		8894	1377		92	putative protein (9.9 kD) (nordar) mRNA.
<a href="#">norfer</a>	<a href="#">norfer.aSep08</a>		7336	400		65	putative protein (norfer) mRNA.
<a href="#">norflo</a>	<a href="#">norflo.aSep08</a>		560	305		39	putative protein (4.5 kD) (norflo) mRNA.
<a href="#">norflu</a>	<a href="#">norflu.aSep08</a>		3271	945		113	CRA a (norflu) mRNA.
<a href="#">norgar</a>	<a href="#">norgar.aSep08</a>		7747	385		128	putative protein, with a transmembrane domain, of eukaryotic origin (norgar) mRNA.
<a href="#">norja</a>	<a href="#">norja.aSep08</a>		12764	368		34	putative protein (3.8 kD) (norja) mRNA.
<a href="#">norjey</a>	<a href="#">norjey.aSep08</a>		1885	243		17	putative protein (norjey) mRNA.
<a href="#">norkee</a>	<a href="#">norkee.aSep08</a>		1813	497		61	CRA b like (6.6 kD) (norkee) mRNA.
<a href="#">norkler</a>	<a href="#">norkler.aSep08</a>		4206	805		52	sarcoma antigen like (norkler) mRNA.
<a href="#">norloy</a>	<a href="#">norloy.aSep08</a>		8941	593		37	putative protein (norloy) mRNA.
<a href="#">normee</a>	<a href="#">normee.aSep08</a>		2082	457		152	neuralized 2 (normee) mRNA.
<a href="#">normer</a>	<a href="#">normer.aSep08</a>		683	332		110	CRA b like (normer) mRNA.
<a href="#">normoy</a>	<a href="#">normoy.aSep08</a>		52519	1388		34	putative protein (3.7 kD) (normoy) mRNA.
<a href="#">norpor</a>	<a href="#">norpor.aSep08</a>		21987	297		48	putative protein (norpor) mRNA.
<a href="#">norsa</a>	<a href="#">norsa.aSep08</a>		3665	408		83	putative protein (norsa) mRNA.
<a href="#">norshee</a>	<a href="#">norshee.bSep08</a>		1270	702	2	77	putative protein (norshee) alternative variant bSep08, mRNA.

<a href="#">norshee</a>	<a href="#">norshee.cSep08</a>		754	523	2	75	putative protein (norshee) alternative variant cSep08, mRNA.
<a href="#">nortu</a>	<a href="#">nortu.aSep08</a>		10764	371		31	putative protein (nortu) mRNA.
<a href="#">norvar</a>	<a href="#">norvar.aSep08</a>		20129	413		137	receptor protein tyrosine phosphatase hPTP-J (norvar) mRNA.
<a href="#">norway</a>	<a href="#">norway.aSep08</a>		2165	245		46	putative protein (norway) mRNA.
<a href="#">Nos1</a>	<a href="#">Nos1.bSep08</a>	<a href="#">24598</a>	6689	589	1	195	nitric oxide synthase 1, neuronal (Nos1) alternative variant bSep08, mRNA.
<a href="#">Nos1ap</a>	<a href="#">Nos1ap.bSep08</a>	<a href="#">192363</a>	1861	407	2	87	nitric oxide synthase 1 (neuronal) adaptor protein (Nos1ap) alternative variant bSep08, mRNA.
<a href="#">Nos2</a>	<a href="#">Nos2.bSep08</a>	<a href="#">24599</a>	4503	299	3	99	nitric oxide synthase 2, inducible, macrophage (Nos2) alternative variant bSep08, mRNA.
<a href="#">Nos3</a>	<a href="#">Nos3.bSep08</a>	<a href="#">24600</a>	3500	625	1	143	nitric oxide synthase 3, endothelial cell (Nos3) alternative variant bSep08, mRNA.
<a href="#">nosa</a>	<a href="#">nosa.aSep08</a>		1622	1006		77	putative protein (7.9 kD) (nosa) mRNA.
<a href="#">noshee</a>	<a href="#">noshee.aSep08</a>		2207	662		65	putative protein (noshee) mRNA.
<a href="#">Nosip</a>	<a href="#">Nosip.aSep08</a>	<a href="#">292894</a>	16556	1078	9	301	nitric oxide synthase interacting protein (33.3 kD) (Nosip) alternative variant aSep08, complete mRNA.
<a href="#">Nosip</a>	<a href="#">Nosip.cSep08</a>	<a href="#">292894</a>	2540	820	6	260	nitric oxide synthase interacting protein (Nosip) alternative variant cSep08, mRNA.
<a href="#">Nosip</a>	<a href="#">Nosip.eSep08</a>	<a href="#">292894</a>	14008	779	3	59	nitric oxide synthase interacting protein (6.5 kD) (Nosip) alternative variant eSep08, mRNA.
<a href="#">Nosip</a>	<a href="#">Nosip.fSep08</a>	<a href="#">292894</a>	3690	695	3	59	nitric oxide synthase interacting protein (6.5 kD) (Nosip) alternative variant fSep08, mRNA.
<a href="#">Nostrin</a>	<a href="#">Nostrin.bSep08</a>	<a href="#">311111</a>	6115	472	3	111	nitric oxide synthase trafficker (Nostrin) alternative variant bSep08, mRNA.
<a href="#">Nostrin</a>	<a href="#">Nostrin.cSep08</a>	<a href="#">311111</a>	7055	378	4	80	nitric oxide synthase trafficker (Nostrin) alternative variant cSep08, mRNA.
<a href="#">Notch1</a>	<a href="#">Notch1.bSep08</a>	<a href="#">25496</a>	7259	3885	6	725	notch gene homolog 1 (Drosophila) (Notch1) alternative variant bSep08, mRNA.
<a href="#">Notch1</a>	<a href="#">Notch1.cSep08</a>	<a href="#">25496</a>	1420	632	2	210	notch gene homolog 1 (Drosophila) (Notch1) alternative variant cSep08, mRNA.
<a href="#">Notch1</a>	<a href="#">Notch1.dSep08</a>	<a href="#">25496</a>	1752	445	4	148	notch gene homolog 1 (Drosophila) (Notch1) alternative variant dSep08, mRNA.
<a href="#">Notch1</a>	<a href="#">Notch1.eSep08</a>	<a href="#">25496</a>	1609	409	3	136	notch gene homolog 1 (Drosophila) (Notch1) alternative variant eSep08, mRNA.
<a href="#">Notch2</a>	<a href="#">Notch2.bSep08</a>	<a href="#">29492</a>	2982	468	3	84	notch homolog 2 (Drosophila) (Notch2) alternative variant bSep08, mRNA.
<a href="#">Notch2</a>	<a href="#">Notch2.cSep08</a>	<a href="#">29492</a>	985	632	2	79	notch homolog 2 (Drosophila) (8.5 kD) (Notch2) alternative variant cSep08, mRNA.
<a href="#">Notch3</a>	<a href="#">Notch3.aSep08</a>	<a href="#">56761</a>	3448	623		207	notch homolog 3 (Drosophila) (Notch3) mRNA.
<a href="#">Notch4</a>	<a href="#">Notch4.bSep08</a>	<a href="#">406162</a>	2849	2226	4	248	notch homolog 4 (26.6 kD) (Notch4) alternative variant bSep08, mRNA.
<a href="#">Notch4</a>	<a href="#">Notch4.cSep08</a>	<a href="#">406162</a>	1588	779	5	228	notch homolog 4 (Notch4) alternative variant cSep08, mRNA.

<a href="#">Notch4</a>	<a href="#">Notch4.dSep08</a>	<a href="#">406162</a>	737	551	2	130	notch homolog 4 (Notch4) alternative variant dSep08, mRNA.
<a href="#">Notch4</a>	<a href="#">Notch4.fSep08</a>	<a href="#">406162</a>	1908	780	3	88	notch homolog 4 CRA f precursor (9.7 kD) (Notch4) alternative variant fSep08, mRNA.
<a href="#">notu</a>	<a href="#">notu.aSep08</a>		1841	1007		181	mutS homolog 6 like (notu) alternative variant aSep08, mRNA.
<a href="#">notu</a>	<a href="#">notu.bSep08</a>		1291	456		151	dna mismatch repair protein msh6 like (notu) alternative variant bSep08, mRNA.
<a href="#">Nova1</a>	<a href="#">Nova1.aSep08</a>	<a href="#">298992</a>	120702	1115	1	371	neuro-oncological ventral antigen 1 (Nova1) alternative variant aSep08, mRNA.
<a href="#">Nova1</a>	<a href="#">Nova1.bSep08</a>	<a href="#">298992</a>	118630	523	2	72	neuro-oncological ventral antigen 1 (Nova1) alternative variant bSep08, mRNA.
<a href="#">novar</a>	<a href="#">novar.aSep08</a>		9603	730		148	putative nuclear protein (16.5 kD) (novar) mRNA.
<a href="#">nowey</a>	<a href="#">nowey.aSep08</a>		2257	284		42	putative protein (nowey) mRNA.
<a href="#">Noxo1</a>	<a href="#">Noxo1.aSep08</a>	<a href="#">302976</a>	2202	1785	4	547	NADPH oxidase organizer 1 (Noxo1) alternative variant aSep08, mRNA.
<a href="#">Noxo1</a>	<a href="#">Noxo1.cSep08</a>	<a href="#">302976</a>	860	419	2	92	NADPH oxidase organizer 1 (Noxo1) alternative variant cSep08, mRNA.
<a href="#">noyby</a>	<a href="#">noyby.aSep08</a>		1103	390		130	mediator of RNA polymerase II transcription homolog (noyby) mRNA.
<a href="#">noychy</a>	<a href="#">noychy.aSep08</a>		5819	323		107	cytoskeleton associated protein 5 (noychy) mRNA.
<a href="#">noydar</a>	<a href="#">noydar.bSep08</a>		3084	465	2	86	putative protein (noydar) alternative variant bSep08, mRNA.
<a href="#">noyfer</a>	<a href="#">noyfer.aSep08</a>		562	277	1	49	putative protein (noyfer) alternative variant aSep08, mRNA.
<a href="#">noyfer</a>	<a href="#">noyfer.bSep08</a>		493	271	1	19	putative protein (noyfer) alternative variant bSep08, mRNA.
<a href="#">noyflo</a>	<a href="#">noyflo.aSep08</a>		491	400		47	putative protein (noyflo) mRNA.
<a href="#">noyflu</a>	<a href="#">noyflu.aSep08</a>		1778	708		45	putative protein (5.4 kD) (noyflu) mRNA.
<a href="#">noygar</a>	<a href="#">noygar.aSep08</a>		1589	706	2	31	putative protein (noygar) alternative variant aSep08, mRNA.
<a href="#">noyja</a>	<a href="#">noyja.aSep08</a>		20867	851		41	putative protein (noyja) mRNA.
<a href="#">noyjey</a>	<a href="#">noyjey.aSep08</a>		1383	421		40	putative protein (4.3 kD) (noyjey) mRNA.
<a href="#">noykee</a>	<a href="#">noykee.aSep08</a>		3344	295		66	putative protein (noykee) mRNA.
<a href="#">noykler</a>	<a href="#">noykler.aSep08</a>		1602	263		69	putative protein (noykler) mRNA.
<a href="#">noyloy</a>	<a href="#">noyloy.aSep08</a>		7861	709		34	putative protein (3.9 kD) (noyloy) mRNA.
<a href="#">noymee</a>	<a href="#">noymee.aSep08</a>		850	402		100	putative protein (noymee) mRNA.
<a href="#">noymer</a>	<a href="#">noymer.aSep08</a>		389	306		101	putative protein of vertebrate origin (noymer) mRNA.
<a href="#">noynoy</a>	<a href="#">noynoy.aSep08</a>		2619	392		130	tensin (noynoy) mRNA.
<a href="#">noypor</a>	<a href="#">noypor.aSep08</a>		23447	349		95	probable E3 ubiquitin-protein ligase Herc1 like (noypor) mRNA.
<a href="#">noysa</a>	<a href="#">noysa.aSep08</a>		8579	671		95	putative cytoplasmic protein (10.8 kD) (noysa) mRNA.
<a href="#">noyshee</a>	<a href="#">noyshee.aSep08</a>		96147	473	4	154	double cortin calcium calmodulin-dependent protein kinase-like 1 CRA a (noyshee) alternative variant aSep08, mRNA.
<a href="#">noyshee</a>	<a href="#">noyshee.bSep08</a>		17659	451	3	101	double kinase-like 1 (noyshee) alternative variant bSep08, mRNA.

<a href="#">noyshee</a>	<a href="#">noyshee.cSep08</a>		4718	398	2	76	activity neurotransmitter-induced early gene protein 4 like (noyshee) alternative variant cSep08, mRNA.
<a href="#">noytu</a>	<a href="#">noytu.aSep08</a>		1881	409		66	putative protein (7.6 kD) (noytu) mRNA.
<a href="#">noyvar</a>	<a href="#">noyvar.aSep08</a>		1222	415		138	receptor protein tyrosine phosphatase hPTP-J (noyvar) mRNA.
<a href="#">noywey</a>	<a href="#">noywey.aSep08</a>		3575	368		67	putative protein (7.3 kD) (noywey) mRNA.
<a href="#">Np</a>	<a href="#">Np.aSep08</a>	<a href="#">290029</a>	7664	1361	6	289	nucleoside phosphorylase (32.3 kD) (Np) alternative variant aSep08, complete mRNA.
<a href="#">Np</a>	<a href="#">Np.bSep08</a>	<a href="#">290029</a>	7648	1726	5	248	nucleoside phosphorylase (27.6 kD) (Np) alternative variant bSep08, complete mRNA.
<a href="#">Np</a>	<a href="#">Np.dSep08</a>	<a href="#">290029</a>	2429	962	5	158	nucleoside phosphorylase (17.6 kD) (Np) alternative variant dSep08, mRNA.
<a href="#">Np</a>	<a href="#">Np.eSep08</a>	<a href="#">290029</a>	3759	264	2	67	nucleoside phosphorylase (Np) alternative variant eSep08, mRNA.
<a href="#">Npal2</a>	<a href="#">Npal2.bSep08</a>	<a href="#">362899</a>	13967	309	3	103	putative protein, with at least 2 transmembrane domains, of fungal and metazoan origin (Npal2) alternative variant bSep08, mRNA.
<a href="#">Npal2</a>	<a href="#">Npal2.cSep08</a>	<a href="#">362899</a>	10307	368	1	25	putative protein (2.8 kD) (Npal2) alternative variant cSep08, mRNA.
<a href="#">Npal3</a>	<a href="#">Npal3.aSep08</a>	<a href="#">502990</a>	14436	585	3	80	putative protein of metazoan origin (Npal3) alternative variant aSep08, mRNA.
<a href="#">Npal3</a>	<a href="#">Npal3.bSep08</a>	<a href="#">502990</a>	4050	285	1	34	putative protein (Npal3) alternative variant bSep08, mRNA.
<a href="#">Npap60</a>	<a href="#">Npap60.bSep08</a>	<a href="#">25497</a>	11505	1035	5	344	nuclear pore associated protein (Npap60) alternative variant bSep08, mRNA.
<a href="#">Npap60</a>	<a href="#">Npap60.cSep08</a>	<a href="#">25497</a>	2265	671	3	189	nuclear pore associated protein (Npap60) alternative variant cSep08, mRNA.
<a href="#">Npas1</a>	<a href="#">Npas1.cSep08</a>	<a href="#">308387</a>	4203	392	3	130	neuronal PAS domain protein 1 (Npas1) alternative variant cSep08, mRNA.
<a href="#">Npas2</a>	<a href="#">Npas2.aSep08</a>	<a href="#">316351</a>	39500	3079	13	599	neuronal PAS domain protein 2 (Npas2) alternative variant aSep08, mRNA.
<a href="#">Npas2</a>	<a href="#">Npas2.cSep08</a>	<a href="#">316351</a>	2738	531	2	169	neuronal PAS domain protein 2 (Npas2) alternative variant cSep08, mRNA.
<a href="#">Npas4</a>	<a href="#">Npas4.bSep08</a>	<a href="#">266734</a>	535	415	1	138	neuronal PAS domain protein 4 (Npas4) alternative variant bSep08, mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.aSep08</a>	<a href="#">296562</a>	5658	1530	7	391	neural proliferation differentiation control 1 CRA d (Npdc1) alternative variant aSep08, complete mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.cSep08</a>	<a href="#">296562</a>	4654	751	6	230	neural proliferation differentiation control 1 (Npdc1) alternative variant cSep08, mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.dSep08</a>	<a href="#">296562</a>	4556	760	5	220	neural proliferation differentiation control 1 (Npdc1) alternative variant dSep08, mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.eSep08</a>	<a href="#">296562</a>	3189	2039	3	163	neural proliferation differentiation control 1 (Npdc1) alternative variant eSep08, mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.fSep08</a>	<a href="#">296562</a>	4333	472	3	106	neural proliferation differentiation control 1 CRA d (Npdc1) alternative variant fSep08, mRNA.
<a href="#">Npdc1</a>	<a href="#">Npdc1.gSep08</a>	<a href="#">296562</a>	1344	1241	2	104	neural proliferation differentiation control 1 CRA d (Npdc1) alternative variant gSep08, mRNA.

<a href="#">Npdc1</a>	<a href="#">Npdc1.hSep08</a>	<a href="#">296562</a>	2996	435	2	98	neural proliferation differentiation control 1 CRA e (10.5 kD) (Npdc1) alternative variant hSep08, mRNA.
<a href="#">Npepl1</a>	<a href="#">Npepl1.aSep08</a>	<a href="#">311671</a>	12559	2131	5	556	aminopeptidase-like 1 CRA a (Npepl1) alternative variant aSep08, mRNA.
<a href="#">Npepl1</a>	<a href="#">Npepl1.bSep08</a>	<a href="#">311671</a>	10704	1106	3	368	aminopeptidase-like 1 CRA a (Npepl1) alternative variant bSep08, mRNA.
<a href="#">Npepl1</a>	<a href="#">Npepl1.cSep08</a>	<a href="#">311671</a>	11400	1359	2	292	aminopeptidase-like 1 CRA a (Npepl1) alternative variant cSep08, mRNA.
<a href="#">Npepl1</a>	<a href="#">Npepl1.dSep08</a>	<a href="#">311671</a>	2387	760	1	102	aminopeptidase-like 1 CRA a (Npepl1) alternative variant dSep08, mRNA.
<a href="#">Npepps</a>	<a href="#">Npepps.fSep08</a>	<a href="#">50558</a>	3709	378	5	20	aminopeptidase puromycin sensitive (2.2 kD) (Npepps) alternative variant fSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.aSep08</a>	<a href="#">296136</a>	50171	2100	18	636	nephronophthisis 1 (Nphp1) alternative variant aSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.aSep08</a>	<a href="#">680233</a>	50171	2100	18	636	nephronophthisis 1 (Nphp1) alternative variant aSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.cSep08</a>	<a href="#">296136</a>	19878	828	10	275	nephronophthisis 1 homolog CRA b (Nphp1) alternative variant cSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.cSep08</a>	<a href="#">680233</a>	19878	828	10	275	nephronophthisis 1 homolog CRA b (Nphp1) alternative variant cSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.dSep08</a>	<a href="#">296136</a>	16144	735	8	244	nephronophthisis 1 (Nphp1) alternative variant dSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.dSep08</a>	<a href="#">680233</a>	16144	735	8	244	nephronophthisis 1 (Nphp1) alternative variant dSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.eSep08</a>	<a href="#">296136</a>	13083	560	6	186	nephronophthisis 1 (Nphp1) alternative variant eSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.eSep08</a>	<a href="#">680233</a>	13083	560	6	186	nephronophthisis 1 (Nphp1) alternative variant eSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.fSep08</a>	<a href="#">296136</a>	7137	733	4	180	nephronophthisis 1 (Nphp1) alternative variant fSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.fSep08</a>	<a href="#">680233</a>	7137	733	4	180	nephronophthisis 1 (Nphp1) alternative variant fSep08, mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.gSep08</a>	<a href="#">296136</a>	857	443	2	81	putative protein (8.5 kD) (Nphp1) alternative variant gSep08, complete mRNA.
<a href="#">Nphp1</a>	<a href="#">Nphp1.gSep08</a>	<a href="#">680233</a>	857	443	2	81	putative protein (8.5 kD) (Nphp1) alternative variant gSep08, complete mRNA.
<a href="#">Nphp3</a>	<a href="#">Nphp3.aSep08</a>	<a href="#">363126</a>	3297	1790		142	nephronophthisis 3 (adolescent) (Nphp3) mRNA.
<a href="#">Nphp4</a>	<a href="#">Nphp4.bSep08</a>	<a href="#">313749</a>	7865	2120	10	530	nephronophthisis 4 (59.7 kD) (Nphp4) alternative variant bSep08, mRNA.
<a href="#">Nphp4</a>	<a href="#">Nphp4.cSep08</a>	<a href="#">313749</a>	9496	572	4	190	nephronophthisis 4 (Nphp4) alternative variant cSep08, mRNA.
<a href="#">Nphp4</a>	<a href="#">Nphp4.dSep08</a>	<a href="#">313749</a>	11130	841	3	114	nephronophthisis 4 (Nphp4) alternative variant dSep08, mRNA.
<a href="#">Nphs1</a>	<a href="#">Nphs1.bSep08</a>	<a href="#">64563</a>	10054	1908		88	nephrosis 1 homolog, nephrin (human) (Nphs1) alternative variant bSep08, mRNA.
<a href="#">Npl</a>	<a href="#">Npl.bSep08</a>	<a href="#">304860</a>	30536	690	7	160	N-acetylneuraminase pyruvate lyase (Npl) alternative variant bSep08, mRNA.

<a href="#">Npl</a>	<a href="#">Npl.cSep08</a>	<a href="#">304860</a>	23866	662	4	106	N-acetylneuraminatase pyruvate lyase (Npl) alternative variant cSep08, mRNA.
<a href="#">Npl</a>	<a href="#">Npl.dSep08</a>	<a href="#">304860</a>	43110	1785	6	46	N-acetylneuraminatase pyruvate lyase (5.0 kD) (Npl) alternative variant dSep08, complete mRNA.
<a href="#">Npm1</a>	<a href="#">Npm1.bSep08</a>	<a href="#">25498</a>	7999	1155	6	257	nucleophosmin 1 (28.4 kD) (Npm1) alternative variant bSep08, complete mRNA.
<a href="#">Npm1</a>	<a href="#">Npm1.cSep08</a>	<a href="#">25498</a>	10634	1169	7	200	nucleophosmin 1 (Npm1) alternative variant cSep08, mRNA.
<a href="#">Npm3</a>	<a href="#">Npm3.aSep08</a>	<a href="#">502389</a>	524	394		71	nucleoplasmin 3 (Npm3) mRNA.
<a href="#">Npr1</a>	<a href="#">Npr1.bSep08</a>	<a href="#">24603</a>	3315	1259	1	297	natriuretic peptide receptor 1 (Npr1) alternative variant bSep08, mRNA.
<a href="#">Npr1</a>	<a href="#">Npr1.cSep08</a>	<a href="#">24603</a>	5044	694	2	212	natriuretic peptide receptor 1 (Npr1) alternative variant cSep08, mRNA.
<a href="#">Npr2</a>	<a href="#">Npr2.bSep08</a>	<a href="#">116564</a>	5405	717	7	238	natriuretic peptide receptor B (Npr2) alternative variant bSep08, mRNA.
<a href="#">Npr2</a>	<a href="#">Npr2.cSep08</a>	<a href="#">116564</a>	1664	1216	4	135	natriuretic peptide receptor 2 (14.8 kD) (Npr2) alternative variant cSep08, mRNA.
<a href="#">Npr2</a>	<a href="#">Npr2.dSep08</a>	<a href="#">116564</a>	983	761	2	89	natriuretic peptide receptor B (Npr2) alternative variant dSep08, mRNA.
<a href="#">Npr3</a>	<a href="#">Npr3.bSep08</a>	<a href="#">25339</a>	6073	1218	3	281	natriuretic peptide receptor 3 (Npr3) alternative variant bSep08, mRNA.
<a href="#">Npr3</a>	<a href="#">Npr3.cSep08</a>	<a href="#">25339</a>	47146	768	4	255	natriuretic peptide receptor 3 (Npr3) alternative variant cSep08, mRNA.
<a href="#">Nptn</a>	<a href="#">Nptn.bSep08</a>	<a href="#">56064</a>	75697	1470	8	281	neuroplastin (31.3 kD) (Nptn) alternative variant bSep08, mRNA.
<a href="#">Nptn</a>	<a href="#">Nptn.cSep08</a>	<a href="#">56064</a>	77301	3061	9	277	neuroplastin (30.8 kD) (Nptn) alternative variant cSep08, complete mRNA.
<a href="#">Nptn</a>	<a href="#">Nptn.eSep08</a>	<a href="#">56064</a>	8173	643	4	45	neuroplastin (Nptn) alternative variant eSep08, mRNA.
<a href="#">Nptx1</a>	<a href="#">Nptx1.aSep08</a>	<a href="#">266777</a>	4929	921		295	neuronal pentraxin 1 (Nptx1) mRNA.
<a href="#">Nptx2</a>	<a href="#">Nptx2.bSep08</a>	<a href="#">288475</a>	2729	1951	2	110	neuronal pentraxin 2 (12.2 kD) (Nptx2) alternative variant bSep08, mRNA.
<a href="#">Npw</a>	<a href="#">Npw.bSep08</a>	<a href="#">259224</a>	1249	397	2	107	neuropeptide W (Npw) alternative variant bSep08, mRNA.
<a href="#">Npy1r</a>	<a href="#">Npy1r.bSep08</a>	<a href="#">29358</a>	8302	752		140	neuropeptide Y receptor Y1 (16.4 kD) (Npy1r) alternative variant bSep08, mRNA.
<a href="#">Nqo2</a>	<a href="#">Nqo2.aSep08</a>	<a href="#">291084</a>	27882	830	2	235	NAD(P)H dehydrogenase, quinone 2 (Nqo2) alternative variant aSep08, mRNA.
<a href="#">Nqo2</a>	<a href="#">Nqo2.cSep08</a>	<a href="#">291084</a>	28301	1352	3	221	NAD(P)H dehydrogenase, quinone 2 (25.1 kD) (Nqo2) alternative variant cSep08, mRNA.
<a href="#">Nr1d1</a>	<a href="#">Nr1d1.bSep08</a>	<a href="#">252917</a>	4498	3366	4	613	nuclear receptor subfamily 1, group D, member 1 (67.7 kD) (Nr1d1) alternative variant bSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.bSep08</a>	<a href="#">58851</a>	4628	1364	9	360	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant bSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.cSep08</a>	<a href="#">58851</a>	2002	711	5	186	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant cSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.dSep08</a>	<a href="#">58851</a>	2126	745	6	159	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant dSep08, mRNA.

<a href="#">Nr1h2</a>	<a href="#">Nr1h2.fSep08</a>	<a href="#">58851</a>	1998	832	5	93	nuclear receptor subfamily 1, group H, member 2 (9.3 kD) (Nr1h2) alternative variant fSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.gSep08</a>	<a href="#">58851</a>	2040	754	6	88	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant gSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.hSep08</a>	<a href="#">58851</a>	1006	669	3	64	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant hSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.iSep08</a>	<a href="#">58851</a>	1802	775	4	38	nuclear receptor subfamily 1, group H, member 2 (Nr1h2) alternative variant iSep08, mRNA.
<a href="#">Nr1h2</a>	<a href="#">Nr1h2.jSep08</a>	<a href="#">58851</a>	3329	1765	6	283	nuclear receptor subfamily 1, group H, member 2 (31.0 kD) (Nr1h2) alternative variant jSep08, mRNA.
<a href="#">Nr1h3</a>	<a href="#">Nr1h3.bSep08</a>	<a href="#">58852</a>	3704	703	1	210	nuclear receptor subfamily 1, group H, member 3 (Nr1h3) alternative variant bSep08, mRNA.
<a href="#">Nr1h3</a>	<a href="#">Nr1h3.cSep08</a>	<a href="#">58852</a>	3724	1100	1	200	nuclear receptor subfamily 1, group H, member 3 (Nr1h3) alternative variant cSep08, mRNA.
<a href="#">Nr1h4</a>	<a href="#">Nr1h4.bSep08</a>	<a href="#">60351</a>	36909	1460	6	300	nuclear receptor subfamily 1, group H, member 4 (34.2 kD) (Nr1h4) alternative variant bSep08, complete mRNA.
<a href="#">Nr1h4</a>	<a href="#">Nr1h4.cSep08</a>	<a href="#">60351</a>	29958	757	4	247	nuclear receptor subfamily 1, group H, member 4 (Nr1h4) alternative variant cSep08, mRNA.
<a href="#">Nr1h4</a>	<a href="#">Nr1h4.dSep08</a>	<a href="#">60351</a>	4710	286	4	95	nuclear receptor subfamily 1, group H, member 4 (Nr1h4) alternative variant dSep08, mRNA.
<a href="#">Nr2c1</a>	<a href="#">Nr2c1.bSep08</a>	<a href="#">252924</a>	16767	930	6	180	nuclear receptor subfamily 2 group C member 1 (20.1 kD) (Nr2c1) alternative variant bSep08, mRNA.
<a href="#">Nr2c1</a>	<a href="#">Nr2c1.cSep08</a>	<a href="#">252924</a>	15769	1774	5	178	nuclear receptor subfamily 2 group C member 1 (18.9 kD) (Nr2c1) alternative variant cSep08, mRNA.
<a href="#">Nr2f2</a>	<a href="#">Nr2f2.bSep08</a>	<a href="#">113984</a>	13810	2720	3	281	nuclear receptor subfamily 2, group F, member 2 (31.5 kD) (Nr2f2) alternative variant bSep08, mRNA.
<a href="#">Nr2f6</a>	<a href="#">Nr2f6.dSep08</a>	<a href="#">245980</a>	2694	583	2	52	nuclear receptor subfamily 2, group F, member 6 and hypothetical protein LOC688751 (Nr2f6) alternative variant dSep08, mRNA.
<a href="#">Nr2f6</a>	<a href="#">Nr2f6.dSep08</a>	<a href="#">688751</a>	2694	583	2	52	nuclear receptor subfamily 2, group F, member 6 and hypothetical protein LOC688751 (Nr2f6) alternative variant dSep08, mRNA.
<a href="#">Nr3c2</a>	<a href="#">Nr3c2.bSep08</a>	<a href="#">25672</a>	47132	780	5	175	mineralocorticoid receptor (Nr3c2) alternative variant bSep08, mRNA.
<a href="#">Nr3c2</a>	<a href="#">Nr3c2.dSep08</a>	<a href="#">25672</a>	5854	337	2	76	putative protein (Nr3c2) alternative variant dSep08, mRNA.
<a href="#">Nr4a1</a>	<a href="#">Nr4a1.bSep08</a>	<a href="#">79240</a>	10008	580	1	163	nuclear receptor subfamily 4, group A, member 1 (Nr4a1) alternative variant bSep08, mRNA.
<a href="#">Nr5a2</a>	<a href="#">Nr5a2.bSep08</a>	<a href="#">60349</a>	4175	553	2	66	nuclear receptor subfamily 5, group A, member 2 (Nr5a2) alternative variant bSep08, mRNA.
<a href="#">Nr6a1</a>	<a href="#">Nr6a1.aSep08</a>	<a href="#">362125</a>	21523	974	5	324	nuclear receptor subfamily 6, group A, member 1 (Nr6a1) alternative variant aSep08, mRNA.
<a href="#">Nr6a1</a>	<a href="#">Nr6a1.bSep08</a>	<a href="#">362125</a>	13306	614	3	204	nuclear receptor subfamily 6, group A, member 1 (Nr6a1) alternative variant bSep08, mRNA.
<a href="#">Nradd</a>	<a href="#">Nradd.bSep08</a>	<a href="#">246143</a>	2161	1090	1	81	neurotrophin receptor associated death domain (Nradd) alternative variant bSep08, mRNA.
<a href="#">Nradd</a>	<a href="#">Nradd.cSep08</a>	<a href="#">246143</a>	2852	559	3	67	neurotrophin receptor associated death domain (Nradd) alternative variant cSep08, mRNA.

<a href="#">Nrap</a>	<a href="#">Nrap.cSep08</a>	<a href="#">307982</a>	7063	414	3	138	nebulin-related anchoring protein (Nrap) alternative variant cSep08, mRNA.
<a href="#">Nrcam</a>	<a href="#">Nrcam.bSep08</a>	<a href="#">497815</a>	245625	1530	11	321	neuron-glia-CAM-related cell adhesion molecule (Nrcam) alternative variant bSep08, mRNA.
<a href="#">Nrcam</a>	<a href="#">Nrcam.cSep08</a>	<a href="#">497815</a>	26298	1142	6	273	neuron-glia-CAM-related cell adhesion molecule (Nrcam) alternative variant cSep08, mRNA.
<a href="#">Nrcam</a>	<a href="#">Nrcam.dSep08</a>	<a href="#">497815</a>	25854	734	7	244	neuron-glia-CAM-related cell adhesion molecule (Nrcam) alternative variant dSep08, mRNA.
<a href="#">Nrcam</a>	<a href="#">Nrcam.eSep08</a>	<a href="#">497815</a>	15923	384	3	127	neuron-glia-CAM-related cell adhesion molecule (Nrcam) alternative variant eSep08, mRNA.
<a href="#">Nrcam</a>	<a href="#">Nrcam.fSep08</a>	<a href="#">497815</a>	81545	241	2	36	neuron-glia-CAM-related cell adhesion molecule (Nrcam) alternative variant fSep08, mRNA.
<a href="#">Nrd1</a>	<a href="#">Nrd1.bSep08</a>	<a href="#">25499</a>	6856	1352	5	202	nardilysin (Nrd1) alternative variant bSep08, mRNA.
<a href="#">Nrd1</a>	<a href="#">Nrd1.cSep08</a>	<a href="#">25499</a>	4050	356	4	106	nardilysin CRA a (Nrd1) alternative variant cSep08, mRNA.
<a href="#">Nrd1</a>	<a href="#">Nrd1.eSep08</a>	<a href="#">25499</a>	2491	663	3	56	nardilysin (Nrd1) alternative variant eSep08, mRNA.
<a href="#">Nrep</a>	<a href="#">Nrep.aSep08</a>	<a href="#">338475</a>	22686	358	3	114	neuronal regeneration related protein (Nrep) alternative variant aSep08, mRNA.
<a href="#">Nrep</a>	<a href="#">Nrep.cSep08</a>	<a href="#">338475</a>	4526	290	2	43	neuronal regeneration related protein (5.3 kD) (Nrep) alternative variant cSep08, mRNA.
<a href="#">Nrf1</a>	<a href="#">Nrf1.bSep08</a>	<a href="#">312195</a>	66225	3279	4	515	nuclear respiratory factor 1 (54.9 kD) (Nrf1) alternative variant bSep08, mRNA.
<a href="#">Nrf1</a>	<a href="#">Nrf1.cSep08</a>	<a href="#">312195</a>	25109	844	2	105	nuclear respiratory factor 1 (10.4 kD) (Nrf1) alternative variant cSep08, mRNA.
<a href="#">Nrg1</a>	<a href="#">Nrg1.bSep08</a>	<a href="#">112400</a>	19576	728	3	242	neuregulin 1 (Nrg1) alternative variant bSep08, mRNA.
<a href="#">Nrg1</a>	<a href="#">Nrg1.cSep08</a>	<a href="#">112400</a>	27792	554	4	184	neuregulin 1 (Nrg1) alternative variant cSep08, mRNA.
<a href="#">Nrg2</a>	<a href="#">Nrg2.aSep08</a>	<a href="#">432361</a>	6322	1770		590	neuregulin 2 (Nrg2) alternative variant aSep08, mRNA.
<a href="#">Nrg2</a>	<a href="#">Nrg2.bSep08</a>	<a href="#">432361</a>	3064	434		144	neuregulin 2 (Nrg2) alternative variant bSep08, mRNA.
<a href="#">Nrg3</a>	<a href="#">Nrg3.aSep08</a>	<a href="#">498596</a>	27344	1913		311	neuregulin 3 (Nrg3) mRNA.
<a href="#">Nrg4</a>	<a href="#">Nrg4.aSep08</a>	<a href="#">690919</a>	36328	800		70	neuregulin 4 (Nrg4) mRNA.
<a href="#">Nrk</a>	<a href="#">Nrk.aSep08</a>	<a href="#">315907</a>	8748	796		264	nik related kinase (Nrk) mRNA.
<a href="#">Nrm</a>	<a href="#">Nrm.bSep08</a>	<a href="#">361791</a>	2150	741	2	189	nurim (nuclear envelope membrane protein) (Nrm) alternative variant bSep08, mRNA.
<a href="#">Nrm</a>	<a href="#">Nrm.cSep08</a>	<a href="#">361791</a>	2589	737	3	188	nurim (nuclear envelope membrane protein) (Nrm) alternative variant cSep08, mRNA.
<a href="#">Nrm</a>	<a href="#">Nrm.dSep08</a>	<a href="#">361791</a>	2630	410	2	136	nurim (nuclear envelope membrane protein) (Nrm) alternative variant dSep08, mRNA.
<a href="#">Nrp1</a>	<a href="#">Nrp1.bSep08</a>	<a href="#">246331</a>	7507	3050	1	154	neuropilin 1 (Nrp1) alternative variant bSep08, mRNA.
<a href="#">Nrp2</a>	<a href="#">Nrp2.bSep08</a>	<a href="#">81527</a>	10638	414	3	116	neuropilin 2 (Nrp2) alternative variant bSep08, mRNA.
<a href="#">Nrsn1</a>	<a href="#">Nrsn1.aSep08</a>	<a href="#">291129</a>	17808	2096	2	196	neurensin 1 (21.7 kD) (Nrsn1) alternative variant aSep08, mRNA.
<a href="#">Nrxn1</a>	<a href="#">Nrxn1.aSep08</a>	<a href="#">60391</a>	85429	960		319	neurexin 1 (Nrxn1) mRNA.
<a href="#">Nrxn2</a>	<a href="#">Nrxn2.bSep08</a>	<a href="#">116595</a>	44661	1449	7	395	neurexin II-alpha-b (Nrxn2) alternative variant bSep08, mRNA.
<a href="#">Nrxn2</a>	<a href="#">Nrxn2.cSep08</a>	<a href="#">116595</a>	19480	753	4	250	neurexin 2 (Nrxn2) alternative variant cSep08, mRNA.



<a href="#">Nrnx2</a>	<a href="#">Nrnx2.dSep08</a>	<a href="#">116595</a>	1311	791	2	207	neurexin II-alpha-b (Nrnx2) alternative variant dSep08, mRNA.
<a href="#">Nrnx3</a>	<a href="#">Nrnx3.cSep08</a>	<a href="#">116508</a>	26962	727	2	33	neurexin 3 (4.0 kD) (Nrnx3) alternative variant cSep08, mRNA.
<a href="#">Ns5atp9</a>	<a href="#">Ns5atp9.bSep08</a>	<a href="#">300795</a>	12633	395	2	115	NS5A (hepatitis C virus) transactivated protein 9 (Ns5atp9) alternative variant bSep08, mRNA.
<a href="#">Nsdhl</a>	<a href="#">Nsdhl.aSep08</a>	<a href="#">309262</a>	23355	2941	6	388	NAD(P) dependent steroid dehydrogenase-like (43.5 kD) (Nsdhl) alternative variant aSep08, mRNA.
<a href="#">Nsf</a>	<a href="#">Nsf.bSep08</a>	<a href="#">60355</a>	9586	308	2	57	N-ethylmaleimide sensitive fusion protein (Nsf) alternative variant bSep08, mRNA.
<a href="#">Nsf1c</a>	<a href="#">Nsf1c.bSep08</a>	<a href="#">83809</a>	13627	997	6	287	NSFL1 (p97) cofactor (p47) (Nsf1c) alternative variant bSep08, mRNA.
<a href="#">Nsf1c</a>	<a href="#">Nsf1c.cSep08</a>	<a href="#">83809</a>	11948	643	3	127	NSFL1 (p97) cofactor (p47) (Nsf1c) alternative variant cSep08, mRNA.
<a href="#">Nsmce1</a>	<a href="#">Nsmce1.bSep08</a>	<a href="#">361645</a>	24859	644	5	200	non-SMC element 1 homolog (S. cerevisiae) (Nsmce1) alternative variant bSep08, mRNA.
<a href="#">Nsmce1</a>	<a href="#">Nsmce1.cSep08</a>	<a href="#">361645</a>	12553	711	5	142	non-SMC element 1 homolog (S. cerevisiae) (Nsmce1) alternative variant cSep08, mRNA.
<a href="#">Nsmce1</a>	<a href="#">Nsmce1.dSep08</a>	<a href="#">361645</a>	19957	782	2	86	non-SMC element 1 homolog (S. cerevisiae) (Nsmce1) alternative variant dSep08, mRNA.
<a href="#">Nsmce2</a>	<a href="#">Nsmce2.bSep08</a>	<a href="#">299957</a>	225905	783	5	158	non-SMC element 2, MMS21 homolog (S. cerevisiae) (Nsmce2) alternative variant bSep08, mRNA.
<a href="#">Nsmce2</a>	<a href="#">Nsmce2.dSep08</a>	<a href="#">299957</a>	9421	483	3	23	non-SMC element 2, MMS21 homolog (S. cerevisiae) (Nsmce2) alternative variant dSep08, mRNA.
<a href="#">Nsmce4a</a>	<a href="#">Nsmce4a.aSep08</a>	<a href="#">293528</a>	14737	1363	10	393	non-smc element 4 homolog a (Nsmce4a) alternative variant aSep08, mRNA.
<a href="#">Nsmce4a</a>	<a href="#">Nsmce4a.cSep08</a>	<a href="#">293528</a>	5821	1471	6	151	non-SMC element 4 homolog a (Nsmce4a) alternative variant cSep08, mRNA.
<a href="#">Nsmce4a</a>	<a href="#">Nsmce4a.dSep08</a>	<a href="#">293528</a>	5891	706	7	144	non-SMC element 4 homolog a (16.8 kD) (Nsmce4a) alternative variant dSep08, mRNA.
<a href="#">Nsun2</a>	<a href="#">Nsun2.bSep08</a>	<a href="#">361191</a>	2142	306	2	87	NOL1/NOP2/Sun domain family, member 2 (Nsun2) alternative variant bSep08, mRNA.
<a href="#">Nsun2</a>	<a href="#">Nsun2.cSep08</a>	<a href="#">361191</a>	2260	742	3	63	NOL1/NOP2/Sun domain family, member 2 (Nsun2) alternative variant cSep08, mRNA.
<a href="#">Nsun4</a>	<a href="#">Nsun4.bSep08</a>	<a href="#">298426</a>	7481	655	1	74	NOL1/NOP2/Sun domain family, member 4 (Nsun4) alternative variant bSep08, mRNA.
<a href="#">Nsun5</a>	<a href="#">Nsun5.aSep08</a>	<a href="#">288595</a>	5615	2196	8	450	NOL1/NOP2/Sun domain family, member 5 (Nsun5) alternative variant aSep08, mRNA.
<a href="#">Nsun5</a>	<a href="#">Nsun5.bSep08</a>	<a href="#">288595</a>	1148	714	1	148	NOL1/NOP2/Sun domain family, member 5 (16.4 kD) (Nsun5) alternative variant bSep08, mRNA.
<a href="#">Nsun6</a>	<a href="#">Nsun6.bSep08</a>	<a href="#">307148</a>	20600	499	6	166	NOL1/NOP2/Sun domain family, member 6 (Nsun6) alternative variant bSep08, mRNA.
<a href="#">Nt5c3</a>	<a href="#">Nt5c3.aSep08</a>	<a href="#">312373</a>	42616	1243	9	331	5'-nucleotidase, cytosolic III (37.3 kD) (Nt5c3) alternative variant aSep08, mRNA.
<a href="#">Nt5c3l</a>	<a href="#">Nt5c3l.bSep08</a>	<a href="#">360629</a>	12894	920	7	266	5'-nucleotidase, cytosolic III-like (Nt5c3l) alternative variant bSep08, mRNA.

<a href="#">Nt5c3l</a>	<a href="#">Nt5c3l.cSep08</a>	<a href="#">360629</a>	11258	731	6	243	5'-nucleotidase, cytosolic III-like (Nt5c3l) alternative variant cSep08, mRNA.
<a href="#">Nt5c3l</a>	<a href="#">Nt5c3l.dSep08</a>	<a href="#">360629</a>	9623	816	7	189	5'-nucleotidase, cytosolic III-like (Nt5c3l) alternative variant dSep08, mRNA.
<a href="#">Nt5c3l</a>	<a href="#">Nt5c3l.eSep08</a>	<a href="#">360629</a>	8479	939	3	156	5'-nucleotidase, cytosolic III-like (Nt5c3l) alternative variant eSep08, mRNA.
<a href="#">Nt5c3l</a>	<a href="#">Nt5c3l.fSep08</a>	<a href="#">360629</a>	8694	416	5	88	5'-nucleotidase, cytosolic III-like (Nt5c3l) alternative variant fSep08, mRNA.
<a href="#">Nt5dc1</a>	<a href="#">Nt5dc1.bSep08</a>	<a href="#">294456</a>	44452	739	3	246	putative protein of ancient origin (Nt5dc1) alternative variant bSep08, mRNA.
<a href="#">Nt5dc1</a>	<a href="#">Nt5dc1.cSep08</a>	<a href="#">294456</a>	18025	616	1	205	putative protein of ancient origin (Nt5dc1) alternative variant cSep08, mRNA.
<a href="#">Nt5dc1</a>	<a href="#">Nt5dc1.dSep08</a>	<a href="#">294456</a>	44564	689	1	179	putative protein of ancient origin (Nt5dc1) alternative variant dSep08, mRNA.
<a href="#">Nt5dc2</a>	<a href="#">Nt5dc2.cSep08</a>	<a href="#">290558</a>	4267	392	3	90	putative protein of eukaryotic origin (Nt5dc2) alternative variant cSep08, mRNA.
<a href="#">Nt5m</a>	<a href="#">Nt5m.aSep08</a>	<a href="#">287368</a>	27576	2254	5	220	5',3'-nucleotidase, mitochondrial (25.5 kD) (Nt5m) alternative variant aSep08, complete mRNA.
<a href="#">Nt5m</a>	<a href="#">Nt5m.cSep08</a>	<a href="#">287368</a>	2210	932	3	152	5',3'-nucleotidase, mitochondrial (15.9 kD) (Nt5m) alternative variant cSep08, mRNA.
<a href="#">Nt5m</a>	<a href="#">Nt5m.dSep08</a>	<a href="#">287368</a>	37808	669	5	54	5',3'-nucleotidase, mitochondrial (Nt5m) alternative variant dSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.bSep08</a>	<a href="#">360462</a>	14993	853	7	238	N-terminal asparagine amidase (Ntan1) alternative variant bSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.cSep08</a>	<a href="#">360462</a>	6270	917	6	205	N-terminal asparagine amidase (23.5 kD) (Ntan1) alternative variant cSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.dSep08</a>	<a href="#">360462</a>	6249	682	6	181	N-terminal asparagine amidase (Ntan1) alternative variant dSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.eSep08</a>	<a href="#">360462</a>	7832	516	5	132	N-terminal asparagine amidase (14.9 kD) (Ntan1) alternative variant eSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.gSep08</a>	<a href="#">360462</a>	1193	411	2	102	N-terminal asparagine amidase (Ntan1) alternative variant gSep08, mRNA.
<a href="#">Ntan1</a>	<a href="#">Ntan1.hSep08</a>	<a href="#">360462</a>	923	582	2	96	N-terminal asparagine amidase (11.1 kD) (Ntan1) alternative variant hSep08, mRNA.
<a href="#">Ntf3</a>	<a href="#">Ntf3.bSep08</a>	<a href="#">81737</a>	70698	820	2	203	neurotrophin 3 (Ntf3) alternative variant bSep08, mRNA.
<a href="#">Nthl1</a>	<a href="#">Nthl1.bSep08</a>	<a href="#">29541</a>	6481	935	6	272	nth (endonuclease III)-like 1 (E.coli) (Nthl1) alternative variant bSep08, mRNA.
<a href="#">Nthl1</a>	<a href="#">Nthl1.cSep08</a>	<a href="#">29541</a>	817	741	2	208	nth (endonuclease III)-like 1 (E.coli) (Nthl1) alternative variant cSep08, mRNA.
<a href="#">Nthl1</a>	<a href="#">Nthl1.dSep08</a>	<a href="#">29541</a>	5103	697	5	193	nth (endonuclease III)-like 1 (E.coli) (Nthl1) alternative variant dSep08, mRNA.
<a href="#">Nthl1</a>	<a href="#">Nthl1.eSep08</a>	<a href="#">29541</a>	5104	713	5	157	nth (endonuclease III)-like 1 (E.coli) (Nthl1) alternative variant eSep08, mRNA.
<a href="#">Ntn2l</a>	<a href="#">Ntn2l.aSep08</a>	<a href="#">114524</a>	2037	1368	5	455	netrin 2-like (chicken) (Ntn2l) alternative variant aSep08, mRNA.
<a href="#">Ntn2l</a>	<a href="#">Ntn2l.bSep08</a>	<a href="#">114524</a>	943	656	3	218	netrin 2-like (chicken) (Ntn2l) alternative variant bSep08, mRNA.

<a href="#">Ntn2l</a>	<a href="#">Ntn2l.cSep08</a>	<a href="#">114524</a>	885	793	1	75	netrin 2-like (chicken) (Ntn2l) alternative variant cSep08, mRNA.
<a href="#">Ntng1</a>	<a href="#">Ntng1.bSep08</a>	<a href="#">295382</a>	153972	568	4	189	netrin G1 (Ntng1) alternative variant bSep08, mRNA.
<a href="#">Ntrk2</a>	<a href="#">Ntrk2.bSep08</a>	<a href="#">25054</a>	78882	5745	6	208	tyrosine kinase receptor (Ntrk2) alternative variant bSep08, mRNA.
<a href="#">Ntrk2</a>	<a href="#">Ntrk2.cSep08</a>	<a href="#">25054</a>	9816	334	4	83	tyrosine kinase receptor (Ntrk2) alternative variant cSep08, mRNA.
<a href="#">Ntrk2</a>	<a href="#">Ntrk2.dSep08</a>	<a href="#">25054</a>	2150	404	2	66	putative protein (Ntrk2) alternative variant dSep08, mRNA.
<a href="#">Ntrk3</a>	<a href="#">Ntrk3.bSep08</a>	<a href="#">29613</a>	53082	783	2	195	neurotrophic tyrosine kinase, receptor, type 3 (Ntrk3) alternative variant bSep08, mRNA.
<a href="#">Ntrk3</a>	<a href="#">Ntrk3.cSep08</a>	<a href="#">29613</a>	52997	623	1	147	neurotrophic tyrosine kinase, receptor, type 3 (Ntrk3) alternative variant cSep08, mRNA.
<a href="#">Nub1</a>	<a href="#">Nub1.bSep08</a>	<a href="#">296731</a>	18432	1174	9	300	negative regulator of ubiquitin-like proteins 1 (Nub1) alternative variant bSep08, mRNA.
<a href="#">Nubp1</a>	<a href="#">Nubp1.aSep08</a>	<a href="#">287042</a>	11582	2308	10	334	nucleotide binding protein 1 like (35.8 kD) (Nubp1) alternative variant aSep08, complete mRNA.
<a href="#">Nubp1</a>	<a href="#">Nubp1.bSep08</a>	<a href="#">287042</a>	10056	961	10	292	nucleotide binding protein 1 like (Nubp1) alternative variant bSep08, mRNA.
<a href="#">Nubp1</a>	<a href="#">Nubp1.cSep08</a>	<a href="#">287042</a>	8343	736	8	164	nucleotide binding protein 1 like (16.8 kD) (Nubp1) alternative variant cSep08, mRNA.
<a href="#">Nubp1</a>	<a href="#">Nubp1.dSep08</a>	<a href="#">287042</a>	3328	2171	2	59	nucleotide binding protein 1 like (6.5 kD) (Nubp1) alternative variant dSep08, mRNA.
<a href="#">Nubp2</a>	<a href="#">Nubp2.bSep08</a>	<a href="#">287125</a>	2547	687	3	163	nucleotide binding protein 2 (Nubp2) alternative variant bSep08, mRNA.
<a href="#">Nubp2</a>	<a href="#">Nubp2.cSep08</a>	<a href="#">287125</a>	3035	475	2	151	nucleotide binding protein 2 (16.0 kD) (Nubp2) alternative variant cSep08, mRNA.
<a href="#">Nubpl</a>	<a href="#">Nubpl.aSep08</a>	<a href="#">299008</a>	211770	1163	3	319	nucleotide binding protein-like (34.1 kD) (Nubpl) alternative variant aSep08, mRNA.
<a href="#">Nubpl</a>	<a href="#">Nubpl.bSep08</a>	<a href="#">299008</a>	14488	254	1	63	nucleotide binding protein-like (Nubpl) alternative variant bSep08, mRNA.
<a href="#">nuby</a>	<a href="#">nuby.aSep08</a>		9217	792		68	putative protein (8.0 kD) (nuby) mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.bSep08</a>	<a href="#">84595</a>	14844	1415	7	264	nucleobindin 1 CRA a precursor (30.9 kD) (Nucb1) alternative variant bSep08, complete mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.cSep08</a>	<a href="#">84595</a>	14275	843	7	251	nucleobindin 1 CRA a (Nucb1) alternative variant cSep08, mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.dSep08</a>	<a href="#">84595</a>	1851	767	6	175	nucleobindin 1 CRA a (20.2 kD) (Nucb1) alternative variant dSep08, mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.eSep08</a>	<a href="#">84595</a>	1645	1153	4	131	nucleobindin 1 CRA a (Nucb1) alternative variant eSep08, mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.fSep08</a>	<a href="#">84595</a>	10698	697	4	128	nucleobindin 1 CRA c precursor (14.4 kD) (Nucb1) alternative variant fSep08, mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.gSep08</a>	<a href="#">84595</a>	3737	527	3	121	nucleobindin 1 CRA a (14.8 kD) (Nucb1) alternative variant gSep08, mRNA.
<a href="#">Nucb1</a>	<a href="#">Nucb1.iSep08</a>	<a href="#">84595</a>	9034	311	3	65	putative protein (Nucb1) alternative variant iSep08, mRNA.
<a href="#">Nucb2</a>	<a href="#">Nucb2.bSep08</a>	<a href="#">59295</a>	24232	794	6	264	nucleobindin 2 (Nucb2) alternative variant bSep08, mRNA.
<a href="#">nuchy</a>	<a href="#">nuchy.aSep08</a>		2114	437		57	putative protein (nuchy) mRNA.

<a href="#">Nucleoporin2.0</a>	<a href="#">Nucleoporin2.0.aSep08</a>		26405	1183	6	394	nucleoporin 98kDa (Nucleoporin2.0) alternative variant aSep08, mRNA.
<a href="#">Nucleoporin2.0</a>	<a href="#">Nucleoporin2.0.bSep08</a>		9356	415	1	138	nucleoporin 98kDa (Nucleoporin2.0) alternative variant bSep08, mRNA.
<a href="#">Nuc_sug_transp.0</a>	<a href="#">Nuc_sug_transp.0.aSep08</a>		6260	435		145	solute carrier family 35 member A5 (Nuc_sug_transp.0) mRNA.
<a href="#">nudar</a>	<a href="#">nudar.aSep08</a>		2659	631		79	putative protein (9.1 kD) (nudar) mRNA.
<a href="#">Nudcd2</a>	<a href="#">Nudcd2.bSep08</a>	<a href="#">287199</a>	4858	690	2	53	putative protein (Nudcd2) alternative variant bSep08, mRNA.
<a href="#">Nudt1</a>	<a href="#">Nudt1.bSep08</a>	<a href="#">117260</a>	6787	508	2	114	nudix (nucleoside diphosphate linked moiety X)-type motif 1 (Nudt1) alternative variant bSep08, mRNA.
<a href="#">Nudt3</a>	<a href="#">Nudt3.bSep08</a>	<a href="#">294292</a>	51322	670	1	131	nudix (nucleotide diphosphate linked moiety X)-type motif 3 (15.1 kD) (Nudt3) alternative variant bSep08, mRNA.
<a href="#">Nudt4</a>	<a href="#">Nudt4.aSep08</a>	<a href="#">94267</a>	14197	900	5	234	nudix (nucleoside diphosphate linked moiety X)-type motif 4 (Nudt4) alternative variant aSep08, mRNA.
<a href="#">Nudt4</a>	<a href="#">Nudt4.cSep08</a>	<a href="#">94267</a>	2571	640	4	151	nudix (nucleoside diphosphate linked moiety X)-type motif 4 (Nudt4) alternative variant cSep08, mRNA.
<a href="#">Nudt4</a>	<a href="#">Nudt4.dSep08</a>	<a href="#">94267</a>	1205	963	2	116	nudix (nucleoside diphosphate linked moiety X)-type motif 4 (13.2 kD) (Nudt4) alternative variant dSep08, complete mRNA.
<a href="#">Nudt4</a>	<a href="#">Nudt4.eSep08</a>	<a href="#">94267</a>	1814	260	3	29	nudix (nucleoside diphosphate linked moiety X)-type motif 4 (Nudt4) alternative variant eSep08, mRNA.
<a href="#">Nudt7</a>	<a href="#">Nudt7.bSep08</a>	<a href="#">361413</a>	26563	431	4	143	nudix (nucleoside diphosphate linked moiety X)-type motif 7 (Nudt7) alternative variant bSep08, mRNA.
<a href="#">Nudt7</a>	<a href="#">Nudt7.cSep08</a>	<a href="#">361413</a>	15077	1043	5	75	nudix (nucleoside diphosphate linked moiety X)-type motif 7 (Nudt7) alternative variant cSep08, mRNA.
<a href="#">Nudt8</a>	<a href="#">Nudt8.aSep08</a>	<a href="#">361692</a>	1712	989		301	nudix (nucleoside diphosphate linked moiety X)-type motif 8 (Nudt8) mRNA.
<a href="#">Nudt9</a>	<a href="#">Nudt9.bSep08</a>	<a href="#">305149</a>	31311	862	8	287	nudix (nucleoside diphosphate linked moiety X)-type motif 9 (Nudt9) alternative variant bSep08, mRNA.
<a href="#">Nudt9</a>	<a href="#">Nudt9.cSep08</a>	<a href="#">305149</a>	23604	619	6	172	nudix (nucleoside diphosphate linked moiety X)-type motif 9 (Nudt9) alternative variant cSep08, mRNA.
<a href="#">Nudt11</a>	<a href="#">Nudt11.aSep08</a>	<a href="#">680248</a>	3565	915		200	nudix (nucleoside diphosphate linked moiety X)-type motif 11 (Nudt11) mRNA.
<a href="#">Nudt15</a>	<a href="#">Nudt15.aSep08</a>	<a href="#">290365</a>	4149	642	1	170	nudix (nucleoside diphosphate linked moiety X)-type motif 15 (19.7 kD) (Nudt15) alternative variant aSep08, mRNA.
<a href="#">Nudt15</a>	<a href="#">Nudt15.bSep08</a>	<a href="#">290365</a>	3674	471	1	137	nudix (nucleoside diphosphate linked moiety X)-type motif 15 (Nudt15) alternative variant bSep08, mRNA.
<a href="#">Nudt16</a>	<a href="#">Nudt16.bSep08</a>	<a href="#">363129</a>	2355	1641	2	171	nudix (nucleoside diphosphate linked moiety X)-type motif 16 (18.8 kD) (Nudt16) alternative variant bSep08, complete mRNA.
<a href="#">Nudt16l1</a>	<a href="#">Nudt16l1.bSep08</a>	<a href="#">497867</a>	979	898	1	151	nudix (nucleoside diphosphate linked moiety X)-type motif 16-like 1 (Nudt16l1) alternative variant bSep08, mRNA.
<a href="#">Nudt18</a>	<a href="#">Nudt18.bSep08</a>	<a href="#">361068</a>	2521	478	2	65	nudix (nucleoside diphosphate linked moiety X)-type motif 18 (6.9 kD) (Nudt18) alternative variant bSep08, mRNA.

<a href="#">Nudt19</a>	<a href="#">Nudt19.aSep08</a>	<a href="#">308518</a>	11691	941	1	201	nudix (nucleoside diphosphate linked moiety X)-type motif 19 (22.6 kD) (Nudt19) alternative variant aSep08, complete mRNA.
<a href="#">Nudt22</a>	<a href="#">Nudt22.bSep08</a>	<a href="#">293703</a>	946	682	2	226	nudix (nucleoside diphosphate linked moiety X)-type motif 22 (Nudt22) alternative variant bSep08, mRNA.
<a href="#">Nudt22</a>	<a href="#">Nudt22.cSep08</a>	<a href="#">293703</a>	2340	652	4	156	nudix (nucleoside diphosphate linked moiety X)-type motif 22 (17.0 kD) (Nudt22) alternative variant cSep08, mRNA.
<a href="#">Nudt22</a>	<a href="#">Nudt22.dSep08</a>	<a href="#">293703</a>	927	569	3	114	nudix (nucleoside diphosphate linked moiety X)-type motif 22 (12.4 kD) (Nudt22) alternative variant dSep08, mRNA.
<a href="#">Nudt22</a>	<a href="#">Nudt22.eSep08</a>	<a href="#">293703</a>	2687	350	3	107	nudix (nucleoside diphosphate linked moiety X)-type motif 22 (Nudt22) alternative variant eSep08, mRNA.
<a href="#">Nuf2</a>	<a href="#">Nuf2.bSep08</a>	<a href="#">304951</a>	5359	814	1	139	NUF2, NDC80 kinetochore complex component, homolog ( <i>S. cerevisiae</i> ) (Nuf2) alternative variant bSep08, mRNA.
<a href="#">nufer</a>	<a href="#">nufer.aSep08</a>		445	396		71	putative protein (nufer) mRNA.
<a href="#">Nufip1</a>	<a href="#">Nufip1.bSep08</a>	<a href="#">364430</a>	8278	1228	3	129	nuclear fragile X mental retardation protein interacting protein 1 (14.6 kD) (Nufip1) alternative variant bSep08, mRNA.
<a href="#">Nufip1</a>	<a href="#">Nufip1.cSep08</a>	<a href="#">364430</a>	10790	849	3	117	nuclear fragile X mental retardation protein interacting protein 1 (Nufip1) alternative variant cSep08, mRNA.
<a href="#">nuflo</a>	<a href="#">nuflo.aSep08</a>		1095	723		78	putative protein (nuflo) mRNA.
<a href="#">nuflu</a>	<a href="#">nuflu.aSep08</a>		3364	198		15	putative protein (nuflu) mRNA.
<a href="#">nugar</a>	<a href="#">nugar.aSep08</a>		2958	634		211	putative protein of vertebrate origin (nugar) mRNA.
<a href="#">nuja</a>	<a href="#">nuja.aSep08</a>		4531	642		89	putative secreted or extracellular protein precursor (10.0 kD) (nuja) mRNA.
<a href="#">nujey</a>	<a href="#">nujey.aSep08</a>		878	412		46	putative protein (5.4 kD) (nujey) mRNA.
<a href="#">nukee</a>	<a href="#">nukee.aSep08</a>		35358	374		124	RalGDS-like (nukee) alternative variant aSep08, mRNA.
<a href="#">nuloy</a>	<a href="#">nuloy.aSep08</a>		459	295		93	putative protein (nuloy) mRNA.
<a href="#">Numa1</a>	<a href="#">Numa1.aSep08</a>	<a href="#">308870</a>	19390	3765	4	1014	nuclear mitotic apparatus protein 1 (Numa1) alternative variant aSep08, mRNA.
<a href="#">Numa1</a>	<a href="#">Numa1.bSep08</a>	<a href="#">308870</a>	2517	1267	3	266	nuclear mitotic apparatus protein 1 (Numa1) alternative variant bSep08, mRNA.
<a href="#">Numb</a>	<a href="#">Numb.aSep08</a>	<a href="#">29419</a>	9161	964	4	207	numb gene homolog ( <i>Drosophila</i> ) (Numb) alternative variant aSep08, mRNA.
<a href="#">Numb</a>	<a href="#">Numb.bSep08</a>	<a href="#">29419</a>	111574	736	8	145	numb gene homolog ( <i>Drosophila</i> ) (Numb) alternative variant bSep08, mRNA.
<a href="#">Numb</a>	<a href="#">Numb.cSep08</a>	<a href="#">29419</a>	69310	389	4	82	numb gene homolog ( <i>Drosophila</i> ) (Numb) alternative variant cSep08, mRNA.
<a href="#">Numbl</a>	<a href="#">Numbl.cSep08</a>	<a href="#">292732</a>	3101	431	2	78	numb-like (Numbl) alternative variant cSep08, mRNA.
<a href="#">numee</a>	<a href="#">numee.aSep08</a>		4427	362		86	putative protein (numee) mRNA.
<a href="#">numer</a>	<a href="#">numer.aSep08</a>		5172	434		48	putative protein (5.4 kD) (numer) mRNA.
<a href="#">nunoy</a>	<a href="#">nunoy.aSep08</a>		2914	2104		94	putative protein of mammalian origin (nunoy) mRNA.
<a href="#">Nup37</a>	<a href="#">Nup37.bSep08</a>	<a href="#">299706</a>	28690	2754	2	293	nucleoporin 37 (33.3 kD) (Nup37) alternative variant bSep08, mRNA.
<a href="#">Nup54</a>	<a href="#">Nup54.bSep08</a>	<a href="#">53372</a>	6473	609	5	203	nucleoporin 54 (Nup54) alternative variant bSep08, mRNA.

<a href="#">Nup54</a>	<a href="#">Nup54.cSep08</a>	<a href="#">53372</a>	3608	895	4	132	nucleoporin 54 (15.2 kD) (Nup54) alternative variant cSep08, mRNA.
<a href="#">Nup54</a>	<a href="#">Nup54.dSep08</a>	<a href="#">53372</a>	2052	719	2	73	nucleoporin 54 (8.5 kD) (Nup54) alternative variant dSep08, mRNA.
<a href="#">Nup62</a>	<a href="#">Nup62.aSep08</a>	<a href="#">65274</a>	16330	2758	2	525	nuclear pore complex glycoprotein p62 (53.4 kD) (Nup62) alternative variant aSep08, mRNA.
<a href="#">Nup62</a>	<a href="#">Nup62.cSep08</a>	<a href="#">65274</a>	22495	785	6	178	interleukin 4 induced 1 like (Nup62) alternative variant cSep08, mRNA.
<a href="#">Nup62cl</a>	<a href="#">Nup62cl.aSep08</a>	<a href="#">300923</a>	60535	2033	9	350	nucleoporin 62 C-terminal like (Nup62cl) alternative variant aSep08, mRNA.
<a href="#">Nup62cl</a>	<a href="#">Nup62cl.bSep08</a>	<a href="#">300923</a>	40434	855	8	225	nucleoporin 62 C-terminal like (Nup62cl) alternative variant bSep08, mRNA.
<a href="#">Nup85</a>	<a href="#">Nup85.bSep08</a>	<a href="#">287830</a>	12625	1433	13	201	nucleoporin 85kDa (Nup85) alternative variant bSep08, mRNA.
<a href="#">Nup85</a>	<a href="#">Nup85.fSep08</a>	<a href="#">287830</a>	3130	1795	3	56	nucleoporin 85kDa (6.3 kD) (Nup85) alternative variant fSep08, mRNA.
<a href="#">Nup88</a>	<a href="#">Nup88.bSep08</a>	<a href="#">113929</a>	7677	2082	2	202	nucleoporin 88 (22.9 kD) (Nup88) alternative variant bSep08, mRNA.
<a href="#">Nup93</a>	<a href="#">Nup93.bSep08</a>	<a href="#">291874</a>	23229	2098	17	645	nucleoporin 93 (Nup93) alternative variant bSep08, mRNA.
<a href="#">Nup93</a>	<a href="#">Nup93.cSep08</a>	<a href="#">291874</a>	78118	696	5	206	nucleoporin 93 (Nup93) alternative variant cSep08, mRNA.
<a href="#">Nup107</a>	<a href="#">Nup107.bSep08</a>	<a href="#">116555</a>	8229	563	1	187	nucleoporin 107 (Nup107) alternative variant bSep08, mRNA.
<a href="#">Nup133.0</a>	<a href="#">Nup133.0.aSep08</a>		48064	3290	25	1066	nucleoporin 133kDa (Nup133.0) alternative variant aSep08, mRNA.
<a href="#">Nup133.0</a>	<a href="#">Nup133.0.bSep08</a>		8780	727	5	242	nucleoporin 133kDa (Nup133.0) alternative variant bSep08, mRNA.
<a href="#">Nup133.0</a>	<a href="#">Nup133.0.cSep08</a>		7383	707	5	169	nucleoporin 133kDa (Nup133.0) alternative variant cSep08, mRNA.
<a href="#">Nup133.0</a>	<a href="#">Nup133.0.dSep08</a>		5300	558	3	159	CRA b like (Nup133.0) alternative variant dSep08, mRNA.
<a href="#">Nup153</a>	<a href="#">Nup153.bSep08</a>	<a href="#">25281</a>	4246	732	3	243	nucleoporin 153 (Nup153) alternative variant bSep08, mRNA.
<a href="#">Nup153</a>	<a href="#">Nup153.cSep08</a>	<a href="#">25281</a>	1010	727	2	96	nucleoporin 153 (Nup153) alternative variant cSep08, mRNA.
<a href="#">Nup155</a>	<a href="#">Nup155.bSep08</a>	<a href="#">117021</a>	14161	1792	3	461	nucleoporin 155 (Nup155) alternative variant bSep08, mRNA.
<a href="#">Nup155</a>	<a href="#">Nup155.cSep08</a>	<a href="#">117021</a>	5632	829	3	226	nucleoporin 155 (25.9 kD) (Nup155) alternative variant cSep08, mRNA.
<a href="#">Nup160</a>	<a href="#">Nup160.bSep08</a>	<a href="#">311182</a>	23439	1698	11	565	nucleoporin 160 (Nup160) alternative variant bSep08, mRNA.
<a href="#">Nup160</a>	<a href="#">Nup160.cSep08</a>	<a href="#">311182</a>	7736	1006	1	118	nucleoporin 160 (Nup160) alternative variant cSep08, mRNA.
<a href="#">Nup160</a>	<a href="#">Nup160.dSep08</a>	<a href="#">311182</a>	11919	1776	4		
<a href="#">Nup188</a>	<a href="#">Nup188.aSep08</a>	<a href="#">366016</a>	11306	3088	16	709	nucleoporin 188 (Nup188) alternative variant aSep08, mRNA.
<a href="#">Nup205</a>	<a href="#">Nup205.bSep08</a>	<a href="#">362335</a>	4962	752	2	206	nucleoporin 205 (Nup205) alternative variant bSep08, mRNA.

<a href="#">Nup214</a>	<a href="#">Nup214.aSep08</a>	<a href="#">296634</a>	9807	1798		599	nucleoporin 214 (Nup214) mRNA.
<a href="#">Nupl1</a>	<a href="#">Nupl1.bSep08</a>	<a href="#">245922</a>	10115	580	5	160	nucleoporin like 1 (Nupl1) alternative variant bSep08, mRNA.
<a href="#">Nupl1</a>	<a href="#">Nupl1.cSep08</a>	<a href="#">245922</a>	6563	271	4	90	nucleoporin like 1 (Nupl1) alternative variant cSep08, mRNA.
<a href="#">Nupl2</a>	<a href="#">Nupl2.cSep08</a>	<a href="#">499974</a>	7153	396	3	17	nucleoporin like 2 (Nupl2) alternative variant cSep08, mRNA.
<a href="#">nupor</a>	<a href="#">nupor.aSep08</a>		1026	347		96	promoter-binding protein like (nupor) mRNA.
<a href="#">Nupr1</a>	<a href="#">Nupr1.bSep08</a>	<a href="#">113900</a>	1986	458	1	53	nuclear protein 1 (Nupr1) alternative variant bSep08, mRNA.
<a href="#">Nus1</a>	<a href="#">Nus1.aSep08</a>	<a href="#">294400</a>	26850	3712		345	nuclear undecaprenyl pyrophosphate synthase 1 homolog ( <i>S. cerevisiae</i> ) (Nus1) mRNA.
<a href="#">nusa</a>	<a href="#">nusa.aSep08</a>		483	400		132	putative protein of mammalian origin (nusa) mRNA.
<a href="#">Nusap1</a>	<a href="#">Nusap1.aSep08</a>	<a href="#">311336</a>	30559	2091	11	416	nucleolar and spindle associated protein 1 (47.4 kD) (Nusap1) alternative variant aSep08, mRNA.
<a href="#">Nusap1</a>	<a href="#">Nusap1.cSep08</a>	<a href="#">311336</a>	13557	1389	6	220	nucleolar and spindle associated protein 1 (Nusap1) alternative variant cSep08, mRNA.
<a href="#">Nusap1</a>	<a href="#">Nusap1.dSep08</a>	<a href="#">311336</a>	12103	698	3	106	nucleolar and spindle associated protein 1 (12.1 kD) (Nusap1) alternative variant dSep08, mRNA.
<a href="#">nushee</a>	<a href="#">nushee.aSep08</a>		4730	631		147	CRA b (16.6 kD) (nushee) mRNA.
<a href="#">nutu</a>	<a href="#">nutu.aSep08</a>		13370	401		72	putative protein (nutu) mRNA.
<a href="#">nuvar</a>	<a href="#">nuvar.aSep08</a>		1187	470		65	putative protein (7.5 kD) (nuvar) mRNA.
<a href="#">nuwey</a>	<a href="#">nuwey.aSep08</a>		2277	656			
<a href="#">Nvl</a>	<a href="#">Nvl.bSep08</a>	<a href="#">289323</a>	12947	763	6	191	nuclear VCP-like CRA b (Nvl) alternative variant bSep08, mRNA.
<a href="#">Nvl</a>	<a href="#">Nvl.cSep08</a>	<a href="#">289323</a>	12649	554	5	137	nuclear VCP-like CRA b (Nvl) alternative variant cSep08, mRNA.
<a href="#">Nvl</a>	<a href="#">Nvl.dSep08</a>	<a href="#">289323</a>	5193	431	3	77	nuclear VCP-like (Nvl) alternative variant dSep08, mRNA.
<a href="#">Nvl</a>	<a href="#">Nvl.eSep08</a>	<a href="#">289323</a>	1649	384	2	51	nuclear VCP-like (Nvl) alternative variant eSep08, mRNA.
<a href="#">Nvl</a>	<a href="#">Nvl.fSep08</a>	<a href="#">289323</a>	2024	739	2	44	nuclear VCP-like (5.1 kD) (Nvl) alternative variant fSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.bSep08</a>	<a href="#">59087</a>	5554	1783	4	310	nuclear export factor (Nxf1) alternative variant bSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.bSep08</a>	<a href="#">690285</a>	5554	1783	4	310	nuclear export factor (Nxf1) alternative variant bSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.cSep08</a>	<a href="#">59087</a>	14634	3060	6	251	nuclear RNA export factor 1 (28.2 kD) (Nxf1) alternative variant cSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.cSep08</a>	<a href="#">690285</a>	14634	3060	6	251	nuclear RNA export factor 1 (28.2 kD) (Nxf1) alternative variant cSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.dSep08</a>	<a href="#">59087</a>	2542	723	2	170	nuclear RNA export factor 1 (Nxf1) alternative variant dSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.dSep08</a>	<a href="#">690285</a>	2542	723	2	170	nuclear RNA export factor 1 (Nxf1) alternative variant dSep08, mRNA.
<a href="#">Nxf1</a>	<a href="#">Nxf1.eSep08</a>	<a href="#">59087</a>	1966	1629	2	77	nuclear RNA export factor 1 (Nxf1) alternative variant eSep08, mRNA.

<a href="#">Nxf1</a>	<a href="#">Nxf1.eSep08</a>	<a href="#">690285</a>	1966	1629	2	77	nuclear RNA export factor 1 (Nxf1) alternative variant eSep08, mRNA.
<a href="#">Nxf3</a>	<a href="#">Nxf3.aSep08</a>	<a href="#">302591</a>	3786	1274		231	nuclear RNA export factor 3 (Nxf3) mRNA.
<a href="#">Nxf7</a>	<a href="#">Nxf7.bSep08</a>	<a href="#">501621</a>	5549	1151	1	184	nuclear RNA export factor 7 (21.3 kD) (Nxf7) alternative variant bSep08, mRNA.
<a href="#">Nxn</a>	<a href="#">Nxn.aSep08</a>	<a href="#">360577</a>	138185	2649	8	451	nucleoredoxin (Nxn) alternative variant aSep08, mRNA.
<a href="#">Nxn</a>	<a href="#">Nxn.cSep08</a>	<a href="#">360577</a>	10721	958	3	147	nucleoredoxin (Nxn) alternative variant cSep08, mRNA.
<a href="#">Nxn</a>	<a href="#">Nxn.dSep08</a>	<a href="#">360577</a>	926	529	2	89	nucleoredoxin (Nxn) alternative variant dSep08, mRNA.
<a href="#">Nxn1</a>	<a href="#">Nxn1.bSep08</a>	<a href="#">306342</a>	21617	422	3	112	nucleoredoxin-like 1 (12.9 kD) (Nxn1) alternative variant bSep08, mRNA.
<a href="#">Nxn1</a>	<a href="#">Nxn1.cSep08</a>	<a href="#">306342</a>	18769	332	2	110	nucleoredoxin-like 1 (Nxn1) alternative variant cSep08, mRNA.
<a href="#">Nxph1</a>	<a href="#">Nxph1.bSep08</a>	<a href="#">25501</a>	95673	449	2	92	neurexophilin 1 (Nxph1) alternative variant bSep08, mRNA.
<a href="#">Nxph3</a>	<a href="#">Nxph3.bSep08</a>	<a href="#">59315</a>	3071	548	2	34	neurexophilin 3 (3.7 kD) (Nxph3) alternative variant bSep08, mRNA.
<a href="#">Nxph4</a>	<a href="#">Nxph4.aSep08</a>	<a href="#">59316</a>	12739	763		254	neurexophilin 4 (Nxph4) mRNA.
<a href="#">Nxt2</a>	<a href="#">Nxt2.bSep08</a>	<a href="#">315352</a>	4389	966	1	134	nuclear transport factor 2-like export factor 2 (15.5 kD) (Nxt2) alternative variant bSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.aSep08</a>	<a href="#">290626</a>	15420	1481	8	462	sarcoma antigen NY-SAR-48 (Ny-sar-48) alternative variant aSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.cSep08</a>	<a href="#">290626</a>	9917	1257	8	201	sarcoma antigen NY-SAR-48 (22.7 kD) (Ny-sar-48) alternative variant cSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.dSep08</a>	<a href="#">290626</a>	4343	700	4	201	sarcoma antigen NY-SAR-48 (Ny-sar-48) alternative variant dSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.eSep08</a>	<a href="#">290626</a>	4330	545	3	149	sarcoma antigen NY-SAR-48 (Ny-sar-48) alternative variant eSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.fSep08</a>	<a href="#">290626</a>	9000	419	5	83	sarcoma antigen NY-SAR-48 (Ny-sar-48) alternative variant fSep08, mRNA.
<a href="#">Ny-sar-48</a>	<a href="#">Ny-sar-48.hSep08</a>	<a href="#">290626</a>	2025	313	2	47	sarcoma antigen NY-SAR-48 (5.0 kD) (Ny-sar-48) alternative variant hSep08, mRNA.
<a href="#">nyby</a>	<a href="#">nyby.aSep08</a>		8230	682		226	polymerase alpha (nyby) mRNA.
<a href="#">nychy</a>	<a href="#">nychy.aSep08</a>		2881	385		128	protein tyrosine phosphatase receptor type J (nychy) mRNA.
<a href="#">nydar</a>	<a href="#">nydar.aSep08</a>		6127	511		36	putative protein (nydar) mRNA.
<a href="#">nyfer</a>	<a href="#">nyfer.aSep08</a>		13809	370		33	putative protein (nyfer) mRNA.
<a href="#">nyflo</a>	<a href="#">nyflo.aSep08</a>		3647	698		90	putative mitochondrial protein (9.8 kD) (nyflo) mRNA.
<a href="#">nyflu</a>	<a href="#">nyflu.aSep08</a>		34660	606	3	140	putative mitochondrial protein (13.8 kD) (nyflu) mRNA.
<a href="#">nygar</a>	<a href="#">nygar.aSep08</a>		13976	583		194	putative protein of vertebrate origin (nygar) mRNA.
<a href="#">nyja</a>	<a href="#">nyja.aSep08</a>		6973	695		58	putative protein (nyja) mRNA.
<a href="#">nyjey</a>	<a href="#">nyjey.aSep08</a>		451	267		74	putative protein (nyjey) mRNA.
<a href="#">nykee</a>	<a href="#">nykee.aSep08</a>		8946	1785	1	20	putative protein (nykee) alternative variant aSep08, mRNA.
<a href="#">nykee</a>	<a href="#">nykee.bSep08</a>		8634	551	2	15	putative protein (nykee) alternative variant bSep08, mRNA.
<a href="#">nyloy</a>	<a href="#">nyloy.aSep08</a>		2210	257		60	vacuolar protein sorting 8 homolog (nyloy) mRNA.
<a href="#">nymee</a>	<a href="#">nymee.aSep08</a>		19233	1527		91	putative protein (nymee) mRNA.



<a href="#">nymer</a>	<a href="#">nymer.aSep08</a>		2574	723		40	putative protein (nymer) mRNA.
<a href="#">nynoy</a>	<a href="#">nynoy.aSep08</a>		859	281		85	putative protein of vertebrate origin (nynoy) mRNA.
<a href="#">nypor</a>	<a href="#">nypor.aSep08</a>		1581	512		109	promoter-binding protein like (nypor) mRNA.
<a href="#">nysa</a>	<a href="#">nysa.aSep08</a>		4249	507		108	putative protein (nysa) mRNA.
<a href="#">nyshee</a>	<a href="#">nyshee.aSep08</a>		4727	385		66	putative protein (nyshee) mRNA.
<a href="#">nytu</a>	<a href="#">nytu.aSep08</a>		1235	246		29	putative protein (nytu) mRNA.
<a href="#">nyvar</a>	<a href="#">nyvar.aSep08</a>		1440	859		84	putative protein (nyvar) mRNA.
<a href="#">Nyw1</a>	<a href="#">Nyw1.bSep08</a>	<a href="#">59319</a>	17537	1945	4	307	ischemia related factor NYW-1 (Nyw1) alternative variant bSep08, mRNA.
<a href="#">Nyw1</a>	<a href="#">Nyw1.cSep08</a>	<a href="#">59319</a>	14632	745	5	197	ischemia related factor NYW-1 (Nyw1) alternative variant cSep08, mRNA.
<a href="#">Nyw1</a>	<a href="#">Nyw1.dSep08</a>	<a href="#">59319</a>	7502	756	3	134	ischemia related factor NYW-1 (Nyw1) alternative variant dSep08, mRNA.
<a href="#">Nyw1</a>	<a href="#">Nyw1.eSep08</a>	<a href="#">59319</a>	18821	331	3	110	ischemia related factor NYW-1 (Nyw1) alternative variant eSep08, mRNA.
<a href="#">nywey</a>	<a href="#">nywey.aSep08</a>		1804	779		151	envelope protein (nywey) mRNA.
<a href="#">Oaf</a>	<a href="#">Oaf.bSep08</a>	<a href="#">315594</a>	3484	1476	1	154	OAF homolog (Drosophila) (17.9 kD) (Oaf) alternative variant bSep08, mRNA.
<a href="#">Oas1a</a>	<a href="#">Oas1a.aSep08</a>	<a href="#">192281</a>	10657	1422	7	379	2'-5' oligoadenylate synthetase 1A (Oas1a) alternative variant aSep08, mRNA.
<a href="#">Oas1a</a>	<a href="#">Oas1a.cSep08</a>	<a href="#">192281</a>	1392	766	2	169	2'-5' oligoadenylate synthetase 1A (Oas1a) alternative variant cSep08, mRNA.
<a href="#">Oas1k</a>	<a href="#">Oas1k.bSep08</a>	<a href="#">494198</a>	5954	724	3	241	2'-5' oligoadenylate synthetase 1K (Oas1k) alternative variant bSep08, mRNA.
<a href="#">Oasl2</a>	<a href="#">Oasl2.bSep08</a>	<a href="#">304549</a>	5282	722	3	240	2'-5' oligoadenylate synthetase-like 2 (Oasl2) alternative variant bSep08, mRNA.
<a href="#">Oasl2</a>	<a href="#">Oasl2.cSep08</a>	<a href="#">304549</a>	3034	677	2	174	2'-5' oligoadenylate synthetase-like 2 (19.9 kD) (Oasl2) alternative variant cSep08, mRNA.
<a href="#">Oasl2</a>	<a href="#">Oasl2.dSep08</a>	<a href="#">304549</a>	882	338	2	106	2'-5' oligoadenylate synthetase-like 2 (Oasl2) alternative variant dSep08, mRNA.
<a href="#">Oat</a>	<a href="#">Oat.bSep08</a>	<a href="#">64313</a>	3175	532	3	143	ornithine aminotransferase (Oat) alternative variant bSep08, mRNA.
<a href="#">Oat</a>	<a href="#">Oat.cSep08</a>	<a href="#">64313</a>	2723	771	2	87	ornithine aminotransferase (Oat) alternative variant cSep08, mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.bSep08</a>	<a href="#">25502</a>	2266	848	5	207	ornithine decarboxylase antizyme 1 (22.6 kD) (Oaz1) alternative variant bSep08, mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.cSep08</a>	<a href="#">25502</a>	2466	1032	5	172	ornithine decarboxylase antizyme 1 (18.6 kD) (Oaz1) alternative variant cSep08, complete mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.dSep08</a>	<a href="#">25502</a>	876	728	3	140	ornithine decarboxylase antizyme 1 (15.8 kD) (Oaz1) alternative variant dSep08, mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.eSep08</a>	<a href="#">25502</a>	2219	661	4	125	ornithine decarboxylase antizyme 1 (14.1 kD) (Oaz1) alternative variant eSep08, mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.fSep08</a>	<a href="#">25502</a>	1196	1118	2	69	ornithine decarboxylase antizyme 1 (7.9 kD) (Oaz1) alternative variant fSep08, mRNA.
<a href="#">Oaz1</a>	<a href="#">Oaz1.gSep08</a>	<a href="#">25502</a>	637	533	2	33	ornithine decarboxylase antizyme 1 (3.6 kD) (Oaz1) alternative variant gSep08, mRNA.

<a href="#">Oaz2</a>	<a href="#">Oaz2.aSep08</a>	<a href="#">501454</a>	12882	862	2	166	ornithine decarboxylase antizyme 2 (18.4 kD) (Oaz2) alternative variant aSep08, mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.bSep08</a>	<a href="#">294025</a>	33817	1337	10	377	oligonucleotide oligosaccharide-binding fold containing 1 like (43.4 kD) (Obfc1) alternative variant bSep08, complete mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.cSep08</a>	<a href="#">294025</a>	33810	1465	10	376	oligonucleotide oligosaccharide-binding fold containing 1 like (43.3 kD) (Obfc1) alternative variant cSep08, complete mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.dSep08</a>	<a href="#">294025</a>	14851	1251	6	312	oligonucleotide oligosaccharide-binding fold containing 1 like (35.5 kD) (Obfc1) alternative variant dSep08, mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.eSep08</a>	<a href="#">294025</a>	12023	1096	4	137	oligonucleotide oligosaccharide-binding fold containing 1 like (15.7 kD) (Obfc1) alternative variant eSep08, mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.fSep08</a>	<a href="#">294025</a>	13285	760	5	108	oligonucleotide oligosaccharide-binding fold containing 1 like (12.3 kD) (Obfc1) alternative variant fSep08, mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.gSep08</a>	<a href="#">294025</a>	7422	593	3	89	oligonucleotide oligosaccharide-binding fold containing 1 like (10.2 kD) (Obfc1) alternative variant gSep08, mRNA.
<a href="#">Obfc1</a>	<a href="#">Obfc1.iSep08</a>	<a href="#">294025</a>	5677	392	2	59	putative protein (Obfc1) alternative variant iSep08, mRNA.
<a href="#">Obfc2a</a>	<a href="#">Obfc2a.bSep08</a>	<a href="#">363227</a>	5947	798	6	118	oligonucleotide/oligosaccharide-binding fold containing 2A (13.1 kD) (Obfc2a) alternative variant bSep08, mRNA.
<a href="#">Obfc2a</a>	<a href="#">Obfc2a.cSep08</a>	<a href="#">363227</a>	5871	1382	5	118	oligonucleotide/oligosaccharide-binding fold containing 2A (13.1 kD) (Obfc2a) alternative variant cSep08, mRNA.
<a href="#">Obfc2b</a>	<a href="#">Obfc2b.aSep08</a>	<a href="#">362813</a>	5711	1082	5	251	oligonucleotide/oligosaccharide-binding fold containing 2B (Obfc2b) alternative variant aSep08, mRNA.
<a href="#">Obfc2b</a>	<a href="#">Obfc2b.cSep08</a>	<a href="#">362813</a>	5486	942	5	211	oligonucleotide/oligosaccharide-binding fold containing 2B (22.4 kD) (Obfc2b) alternative variant cSep08, mRNA.
<a href="#">Obfc2b</a>	<a href="#">Obfc2b.dSep08</a>	<a href="#">362813</a>	1210	539	2	80	oligonucleotide/oligosaccharide-binding fold containing 2B (Obfc2b) alternative variant dSep08, mRNA.
<a href="#">Obp3</a>	<a href="#">Obp3.aSep08</a>	<a href="#">259247</a>	3466	885	1	206	alpha-2u globulin PGCL4 (Obp3) alternative variant aSep08, mRNA.
<a href="#">Obp3</a>	<a href="#">Obp3.dSep08</a>	<a href="#">259247</a>	3341	772	1	195	alpha-2u globulin PGCL4 (Obp3) alternative variant dSep08, mRNA.
<a href="#">Obscn</a>	<a href="#">Obscn.aSep08</a>	<a href="#">338458</a>	19448	376	4	104	obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF (Obscn) alternative variant aSep08, mRNA.
<a href="#">Obscnl</a>	<a href="#">Obscnl.aSep08</a>	<a href="#">287353</a>	3005	773		257	obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF-like (Obscnl) mRNA.
<a href="#">Obsl1</a>	<a href="#">Obsl1.aSep08</a>	<a href="#">363259</a>	8378	2732	8	821	obscurin-like 1 (Obsl1) alternative variant aSep08, mRNA.
<a href="#">OCIA.0</a>	<a href="#">OCIA.0.aSep08</a>		18876	1980	3	283	ovarian carcinoma immunoreactive antigen (31.1 kD) (OCIA.0) alternative variant aSep08, mRNA.
<a href="#">OCIA.0</a>	<a href="#">OCIA.0.bSep08</a>		16313	1667	1	194	ovarian carcinoma immunoreactive antigen (OCIA.0) alternative variant bSep08, mRNA.
<a href="#">OCIA.0</a>	<a href="#">OCIA.0.cSep08</a>		27011	752	1	119	ovarian carcinoma immunoreactive antigen (OCIA.0) alternative variant cSep08, mRNA.
<a href="#">Ociad1</a>	<a href="#">Ociad1.bSep08</a>	<a href="#">289590</a>	13858	685	6	179	ovarian carcinoma immunoreactive antigen (Ociad1) alternative variant bSep08, mRNA.
<a href="#">Ociad1</a>	<a href="#">Ociad1.cSep08</a>	<a href="#">289590</a>	13855	772	7	163	ovarian carcinoma immunoreactive antigen (Ociad1) alternative variant cSep08, mRNA.

<a href="#">Ociad1</a>	<a href="#">Ociad1.dSep08</a>	<a href="#">289590</a>	7143	557	2	36	putative protein of mammalian origin (Ociad1) alternative variant dSep08, mRNA.
<a href="#">Ociad1</a>	<a href="#">Ociad1.fSep08</a>	<a href="#">289590</a>	7197	655	2	63	putative protein of mammalian origin (Ociad1) alternative variant fSep08, mRNA.
<a href="#">Ocln</a>	<a href="#">Ocln.aSep08</a>	<a href="#">83497</a>	49323	3080	9	523	occludin (59.2 kD) (Ocln) alternative variant aSep08, mRNA.
<a href="#">Ocln</a>	<a href="#">Ocln.bSep08</a>	<a href="#">83497</a>	12545	776	3	229	occludin (Ocln) alternative variant bSep08, mRNA.
<a href="#">Ocln</a>	<a href="#">Ocln.cSep08</a>	<a href="#">83497</a>	12457	732	3	195	occludin (Ocln) alternative variant cSep08, mRNA.
<a href="#">Ocln</a>	<a href="#">Ocln.dSep08</a>	<a href="#">83497</a>	7976	702	5	188	occludin (Ocln) alternative variant dSep08, mRNA.
<a href="#">Ocm</a>	<a href="#">Ocm.aSep08</a>	<a href="#">25503</a>	6856	637	1	127	oncomodulin (Ocm) alternative variant aSep08, mRNA.
<a href="#">Ocri</a>	<a href="#">Ocri.bSep08</a>	<a href="#">317576</a>	13606	743	7	210	oculocerebrorrenal syndrome of Lowe (Ocri) alternative variant bSep08, mRNA.
<a href="#">Odc1</a>	<a href="#">Odc1.bSep08</a>	<a href="#">24609</a>	3304	746	3	248	ornithine decarboxylase 1 (Odc1) alternative variant bSep08, mRNA.
<a href="#">Odc1</a>	<a href="#">Odc1.cSep08</a>	<a href="#">24609</a>	3477	807	4	205	ornithine decarboxylase 1 (Odc1) alternative variant cSep08, mRNA.
<a href="#">Odc1</a>	<a href="#">Odc1.dSep08</a>	<a href="#">24609</a>	3360	886	4	194	ornithine decarboxylase 1 (Odc1) alternative variant dSep08, mRNA.
<a href="#">Odc1</a>	<a href="#">Odc1.eSep08</a>	<a href="#">24609</a>	3666	1363	3	150	ornithine decarboxylase 1 (16.6 kD) (Odc1) alternative variant eSep08, mRNA.
<a href="#">Odc1</a>	<a href="#">Odc1.fSep08</a>	<a href="#">24609</a>	2601	406	2	45	ornithine decarboxylase 1 (Odc1) alternative variant fSep08, mRNA.
<a href="#">Odf1</a>	<a href="#">Odf1.aSep08</a>	<a href="#">24610</a>	6121	636	1	171	outer dense fiber of sperm tails 1 (Odf1) alternative variant aSep08, mRNA.
<a href="#">Odf1</a>	<a href="#">Odf1.bSep08</a>	<a href="#">24610</a>	12186	727	1	169	outer dense fiber of sperm tails 1 (18.3 kD) (Odf1) alternative variant bSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.bSep08</a>	<a href="#">29479</a>	18109	1441	13	449	outer dense fiber 2 (Odf2) alternative variant bSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.cSep08</a>	<a href="#">29479</a>	17076	901	7	300	outer dense fiber of sperm tails 2 CRA e (Odf2) alternative variant cSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.dSep08</a>	<a href="#">29479</a>	13802	867	6	276	outer dense fiber of sperm tails 2 CRA a (Odf2) alternative variant dSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.eSep08</a>	<a href="#">29479</a>	10085	778	7	225	outer dense fiber 2 (Odf2) alternative variant eSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.fSep08</a>	<a href="#">29479</a>	10061	764	7	224	outer dense fiber 2 (Odf2) alternative variant fSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.gSep08</a>	<a href="#">29479</a>	11120	2074	4	186	outer dense fiber 2 (21.6 kD) (Odf2) alternative variant gSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.hSep08</a>	<a href="#">29479</a>	7397	496	3	150	outer dense fiber 2 (Odf2) alternative variant hSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.iSep08</a>	<a href="#">29479</a>	15000	764	6	148	outer dense fiber 2 (Odf2) alternative variant iSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.jSep08</a>	<a href="#">29479</a>	40987	483	4	142	outer dense fiber of sperm tails 2 CRA a (Odf2) alternative variant jSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.kSep08</a>	<a href="#">29479</a>	9480	777	5	136	outer dense fiber 2 (Odf2) alternative variant kSep08, mRNA.

<a href="#">Odf2</a>	<a href="#">Odf2.lSep08</a>	<a href="#">29479</a>	2128	718	3	82	outer dense fiber 2 (Odf2) alternative variant lSep08, mRNA.
<a href="#">Odf2</a>	<a href="#">Odf2.mSep08</a>	<a href="#">29479</a>	7501	638	3	82	outer dense fiber of sperm tails 2 CRA a (Odf2) alternative variant mSep08, mRNA.
<a href="#">Odf2l</a>	<a href="#">Odf2l.bSep08</a>	<a href="#">685425</a>	36713	2516	15	379	outer dense fiber of sperm tails 2-like (Odf2l) alternative variant bSep08, mRNA.
<a href="#">Odf2l</a>	<a href="#">Odf2l.cSep08</a>	<a href="#">685425</a>	7310	367	4	122	outer dense fiber of sperm tails 2-like (Odf2l) alternative variant cSep08, mRNA.
<a href="#">Odf2l</a>	<a href="#">Odf2l.dSep08</a>	<a href="#">685425</a>	1375	406	2	68	outer dense fiber of sperm tails 2-like (8.2 kD) (Odf2l) alternative variant dSep08, mRNA.
<a href="#">Odf3</a>	<a href="#">Odf3.bSep08</a>	<a href="#">365387</a>	1125	717	1	101	outer dense fiber of sperm tails 3 (11.0 kD) (Odf3) alternative variant bSep08, mRNA.
<a href="#">Odf4</a>	<a href="#">Odf4.aSep08</a>	<a href="#">303236</a>	5461	1345	5	326	outer dense fiber of sperm tails 4 (Odf4) alternative variant aSep08, mRNA.
<a href="#">Odf4</a>	<a href="#">Odf4.cSep08</a>	<a href="#">303236</a>	4521	791	4	211	outer dense fiber of sperm tails 4 (Odf4) alternative variant cSep08, mRNA.
<a href="#">Odf4</a>	<a href="#">Odf4.dSep08</a>	<a href="#">303236</a>	1270	940	2	57	outer dense fiber of sperm tails 4 (6.0 kD) (Odf4) alternative variant dSep08, mRNA.
<a href="#">Odz2</a>	<a href="#">Odz2.aSep08</a>	<a href="#">117242</a>	13443	993		330	odd Oz/ten-m homolog 2 (Drosophila) (Odz2) mRNA.
<a href="#">Odz3</a>	<a href="#">Odz3.aSep08</a>	<a href="#">306451</a>	15494	742		247	odd Oz/ten-m homolog 3 (Drosophila) (Odz3) mRNA.
<a href="#">Odz4</a>	<a href="#">Odz4.aSep08</a>	<a href="#">308831</a>	26080	395		131	odd Oz/ten-m homolog 4 (Drosophila) (Odz4) mRNA.
<a href="#">Ofd1</a>	<a href="#">Ofd1.bSep08</a>	<a href="#">302661</a>	8409	431	4	143	oral-facial-digital syndrome 1 gene homolog (human) (Ofd1) alternative variant bSep08, mRNA.
<a href="#">Ogdh</a>	<a href="#">Ogdh.bSep08</a>	<a href="#">360975</a>	44485	764	6	254	oxoglutarate dehydrogenase (lipoamide) (Ogdh) alternative variant bSep08, mRNA.
<a href="#">Ogdh</a>	<a href="#">Ogdh.cSep08</a>	<a href="#">360975</a>	19068	574	3	190	oxoglutarate dehydrogenase (lipoamide) (Ogdh) alternative variant cSep08, mRNA.
<a href="#">Ogdh</a>	<a href="#">Ogdh.dSep08</a>	<a href="#">360975</a>	1373	833	2	186	oxoglutarate dehydrogenase (lipoamide) (20.6 kD) (Ogdh) alternative variant dSep08, mRNA.
<a href="#">Ogdh</a>	<a href="#">Ogdh.eSep08</a>	<a href="#">360975</a>	4903	701	3	160	oxoglutarate dehydrogenase (lipoamide) (Ogdh) alternative variant eSep08, mRNA.
<a href="#">Ogdh</a>	<a href="#">Ogdh.fSep08</a>	<a href="#">360975</a>	20165	495	3	119	oxoglutarate dehydrogenase (lipoamide) (Ogdh) alternative variant fSep08, mRNA.
<a href="#">Ogdhl</a>	<a href="#">Ogdhl.bSep08</a>	<a href="#">290566</a>	10035	930	6	297	oxoglutarate dehydrogenase-like (Ogdhl) alternative variant bSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.cSep08</a>	<a href="#">288656</a>	5008	3293	5	229	putative protein of ancient origin (25.4 kD) (Ogfod2andArl6ip4) alternative variant cSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.cSep08</a>	<a href="#">288657</a>	5008	3293	5	229	putative protein of ancient origin (25.4 kD) (Ogfod2andArl6ip4) alternative variant cSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.dSep08</a>	<a href="#">288656</a>	966	529	3	175	ADP-ribosylation factor-like 6 interacting protein 4 (Ogfod2andArl6ip4) alternative variant dSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.dSep08</a>	<a href="#">288657</a>	966	529	3	175	ADP-ribosylation factor-like 6 interacting protein 4 (Ogfod2andArl6ip4) alternative variant dSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.eSep08</a>	<a href="#">288656</a>	1043	532	3	93	ADP-ribosylation-like factor 6 interacting protein 4 (Ogfod2andArl6ip4) alternative variant eSep08, mRNA.
<a href="#">Ogfod2andArl6ip4</a>	<a href="#">Ogfod2andArl6ip4.eSep08</a>	<a href="#">288657</a>	1043	532	3	93	ADP-ribosylation-like factor 6 interacting protein 4 (Ogfod2andArl6ip4) alternative variant eSep08, mRNA.

<a href="#">Ogfr</a>	<a href="#">Ogfr.aSep08</a>	<a href="#">83525</a>	6291	2061	7	571	opioid growth factor receptor (63.8 kD) (Ogfr) alternative variant aSep08, complete mRNA.
<a href="#">Ogfr1</a>	<a href="#">Ogfr1.cSep08</a>	<a href="#">316290</a>	4558	534	3	154	opioid growth factor receptor-like 1 (Ogfr1) alternative variant cSep08, mRNA.
<a href="#">Ogg1</a>	<a href="#">Ogg1.bSep08</a>	<a href="#">81528</a>	1742	446	1	55	8-oxoguanine DNA-glycosylase 1 (6.1 kD) (Ogg1) alternative variant bSep08, mRNA.
<a href="#">Ogg1</a>	<a href="#">Ogg1.cSep08</a>	<a href="#">81528</a>	9516	1189	6	137	8-oxoguanine DNA-glycosylase 1 (15.6 kD) (Ogg1) alternative variant cSep08, mRNA.
<a href="#">Ogn</a>	<a href="#">Ogn.bSep08</a>	<a href="#">291015</a>	20474	1367	9	298	osteoglycin (34.1 kD) (Ogn) alternative variant bSep08, mRNA.
<a href="#">Ogn</a>	<a href="#">Ogn.cSep08</a>	<a href="#">291015</a>	13018	703	4	234	osteoglycin (Ogn) alternative variant cSep08, mRNA.
<a href="#">Ogn</a>	<a href="#">Ogn.dSep08</a>	<a href="#">291015</a>	2402	325	2	30	osteoglycin (3.7 kD) (Ogn) alternative variant dSep08, complete mRNA.
<a href="#">Ogt</a>	<a href="#">Ogt.bSep08</a>	<a href="#">26295</a>	12197	1806	4	405	O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (Ogt) alternative variant bSep08, mRNA.
<a href="#">Ogt</a>	<a href="#">Ogt.cSep08</a>	<a href="#">26295</a>	27203	736	6	194	O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (Ogt) alternative variant cSep08, mRNA.
<a href="#">Ogt</a>	<a href="#">Ogt.eSep08</a>	<a href="#">26295</a>	3879	388	3	69	O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (Ogt) alternative variant eSep08, mRNA.
<a href="#">Oit1</a>	<a href="#">Oit1.aSep08</a>	<a href="#">289949</a>	33667	1785	8	543	oncoprotein induced transcript 1 (Oit1) alternative variant aSep08, mRNA.
<a href="#">Oit1</a>	<a href="#">Oit1.cSep08</a>	<a href="#">289949</a>	31160	769	9	170	oncoprotein induced transcript 1 (Oit1) alternative variant cSep08, mRNA.
<a href="#">Oit1</a>	<a href="#">Oit1.dSep08</a>	<a href="#">289949</a>	6295	703	4	119	oncoprotein induced transcript 1 (Oit1) alternative variant dSep08, mRNA.
<a href="#">Olfm1</a>	<a href="#">Olfm1.bSep08</a>	<a href="#">93667</a>	22002	980	4	184	olfactomedin 1 (Olfm1) alternative variant bSep08, mRNA.
<a href="#">Olfm1</a>	<a href="#">Olfm1.cSep08</a>	<a href="#">93667</a>	21230	720	4	130	olfactomedin 1 (Olfm1) alternative variant cSep08, mRNA.
<a href="#">Olfm1</a>	<a href="#">Olfm1.dSep08</a>	<a href="#">93667</a>	977	887	2	75	olfactomedin 1 (Olfm1) alternative variant dSep08, mRNA.
<a href="#">Olfm2</a>	<a href="#">Olfm2.bSep08</a>	<a href="#">313783</a>	3768	749	2	249	olfactomedin 2 (Olfm2) alternative variant bSep08, mRNA.
<a href="#">Olfm2</a>	<a href="#">Olfm2.cSep08</a>	<a href="#">313783</a>	75453	391	4	130	olfactomedin 2 (Olfm2) alternative variant cSep08, mRNA.
<a href="#">Olfml2b</a>	<a href="#">Olfml2b.bSep08</a>	<a href="#">304960</a>	4465	487	1	124	olfactomedin-like 2B (Olfml2b) alternative variant bSep08, mRNA.
<a href="#">Olfml3</a>	<a href="#">Olfml3.cSep08</a>	<a href="#">310743</a>	910	398	2	74	olfactomedin-like 3 (Olfml3) alternative variant cSep08, mRNA.
<a href="#">Olr35</a>	<a href="#">Olr35.bSep08</a>	<a href="#">293140</a>	1404	746	3	79	olfactory receptor 35 (8.7 kD) (Olr35) alternative variant bSep08, mRNA.
<a href="#">Olr738-ps</a>	<a href="#">Olr738-ps.aSep08</a>	<a href="#">405500</a>	9290	289		84	olfactory receptor pseudogene 738 (Olr738-ps) mRNA.
<a href="#">Olr1366</a>	<a href="#">Olr1366.bSep08</a>	<a href="#">405335</a>	9397	777	5	105	olfactory receptor 1366 (11.5 kD) (Olr1366) alternative variant bSep08, mRNA.
<a href="#">Olr1366</a>	<a href="#">Olr1366.cSep08</a>	<a href="#">405335</a>	4248	341	3	79	olfactory receptor 1366 (Olr1366) alternative variant cSep08, mRNA.
<a href="#">Olr1705</a>	<a href="#">Olr1705.bSep08</a>	<a href="#">405188</a>	10030	444	4	116	olfactory receptor 1705 (Olr1705) alternative variant bSep08, mRNA.

<a href="#">Omg</a>	<a href="#">Omg.bSep08</a>	<a href="#">450224</a>	2718	1485	2	331	oligodendrocyte-myelin glycoprotein (36.6 kD) (Omg) alternative variant bSep08, mRNA.
<a href="#">Oosp1</a>	<a href="#">Oosp1.aSep08</a>	<a href="#">690893</a>	15213	308		64	oocyte secreted protein 1 (Oosp1) mRNA.
<a href="#">Opa1</a>	<a href="#">Opa1.bSep08</a>	<a href="#">171116</a>	1614	780	2	52	optic atrophy 1 homolog (human) (Opa1) alternative variant bSep08, mRNA.
<a href="#">Opa1</a>	<a href="#">Opa1.cSep08</a>	<a href="#">171116</a>	1398	237	3	43	optic atrophy 1 homolog (human) (Opa1) alternative variant cSep08, mRNA.
<a href="#">Opalin</a>	<a href="#">Opalin.bSep08</a>	<a href="#">361757</a>	8182	460	1	41	oligodendrocytic myelin paranodal and inner loop protein (4.3 kD) (Opalin) alternative variant bSep08, complete mRNA.
<a href="#">Opalin</a>	<a href="#">Opalin.cSep08</a>	<a href="#">361757</a>	15321	584	5	80	oligodendrocytic myelin paranodal and inner loop protein (Opalin) alternative variant cSep08, mRNA.
<a href="#">Ophn1</a>	<a href="#">Ophn1.bSep08</a>	<a href="#">312108</a>	8149	525	1	105	oligophrenin 1 (Ophn1) alternative variant bSep08, mRNA.
<a href="#">Oplah</a>	<a href="#">Oplah.bSep08</a>	<a href="#">116684</a>	3997	1349	4	239	5-oxoprolinase (ATP-hydrolysing) (Oplah) alternative variant bSep08, mRNA.
<a href="#">Oplah</a>	<a href="#">Oplah.cSep08</a>	<a href="#">116684</a>	1800	902	5	189	5-oxoprolinase (ATP-hydrolysing) (Oplah) alternative variant cSep08, mRNA.
<a href="#">Oplah</a>	<a href="#">Oplah.dSep08</a>	<a href="#">116684</a>	2076	717	5	164	5-oxoprolinase (ATP-hydrolysing) (Oplah) alternative variant dSep08, mRNA.
<a href="#">Oplah</a>	<a href="#">Oplah.eSep08</a>	<a href="#">116684</a>	489	371	2	115	5-oxoprolinase (ATP-hydrolysing) (Oplah) alternative variant eSep08, mRNA.
<a href="#">Oplah</a>	<a href="#">Oplah.fSep08</a>	<a href="#">116684</a>	42099	410	3	86	5-oxoprolinase (ATP-hydrolysing) (Oplah) alternative variant fSep08, mRNA.
<a href="#">Opn3</a>	<a href="#">Opn3.aSep08</a>	<a href="#">498289</a>	27717	604	1	201	opsin 3 (Opn3) alternative variant aSep08, mRNA.
<a href="#">Opn3</a>	<a href="#">Opn3.bSep08</a>	<a href="#">498289</a>	628	269	1	69	opsin 3 (Opn3) alternative variant bSep08, mRNA.
<a href="#">Opn4</a>	<a href="#">Opn4.bSep08</a>	<a href="#">192223</a>	5475	1750	1	473	opsin 4 (melanopsin) (Opn4) alternative variant bSep08, mRNA.
<a href="#">Opn5</a>	<a href="#">Opn5.bSep08</a>	<a href="#">316259</a>	13724	745	3	188	opsin 5 (20.6 kD) (Opn5) alternative variant bSep08, mRNA.
<a href="#">Opn5</a>	<a href="#">Opn5.cSep08</a>	<a href="#">316259</a>	10910	360	3	38	opsin 5 (Opn5) alternative variant cSep08, mRNA.
<a href="#">Opr1</a>	<a href="#">Opr1.bSep08</a>	<a href="#">29256</a>	3895	719	2	171	opioid receptor-like 1 (Opr1) alternative variant bSep08, mRNA.
<a href="#">Oprs1</a>	<a href="#">Oprs1.bSep08</a>	<a href="#">29336</a>	1996	905	1	151	opioid receptor, sigma 1 (17.1 kD) (Oprs1) alternative variant bSep08, mRNA.
<a href="#">Oprs1</a>	<a href="#">Oprs1.cSep08</a>	<a href="#">29336</a>	2233	962	2	106	opioid receptor, sigma 1 (Oprs1) alternative variant cSep08, mRNA.
<a href="#">Optn</a>	<a href="#">Optn.aSep08</a>	<a href="#">246294</a>	50649	4047	13	585	optineurin (67.0 kD) (Optn) alternative variant aSep08, mRNA.
<a href="#">Optn</a>	<a href="#">Optn.bSep08</a>	<a href="#">246294</a>	19019	746	5	232	optineurin (Optn) alternative variant bSep08, mRNA.
<a href="#">Optn</a>	<a href="#">Optn.cSep08</a>	<a href="#">246294</a>	7194	360	1	96	optineurin (Optn) alternative variant cSep08, mRNA.
<a href="#">Orai2</a>	<a href="#">Orai2.aSep08</a>	<a href="#">304592</a>	17741	1668		313	ORAI calcium release-activated calcium modulator 2 (34.5 kD) (Orai2) mRNA.
<a href="#">Oraov1</a>	<a href="#">Oraov1.bSep08</a>	<a href="#">309136</a>	4875	622	2	173	oral cancer overexpressed 1 (Oraov1) alternative variant bSep08, mRNA.
<a href="#">Oraov1</a>	<a href="#">Oraov1.cSep08</a>	<a href="#">309136</a>	4873	617	2	172	oral cancer overexpressed 1 (Oraov1) alternative variant cSep08, mRNA.

<a href="#">Orc2l</a>	<a href="#">Orc2l.bSep08</a>	<a href="#">301430</a>	9742	2247	7	274	origin recognition complex, subunit 2-like ( <i>S. cerevisiae</i> ) (Orc2l) alternative variant bSep08, mRNA.
<a href="#">Orc2l</a>	<a href="#">Orc2l.cSep08</a>	<a href="#">301430</a>	20615	703	7	169	origin recognition complex, subunit 2-like ( <i>S. cerevisiae</i> ) (Orc2l) alternative variant cSep08, mRNA.
<a href="#">Orc2l</a>	<a href="#">Orc2l.dSep08</a>	<a href="#">301430</a>	14142	585	6	137	origin recognition complex, subunit 2-like ( <i>S. cerevisiae</i> ) (Orc2l) alternative variant dSep08, mRNA.
<a href="#">Orc3l</a>	<a href="#">Orc3l.bSep08</a>	<a href="#">313138</a>	2414	935	1	91	origin recognition complex, subunit 3-like ( <i>S. cerevisiae</i> ) (Orc3l) alternative variant bSep08, mRNA.
<a href="#">Orc4l</a>	<a href="#">Orc4l.bSep08</a>	<a href="#">295596</a>	10479	846	3	139	origin recognition complex, subunit 4-like ( <i>S. cerevisiae</i> ) (Orc4l) alternative variant bSep08, mRNA.
<a href="#">Orc4l</a>	<a href="#">Orc4l.cSep08</a>	<a href="#">295596</a>	10808	637	5	130	origin recognition complex, subunit 4-like ( <i>S. cerevisiae</i> ) (Orc4l) alternative variant cSep08, mRNA.
<a href="#">Orc4l</a>	<a href="#">Orc4l.dSep08</a>	<a href="#">295596</a>	76271	437	1	69	origin recognition complex, subunit 4-like ( <i>S. cerevisiae</i> ) (Orc4l) alternative variant dSep08, mRNA.
<a href="#">Orc5l</a>	<a href="#">Orc5l.bSep08</a>	<a href="#">362304</a>	29059	698	6	221	origin recognition complex (Orc5l) alternative variant bSep08, mRNA.
<a href="#">Orc5l</a>	<a href="#">Orc5l.cSep08</a>	<a href="#">362304</a>	39718	1313	7	171	origin recognition complex (19.7 kD) (Orc5l) alternative variant cSep08, mRNA.
<a href="#">Orc5l</a>	<a href="#">Orc5l.dSep08</a>	<a href="#">362304</a>	9603	625	6	152	origin recognition complex (Orc5l) alternative variant dSep08, mRNA.
<a href="#">Orc5l</a>	<a href="#">Orc5l.eSep08</a>	<a href="#">362304</a>	12970	429	5	143	origin recognition complex (Orc5l) alternative variant eSep08, mRNA.
<a href="#">Orc6l</a>	<a href="#">Orc6l.aSep08</a>	<a href="#">291927</a>	7484	1777	5	545	origin recognition complex, subunit 6-like ( <i>S. cerevisiae</i> ) (Orc6l) alternative variant aSep08, mRNA.
<a href="#">Orc6l</a>	<a href="#">Orc6l.cSep08</a>	<a href="#">291927</a>	2854	462	4	139	origin recognition complex, subunit 6-like ( <i>S. cerevisiae</i> ) (Orc6l) alternative variant cSep08, mRNA.
<a href="#">Orc6l</a>	<a href="#">Orc6l.fSep08</a>	<a href="#">291927</a>	2236	186	3	31	origin recognition complex, subunit 6-like ( <i>S. cerevisiae</i> ) (3.7 kD) (Orc6l) alternative variant fSep08, mRNA.
<a href="#">ORF19</a>	<a href="#">ORF19.bSep08</a>	<a href="#">367328</a>	7048	647	6	215	protein family homolog (ORF19) alternative variant bSep08, mRNA.
<a href="#">ORF19</a>	<a href="#">ORF19.cSep08</a>	<a href="#">367328</a>	1083	431	2	143	putative protein of vertebrate origin (ORF19) alternative variant cSep08, mRNA.
<a href="#">ORF19</a>	<a href="#">ORF19.dSep08</a>	<a href="#">367328</a>	1183	474	2	52	putative protein (ORF19) alternative variant dSep08, mRNA.
<a href="#">ORF19</a>	<a href="#">ORF19.fSep08</a>	<a href="#">367328</a>	4239	366	2	40	putative protein human specific (4.4 kD) (ORF19) alternative variant fSep08, mRNA.
<a href="#">Ormdl2</a>	<a href="#">Ormdl2.aSep08</a>	<a href="#">288783</a>	3134	986	4	162	ORM1-like 2 ( <i>S. cerevisiae</i> ) (Ormdl2) alternative variant aSep08, mRNA.
<a href="#">Ormdl2</a>	<a href="#">Ormdl2.bSep08</a>	<a href="#">288783</a>	1411	475	3	158	ORM1-like 2 ( <i>S. cerevisiae</i> ) (Ormdl2) alternative variant bSep08, mRNA.
<a href="#">Ormdl2</a>	<a href="#">Ormdl2.dSep08</a>	<a href="#">288783</a>	1763	552	3	109	ORM1-like 2 ( <i>S. cerevisiae</i> ) (12.3 kD) (Ormdl2) alternative variant dSep08, mRNA.
<a href="#">Ormdl3</a>	<a href="#">Ormdl3.aSep08</a>	<a href="#">360618</a>	4881	1232	3	347	ORM1-like 3 ( <i>S. cerevisiae</i> ) (Ormdl3) alternative variant aSep08, mRNA.
<a href="#">Ormdl3</a>	<a href="#">Ormdl3.cSep08</a>	<a href="#">360618</a>	5923	1974	3	180	ORM1-like 3 ( <i>S. cerevisiae</i> ) (Ormdl3) alternative variant cSep08, mRNA.

<a href="#">Os9</a>	<a href="#">Os9.aSep08</a>	<a href="#">362891</a>	27017	3080	15	684	amplified in osteosarcoma (Os9) alternative variant aSep08, mRNA.
<a href="#">Os9</a>	<a href="#">Os9.bSep08</a>	<a href="#">362891</a>	3818	1436	6	281	amplified in osteosarcoma (Os9) alternative variant bSep08, mRNA.
<a href="#">Os9</a>	<a href="#">Os9.cSep08</a>	<a href="#">362891</a>	1891	780	5	260	amplified in osteosarcoma (Os9) alternative variant cSep08, mRNA.
<a href="#">Osbp</a>	<a href="#">Osbp.bSep08</a>	<a href="#">365410</a>	2137	687	5	167	oxysterol binding protein (Osbp) alternative variant bSep08, mRNA.
<a href="#">Osbp</a>	<a href="#">Osbp.cSep08</a>	<a href="#">365410</a>	1607	756	4	146	oxysterol binding protein (Osbp) alternative variant cSep08, mRNA.
<a href="#">Osbp</a>	<a href="#">Osbp.dSep08</a>	<a href="#">365410</a>	1657	718	3	126	oxysterol binding protein (15.0 kD) (Osbp) alternative variant dSep08, mRNA.
<a href="#">Osbp2</a>	<a href="#">Osbp2.bSep08</a>	<a href="#">305475</a>	3779	660	6	199	oxysterol binding protein 2 (Osbp2) alternative variant bSep08, mRNA.
<a href="#">Osbp2</a>	<a href="#">Osbp2.dSep08</a>	<a href="#">305475</a>	446	215	2	58	oxysterol binding protein 2 (Osbp2) alternative variant dSep08, mRNA.
<a href="#">Osbpl2</a>	<a href="#">Osbpl2.bSep08</a>	<a href="#">296461</a>	9517	777	7	258	oxysterol-binding protein-like protein 2 (Osbpl2) alternative variant bSep08, mRNA.
<a href="#">Osbpl2</a>	<a href="#">Osbpl2.cSep08</a>	<a href="#">296461</a>	3736	681	3	125	oxysterol binding protein-like 2 (Osbpl2) alternative variant cSep08, mRNA.
<a href="#">Osbpl2</a>	<a href="#">Osbpl2.dSep08</a>	<a href="#">296461</a>	1713	762	2	100	putative mitochondrial protein (11.2 kD) (Osbpl2) alternative variant dSep08, mRNA.
<a href="#">Osbpl2</a>	<a href="#">Osbpl2.eSep08</a>	<a href="#">296461</a>	12764	400	4	46	oxysterol-binding protein-like protein 2 (Osbpl2) alternative variant eSep08, mRNA.
<a href="#">Osbpl2</a>	<a href="#">Osbpl2.gSep08</a>	<a href="#">296461</a>	895	760	2	61	putative protein (7.5 kD) (Osbpl2) alternative variant gSep08, mRNA.
<a href="#">Osbpl3</a>	<a href="#">Osbpl3.aSep08</a>	<a href="#">362360</a>	15077	676	5	225	oxysterol binding protein-like 3 (Osbpl3) alternative variant aSep08, mRNA.
<a href="#">Osbpl5</a>	<a href="#">Osbpl5.bSep08</a>	<a href="#">361686</a>	2653	720	4	195	oxysterol binding protein-like 5 (Osbpl5) alternative variant bSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.bSep08</a>	<a href="#">311129</a>	11908	499	5	165	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant bSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.bSep08</a>	<a href="#">679513</a>	11908	499	5	165	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant bSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.cSep08</a>	<a href="#">311129</a>	121551	546	3	127	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant cSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.cSep08</a>	<a href="#">679513</a>	121551	546	3	127	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant cSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.dSep08</a>	<a href="#">311129</a>	2751	430	3	83	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant dSep08, mRNA.
<a href="#">Osbpl6</a>	<a href="#">Osbpl6.dSep08</a>	<a href="#">679513</a>	2751	430	3	83	oxysterol binding protein-like 6 and hypothetical protein LOC679513 (Osbpl6) alternative variant dSep08, mRNA.
<a href="#">Osbpl7</a>	<a href="#">Osbpl7.bSep08</a>	<a href="#">303497</a>	3638	680	7	174	oxysterol binding protein-like 7 (Osbpl7) alternative variant bSep08, mRNA.
<a href="#">Osbpl9</a>	<a href="#">Osbpl9.aSep08</a>	<a href="#">298369</a>	36427	2312	17	572	oxysterol binding protein-like 9 (Osbpl9) alternative variant aSep08, mRNA.



<a href="#">Osbp19</a>	<a href="#">Osbp19.bSep08</a>	<a href="#">298369</a>	2943	1161	4	108	oxysterol binding protein-like 9 (13.1 kD) (Osbp19) alternative variant bSep08, mRNA.
<a href="#">Osbp19</a>	<a href="#">Osbp19.cSep08</a>	<a href="#">298369</a>	2637	470	4	75	oxysterol binding protein-like 9 (Osbp19) alternative variant cSep08, mRNA.
<a href="#">Osbp10</a>	<a href="#">Osbp10.aSep08</a>	<a href="#">316039</a>	155756	1722	4	477	OSBP related protein 10 (Osbp10) alternative variant aSep08, mRNA.
<a href="#">Osbp10</a>	<a href="#">Osbp10.bSep08</a>	<a href="#">316039</a>	140442	966	6	300	oxysterol binding protein-like 10 CRA b (Osbp10) alternative variant bSep08, mRNA.
<a href="#">Osbp10</a>	<a href="#">Osbp10.cSep08</a>	<a href="#">316039</a>	93331	705	5	235	oxysterol-binding protein-like protein 10 (Osbp10) alternative variant cSep08, mRNA.
<a href="#">Osbp10</a>	<a href="#">Osbp10.dSep08</a>	<a href="#">316039</a>	14576	770	3	212	oxysterol-binding protein-like protein 10 (Osbp10) alternative variant dSep08, mRNA.
<a href="#">Osbp10</a>	<a href="#">Osbp10.eSep08</a>	<a href="#">316039</a>	28271	749	3	138	oxysterol-binding protein-like protein 10 (Osbp10) alternative variant eSep08, mRNA.
<a href="#">Oscar</a>	<a href="#">Oscar.aSep08</a>	<a href="#">292537</a>	671	521		146	osteoclast associated receptor (Oscar) mRNA.
<a href="#">Osgep</a>	<a href="#">Osgep.bSep08</a>	<a href="#">290028</a>	6500	1036	4	186	O-sialoglycoprotein endopeptidase (19.8 kD) (Osgep) alternative variant bSep08, mRNA.
<a href="#">Osgep</a>	<a href="#">Osgep.cSep08</a>	<a href="#">290028</a>	5816	689	3	122	O-sialoglycoprotein endopeptidase (Osgep) alternative variant cSep08, mRNA.
<a href="#">Osgepl1</a>	<a href="#">Osgepl1.bSep08</a>	<a href="#">314548</a>	10961	1084	1	196	O-sialoglycoprotein endopeptidase-like 1 (21.4 kD) (Osgepl1) alternative variant bSep08, complete mRNA.
<a href="#">Osgin1</a>	<a href="#">Osgin1.bSep08</a>	<a href="#">171493</a>	7254	901	4	245	oxidative stress induced growth inhibitor 1 (Osgin1) alternative variant bSep08, mRNA.
<a href="#">Osgin1</a>	<a href="#">Osgin1.cSep08</a>	<a href="#">171493</a>	23474	757	5	105	oxidative stress induced growth inhibitor 1 (Osgin1) alternative variant cSep08, mRNA.
<a href="#">Osgin1</a>	<a href="#">Osgin1.dSep08</a>	<a href="#">171493</a>	24484	750	5	86	oxidative stress induced growth inhibitor 1 (Osgin1) alternative variant dSep08, mRNA.
<a href="#">Osgin2</a>	<a href="#">Osgin2.aSep08</a>	<a href="#">313085</a>	18853	1797	3	417	oxidative stress induced growth inhibitor family member 2 (Osgin2) alternative variant aSep08, mRNA.
<a href="#">Osgin2</a>	<a href="#">Osgin2.bSep08</a>	<a href="#">313085</a>	4594	1795	2	347	oxidative stress induced growth inhibitor family member 2 (Osgin2) alternative variant bSep08, mRNA.
<a href="#">Osgin2</a>	<a href="#">Osgin2.cSep08</a>	<a href="#">313085</a>	10051	426	1	141	oxidative stress induced growth inhibitor family member 2 (Osgin2) alternative variant cSep08, mRNA.
<a href="#">Osr2</a>	<a href="#">Osr2.aSep08</a>	<a href="#">315039</a>	3573	1924	2	312	odd-skipped related 2 (Drosophila) (35.5 kD) (Osr2) alternative variant aSep08, mRNA.
<a href="#">Ostb</a>	<a href="#">Ostb.aSep08</a>	<a href="#">300790</a>	8114	711		104	organic solute transporter beta (11.3 kD) (Ostb) mRNA.
<a href="#">Ostf1</a>	<a href="#">Ostf1.bSep08</a>	<a href="#">259275</a>	8772	1502	5	137	osteoclast stimulating factor 1 (14.9 kD) (Ostf1) alternative variant bSep08, mRNA.
<a href="#">Ostf1</a>	<a href="#">Ostf1.cSep08</a>	<a href="#">259275</a>	3095	694	2	36	osteoclast stimulating factor 1 (3.8 kD) (Ostf1) alternative variant cSep08, mRNA.
<a href="#">Otc</a>	<a href="#">Otc.bSep08</a>	<a href="#">25611</a>	9819	398	1	132	ornithine transcarbamylase (Otc) alternative variant bSep08, mRNA.
<a href="#">Otoa</a>	<a href="#">Otoa.aSep08</a>	<a href="#">499260</a>	15746	782		260	otoancorin (Otoa) mRNA.
<a href="#">Otof</a>	<a href="#">Otof.aSep08</a>	<a href="#">84573</a>	43478	385		92	otoferlin (Otof) mRNA.
<a href="#">Otos</a>	<a href="#">Otos.aSep08</a>	<a href="#">246044</a>	1155	338		89	otospiralin (10.1 kD) (Otos) mRNA.

<a href="#">Otp</a>	<a href="#">Otp.aSep08</a>	<a href="#">294640</a>	1878	624	2	159	orthopedia homolog (Drosophila) (Otp) alternative variant aSep08, mRNA.
<a href="#">Otp</a>	<a href="#">Otp.bSep08</a>	<a href="#">294640</a>	6033	477	2	158	orthopedia homolog (Drosophila) (Otp) alternative variant bSep08, mRNA.
<a href="#">OTU.0</a>	<a href="#">OTU.0.aSep08</a>		11111	2233		278	ovarian tumour, otubain (OTU.0) mRNA.
<a href="#">Otub2</a>	<a href="#">Otub2.bSep08</a>	<a href="#">314405</a>	18488	2695	5	219	OTU domain, ubiquitin aldehyde binding 2 (25.0 kD) (Otub2) alternative variant bSep08, mRNA.
<a href="#">Otub2</a>	<a href="#">Otub2.cSep08</a>	<a href="#">314405</a>	1186	854	1	75	OTU domain, ubiquitin aldehyde binding 2 (Otub2) alternative variant cSep08, mRNA.
<a href="#">Otud3</a>	<a href="#">Otud3.aSep08</a>	<a href="#">500572</a>	19457	616	3	204	ovarian tumour, otubain (Otud3) alternative variant aSep08, mRNA.
<a href="#">Otud3</a>	<a href="#">Otud3.bSep08</a>	<a href="#">500572</a>	19426	472	2	128	CRA a like (Otud3) alternative variant bSep08, mRNA.
<a href="#">Otud3</a>	<a href="#">Otud3.cSep08</a>	<a href="#">500572</a>	19446	430	2	109	putative protein of eukaryotic origin (Otud3) alternative variant cSep08, mRNA.
<a href="#">Otud3</a>	<a href="#">Otud3.dSep08</a>	<a href="#">500572</a>	14217	516	2	89	putative protein of eukaryotic origin (Otud3) alternative variant dSep08, mRNA.
<a href="#">Otud4</a>	<a href="#">Otud4.aSep08</a>	<a href="#">307774</a>	35158	1776		515	protein hin-1 (Otud4) alternative variant aSep08, mRNA.
<a href="#">Otud4</a>	<a href="#">Otud4.bSep08</a>	<a href="#">307774</a>	8901	772		138	protein hin-1 (Otud4) alternative variant bSep08, mRNA.
<a href="#">Otud5</a>	<a href="#">Otud5.bSep08</a>	<a href="#">363452</a>	25261	1110	5	364	ovarian tumour, otubain (Otud5) alternative variant bSep08, mRNA.
<a href="#">Otud5</a>	<a href="#">Otud5.dSep08</a>	<a href="#">363452</a>	1029	750	2	67	putative protein (Otud5) alternative variant dSep08, mRNA.
<a href="#">Otud6b</a>	<a href="#">Otud6b.bSep08</a>	<a href="#">297911</a>	11109	844	6	108	CRA c (Otud6b) alternative variant bSep08, mRNA.
<a href="#">Otud7a</a>	<a href="#">Otud7a.aSep08</a>	<a href="#">309252</a>	75920	683		150	putative protein of vertebrate origin (Otud7a) mRNA.
<a href="#">Oxa1l</a>	<a href="#">Oxa1l.aSep08</a>	<a href="#">691393</a>	7628	1531	10	433	oxidase assembly 1-like (48.0 kD) (Oxa1l) alternative variant aSep08, complete mRNA.
<a href="#">Oxa1l</a>	<a href="#">Oxa1l.bSep08</a>	<a href="#">691393</a>	6222	741	6	247	oxidase assembly 1-like (Oxa1l) alternative variant bSep08, mRNA.
<a href="#">Oxa1l</a>	<a href="#">Oxa1l.cSep08</a>	<a href="#">691393</a>	1219	852	3	156	oxidase assembly 1-like (17.9 kD) (Oxa1l) alternative variant cSep08, mRNA.
<a href="#">Oxa1l</a>	<a href="#">Oxa1l.dSep08</a>	<a href="#">691393</a>	2852	935	3	146	oxidase assembly 1-like (15.2 kD) (Oxa1l) alternative variant dSep08, complete mRNA.
<a href="#">Oxa1l</a>	<a href="#">Oxa1l.eSep08</a>	<a href="#">691393</a>	930	736	2	111	oxidase assembly 1-like (12.8 kD) (Oxa1l) alternative variant eSep08, mRNA.
<a href="#">Oxct1</a>	<a href="#">Oxct1.aSep08</a>	<a href="#">690163</a>	94860	1501		500	3-oxoacid CoA transferase 1 (Oxct1) mRNA.
<a href="#">Oxnad1</a>	<a href="#">Oxnad1.bSep08</a>	<a href="#">306270</a>	25868	977	7	241	putative protein of ancient origin (Oxnad1) alternative variant bSep08, mRNA.
<a href="#">Oxnad1</a>	<a href="#">Oxnad1.cSep08</a>	<a href="#">306270</a>	11121	1057	7	229	putative mitochondrial protein of ancient origin (25.6 kD) (Oxnad1) alternative variant cSep08, complete mRNA.
<a href="#">Oxnad1</a>	<a href="#">Oxnad1.dSep08</a>	<a href="#">306270</a>	18521	1447	4	220	ferric reductase, NAD binding and oxidoreductase FAD/NAD(P)-binding (Oxnad1) alternative variant dSep08, mRNA.
<a href="#">Oxnad1</a>	<a href="#">Oxnad1.eSep08</a>	<a href="#">306270</a>	24018	887	6	216	putative protein of ancient origin (Oxnad1) alternative variant eSep08, mRNA.
<a href="#">Oxnad1</a>	<a href="#">Oxnad1.fSep08</a>	<a href="#">306270</a>	23929	905	7	181	putative protein of ancient origin (Oxnad1) alternative variant fSep08, mRNA.

<a href="#">Oxnad1</a>	<a href="#">Oxnad1.gSep08</a>	<a href="#">306270</a>	8282	757	5	78	putative mitochondrial protein of mammalian origin (8.7 kD) (Oxnad1) alternative variant gSep08, mRNA.
<a href="#">Oxr1</a>	<a href="#">Oxr1.bSep08</a>	<a href="#">117520</a>	35253	743	7	228	oxidation resistance 1 (Oxr1) alternative variant bSep08, mRNA.
<a href="#">Oxr1</a>	<a href="#">Oxr1.cSep08</a>	<a href="#">117520</a>	37110	2519	6	216	oxidation resistance 1 (24.4 kD) (Oxr1) alternative variant cSep08, mRNA.
<a href="#">Oxr1</a>	<a href="#">Oxr1.dSep08</a>	<a href="#">117520</a>	36738	591	6	197	oxidation resistance 1 (Oxr1) alternative variant dSep08, mRNA.
<a href="#">Oxr1</a>	<a href="#">Oxr1.eSep08</a>	<a href="#">117520</a>	8809	790	5	189	oxidation resistance 1 (Oxr1) alternative variant eSep08, mRNA.
<a href="#">Oxr1</a>	<a href="#">Oxr1.fSep08</a>	<a href="#">117520</a>	7797	285	4	94	oxidation resistance 1 (Oxr1) alternative variant fSep08, mRNA.
<a href="#">Oxsm</a>	<a href="#">Oxsm.aSep08</a>	<a href="#">289934</a>	3074	2071	2	316	3-oxoacyl-ACP synthase, mitochondrial (Oxsm) alternative variant aSep08, mRNA.
<a href="#">Oxsr1</a>	<a href="#">Oxsr1.bSep08</a>	<a href="#">316064</a>	25552	757	1	211	oxidative-stress responsive 1 (Oxsr1) alternative variant bSep08, mRNA.
<a href="#">Oxt</a>	<a href="#">Oxt.aSep08</a>	<a href="#">25504</a>	829	522		134	oxytocin (Oxt) mRNA.
<a href="#">P2rx1</a>	<a href="#">P2rx1.bSep08</a>	<a href="#">25505</a>	4418	825	6	186	purinergic receptor P2X, ligand-gated ion channel, 1 (21.2 kD) (P2rx1) alternative variant bSep08, mRNA.
<a href="#">P2rx2</a>	<a href="#">P2rx2.aSep08</a>	<a href="#">114115</a>	3172	1783	2	547	purinergic receptor P2X, ligand-gated ion channel, 2 (P2rx2) alternative variant aSep08, mRNA.
<a href="#">P2rx2</a>	<a href="#">P2rx2.cSep08</a>	<a href="#">114115</a>	726	519	2	66	purinergic receptor P2X, ligand-gated ion channel, 2 (P2rx2) alternative variant cSep08, mRNA.
<a href="#">P2rx3</a>	<a href="#">P2rx3.bSep08</a>	<a href="#">81739</a>	25898	1928	6	172	purinergic receptor P2X, ligand-gated ion channel, 3 (P2rx3) alternative variant bSep08, mRNA.
<a href="#">P2rx3</a>	<a href="#">P2rx3.cSep08</a>	<a href="#">81739</a>	15226	594	2	165	purinergic receptor P2X, ligand-gated ion channel, 3 (P2rx3) alternative variant cSep08, mRNA.
<a href="#">P2rx4</a>	<a href="#">P2rx4.bSep08</a>	<a href="#">29659</a>	3033	761	5	226	purinergic receptor P2X, ligand-gated ion channel 4 (P2rx4) alternative variant bSep08, mRNA.
<a href="#">P2rx4</a>	<a href="#">P2rx4.cSep08</a>	<a href="#">29659</a>	8935	824	5	73	purinergic receptor P2X, ligand-gated ion channel 4 (P2rx4) alternative variant cSep08, mRNA.
<a href="#">P2rx6</a>	<a href="#">P2rx6.bSep08</a>	<a href="#">25041</a>	6378	691	6	230	purinergic receptor P2X, ligand-gated ion channel, 6 (P2rx6) alternative variant bSep08, mRNA.
<a href="#">P2rx7</a>	<a href="#">P2rx7.cSep08</a>	<a href="#">29665</a>	17981	579	3	187	purinergic receptor P2X, ligand-gated ion channel, 7 (P2rx7) alternative variant cSep08, mRNA.
<a href="#">P2ry1</a>	<a href="#">P2ry1.bSep08</a>	<a href="#">25265</a>	5584	1637	1	58	purinergic receptor P2Y, G-protein coupled 1 (6.6 kD) (P2ry1) alternative variant bSep08, mRNA.
<a href="#">P2ry2</a>	<a href="#">P2ry2.aSep08</a>	<a href="#">29597</a>	13635	1443		318	purinergic receptor P2Y, G-protein coupled 2 (P2ry2) mRNA.
<a href="#">P2ry6</a>	<a href="#">P2ry6.bSep08</a>	<a href="#">117264</a>	23520	429	3	17	pyrimidinergic receptor P2Y, G-protein coupled, 6 (P2ry6) alternative variant bSep08, mRNA.
<a href="#">P2ry6</a>	<a href="#">P2ry6.cSep08</a>	<a href="#">117264</a>	2972	363	3	40	pyrimidinergic receptor P2Y, G-protein coupled, 6 (P2ry6) alternative variant cSep08, mRNA.
<a href="#">P2ry10</a>	<a href="#">P2ry10.aSep08</a>	<a href="#">317219</a>	14113	2923		328	purinergic receptor P2Y, G-protein coupled 10 (37.3 kD) (P2ry10) mRNA.
<a href="#">P2ry13</a>	<a href="#">P2ry13.aSep08</a>	<a href="#">310444</a>	2445	1793	3	536	purinergic receptor P2Y, G-protein coupled, 13 (P2ry13) alternative variant aSep08, mRNA.

<a href="#">P4ha1</a>	<a href="#">P4ha1.bSep08</a>	<a href="#">64475</a>	10726	1866	6	136	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha 1 polypeptide (15.2 kD) (P4ha1) alternative variant bSep08, mRNA.
<a href="#">P4ha1</a>	<a href="#">P4ha1.cSep08</a>	<a href="#">64475</a>	2426	850	3	78	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha 1 polypeptide (8.6 kD) (P4ha1) alternative variant cSep08, mRNA.
<a href="#">P4ha2</a>	<a href="#">P4ha2.aSep08</a>	<a href="#">360526</a>	28825	2230	16	597	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide (P4ha2) alternative variant aSep08, mRNA.
<a href="#">P4ha2</a>	<a href="#">P4ha2.cSep08</a>	<a href="#">360526</a>	8707	752	5	250	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide (P4ha2) alternative variant cSep08, mRNA.
<a href="#">P4ha2</a>	<a href="#">P4ha2.eSep08</a>	<a href="#">360526</a>	2821	953	3	100	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide (10.9 kD) (P4ha2) alternative variant eSep08, mRNA.
<a href="#">P4ha3</a>	<a href="#">P4ha3.bSep08</a>	<a href="#">361612</a>	1897	829	2	112	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide III (P4ha3) alternative variant bSep08, mRNA.
<a href="#">P4ha3</a>	<a href="#">P4ha3.cSep08</a>	<a href="#">361612</a>	4398	351	3	108	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide III (P4ha3) alternative variant cSep08, mRNA.
<a href="#">P4hb</a>	<a href="#">P4hb.bSep08</a>	<a href="#">25506</a>	2603	771	6	230	prolyl 4-hydroxylase, beta polypeptide (P4hb) alternative variant bSep08, mRNA.
<a href="#">P4hb</a>	<a href="#">P4hb.cSep08</a>	<a href="#">25506</a>	823	714	2	104	prolyl 4-hydroxylase, beta polypeptide (11.4 kD) (P4hb) alternative variant cSep08, mRNA.
<a href="#">P15rs</a>	<a href="#">P15rs.aSep08</a>	<a href="#">291736</a>	44867	1290	6	364	cyclin-dependent kinase 2B-inhibitor-related protein (P15rs) alternative variant aSep08, mRNA.
<a href="#">P15rs</a>	<a href="#">P15rs.bSep08</a>	<a href="#">291736</a>	2697	496	4	155	cyclin-dependent kinase 2B-inhibitor-related protein (P15rs) alternative variant bSep08, mRNA.
<a href="#">P15rs</a>	<a href="#">P15rs.cSep08</a>	<a href="#">291736</a>	1649	1546	1	103	cyclin-dependent kinase 2B-inhibitor-related protein (P15rs) alternative variant cSep08, mRNA.
<a href="#">P22k15</a>	<a href="#">P22k15.bSep08</a>	<a href="#">296229</a>	2554	308	2	43	cystatin related protein 2 (P22k15) alternative variant bSep08, mRNA.
<a href="#">P76</a>	<a href="#">P76.bSep08</a>	<a href="#">246120</a>	2346	1724	2	115	mannose-6-phosphate protein p76 (12.9 kD) (P76) alternative variant bSep08, mRNA.
<a href="#">p450.0</a>	<a href="#">p450.0.aSep08</a>	<a href="#">690021</a>	7350	1378	11	449	cytochrome P450 family 4 subfamily a polypeptide (51.7 kD) (p450.0) alternative variant aSep08, mRNA.
<a href="#">p450.0</a>	<a href="#">p450.0.bSep08</a>	<a href="#">690021</a>	2925	774	4	104	putative mitochondrial protein (12.3 kD) (p450.0) alternative variant bSep08, mRNA.
<a href="#">p450.1</a>	<a href="#">p450.1.aSep08</a>		50280	1003	5	331	p450 2c2 (p450.1) alternative variant aSep08, mRNA.
<a href="#">p450.1</a>	<a href="#">p450.1.bSep08</a>		6007	780	4	152	cytochrome (17.4 kD) (p450.1) alternative variant bSep08, mRNA.
<a href="#">p450.1</a>	<a href="#">p450.1.cSep08</a>		4824	675	3	123	cytochrome P450f (13.9 kD) (p450.1) alternative variant cSep08, complete mRNA.
<a href="#">p450.1</a>	<a href="#">p450.1.dSep08</a>		1948	931	2	85	cytochrome (9.6 kD) (p450.1) alternative variant dSep08, mRNA.
<a href="#">p450.1</a>	<a href="#">p450.1.eSep08</a>		3464	419	2	83	cytochrome (p450.1) alternative variant eSep08, mRNA.

<a href="#">p450.2</a>	<a href="#">p450.2.aSep08</a>		6815	856	2	274	cytochrome P-450 IIC13 (p450.2) alternative variant aSep08, mRNA.
<a href="#">p450.2</a>	<a href="#">p450.2.bSep08</a>		6568	296		98	cytochrome P-450 IIC13 (p450.2) alternative variant bSep08, mRNA.
<a href="#">p450.3</a>	<a href="#">p450.3.aSep08</a>		47628	1802	6	505	cytochrome p450 family 2 subfamily c polypeptide (p450.3) alternative variant aSep08, complete mRNA.
<a href="#">p450.3</a>	<a href="#">p450.3.bSep08</a>		47459	1210	7	377	cytochrome p450 family 2 subfamily c polypeptide (p450.3) alternative variant bSep08, mRNA.
<a href="#">p450.3</a>	<a href="#">p450.3.cSep08</a>		2525	288	2	95	cytochrome P450f (p450.3) alternative variant cSep08, mRNA.
<a href="#">p450.3</a>	<a href="#">p450.3.dSep08</a>		16214	308	3	92	p450 2c2 (p450.3) alternative variant dSep08, mRNA.
<a href="#">p450.3</a>	<a href="#">p450.3.eSep08</a>		1166	261	2	61	putative protein (6.9 kD) (p450.3) alternative variant eSep08, complete mRNA.
<a href="#">Pa2g4</a>	<a href="#">Pa2g4.bSep08</a>	<a href="#">288778</a>	958	429	1	142	proliferation-associated 2G4 (Pa2g4) alternative variant bSep08, mRNA.
<a href="#">PAAD_DAPIN.0</a>	<a href="#">PAAD_DAPIN.0.aSep08</a>		1499	474	3	118	pyrin (PAAD_DAPIN.0) alternative variant aSep08, mRNA.
<a href="#">PAAD_DAPIN.0</a>	<a href="#">PAAD_DAPIN.0.bSep08</a>		643	433	1	83	putative protein of mammalian origin (PAAD_DAPIN.0) alternative variant bSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.bSep08</a>	<a href="#">171350</a>	9527	1942	11	499	poly binding protein cytoplasmic 1 like (Pabpc1) alternative variant bSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.cSep08</a>	<a href="#">171350</a>	8196	2306	6	486	poly binding protein cytoplasmic 1 like (Pabpc1) alternative variant cSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.dSep08</a>	<a href="#">171350</a>	5136	881	4	225	poly binding protein cytoplasmic 1 like (Pabpc1) alternative variant dSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.eSep08</a>	<a href="#">171350</a>	2767	874	2	122	poly binding protein cytoplasmic 1 like (Pabpc1) alternative variant eSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.fSep08</a>	<a href="#">171350</a>	2302	739	5	117	poly binding protein cytoplasmic 1 like (Pabpc1) alternative variant fSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.gSep08</a>	<a href="#">171350</a>	1854	593	5	88	poly binding protein cytoplasmic 1 like (9.4 kD) (Pabpc1) alternative variant gSep08, mRNA.
<a href="#">Pabpc1</a>	<a href="#">Pabpc1.hSep08</a>	<a href="#">171350</a>	1331	546	3	64	poly Binding Protein like (6.8 kD) (Pabpc1) alternative variant hSep08, mRNA.
<a href="#">Pabpc4</a>	<a href="#">Pabpc4.bSep08</a>	<a href="#">298510</a>	7733	1644	12	444	poly A binding protein, cytoplasmic 4 (Pabpc4) alternative variant bSep08, mRNA.
<a href="#">Pabpc4</a>	<a href="#">Pabpc4.cSep08</a>	<a href="#">298510</a>	3900	862	6	184	poly A binding protein, cytoplasmic 4 (Pabpc4) alternative variant cSep08, mRNA.
<a href="#">Pabpc4</a>	<a href="#">Pabpc4.dSep08</a>	<a href="#">298510</a>	3150	685	5	133	poly A binding protein, cytoplasmic 4 (Pabpc4) alternative variant dSep08, mRNA.
<a href="#">Pabpn1</a>	<a href="#">Pabpn1.aSep08</a>	<a href="#">116697</a>	4600	1776	7	302	poly(A) binding protein, nuclear 1 (Pabpn1) alternative variant aSep08, mRNA.
<a href="#">Pabpn1</a>	<a href="#">Pabpn1.bSep08</a>	<a href="#">116697</a>	4336	2238	6	236	poly(A) binding protein, nuclear 1 (Pabpn1) alternative variant bSep08, mRNA.
<a href="#">Pabpn1</a>	<a href="#">Pabpn1.cSep08</a>	<a href="#">116697</a>	3385	736	7	209	poly(A) binding protein, nuclear 1 (Pabpn1) alternative variant cSep08, mRNA.
<a href="#">Pabpn1</a>	<a href="#">Pabpn1.dSep08</a>	<a href="#">116697</a>	2697	774	6	201	poly(A) binding protein, nuclear 1 (Pabpn1) alternative variant dSep08, mRNA.

<a href="#">paby</a>	<a href="#">paby.aSep08</a>		1093	399		133	mediator of RNA polymerase II transcription homolog (paby) mRNA.
<a href="#">pachy</a>	<a href="#">pachy.aSep08</a>		6329	350		116	cytoskeleton associated protein 5 CRA a (pachy) mRNA.
<a href="#">Pacrg</a>	<a href="#">Pacrg.bSep08</a>	<a href="#">499021</a>	274968	845	4	139	park2 co-regulated (15.9 kD) (Pacrg) alternative variant bSep08, mRNA.
<a href="#">Pacrg</a>	<a href="#">Pacrg.cSep08</a>	<a href="#">499021</a>	74797	658	2	90	park2 co-regulated (10.0 kD) (Pacrg) alternative variant cSep08, mRNA.
<a href="#">Pacs-1.0</a>	<a href="#">Pacs-1.0.aSep08</a>		6515	559		186	phosphofurin acidic cluster sorting protein 2 (Pacs-1.0) mRNA.
<a href="#">Pacs-1.1</a>	<a href="#">Pacs-1.1.aSep08</a>		3049	1363		167	phosphofurin acidic cluster sorting protein 2 (Pacs-1.1) mRNA.
<a href="#">Pacsin2</a>	<a href="#">Pacsin2.bSep08</a>	<a href="#">124461</a>	83820	1229	7	314	protein kinase C and casein kinase substrate in neurons 2 (36.8 kD) (Pacsin2) alternative variant bSep08, mRNA.
<a href="#">Pacsin2</a>	<a href="#">Pacsin2.cSep08</a>	<a href="#">124461</a>	92212	1573	9	223	protein kinase C and casein kinase substrate in neurons 2 (25.5 kD) (Pacsin2) alternative variant cSep08, mRNA.
<a href="#">Pacsin2</a>	<a href="#">Pacsin2.dSep08</a>	<a href="#">124461</a>	79170	710	5	188	protein kinase C and casein kinase substrate in neurons 2 (Pacsin2) alternative variant dSep08, mRNA.
<a href="#">Pacsin2</a>	<a href="#">Pacsin2.eSep08</a>	<a href="#">124461</a>	4707	782	3	70	protein kinase C and casein kinase substrate in neurons 2 (Pacsin2) alternative variant eSep08, mRNA.
<a href="#">Pacsin2</a>	<a href="#">Pacsin2.fSep08</a>	<a href="#">124461</a>	4480	1145	3	151	protein kinase C and casein kinase substrate in neurons 2 (Pacsin2) alternative variant fSep08, mRNA.
<a href="#">Pacsin3</a>	<a href="#">Pacsin3.bSep08</a>	<a href="#">311187</a>	3058	848	4	282	protein kinase C and casein kinase substrate in neurons 3 (Pacsin3) alternative variant bSep08, mRNA.
<a href="#">Pacsin3</a>	<a href="#">Pacsin3.cSep08</a>	<a href="#">311187</a>	7539	1528	7	266	protein kinase C and casein kinase substrate in neurons 3 (30.9 kD) (Pacsin3) alternative variant cSep08, mRNA.
<a href="#">Pacsin3</a>	<a href="#">Pacsin3.dSep08</a>	<a href="#">311187</a>	7009	795	3	190	protein kinase C and casein kinase substrate in neurons 3 (22.4 kD) (Pacsin3) alternative variant dSep08, mRNA.
<a href="#">Pacsin3</a>	<a href="#">Pacsin3.eSep08</a>	<a href="#">311187</a>	4419	579	3	106	protein kinase C and casein kinase substrate in neurons 3 (Pacsin3) alternative variant eSep08, mRNA.
<a href="#">padar</a>	<a href="#">padar.bSep08</a>		857	533	2	107	putative mitochondrial protein (11.2 kD) (padar) alternative variant bSep08, mRNA.
<a href="#">Padi2</a>	<a href="#">Padi2.aSep08</a>	<a href="#">29511</a>	18865	1160		260	peptidyl arginine deiminase, type II (Padi2) mRNA.
<a href="#">PAD_N.0</a>	<a href="#">PAD_N.0.aSep08</a>		10796	428		114	peptidyl arginine deiminase type II (PAD_N.0) mRNA.
<a href="#">Paf1</a>	<a href="#">Paf1.bSep08</a>	<a href="#">361531</a>	778	677	2	98	paf1, RNA polymerase II associated factor, homolog (S. cerevisiae) (Paf1) alternative variant bSep08, mRNA.
<a href="#">Paf1</a>	<a href="#">Paf1.cSep08</a>	<a href="#">361531</a>	536	389	3	69	paf1, RNA polymerase II associated factor, homolog (S. cerevisiae) (Paf1) alternative variant cSep08, mRNA.
<a href="#">Paf1</a>	<a href="#">Paf1.dSep08</a>	<a href="#">361531</a>	930	836	2	39	paf1, RNA polymerase II associated factor, homolog (S. cerevisiae) (Paf1) alternative variant dSep08, mRNA.
<a href="#">Pafah1b1</a>	<a href="#">Pafah1b1.bSep08</a>	<a href="#">83572</a>	10863	3476	6	239	platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit 45kDa (27.1 kD) (Pafah1b1) alternative variant bSep08, mRNA.
<a href="#">Pafah1b2</a>	<a href="#">Pafah1b2.aSep08</a>	<a href="#">64189</a>	17698	1782	3	423	platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit (Pafah1b2) alternative variant aSep08, complete mRNA.

<a href="#">Pafah1b3</a>	<a href="#">Pafah1b3.cSep08</a>	<a href="#">114113</a>	2511	1375	5	103	platelet-activating factor acetylhydrolase, isoform 1b, alpha1 subunit (Pafah1b3) alternative variant cSep08, mRNA.
<a href="#">pafer</a>	<a href="#">pafer.aSep08</a>		1220	568		26	putative protein (2.9 kD) (pafer) mRNA.
<a href="#">paflo</a>	<a href="#">paflo.aSep08</a>		7116	251		83	CRA a like (paflo) mRNA.
<a href="#">paflu</a>	<a href="#">paflu.aSep08</a>		847	712		31	putative protein (paflu) mRNA.
<a href="#">pagar</a>	<a href="#">pagar.aSep08</a>		5418	532		176	polybromo 1 CRA c (pagar) mRNA.
<a href="#">Pah</a>	<a href="#">Pah.bSep08</a>	<a href="#">24616</a>	49476	815	1	217	phenylalanine hydroxylase (Pah) alternative variant bSep08, mRNA.
<a href="#">Paics</a>	<a href="#">Paics.bSep08</a>	<a href="#">140946</a>	11181	866	2	288	phosphoribosylaminoimidazole carboxylase, phosphoribosylaminoimidazole succinocarboxamide synthetase (Paics) alternative variant bSep08, mRNA.
<a href="#">Paics</a>	<a href="#">Paics.cSep08</a>	<a href="#">140946</a>	11174	793	2	264	phosphoribosylaminoimidazole carboxylase, phosphoribosylaminoimidazole succinocarboxamide synthetase (Paics) alternative variant cSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.aSep08</a>	<a href="#">365684</a>	28145	2829	11	460	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant aSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.aSep08</a>	<a href="#">689651</a>	28145	2829	11	460	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant aSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.bSep08</a>	<a href="#">365684</a>	18715	983	7	327	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant bSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.bSep08</a>	<a href="#">689651</a>	18715	983	7	327	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant bSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.cSep08</a>	<a href="#">365684</a>	7749	465	4	143	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant cSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.cSep08</a>	<a href="#">689651</a>	7749	465	4	143	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (Paip1) alternative variant cSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.dSep08</a>	<a href="#">365684</a>	6447	949	4	102	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (12.5 kD) (Paip1) alternative variant dSep08, mRNA.
<a href="#">Paip1</a>	<a href="#">Paip1.dSep08</a>	<a href="#">689651</a>	6447	949	4	102	polyadenylate binding protein-interacting protein 1 and hypothetical protein LOC689651 (12.5 kD) (Paip1) alternative variant dSep08, mRNA.
<a href="#">Paip2</a>	<a href="#">Paip2.aSep08</a>	<a href="#">361309</a>	17723	775	4	147	polyadenylate-binding protein interacting protein 2 (17.2 kD) (Paip2) alternative variant aSep08, mRNA.
<a href="#">Paip2</a>	<a href="#">Paip2.bSep08</a>	<a href="#">361309</a>	17856	836	4	124	polyadenylate-binding protein interacting protein 2 (14.7 kD) (Paip2) alternative variant bSep08, complete mRNA.
<a href="#">Paip2</a>	<a href="#">Paip2.dSep08</a>	<a href="#">361309</a>	18511	1418	3	124	polyadenylate-binding protein interacting protein 2 (14.7 kD) (Paip2) alternative variant dSep08, mRNA.
<a href="#">Paip2</a>	<a href="#">Paip2.eSep08</a>	<a href="#">361309</a>	17587	625	4	114	polyadenylate-binding protein interacting protein 2 (Paip2) alternative variant eSep08, mRNA.

<a href="#">Paip2</a>	<a href="#">Paip2.fSep08</a>	<a href="#">361309</a>	18200	1111	3	101	polyadenylate-binding protein interacting protein 2 (12.0 kD) (Paip2) alternative variant fSep08, mRNA.
<a href="#">Paip2</a>	<a href="#">Paip2.gSep08</a>	<a href="#">361309</a>	14817	525	3	101	polyadenylate-binding protein interacting protein 2 (Paip2) alternative variant gSep08, mRNA.
<a href="#">Paip2b</a>	<a href="#">Paip2b.aSep08</a>	<a href="#">312490</a>	27463	587	1	195	poly(A) binding protein interacting protein 2B (Paip2b) alternative variant aSep08, mRNA.
<a href="#">Paip2b</a>	<a href="#">Paip2b.bSep08</a>	<a href="#">312490</a>	29060	575	3	167	poly(A) binding protein interacting protein 2B (Paip2b) alternative variant bSep08, mRNA.
<a href="#">paja</a>	<a href="#">paja.aSep08</a>		823	369		30	putative protein (paja) mRNA.
<a href="#">pajey</a>	<a href="#">pajey.aSep08</a>		1115	500		51	putative protein (pajey) mRNA.
<a href="#">Pak1</a>	<a href="#">Pak1.bSep08</a>	<a href="#">29431</a>	4747	758	2	60	p21 (CDKN1A)-activated kinase 1 (6.8 kD) (Pak1) alternative variant bSep08, mRNA.
<a href="#">Pak1ip1</a>	<a href="#">Pak1ip1.bSep08</a>	<a href="#">361232</a>	7874	943	7	129	PAK1 interacting protein 1 (Pak1ip1) alternative variant bSep08, mRNA.
<a href="#">Pak2</a>	<a href="#">Pak2.cSep08</a>	<a href="#">29432</a>	6287	708	3		
<a href="#">Pak3</a>	<a href="#">Pak3.bSep08</a>	<a href="#">29433</a>	26785	383	3	106	putative protein human specific (Pak3) alternative variant bSep08, mRNA.
<a href="#">Pak3</a>	<a href="#">Pak3.dSep08</a>	<a href="#">29433</a>	26647	254	3	59	putative protein human specific (Pak3) alternative variant dSep08, mRNA.
<a href="#">pakee</a>	<a href="#">pakee.aSep08</a>		2881	292		82	uncharacterized protein like (pakee) mRNA.
<a href="#">pakler</a>	<a href="#">pakler.aSep08</a>	<a href="#">363309</a>	40614	2075	21	613	tubulin-specific chaperone d CRA c (pakler) alternative variant aSep08, mRNA.
<a href="#">Palb2</a>	<a href="#">Palb2.aSep08</a>	<a href="#">293452</a>	3164	863	1	181	partner and localizer of BRCA2 (19.3 kD) (Palb2) alternative variant aSep08, mRNA.
<a href="#">Palb2</a>	<a href="#">Palb2.bSep08</a>	<a href="#">293452</a>	3998	652	2	137	partner and localizer of BRCA2 (Palb2) alternative variant bSep08, mRNA.
<a href="#">Pald</a>	<a href="#">Pald.bSep08</a>	<a href="#">294508</a>	7444	872	2	104	paladin (12.5 kD) (Pald) alternative variant bSep08, mRNA.
<a href="#">Palm</a>	<a href="#">Palm.aSep08</a>	<a href="#">170673</a>	26559	2652	1	383	paralemmin (41.9 kD) (Palm) alternative variant aSep08, mRNA.
<a href="#">Palm</a>	<a href="#">Palm.bSep08</a>	<a href="#">170673</a>	25289	1250		339	paralemmin (37.0 kD) (Palm) alternative variant bSep08, mRNA.
<a href="#">paloy</a>	<a href="#">paloy.aSep08</a>		2112	613		87	putative protein (9.7 kD) (paloy) mRNA.
<a href="#">Pam</a>	<a href="#">Pam.bSep08</a>	<a href="#">25508</a>	22641	1375	6	233	peptidylglycine alpha-amidating monooxygenase (Pam) alternative variant bSep08, mRNA.
<a href="#">Pam</a>	<a href="#">Pam.cSep08</a>	<a href="#">25508</a>	22035	565	5	188	peptidylglycine alpha-amidating monooxygenase (Pam) alternative variant cSep08, mRNA.
<a href="#">Pam</a>	<a href="#">Pam.dSep08</a>	<a href="#">25508</a>	35700	479	5	159	peptidylglycine alpha-amidating monooxygenase (Pam) alternative variant dSep08, mRNA.
<a href="#">Pam</a>	<a href="#">Pam.eSep08</a>	<a href="#">25508</a>	18455	454	5	151	peptidylglycine alpha-amidating monooxygenase (Pam) alternative variant eSep08, mRNA.
<a href="#">Pam</a>	<a href="#">Pam.fSep08</a>	<a href="#">25508</a>	16075	682	4	144	peptidylglycine alpha-amidating monooxygenase (Pam) alternative variant fSep08, mRNA.
<a href="#">pamee</a>	<a href="#">pamee.aSep08</a>		978	555		119	serine threonine kinase 10 (13.2 kD) (pamee) mRNA.
<a href="#">pamer</a>	<a href="#">pamer.aSep08</a>		15627	538	4	179	acyl-Coenzyme A Oxidase 1 palmitoyl (pamer) alternative variant aSep08, mRNA.



<a href="#">Pan3</a>	<a href="#">Pan3.aSep08</a>	<a href="#">360760</a>	10641	2581	6	177	PAN3 polyA specific ribonuclease subunit homolog ( <i>S. cerevisiae</i> ) (Pan3) alternative variant aSep08, mRNA.
<a href="#">Pan3</a>	<a href="#">Pan3.bSep08</a>	<a href="#">360760</a>	5375	655	4	149	PAN3 polyA specific ribonuclease subunit homolog ( <i>S. cerevisiae</i> ) (17.0 kD) (Pan3) alternative variant bSep08, mRNA.
<a href="#">Pank1</a>	<a href="#">Pank1.aSep08</a>	<a href="#">294088</a>	50172	937	5	283	pantothenate kinase 1 (Pank1) alternative variant aSep08, mRNA.
<a href="#">Pank2</a>	<a href="#">Pank2.aSep08</a>	<a href="#">296167</a>	21695	1320	6	262	pantothenate kinase 2 (Pank2) alternative variant aSep08, mRNA.
<a href="#">Pank2</a>	<a href="#">Pank2.cSep08</a>	<a href="#">296167</a>	13371	511	3	106	CRA b (Pank2) alternative variant cSep08, mRNA.
<a href="#">Pank2</a>	<a href="#">Pank2.dSep08</a>	<a href="#">296167</a>	9706	551	3	106	pantothenate kinase 2 (Pank2) alternative variant dSep08, mRNA.
<a href="#">Pank2</a>	<a href="#">Pank2.eSep08</a>	<a href="#">296167</a>	3353	827	2	40	pantothenate kinase 2 (4.5 kD) (Pank2) alternative variant eSep08, mRNA.
<a href="#">Pank4</a>	<a href="#">Pank4.bSep08</a>	<a href="#">171053</a>	9507	1649	14	257	pantothenate kinase 4 (28.4 kD) (Pank4) alternative variant bSep08, mRNA.
<a href="#">Pank4</a>	<a href="#">Pank4.cSep08</a>	<a href="#">171053</a>	8439	1426	13	236	pantothenate kinase 4 (Pank4) alternative variant cSep08, mRNA.
<a href="#">Pank4</a>	<a href="#">Pank4.dSep08</a>	<a href="#">171053</a>	3388	745	7	115	pantothenate kinase 4 (13.3 kD) (Pank4) alternative variant dSep08, mRNA.
<a href="#">Pank4</a>	<a href="#">Pank4.eSep08</a>	<a href="#">171053</a>	3737	754	4	104	pantothenate kinase 4 (Pank4) alternative variant eSep08, mRNA.
<a href="#">panoy</a>	<a href="#">panoy.aSep08</a>		15007	682		227	tensin (panoy) mRNA.
<a href="#">Panx2</a>	<a href="#">Panx2.bSep08</a>	<a href="#">362979</a>	8399	1893	3	369	pannexin 2 (Panx2) alternative variant bSep08, mRNA.
<a href="#">Panx2</a>	<a href="#">Panx2.cSep08</a>	<a href="#">362979</a>	3723	331	2	79	pannexin 2 (Panx2) alternative variant cSep08, mRNA.
<a href="#">Panx3</a>	<a href="#">Panx3.bSep08</a>	<a href="#">315567</a>	4787	834	1	119	pannexin 3 (Panx3) alternative variant bSep08, mRNA.
<a href="#">Pap2d</a>	<a href="#">Pap2d.bSep08</a>	<a href="#">310812</a>	4202	892	2	139	phosphatidic acid phosphatase type 2 (15.4 kD) (Pap2d) alternative variant bSep08, mRNA.
<a href="#">Pap2d</a>	<a href="#">Pap2d.cSep08</a>	<a href="#">310812</a>	13021	425	2	76	phosphatidic acid phosphatase type 2 (Pap2d) alternative variant cSep08, mRNA.
<a href="#">Papd1</a>	<a href="#">Papd1.bSep08</a>	<a href="#">307050</a>	9452	1011	5	332	putative protein of metazoan origin (Papd1) alternative variant bSep08, mRNA.
<a href="#">Papd1</a>	<a href="#">Papd1.cSep08</a>	<a href="#">307050</a>	6686	828	4	246	putative mitochondrial protein of vertebrate origin (27.6 kD) (Papd1) alternative variant cSep08, mRNA.
<a href="#">Papd1</a>	<a href="#">Papd1.dSep08</a>	<a href="#">307050</a>	6615	699	4	232	putative protein of metazoan origin (Papd1) alternative variant dSep08, mRNA.
<a href="#">Papd1</a>	<a href="#">Papd1.eSep08</a>	<a href="#">307050</a>	12021	2806	4	200	PAP/25A-associated (22.4 kD) (Papd1) alternative variant eSep08, mRNA.
<a href="#">Papd1</a>	<a href="#">Papd1.fSep08</a>	<a href="#">307050</a>	6505	766	5	181	putative protein of bilateral origin (Papd1) alternative variant fSep08, mRNA.
<a href="#">Papd4</a>	<a href="#">Papd4.bSep08</a>	<a href="#">361878</a>	51942	2534	6	480	PAP/25A-associated (55.5 kD) (Papd4) alternative variant bSep08, mRNA.
<a href="#">Papd4</a>	<a href="#">Papd4.cSep08</a>	<a href="#">361878</a>	44723	1784	2	214	putative protein of ancient origin (Papd4) alternative variant cSep08, mRNA.
<a href="#">Papd4</a>	<a href="#">Papd4.dSep08</a>	<a href="#">361878</a>	3709	2490	2	32	putative protein of mammalian origin (3.9 kD) (Papd4) alternative variant dSep08, mRNA.

<a href="#">Papd5</a>	<a href="#">Papd5.aSep08</a>	<a href="#">307745</a>	50702	1623	10	540	DNA polymerase, beta-like region and PAP/25A-associated (Papd5) alternative variant aSep08, mRNA.
<a href="#">Papln</a>	<a href="#">Papln.bSep08</a>	<a href="#">314297</a>	4641	608	1	59	papilin (6.6 kD) (Papln) alternative variant bSep08, mRNA.
<a href="#">Papola</a>	<a href="#">Papola.bSep08</a>	<a href="#">314417</a>	25244	1369	10	317	poly (A) polymerase alpha (36.3 kD) (Papola) alternative variant bSep08, complete mRNA.
<a href="#">Papola</a>	<a href="#">Papola.cSep08</a>	<a href="#">314417</a>	10450	719	5	230	poly (A) polymerase alpha (Papola) alternative variant cSep08, mRNA.
<a href="#">Papola</a>	<a href="#">Papola.dSep08</a>	<a href="#">314417</a>	9675	683	7	227	poly (A) polymerase alpha (Papola) alternative variant dSep08, mRNA.
<a href="#">Papola</a>	<a href="#">Papola.eSep08</a>	<a href="#">314417</a>	13022	731	6	216	poly (A) polymerase alpha (Papola) alternative variant eSep08, mRNA.
<a href="#">Papolg</a>	<a href="#">Papolg.bSep08</a>	<a href="#">305586</a>	2237	393	2	60	poly(A) polymerase gamma (Papolg) alternative variant bSep08, mRNA.
<a href="#">papor</a>	<a href="#">papor.aSep08</a>		6593	449	3	149	probable E3 ubiquitin-protein ligase herc1 (papor) alternative variant aSep08, mRNA.
<a href="#">papor</a>	<a href="#">papor.bSep08</a>		634	132	1	43	guanine nucleotide exchange factor p532 like (papor) alternative variant bSep08, mRNA.
<a href="#">Papss1</a>	<a href="#">Papss1.bSep08</a>	<a href="#">295443</a>	19235	1019	1	237	3'-phosphoadenosine 5'-phosphosulfate synthase 1 (Papss1) alternative variant bSep08, mRNA.
<a href="#">Papss2</a>	<a href="#">Papss2.bSep08</a>	<a href="#">294103</a>	84984	3509	7	614	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (Papss2) alternative variant bSep08, complete mRNA.
<a href="#">Papss2</a>	<a href="#">Papss2.cSep08</a>	<a href="#">294103</a>	69657	783	1	241	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (Papss2) alternative variant cSep08, mRNA.
<a href="#">Paqr4</a>	<a href="#">Paqr4.bSep08</a>	<a href="#">302967</a>	1627	415	2	76	progesterin and adipoQ receptor family member IV (Paqr4) alternative variant bSep08, mRNA.
<a href="#">Paqr5</a>	<a href="#">Paqr5.bSep08</a>	<a href="#">315741</a>	37168	712	3	155	progesterin and adipoQ receptor family member V (Paqr5) alternative variant bSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.aSep08</a>	<a href="#">681021</a>	3263	1034	7	344	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant aSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.bSep08</a>	<a href="#">681021</a>	3199	713	7	237	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant bSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.cSep08</a>	<a href="#">681021</a>	2414	1237	5	169	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant cSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.dSep08</a>	<a href="#">681021</a>	1603	610	5	144	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant dSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.eSep08</a>	<a href="#">681021</a>	2427	316	4	87	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant eSep08, mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.fSep08</a>	<a href="#">681021</a>	3907	1176	5	75	progesterin and adipoQ receptor family member VI (8.2 kD) (Paqr6) alternative variant fSep08, complete mRNA.
<a href="#">Paqr6</a>	<a href="#">Paqr6.gSep08</a>	<a href="#">681021</a>	1165	669	2	64	progesterin and adipoQ receptor family member VI (Paqr6) alternative variant gSep08, mRNA.
<a href="#">Paqr8</a>	<a href="#">Paqr8.aSep08</a>	<a href="#">316275</a>	50537	5027	3	354	progesterin adipoQ receptor family member VIII (40.6 kD) (Paqr8) alternative variant aSep08, mRNA.
<a href="#">Paqr8</a>	<a href="#">Paqr8.cSep08</a>	<a href="#">316275</a>	43587	317	3	97	putative protein (Paqr8) alternative variant cSep08, mRNA.
<a href="#">Paqr8</a>	<a href="#">Paqr8.dSep08</a>	<a href="#">316275</a>	2623	393	3	50	putative protein (Paqr8) alternative variant dSep08, mRNA.
<a href="#">Paqr8</a>	<a href="#">Paqr8.eSep08</a>	<a href="#">316275</a>	44618	649	3	24	putative protein (Paqr8) alternative variant eSep08, mRNA.

<a href="#">Paqr8</a>	<a href="#">Paqr8.fSep08</a>	<a href="#">316275</a>	2087	296	2	53	putative protein (Paqr8) alternative variant fSep08, mRNA.
<a href="#">Paralemmin.0</a>	<a href="#">Paralemmin.0.aSep08</a>		136929	1102	4	298	paralemmin 2 (Paralemmin.0) alternative variant aSep08, mRNA.
<a href="#">Paralemmin.0</a>	<a href="#">Paralemmin.0.bSep08</a>		72126	387	2	24	putative protein (Paralemmin.0) alternative variant bSep08, mRNA.
<a href="#">parby</a>	<a href="#">parby.aSep08</a>		551	280		30	putative protein (parby) mRNA.
<a href="#">Parc</a>	<a href="#">Parc.aSep08</a>	<a href="#">316228</a>	38798	1790	10	596	p53-associated parkin-like cytoplasmic protein (Parc) alternative variant aSep08, mRNA.
<a href="#">Parc</a>	<a href="#">Parc.bSep08</a>	<a href="#">316228</a>	9354	1813	12	562	p53-associated parkin-like cytoplasmic protein (Parc) alternative variant bSep08, mRNA.
<a href="#">Parc</a>	<a href="#">Parc.cSep08</a>	<a href="#">316228</a>	2601	1790	4	555	p53-associated parkin-like cytoplasmic protein (Parc) alternative variant cSep08, mRNA.
<a href="#">Parc</a>	<a href="#">Parc.dSep08</a>	<a href="#">316228</a>	3556	1681	5	372	p53-associated parkin-like cytoplasmic protein (41.1 kD) (Parc) alternative variant dSep08, mRNA.
<a href="#">Parc</a>	<a href="#">Parc.eSep08</a>	<a href="#">316228</a>	29006	1122	5	128	p53-associated parkin-like cytoplasmic protein (Parc) alternative variant eSep08, mRNA.
<a href="#">Parc</a>	<a href="#">Parc.fSep08</a>	<a href="#">316228</a>	537	463	2	111	p53-associated parkin-like cytoplasmic protein (Parc) alternative variant fSep08, mRNA.
<a href="#">parchy</a>	<a href="#">parchy.aSep08</a>		5946	233		21	putative protein (2.4 kD) (parchy) mRNA.
<a href="#">Pard3</a>	<a href="#">Pard3.bSep08</a>	<a href="#">81918</a>	147854	2222	4	219	partitioning-defective protein 3 (Pard3) alternative variant bSep08, mRNA.
<a href="#">Pard3</a>	<a href="#">Pard3.cSep08</a>	<a href="#">81918</a>	13225	579	2	158	partitioning-defective 3 homolog (17.7 kD) (Pard3) alternative variant cSep08, mRNA.
<a href="#">Pard3</a>	<a href="#">Pard3.dSep08</a>	<a href="#">81918</a>	19327	735	3	136	partitioning-defective 3 homolog (Pard3) alternative variant dSep08, mRNA.
<a href="#">Pard6a</a>	<a href="#">Pard6a.aSep08</a>	<a href="#">307799</a>	1779	1192	2	357	par-6 homolog alpha (Pard6a) alternative variant aSep08, mRNA.
<a href="#">Pard6g</a>	<a href="#">Pard6g.bSep08</a>	<a href="#">307237</a>	15481	1718	3	119	putative protein (12.9 kD) (Pard6g) alternative variant bSep08, mRNA.
<a href="#">pardar</a>	<a href="#">pardar.aSep08</a>		6279	1018		46	putative protein (pardar) mRNA.
<a href="#">parfer</a>	<a href="#">parfer.aSep08</a>		56558	546		133	putative protein of metazoan origin (parfer) mRNA.
<a href="#">parflo</a>	<a href="#">parflo.aSep08</a>		13669	422		113	putative protein (parflo) mRNA.
<a href="#">parflu</a>	<a href="#">parflu.aSep08</a>		55082	410		23	putative protein (parflu) mRNA.
<a href="#">Parg</a>	<a href="#">Parg.bSep08</a>	<a href="#">83507</a>	61059	699	8	232	poly (ADP-ribose) glycohydrolase (Parg) alternative variant bSep08, mRNA.
<a href="#">Parg</a>	<a href="#">Parg.cSep08</a>	<a href="#">83507</a>	13172	715	3	137	poly (ADP-ribose) glycohydrolase (Parg) alternative variant cSep08, mRNA.
<a href="#">Parg</a>	<a href="#">Parg.dSep08</a>	<a href="#">83507</a>	21732	733	5	69	poly (ADP-ribose) glycohydrolase (8.1 kD) (Parg) alternative variant dSep08, mRNA.
<a href="#">pargar</a>	<a href="#">pargar.aSep08</a>		3872	460		53	CRA a like (6.1 kD) (pargar) mRNA.
<a href="#">parja</a>	<a href="#">parja.aSep08</a>		1022	488	2	118	putative protein (parja) alternative variant aSep08, mRNA.
<a href="#">parjey</a>	<a href="#">parjey.aSep08</a>		7843	432		143	extracellular matrix protein Fras1 (parjey) mRNA.
<a href="#">Park2</a>	<a href="#">Park2.aSep08</a>	<a href="#">56816</a>	27620	742		88	parkin (9.9 kD) (Park2) mRNA.
<a href="#">Park7</a>	<a href="#">Park7.aSep08</a>	<a href="#">117287</a>	18467	1391	7	214	parkinson disease (autosomal recessive, early onset) 7 (22.5 kD) (Park7) alternative variant aSep08, complete mRNA.

<a href="#">Park7</a>	<a href="#">Park7.bSep08</a>	<a href="#">117287</a>	11684	797	7	189	parkinson disease (autosomal recessive, early onset) 7 (20.0 kD) (Park7) alternative variant bSep08, complete mRNA.
<a href="#">Park7</a>	<a href="#">Park7.cSep08</a>	<a href="#">117287</a>	11695	868	7	189	parkinson disease (autosomal recessive, early onset) 7 (20.0 kD) (Park7) alternative variant cSep08, complete mRNA.
<a href="#">Park7</a>	<a href="#">Park7.eSep08</a>	<a href="#">117287</a>	6340	731	5	175	parkinson disease (autosomal recessive, early onset) 7 (Park7) alternative variant eSep08, mRNA.
<a href="#">Park7</a>	<a href="#">Park7.fSep08</a>	<a href="#">117287</a>	15949	533	7	140	parkinson disease (autosomal recessive, early onset) 7 (Park7) alternative variant fSep08, mRNA.
<a href="#">Park7</a>	<a href="#">Park7.iSep08</a>	<a href="#">117287</a>	2017	638	2	64	parkinson disease (autosomal recessive, early onset) 7 (Park7) alternative variant iSep08, mRNA.
<a href="#">parkee</a>	<a href="#">parkee.aSep08</a>		23801	588		128	putative protein (parkee) mRNA.
<a href="#">parkler</a>	<a href="#">parkler.aSep08</a>		3198	199	2	63	gag protein like (parkler) alternative variant aSep08, mRNA.
<a href="#">Parl</a>	<a href="#">Parl.bSep08</a>	<a href="#">287979</a>	19516	783	6	249	presenilin associated, rhomboid-like (Parl) alternative variant bSep08, mRNA.
<a href="#">Parl</a>	<a href="#">Parl.cSep08</a>	<a href="#">287979</a>	9868	687	5	177	presenilin associated, rhomboid-like (Parl) alternative variant cSep08, mRNA.
<a href="#">Parl</a>	<a href="#">Parl.dSep08</a>	<a href="#">287979</a>	10508	598	4	171	presenilin associated, rhomboid-like (19.5 kD) (Parl) alternative variant dSep08, complete mRNA.
<a href="#">Parl</a>	<a href="#">Parl.eSep08</a>	<a href="#">287979</a>	5011	595	3	131	presenilin associated, rhomboid-like (Parl) alternative variant eSep08, mRNA.
<a href="#">parloy</a>	<a href="#">parloy.aSep08</a>		7904	1082		106	apoptosis-inducing factor like (11.1 kD) (parloy) mRNA.
<a href="#">parmee</a>	<a href="#">parmee.aSep08</a>		36952	386		29	putative protein (3.4 kD) (parmee) mRNA.
<a href="#">parmer</a>	<a href="#">parmer.aSep08</a>		4773	710	6	118	putative protein (12.8 kD) (parmer) alternative variant aSep08, mRNA.
<a href="#">parmer</a>	<a href="#">parmer.cSep08</a>		943	712	2	23	putative protein (2.7 kD) (parmer) alternative variant cSep08, mRNA.
<a href="#">Parn</a>	<a href="#">Parn.aSep08</a>	<a href="#">360464</a>	103883	1924		338	poly(A)-specific ribonuclease (deadenylation nuclease) (Parn) mRNA.
<a href="#">parnoy</a>	<a href="#">parnoy.aSep08</a>		2564	345		66	putative protein (parnoy) mRNA.
<a href="#">PARP.0</a>	<a href="#">PARP.0.aSep08</a>		10119	391		129	poly polymerase 4 (PARP.0) mRNA.
<a href="#">PARP.1</a>	<a href="#">PARP.1.aSep08</a>		6478	1282	2	330	poly polymerase family member 10 (PARP.1) alternative variant aSep08, mRNA.
<a href="#">PARP.1</a>	<a href="#">PARP.1.bSep08</a>		1035	698	1	54	poly polymerase family member 10 (PARP.1) alternative variant bSep08, mRNA.
<a href="#">Parp1</a>	<a href="#">Parp1.bSep08</a>	<a href="#">25591</a>	6619	721	4	177	poly (ADP-ribose) polymerase family, member 1 (Parp1) alternative variant bSep08, mRNA.
<a href="#">Parp1</a>	<a href="#">Parp1.cSep08</a>	<a href="#">25591</a>	2613	887	4	74	poly (ADP-ribose) polymerase family, member 1 (Parp1) alternative variant cSep08, mRNA.
<a href="#">Parp1</a>	<a href="#">Parp1.dSep08</a>	<a href="#">25591</a>	1953	1704	2	122	poly (ADP-ribose) polymerase family, member 1 (13.2 kD) (Parp1) alternative variant dSep08, mRNA.
<a href="#">Parp1</a>	<a href="#">Parp1.gSep08</a>	<a href="#">25591</a>	1141	413	2	40	poly (ADP-ribose) polymerase family, member 1 (4.2 kD) (Parp1) alternative variant gSep08, mRNA.
<a href="#">Parp2</a>	<a href="#">Parp2.bSep08</a>	<a href="#">290027</a>	1580	692	4	160	poly polymerase family member 2 (17.6 kD) (Parp2) alternative variant bSep08, mRNA.

<a href="#">Parp2</a>	<a href="#">Parp2.cSep08</a>	<a href="#">290027</a>	923	580	3	157	poly polymerase family member 2 (Parp2) alternative variant cSep08, mRNA.
<a href="#">Parp2</a>	<a href="#">Parp2.dSep08</a>	<a href="#">290027</a>	8870	3745	6	139	poly polymerase family member 2 (15.7 kD) (Parp2) alternative variant dSep08, mRNA.
<a href="#">Parp2</a>	<a href="#">Parp2.eSep08</a>	<a href="#">290027</a>	5428	374	5	113	poly polymerase family member 2 (Parp2) alternative variant eSep08, mRNA.
<a href="#">Parp3</a>	<a href="#">Parp3.bSep08</a>	<a href="#">300985</a>	1801	796	2	254	poly (ADP-ribose) polymerase family, member 3 (Parp3) alternative variant bSep08, mRNA.
<a href="#">Parp4</a>	<a href="#">Parp4.aSep08</a>	<a href="#">361046</a>	12684	1097	4	365	poly (ADP-ribose) polymerase family, member 4 (Parp4) alternative variant aSep08, mRNA.
<a href="#">Parp4</a>	<a href="#">Parp4.bSep08</a>	<a href="#">361046</a>	12818	1192	4	279	poly (ADP-ribose) polymerase family, member 4 (Parp4) alternative variant bSep08, mRNA.
<a href="#">Parp4</a>	<a href="#">Parp4.cSep08</a>	<a href="#">361046</a>	48591	613	6	157	poly (ADP-ribose) polymerase family, member 4 (Parp4) alternative variant cSep08, mRNA.
<a href="#">Parp6</a>	<a href="#">Parp6.bSep08</a>	<a href="#">300759</a>	16884	1580	16	479	poly (ADP-ribose) polymerase family, member 6 (Parp6) alternative variant bSep08, mRNA.
<a href="#">Parp6</a>	<a href="#">Parp6.cSep08</a>	<a href="#">300759</a>	14350	849	8	144	poly (ADP-ribose) polymerase family, member 6 (Parp6) alternative variant cSep08, mRNA.
<a href="#">Parp6</a>	<a href="#">Parp6.dSep08</a>	<a href="#">300759</a>	4810	548	5	134	poly (ADP-ribose) polymerase family, member 6 (Parp6) alternative variant dSep08, mRNA.
<a href="#">Parp6</a>	<a href="#">Parp6.eSep08</a>	<a href="#">300759</a>	1003	540	3	101	poly (ADP-ribose) polymerase family, member 6 (Parp6) alternative variant eSep08, mRNA.
<a href="#">Parp8</a>	<a href="#">Parp8.aSep08</a>	<a href="#">294762</a>	83039	2688	24	804	poly (ADP-ribose) polymerase family, member 8 (Parp8) alternative variant aSep08, mRNA.
<a href="#">Parp8</a>	<a href="#">Parp8.bSep08</a>	<a href="#">294762</a>	10469	397	3	131	poly (ADP-ribose) polymerase family, member 8 (Parp8) alternative variant bSep08, mRNA.
<a href="#">Parp8</a>	<a href="#">Parp8.dSep08</a>	<a href="#">294762</a>	1647	537	2	60	poly (ADP-ribose) polymerase family, member 8 (6.4 kD) (Parp8) alternative variant dSep08, mRNA.
<a href="#">Parp11</a>	<a href="#">Parp11.aSep08</a>	<a href="#">500323</a>	71530	1785	8	473	poly (ADP-ribose) polymerase family, member 11 (Parp11) alternative variant aSep08, mRNA.
<a href="#">Parp11</a>	<a href="#">Parp11.bSep08</a>	<a href="#">500323</a>	59046	1061	8	196	poly (ADP-ribose) polymerase family, member 11 (23.2 kD) (Parp11) alternative variant bSep08, mRNA.
<a href="#">Parp12</a>	<a href="#">Parp12.aSep08</a>	<a href="#">362343</a>	30033	809		269	poly (ADP-ribose) polymerase family, member 12 (Parp12) mRNA.
<a href="#">Parp14</a>	<a href="#">Parp14.aSep08</a>	<a href="#">303903</a>	6927	2579		284	poly (ADP-ribose) polymerase family, member 14 (Parp14) mRNA.
<a href="#">Parp16</a>	<a href="#">Parp16.bSep08</a>	<a href="#">315760</a>	9781	871	1	168	poly (ADP-ribose) polymerase family, member 16 (Parp16) alternative variant bSep08, mRNA.
<a href="#">parpor</a>	<a href="#">parpor.aSep08</a>		2406	278		92	guanine nucleotide exchange factor p532 (parpor) mRNA.
<a href="#">parsa</a>	<a href="#">parsa.aSep08</a>		3673	694		122	putative protein (13.8 kD) (parsa) mRNA.
<a href="#">parshee</a>	<a href="#">parshee.aSep08</a>		22334	599		199	neurobeachin CRA a (parshee) mRNA.
<a href="#">partu</a>	<a href="#">partu.aSep08</a>		9979	622		176	baculoviral IAP repeat-containing 6 (partu) mRNA.
<a href="#">Parva</a>	<a href="#">Parva.aSep08</a>	<a href="#">57341</a>	39806	1788	4	519	parvin, alpha (Parva) alternative variant aSep08, mRNA.
<a href="#">Parva</a>	<a href="#">Parva.cSep08</a>	<a href="#">57341</a>	1279	539	2	35	parvin, alpha (Parva) alternative variant cSep08, mRNA.
<a href="#">parvar</a>	<a href="#">parvar.aSep08</a>		542	441		65	putative protein (parvar) mRNA.
<a href="#">Parvb</a>	<a href="#">Parvb.bSep08</a>	<a href="#">362973</a>	13670	435	5	138	parvin, beta (Parvb) alternative variant bSep08, mRNA.

<a href="#">Parvb</a>	<a href="#">Parvb.cSep08</a>	<a href="#">362973</a>	9178	397	3	77	parvin, beta (Parvb) alternative variant cSep08, mRNA.
<a href="#">Parvb</a>	<a href="#">Parvb.dSep08</a>	<a href="#">362973</a>	62325	418	5	39	parvin, beta (Parvb) alternative variant dSep08, mRNA.
<a href="#">Parvg</a>	<a href="#">Parvg.bSep08</a>	<a href="#">689069</a>	8506	464	2	75	parvin, gamma (Parvg) alternative variant bSep08, mRNA.
<a href="#">parwey</a>	<a href="#">parwey.aSep08</a>		3699	2095		137	putative protein of vertebrate origin (parwey) mRNA.
<a href="#">pasa</a>	<a href="#">pasa.aSep08</a>		38284	851	1	125	putative protein (pasa) alternative variant aSep08, mRNA.
<a href="#">pasa</a>	<a href="#">pasa.bSep08</a>		38310	1783	1	44	putative protein (pasa) alternative variant bSep08, mRNA.
<a href="#">pashee</a>	<a href="#">pashee.aSep08</a>		7186	2241		179	neurobeachin (pashee) mRNA.
<a href="#">Pask</a>	<a href="#">Pask.bSep08</a>	<a href="#">301617</a>	3837	535	1	140	pask protein (Pask) alternative variant bSep08, mRNA.
<a href="#">Patatin.0</a>	<a href="#">Patatin.0.aSep08</a>		5479	733		244	patatin (Patatin.0) mRNA.
<a href="#">Pat1</a>	<a href="#">Pat1.bSep08</a>	<a href="#">361736</a>	19582	3067	12	500	protein associated with topoisomerase II homolog 1 (yeast) (Pat1) alternative variant bSep08, mRNA.
<a href="#">Pat1</a>	<a href="#">Pat1.cSep08</a>	<a href="#">361736</a>	5752	2961	3	116	protein associated with topoisomerase II homolog 1 (yeast) (Pat1) alternative variant cSep08, mRNA.
<a href="#">Pat1</a>	<a href="#">Pat1.eSep08</a>	<a href="#">361736</a>	7966	579	6	61	protein associated with topoisomerase II homolog 1 (yeast) (7.0 kD) (Pat1) alternative variant eSep08, mRNA.
<a href="#">patu</a>	<a href="#">patu.aSep08</a>		5623	1368		286	neurexin 1 CRA c (patu) alternative variant aSep08, mRNA.
<a href="#">pavar</a>	<a href="#">pavar.aSep08</a>		17355	394		130	protein tyrosine phosphatase receptor type U CRA a (pavar) mRNA.
<a href="#">pawby</a>	<a href="#">pawby.aSep08</a>		6463	601		200	taf1 RNA polymerase II TATA box binding protein - associated factor 250kDa CRA a like (pawby) mRNA.
<a href="#">pawchy</a>	<a href="#">pawchy.aSep08</a>		6573	430		43	putative protein (4.9 kD) (pawchy) mRNA.
<a href="#">pawdar</a>	<a href="#">pawdar.aSep08</a>		1465	425		32	putative protein (3.7 kD) (pawdar) mRNA.
<a href="#">pawey</a>	<a href="#">pawey.aSep08</a>		2675	728	1	45	putative protein (pawey) alternative variant aSep08, mRNA.
<a href="#">pawey</a>	<a href="#">pawey.bSep08</a>		2148	785	1	14	putative protein (1.4 kD) (pawey) alternative variant bSep08, mRNA.
<a href="#">pawfer</a>	<a href="#">pawfer.aSep08</a>		37526	362		120	putative protein of mammalian origin (pawfer) mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.bSep08</a>		3886	1225	3	85	putative cytoplasmic protein (9.8 kD) (pawflo) alternative variant bSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.cSep08</a>		2937	993	2	71	putative protein (7.9 kD) (pawflo) alternative variant cSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.dSep08</a>		2595	907	2	101	putative protein (pawflo) alternative variant dSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.eSep08</a>		2142	793	2	118	putative nuclear protein (12.5 kD) (pawflo) alternative variant eSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.fSep08</a>		3986	730	3	85	putative cytoplasmic protein (9.8 kD) (pawflo) alternative variant fSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.gSep08</a>		2249	644	2	111	putative protein (pawflo) alternative variant gSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.jSep08</a>		1312	555	2	49	putative protein (5.0 kD) (pawflo) alternative variant jSep08, mRNA.
<a href="#">pawflo</a>	<a href="#">pawflo.kSep08</a>		3682	467	3	40	putative protein (pawflo) alternative variant kSep08, mRNA.
<a href="#">pawflu</a>	<a href="#">pawflu.aSep08</a>		2544	938	1	47	putative protein (5.3 kD) (pawflu) alternative variant aSep08, mRNA.

<a href="#">pawflu</a>	<a href="#">pawflu.bSep08</a>		6316	686	2	79	putative protein (pawflu) alternative variant bSep08, mRNA.
<a href="#">pawgar</a>	<a href="#">pawgar.aSep08</a>		18864	539		69	putative protein (7.5 kD) (pawgar) mRNA.
<a href="#">pawja</a>	<a href="#">pawja.aSep08</a>		4934	715		74	putative protein (8.6 kD) (pawja) mRNA.
<a href="#">pawjey</a>	<a href="#">pawjey.aSep08</a>		7321	1780		14	putative protein (1.7 kD) (pawjey) mRNA.
<a href="#">pawkler</a>	<a href="#">pawkler.aSep08</a>		8560	703	1	234	carboxylesterase (pawkler) alternative variant aSep08, mRNA.
<a href="#">pawkler</a>	<a href="#">pawkler.bSep08</a>		8240	703	1	201	carboxylesterase (pawkler) alternative variant bSep08, mRNA.
<a href="#">pawloy</a>	<a href="#">pawloy.aSep08</a>		512	344		90	putative protein (pawloy) mRNA.
<a href="#">pawmee</a>	<a href="#">pawmee.aSep08</a>		10919	437		78	putative protein (9.0 kD) (pawmee) mRNA.
<a href="#">pawmer</a>	<a href="#">pawmer.aSep08</a>		1538	920		68	putative protein (7.4 kD) (pawmer) mRNA.
<a href="#">pawnoy</a>	<a href="#">pawnoy.aSep08</a>		8416	528		175	ubiquitin specific peptidase 37 (pawnoy) mRNA.
<a href="#">pawpor</a>	<a href="#">pawpor.aSep08</a>		4087	541		179	guanine nucleotide exchange factor p532 (pawpor) mRNA.
<a href="#">pawsa</a>	<a href="#">pawsa.aSep08</a>		6132	480		94	putative protein (pawsa) mRNA.
<a href="#">pawshee</a>	<a href="#">pawshee.aSep08</a>		20272	421		140	neurobeachin (pawshee) mRNA.
<a href="#">pawtu</a>	<a href="#">pawtu.aSep08</a>		6314	425		141	baculoviral IAP repeat-containing 6 (pawtu) mRNA.
<a href="#">pawvar</a>	<a href="#">pawvar.aSep08</a>		36624	314		104	putative protein (pawvar) mRNA.
<a href="#">pawwey</a>	<a href="#">pawwey.aSep08</a>		2235	283		93	putative protein, with a coiled coil domain, of mammalian origin (pawwey) mRNA.
<a href="#">Pax3</a>	<a href="#">Pax3.aSep08</a>	<a href="#">114502</a>	32750	1730	6	230	paired box gene 3 (Pax3) alternative variant aSep08, mRNA.
<a href="#">Pax4</a>	<a href="#">Pax4.bSep08</a>	<a href="#">83630</a>	3965	462	1	139	paired box gene 4 (Pax4) alternative variant bSep08, mRNA.
<a href="#">Pax7</a>	<a href="#">Pax7.aSep08</a>	<a href="#">500574</a>	55984	314		104	paired box gene 7 (Pax7) mRNA.
<a href="#">Pax8</a>	<a href="#">Pax8.bSep08</a>	<a href="#">81819</a>	2009	689	1	78	paired box gene 8 (Pax8) alternative variant bSep08, mRNA.
<a href="#">Paxip1</a>	<a href="#">Paxip1.bSep08</a>	<a href="#">311944</a>	9047	717	7	239	PAX interacting (with transcription-activation domain) protein 1 (Paxip1) alternative variant bSep08, mRNA.
<a href="#">Paxip1</a>	<a href="#">Paxip1.cSep08</a>	<a href="#">311944</a>	9656	902	5	205	PAX interacting (with transcription-activation domain) protein 1 (Paxip1) alternative variant cSep08, mRNA.
<a href="#">PB1.0</a>	<a href="#">PB1.0.aSep08</a>		17645	1795	3	540	neighbor of Brca1 gene 1 (PB1.0) alternative variant aSep08, mRNA.
<a href="#">PB1.0</a>	<a href="#">PB1.0.bSep08</a>		3161	706		234	neighbor of Brca1 gene 1 CRA a (PB1.0) alternative variant bSep08, mRNA.
<a href="#">PB1.0</a>	<a href="#">PB1.0.cSep08</a>		13107	753	2	191	neighbor of Brca1 gene 1 (PB1.0) alternative variant cSep08, mRNA.
<a href="#">PB1.0</a>	<a href="#">PB1.0.dSep08</a>		13488	700	2	185	neighbor of Brca1 gene 1 (PB1.0) alternative variant dSep08, mRNA.
<a href="#">PBD.0</a>	<a href="#">PBD.0.aSep08</a>		5090	391		130	CRA a (PBD.0) mRNA.
<a href="#">Pbld</a>	<a href="#">Pbld.bSep08</a>	<a href="#">171564</a>	13556	1206	6	278	MAWD binding protein like (30.5 kD) (Pbld) alternative variant bSep08, complete mRNA.
<a href="#">Pbld</a>	<a href="#">Pbld.cSep08</a>	<a href="#">171564</a>	6824	678	4	189	MAWD binding protein like (Pbld) alternative variant cSep08, mRNA.

<a href="#">Pbld</a>	<a href="#">Pbld.dSep08</a>	<a href="#">171564</a>	840	218	1	72	MAWD binding protein like (Pbld) alternative variant dSep08, mRNA.
<a href="#">Pbrm1</a>	<a href="#">Pbrm1.aSep08</a>	<a href="#">306254</a>	16170	3821	5	361	polybromo 1 (Pbrm1) alternative variant aSep08, mRNA.
<a href="#">Pbrm1</a>	<a href="#">Pbrm1.bSep08</a>	<a href="#">306254</a>	13066	1091	3	228	polybromo 1 (Pbrm1) alternative variant bSep08, mRNA.
<a href="#">Pbrm1</a>	<a href="#">Pbrm1.cSep08</a>	<a href="#">306254</a>	9680	417	2	139	polybromo 1 (Pbrm1) alternative variant cSep08, mRNA.
<a href="#">Pbsn</a>	<a href="#">Pbsn.aSep08</a>	<a href="#">54193</a>	38654	755	1	181	probasin (21.2 kD) (Pbsn) alternative variant aSep08, mRNA.
<a href="#">Pbx1</a>	<a href="#">Pbx1.bSep08</a>	<a href="#">304947</a>	275632	2154	7	347	pre-B-cell leukemia transcription factor 1 (38.4 kD) (Pbx1) alternative variant bSep08, mRNA.
<a href="#">Pbx1</a>	<a href="#">Pbx1.cSep08</a>	<a href="#">304947</a>	5378	1936	2	108	pre-B-cell leukemia transcription factor 1 (Pbx1) alternative variant cSep08, mRNA.
<a href="#">Pbx2</a>	<a href="#">Pbx2.bSep08</a>	<a href="#">406164</a>	432	348	1	115	pre-B-cell leukemia transcription factor 2 (Pbx2) alternative variant bSep08, mRNA.
<a href="#">Pbx3</a>	<a href="#">Pbx3.bSep08</a>	<a href="#">311876</a>	29134	1814	4	84	pre B-cell leukemia transcription factor 3 (Pbx3) alternative variant bSep08, mRNA.
<a href="#">Pbx3</a>	<a href="#">Pbx3.cSep08</a>	<a href="#">311876</a>	170000	1763	3		
<a href="#">Pbx4</a>	<a href="#">Pbx4.bSep08</a>	<a href="#">361131</a>	5171	746	1	169	pre-B-cell leukemia homeobox 4 (18.4 kD) (Pbx4) alternative variant bSep08, mRNA.
<a href="#">Pbxip1</a>	<a href="#">Pbxip1.bSep08</a>	<a href="#">310644</a>	6679	1130	1	234	pre-B-cell leukemia transcription factor interacting protein 1 (24.7 kD) (Pbxip1) alternative variant bSep08, mRNA.
<a href="#">Pcaf</a>	<a href="#">Pcaf.aSep08</a>	<a href="#">301164</a>	36744	1689	13	548	p300/CBP-associated factor (Pcaf) alternative variant aSep08, mRNA.
<a href="#">Pcaf</a>	<a href="#">Pcaf.bSep08</a>	<a href="#">301164</a>	12642	772	5	183	p300/CBP-associated factor (Pcaf) alternative variant bSep08, mRNA.
<a href="#">Pcbd1</a>	<a href="#">Pcbd1.bSep08</a>	<a href="#">29700</a>	1464	925	1	87	pterin 4 alpha carbinolamine dehydratase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1) 1 (Pcbd1) alternative variant bSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.aSep08</a>	<a href="#">363005</a>	13611	1110	11	324	poly(rC) binding protein 2 (Pcbp2) alternative variant aSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.bSep08</a>	<a href="#">363005</a>	14260	1018	9	218	poly(rC) binding protein 2 (Pcbp2) alternative variant bSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.cSep08</a>	<a href="#">363005</a>	14286	951	8	206	poly(rC) binding protein 2 (Pcbp2) alternative variant cSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.dSep08</a>	<a href="#">363005</a>	12081	1381	7	151	poly(rC) binding protein 2 (Pcbp2) alternative variant dSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.eSep08</a>	<a href="#">363005</a>	11517	778	6	142	poly(rC) binding protein 2 (Pcbp2) alternative variant eSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.fSep08</a>	<a href="#">363005</a>	3066	524	3	128	poly(rC) binding protein 2 (Pcbp2) alternative variant fSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.gSep08</a>	<a href="#">363005</a>	8597	2119	2	21	poly(rC) binding protein 2 (2.4 kD) (Pcbp2) alternative variant gSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.hSep08</a>	<a href="#">363005</a>	8590	1478	3	60	poly(rC) binding protein 2 (6.7 kD) (Pcbp2) alternative variant hSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.iSep08</a>	<a href="#">363005</a>	4094	730	3	60	poly(rC) binding protein 2 (6.7 kD) (Pcbp2) alternative variant iSep08, mRNA.
<a href="#">Pcbp2</a>	<a href="#">Pcbp2.jSep08</a>	<a href="#">363005</a>	3409	565	5	24	poly(rC) binding protein 2 (Pcbp2) alternative variant jSep08, mRNA.



<a href="#">Pcbp2</a>	<a href="#">Pcbp2.kSep08</a>	<a href="#">363005</a>	491	400	2	27	poly(rC) binding protein 2 (Pcbp2) alternative variant kSep08, mRNA.
<a href="#">Pcbp3</a>	<a href="#">Pcbp3.aSep08</a>	<a href="#">294336</a>	201107	2118	4	371	poly(rC) binding protein 3 (39.3 kD) (Pcbp3) alternative variant aSep08, complete mRNA.
<a href="#">Pcbp3</a>	<a href="#">Pcbp3.bSep08</a>	<a href="#">294336</a>	48734	1100	4	245	poly(rC) binding protein 3 (Pcbp3) alternative variant bSep08, mRNA.
<a href="#">Pcbp4</a>	<a href="#">Pcbp4.aSep08</a>	<a href="#">363133</a>	10166	1951	3	403	poly(rC) binding protein 4 (41.5 kD) (Pcbp4) alternative variant aSep08, complete mRNA.
<a href="#">Pcbp4</a>	<a href="#">Pcbp4.bSep08</a>	<a href="#">363133</a>	1524	1418	1	220	poly(rC) binding protein 4 (23.2 kD) (Pcbp4) alternative variant bSep08, mRNA.
<a href="#">Pcbp4</a>	<a href="#">Pcbp4.cSep08</a>	<a href="#">363133</a>	1647	564	1	187	poly(rC) binding protein 4 (Pcbp4) alternative variant cSep08, mRNA.
<a href="#">Pcbp4</a>	<a href="#">Pcbp4.dSep08</a>	<a href="#">363133</a>	1537	840	1	73	poly(rC) binding protein 4 (Pcbp4) alternative variant dSep08, mRNA.
<a href="#">Pcca</a>	<a href="#">Pcca.bSep08</a>	<a href="#">687008</a>	193510	1401	12	374	propionyl-coenzyme A carboxylase, alpha polypeptide (Pcca) alternative variant bSep08, mRNA.
<a href="#">Pcca</a>	<a href="#">Pcca.cSep08</a>	<a href="#">687008</a>	13140	599	4	68	propionyl-coenzyme A carboxylase, alpha polypeptide (Pcca) alternative variant cSep08, mRNA.
<a href="#">Pcca</a>	<a href="#">Pcca.dSep08</a>	<a href="#">687008</a>	17157	516	2	63	propionyl-coenzyme A carboxylase, alpha polypeptide (Pcca) alternative variant dSep08, mRNA.
<a href="#">Pcca</a>	<a href="#">Pcca.eSep08</a>	<a href="#">687008</a>	34677	1298	6	71	propionyl-coenzyme A carboxylase, alpha polypeptide (Pcca) alternative variant eSep08, complete mRNA.
<a href="#">Pcdh1</a>	<a href="#">Pcdh1.aSep08</a>	<a href="#">307481</a>	9195	1995	3	664	protocadherin 1 (Pcdh1) alternative variant aSep08, mRNA.
<a href="#">Pcdh8</a>	<a href="#">Pcdh8.bSep08</a>	<a href="#">64865</a>	3036	1293	2	303	protocadherin 8 (Pcdh8) alternative variant bSep08, mRNA.
<a href="#">Pcdh9</a>	<a href="#">Pcdh9.aSep08</a>	<a href="#">306091</a>	55526	364		69	protocadherin 9 (Pcdh9) mRNA.
<a href="#">Pcdh10</a>	<a href="#">Pcdh10.aSep08</a>	<a href="#">361943</a>	19448	1253	5	339	protocadherin 10 (Pcdh10) alternative variant aSep08, mRNA.
<a href="#">Pcdh10</a>	<a href="#">Pcdh10.bSep08</a>	<a href="#">361943</a>	28577	754	2	105	protocadherin 10 (Pcdh10) alternative variant bSep08, mRNA.
<a href="#">Pcdh12</a>	<a href="#">Pcdh12.aSep08</a>	<a href="#">116808</a>	9139	891	3	192	protocadherin 12 (Pcdh12) alternative variant aSep08, mRNA.
<a href="#">Pcdh15</a>	<a href="#">Pcdh15.aSep08</a>	<a href="#">690865</a>	19880	362		120	protocadherin 15 (Pcdh15) mRNA.
<a href="#">Pcdh17</a>	<a href="#">Pcdh17.bSep08</a>	<a href="#">306055</a>	30703	390	3	129	protocadherin 17 (Pcdh17) alternative variant bSep08, mRNA.
<a href="#">Pcdh18</a>	<a href="#">Pcdh18.aSep08</a>	<a href="#">295027</a>	13894	5098		1135	protocadherin 18 (125.6 kD) (Pcdh18) mRNA.
<a href="#">Pcdh19</a>	<a href="#">Pcdh19.aSep08</a>	<a href="#">317183</a>	98770	2595		633	protocadherin 19 (Pcdh19) mRNA.
<a href="#">Pcdh24</a>	<a href="#">Pcdh24.aSep08</a>	<a href="#">291002</a>	8406	1685	13	530	protocadherin 24 (Pcdh24) alternative variant aSep08, mRNA.
<a href="#">Pcdh24</a>	<a href="#">Pcdh24.bSep08</a>	<a href="#">291002</a>	6835	1092	5	364	protocadherin 24 (Pcdh24) alternative variant bSep08, mRNA.
<a href="#">Pcf11</a>	<a href="#">Pcf11.aSep08</a>	<a href="#">361605</a>	20200	5162	13	1356	cleavage and polyadenylation factor subunit homolog (S. cerevisiae) (Pcf11) alternative variant aSep08, mRNA.
<a href="#">Pcf11</a>	<a href="#">Pcf11.bSep08</a>	<a href="#">361605</a>	2824	968	3	112	cleavage and polyadenylation factor subunit homolog (S. cerevisiae) (13.3 kD) (Pcf11) alternative variant bSep08, mRNA.

<a href="#">Pcf11</a>	<a href="#">Pcf11.cSep08</a>	<a href="#">361605</a>	1317	504	2	72	cleavage and polyadenylation factor subunit homolog ( <i>S. cerevisiae</i> ) (Pcf11) alternative variant cSep08, mRNA.
<a href="#">Pcf11</a>	<a href="#">Pcf11.dSep08</a>	<a href="#">361605</a>	618	179	2	38	cleavage and polyadenylation factor subunit homolog ( <i>S. cerevisiae</i> ) (Pcf11) alternative variant dSep08, mRNA.
<a href="#">Pcgf1</a>	<a href="#">Pcgf1.aSep08</a>	<a href="#">312480</a>	2660	834	9	224	polycomb group ring finger 1 (Pcgf1) alternative variant aSep08, mRNA.
<a href="#">Pcgf1</a>	<a href="#">Pcgf1.bSep08</a>	<a href="#">312480</a>	2595	1076	9	176	polycomb group ring finger 1 (20.9 kD) (Pcgf1) alternative variant bSep08, mRNA.
<a href="#">Pcgf1</a>	<a href="#">Pcgf1.dSep08</a>	<a href="#">312480</a>	705	619	2	73	polycomb group ring finger 1 (8.2 kD) (Pcgf1) alternative variant dSep08, mRNA.
<a href="#">Pcgf3</a>	<a href="#">Pcgf3.aSep08</a>	<a href="#">305624</a>	55159	1779	11	241	ring finger 3 (28.0 kD) (Pcgf3) alternative variant aSep08, mRNA.
<a href="#">Pcgf3</a>	<a href="#">Pcgf3.bSep08</a>	<a href="#">305624</a>	39124	763	6	192	ring finger 3 (Pcgf3) alternative variant bSep08, mRNA.
<a href="#">Pcgf5</a>	<a href="#">Pcgf5.aSep08</a>	<a href="#">681178</a>	40197	824	1	174	polycomb group ring finger 5 (Pcgf5) alternative variant aSep08, mRNA.
<a href="#">Pcgf5</a>	<a href="#">Pcgf5.bSep08</a>	<a href="#">681178</a>	88426	760	2	155	polycomb group ring finger 5 (Pcgf5) alternative variant bSep08, mRNA.
<a href="#">Pcgf6</a>	<a href="#">Pcgf6.aSep08</a>	<a href="#">309457</a>	4969	838	4	278	polycomb group ring finger 6 (Pcgf6) alternative variant aSep08, mRNA.
<a href="#">Pcgf6</a>	<a href="#">Pcgf6.bSep08</a>	<a href="#">309457</a>	1901	780	1	200	polycomb group ring finger 6 (Pcgf6) alternative variant bSep08, mRNA.
<a href="#">Pcif1</a>	<a href="#">Pcif1.bSep08</a>	<a href="#">362269</a>	5452	2423	3	126	PDX1 C-terminal inhibiting factor 1 (14.2 kD) (Pcif1) alternative variant bSep08, mRNA.
<a href="#">Pck1</a>	<a href="#">Pck1.bSep08</a>	<a href="#">362282</a>	2952	2253	3	181	phosphoenolpyruvate carboxykinase 1, cytosolic (Pck1) alternative variant bSep08, mRNA.
<a href="#">Pck1</a>	<a href="#">Pck1.cSep08</a>	<a href="#">362282</a>	1308	764	3	137	phosphoenolpyruvate carboxykinase 1, cytosolic (15.6 kD) (Pck1) alternative variant cSep08, mRNA.
<a href="#">Pck2</a>	<a href="#">Pck2.bSep08</a>	<a href="#">361042</a>	2773	1346	5	349	phosphoenolpyruvate carboxykinase 2 (mitochondrial) (38.2 kD) (Pck2) alternative variant bSep08, mRNA.
<a href="#">Pcm1</a>	<a href="#">Pcm1.aSep08</a>	<a href="#">81740</a>	39776	2668	17	781	pericentriolar material 1 (Pcm1) alternative variant aSep08, mRNA.
<a href="#">Pcm1</a>	<a href="#">Pcm1.bSep08</a>	<a href="#">81740</a>	35706	1930	11	467	pericentriolar material 1 (Pcm1) alternative variant bSep08, mRNA.
<a href="#">Pcm1</a>	<a href="#">Pcm1.cSep08</a>	<a href="#">81740</a>	1163	413	2	122	pericentriolar material 1 (Pcm1) alternative variant cSep08, mRNA.
<a href="#">Pcm1</a>	<a href="#">Pcm1.dSep08</a>	<a href="#">81740</a>	2811	703	3	79	pericentriolar material 1 (Pcm1) alternative variant dSep08, mRNA.
<a href="#">Pcmt1</a>	<a href="#">Pcmt1.bSep08</a>	<a href="#">25604</a>	14259	963	6	170	protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 (Pcmt1) alternative variant bSep08, mRNA.
<a href="#">Pcmt1</a>	<a href="#">Pcmt1.cSep08</a>	<a href="#">25604</a>	6770	643	4	163	protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 (Pcmt1) alternative variant cSep08, mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.bSep08</a>	<a href="#">288165</a>	14520	2215	5	178	PEST proteolytic signal containing nuclear protein (19.0 kD) (Pcnp) alternative variant bSep08, complete mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.cSep08</a>	<a href="#">288165</a>	13100	743	5	162	PEST proteolytic signal containing nuclear protein (Pcnp) alternative variant cSep08, mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.dSep08</a>	<a href="#">288165</a>	13119	1014	5	111	PEST proteolytic signal containing nuclear protein (12.1 kD) (Pcnp) alternative variant dSep08, complete mRNA.

<a href="#">Pcnp</a>	<a href="#">Pcnp.eSep08</a>	<a href="#">288165</a>	8495	1897	4	110	PEST proteolytic signal containing nuclear protein (12.0 kD) (Pcnp) alternative variant eSep08, mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.fSep08</a>	<a href="#">288165</a>	13192	672	4	105	PEST proteolytic signal containing nuclear protein (Pcnp) alternative variant fSep08, mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.hSep08</a>	<a href="#">288165</a>	5729	1379	3	90	PEST proteolytic signal containing nuclear protein (10.1 kD) (Pcnp) alternative variant hSep08, mRNA.
<a href="#">Pcnp</a>	<a href="#">Pcnp.iSep08</a>	<a href="#">288165</a>	2065	869	2	55	PEST proteolytic signal containing nuclear protein (6.2 kD) (Pcnp) alternative variant iSep08, mRNA.
<a href="#">Pcnx</a>	<a href="#">Pcnx.aSep08</a>	<a href="#">314288</a>	14760	3352		764	pecanex homolog (Drosophila) (Pcnx) mRNA.
<a href="#">Pcnxl2</a>	<a href="#">Pcnxl2.aSep08</a>	<a href="#">307949</a>	19126	356		118	pecanex-like 2 (Drosophila) (Pcnxl2) mRNA.
<a href="#">Pcnxl3</a>	<a href="#">Pcnxl3.aSep08</a>	<a href="#">309167</a>	12418	3309	4	965	pecanex-like 3 (Drosophila) (Pcnxl3) alternative variant aSep08, mRNA.
<a href="#">Pcnxl3</a>	<a href="#">Pcnxl3.bSep08</a>	<a href="#">309167</a>	2329	1391	5	463	pecanex-like 3 (Drosophila) (Pcnxl3) alternative variant bSep08, mRNA.
<a href="#">Pcnxl3</a>	<a href="#">Pcnxl3.cSep08</a>	<a href="#">309167</a>	850	754	2	251	pecanex-like 3 (Drosophila) (Pcnxl3) alternative variant cSep08, mRNA.
<a href="#">Pcnxl3</a>	<a href="#">Pcnxl3.dSep08</a>	<a href="#">309167</a>	1154	745	3	148	pecanex-like 3 (Drosophila) (Pcnxl3) alternative variant dSep08, mRNA.
<a href="#">Pcolce</a>	<a href="#">Pcolce.bSep08</a>	<a href="#">29569</a>	4227	279	2	56	procollagen C-endopeptidase enhancer protein (Pcolce) alternative variant bSep08, mRNA.
<a href="#">Pcolce</a>	<a href="#">Pcolce.dSep08</a>	<a href="#">29569</a>	891	795	2	47	procollagen C-endopeptidase enhancer protein (Pcolce) alternative variant dSep08, mRNA.
<a href="#">Pcp2</a>	<a href="#">Pcp2.bSep08</a>	<a href="#">304195</a>	1749	452	1	129	purkinje cell protein 2 (L7) (Pcp2) alternative variant bSep08, mRNA.
<a href="#">Pcp2</a>	<a href="#">Pcp2.cSep08</a>	<a href="#">304195</a>	1733	460	1	78	purkinje cell protein 2 (L7) (Pcp2) alternative variant cSep08, mRNA.
<a href="#">Pcp4l1</a>	<a href="#">Pcp4l1.aSep08</a>	<a href="#">685448</a>	22721	618	1	85	purkinje cell protein 4-like 1 (9.9 kD) (Pcp4l1) alternative variant aSep08, mRNA.
<a href="#">Pcsk2</a>	<a href="#">Pcsk2.bSep08</a>	<a href="#">25121</a>	240841	1555	2	237	proprotein convertase subtilisin/kexin type 2 (Pcsk2) alternative variant bSep08, mRNA.
<a href="#">Pcsk4</a>	<a href="#">Pcsk4.bSep08</a>	<a href="#">171085</a>	858	677	2	105	proprotein convertase subtilisin/kexin type 4 (Pcsk4) alternative variant bSep08, mRNA.
<a href="#">Pcsk4</a>	<a href="#">Pcsk4.cSep08</a>	<a href="#">171085</a>	814	742	2	66	proprotein convertase subtilisin/kexin type 4 (Pcsk4) alternative variant cSep08, mRNA.
<a href="#">Pcsk5</a>	<a href="#">Pcsk5.aSep08</a>	<a href="#">116548</a>	99514	771		257	proprotein convertase subtilisin/kexin type 5 (Pcsk5) mRNA.
<a href="#">Pcsk6</a>	<a href="#">Pcsk6.bSep08</a>	<a href="#">25507</a>	125527	2332	18	776	proprotein convertase subtilisin/kexin type 6 (Pcsk6) alternative variant bSep08, mRNA.
<a href="#">Pcsk6</a>	<a href="#">Pcsk6.cSep08</a>	<a href="#">25507</a>	25085	2396	9	350	proprotein convertase subtilisin/kexin type 6 (Pcsk6) alternative variant cSep08, mRNA.
<a href="#">Pcsk6</a>	<a href="#">Pcsk6.dSep08</a>	<a href="#">25507</a>	56741	880	5	293	proprotein convertase subtilisin/kexin type 6 (Pcsk6) alternative variant dSep08, mRNA.
<a href="#">Pcsk6</a>	<a href="#">Pcsk6.eSep08</a>	<a href="#">25507</a>	25574	791	6	212	proprotein convertase subtilisin/kexin type 6 (Pcsk6) alternative variant eSep08, mRNA.
<a href="#">Pcsk7</a>	<a href="#">Pcsk7.bSep08</a>	<a href="#">29606</a>	917	409	3	136	proprotein convertase subtilisin/kexin type 7 (Pcsk7) alternative variant bSep08, mRNA.

<a href="#">Pctk1</a>	<a href="#">Pctk1.aSep08</a>	<a href="#">81741</a>	6304	1291	5	298	pctaire protein kinase 1 (31.8 kD) (Pctk1) alternative variant aSep08, mRNA.
<a href="#">Pctk1</a>	<a href="#">Pctk1.bSep08</a>	<a href="#">81741</a>	2014	815	6	271	pctaire protein kinase 1 (Pctk1) alternative variant bSep08, mRNA.
<a href="#">Pctk1</a>	<a href="#">Pctk1.cSep08</a>	<a href="#">81741</a>	1447	712	4	237	pctaire protein kinase 1 (Pctk1) alternative variant cSep08, mRNA.
<a href="#">Pctk1</a>	<a href="#">Pctk1.fSep08</a>	<a href="#">81741</a>	2294	742	5	56	pctaire protein kinase 1 (6.4 kD) (Pctk1) alternative variant fSep08, mRNA.
<a href="#">Pctk1</a>	<a href="#">Pctk1.gSep08</a>	<a href="#">81741</a>	1909	522	3	39	putative protein (Pctk1) alternative variant gSep08, mRNA.
<a href="#">Pctk2</a>	<a href="#">Pctk2.bSep08</a>	<a href="#">314743</a>	6402	707	6	178	PCTAIRE-motif protein kinase 2 (Pctk2) alternative variant bSep08, mRNA.
<a href="#">Pctk2</a>	<a href="#">Pctk2.dSep08</a>	<a href="#">314743</a>	5016	412	3	103	PCTAIRE-motif protein kinase 2 (Pctk2) alternative variant dSep08, mRNA.
<a href="#">Pctk3</a>	<a href="#">Pctk3.bSep08</a>	<a href="#">289019</a>	2307	400	4	98	PCTAIRE-motif protein kinase 3 (Pctk3) alternative variant bSep08, mRNA.
<a href="#">Pcx</a>	<a href="#">Pcx.aSep08</a>	<a href="#">25104</a>	99912	4036	21	1179	pyruvate carboxylase (129.9 kD) (Pcx) alternative variant aSep08, mRNA.
<a href="#">Pcx</a>	<a href="#">Pcx.cSep08</a>	<a href="#">25104</a>	1636	858	4	286	pyruvate carboxylase (Pcx) alternative variant cSep08, mRNA.
<a href="#">Pcx</a>	<a href="#">Pcx.dSep08</a>	<a href="#">25104</a>	4667	388	3	129	pyruvate carboxylase (Pcx) alternative variant dSep08, mRNA.
<a href="#">Pcyox1l</a>	<a href="#">Pcyox1l.bSep08</a>	<a href="#">307396</a>	8369	621	1	80	prenylcysteine oxidase 1 like (Pcyox1l) alternative variant bSep08, mRNA.
<a href="#">Pcyt1a</a>	<a href="#">Pcyt1a.bSep08</a>	<a href="#">140544</a>	31741	524	5	162	phosphate cytidyltransferase 1, choline, alpha isoform (Pcyt1a) alternative variant bSep08, mRNA.
<a href="#">Pcyt1a</a>	<a href="#">Pcyt1a.cSep08</a>	<a href="#">140544</a>	3361	356	2	103	phosphate cytidyltransferase 1, choline, alpha isoform (Pcyt1a) alternative variant cSep08, mRNA.
<a href="#">Pcyt1a</a>	<a href="#">Pcyt1a.fSep08</a>	<a href="#">140544</a>	1291	385	2	45	phosphate cytidyltransferase 1, choline, alpha isoform (4.9 kD) (Pcyt1a) alternative variant fSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.bSep08</a>	<a href="#">89841</a>	6681	1158	13	385	phosphate cytidyltransferase 2 ethanolamine CRA b (Pcyt2) alternative variant bSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.cSep08</a>	<a href="#">89841</a>	2483	807	8	236	phosphate cytidyltransferase 2 ethanolamine (Pcyt2) alternative variant cSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.dSep08</a>	<a href="#">89841</a>	4531	771	7	233	phosphate cytidyltransferase 2 ethanolamine (25.8 kD) (Pcyt2) alternative variant dSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.eSep08</a>	<a href="#">89841</a>	4904	617	7	180	phosphate cytidyltransferase 2 ethanolamine (Pcyt2) alternative variant eSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.fSep08</a>	<a href="#">89841</a>	1469	910	5	121	phosphate cytidyltransferase 2 ethanolamine (13.8 kD) (Pcyt2) alternative variant fSep08, mRNA.
<a href="#">Pcyt2</a>	<a href="#">Pcyt2.gSep08</a>	<a href="#">89841</a>	1488	627	6	103	phosphate cytidyltransferase 2 ethanolamine CRA c (Pcyt2) alternative variant gSep08, mRNA.
<a href="#">Pdcd2</a>	<a href="#">Pdcd2.aSep08</a>	<a href="#">58934</a>	5760	1403		359	programmed cell death 2 (Pdcd2) mRNA.
<a href="#">Pdcd2l</a>	<a href="#">Pdcd2l.bSep08</a>	<a href="#">689637</a>	10708	1781	6	222	programmed cell death 2-like (Pdcd2l) alternative variant bSep08, mRNA.
<a href="#">Pdcd2l</a>	<a href="#">Pdcd2l.cSep08</a>	<a href="#">689637</a>	4501	396	2	131	programmed cell death 2-like (Pdcd2l) alternative variant cSep08, mRNA.

<a href="#">Pdcd4</a>	<a href="#">Pdcd4.bSep08</a>	<a href="#">64031</a>	3907	522	5	110	programmed cell death 4 (Pdcd4) alternative variant bSep08, mRNA.
<a href="#">Pdcd5</a>	<a href="#">Pdcd5.cSep08</a>	<a href="#">292814</a>	4143	711	4	101	programmed cell death 5 (Pdcd5) alternative variant cSep08, mRNA.
<a href="#">Pdcd5</a>	<a href="#">Pdcd5.dSep08</a>	<a href="#">292814</a>	5278	584	6	84	programmed cell death 5 (9.4 kD) (Pdcd5) alternative variant dSep08, complete mRNA.
<a href="#">Pdcd6</a>	<a href="#">Pdcd6.bSep08</a>	<a href="#">308061</a>	15248	1106	3	189	programmed cell death 6 (21.7 kD) (Pdcd6) alternative variant bSep08, mRNA.
<a href="#">Pdcd6</a>	<a href="#">Pdcd6.cSep08</a>	<a href="#">308061</a>	14846	648	2	183	programmed cell death 6 (Pdcd6) alternative variant cSep08, mRNA.
<a href="#">Pdcd6</a>	<a href="#">Pdcd6.dSep08</a>	<a href="#">308061</a>	15022	778	3	155	programmed cell death 6 (17.6 kD) (Pdcd6) alternative variant dSep08, complete mRNA.
<a href="#">Pdcd6ip</a>	<a href="#">Pdcd6ip.bSep08</a>	<a href="#">501083</a>	1806	633			
<a href="#">Pdcd7</a>	<a href="#">Pdcd7.aSep08</a>	<a href="#">363082</a>	15079	2271		454	programmed cell death protein 7 (Pdcd7) mRNA.
<a href="#">Pdcd10</a>	<a href="#">Pdcd10.aSep08</a>	<a href="#">494345</a>	43120	1907	2	133	programmed cell death 10 (15.2 kD) (Pdcd10) alternative variant aSep08, complete mRNA.
<a href="#">Pdcd11</a>	<a href="#">Pdcd11.bSep08</a>	<a href="#">309458</a>	6355	760	4	188	programmed cell death 11 (Pdcd11) alternative variant bSep08, mRNA.
<a href="#">Pdcd11</a>	<a href="#">Pdcd11.cSep08</a>	<a href="#">309458</a>	6575	727	6	159	programmed cell death 11 (Pdcd11) alternative variant cSep08, mRNA.
<a href="#">Pdcd11</a>	<a href="#">Pdcd11.eSep08</a>	<a href="#">309458</a>	633	474	2	94	programmed cell death 11 (Pdcd11) alternative variant eSep08, mRNA.
<a href="#">Pdcl</a>	<a href="#">Pdcl.bSep08</a>	<a href="#">64013</a>	11449	558	4	119	phosducin-like (13.7 kD) (Pdcl) alternative variant bSep08, mRNA.
<a href="#">Pdcl2</a>	<a href="#">Pdcl2.aSep08</a>	<a href="#">498352</a>	14889	749		238	phosducin-like 2 (Pdcl2) mRNA.
<a href="#">Pdcd1</a>	<a href="#">Pdcd1.bSep08</a>	<a href="#">309110</a>	655	531	1	86	putative protein (9.3 kD) (Pdcd1) alternative variant bSep08, mRNA.
<a href="#">Pde1a</a>	<a href="#">Pde1a.bSep08</a>	<a href="#">81529</a>	21020	772	5	191	phosphodiesterase 1A, calmodulin-dependent (Pde1a) alternative variant bSep08, mRNA.
<a href="#">Pde1a</a>	<a href="#">Pde1a.cSep08</a>	<a href="#">81529</a>	134125	874	5	174	phosphodiesterase 1A, calmodulin-dependent (Pde1a) alternative variant cSep08, mRNA.
<a href="#">Pde1a</a>	<a href="#">Pde1a.dSep08</a>	<a href="#">81529</a>	7828	593	4	130	phosphodiesterase 1A, calmodulin-dependent (Pde1a) alternative variant dSep08, mRNA.
<a href="#">Pde2a</a>	<a href="#">Pde2a.bSep08</a>	<a href="#">81743</a>	33834	326	4	108	phosphodiesterase 2A, cGMP-stimulated (Pde2a) alternative variant bSep08, mRNA.
<a href="#">Pde2a</a>	<a href="#">Pde2a.cSep08</a>	<a href="#">81743</a>	63997	451	4	86	phosphodiesterase 2A, cGMP-stimulated (Pde2a) alternative variant cSep08, mRNA.
<a href="#">Pde2a</a>	<a href="#">Pde2a.dSep08</a>	<a href="#">81743</a>	2259	675	4	84	phosphodiesterase 2A, cGMP-stimulated (9.9 kD) (Pde2a) alternative variant dSep08, mRNA.
<a href="#">Pde2a</a>	<a href="#">Pde2a.eSep08</a>	<a href="#">81743</a>	1046	545	2	44	phosphodiesterase 2A, cGMP-stimulated (Pde2a) alternative variant eSep08, mRNA.
<a href="#">Pde4a</a>	<a href="#">Pde4a.bSep08</a>	<a href="#">25638</a>	2809	760	4	253	phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) (Pde4a) alternative variant bSep08, mRNA.
<a href="#">Pde4a</a>	<a href="#">Pde4a.cSep08</a>	<a href="#">25638</a>	2684	681	4	222	phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) (Pde4a) alternative variant cSep08, mRNA.

<a href="#">Pde4b</a>	<a href="#">Pde4b.bSep08</a>	<a href="#">24626</a>	28218	755	5	251	phosphodiesterase 4B, cAMP specific (Pde4b) alternative variant bSep08, mRNA.
<a href="#">Pde4b</a>	<a href="#">Pde4b.cSep08</a>	<a href="#">24626</a>	29483	521	5	173	phosphodiesterase 4B, cAMP specific (Pde4b) alternative variant cSep08, mRNA.
<a href="#">Pde4b</a>	<a href="#">Pde4b.eSep08</a>	<a href="#">24626</a>	1320	593	2	75	phosphodiesterase 4B, cAMP specific (8.6 kD) (Pde4b) alternative variant eSep08, mRNA.
<a href="#">Pde4c</a>	<a href="#">Pde4c.aSep08</a>	<a href="#">290646</a>	3751	1311	4	235	phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila) (24.5 kD) (Pde4c) mRNA.
<a href="#">Pde4d</a>	<a href="#">Pde4d.dSep08</a>	<a href="#">24627</a>	170847	521	1	84	phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila) (Pde4d) alternative variant dSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.bSep08</a>	<a href="#">64183</a>	7915	2163	7	344	phosphodiesterase 4D interacting protein precursor (37.4 kD) (Pde4dip) alternative variant bSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.cSep08</a>	<a href="#">64183</a>	8917	3401	6	232	phosphodiesterase 4D interacting protein (Pde4dip) alternative variant cSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.dSep08</a>	<a href="#">64183</a>	8876	551	5	183	phosphodiesterase 4D interacting protein (Pde4dip) alternative variant dSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.eSep08</a>	<a href="#">64183</a>	22221	366	3	70	phosphodiesterase 4D interacting protein (Pde4dip) alternative variant eSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.fSep08</a>	<a href="#">64183</a>	11233	978	3	63	phosphodiesterase 4D interacting protein (Pde4dip) alternative variant fSep08, mRNA.
<a href="#">Pde4dip</a>	<a href="#">Pde4dip.gSep08</a>	<a href="#">64183</a>	1657	955	2	46	phosphodiesterase 4D interacting protein (Pde4dip) alternative variant gSep08, mRNA.
<a href="#">Pde5a</a>	<a href="#">Pde5a.bSep08</a>	<a href="#">171115</a>	23180	785	6	221	phosphodiesterase 5A, cGMP-specific (Pde5a) alternative variant bSep08, mRNA.
<a href="#">Pde5a</a>	<a href="#">Pde5a.cSep08</a>	<a href="#">171115</a>	9084	613	4	119	phosphodiesterase 5A, cGMP-specific (Pde5a) alternative variant cSep08, mRNA.
<a href="#">Pde6a</a>	<a href="#">Pde6a.bSep08</a>	<a href="#">307401</a>	4961	998	2	60	phosphodiesterase 6A, cGMP-specific, rod, alpha (7.0 kD) (Pde6a) alternative variant bSep08, mRNA.
<a href="#">Pde6b</a>	<a href="#">Pde6b.bSep08</a>	<a href="#">289878</a>	3687	555		148	phosphodiesterase 6B, cGMP-specific, rod, beta (Pde6b) alternative variant bSep08, mRNA.
<a href="#">Pde6h</a>	<a href="#">Pde6h.aSep08</a>	<a href="#">114248</a>	10853	577	2	83	phosphodiesterase 6H, cGMP-specific, cone, gamma (9.0 kD) (Pde6h) alternative variant aSep08, mRNA.
<a href="#">Pde8a</a>	<a href="#">Pde8a.aSep08</a>	<a href="#">308776</a>	84380	769		202	phosphodiesterase 8A (Pde8a) mRNA.
<a href="#">Pde8b</a>	<a href="#">Pde8b.bSep08</a>	<a href="#">309962</a>	9400	378	3	95	phosphodiesterase 8B (Pde8b) alternative variant bSep08, mRNA.
<a href="#">Pde9a</a>	<a href="#">Pde9a.bSep08</a>	<a href="#">191569</a>	33694	853	10	212	phosphodiesterase 9A (Pde9a) alternative variant bSep08, mRNA.
<a href="#">Pde9a</a>	<a href="#">Pde9a.cSep08</a>	<a href="#">191569</a>	69510	670	10	200	phosphodiesterase 9A (Pde9a) alternative variant cSep08, mRNA.
<a href="#">Pde9a</a>	<a href="#">Pde9a.dSep08</a>	<a href="#">191569</a>	15246	786	5	144	phosphodiesterase 9A (Pde9a) alternative variant dSep08, mRNA.
<a href="#">Pde9a</a>	<a href="#">Pde9a.eSep08</a>	<a href="#">191569</a>	5906	585	5	114	phosphodiesterase 9A (Pde9a) alternative variant eSep08, mRNA.
<a href="#">Pde9a</a>	<a href="#">Pde9a.gSep08</a>	<a href="#">191569</a>	2167	484	2	28	phosphodiesterase 9A (3.3 kD) (Pde9a) alternative variant gSep08, mRNA.

<a href="#">Pde10a</a>	<a href="#">Pde10a.aSep08</a>	<a href="#">63885</a>	196134	2142		703	phosphodiesterase 10A (Pde10a) alternative variant aSep08, mRNA.
<a href="#">Pde12</a>	<a href="#">Pde12.bSep08</a>	<a href="#">306231</a>	6168	4076	1	608	phosphodiesterase 12 (67.2 kD) (Pde12) alternative variant bSep08, mRNA.
<a href="#">PDEase_I.0</a>	<a href="#">PDEase_I.0.aSep08</a>		32494	1419	9	473	phosphodiesterase 10A CRA a (PDEase_I.0) alternative variant aSep08, mRNA.
<a href="#">PDEase_I.0</a>	<a href="#">PDEase_I.0.bSep08</a>		2487	432	1	95	phosphodiesterase 10A CRA c (PDEase_I.0) alternative variant bSep08, mRNA.
<a href="#">PDEase_I.1</a>	<a href="#">PDEase_I.1.aSep08</a>		25781	2442	11	484	phosphodiesterase 8A CRA b (PDEase_I.1) alternative variant aSep08, mRNA.
<a href="#">PDEase_I.1</a>	<a href="#">PDEase_I.1.bSep08</a>		6540	809	4	136	phosphodiesterase 8A CRA b (PDEase_I.1) alternative variant bSep08, mRNA.
<a href="#">PDEase_I.1</a>	<a href="#">PDEase_I.1.cSep08</a>		361	240	2	28	putative protein (PDEase_I.1) alternative variant cSep08, mRNA.
<a href="#">Pdgfa</a>	<a href="#">Pdgfa.bSep08</a>	<a href="#">25266</a>	17869	1791	6	190	platelet-derived growth factor alpha polypeptide (Pdgfa) alternative variant bSep08, mRNA.
<a href="#">Pdgfa</a>	<a href="#">Pdgfa.cSep08</a>	<a href="#">25266</a>	17507	1359	5	175	platelet-derived growth factor alpha polypeptide (Pdgfa) alternative variant cSep08, mRNA.
<a href="#">Pdgfa</a>	<a href="#">Pdgfa.dSep08</a>	<a href="#">25266</a>	16779	627	5	174	platelet-derived growth factor alpha polypeptide (Pdgfa) alternative variant dSep08, mRNA.
<a href="#">Pdgfa</a>	<a href="#">Pdgfa.eSep08</a>	<a href="#">25266</a>	2559	404	2	105	platelet-derived growth factor alpha polypeptide (Pdgfa) alternative variant eSep08, mRNA.
<a href="#">Pdgfc</a>	<a href="#">Pdgfc.bSep08</a>	<a href="#">79429</a>	117704	755	4	251	platelet derived growth factor C (Pdgfc) alternative variant bSep08, mRNA.
<a href="#">Pdgfc</a>	<a href="#">Pdgfc.cSep08</a>	<a href="#">79429</a>	93744	432	2	57	platelet derived growth factor C (6.5 kD) (Pdgfc) alternative variant cSep08, mRNA.
<a href="#">Pdgfra</a>	<a href="#">Pdgfra.aSep08</a>	<a href="#">25267</a>	22111	4803	13	562	platelet derived growth factor receptor, alpha polypeptide (Pdgfra) alternative variant aSep08, mRNA.
<a href="#">Pdgfrb</a>	<a href="#">Pdgfrb.aSep08</a>	<a href="#">24629</a>	17648	4110	17	798	platelet derived growth factor receptor, beta polypeptide (Pdgfrb) alternative variant aSep08, mRNA.
<a href="#">Pdgfrb</a>	<a href="#">Pdgfrb.bSep08</a>	<a href="#">24629</a>	5683	907	6	302	platelet derived growth factor receptor, beta polypeptide (Pdgfrb) alternative variant bSep08, mRNA.
<a href="#">Pdgfrb</a>	<a href="#">Pdgfrb.cSep08</a>	<a href="#">24629</a>	1996	859	2	132	platelet derived growth factor receptor, beta polypeptide (14.3 kD) (Pdgfrb) alternative variant cSep08, mRNA.
<a href="#">Pdgfrl</a>	<a href="#">Pdgfrl.bSep08</a>	<a href="#">290771</a>	16658	721	2	117	platelet-derived growth factor receptor-like (Pdgfrl) alternative variant bSep08, mRNA.
<a href="#">Pdgfrl</a>	<a href="#">Pdgfrl.cSep08</a>	<a href="#">290771</a>	2193	808	2	99	platelet-derived growth factor receptor-like (11.1 kD) (Pdgfrl) alternative variant cSep08, mRNA.
<a href="#">Pdha1</a>	<a href="#">Pdha1.aSep08</a>	<a href="#">29554</a>	14376	3361	3	390	pyruvate dehydrogenase E1 alpha 1 (43.2 kD) (Pdha1) alternative variant aSep08, mRNA.
<a href="#">Pdha1</a>	<a href="#">Pdha1.bSep08</a>	<a href="#">29554</a>	8340	758	3	252	pyruvate dehydrogenase E1 alpha 1 (Pdha1) alternative variant bSep08, mRNA.
<a href="#">Pdha1</a>	<a href="#">Pdha1.cSep08</a>	<a href="#">29554</a>	1818	784	1	101	pyruvate dehydrogenase E1 alpha 1 (Pdha1) alternative variant cSep08, mRNA.
<a href="#">Pdhb</a>	<a href="#">Pdhb.bSep08</a>	<a href="#">289950</a>	2811	1051	6	114	pyruvate dehydrogenase E1-beta (12.7 kD) (Pdhb) alternative variant bSep08, mRNA.

<a href="#">Pdhb</a>	<a href="#">Pdhb.cSep08</a>	<a href="#">289950</a>	1043	804	2	86	pyruvate dehydrogenase (9.3 kD) (Pdhb) alternative variant cSep08, mRNA.
<a href="#">Pdhx</a>	<a href="#">Pdhx.bSep08</a>	<a href="#">311254</a>	65897	1757	11	510	pyruvate dehydrogenase complex, component X (Pdhx) alternative variant bSep08, mRNA.
<a href="#">Pdhx</a>	<a href="#">Pdhx.cSep08</a>	<a href="#">311254</a>	11540	1491	3	128	pyruvate dehydrogenase complex, component X (Pdhx) alternative variant cSep08, mRNA.
<a href="#">Pdia3</a>	<a href="#">Pdia3.bSep08</a>	<a href="#">29468</a>	13597	553	4	91	protein disulfide isomerase associated 3 (Pdia3) alternative variant bSep08, mRNA.
<a href="#">Pdia4</a>	<a href="#">Pdia4.bSep08</a>	<a href="#">116598</a>	5353	732	4	243	protein disulfide isomerase associated 4 (Pdia4) alternative variant bSep08, mRNA.
<a href="#">Pdia5</a>	<a href="#">Pdia5.bSep08</a>	<a href="#">360722</a>	80806	1119	12	180	protein disulfide isomerase-associated 5 (Pdia5) alternative variant bSep08, mRNA.
<a href="#">Pdia5</a>	<a href="#">Pdia5.cSep08</a>	<a href="#">360722</a>	1105	383	2	24	protein disulfide isomerase-associated 5 (2.7 kD) (Pdia5) alternative variant cSep08, mRNA.
<a href="#">Pdia6</a>	<a href="#">Pdia6.bSep08</a>	<a href="#">286906</a>	5658	1099	4	94	protein disulfide isomerase associated 6 CRA a (10.3 kD) (Pdia6) alternative variant bSep08, mRNA.
<a href="#">Pdia6</a>	<a href="#">Pdia6.cSep08</a>	<a href="#">286906</a>	1079	327	3	92	protein disulfide isomerase associated 6 CRA b (Pdia6) alternative variant cSep08, mRNA.
<a href="#">Pdia6</a>	<a href="#">Pdia6.dSep08</a>	<a href="#">286906</a>	3516	1250	3	82	protein disulfide isomerase (9.0 kD) (Pdia6) alternative variant dSep08, mRNA.
<a href="#">Pdk1</a>	<a href="#">Pdk1.bSep08</a>	<a href="#">116551</a>	836	393	3	68	pyruvate dehydrogenase kinase, isoenzyme 1 (Pdk1) alternative variant bSep08, mRNA.
<a href="#">Pdk2</a>	<a href="#">Pdk2.bSep08</a>	<a href="#">81530</a>	12617	1533	9	268	pyruvate dehydrogenase kinase, isoenzyme 2 (30.7 kD) (Pdk2) alternative variant bSep08, mRNA.
<a href="#">Pdk2</a>	<a href="#">Pdk2.cSep08</a>	<a href="#">81530</a>	11082	656	5	192	pyruvate dehydrogenase kinase, isoenzyme 2 (Pdk2) alternative variant cSep08, mRNA.
<a href="#">Pdk2</a>	<a href="#">Pdk2.dSep08</a>	<a href="#">81530</a>	1623	936	4	117	pyruvate dehydrogenase kinase, isoenzyme 2 (12.7 kD) (Pdk2) alternative variant dSep08, mRNA.
<a href="#">Pdk3</a>	<a href="#">Pdk3.aSep08</a>	<a href="#">296849</a>	66701	2078	6	415	pyruvate dehydrogenase kinase, isoenzyme 3 (47.9 kD) (Pdk3) alternative variant aSep08, mRNA.
<a href="#">Pdk3</a>	<a href="#">Pdk3.bSep08</a>	<a href="#">296849</a>	24339	557	1	185	pyruvate dehydrogenase kinase, isoenzyme 3 (Pdk3) alternative variant bSep08, mRNA.
<a href="#">Pdlim1</a>	<a href="#">Pdlim1.bSep08</a>	<a href="#">54133</a>	20918	640	4	213	PDZ and LIM domain 1 (elfin) (Pdlim1) alternative variant bSep08, mRNA.
<a href="#">Pdlim1</a>	<a href="#">Pdlim1.cSep08</a>	<a href="#">54133</a>	27039	808	5	176	PDZ and LIM domain 1 (elfin) (Pdlim1) alternative variant cSep08, mRNA.
<a href="#">Pdlim1</a>	<a href="#">Pdlim1.dSep08</a>	<a href="#">54133</a>	6929	753	3	95	PDZ and LIM domain 1 (elfin) (Pdlim1) alternative variant dSep08, mRNA.
<a href="#">Pdlim2</a>	<a href="#">Pdlim2.bSep08</a>	<a href="#">290354</a>	4115	705	5	156	PDZ and LIM domain 2 (16.5 kD) (Pdlim2) alternative variant bSep08, mRNA.
<a href="#">Pdlim2</a>	<a href="#">Pdlim2.cSep08</a>	<a href="#">290354</a>	4055	420	4	104	PDZ and LIM domain 2 (Pdlim2) alternative variant cSep08, mRNA.
<a href="#">Pdlim3</a>	<a href="#">Pdlim3.bSep08</a>	<a href="#">114108</a>	31721	1417	7	316	LIM protein (34.4 kD) (Pdlim3) alternative variant bSep08, complete mRNA.
<a href="#">Pdlim3</a>	<a href="#">Pdlim3.cSep08</a>	<a href="#">114108</a>	30128	997	6	292	LIM protein (31.5 kD) (Pdlim3) alternative variant cSep08, complete mRNA.
<a href="#">Pdlim3</a>	<a href="#">Pdlim3.dSep08</a>	<a href="#">114108</a>	29961	710	6	213	LIM protein (Pdlim3) alternative variant dSep08, mRNA.



<a href="#">Pdlim3</a>	<a href="#">Pdlim3.eSep08</a>	<a href="#">114108</a>	3219	896	2	75	lim protein (Pdlim3) alternative variant eSep08, mRNA.
<a href="#">Pdlim3</a>	<a href="#">Pdlim3.fSep08</a>	<a href="#">114108</a>	1642	517	2	54	LIM protein (Pdlim3) alternative variant fSep08, mRNA.
<a href="#">Pdlim5</a>	<a href="#">Pdlim5.bSep08</a>	<a href="#">64353</a>	54244	1621	8	207	PDZ LIM domain 5 (Pdlim5) alternative variant bSep08, mRNA.
<a href="#">Pdlim5</a>	<a href="#">Pdlim5.cSep08</a>	<a href="#">64353</a>	53562	924	7	202	PDZ LIM domain 5 (Pdlim5) alternative variant cSep08, mRNA.
<a href="#">Pdlim5</a>	<a href="#">Pdlim5.dSep08</a>	<a href="#">64353</a>	34815	596	5	151	PDZ LIM domain 5 (Pdlim5) alternative variant dSep08, mRNA.
<a href="#">Pdlim5</a>	<a href="#">Pdlim5.eSep08</a>	<a href="#">64353</a>	12485	908	7	124	PDZ LIM domain 5 (Pdlim5) alternative variant eSep08, mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.bSep08</a>	<a href="#">286908</a>	7537	768	9	232	PDZ and LIM domain 7 (Pdlim7) alternative variant bSep08, mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.cSep08</a>	<a href="#">286908</a>	6758	1000	8	222	PDZ and LIM domain 7 (24.5 kD) (Pdlim7) alternative variant cSep08, complete mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.dSep08</a>	<a href="#">286908</a>	10664	1522	11	213	PDZ and LIM domain 7 (23.1 kD) (Pdlim7) alternative variant dSep08, mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.eSep08</a>	<a href="#">286908</a>	2217	661	6	134	PDZ and LIM domain 7 (Pdlim7) alternative variant eSep08, mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.gSep08</a>	<a href="#">286908</a>	9842	763	7	103	PDZ and LIM domain 7 (Pdlim7) alternative variant gSep08, mRNA.
<a href="#">Pdlim7</a>	<a href="#">Pdlim7.hSep08</a>	<a href="#">286908</a>	3522	675	7	90	PDZ and LIM domain 7 (10.2 kD) (Pdlim7) alternative variant hSep08, mRNA.
<a href="#">Pdpk1</a>	<a href="#">Pdpk1.bSep08</a>	<a href="#">81745</a>	50395	714	6	212	3-phosphoinositide dependent protein kinase-1 (Pdpk1) alternative variant bSep08, mRNA.
<a href="#">Pdpk1</a>	<a href="#">Pdpk1.cSep08</a>	<a href="#">81745</a>	49654	719	5	179	3-phosphoinositide dependent protein kinase-1 (Pdpk1) alternative variant cSep08, mRNA.
<a href="#">Pdpk1</a>	<a href="#">Pdpk1.dSep08</a>	<a href="#">81745</a>	10666	1114	4	171	3-phosphoinositide dependent protein kinase-1 (19.8 kD) (Pdpk1) alternative variant dSep08, mRNA.
<a href="#">Pdpr</a>	<a href="#">Pdpr.bSep08</a>	<a href="#">307852</a>	8495	1784	5	231	pyruvate dehydrogenase phosphatase regulatory subunit (Pdpr) alternative variant bSep08, mRNA.
<a href="#">Pds5a</a>	<a href="#">Pds5a.bSep08</a>	<a href="#">305343</a>	4612	412	3	137	PDS5, regulator of cohesion maintenance, homolog A (S. cerevisiae) (Pds5a) alternative variant bSep08, mRNA.
<a href="#">Pds5b</a>	<a href="#">Pds5b.cSep08</a>	<a href="#">304218</a>	7833	678	3	225	PDS5, regulator of cohesion maintenance, homolog B (S. cerevisiae) (Pds5b) alternative variant cSep08, mRNA.
<a href="#">Pds5b</a>	<a href="#">Pds5b.dSep08</a>	<a href="#">304218</a>	2527	1088	2	39	PDS5, regulator of cohesion maintenance, homolog B (S. cerevisiae) (4.4 kD) (Pds5b) alternative variant dSep08, mRNA.
<a href="#">Pdss2</a>	<a href="#">Pdss2.aSep08</a>	<a href="#">365592</a>	901900	1793	3	597	prenyl (solanesyl) diphosphate synthase, subunit 2 (Pdss2) alternative variant aSep08, mRNA.
<a href="#">Pdss2</a>	<a href="#">Pdss2.bSep08</a>	<a href="#">365592</a>	34998	813	1	150	prenyl (solanesyl) diphosphate synthase, subunit 2 (17.4 kD) (Pdss2) alternative variant bSep08, mRNA.
<a href="#">Pdss2</a>	<a href="#">Pdss2.dSep08</a>	<a href="#">365592</a>	36717	834	2	63	prenyl (solanesyl) diphosphate synthase, subunit 2 (Pdss2) alternative variant dSep08, mRNA.
<a href="#">Pdxdc1</a>	<a href="#">Pdxdc1.aSep08</a>	<a href="#">304721</a>	84638	2699	23	807	putative protein, with a coiled coil domain, of ancient origin (Pdxdc1) alternative variant aSep08, mRNA.
<a href="#">Pdxdc1</a>	<a href="#">Pdxdc1.bSep08</a>	<a href="#">304721</a>	9202	793	3	109	putative protein of mammalian origin (Pdxdc1) alternative variant bSep08, mRNA.

<a href="#">PDZ.0</a>	<a href="#">PDZ.0.bSep08</a>		25938	966	10	312	LIM domain binding 3 like (PDZ.0) alternative variant bSep08, mRNA.
<a href="#">PDZ.0</a>	<a href="#">PDZ.0.cSep08</a>		11773	740	5	246	LIM domain binding 3 like (PDZ.0) alternative variant cSep08, mRNA.
<a href="#">PDZ.1</a>	<a href="#">PDZ.1.aSep08</a>		14631	1230	7	400	post-synaptic density 95 (PDZ.1) alternative variant aSep08, mRNA.
<a href="#">PDZ.1</a>	<a href="#">PDZ.1.bSep08</a>		564	444	2	105	discs large homolog 4 (PDZ.1) alternative variant bSep08, mRNA.
<a href="#">PDZ.1</a>	<a href="#">PDZ.1.cSep08</a>		578	224	3	74	discs large homolog 4 (PDZ.1) alternative variant cSep08, mRNA.
<a href="#">PDZ.2</a>	<a href="#">PDZ.2.aSep08</a>		9482	639	6	212	rho guanine nucleotide exchange factor 12 (PDZ.2) alternative variant aSep08, mRNA.
<a href="#">PDZ.2</a>	<a href="#">PDZ.2.bSep08</a>		6766	473	4	138	rho guanine nucleotide exchange factor 12 (PDZ.2) alternative variant bSep08, mRNA.
<a href="#">PDZ.3</a>	<a href="#">PDZ.3.aSep08</a>		15513	2462	12	435	multiple PDZ domain protein (PDZ.3) alternative variant aSep08, mRNA.
<a href="#">PDZ.3</a>	<a href="#">PDZ.3.bSep08</a>		3464	508	3	91	multiple PDZ domain protein (PDZ.3) alternative variant bSep08, mRNA.
<a href="#">PDZ.3</a>	<a href="#">PDZ.3.dSep08</a>		3542	617	3	45	putative protein (5.0 kD) (PDZ.3) alternative variant dSep08, mRNA.
<a href="#">PDZ.4</a>	<a href="#">PDZ.4.aSep08</a>		3283	383		127	multiple PDZ domain protein (PDZ.4) mRNA.
<a href="#">PDZ.5</a>	<a href="#">PDZ.5.aSep08</a>		27041	891		256	InaD-like (PDZ.5) mRNA.
<a href="#">PDZ.6</a>	<a href="#">PDZ.6.aSep08</a>		63356	631	4	128	InaD-like (PDZ.6) alternative variant aSep08, mRNA.
<a href="#">PDZ.7</a>	<a href="#">PDZ.7.aSep08</a>		12869	1172		390	signal-induced proliferation-associated 1 like 3 (PDZ.7) mRNA.
<a href="#">Pdzd3</a>	<a href="#">Pdzd3.aSep08</a>	<a href="#">500986</a>	1500	660	5	211	PDZ protein (Pdzd3) alternative variant aSep08, mRNA.
<a href="#">Pdzd3</a>	<a href="#">Pdzd3.bSep08</a>	<a href="#">500986</a>	846	714	2	71	PDZ protein like (Pdzd3) alternative variant bSep08, mRNA.
<a href="#">Pdzd7</a>	<a href="#">Pdzd7.aSep08</a>	<a href="#">293996</a>	2009	1653		155	putative protein of metazoan origin (Pdzd7) mRNA.
<a href="#">Pdzd11</a>	<a href="#">Pdzd11.bSep08</a>	<a href="#">302422</a>	2988	1023	6	140	PDZ/DHR/GLGF (16.2 kD) (Pdzd11) alternative variant bSep08, mRNA.
<a href="#">Pdzd11</a>	<a href="#">Pdzd11.cSep08</a>	<a href="#">302422</a>	3033	984	5	112	PDZ/DHR/GLGF (12.9 kD) (Pdzd11) alternative variant cSep08, mRNA.
<a href="#">Pdzd11</a>	<a href="#">Pdzd11.dSep08</a>	<a href="#">302422</a>	2425	1181	3	89	PDZ/DHR/GLGF (9.8 kD) (Pdzd11) alternative variant dSep08, mRNA.
<a href="#">Pdzd11</a>	<a href="#">Pdzd11.eSep08</a>	<a href="#">302422</a>	1885	416	4	50	putative protein of vertebrate origin (5.9 kD) (Pdzd11) alternative variant eSep08, mRNA.
<a href="#">Pdzk1</a>	<a href="#">Pdzk1.bSep08</a>	<a href="#">65144</a>	7242	1401	4	201	PDZ/DHR/GLGF (Pdzk1) alternative variant bSep08, mRNA.
<a href="#">Pdzk1ip1</a>	<a href="#">Pdzk1ip1.bSep08</a>	<a href="#">81916</a>	3705	399	1	132	PDZK1 interacting protein 1 (Pdzk1ip1) alternative variant bSep08, mRNA.
<a href="#">Pdzk1ip1</a>	<a href="#">Pdzk1ip1.cSep08</a>	<a href="#">81916</a>	3021	694	2	84	PDZK1 interacting protein 1 (9.2 kD) (Pdzk1ip1) alternative variant cSep08, mRNA.
<a href="#">Pdzm3</a>	<a href="#">Pdzm3.aSep08</a>	<a href="#">312607</a>	19172	1820		606	domain-containing ring finger protein 3 (Pdzm3) alternative variant aSep08, mRNA.

<a href="#">Pdzn3</a>	<a href="#">Pdzn3.bSep08</a>	<a href="#">312607</a>	20540	1685		330	PDZ/DHR/GLGF (Pdzn3) alternative variant bSep08, mRNA.
<a href="#">Pear1andRGD1309453</a>	<a href="#">Pear1andRGD1309453.aSep08</a>	<a href="#">295293</a>	21044	5738	25	1033	platelet endothelial aggregation receptor 1 (110.4 kD) (Pear1andRGD1309453) alternative variant aSep08, mRNA.
<a href="#">Pear1andRGD1309453</a>	<a href="#">Pear1andRGD1309453.aSep08</a>	<a href="#">310689</a>	21044	5738	25	1033	platelet endothelial aggregation receptor 1 (110.4 kD) (Pear1andRGD1309453) alternative variant aSep08, mRNA.
<a href="#">Pear1andRGD1309453</a>	<a href="#">Pear1andRGD1309453.cSep08</a>	<a href="#">295293</a>	3924	1086	6	206	CRA c (22.7 kD) (Pear1andRGD1309453) alternative variant cSep08, mRNA.
<a href="#">Pear1andRGD1309453</a>	<a href="#">Pear1andRGD1309453.cSep08</a>	<a href="#">310689</a>	3924	1086	6	206	CRA c (22.7 kD) (Pear1andRGD1309453) alternative variant cSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.aSep08</a>	<a href="#">291075</a>	36316	1766	5	445	peroxisomal isomerase CRA c (Peci) alternative variant aSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.cSep08</a>	<a href="#">291075</a>	27723	762	7	254	peroxisomal isomerase CRA e (Peci) alternative variant cSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.dSep08</a>	<a href="#">291075</a>	7961	821	7	229	peroxisomal isomerase CRA e (Peci) alternative variant dSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.eSep08</a>	<a href="#">291075</a>	6488	683	6	221	peroxisomal isomerase CRA e (Peci) alternative variant eSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.fSep08</a>	<a href="#">291075</a>	6444	826	6	178	peroxisomal isomerase CRA e (Peci) alternative variant fSep08, mRNA.
<a href="#">Peci</a>	<a href="#">Peci.gSep08</a>	<a href="#">291075</a>	1380	762	2	96	peroxisomal delta3 delta2-enoyl-Coenzyme a isomerase (11.2 kD) (Peci) alternative variant gSep08, mRNA.
<a href="#">Pecr</a>	<a href="#">Pecr.bSep08</a>	<a href="#">113956</a>	10188	877	4	132	peroxisomal trans-2-enoyl-CoA reductase (Pecr) alternative variant bSep08, mRNA.
<a href="#">Pecr</a>	<a href="#">Pecr.cSep08</a>	<a href="#">113956</a>	12059	519	2	121	peroxisomal trans-2-enoyl-CoA reductase (Pecr) alternative variant cSep08, mRNA.
<a href="#">Pecr</a>	<a href="#">Pecr.dSep08</a>	<a href="#">113956</a>	2971	424	2	33	peroxisomal trans-2-enoyl-CoA reductase (3.9 kD) (Pecr) alternative variant dSep08, mRNA.
<a href="#">peeby</a>	<a href="#">peeby.aSep08</a>		1411	454		151	taf1 RNA polymerase II TATA box binding protein - associated factor 250kDa like (peeby) mRNA.
<a href="#">peedar</a>	<a href="#">peedar.aSep08</a>		4080	1926		132	ATP-binding cassette sub-family e member 1 like (peedar) mRNA.
<a href="#">peefer</a>	<a href="#">peefer.aSep08</a>		24654	538		179	putative protein of metazoan origin (peefer) mRNA.
<a href="#">peeflo</a>	<a href="#">peeflo.aSep08</a>		1308	791	2	48	putative protein (5.3 kD) (peeflo) alternative variant aSep08, mRNA.
<a href="#">peeflu</a>	<a href="#">peeflu.aSep08</a>		5226	531		102	putative protein (10.9 kD) (peeflu) mRNA.
<a href="#">peegar</a>	<a href="#">peegar.bSep08</a>		1065	189	2	62	putative protein (peegar) alternative variant bSep08, mRNA.
<a href="#">peeja</a>	<a href="#">peeja.aSep08</a>		11660	288		60	putative protein (peeja) mRNA.
<a href="#">peeje</a>	<a href="#">peeje.aSep08</a>		3848	295		24	putative protein (3.0 kD) (peeje) mRNA.
<a href="#">peekee</a>	<a href="#">peekee.aSep08</a>		3465	526	2	46	CRA b like (peekee) alternative variant aSep08, mRNA.
<a href="#">peekee</a>	<a href="#">peekee.cSep08</a>		9186	465	4	45	CRA b like (peekee) alternative variant cSep08, mRNA.
<a href="#">peekee</a>	<a href="#">peekee.dSep08</a>		13631	403	3	44	putative protein (peekee) alternative variant dSep08, mRNA.

<a href="#">peekler</a>	<a href="#">peekler.aSep08</a>		23069	553	3	93	putative protein (peekler) alternative variant aSep08, mRNA.
<a href="#">peekler</a>	<a href="#">peekler.bSep08</a>		75080	597	4	81	putative protein (peekler) alternative variant bSep08, mRNA.
<a href="#">peeloy</a>	<a href="#">peeloy.aSep08</a>		8030	1545	8	478	putative protein of ancient origin (peeloy) alternative variant aSep08, mRNA.
<a href="#">peeloy</a>	<a href="#">peeloy.bSep08</a>		1058	383	3	111	putative protein of ancient origin (peeloy) alternative variant bSep08, mRNA.
<a href="#">peeloy</a>	<a href="#">peeloy.cSep08</a>		950	405	1	71	putative protein of eukaryotic origin (peeloy) alternative variant cSep08, mRNA.
<a href="#">peemee</a>	<a href="#">peemee.aSep08</a>		62579	382		127	dedicator of 2 (peemee) mRNA.
<a href="#">peemer</a>	<a href="#">peemer.aSep08</a>		2149	549		44	putative protein (5.1 kD) (peemer) mRNA.
<a href="#">peenoy</a>	<a href="#">peenoy.aSep08</a>		4923	698		232	ubiquitin specific peptidase 37 (peenoy) mRNA.
<a href="#">peepor</a>	<a href="#">peepor.aSep08</a>		2029	481		68	putative protein (7.8 kD) (peepor) alternative variant aSep08, mRNA.
<a href="#">peepor</a>	<a href="#">peepor.bSep08</a>		3233	718		54	putative protein (peepor) alternative variant bSep08, mRNA.
<a href="#">peesaa</a>	<a href="#">peesaa.aSep08</a>		5775	701		82	putative protein (9.3 kD) (peesaa) mRNA.
<a href="#">peeshee</a>	<a href="#">peeshee.aSep08</a>		9406	641		38	putative protein (4.4 kD) (peeshee) mRNA.
<a href="#">peetu</a>	<a href="#">peetu.aSep08</a>		8358	983		327	baculoviral IAP repeat-containing 6 (peetu) mRNA.
<a href="#">peevar</a>	<a href="#">peevar.aSep08</a>		3936	426	2	103	zinc finger DHHC-type containing 18 CRA b (peevar) alternative variant aSep08, mRNA.
<a href="#">peevar</a>	<a href="#">peevar.bSep08</a>		4679	895	2	98	containing 18 (peevar) alternative variant bSep08, mRNA.
<a href="#">peevar</a>	<a href="#">peevar.cSep08</a>		3008	474	2	84	containing Zinc finger DHHC domain 1 (peevar) alternative variant cSep08, mRNA.
<a href="#">peewey</a>	<a href="#">peewey.aSep08</a>		4473	411		85	calcium channel voltage-dependent (peewey) mRNA.
<a href="#">Pef1</a>	<a href="#">Pef1.bSep08</a>	<a href="#">297900</a>	6648	1336	4	229	calcium-binding EF-hand containing protein (Pef1) alternative variant bSep08, mRNA.
<a href="#">Peli1</a>	<a href="#">Peli1.bSep08</a>	<a href="#">305549</a>	7970	403	4	106	pellino homolog 1 (Drosophila) (Peli1) alternative variant bSep08, mRNA.
<a href="#">Peli1</a>	<a href="#">Peli1.cSep08</a>	<a href="#">305549</a>	17523	410	4	87	pellino homolog 1 (Drosophila) (Peli1) alternative variant cSep08, mRNA.
<a href="#">Penk1</a>	<a href="#">Penk1.bSep08</a>	<a href="#">29237</a>	5336	1294	2	269	proenkephalin 1 (30.9 kD) (Penk1) alternative variant bSep08, complete mRNA.
<a href="#">Peo1</a>	<a href="#">Peo1.bSep08</a>	<a href="#">309441</a>	7228	1472	6	231	twinkle (26.1 kD) (Peo1) alternative variant bSep08, mRNA.
<a href="#">Peo1</a>	<a href="#">Peo1.cSep08</a>	<a href="#">309441</a>	3336	350	2	116	twinkle (Peo1) alternative variant cSep08, mRNA.
<a href="#">Peo1</a>	<a href="#">Peo1.dSep08</a>	<a href="#">309441</a>	3271	272	2	61	putative protein (Peo1) alternative variant dSep08, mRNA.
<a href="#">Pepd</a>	<a href="#">Pepd.bSep08</a>	<a href="#">292808</a>	66436	1312		330	peptidase D (Pepd) alternative variant bSep08, mRNA.
<a href="#">Peptidase_C1.0</a>	<a href="#">Peptidase_C1.0.aSep08</a>		1170	393		130	cathepsin M (Peptidase_C1.0) mRNA.
<a href="#">Peptidase_C48.0</a>	<a href="#">Peptidase_C48.0.aSep08</a>		10038	2293		225	specific 7 (Peptidase_C48.0) mRNA.
<a href="#">Peptidase_M43.0</a>	<a href="#">Peptidase_M43.0.aSep08</a>		9713	690		230	pappalysin 2 (Peptidase_M43.0) mRNA.

<a href="#">Peptidase_M49.0</a>	<a href="#">Peptidase_M49.0.aSep08</a>		12777	1936	12	516	dipeptidyl peptidase III (Peptidase_M49.0) alternative variant aSep08, mRNA.
<a href="#">Peptidase_M49.0</a>	<a href="#">Peptidase_M49.0.bSep08</a>		3044	709	2	105	dipeptidylpeptidase 3 (11.7 kD) (Peptidase_M49.0) alternative variant bSep08, mRNA.
<a href="#">Peptidase_S9.0</a>	<a href="#">Peptidase_S9.0.aSep08</a>		5013	1521		307	dipeptidylpeptidase 9 (Peptidase_S9.0) mRNA.
<a href="#">Per1</a>	<a href="#">Per1.bSep08</a>	<a href="#">287422</a>	3101	1859	2	270	period homolog 1 (Drosophila) (27.5 kD) (Per1) alternative variant bSep08, mRNA.
<a href="#">Per2</a>	<a href="#">Per2.bSep08</a>	<a href="#">63840</a>	14482	1099	1	297	period homolog 2 (Drosophila) (32.4 kD) (Per2) alternative variant bSep08, mRNA.
<a href="#">Per3</a>	<a href="#">Per3.bSep08</a>	<a href="#">78962</a>	7295	822	4	187	period homolog 3 (Per3) alternative variant bSep08, mRNA.
<a href="#">Per3</a>	<a href="#">Per3.cSep08</a>	<a href="#">78962</a>	1706	491	3	149	period homolog 3 (Per3) alternative variant cSep08, mRNA.
<a href="#">Per3</a>	<a href="#">Per3.dSep08</a>	<a href="#">78962</a>	7733	559	4	125	period homolog 3 (Per3) alternative variant dSep08, mRNA.
<a href="#">Per3</a>	<a href="#">Per3.fSep08</a>	<a href="#">78962</a>	627	396	2	40	period homolog 3 (Per3) alternative variant fSep08, mRNA.
<a href="#">perby</a>	<a href="#">perby.aSep08</a>		5424	731		243	taf1 RNA polymerase II TATA box binding protein - associated factor 250kDa like (perby) mRNA.
<a href="#">perchy</a>	<a href="#">perchy.aSep08</a>		4305	716		56	putative protein (5.9 kD) (perchy) mRNA.
<a href="#">perdar</a>	<a href="#">perdar.aSep08</a>		5416	1145		381	protein hin-1 (perdar) mRNA.
<a href="#">perfer</a>	<a href="#">perfer.aSep08</a>		11326	578		73	CRA a like (perfer) mRNA.
<a href="#">perflo</a>	<a href="#">perflo.aSep08</a>		1971	623		110	putative protein of mammalian origin (perflo) alternative variant aSep08, mRNA.
<a href="#">perflu</a>	<a href="#">perflu.aSep08</a>		2314	427		35	putative protein (perflu) mRNA.
<a href="#">pergar</a>	<a href="#">pergar.aSep08</a>		1689	661		80	putative protein (9.1 kD) (pergar) mRNA.
<a href="#">perja</a>	<a href="#">perja.aSep08</a>		1301	1199	2	88	putative endoplasmic reticulum protein (9.9 kD) (perja) alternative variant aSep08, mRNA.
<a href="#">perjey</a>	<a href="#">perjey.aSep08</a>		892	692		43	putative protein (5.1 kD) (perjey) mRNA.
<a href="#">perkee</a>	<a href="#">perkee.aSep08</a>		53967	755		25	putative protein (2.8 kD) (perkee) mRNA.
<a href="#">perkler</a>	<a href="#">perkler.aSep08</a>		1351	522		69	putative protein (perkler) mRNA.
<a href="#">perloy</a>	<a href="#">perloy.aSep08</a>		302	195		64	putative protein (perloy) mRNA.
<a href="#">permee</a>	<a href="#">permee.aSep08</a>		3681	529	1	62	putative protein (permee) alternative variant aSep08, mRNA.
<a href="#">permee</a>	<a href="#">permee.bSep08</a>		10947	364	1	51	putative protein (permee) alternative variant bSep08, mRNA.
<a href="#">permer</a>	<a href="#">permer.aSep08</a>		8249	904		217	putative protein (permer) mRNA.
<a href="#">pernoy</a>	<a href="#">pernoy.aSep08</a>		4585	308		19	putative protein (pernoy) mRNA.
<a href="#">perpor</a>	<a href="#">perpor.aSep08</a>		1783	294		35	putative protein (perpor) mRNA.
<a href="#">persa</a>	<a href="#">persa.aSep08</a>		1457	388		36	putative protein (persa) mRNA.
<a href="#">pershee</a>	<a href="#">pershee.aSep08</a>		28637	743		37	putative protein (3.9 kD) (pershee) mRNA.
<a href="#">pertu</a>	<a href="#">pertu.aSep08</a>		567	416		132	baculoviral IAP repeat-containing 6 CRA c (pertu) mRNA.
<a href="#">pervar</a>	<a href="#">pervar.aSep08</a>	<a href="#">362616</a>	3826	528	3	128	CD164 sialomucin-like 2 (pervar) alternative variant aSep08, mRNA.

<a href="#">pervar</a>	<a href="#">pervar.bSep08</a>	<a href="#">362616</a>	3298	679	5	108	cd164 sialomucin-like 2 (11.8 kD) (pervar) alternative variant bSep08, mRNA.
<a href="#">perwey</a>	<a href="#">perwey.aSep08</a>		410	340		45	putative protein (5.1 kD) (perwey) mRNA.
<a href="#">PET.0</a>	<a href="#">PET.0.aSep08</a>		91206	991	7	305	prickle 1 (PET.0) alternative variant aSep08, mRNA.
<a href="#">PET.0</a>	<a href="#">PET.0.cSep08</a>		48043	550	4	82	putative protein (9.4 kD) (PET.0) alternative variant cSep08, mRNA.
<a href="#">PET.0</a>	<a href="#">PET.0.dSep08</a>		946	445	2	49	putative protein (PET.0) alternative variant dSep08, mRNA.
<a href="#">Pet112l</a>	<a href="#">Pet112l.aSep08</a>	<a href="#">361974</a>	25697	609		116	PET112-like (yeast) (Pet112l) mRNA.
<a href="#">Pex1</a>	<a href="#">Pex1.bSep08</a>	<a href="#">500006</a>	8815	1343	7	394	biogenesis factor 1 (Pex1) alternative variant bSep08, mRNA.
<a href="#">Pex1</a>	<a href="#">Pex1.cSep08</a>	<a href="#">500006</a>	3638	583	3	137	peroxisome biogenesis factor 1 (Pex1) alternative variant cSep08, mRNA.
<a href="#">Pex3</a>	<a href="#">Pex3.bSep08</a>	<a href="#">83519</a>	18554	743	6	117	peroxisomal biogenesis factor 3 (13.7 kD) (Pex3) alternative variant bSep08, mRNA.
<a href="#">Pex3</a>	<a href="#">Pex3.cSep08</a>	<a href="#">83519</a>	18506	504	6	97	peroxisomal biogenesis factor 3 (Pex3) alternative variant cSep08, mRNA.
<a href="#">Pex3</a>	<a href="#">Pex3.fSep08</a>	<a href="#">83519</a>	6680	354	2	40	peroxisomal biogenesis factor 3 (Pex3) alternative variant fSep08, mRNA.
<a href="#">Pex5</a>	<a href="#">Pex5.aSep08</a>	<a href="#">312703</a>	28709	3153	15	521	peroxisome biogenesis factor 5 (58.1 kD) (Pex5) alternative variant aSep08, mRNA.
<a href="#">Pex5</a>	<a href="#">Pex5.bSep08</a>	<a href="#">312703</a>	754	364	2	93	peroxisome biogenesis factor 5 (Pex5) alternative variant bSep08, mRNA.
<a href="#">Pex5</a>	<a href="#">Pex5.dSep08</a>	<a href="#">312703</a>	1603	391	2	64	peroxisome biogenesis factor 5 (Pex5) alternative variant dSep08, mRNA.
<a href="#">Pex5l</a>	<a href="#">Pex5l.bSep08</a>	<a href="#">286937</a>	56063	984	7	327	peroxisomal biogenesis factor 5-like (Pex5l) alternative variant bSep08, mRNA.
<a href="#">Pex5l</a>	<a href="#">Pex5l.cSep08</a>	<a href="#">286937</a>	69148	1398	9	308	peroxisomal biogenesis factor 5-like (Pex5l) alternative variant cSep08, mRNA.
<a href="#">Pex6</a>	<a href="#">Pex6.aSep08</a>	<a href="#">117265</a>	6897	1214		404	peroxisomal biogenesis factor 6 (Pex6) mRNA.
<a href="#">Pex11b</a>	<a href="#">Pex11b.aSep08</a>	<a href="#">310682</a>	9226	1742	2	450	peroxisomal biogenesis factor 11b (Pex11b) alternative variant aSep08, mRNA.
<a href="#">Pex11b</a>	<a href="#">Pex11b.cSep08</a>	<a href="#">310682</a>	8550	895	1	201	peroxisomal biogenesis factor 11b (22.2 kD) (Pex11b) alternative variant cSep08, mRNA.
<a href="#">Pex11c</a>	<a href="#">Pex11c.bSep08</a>	<a href="#">288369</a>	3970	711	1	71	peroxisomal biogenesis factor 11c (Pex11c) alternative variant bSep08, mRNA.
<a href="#">Pex14</a>	<a href="#">Pex14.bSep08</a>	<a href="#">64460</a>	111744	721	7	216	peroxisomal biogenesis factor 14 (Pex14) alternative variant bSep08, mRNA.
<a href="#">Pex14</a>	<a href="#">Pex14.cSep08</a>	<a href="#">64460</a>	85312	563	5	151	peroxisomal biogenesis factor 14 (Pex14) alternative variant cSep08, mRNA.
<a href="#">Pex14</a>	<a href="#">Pex14.dSep08</a>	<a href="#">64460</a>	21146	699	6	151	peroxisomal biogenesis factor 14 (Pex14) alternative variant dSep08, mRNA.
<a href="#">Pex14</a>	<a href="#">Pex14.eSep08</a>	<a href="#">64460</a>	87184	404	3	41	peroxisomal biogenesis factor 14 (Pex14) alternative variant eSep08, mRNA.
<a href="#">Pex16</a>	<a href="#">Pex16.aSep08</a>	<a href="#">311203</a>	3422	913	9	304	peroxisome biogenesis factor 16 (Pex16) alternative variant aSep08, mRNA.

<a href="#">Pex19</a>	<a href="#">Pex19.cSep08</a>	<a href="#">289233</a>	16332	1970	9	293	peroxisome biogenesis factor 19 (Pex19) alternative variant cSep08, mRNA.
<a href="#">Pex19</a>	<a href="#">Pex19.dSep08</a>	<a href="#">289233</a>	2177	1332	3	116	peroxisome biogenesis factor 19 (13.2 kD) (Pex19) alternative variant dSep08, mRNA.
<a href="#">peyby</a>	<a href="#">peyby.aSep08</a>		3755	370		123	taf1 RNA polymerase II TATA box binding protein - associated factor 250kDa CRA c like (peyby) mRNA.
<a href="#">peychy</a>	<a href="#">peychy.aSep08</a>		1676	433		77	putative protein (peychy) mRNA.
<a href="#">peydar</a>	<a href="#">peydar.aSep08</a>		68416	412		137	solute carrier family 10 member 7 (peydar) mRNA.
<a href="#">peyfer</a>	<a href="#">peyfer.aSep08</a>		651	334		49	putative protein (peyfer) mRNA.
<a href="#">peyflo</a>	<a href="#">peyflo.aSep08</a>		1468	433		53	putative protein (peyflo) mRNA.
<a href="#">peyflu</a>	<a href="#">peyflu.bSep08</a>		2424	790	2	49	putative protein (peyflu) alternative variant bSep08, mRNA.
<a href="#">peygar</a>	<a href="#">peygar.aSep08</a>		18855	433		40	putative protein (peygar) mRNA.
<a href="#">peyja</a>	<a href="#">peyja.aSep08</a>		8429	255		39	putative protein (peyja) mRNA.
<a href="#">peyjey</a>	<a href="#">peyjey.aSep08</a>		3379	227		62	putative protein (7.0 kD) (peyjey) mRNA.
<a href="#">peykee</a>	<a href="#">peykee.aSep08</a>		13085	520		172	astrotactin (peykee) mRNA.
<a href="#">peykler</a>	<a href="#">peykler.bSep08</a>		5759	878	2	72	ab2-143 like (8.0 kD) (peykler) alternative variant bSep08, mRNA.
<a href="#">peylo</a>	<a href="#">peylo.aSep08</a>		977	710		34	putative protein (3.8 kD) (peylo) mRNA.
<a href="#">peyloy</a>	<a href="#">peyloy.bSep08</a>		6819	1859	2	72	ab2-143 like (8.0 kD) (peyloy) alternative variant bSep08, mRNA.
<a href="#">peymee</a>	<a href="#">peymee.aSep08</a>		6270	183		50	putative protein (peymee) mRNA.
<a href="#">peymer</a>	<a href="#">peymer.aSep08</a>		554	354		51	putative protein (5.5 kD) (peymer) mRNA.
<a href="#">peynoy</a>	<a href="#">peynoy.aSep08</a>		4280	320		62	putative protein (peynoy) mRNA.
<a href="#">peypor</a>	<a href="#">peypor.aSep08</a>		8741	634		32	putative protein (peypor) mRNA.
<a href="#">peysa</a>	<a href="#">peysa.aSep08</a>		16483	723	2	124	putative protein (peysa) alternative variant aSep08, mRNA.
<a href="#">peysa</a>	<a href="#">peysa.bSep08</a>		7019	611	2	73	putative protein (peysa) alternative variant bSep08, mRNA.
<a href="#">peyshee</a>	<a href="#">peyshee.aSep08</a>		44904	465		94	mediator of RNA polymerase II transcription (peyshee) mRNA.
<a href="#">peytu</a>	<a href="#">peytu.aSep08</a>		18408	479		53	putative protein (6.1 kD) (peytu) mRNA.
<a href="#">peyvar</a>	<a href="#">peyvar.aSep08</a>		15921	394		75	putative protein (8.2 kD) (peyvar) mRNA.
<a href="#">peywey</a>	<a href="#">peywey.aSep08</a>		3506	1211		114	calcium channel voltage-dependent (peywey) mRNA.
<a href="#">Pfas</a>	<a href="#">Pfas.bSep08</a>	<a href="#">287420</a>	9528	401	4	133	phosphoribosylformylglycinamide synthase (FGAR amidotransferase) (Pfas) alternative variant bSep08, mRNA.
<a href="#">Pfas</a>	<a href="#">Pfas.cSep08</a>	<a href="#">287420</a>	788	392	3	130	phosphoribosylformylglycinamide synthase (FGAR amidotransferase) (Pfas) alternative variant cSep08, mRNA.
<a href="#">Pfas</a>	<a href="#">Pfas.dSep08</a>	<a href="#">287420</a>	547	422	2	95	phosphoribosylformylglycinamide synthase (FGAR amidotransferase) (Pfas) alternative variant dSep08, mRNA.
<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.cSep08</a>	<a href="#">300257</a>	4019	752	3	140	melanocyte proliferating gene 1 CRA a like (Pfdn5andMyg1) alternative variant cSep08, mRNA.
<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.cSep08</a>	<a href="#">300258</a>	4019	752	3	140	melanocyte proliferating gene 1 CRA a like (Pfdn5andMyg1) alternative variant cSep08, mRNA.

<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.dSep08</a>	<a href="#">300257</a>	888	751	2	110	melanocyte proliferating gene 1 like (12.6 kD) (Pfdn5andMyg1) alternative variant dSep08, complete mRNA.
<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.dSep08</a>	<a href="#">300258</a>	888	751	2	110	melanocyte proliferating gene 1 like (12.6 kD) (Pfdn5andMyg1) alternative variant dSep08, complete mRNA.
<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.eSep08</a>	<a href="#">300257</a>	6254	2445	5	85	prefoldin (9.6 kD) (Pfdn5andMyg1) alternative variant eSep08, complete mRNA.
<a href="#">Pfdn5andMyg1</a>	<a href="#">Pfdn5andMyg1.eSep08</a>	<a href="#">300258</a>	6254	2445	5	85	prefoldin (9.6 kD) (Pfdn5andMyg1) alternative variant eSep08, complete mRNA.
<a href="#">Pfkfb1</a>	<a href="#">Pfkfb1.bSep08</a>	<a href="#">24638</a>	32055	887	9	264	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1 (Pfkfb1) alternative variant bSep08, mRNA.
<a href="#">Pfkfb1</a>	<a href="#">Pfkfb1.cSep08</a>	<a href="#">24638</a>	34832	858	9	207	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1 (Pfkfb1) alternative variant cSep08, mRNA.
<a href="#">Pfkfb1</a>	<a href="#">Pfkfb1.dSep08</a>	<a href="#">24638</a>	1598	244	3	28	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1 (Pfkfb1) alternative variant dSep08, mRNA.
<a href="#">Pfkfb2</a>	<a href="#">Pfkfb2.dSep08</a>	<a href="#">24640</a>	1390	383	3	104	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2 (Pfkfb2) alternative variant dSep08, mRNA.
<a href="#">Pfkfb3</a>	<a href="#">Pfkfb3.bSep08</a>	<a href="#">117276</a>	26992	1712	14	462	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (53.6 kD) (Pfkfb3) alternative variant bSep08, mRNA.
<a href="#">Pfkfb3</a>	<a href="#">Pfkfb3.cSep08</a>	<a href="#">117276</a>	22309	1802	7	376	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (Pfkfb3) alternative variant cSep08, mRNA.
<a href="#">Pfkfb3</a>	<a href="#">Pfkfb3.eSep08</a>	<a href="#">117276</a>	65894	418	5	138	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (Pfkfb3) alternative variant eSep08, mRNA.
<a href="#">Pfkfb3</a>	<a href="#">Pfkfb3.gSep08</a>	<a href="#">117276</a>	2142	476	3	103	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (Pfkfb3) alternative variant gSep08, mRNA.
<a href="#">Pfkfb3</a>	<a href="#">Pfkfb3.iSep08</a>	<a href="#">117276</a>	6210	385	2	28	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (Pfkfb3) alternative variant iSep08, mRNA.
<a href="#">Pfkfb4</a>	<a href="#">Pfkfb4.bSep08</a>	<a href="#">54283</a>	38525	2685	13	419	6-phosphofructo-2-kinase fructose-2 (48.3 kD) (Pfkfb4) alternative variant bSep08, mRNA.
<a href="#">Pfkfb4</a>	<a href="#">Pfkfb4.cSep08</a>	<a href="#">54283</a>	37713	1757	14	379	6-phosphofructo-2-kinase fructose-2 4 (44.0 kD) (Pfkfb4) alternative variant cSep08, mRNA.
<a href="#">Pfkfb4</a>	<a href="#">Pfkfb4.dSep08</a>	<a href="#">54283</a>	37540	1413	12	372	6-phosphofructo-2-kinase fructose-2 4 (43.2 kD) (Pfkfb4) alternative variant dSep08, mRNA.
<a href="#">Pfkfb4</a>	<a href="#">Pfkfb4.eSep08</a>	<a href="#">54283</a>	19224	1339	8	151	6-phosphofructo-2-kinase fructose-2 transcript (Pfkfb4) alternative variant eSep08, mRNA.
<a href="#">Pfkfb4</a>	<a href="#">Pfkfb4.fSep08</a>	<a href="#">54283</a>	14672	1204	3	132	6-phosphofructo-2-kinase fructose-2 6 (Pfkfb4) alternative variant fSep08, mRNA.
<a href="#">Pfkl</a>	<a href="#">Pfkl.bSep08</a>	<a href="#">25741</a>	7156	888	9	235	phosphofructokinase, liver, B-type (Pfkl) alternative variant bSep08, mRNA.
<a href="#">Pfkm</a>	<a href="#">Pfkm.bSep08</a>	<a href="#">65152</a>	3231	562	5	171	phosphofructokinase, muscle (Pfkm) alternative variant bSep08, mRNA.
<a href="#">Pfkm</a>	<a href="#">Pfkm.cSep08</a>	<a href="#">65152</a>	7609	417	4	115	phosphofructokinase, muscle (Pfkm) alternative variant cSep08, mRNA.
<a href="#">Pfkm</a>	<a href="#">Pfkm.eSep08</a>	<a href="#">65152</a>	826	731	2	48	phosphofructokinase, muscle (5.6 kD) (Pfkm) alternative variant eSep08, mRNA.



<a href="#">Pfkp</a>	<a href="#">Pfkp.aSep08</a>	<a href="#">60416</a>	62766	3346	21	465	phosphofructokinase, platelet (Pfkp) alternative variant aSep08, mRNA.
<a href="#">Pfkp</a>	<a href="#">Pfkp.bSep08</a>	<a href="#">60416</a>	24674	1168	12	254	phosphofructokinase, platelet (Pfkp) alternative variant bSep08, mRNA.
<a href="#">Pfkp</a>	<a href="#">Pfkp.cSep08</a>	<a href="#">60416</a>	2000	778	1	54	phosphofructokinase, platelet (Pfkp) alternative variant cSep08, mRNA.
<a href="#">Pfn1</a>	<a href="#">Pfn1.cSep08</a>	<a href="#">64303</a>	1335	814	2	59	profilin 1 (6.8 kD) (Pfn1) alternative variant cSep08, mRNA.
<a href="#">Pfn1</a>	<a href="#">Pfn1.dSep08</a>	<a href="#">64303</a>	2519	627	3	59	profilin 1 (6.8 kD) (Pfn1) alternative variant dSep08, mRNA.
<a href="#">Pftk1</a>	<a href="#">Pftk1.bSep08</a>	<a href="#">362316</a>	226422	923	6	123	kinase (13.9 kD) (Pftk1) alternative variant bSep08, mRNA.
<a href="#">Pftk1</a>	<a href="#">Pftk1.cSep08</a>	<a href="#">362316</a>	32515	665	3	109	kinase (Pftk1) alternative variant cSep08, mRNA.
<a href="#">Pftk1</a>	<a href="#">Pftk1.dSep08</a>	<a href="#">362316</a>	7112	482	2	43	kinase (4.9 kD) (Pftk1) alternative variant dSep08, complete mRNA.
<a href="#">Pgam5</a>	<a href="#">Pgam5.bSep08</a>	<a href="#">288731</a>	6593	1520	4	296	phosphoglycerate mutase family member 5 (Pgam5) alternative variant bSep08, mRNA.
<a href="#">Pgam5</a>	<a href="#">Pgam5.cSep08</a>	<a href="#">288731</a>	3382	802	3	200	phosphoglycerate mutase family member 5 (Pgam5) alternative variant cSep08, mRNA.
<a href="#">Pgam5</a>	<a href="#">Pgam5.dSep08</a>	<a href="#">288731</a>	3199	713	2	138	phosphoglycerate mutase family member 5 (Pgam5) alternative variant dSep08, mRNA.
<a href="#">Pgcp</a>	<a href="#">Pgcp.bSep08</a>	<a href="#">58952</a>	277055	1374	5	327	plasma glutamate carboxypeptidase (35.9 kD) (Pgcp) alternative variant bSep08, mRNA.
<a href="#">Pgcp</a>	<a href="#">Pgcp.cSep08</a>	<a href="#">58952</a>	176184	1165	5	324	plasma glutamate carboxypeptidase (35.6 kD) (Pgcp) alternative variant cSep08, mRNA.
<a href="#">Pggt1b</a>	<a href="#">Pggt1b.bSep08</a>	<a href="#">81746</a>	29802	726	6	241	geranylgeranyltransferase type I (Pggt1b) alternative variant bSep08, mRNA.
<a href="#">Pggt1b</a>	<a href="#">Pggt1b.cSep08</a>	<a href="#">81746</a>	27119	2815	5	230	protein geranylgeranyltransferase type I (26.1 kD) (Pggt1b) alternative variant cSep08, mRNA.
<a href="#">Pggt1b</a>	<a href="#">Pggt1b.dSep08</a>	<a href="#">81746</a>	16293	437	4	118	geranylgeranyltransferase type I (13.6 kD) (Pggt1b) alternative variant dSep08, mRNA.
<a href="#">Pgk1</a>	<a href="#">Pgk1.bSep08</a>	<a href="#">24644</a>	4634	870	4	213	phosphoglycerate kinase 1 (Pgk1) alternative variant bSep08, mRNA.
<a href="#">Pgk1</a>	<a href="#">Pgk1.cSep08</a>	<a href="#">24644</a>	1125	564	2	56	phosphoglycerate kinase 1 (Pgk1) alternative variant cSep08, mRNA.
<a href="#">Pgls</a>	<a href="#">Pgls.bSep08</a>	<a href="#">290636</a>	3408	571	5	132	6-phosphogluconolactonase (Pgls) alternative variant bSep08, mRNA.
<a href="#">Pglyrp1</a>	<a href="#">Pglyrp1.bSep08</a>	<a href="#">84387</a>	13745	611	2	105	peptidoglycan recognition protein 1 (11.8 kD) (Pglyrp1) alternative variant bSep08, mRNA.
<a href="#">Pglyrp3</a>	<a href="#">Pglyrp3.aSep08</a>	<a href="#">499658</a>	5076	487		162	peptidoglycan recognition protein 3 (Pglyrp3) mRNA.
<a href="#">Pgm2</a>	<a href="#">Pgm2.aSep08</a>	<a href="#">289632</a>	35142	2109	14	636	phosphoglucomutase 2 (Pgm2) alternative variant aSep08, mRNA.
<a href="#">Pgm2</a>	<a href="#">Pgm2.dSep08</a>	<a href="#">289632</a>	23001	594	4	60	phosphoglucomutase 2 (Pgm2) alternative variant dSep08, mRNA.
<a href="#">Pgm2l1</a>	<a href="#">Pgm2l1.bSep08</a>	<a href="#">685076</a>	4988	1319	3	87	phosphoglucomutase 2-like 1 (Pgm2l1) alternative variant bSep08, mRNA.
<a href="#">Pgm3</a>	<a href="#">Pgm3.aSep08</a>	<a href="#">363109</a>	18242	2587	13	555	phosphoglucomutase 3 (Pgm3) alternative variant aSep08, mRNA.

<a href="#">Pgm5</a>	<a href="#">Pgm5.aSep08</a>	<a href="#">679990</a>	68481	1815	3	222	phosphoglucomutase 5 (Pgm5) alternative variant aSep08, mRNA.
<a href="#">Pgm5</a>	<a href="#">Pgm5.bSep08</a>	<a href="#">679990</a>	45549	297	1	78	phosphoglucomutase 5 (Pgm5) alternative variant bSep08, mRNA.
<a href="#">PGM_PMM_I.0</a>	<a href="#">PGM_PMM_I.0.aSep08</a>		41887	610		203	phosphoglucomutase (PGM_PMM_I.0) mRNA.
<a href="#">Pgrmc1</a>	<a href="#">Pgrmc1.bSep08</a>	<a href="#">291948</a>	6859	419	1	119	progesterone receptor membrane component 1 (Pgrmc1) alternative variant bSep08, mRNA.
<a href="#">Pgs1</a>	<a href="#">Pgs1.aSep08</a>	<a href="#">303698</a>	35625	2265	10	556	phosphatidylglycerophosphate synthase 1 (Pgs1) alternative variant aSep08, mRNA.
<a href="#">Pgs1</a>	<a href="#">Pgs1.cSep08</a>	<a href="#">303698</a>	4284	621	2	93	phosphatidylglycerophosphate synthase 1 (9.8 kD) (Pgs1) alternative variant cSep08, mRNA.
<a href="#">Pgs1</a>	<a href="#">Pgs1.eSep08</a>	<a href="#">303698</a>	4366	440	2	64	phosphatidylglycerophosphate synthase 1 (Pgs1) alternative variant eSep08, mRNA.
<a href="#">Pgs1</a>	<a href="#">Pgs1.fSep08</a>	<a href="#">303698</a>	4431	543	2	48	phosphatidylglycerophosphate synthase 1 (5.2 kD) (Pgs1) alternative variant fSep08, mRNA.
<a href="#">Pgs1</a>	<a href="#">Pgs1.gSep08</a>	<a href="#">303698</a>	6369	370	3	55	phosphatidylglycerophosphate synthase 1 (Pgs1) alternative variant gSep08, mRNA.
<a href="#">PG_binding_1.0</a>	<a href="#">PG_binding_1.0.aSep08</a>		1519	524		174	matrix 13 (PG_binding_1.0) mRNA.
<a href="#">Ph-4</a>	<a href="#">Ph-4.aSep08</a>	<a href="#">301008</a>	17839	1736	8	507	hypoxia-inducible factor prolyl 4-hydroxylase (Ph-4) alternative variant aSep08, mRNA.
<a href="#">Ph-4</a>	<a href="#">Ph-4.bSep08</a>	<a href="#">301008</a>	2174	428	1	118	hypoxia-inducible factor prolyl 4-hydroxylase (Ph-4) alternative variant bSep08, mRNA.
<a href="#">Ph-4</a>	<a href="#">Ph-4.cSep08</a>	<a href="#">301008</a>	11632	382	2	30	hypoxia-inducible factor prolyl 4-hydroxylase (3.1 kD) (Ph-4) alternative variant cSep08, mRNA.
<a href="#">PH.0</a>	<a href="#">PH.0.aSep08</a>		53054	383		109	rho GTPase activating protein 2 (PH.0) mRNA.
<a href="#">PH.1</a>	<a href="#">PH.1.aSep08</a>		5813	488		162	intersectin 1 CRA a (PH.1) mRNA.
<a href="#">PH.2</a>	<a href="#">PH.2.aSep08</a>		3847	1253		143	pleckstrin homology-like domain family B member 2 (PH.2) mRNA.
<a href="#">PH.3</a>	<a href="#">PH.3.bSep08</a>		47676	830	4	233	1 -specific guanine nucleotide-releasing factor (PH.3) alternative variant bSep08, mRNA.
<a href="#">PH.4</a>	<a href="#">PH.4.aSep08</a>		13641	657	6	218	oxysterol-binding protein-like protein 8 (PH.4) alternative variant aSep08, mRNA.
<a href="#">PH.5</a>	<a href="#">PH.5.aSep08</a>		4504	514	2	150	spectrin beta (PH.5) alternative variant aSep08, mRNA.
<a href="#">PH.5</a>	<a href="#">PH.5.bSep08</a>		801	349	1	79	spectrin beta (PH.5) alternative variant bSep08, mRNA.
<a href="#">PH.6</a>	<a href="#">PH.6.aSep08</a>		160202	1072	8	357	activator protein for secretion 2 CRA b (PH.6) alternative variant aSep08, mRNA.
<a href="#">PH.6</a>	<a href="#">PH.6.bSep08</a>		1757	374	1	109	activator protein for secretion 2 (PH.6) alternative variant bSep08, mRNA.
<a href="#">PH.7</a>	<a href="#">PH.7.aSep08</a>		8872	1137	8	273	pleckstrin homology-like domain family B member 3 (PH.7) alternative variant aSep08, mRNA.
<a href="#">PH.7</a>	<a href="#">PH.7.bSep08</a>		1206	738	2	89	pleckstrin homology-like domain family B member 3 CRA c (9.8 kD) (PH.7) alternative variant bSep08, mRNA.
<a href="#">PH.8</a>	<a href="#">PH.8.aSep08</a>		8561	487	1	161	homology domain-containing family A member 7 (PH.8) alternative variant aSep08, mRNA.

<a href="#">PH.8</a>	<a href="#">PH.8.bSep08</a>		15559	504	4	158	CRA b (PH.8) alternative variant bSep08, mRNA.
<a href="#">Phactr1</a>	<a href="#">Phactr1.bSep08</a>	<a href="#">306844</a>	162743	818	5	272	phosphatase and actin regulator 1 (Phactr1) alternative variant bSep08, mRNA.
<a href="#">Phactr3</a>	<a href="#">Phactr3.bSep08</a>	<a href="#">362284</a>	9346	1459	5	128	phosphatase and actin regulator 3 (15.6 kD) (Phactr3) alternative variant bSep08, mRNA.
<a href="#">Phb2</a>	<a href="#">Phb2.bSep08</a>	<a href="#">114766</a>	3600	2152	6	206	prohibitin 2 (22.9 kD) (Phb2) alternative variant bSep08, mRNA.
<a href="#">Phb2</a>	<a href="#">Phb2.cSep08</a>	<a href="#">114766</a>	2702	844	6	183	prohibitin 2 (Phb2) alternative variant cSep08, mRNA.
<a href="#">Phc1</a>	<a href="#">Phc1.bSep08</a>	<a href="#">312690</a>	14184	1644	4	547	polyhomeotic-like 1 (Phc1) alternative variant bSep08, mRNA.
<a href="#">Phc1</a>	<a href="#">Phc1.cSep08</a>	<a href="#">312690</a>	13894	2438	4	497	polyhomeotic-like 1 CRA c (Phc1) alternative variant cSep08, mRNA.
<a href="#">Phc1</a>	<a href="#">Phc1.eSep08</a>	<a href="#">312690</a>	967	796	2	100	polyhomeotic-like 1 CRA a (Phc1) alternative variant eSep08, mRNA.
<a href="#">Phc2</a>	<a href="#">Phc2.bSep08</a>	<a href="#">313038</a>	24230	2446	7	425	polyhomeotic-like 2 (Drosophila) (Phc2) alternative variant bSep08, mRNA.
<a href="#">Phc2</a>	<a href="#">Phc2.cSep08</a>	<a href="#">313038</a>	20369	1565	6	232	polyhomeotic-like 2 (Drosophila) (26.0 kD) (Phc2) alternative variant cSep08, mRNA.
<a href="#">Phc2</a>	<a href="#">Phc2.dSep08</a>	<a href="#">313038</a>	5830	784	4	178	polyhomeotic-like 2 (Drosophila) (19.8 kD) (Phc2) alternative variant dSep08, mRNA.
<a href="#">Phca</a>	<a href="#">Phca.aSep08</a>	<a href="#">499210</a>	6580	1216	4	82	phytoceramidase, alkaline (Phca) alternative variant aSep08, mRNA.
<a href="#">Phca</a>	<a href="#">Phca.bSep08</a>	<a href="#">499210</a>	5720	1034	3	72	phytoceramidase, alkaline (Phca) alternative variant bSep08, mRNA.
<a href="#">PHD.0</a>	<a href="#">PHD.0.aSep08</a>		1611	934		311	jumonji AT rich interactive domain 1C (PHD.0) mRNA.
<a href="#">PHD.1</a>	<a href="#">PHD.1.aSep08</a>		36167	3818		645	taf3 RNA polymerase II tata box binding protein -associated factor like (PHD.1) mRNA.
<a href="#">PHD.2</a>	<a href="#">PHD.2.aSep08</a>		8636	370		123	myst histone acetyltransferase 3 (PHD.2) mRNA.
<a href="#">PHD.3</a>	<a href="#">PHD.3.aSep08</a>		3020	376	3	125	PHD finger protein 20 (PHD.3) alternative variant aSep08, mRNA.
<a href="#">PHD.4</a>	<a href="#">PHD.4.aSep08</a>		12177	717	5	238	bromodomain adjacent zinc finger domain 1A (PHD.4) alternative variant aSep08, mRNA.
<a href="#">PHD.5</a>	<a href="#">PHD.5.aSep08</a>		44943	1549	9	351	myeloid lymphoid mixed-lineage leukemia 5 CRA a like (38.9 kD) (PHD.5) alternative variant aSep08, mRNA.
<a href="#">PHD.5</a>	<a href="#">PHD.5.bSep08</a>		16197	433	3	85	putative protein (PHD.5) alternative variant bSep08, mRNA.
<a href="#">PHD.5</a>	<a href="#">PHD.5.cSep08</a>		16653	816	2	75	putative protein (PHD.5) alternative variant cSep08, mRNA.
<a href="#">PHD.6</a>	<a href="#">PHD.6.aSep08</a>		1762	561		186	myeloid lymphoid mixed-lineage leukemia like (PHD.6) mRNA.
<a href="#">PHD.7</a>	<a href="#">PHD.7.aSep08</a>		14907	1046	7	258	spacing factor 1 (PHD.7) alternative variant aSep08, mRNA.
<a href="#">Phf1</a>	<a href="#">Phf1.bSep08</a>	<a href="#">294287</a>	2310	742	6	247	PHD finger protein 1 (Phf1) alternative variant bSep08, mRNA.
<a href="#">Phf1</a>	<a href="#">Phf1.cSep08</a>	<a href="#">294287</a>	1219	375	4	125	PHD finger protein 1 (Phf1) alternative variant cSep08, mRNA.

<a href="#">Phf1</a>	<a href="#">Phf1.dSep08</a>	<a href="#">294287</a>	1471	732	6	115	PHD finger protein 1 (Phf1) alternative variant dSep08, mRNA.
<a href="#">Phf2</a>	<a href="#">Phf2.aSep08</a>	<a href="#">306814</a>	11441	1120	6	373	PHD finger protein 2 (Phf2) alternative variant aSep08, mRNA.
<a href="#">Phf3</a>	<a href="#">Phf3.bSep08</a>	<a href="#">363210</a>	10126	1799	1	469	PHD finger protein 3 (Phf3) alternative variant bSep08, mRNA.
<a href="#">Phf5a</a>	<a href="#">Phf5a.bSep08</a>	<a href="#">192246</a>	6045	838	1	88	PHD finger protein 5A (Phf5a) alternative variant bSep08, mRNA.
<a href="#">Phf7</a>	<a href="#">Phf7.bSep08</a>	<a href="#">364510</a>	12788	1387	9	221	PHD finger protein 7 (25.1 kD) (Phf7) alternative variant bSep08, mRNA.
<a href="#">Phf7</a>	<a href="#">Phf7.cSep08</a>	<a href="#">364510</a>	5510	811	5	72	PHD finger protein 7 (8.3 kD) (Phf7) alternative variant cSep08, mRNA.
<a href="#">Phf10</a>	<a href="#">Phf10.bSep08</a>	<a href="#">292404</a>	6740	737	7	245	PHD finger protein 10 (Phf10) alternative variant bSep08, mRNA.
<a href="#">Phf10</a>	<a href="#">Phf10.cSep08</a>	<a href="#">292404</a>	2202	532	4	148	PHD finger protein 10 (Phf10) alternative variant cSep08, mRNA.
<a href="#">Phf10</a>	<a href="#">Phf10.dSep08</a>	<a href="#">292404</a>	4914	1750	3	136	PHD finger protein 10 (14.9 kD) (Phf10) alternative variant dSep08, mRNA.
<a href="#">Phf10</a>	<a href="#">Phf10.gSep08</a>	<a href="#">292404</a>	1563	1301	2	81	PHD finger protein 10 (9.2 kD) (Phf10) alternative variant gSep08, mRNA.
<a href="#">Phf11</a>	<a href="#">Phf11.aSep08</a>	<a href="#">361051</a>	31502	719	7	239	PHD finger protein 11 (Phf11) alternative variant aSep08, mRNA.
<a href="#">Phf11</a>	<a href="#">Phf11.bSep08</a>	<a href="#">361051</a>	56899	736	2	110	PHD finger protein 11 (12.4 kD) (Phf11) alternative variant bSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.bSep08</a>	<a href="#">303274</a>	13931	3075	10	725	PHD finger protein 12 (Phf12) alternative variant bSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.cSep08</a>	<a href="#">303274</a>	5925	1844	6	236	PHD finger protein 12 (25.4 kD) (Phf12) alternative variant cSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.dSep08</a>	<a href="#">303274</a>	3664	404	6	134	PHD finger protein 12 (Phf12) alternative variant dSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.eSep08</a>	<a href="#">303274</a>	8626	673	5	127	PHD finger protein 12 (Phf12) alternative variant eSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.fSep08</a>	<a href="#">303274</a>	3007	370	4	123	PHD finger protein 12 (Phf12) alternative variant fSep08, mRNA.
<a href="#">Phf12</a>	<a href="#">Phf12.iSep08</a>	<a href="#">303274</a>	1490	338	3	28	putative protein (Phf12) alternative variant iSep08, mRNA.
<a href="#">Phf14</a>	<a href="#">Phf14.bSep08</a>	<a href="#">500030</a>	104699	3511	16	877	PHD finger protein 14 (98.7 kD) (Phf14) alternative variant bSep08, complete mRNA.
<a href="#">Phf14</a>	<a href="#">Phf14.cSep08</a>	<a href="#">500030</a>	120222	788	6	218	PHD finger protein 14 (Phf14) alternative variant cSep08, mRNA.
<a href="#">Phf14</a>	<a href="#">Phf14.dSep08</a>	<a href="#">500030</a>	22191	435	3	100	PHD finger protein 14 (Phf14) alternative variant dSep08, mRNA.
<a href="#">Phf14</a>	<a href="#">Phf14.eSep08</a>	<a href="#">500030</a>	29495	689	4	97	PHD finger protein 14 (Phf14) alternative variant eSep08, mRNA.
<a href="#">Phf14</a>	<a href="#">Phf14.fSep08</a>	<a href="#">500030</a>	92148	500	4	83	PHD finger protein 14 (Phf14) alternative variant fSep08, mRNA.
<a href="#">Phf15</a>	<a href="#">Phf15.bSep08</a>	<a href="#">303113</a>	43462	2297	2	671	PHD finger protein 15 (Phf15) alternative variant bSep08, mRNA.

<a href="#">Phf17</a>	<a href="#">Phf17.bSep08</a>	<a href="#">310352</a>	8083	845	4	281	PHD finger protein 17 (Phf17) alternative variant bSep08, mRNA.
<a href="#">Phf17</a>	<a href="#">Phf17.cSep08</a>	<a href="#">310352</a>	27909	492	5	163	PHD finger protein 17 (Phf17) alternative variant cSep08, mRNA.
<a href="#">Phf19</a>	<a href="#">Phf19.bSep08</a>	<a href="#">296653</a>	3537	2400		132	PHD finger protein 19 (Phf19) alternative variant bSep08, mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.aSep08</a>	<a href="#">314964</a>	32824	1045	4	327	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (Phf201) alternative variant aSep08, mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.aSep08</a>	<a href="#">690663</a>	32824	1045	4	327	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (Phf201) alternative variant aSep08, mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.bSep08</a>	<a href="#">314964</a>	30611	2521	4	316	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (35.8 kD) (Phf201) alternative variant bSep08, complete mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.bSep08</a>	<a href="#">690663</a>	30611	2521	4	316	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (35.8 kD) (Phf201) alternative variant bSep08, complete mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.cSep08</a>	<a href="#">314964</a>	14673	1324	3	285	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (Phf201) alternative variant cSep08, mRNA.
<a href="#">Phf201</a>	<a href="#">Phf201.cSep08</a>	<a href="#">690663</a>	14673	1324	3	285	PHD finger protein 20-like 1 and hypothetical protein LOC690663 (Phf201) alternative variant cSep08, mRNA.
<a href="#">Phf21b</a>	<a href="#">Phf21b.bSep08</a>	<a href="#">300117</a>	62783	1802	5	400	PHD finger protein 21B (Phf21b) alternative variant bSep08, mRNA.
<a href="#">Phf21b</a>	<a href="#">Phf21b.cSep08</a>	<a href="#">300117</a>	2513	407	2	113	PHD finger protein 21B (Phf21b) alternative variant cSep08, mRNA.
<a href="#">Phgdh</a>	<a href="#">Phgdh.aSep08</a>	<a href="#">58835</a>	29018	1832	12	533	3-phosphoglycerate dehydrogenase (56.5 kD) (Phgdh) alternative variant aSep08, mRNA.
<a href="#">Phgdh</a>	<a href="#">Phgdh.bSep08</a>	<a href="#">58835</a>	3248	1382	5	429	3-phosphoglycerate dehydrogenase (Phgdh) alternative variant bSep08, mRNA.
<a href="#">Phgdh</a>	<a href="#">Phgdh.cSep08</a>	<a href="#">58835</a>	12211	767	4	92	3-phosphoglycerate dehydrogenase (10.1 kD) (Phgdh) alternative variant cSep08, mRNA.
<a href="#">Phgdh1</a>	<a href="#">Phgdh1.aSep08</a>	<a href="#">361094</a>	146141	1342	7	345	phosphoglycerate dehydrogenase like 1 (39.1 kD) (Phgdh1) alternative variant aSep08, mRNA.
<a href="#">Phgdh1</a>	<a href="#">Phgdh1.bSep08</a>	<a href="#">361094</a>	107051	820	4	187	phosphoglycerate dehydrogenase like 1 (21.4 kD) (Phgdh1) alternative variant bSep08, mRNA.
<a href="#">Phka1</a>	<a href="#">Phka1.aSep08</a>	<a href="#">64561</a>	25323	706	3	235	phosphorylase kinase alpha 1 (Phka1) alternative variant aSep08, mRNA.
<a href="#">Phka1</a>	<a href="#">Phka1.bSep08</a>	<a href="#">64561</a>	16874	403	1	134	phosphorylase kinase alpha 1 (Phka1) alternative variant bSep08, mRNA.
<a href="#">Phka2</a>	<a href="#">Phka2.aSep08</a>	<a href="#">678739</a>	3603	636		201	phosphorylase kinase alpha 2 (Phka2) mRNA.
<a href="#">Phkb</a>	<a href="#">Phkb.bSep08</a>	<a href="#">361377</a>	190210	1825	8	384	phosphorylase kinase, beta (Phkb) alternative variant bSep08, complete mRNA.
<a href="#">Phkb</a>	<a href="#">Phkb.cSep08</a>	<a href="#">361377</a>	27585	784	7	260	phosphorylase kinase, beta (Phkb) alternative variant cSep08, mRNA.
<a href="#">Phkb</a>	<a href="#">Phkb.dSep08</a>	<a href="#">361377</a>	30232	2680	6	223	phosphorylase kinase, beta (25.7 kD) (Phkb) alternative variant dSep08, mRNA.
<a href="#">Phkg2</a>	<a href="#">Phkg2.bSep08</a>	<a href="#">140671</a>	12053	1230	8	286	phosphorylase kinase gamma (32.9 kD) (Phkg2) alternative variant bSep08, mRNA.

<a href="#">Phkg2</a>	<a href="#">Phkg2.cSep08</a>	<a href="#">140671</a>	11629	778	7	207	phosphorylase kinase gamma (Phkg2) alternative variant cSep08, mRNA.
<a href="#">Phkg2</a>	<a href="#">Phkg2.dSep08</a>	<a href="#">140671</a>	3850	1433	3	133	phosphorylase kinase gamma 2 (Phkg2) alternative variant dSep08, mRNA.
<a href="#">Phkg2</a>	<a href="#">Phkg2.eSep08</a>	<a href="#">140671</a>	8888	702	4	118	phosphorylase kinase gamma 2 (Phkg2) alternative variant eSep08, mRNA.
<a href="#">Phkg2</a>	<a href="#">Phkg2.fSep08</a>	<a href="#">140671</a>	11320	795	6	96	phosphorylase kinase gamma 2 (Phkg2) alternative variant fSep08, mRNA.
<a href="#">Phkg2</a>	<a href="#">Phkg2.hSep08</a>	<a href="#">140671</a>	564	483	2	63	phosphorylase kinase gamma 2 (Phkg2) alternative variant hSep08, mRNA.
<a href="#">Phlda1</a>	<a href="#">Phlda1.bSep08</a>	<a href="#">29380</a>	1887	1479	2	114	pleckstrin homology-like domain, family A, member 1 and hypothetical protein LOC688472 (11.0 kD) (Phlda1) alternative variant bSep08, complete mRNA.
<a href="#">Phlda1</a>	<a href="#">Phlda1.bSep08</a>	<a href="#">688472</a>	1887	1479	2	114	pleckstrin homology-like domain, family A, member 1 and hypothetical protein LOC688472 (11.0 kD) (Phlda1) alternative variant bSep08, complete mRNA.
<a href="#">Phlda2</a>	<a href="#">Phlda2.aSep08</a>	<a href="#">293637</a>	949	651		129	pleckstrin homology-like domain, family A, member 2 (Phlda2) mRNA.
<a href="#">Phldb1</a>	<a href="#">Phldb1.aSep08</a>	<a href="#">171434</a>	9547	1865	6	203	pleckstrin homology-like domain, family B, member 1 (Phldb1) alternative variant aSep08, mRNA.
<a href="#">Phldb1</a>	<a href="#">Phldb1.bSep08</a>	<a href="#">171434</a>	1787	872	2	72	pleckstrin homology-like domain, family B, member 1 (Phldb1) alternative variant bSep08, mRNA.
<a href="#">Phldb2</a>	<a href="#">Phldb2.aSep08</a>	<a href="#">685611</a>	15345	665	5	221	pleckstrin homology-like domain, family B, member 2 (Phldb2) alternative variant aSep08, mRNA.
<a href="#">Phldb3</a>	<a href="#">Phldb3.aSep08</a>	<a href="#">308431</a>	5983	961		225	pleckstrin homology-like domain, family B, member 3 (Phldb3) mRNA.
<a href="#">Phlpb</a>	<a href="#">Phlpb.bSep08</a>	<a href="#">192259</a>	20916	1096	10	232	phospholipase B CRA c (26.7 kD) (Phlpb) alternative variant bSep08, mRNA.
<a href="#">Phlpb</a>	<a href="#">Phlpb.cSep08</a>	<a href="#">192259</a>	4277	753	5	102	phospholipase B CRA b (Phlpb) alternative variant cSep08, mRNA.
<a href="#">Phlpb</a>	<a href="#">Phlpb.dSep08</a>	<a href="#">192259</a>	3380	605	3	93	putative protein (Phlpb) alternative variant dSep08, mRNA.
<a href="#">Phlpb</a>	<a href="#">Phlpb.eSep08</a>	<a href="#">192259</a>	2110	465	3	62	phospholipase B CRA b (Phlpb) alternative variant eSep08, mRNA.
<a href="#">Phlpb</a>	<a href="#">Phlpb.fSep08</a>	<a href="#">192259</a>	1885	320	3	54	phospholipase B CRA b (Phlpb) alternative variant fSep08, mRNA.
<a href="#">Phlppi</a>	<a href="#">Phlppi.bSep08</a>	<a href="#">498949</a>	1648	506	1	89	PH domain and leucine rich repeat protein phosphatase-like (Phlppi) alternative variant bSep08, mRNA.
<a href="#">Phpt1</a>	<a href="#">Phpt1.bSep08</a>	<a href="#">296571</a>	1374	672	3	115	phosphohistidine phosphatase 1 (13.0 kD) (Phpt1) alternative variant bSep08, complete mRNA.
<a href="#">Phtf1</a>	<a href="#">Phtf1.aSep08</a>	<a href="#">252962</a>	14010	1778	10	487	putative homeodomain transcription factor 1 (Phtf1) alternative variant aSep08, mRNA.
<a href="#">Phtf1</a>	<a href="#">Phtf1.bSep08</a>	<a href="#">252962</a>	7506	919	5	145	putative homeodomain transcription factor 1 (Phtf1) alternative variant bSep08, mRNA.
<a href="#">Phtf1</a>	<a href="#">Phtf1.cSep08</a>	<a href="#">252962</a>	1599	755	2	73	putative homeodomain transcription factor 1 (Phtf1) alternative variant cSep08, mRNA.
<a href="#">Phtf2</a>	<a href="#">Phtf2.aSep08</a>	<a href="#">296762</a>	25988	943		314	putative homeodomain transcription factor 2 (Phtf2) mRNA.

<a href="#">Phyh</a>	<a href="#">Phyh.bSep08</a>	<a href="#">114209</a>	9980	1493	6	241	phytanoyl-CoA (27.6 kD) (Phyh) alternative variant bSep08, mRNA.
<a href="#">Phyh</a>	<a href="#">Phyh.cSep08</a>	<a href="#">114209</a>	6404	520	3	115	phytanoyl-CoA (Phyh) alternative variant cSep08, mRNA.
<a href="#">Phyh</a>	<a href="#">Phyh.dSep08</a>	<a href="#">114209</a>	6431	959	3	104	phytanoyl-CoA hydroxylase (11.8 kD) (Phyh) alternative variant dSep08, mRNA.
<a href="#">Phyhd1</a>	<a href="#">Phyhd1.bSep08</a>	<a href="#">296621</a>	9292	415	6	138	putative protein of ancient origin (Phyhd1) alternative variant bSep08, mRNA.
<a href="#">Phyhipl</a>	<a href="#">Phyhipl.aSep08</a>	<a href="#">309901</a>	41741	2680	5	375	phytanoyl-CoA hydroxylase interacting protein-like (42.4 kD) (Phyhipl) alternative variant aSep08, mRNA.
<a href="#">Pi4k2a</a>	<a href="#">Pi4k2a.bSep08</a>	<a href="#">114554</a>	16556	1700	8	309	phosphatidylinositol 4-kinase type 2 alpha (Pi4k2a) alternative variant bSep08, mRNA.
<a href="#">Pi4k2a</a>	<a href="#">Pi4k2a.cSep08</a>	<a href="#">114554</a>	5560	2016	3	126	phosphatidylinositol 4-kinase type 2 alpha (Pi4k2a) alternative variant cSep08, mRNA.
<a href="#">Pi4k2a</a>	<a href="#">Pi4k2a.dSep08</a>	<a href="#">114554</a>	7851	465	5	122	phosphatidylinositol 4-kinase type 2 alpha (Pi4k2a) alternative variant dSep08, mRNA.
<a href="#">Pi4k2a</a>	<a href="#">Pi4k2a.eSep08</a>	<a href="#">114554</a>	2857	355	3	118	phosphatidylinositol 4-kinase type 2 alpha (Pi4k2a) alternative variant eSep08, mRNA.
<a href="#">Pi4ka</a>	<a href="#">Pi4ka.bSep08</a>	<a href="#">64161</a>	2030	408	4	135	phosphatidylinositol 4-kinase, catalytic, alpha (Pi4ka) alternative variant bSep08, mRNA.
<a href="#">Pi4kb</a>	<a href="#">Pi4kb.bSep08</a>	<a href="#">81747</a>	8198	1332	5	217	phosphatidylinositol 4-kinase, catalytic, beta polypeptide (Pi4kb) alternative variant bSep08, mRNA.
<a href="#">Pi4kb</a>	<a href="#">Pi4kb.cSep08</a>	<a href="#">81747</a>	1391	683	2	84	phosphatidylinositol 4-kinase, catalytic, beta polypeptide (Pi4kb) alternative variant cSep08, mRNA.
<a href="#">Pi16</a>	<a href="#">Pi16.aSep08</a>	<a href="#">294312</a>	8231	1699	5	535	peptidase inhibitor 16 (Pi16) alternative variant aSep08, mRNA.
<a href="#">Pi16</a>	<a href="#">Pi16.bSep08</a>	<a href="#">294312</a>	8574	858	7	264	peptidase inhibitor 16 (29.3 kD) (Pi16) alternative variant bSep08, mRNA.
<a href="#">Pi16</a>	<a href="#">Pi16.cSep08</a>	<a href="#">294312</a>	1180	653	3	183	peptidase inhibitor 16 (Pi16) alternative variant cSep08, mRNA.
<a href="#">Pi16</a>	<a href="#">Pi16.dSep08</a>	<a href="#">294312</a>	7388	852	5	175	peptidase inhibitor 16 (19.7 kD) (Pi16) alternative variant dSep08, mRNA.
<a href="#">Pias1</a>	<a href="#">Pias1.bSep08</a>	<a href="#">300772</a>	59072	728	5	242	protein inhibitor of activated STAT 1 (Pias1) alternative variant bSep08, mRNA.
<a href="#">Pias1</a>	<a href="#">Pias1.cSep08</a>	<a href="#">300772</a>	11137	881	4	140	protein inhibitor of activated STAT 1 (Pias1) alternative variant cSep08, mRNA.
<a href="#">Pias2</a>	<a href="#">Pias2.bSep08</a>	<a href="#">83422</a>	18187	924	7	231	protein inhibitor of activated STAT 2 (Pias2) alternative variant bSep08, mRNA.
<a href="#">Pias3</a>	<a href="#">Pias3.bSep08</a>	<a href="#">83614</a>	4890	1089	7	328	protein inhibitor of activated STAT 3 (Pias3) alternative variant bSep08, mRNA.
<a href="#">Pias3</a>	<a href="#">Pias3.cSep08</a>	<a href="#">83614</a>	2626	580	2	190	protein inhibitor of activated STAT 3 (Pias3) alternative variant cSep08, mRNA.
<a href="#">Pias4</a>	<a href="#">Pias4.bSep08</a>	<a href="#">362827</a>	1098	992	2	108	protein inhibitor of activated STAT, 4 (Pias4) alternative variant bSep08, mRNA.
<a href="#">Pib5pa</a>	<a href="#">Pib5pa.bSep08</a>	<a href="#">171088</a>	1247	1135	1	319	phosphatidylinositol (4,5) bisphosphate 5-phosphatase, A (Pib5pa) alternative variant bSep08, mRNA.
<a href="#">Pibf1</a>	<a href="#">Pibf1.aSep08</a>	<a href="#">306104</a>	56949	867		181	progesterone immunomodulatory binding factor 1 (Pibf1) mRNA.

<a href="#">Picalm</a>	<a href="#">Picalm.bSep08</a>	<a href="#">89816</a>	23617	1684	10	306	phosphatidylinositol binding clathrin assembly protein (Picalm) alternative variant bSep08, mRNA.
<a href="#">Picalm</a>	<a href="#">Picalm.cSep08</a>	<a href="#">89816</a>	33189	1265	11	288	phosphatidylinositol binding clathrin assembly protein (Picalm) alternative variant cSep08, mRNA.
<a href="#">Picalm</a>	<a href="#">Picalm.dSep08</a>	<a href="#">89816</a>	32417	711	9	224	phosphatidylinositol binding clathrin assembly protein (Picalm) alternative variant dSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.aSep08</a>	<a href="#">84591</a>	19699	1791	6	452	protein -binding like (Pick1) alternative variant aSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.cSep08</a>	<a href="#">84591</a>	19640	1840	12	288	arfaptin-like (32.9 kD) (Pick1) alternative variant cSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.dSep08</a>	<a href="#">84591</a>	16323	779	9	259	protein -binding like (Pick1) alternative variant dSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.eSep08</a>	<a href="#">84591</a>	16597	1000	9	228	protein interacting with 1 (Pick1) alternative variant eSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.fSep08</a>	<a href="#">84591</a>	13472	628	6	135	protein -binding like (Pick1) alternative variant fSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.gSep08</a>	<a href="#">84591</a>	1105	751	2	96	putative cytoplasmic protein (10.4 kD) (Pick1) alternative variant gSep08, mRNA.
<a href="#">Pick1</a>	<a href="#">Pick1.hSep08</a>	<a href="#">84591</a>	2610	742	4	88	protein -binding like (Pick1) alternative variant hSep08, mRNA.
<a href="#">PID.0</a>	<a href="#">PID.0.aSep08</a>		24686	4766	2	382	adaptor protein containing pH domain leucine zipper 1 (PID.0) alternative variant aSep08, mRNA.
<a href="#">PID.0</a>	<a href="#">PID.0.bSep08</a>		14594	741	1	247	adaptor protein containing pH domain leucine zipper 1 (PID.0) alternative variant bSep08, mRNA.
<a href="#">Pid1</a>	<a href="#">Pid1.aSep08</a>	<a href="#">501174</a>	219026	2627	3	293	putative protein of bilateral origin (Pid1) alternative variant aSep08, mRNA.
<a href="#">Piga</a>	<a href="#">Piga.bSep08</a>	<a href="#">363464</a>	5741	744	2	41	phosphatidylinositol glycan anchor biosynthesis, class A (Piga) alternative variant bSep08, mRNA.
<a href="#">Pigc</a>	<a href="#">Pigc.aSep08</a>	<a href="#">364032</a>	2473	1558	1	297	phosphatidylinositol glycan anchor biosynthesis, class C (33.7 kD) (Pigc) alternative variant aSep08, mRNA.
<a href="#">Pigg</a>	<a href="#">Pigg.aSep08</a>	<a href="#">305626</a>	5737	508		169	phosphatidylinositol glycan anchor biosynthesis, class G (Pigg) mRNA.
<a href="#">Pigh</a>	<a href="#">Pigh.bSep08</a>	<a href="#">362756</a>	2165	362	2	114	phosphatidylinositol glycan anchor biosynthesis, class H (Pigh) alternative variant bSep08, mRNA.
<a href="#">Pigh</a>	<a href="#">Pigh.cSep08</a>	<a href="#">362756</a>	2041	399	3	81	phosphatidylinositol glycan anchor biosynthesis, class H (Pigh) alternative variant cSep08, mRNA.
<a href="#">Pigl</a>	<a href="#">Pigl.bSep08</a>	<a href="#">192263</a>	1866	1307	2	83	phosphatidylinositol glycan anchor biosynthesis, class L (Pigl) alternative variant bSep08, mRNA.
<a href="#">Pigo</a>	<a href="#">Pigo.aSep08</a>	<a href="#">313341</a>	3331	1793	4	547	phosphatidylinositol glycan anchor biosynthesis, class O (Pigo) alternative variant aSep08, mRNA.
<a href="#">Pigo</a>	<a href="#">Pigo.bSep08</a>	<a href="#">313341</a>	1338	893	2	297	phosphatidylinositol glycan anchor biosynthesis, class O (Pigo) alternative variant bSep08, mRNA.
<a href="#">Pigo</a>	<a href="#">Pigo.cSep08</a>	<a href="#">313341</a>	566	448	2	61	phosphatidylinositol glycan anchor biosynthesis, class O (Pigo) alternative variant cSep08, mRNA.
<a href="#">Pigp</a>	<a href="#">Pigp.bSep08</a>	<a href="#">288238</a>	5978	633	5	132	phosphatidylinositol glycan anchor biosynthesis, class P (15.2 kD) (Pigp) alternative variant bSep08, mRNA.



<a href="#">Pigp</a>	<a href="#">Pigp.dSep08</a>	<a href="#">288238</a>	4265	392	4	55	phosphatidylinositol glycan anchor biosynthesis, class P (6.4 kD) (Pigp) alternative variant dSep08, mRNA.
<a href="#">Pigq</a>	<a href="#">Pigq.bSep08</a>	<a href="#">287159</a>	7364	819	8	259	phosphatidylinositol glycan anchor biosynthesis, class Q (Pigq) alternative variant bSep08, mRNA.
<a href="#">Pigq</a>	<a href="#">Pigq.cSep08</a>	<a href="#">287159</a>	4913	760	2	253	phosphatidylinositol glycan anchor biosynthesis, class Q (Pigq) alternative variant cSep08, mRNA.
<a href="#">Pigs</a>	<a href="#">Pigs.bSep08</a>	<a href="#">303277</a>	8823	556	5	185	phosphatidylinositol glycan anchor biosynthesis, class S (Pigs) alternative variant bSep08, mRNA.
<a href="#">Pigt</a>	<a href="#">Pigt.bSep08</a>	<a href="#">296360</a>	8443	2015	2	233	phosphatidylinositol glycan anchor biosynthesis, class T (Pigt) alternative variant bSep08, mRNA.
<a href="#">Pigv</a>	<a href="#">Pigv.bSep08</a>	<a href="#">366478</a>	8359	2075	4	399	phosphatidylinositol glycan anchor biosynthesis, class V (43.7 kD) (Pigv) alternative variant bSep08, complete mRNA.
<a href="#">Pigv</a>	<a href="#">Pigv.cSep08</a>	<a href="#">366478</a>	7322	873	2	250	phosphatidylinositol glycan anchor biosynthesis, class V (Pigv) alternative variant cSep08, mRNA.
<a href="#">Pigv</a>	<a href="#">Pigv.dSep08</a>	<a href="#">366478</a>	11841	1680	2	118	phosphatidylinositol glycan anchor biosynthesis, class V (13.6 kD) (Pigv) alternative variant dSep08, mRNA.
<a href="#">Pigx</a>	<a href="#">Pigx.aSep08</a>	<a href="#">288041</a>	17906	1496	7	315	phosphatidylinositol glycan anchor biosynthesis, class X (Pigx) alternative variant aSep08, mRNA.
<a href="#">Pigx</a>	<a href="#">Pigx.cSep08</a>	<a href="#">288041</a>	1470	847	2	52	phosphatidylinositol glycan anchor biosynthesis, class X (5.7 kD) (Pigx) alternative variant cSep08, mRNA.
<a href="#">Pih1d1</a>	<a href="#">Pih1d1.aSep08</a>	<a href="#">292898</a>	4208	1785	6	568	putative protein of eukaryotic origin (Pih1d1) alternative variant aSep08, mRNA.
<a href="#">Pih1d1</a>	<a href="#">Pih1d1.cSep08</a>	<a href="#">292898</a>	3035	813	4	87	putative protein of eukaryotic origin (Pih1d1) alternative variant cSep08, mRNA.
<a href="#">Pih1d2</a>	<a href="#">Pih1d2.aSep08</a>	<a href="#">315645</a>	4877	1061	6	235	putative protein of eukaryotic origin (Pih1d2) alternative variant aSep08, mRNA.
<a href="#">Pih1d2</a>	<a href="#">Pih1d2.bSep08</a>	<a href="#">315645</a>	6004	1119	4	185	putative protein of eukaryotic origin (Pih1d2) alternative variant bSep08, mRNA.
<a href="#">Pih1d2</a>	<a href="#">Pih1d2.cSep08</a>	<a href="#">315645</a>	1989	1070	2	82	putative protein (4.9 kD) (Pih1d2) alternative variant cSep08, mRNA.
<a href="#">Pik3ap1</a>	<a href="#">Pik3ap1.aSep08</a>	<a href="#">294048</a>	60447	4114	15	668	phosphoinositide-3-kinase adaptor protein 1 (Pik3ap1) alternative variant aSep08, mRNA.
<a href="#">Pik3ap1</a>	<a href="#">Pik3ap1.cSep08</a>	<a href="#">294048</a>	8740	445	3	90	phosphoinositide-3-kinase adaptor protein 1 (Pik3ap1) alternative variant cSep08, mRNA.
<a href="#">Pik3c2g</a>	<a href="#">Pik3c2g.bSep08</a>	<a href="#">116720</a>	19587	672	4	206	phox-like (Pik3c2g) alternative variant bSep08, mRNA.
<a href="#">Pik3ca</a>	<a href="#">Pik3ca.aSep08</a>	<a href="#">170911</a>	16328	4699	11	495	phosphatidylinositol 3-kinase, catalytic, alpha polypeptide (Pik3ca) alternative variant aSep08, mRNA.
<a href="#">Pik3ca</a>	<a href="#">Pik3ca.bSep08</a>	<a href="#">170911</a>	1930	1168	3	188	phosphatidylinositol 3-kinase, catalytic, alpha polypeptide (22.0 kD) (Pik3ca) alternative variant bSep08, mRNA.
<a href="#">Pik3ca</a>	<a href="#">Pik3ca.cSep08</a>	<a href="#">170911</a>	606	506	2	66	phosphatidylinositol 3-kinase, catalytic, alpha polypeptide (Pik3ca) alternative variant cSep08, mRNA.
<a href="#">Pik3cb</a>	<a href="#">Pik3cb.bSep08</a>	<a href="#">85243</a>	24611	2448	9	377	phosphatidylinositol 3-kinase, catalytic, beta polypeptide (Pik3cb) alternative variant bSep08, mRNA.
<a href="#">Pik3cb</a>	<a href="#">Pik3cb.cSep08</a>	<a href="#">85243</a>	1788	738	2	106	phosphatidylinositol 3-kinase, catalytic, beta polypeptide (Pik3cb) alternative variant cSep08, mRNA.

<a href="#">Pik3cd</a>	<a href="#">Pik3cd.bSep08</a>	<a href="#">366508</a>	3270	1113	5	197	phosphatidylinositol 3-kinase catalytic delta polypeptide (Pik3cd) alternative variant bSep08, mRNA.
<a href="#">Pik3cg</a>	<a href="#">Pik3cg.aSep08</a>	<a href="#">298947</a>	21170	1414		264	phosphoinositide-3-kinase, catalytic, gamma polypeptide (Pik3cg) mRNA.
<a href="#">Pik3ip1</a>	<a href="#">Pik3ip1.aSep08</a>	<a href="#">305472</a>	11981	2336	6	267	interacting protein 1 (29.1 kD) (Pik3ip1) alternative variant aSep08, mRNA.
<a href="#">Pik3ip1</a>	<a href="#">Pik3ip1.bSep08</a>	<a href="#">305472</a>	5271	773	2	112	putative mitochondrial protein (12.0 kD) (Pik3ip1) alternative variant bSep08, mRNA.
<a href="#">Pik3ip1</a>	<a href="#">Pik3ip1.cSep08</a>	<a href="#">305472</a>	8202	735	2	112	putative mitochondrial protein (12.0 kD) (Pik3ip1) alternative variant cSep08, mRNA.
<a href="#">Pik3ip1</a>	<a href="#">Pik3ip1.eSep08</a>	<a href="#">305472</a>	1686	757	4	88	putative protein (Pik3ip1) alternative variant eSep08, mRNA.
<a href="#">Pik3ip1</a>	<a href="#">Pik3ip1.gSep08</a>	<a href="#">305472</a>	1272	278	2	25	putative protein (Pik3ip1) alternative variant gSep08, mRNA.
<a href="#">Pik3r1</a>	<a href="#">Pik3r1.bSep08</a>	<a href="#">25513</a>	5860	2545	8	465	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha) (Pik3r1) alternative variant bSep08, mRNA.
<a href="#">Pik3r1</a>	<a href="#">Pik3r1.cSep08</a>	<a href="#">25513</a>	19005	1559	3	126	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha) (13.9 kD) (Pik3r1) alternative variant cSep08, mRNA.
<a href="#">Pik3r1</a>	<a href="#">Pik3r1.dSep08</a>	<a href="#">25513</a>	1505	351	2	116	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha) (Pik3r1) alternative variant dSep08, mRNA.
<a href="#">Pik3r2</a>	<a href="#">Pik3r2.bSep08</a>	<a href="#">29741</a>	2798	1566	7	322	regulatory (37.3 kD) (Pik3r2) alternative variant bSep08, mRNA.
<a href="#">Pik3r2</a>	<a href="#">Pik3r2.cSep08</a>	<a href="#">29741</a>	2226	1257	7	200	regulatory (23.8 kD) (Pik3r2) alternative variant cSep08, mRNA.
<a href="#">Pik3r2</a>	<a href="#">Pik3r2.dSep08</a>	<a href="#">29741</a>	1159	732	4	192	regulatory (Pik3r2) alternative variant dSep08, mRNA.
<a href="#">Pik3r2</a>	<a href="#">Pik3r2.eSep08</a>	<a href="#">29741</a>	6899	484	4	157	regulatory (Pik3r2) alternative variant eSep08, mRNA.
<a href="#">Pik3r2</a>	<a href="#">Pik3r2.fSep08</a>	<a href="#">29741</a>	1125	737	4	131	regulatory (Pik3r2) alternative variant fSep08, mRNA.
<a href="#">Pik3r3</a>	<a href="#">Pik3r3.bSep08</a>	<a href="#">60664</a>	12601	1782	5	184	phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 3 (p55) (Pik3r3) alternative variant bSep08, mRNA.
<a href="#">Pik3r3</a>	<a href="#">Pik3r3.cSep08</a>	<a href="#">60664</a>	45400	512	4	106	phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 3 (p55) (Pik3r3) alternative variant cSep08, mRNA.
<a href="#">Pik3r4</a>	<a href="#">Pik3r4.aSep08</a>	<a href="#">363131</a>	49015	4998	20	1358	phosphoinositide-3-kinase, regulatory subunit 4, p150 (152.4 kD) (Pik3r4) alternative variant aSep08, mRNA.
<a href="#">Pik3r5</a>	<a href="#">Pik3r5.aSep08</a>	<a href="#">497931</a>	2332	668		222	phosphoinositide-3-kinase, regulatory subunit 5, p101 (Pik3r5) mRNA.
<a href="#">Pik3r6</a>	<a href="#">Pik3r6.bSep08</a>	<a href="#">497932</a>	19790	1419	8	259	phosphoinositide-3-kinase, regulatory subunit 6 (Pik3r6) alternative variant bSep08, mRNA.
<a href="#">Pik3r6</a>	<a href="#">Pik3r6.cSep08</a>	<a href="#">497932</a>	49400	825	8	228	phosphoinositide-3-kinase, regulatory subunit 6 (Pik3r6) alternative variant cSep08, mRNA.
<a href="#">Pik3r6</a>	<a href="#">Pik3r6.dSep08</a>	<a href="#">497932</a>	1840	338	3	73	phosphoinositide-3-kinase, regulatory subunit 6 (Pik3r6) alternative variant dSep08, mRNA.

<a href="#">Pilra</a>	<a href="#">Pilra.aSep08</a>	<a href="#">288568</a>	11477	1458	3	199	paired immunoglobulin-like type 2 receptor alpha (Pilra) alternative variant aSep08, mRNA.
<a href="#">Pilra</a>	<a href="#">Pilra.bSep08</a>	<a href="#">288568</a>	5658	446	1	109	paired immunoglobulin-like type 2 receptor alpha (Pilra) alternative variant bSep08, mRNA.
<a href="#">Pim1</a>	<a href="#">Pim1.cSep08</a>	<a href="#">24649</a>	4545	241	3	45	proviral integration site 1 (Pim1) alternative variant cSep08, mRNA.
<a href="#">Pim3</a>	<a href="#">Pim3.aSep08</a>	<a href="#">64534</a>	1412	1337		121	proviral integration site 3 (Pim3) mRNA.
<a href="#">Pin1</a>	<a href="#">Pin1.bSep08</a>	<a href="#">298696</a>	12603	1104	3	89	protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1 (Pin1) alternative variant bSep08, mRNA.
<a href="#">Pin1</a>	<a href="#">Pin1.cSep08</a>	<a href="#">298696</a>	8015	388	2	73	protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1 (Pin1) alternative variant cSep08, mRNA.
<a href="#">Pink1</a>	<a href="#">Pink1.aSep08</a>	<a href="#">298575</a>	12641	2837	8	580	PTEN induced putative kinase 1 (Pink1) alternative variant aSep08, complete mRNA.
<a href="#">Pinx1</a>	<a href="#">Pinx1.bSep08</a>	<a href="#">305963</a>	37717	764	3	79	PIN2-interacting protein 1 (9.2 kD) (Pinx1) alternative variant bSep08, mRNA.
<a href="#">Pip4k2a</a>	<a href="#">Pip4k2a.bSep08</a>	<a href="#">116723</a>	25468	2757		193	phosphatidylinositol-5-phosphate 4-kinase, type II, alpha (Pip4k2a) alternative variant bSep08, mRNA.
<a href="#">Pip4k2c</a>	<a href="#">Pip4k2c.cSep08</a>	<a href="#">140607</a>	1128	1079	2	38	phosphatidylinositol-5-phosphate 4-kinase, type II, gamma (4.1 kD) (Pip4k2c) alternative variant cSep08, mRNA.
<a href="#">Pip5k1a</a>	<a href="#">Pip5k1a.bSep08</a>	<a href="#">365865</a>	27619	778	1	105	phosphatidylinositol-4-phosphate 5-kinase, type 1, alpha (Pip5k1a) alternative variant bSep08, mRNA.
<a href="#">Pip5k1b</a>	<a href="#">Pip5k1b.bSep08</a>	<a href="#">309419</a>	9520	858	1	11	phosphatidylinositol-4-phosphate 5-kinase, type 1, beta (8.2 kD) (Pip5k1b) alternative variant bSep08, mRNA.
<a href="#">Pip5k1c</a>	<a href="#">Pip5k1c.aSep08</a>	<a href="#">314641</a>	27892	4184	17	636	phosphatidylinositol-4-phosphate 5-kinase, type 1 gamma (69.7 kD) (Pip5k1c) alternative variant aSep08, complete mRNA.
<a href="#">Pip5k1c</a>	<a href="#">Pip5k1c.bSep08</a>	<a href="#">314641</a>	5091	972	7	192	phosphatidylinositol-4-phosphate 5-kinase, type 1 gamma (Pip5k1c) alternative variant bSep08, mRNA.
<a href="#">Pip5k1c</a>	<a href="#">Pip5k1c.cSep08</a>	<a href="#">314641</a>	14747	676	5	176	phosphatidylinositol-4-phosphate 5-kinase, type 1 gamma (18.9 kD) (Pip5k1c) alternative variant cSep08, mRNA.
<a href="#">Pip5k1c</a>	<a href="#">Pip5k1c.dSep08</a>	<a href="#">314641</a>	2182	399	2	122	phosphatidylinositol-4-phosphate 5-kinase, type 1 gamma (Pip5k1c) alternative variant dSep08, mRNA.
<a href="#">Pip5k3</a>	<a href="#">Pip5k3.aSep08</a>	<a href="#">316457</a>	8707	631		210	phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinase, type III (Pip5k3) mRNA.
<a href="#">Pirb</a>	<a href="#">Pirb.bSep08</a>	<a href="#">65146</a>	3078	741	2	136	paired-Ig-like receptor B (Pirb) alternative variant bSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.aSep08</a>	<a href="#">681361</a>	49017	2179	8	406	phosphatidylserine decarboxylase (Pisd) alternative variant aSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.bSep08</a>	<a href="#">681361</a>	46862	717	6	238	phosphatidylserine decarboxylase (Pisd) alternative variant bSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.cSep08</a>	<a href="#">681361</a>	7619	992	6	227	phosphatidylserine decarboxylase (Pisd) alternative variant cSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.dSep08</a>	<a href="#">681361</a>	8014	738	5	168	phosphatidylserine decarboxylase (Pisd) alternative variant dSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.eSep08</a>	<a href="#">681361</a>	22723	377	3	117	CRA a (Pisd) alternative variant eSep08, mRNA.
<a href="#">Pisd</a>	<a href="#">Pisd.fSep08</a>	<a href="#">681361</a>	6878	394	3	79	phosphatidylserine decarboxylase (Pisd) alternative variant fSep08, mRNA.

<a href="#">Pisd</a>	<a href="#">Pisd.gSep08</a>	<a href="#">681361</a>	5367	417	3	44	phosphatidylserine decarboxylase (Pisd) alternative variant gSep08, mRNA.
<a href="#">Pitpna</a>	<a href="#">Pitpna.bSep08</a>	<a href="#">29525</a>	10181	389	3	129	phosphatidylinositol transfer protein, alpha (Pitpna) alternative variant bSep08, mRNA.
<a href="#">Pitpnb</a>	<a href="#">Pitpnb.aSep08</a>	<a href="#">114561</a>	33190	2348	7	170	phosphatidylinositol transfer protein, beta (Pitpnb) alternative variant aSep08, mRNA.
<a href="#">Pitpnb</a>	<a href="#">Pitpnb.bSep08</a>	<a href="#">114561</a>	31640	714	6	154	phosphatidylinositol transfer protein, beta (Pitpnb) alternative variant bSep08, mRNA.
<a href="#">Pitpnb</a>	<a href="#">Pitpnb.cSep08</a>	<a href="#">114561</a>	30907	1543	5	83	phosphatidylinositol transfer protein, beta (10.1 kD) (Pitpnb) alternative variant cSep08, mRNA.
<a href="#">Pitpnb</a>	<a href="#">Pitpnb.dSep08</a>	<a href="#">114561</a>	7991	1182	5	82	phosphatidylinositol transfer protein, beta (9.9 kD) (Pitpnb) alternative variant dSep08, mRNA.
<a href="#">Pitpnm1</a>	<a href="#">Pitpnm1.bSep08</a>	<a href="#">361694</a>	1811	411	2	136	phosphatidylinositol transfer protein, membrane-associated 1 (Pitpnm1) alternative variant bSep08, mRNA.
<a href="#">Pitpnm1</a>	<a href="#">Pitpnm1.dSep08</a>	<a href="#">361694</a>	1718	386	2	51	phosphatidylinositol transfer protein, membrane-associated 1 (Pitpnm1) alternative variant dSep08, mRNA.
<a href="#">Pitpnm2</a>	<a href="#">Pitpnm2.aSep08</a>	<a href="#">304474</a>	10723	1307		411	phosphatidylinositol transfer protein, membrane-associated 2 (Pitpnm2) mRNA.
<a href="#">Pitpnm3</a>	<a href="#">Pitpnm3.aSep08</a>	<a href="#">287467</a>	3544	422		140	PITPNM family member 3 (Pitpnm3) mRNA.
<a href="#">Pitrm1</a>	<a href="#">Pitrm1.bSep08</a>	<a href="#">307081</a>	7152	831	8	276	pitriysin 1 (Pitrm1) alternative variant bSep08, mRNA.
<a href="#">Pitrm1</a>	<a href="#">Pitrm1.cSep08</a>	<a href="#">307081</a>	6627	704	7	132	pitriysin 1 (Pitrm1) alternative variant cSep08, mRNA.
<a href="#">Pitrm1</a>	<a href="#">Pitrm1.dSep08</a>	<a href="#">307081</a>	1965	779	2	87	pitriysin 1 (Pitrm1) alternative variant dSep08, mRNA.
<a href="#">Pitx1</a>	<a href="#">Pitx1.bSep08</a>	<a href="#">113983</a>	5390	606	1	198	paired-like homeodomain transcription factor 1 (Pitx1) alternative variant bSep08, mRNA.
<a href="#">Pitx3</a>	<a href="#">Pitx3.bSep08</a>	<a href="#">29609</a>	991	597	2	115	paired-like homeodomain transcription factor 3 (Pitx3) alternative variant bSep08, mRNA.
<a href="#">Pitx3</a>	<a href="#">Pitx3.cSep08</a>	<a href="#">29609</a>	4205	334	3	111	paired-like homeodomain transcription factor 3 (Pitx3) alternative variant cSep08, mRNA.
<a href="#">Piwil4</a>	<a href="#">Piwil4.aSep08</a>	<a href="#">689972</a>	16446	2015		245	piwi-like 4 (Drosophila) (Piwil4) mRNA.
<a href="#">Pja2</a>	<a href="#">Pja2.bSep08</a>	<a href="#">192256</a>	19455	600	4	199	pja2, RING-H2 motif containing (Pja2) alternative variant bSep08, mRNA.
<a href="#">Pja2</a>	<a href="#">Pja2.cSep08</a>	<a href="#">192256</a>	24214	700	5	135	pja2, RING-H2 motif containing (Pja2) alternative variant cSep08, mRNA.
<a href="#">PKD.0</a>	<a href="#">PKD.0.aSep08</a>		608	399		93	polycystic kidney disease 1 like (PKD.0) mRNA.
<a href="#">PKD2</a>	<a href="#">PKD2.aSep08</a>	<a href="#">498328</a>	9103	1818	6	295	polycystic kidney disease 2 (PKD2) alternative variant aSep08, mRNA.
<a href="#">PKD2</a>	<a href="#">PKD2.bSep08</a>	<a href="#">498328</a>	2721	1788	3	93	polycystic kidney disease 2 (PKD2) alternative variant bSep08, mRNA.
<a href="#">Pkd2l1</a>	<a href="#">Pkd2l1.bSep08</a>	<a href="#">293937</a>	8547	726	2	198	polycystic kidney disease 2-like 1 (Pkd2l1) alternative variant bSep08, mRNA.
<a href="#">PKD_channel.0</a>	<a href="#">PKD_channel.0.aSep08</a>		3253	2588	3	550	polycystin 1 (PKD_channel.0) alternative variant aSep08, mRNA.
<a href="#">PKD_channel.0</a>	<a href="#">PKD_channel.0.bSep08</a>		1114	1028	1	215	polycystic kidney disease 1 protein like (PKD_channel.0) alternative variant bSep08, mRNA.
<a href="#">Pkia</a>	<a href="#">Pkia.aSep08</a>	<a href="#">114906</a>	15017	3678	3	136	protein kinase inhibitor, alpha (Pkia) alternative variant aSep08, mRNA.

<a href="#">Pkia</a>	<a href="#">Pkia.cSep08</a>	<a href="#">114906</a>	68155	498	3	82	protein kinase inhibitor, alpha (Pkia) alternative variant cSep08, mRNA.
<a href="#">Pkia</a>	<a href="#">Pkia.dSep08</a>	<a href="#">114906</a>	72013	422	3	42	protein kinase inhibitor, alpha (Pkia) alternative variant dSep08, mRNA.
<a href="#">Pkib</a>	<a href="#">Pkib.bSep08</a>	<a href="#">24678</a>	79161	352	5	80	protein kinase inhibitor beta, (cAMP-dependent, catalytic) inhibitor beta (Pkib) alternative variant bSep08, mRNA.
<a href="#">Pkib</a>	<a href="#">Pkib.cSep08</a>	<a href="#">24678</a>	5890	282	3	72	protein kinase inhibitor beta, (cAMP-dependent, catalytic) inhibitor beta (Pkib) alternative variant cSep08, mRNA.
<a href="#">Pkib</a>	<a href="#">Pkib.eSep08</a>	<a href="#">24678</a>	94480	1132	5	78	protein kinase inhibitor beta, (cAMP-dependent, catalytic) inhibitor beta (8.4 kD) (Pkib) alternative variant eSep08, mRNA.
<a href="#">Pkib</a>	<a href="#">Pkib.fSep08</a>	<a href="#">24678</a>	119855	656	6	100	protein kinase inhibitor beta, (cAMP-dependent, catalytic) inhibitor beta (Pkib) alternative variant fSep08, mRNA.
<a href="#">Pkig</a>	<a href="#">Pkig.aSep08</a>	<a href="#">266709</a>	30759	536	3	120	protein kinase inhibitor, gamma (Pkig) alternative variant aSep08, mRNA.
<a href="#">Pkig</a>	<a href="#">Pkig.dSep08</a>	<a href="#">266709</a>	30736	582	4	76	protein kinase inhibitor, gamma (7.9 kD) (Pkig) alternative variant dSep08, mRNA.
<a href="#">Pkig</a>	<a href="#">Pkig.eSep08</a>	<a href="#">266709</a>	66730	536	5	76	protein kinase inhibitor, gamma (7.9 kD) (Pkig) alternative variant eSep08, mRNA.
<a href="#">Pkinase.0</a>	<a href="#">Pkinase.0.aSep08</a>		3960	813	4	270	protein-serine threonine kinase (Pkinase.0) alternative variant aSep08, mRNA.
<a href="#">Pkinase.0</a>	<a href="#">Pkinase.0.bSep08</a>		1544	888	2	244	protein-serine threonine kinase (Pkinase.0) alternative variant bSep08, mRNA.
<a href="#">Pkinase.1</a>	<a href="#">Pkinase.1.aSep08</a>		4424	1020	7	224	CDC-like kinase 4 (Pkinase.1) alternative variant aSep08, mRNA.
<a href="#">Pkinase.1</a>	<a href="#">Pkinase.1.bSep08</a>		2773	852	4	72	CDC like kinase 4 CRA a (Pkinase.1) alternative variant bSep08, mRNA.
<a href="#">Pkinase.1</a>	<a href="#">Pkinase.1.dSep08</a>		693	447	2	60	putative protein (7.0 kD) (Pkinase.1) alternative variant dSep08, mRNA.
<a href="#">Pkinase.2</a>	<a href="#">Pkinase.2.aSep08</a>		5049	2673		790	protein kinase 7 (Pkinase.2) mRNA.
<a href="#">Pkinase.4</a>	<a href="#">Pkinase.4.aSep08</a>		3230	2796		931	titin CRA a (Pkinase.4) mRNA.
<a href="#">Pkinase.5</a>	<a href="#">Pkinase.5.aSep08</a>		26460	2634	10	409	protein kinase, C-terminal and tyrosine protein kinase (Pkinase.5) alternative variant aSep08, mRNA.
<a href="#">Pkinase.5</a>	<a href="#">Pkinase.5.cSep08</a>		1896	552	2	68	putative protein (8.0 kD) (Pkinase.5) alternative variant cSep08, mRNA.
<a href="#">Pkinase_Tyr.0</a>	<a href="#">Pkinase_Tyr.0.aSep08</a>		5942	474	4	158	eph receptor A4 (Pkinase_Tyr.0) alternative variant aSep08, mRNA.
<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.aSep08</a>		10806	2017	10	497	tyrosine kinase 2 (Pkinase_Tyr.1) alternative variant aSep08, mRNA.
<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.bSep08</a>		3812	991	8	303	tyrosine kinase (Pkinase_Tyr.1) alternative variant bSep08, mRNA.
<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.cSep08</a>		4539	1606	4	209	tyrosine kinase 2 CRA a (Pkinase_Tyr.1) alternative variant cSep08, mRNA.
<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.dSep08</a>		1272	422	3	131	tyrosine kinase 2 (Pkinase_Tyr.1) alternative variant dSep08, mRNA.
<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.fSep08</a>		464	388	2	69	tyrosine kinase 2 CRA a (8.1 kD) (Pkinase_Tyr.1) alternative variant fSep08, mRNA.

<a href="#">Pkinase_Tyr.1</a>	<a href="#">Pkinase_Tyr.1.gSep08</a>		3097	457	3	13	putative protein (Pkinase_Tyr.1) alternative variant gSep08, mRNA.
<a href="#">Pklr</a>	<a href="#">Pklr.bSep08</a>	<a href="#">24651</a>	3893	696	5	231	pyruvate kinase, liver and red blood cell (Pklr) alternative variant bSep08, mRNA.
<a href="#">Pklr</a>	<a href="#">Pklr.cSep08</a>	<a href="#">24651</a>	3794	1415	5	202	pyruvate kinase, liver and red blood cell (22.0 kD) (Pklr) alternative variant cSep08, mRNA.
<a href="#">Pkm2</a>	<a href="#">Pkm2.bSep08</a>	<a href="#">25630</a>	19849	2331	12	600	pyruvate kinase, muscle (Pkm2) alternative variant bSep08, mRNA.
<a href="#">Pkm2</a>	<a href="#">Pkm2.cSep08</a>	<a href="#">25630</a>	7455	632	5	181	pyruvate kinase, muscle (Pkm2) alternative variant cSep08, mRNA.
<a href="#">Pkm2</a>	<a href="#">Pkm2.dSep08</a>	<a href="#">25630</a>	2529	1975	2	38	pyruvate kinase, muscle (4.1 kD) (Pkm2) alternative variant dSep08, mRNA.
<a href="#">Pkmyt1</a>	<a href="#">Pkmyt1.bSep08</a>	<a href="#">287101</a>	11956	2719	7	401	protein kinase, membrane associated tyrosine/threonine 1 (44.5 kD) (Pkmyt1) alternative variant bSep08, mRNA.
<a href="#">Pkmyt1</a>	<a href="#">Pkmyt1.cSep08</a>	<a href="#">287101</a>	9558	755	4	251	protein kinase, membrane associated tyrosine/threonine 1 (Pkmyt1) alternative variant cSep08, mRNA.
<a href="#">Pkn1</a>	<a href="#">Pkn1.bSep08</a>	<a href="#">29355</a>	5992	752	8	250	tyrosine protein kinase (Pkn1) alternative variant bSep08, mRNA.
<a href="#">Pkn1</a>	<a href="#">Pkn1.cSep08</a>	<a href="#">29355</a>	2235	1373	6	190	protein kinase and tyrosine protein kinase (Pkn1) alternative variant cSep08, mRNA.
<a href="#">Pkn1</a>	<a href="#">Pkn1.dSep08</a>	<a href="#">29355</a>	7575	2635	9	177	protein kinase, C-terminal (20.0 kD) (Pkn1) alternative variant dSep08, mRNA.
<a href="#">Pkn1</a>	<a href="#">Pkn1.eSep08</a>	<a href="#">29355</a>	1258	882	3	117	putative protein of eukaryotic origin (Pkn1) alternative variant eSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.bSep08</a>	<a href="#">296619</a>	10473	2040	18	679	PKN/rhopilin/rhotekin rho-binding repeat containing protein (Pkn3) alternative variant bSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.cSep08</a>	<a href="#">296619</a>	2866	753	3	186	putative protein (Pkn3) alternative variant cSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.dSep08</a>	<a href="#">296619</a>	8289	1181	5	181	set translocation (20.9 kD) (Pkn3) alternative variant dSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.eSep08</a>	<a href="#">296619</a>	8176	736	6	175	set translocation (Pkn3) alternative variant eSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.fSep08</a>	<a href="#">296619</a>	3022	838	5	164	set translocation (Pkn3) alternative variant fSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.gSep08</a>	<a href="#">296619</a>	3730	1066	8	150	protein kinase N2 CRA d (16.8 kD) (Pkn3) alternative variant gSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.hSep08</a>	<a href="#">296619</a>	6204	527	5	147	SET translocation (Pkn3) alternative variant hSep08, mRNA.
<a href="#">Pkn3</a>	<a href="#">Pkn3.iSep08</a>	<a href="#">296619</a>	2040	359	4	119	SET translocation (Pkn3) alternative variant iSep08, mRNA.
<a href="#">Pknox1</a>	<a href="#">Pknox1.bSep08</a>	<a href="#">294322</a>	11114	787	6	188	pbx/knotted 1 homeobox (Pknox1) alternative variant bSep08, mRNA.
<a href="#">Pkp1</a>	<a href="#">Pkp1.bSep08</a>	<a href="#">304822</a>	4016	603	1	200	plakophilin 1 (Pkp1) alternative variant bSep08, mRNA.
<a href="#">Pkp2</a>	<a href="#">Pkp2.aSep08</a>	<a href="#">287925</a>	31618	2257		378	plakophilin 2 (Pkp2) mRNA.
<a href="#">Pkp3</a>	<a href="#">Pkp3.bSep08</a>	<a href="#">293619</a>	3359	1019	1	339	plakophilin 3 (Pkp3) alternative variant bSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.bSep08</a>	<a href="#">295625</a>	15884	1631	8	411	plakophilin 4 (Pkp4) alternative variant bSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.cSep08</a>	<a href="#">295625</a>	9393	1051	5	256	plakophilin 4 CRA a (Pkp4) alternative variant cSep08, mRNA.

<a href="#">Pkp4</a>	<a href="#">Pkp4.dSep08</a>	<a href="#">295625</a>	10174	513	4	170	plakophilin 4 CRA b (Pkp4) alternative variant dSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.eSep08</a>	<a href="#">295625</a>	7888	377	4	125	plakophilin 4 CRA a (Pkp4) alternative variant eSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.fSep08</a>	<a href="#">295625</a>	89035	611	6	115	plakophilin 4 (Pkp4) alternative variant fSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.gSep08</a>	<a href="#">295625</a>	4665	659	3	97	plakophilin 4 CRA i (Pkp4) alternative variant gSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.hSep08</a>	<a href="#">295625</a>	83783	499	4	96	plakophilin 4 CRA a (Pkp4) alternative variant hSep08, mRNA.
<a href="#">Pkp4</a>	<a href="#">Pkp4.iSep08</a>	<a href="#">295625</a>	33933	522	4	36	plakophilin 4 CRA i (Pkp4) alternative variant iSep08, mRNA.
<a href="#">Pla2g2a</a>	<a href="#">Pla2g2a.aSep08</a>	<a href="#">29692</a>	1951	739	3	156	phospholipase A2, group IIA (platelets, synovial fluid) (Pla2g2a) alternative variant aSep08, mRNA.
<a href="#">Pla2g2a</a>	<a href="#">Pla2g2a.cSep08</a>	<a href="#">29692</a>	2563	606	2	31	phospholipase A2, group IIA (platelets, synovial fluid) (Pla2g2a) alternative variant cSep08, mRNA.
<a href="#">Pla2g2a</a>	<a href="#">Pla2g2a.dSep08</a>	<a href="#">29692</a>	2560	1357	3	32	phospholipase A2, group IIA (platelets, synovial fluid) (Pla2g2a) alternative variant dSep08, mRNA.
<a href="#">Pla2g2c</a>	<a href="#">Pla2g2c.bSep08</a>	<a href="#">29387</a>	975	323	2	64	phospholipase A2, group IIC (Pla2g2c) alternative variant bSep08, mRNA.
<a href="#">Pla2g2e</a>	<a href="#">Pla2g2e.bSep08</a>	<a href="#">298581</a>	5237	750	1	137	phospholipase A2, group IIE (Pla2g2e) alternative variant bSep08, mRNA.
<a href="#">Pla2g4a</a>	<a href="#">Pla2g4a.bSep08</a>	<a href="#">24653</a>	81055	542		127	phospholipase A2, group IVA (cytosolic, calcium-dependent) (Pla2g4a) alternative variant bSep08, mRNA.
<a href="#">Pla2g4b</a>	<a href="#">Pla2g4b.bSep08</a>	<a href="#">311341</a>	2903	739	1	212	phospholipase A2, group IVB (cytosolic) (Pla2g4b) alternative variant bSep08, mRNA.
<a href="#">Pla2g4d</a>	<a href="#">Pla2g4d.aSep08</a>	<a href="#">691905</a>	4869	746		248	phospholipase A2, group IVD (Pla2g4d) mRNA.
<a href="#">Pla2g4e</a>	<a href="#">Pla2g4e.aSep08</a>	<a href="#">296091</a>	3399	180		59	phospholipase A2, group IVE (Pla2g4e) mRNA.
<a href="#">Pla2g5</a>	<a href="#">Pla2g5.aSep08</a>	<a href="#">29354</a>	20257	808	5	137	phospholipase A2, group V (16.0 kD) (Pla2g5) alternative variant aSep08, mRNA.
<a href="#">Pla2g5</a>	<a href="#">Pla2g5.bSep08</a>	<a href="#">29354</a>	64821	749	7	112	phospholipase A2, group V (Pla2g5) alternative variant bSep08, mRNA.
<a href="#">Pla2g5</a>	<a href="#">Pla2g5.cSep08</a>	<a href="#">29354</a>	7239	464	2	37	phospholipase A2, group V (Pla2g5) alternative variant cSep08, mRNA.
<a href="#">Pla2g6</a>	<a href="#">Pla2g6.aSep08</a>	<a href="#">360426</a>	40362	3314	19	831	phospholipase A2, group VI (92.1 kD) (Pla2g6) alternative variant aSep08, mRNA.
<a href="#">Pla2g6</a>	<a href="#">Pla2g6.bSep08</a>	<a href="#">360426</a>	30645	1838	10	457	phospholipase A2, group VI (Pla2g6) alternative variant bSep08, mRNA.
<a href="#">Pla2g6</a>	<a href="#">Pla2g6.cSep08</a>	<a href="#">360426</a>	1126	413	2	77	phospholipase A2, group VI (Pla2g6) alternative variant cSep08, mRNA.
<a href="#">Pla2g6</a>	<a href="#">Pla2g6.eSep08</a>	<a href="#">360426</a>	10096	265	2	69	phospholipase A2, group VI (Pla2g6) alternative variant eSep08, mRNA.
<a href="#">Pla2g12a</a>	<a href="#">Pla2g12a.aSep08</a>	<a href="#">362039</a>	17027	1513	2	192	phospholipase A2, group XIIA (20.9 kD) (Pla2g12a) alternative variant aSep08, complete mRNA.
<a href="#">Pla2g12a</a>	<a href="#">Pla2g12a.bSep08</a>	<a href="#">362039</a>	6755	737	2	119	phospholipase A2, group XIIA (Pla2g12a) alternative variant bSep08, mRNA.
<a href="#">Pla2g12a</a>	<a href="#">Pla2g12a.cSep08</a>	<a href="#">362039</a>	6711	656	2	105	phospholipase A2, group XIIA (Pla2g12a) alternative variant cSep08, mRNA.

<a href="#">Pla2g12b</a>	<a href="#">Pla2g12b.aSep08</a>	<a href="#">367415</a>	18461	1094	4	224	phospholipase A2, group XIIB (Pla2g12b) alternative variant aSep08, mRNA.
<a href="#">Pla2g12b</a>	<a href="#">Pla2g12b.bSep08</a>	<a href="#">367415</a>	2006	216	2	49	phospholipase A2, group XIIB (Pla2g12b) alternative variant bSep08, mRNA.
<a href="#">Pla2r1</a>	<a href="#">Pla2r1.aSep08</a>	<a href="#">295631</a>	11863	730		182	phospholipase A2 receptor 1 (Pla2r1) mRNA.
<a href="#">Plaa</a>	<a href="#">Plaa.bSep08</a>	<a href="#">116645</a>	3209	1241	2	251	phospholipase A2, activating protein (28.0 kD) (Plaa) alternative variant bSep08, mRNA.
<a href="#">plabor</a>	<a href="#">plabor.aSep08</a>		982	334		46	putative protein (5.0 kD) (plabor) mRNA.
<a href="#">PLAC.0</a>	<a href="#">PLAC.0.aSep08</a>		20501	1136	8	315	proprotein convertase subtilisin kexin type 5 (PLAC.0) alternative variant aSep08, mRNA.
<a href="#">Plac1</a>	<a href="#">Plac1.bSep08</a>	<a href="#">317316</a>	138704	539	3	56	placenta-specific 1 (6.3 kD) (Plac1) alternative variant bSep08, mRNA.
<a href="#">Plac8</a>	<a href="#">Plac8.bSep08</a>	<a href="#">360914</a>	8224	482	3	69	placenta-specific 8 (Plac8) alternative variant bSep08, mRNA.
<a href="#">plachy</a>	<a href="#">plachy.aSep08</a>		2539	488		51	putative protein (plachy) mRNA.
<a href="#">pladoy</a>	<a href="#">pladoy.aSep08</a>		2409	400		37	putative protein (pladoy) mRNA.
<a href="#">plafee</a>	<a href="#">plafee.aSep08</a>		1196	184		42	cathepsin Q2 (plafee) mRNA.
<a href="#">plaflu</a>	<a href="#">plaflu.aSep08</a>		3897	1163	3	113	putative endoplasmic reticulum protein, with a transmembrane domain, of mammalian origin (12.4 kD) (plaflu) alternative variant aSep08, mRNA.
<a href="#">plafly</a>	<a href="#">plafly.aSep08</a>		2895	754		95	putative protein (10.7 kD) (plafly) mRNA.
<a href="#">plagar</a>	<a href="#">plagar.aSep08</a>		1765	1673	2	124	putative protein (14.5 kD) (plagar) alternative variant aSep08, mRNA.
<a href="#">Plagl1</a>	<a href="#">Plagl1.aSep08</a>	<a href="#">25157</a>	32704	409		41	pleiomorphic adenoma gene-like 1 (Plagl1) mRNA.
<a href="#">Plagl2</a>	<a href="#">Plagl2.aSep08</a>	<a href="#">296281</a>	3016	327	2	109	pleiomorphic adenoma gene-like 2 (Plagl2) alternative variant aSep08, mRNA.
<a href="#">plaja</a>	<a href="#">plaja.aSep08</a>		68475	434		123	diaphanous homolog 3 (plaja) mRNA.
<a href="#">plajey</a>	<a href="#">plajey.aSep08</a>		1091	420		43	putative protein (5.1 kD) (plajey) mRNA.
<a href="#">plalo</a>	<a href="#">plalo.aSep08</a>		7299	625	2	100	putative protein (plalo) alternative variant aSep08, mRNA.
<a href="#">plalo</a>	<a href="#">plalo.bSep08</a>		4823	411		46	putative protein (4.8 kD) (plalo) alternative variant bSep08, mRNA.
<a href="#">plamee</a>	<a href="#">plamee.aSep08</a>		18503	875	7	274	myosin XVIIIa (plamee) alternative variant aSep08, mRNA.
<a href="#">plamee</a>	<a href="#">plamee.bSep08</a>		40419	731	6	243	myosin XVIIIa (plamee) alternative variant bSep08, mRNA.
<a href="#">plamee</a>	<a href="#">plamee.cSep08</a>		11104	746	4	200	putative protein (plamee) alternative variant cSep08, mRNA.
<a href="#">plapey</a>	<a href="#">plapey.aSep08</a>		3229	296		28	putative protein (plapey) mRNA.
<a href="#">plapor</a>	<a href="#">plapor.aSep08</a>		671	571		78	putative protein (plapor) mRNA.
<a href="#">plarbor</a>	<a href="#">plarbor.aSep08</a>		987	392		35	putative protein (4.3 kD) (plarbor) mRNA.
<a href="#">plarchy</a>	<a href="#">plarchy.aSep08</a>		6188	371		123	retinoblastoma-like 1 (plarchy) mRNA.
<a href="#">plardoy</a>	<a href="#">plardoy.aSep08</a>		4062	740		44	putative protein (4.9 kD) (plardoy) mRNA.
<a href="#">plarfee</a>	<a href="#">plarfee.aSep08</a>		15727	619	7	206	putative protein of eukaryotic origin (plarfee) alternative variant aSep08, mRNA.
<a href="#">plarflu</a>	<a href="#">plarflu.aSep08</a>		11279	295		46	putative protein (5.0 kD) (plarflu) mRNA.
<a href="#">plarfly</a>	<a href="#">plarfly.aSep08</a>		5954	519		173	thrombospondin 2 (plarfly) mRNA.



<a href="#">plarja</a>	<a href="#">plarja.aSep08</a>		12627	725		241	progesterone-induced blocking factor 1 like (plarja) mRNA.
<a href="#">plarjey</a>	<a href="#">plarjey.aSep08</a>		24691	922		306	ubiquitin 34 (plarjey) mRNA.
<a href="#">parlo</a>	<a href="#">parlo.aSep08</a>		10920	299		37	putative protein (parlo) mRNA.
<a href="#">plarmee</a>	<a href="#">plarmee.aSep08</a>		547	407	2	76	putative protein (8.3 kD) (plarmee) alternative variant aSep08, mRNA.
<a href="#">plarmee</a>	<a href="#">plarmee.bSep08</a>		11213	537	3	76	putative protein (8.7 kD) (plarmee) alternative variant bSep08, mRNA.
<a href="#">plarmee</a>	<a href="#">plarmee.cSep08</a>		16035	784	6	76	putative protein (8.7 kD) (plarmee) alternative variant cSep08, mRNA.
<a href="#">plaroy</a>	<a href="#">plaroy.aSep08</a>		26802	884		43	protein CRA a like (plaroy) alternative variant aSep08, mRNA.
<a href="#">plarpey</a>	<a href="#">plarpey.aSep08</a>		839	425		34	putative protein (3.8 kD) (plarpey) mRNA.
<a href="#">plarpor</a>	<a href="#">plarpor.aSep08</a>		3441	468		156	golgi autoantigen golgin subfamily a 4 CRA a like (plarpor) mRNA.
<a href="#">plarroy</a>	<a href="#">plarroy.aSep08</a>		2843	1655	2	184	uncharacterized protein like (plarroy) alternative variant aSep08, mRNA.
<a href="#">plarroy</a>	<a href="#">plarroy.bSep08</a>		8438	185	2	61	putative protein (plarroy) alternative variant bSep08, mRNA.
<a href="#">plarshaw</a>	<a href="#">plarshaw.aSep08</a>		6499	564	1	136	putative protein (plarshaw) alternative variant aSep08, mRNA.
<a href="#">plarshaw</a>	<a href="#">plarshaw.bSep08</a>		71338	333	2	63	putative protein (plarshaw) alternative variant bSep08, mRNA.
<a href="#">plarshee</a>	<a href="#">plarshee.aSep08</a>		9386	361		97	CRA a like (plarshee) mRNA.
<a href="#">plartu</a>	<a href="#">plartu.aSep08</a>		34310	383		50	putative protein (plartu) mRNA.
<a href="#">plarvo</a>	<a href="#">plarvo.aSep08</a>		7955	2862		230	putative protein of eukaryotic origin (plarvo) mRNA.
<a href="#">plarwer</a>	<a href="#">plarwer.aSep08</a>		9355	766		122	putative protein (12.8 kD) (plarwer) mRNA.
<a href="#">plashaw</a>	<a href="#">plashaw.aSep08</a>		4653	855		58	putative protein (6.6 kD) (plashaw) mRNA.
<a href="#">plashee</a>	<a href="#">plashee.aSep08</a>		1768	719		54	putative protein (plashee) mRNA.
<a href="#">Plat</a>	<a href="#">Plat.bSep08</a>	<a href="#">25692</a>	12431	446	6	148	plasminogen activator, tissue (Plat) alternative variant bSep08, mRNA.
<a href="#">Plat</a>	<a href="#">Plat.cSep08</a>	<a href="#">25692</a>	4424	346	4	115	plasminogen activator, tissue (Plat) alternative variant cSep08, mRNA.
<a href="#">platu</a>	<a href="#">platu.aSep08</a>		5847	937		82	putative nuclear protein (9.3 kD) (platu) mRNA.
<a href="#">Plau</a>	<a href="#">Plau.bSep08</a>	<a href="#">25619</a>	2377	800	5	152	plasminogen activator, urokinase (Plau) alternative variant bSep08, mRNA.
<a href="#">Plau</a>	<a href="#">Plau.cSep08</a>	<a href="#">25619</a>	1706	381	5	66	plasminogen activator, urokinase (Plau) alternative variant cSep08, mRNA.
<a href="#">Plaur</a>	<a href="#">Plaur.aSep08</a>	<a href="#">50692</a>	4237	288		84	plasminogen activator, urokinase receptor (Plaur) mRNA.
<a href="#">plavo</a>	<a href="#">plavo.aSep08</a>		4964	434		54	putative protein (5.7 kD) (plavo) mRNA.
<a href="#">plawchy</a>	<a href="#">plawchy.aSep08</a>		3531	327		40	putative protein (plawchy) mRNA.
<a href="#">plawdoy</a>	<a href="#">plawdoy.aSep08</a>		840	751		59	putative protein (6.9 kD) (plawdoy) mRNA.
<a href="#">plawer</a>	<a href="#">plawer.aSep08</a>		2294	338		34	putative protein (plawer) mRNA.
<a href="#">plawfee</a>	<a href="#">plawfee.aSep08</a>		1808	426		56	CRA a like (6.2 kD) (plawfee) mRNA.
<a href="#">plawflu</a>	<a href="#">plawflu.aSep08</a>		563	426		54	putative protein (plawflu) mRNA.

<a href="#">plawfly</a>	<a href="#">plawfly.aSep08</a>		4046	580		118	thrombospondin 2 (plawfly) mRNA.
<a href="#">plawja</a>	<a href="#">plawja.aSep08</a>		61817	820		244	progesterone-induced blocking factor 1 like (29.2 kD) (plawja) mRNA.
<a href="#">plawjey</a>	<a href="#">plawjey.aSep08</a>		8754	997		318	ubiquitin 34 (plawjey) mRNA.
<a href="#">plawlo</a>	<a href="#">plawlo.aSep08</a>		1489	387		128	gcn1 general control of amino-acid synthesis 1-like 1 (plawlo) mRNA.
<a href="#">plawmee</a>	<a href="#">plawmee.aSep08</a>		1761	282		93	CRA a like (plawmee) mRNA.
<a href="#">plawpey</a>	<a href="#">plawpey.aSep08</a>		1005	409		48	putative protein (plawpey) mRNA.
<a href="#">plawpor</a>	<a href="#">plawpor.aSep08</a>		137959	734		244	integrin alpha 9 (plawpor) mRNA.
<a href="#">plawroy</a>	<a href="#">plawroy.aSep08</a>		1296	263		53	putative protein (5.5 kD) (plawroy) mRNA.
<a href="#">plawshaw</a>	<a href="#">plawshaw.aSep08</a>		1191	770		13	putative protein (plawshaw) mRNA.
<a href="#">plawshee</a>	<a href="#">plawshee.aSep08</a>		11819	376		78	putative protein (plawshee) mRNA.
<a href="#">plawtu</a>	<a href="#">plawtu.aSep08</a>		919	221		46	putative protein (plawtu) mRNA.
<a href="#">plawvo</a>	<a href="#">plawvo.aSep08</a>		6537	532		176	F-box protein 10 (plawvo) mRNA.
<a href="#">plawwer</a>	<a href="#">plawwer.aSep08</a>		753	293		80	putative protein (plawwer) mRNA.
<a href="#">Plcb2</a>	<a href="#">Plcb2.aSep08</a>	<a href="#">85240</a>	930	393		87	phospholipase C, beta 2 (Plcb2) mRNA.
<a href="#">Plcb3</a>	<a href="#">Plcb3.bSep08</a>	<a href="#">29322</a>	4044	1771	1	491	phospholipase C, beta 3 (Plcb3) alternative variant bSep08, mRNA.
<a href="#">Plcb4</a>	<a href="#">Plcb4.bSep08</a>	<a href="#">25031</a>	20919	1338	8	242	phospholipase C, beta 4 (Plcb4) alternative variant bSep08, mRNA.
<a href="#">Plcb4</a>	<a href="#">Plcb4.cSep08</a>	<a href="#">25031</a>	149554	440	5	74	phospholipase C, beta 4 (Plcb4) alternative variant cSep08, mRNA.
<a href="#">Plcd1</a>	<a href="#">Plcd1.bSep08</a>	<a href="#">24655</a>	28609	2352	1	783	phospholipase C, delta 1 (88.4 kD) (Plcd1) alternative variant bSep08, mRNA.
<a href="#">Plcd3</a>	<a href="#">Plcd3.bSep08</a>	<a href="#">287745</a>	571	482	2	137	phospholipase C, delta 3 (Plcd3) alternative variant bSep08, mRNA.
<a href="#">Plcd3</a>	<a href="#">Plcd3.cSep08</a>	<a href="#">287745</a>	30448	502	3	100	phospholipase C, delta 3 (Plcd3) alternative variant cSep08, mRNA.
<a href="#">Plcd4</a>	<a href="#">Plcd4.bSep08</a>	<a href="#">140693</a>	7883	790	6	262	phospholipase C, delta 4 (Plcd4) alternative variant bSep08, mRNA.
<a href="#">Plcd4</a>	<a href="#">Plcd4.cSep08</a>	<a href="#">140693</a>	7920	624	5	141	phospholipase C, delta 4 (16.7 kD) (Plcd4) alternative variant cSep08, mRNA.
<a href="#">Plcg1</a>	<a href="#">Plcg1.bSep08</a>	<a href="#">25738</a>	19202	692	7	230	phospholipase C, gamma 1 (Plcg1) alternative variant bSep08, mRNA.
<a href="#">Plcg1</a>	<a href="#">Plcg1.cSep08</a>	<a href="#">25738</a>	1369	396	3	131	phospholipase C, gamma 1 (Plcg1) alternative variant cSep08, mRNA.
<a href="#">Plcg1</a>	<a href="#">Plcg1.dSep08</a>	<a href="#">25738</a>	891	386	3	128	phospholipase C, gamma 1 (Plcg1) alternative variant dSep08, mRNA.
<a href="#">Plcg1</a>	<a href="#">Plcg1.eSep08</a>	<a href="#">25738</a>	890	797	2	91	phospholipase C, gamma 1 (Plcg1) alternative variant eSep08, mRNA.
<a href="#">Plcg2</a>	<a href="#">Plcg2.bSep08</a>	<a href="#">29337</a>	30914	1617	10	436	phospholipase C, gamma 2 (Plcg2) alternative variant bSep08, mRNA.
<a href="#">Plcg2</a>	<a href="#">Plcg2.cSep08</a>	<a href="#">29337</a>	1630	776	1	50	phospholipase C, gamma 2 (5.5 kD) (Plcg2) alternative variant cSep08, mRNA.
<a href="#">Plch1</a>	<a href="#">Plch1.aSep08</a>	<a href="#">310463</a>	2718	356		118	phospholipase C, eta 1 (Plch1) mRNA.

<a href="#">Plch2</a>	<a href="#">Plch2.aSep08</a>	<a href="#">313756</a>	1174	774		206	phospholipase C, eta 2 (Plch2) mRNA.
<a href="#">Plcz1</a>	<a href="#">Plcz1.bSep08</a>	<a href="#">497197</a>	30832	1058	5	307	phospholipase C, zeta 1 (Plcz1) alternative variant bSep08, mRNA.
<a href="#">Plcz1</a>	<a href="#">Plcz1.cSep08</a>	<a href="#">497197</a>	13513	693	4	231	phospholipase C, zeta 1 (Plcz1) alternative variant cSep08, mRNA.
<a href="#">Plcz1</a>	<a href="#">Plcz1.dSep08</a>	<a href="#">497197</a>	17245	750	5	188	phospholipase C, zeta 1 (Plcz1) alternative variant dSep08, mRNA.
<a href="#">Pld1</a>	<a href="#">Pld1.aSep08</a>	<a href="#">25096</a>	57940	2555		355	phospholipase D1 (Pld1) mRNA.
<a href="#">Pld3</a>	<a href="#">Pld3.bSep08</a>	<a href="#">361527</a>	14688	1143	8	307	phospholipase D family, member 3 (Pld3) alternative variant bSep08, mRNA.
<a href="#">Pld3</a>	<a href="#">Pld3.cSep08</a>	<a href="#">361527</a>	12135	785	7	224	phospholipase D family, member 3 (Pld3) alternative variant cSep08, mRNA.
<a href="#">Pld3</a>	<a href="#">Pld3.dSep08</a>	<a href="#">361527</a>	3729	605	6	169	phospholipase D family, member 3 (Pld3) alternative variant dSep08, mRNA.
<a href="#">Pld5</a>	<a href="#">Pld5.aSep08</a>	<a href="#">289270</a>	111943	658		219	phospholipase D family, member 5 (Pld5) mRNA.
<a href="#">PLDc.0</a>	<a href="#">PLDc.0.aSep08</a>		27520	1253		397	phospholipase D1 (PLDc.0) mRNA.
<a href="#">Pldn</a>	<a href="#">Pldn.bSep08</a>	<a href="#">317630</a>	8552	659	5	158	pallidin (Pldn) alternative variant bSep08, mRNA.
<a href="#">Plec1</a>	<a href="#">Plec1.bSep08</a>	<a href="#">64204</a>	5274	410	2	58	plectin 1 (Plec1) alternative variant bSep08, mRNA.
<a href="#">Plectin.0</a>	<a href="#">Plectin.0.aSep08</a>		2375	1310	1	436	desmoplakin CRA b (Plectin.0) alternative variant aSep08, mRNA.
<a href="#">Plectin.0</a>	<a href="#">Plectin.0.bSep08</a>		3431	569	1	189	desmoplakin II (Plectin.0) alternative variant bSep08, mRNA.
<a href="#">pleebor</a>	<a href="#">pleebor.aSep08</a>		115700	1671		184	catenin Alpha-3 (pleebor) mRNA.
<a href="#">pleechy</a>	<a href="#">pleechy.aSep08</a>		7183	836		278	putative protein (pleechy) mRNA.
<a href="#">pleedoy</a>	<a href="#">pleedoy.aSep08</a>		11672	357		83	putative protein (pleedoy) mRNA.
<a href="#">pleeflu</a>	<a href="#">pleeflu.aSep08</a>		1757	276		46	putative protein (pleeflu) mRNA.
<a href="#">pleefly</a>	<a href="#">pleefly.aSep08</a>		3088	763		84	putative secreted or extracellular protein precursor (9.0 kD) (pleefly) mRNA.
<a href="#">pleeja</a>	<a href="#">pleeja.aSep08</a>		1285	858		52	putative protein (pleeja) mRNA.
<a href="#">pleejey</a>	<a href="#">pleejey.aSep08</a>		8721	394		119	ubiquitin 34 (pleejey) mRNA.
<a href="#">pleelo</a>	<a href="#">pleelo.aSep08</a>		7214	622		207	gcn1 1 (pleelo) mRNA.
<a href="#">pleemee</a>	<a href="#">pleemee.aSep08</a>		342	191		63	CRA a like (pleemee) mRNA.
<a href="#">pleepey</a>	<a href="#">pleepey.aSep08</a>		1915	735		72	putative protein (7.8 kD) (pleepey) mRNA.
<a href="#">pleepor</a>	<a href="#">pleepor.aSep08</a>		1358	633		31	putative protein (pleepor) mRNA.
<a href="#">pleeroy</a>	<a href="#">pleeroy.aSep08</a>		6252	893		92	putative protein (10.2 kD) (pleeroy) mRNA.
<a href="#">pleeshaw</a>	<a href="#">pleeshaw.aSep08</a>		7688	468		34	putative protein (pleeshaw) mRNA.
<a href="#">pleeshee</a>	<a href="#">pleeshee.aSep08</a>		1602	196		65	centromere protein E CRA b (pleeshee) mRNA.
<a href="#">pleetu</a>	<a href="#">pleetu.aSep08</a>		685	384		53	putative protein (pleetu) mRNA.
<a href="#">pleevo</a>	<a href="#">pleevo.aSep08</a>		5150	643		189	putative protein of metazoan origin (pleevo) mRNA.
<a href="#">pleewer</a>	<a href="#">pleewer.aSep08</a>		687	262		39	putative protein (pleewer) mRNA.
<a href="#">Plek2</a>	<a href="#">Plek2.bSep08</a>	<a href="#">314260</a>	1744	406	2	83	pleckstrin 2 (Plek2) alternative variant bSep08, mRNA.
<a href="#">Plek2</a>	<a href="#">Plek2.cSep08</a>	<a href="#">314260</a>	1280	656	1	36	pleckstrin 2 (3.9 kD) (Plek2) alternative variant cSep08, mRNA.

<a href="#">Plekha1</a>	<a href="#">Plekha1.bSep08</a>	<a href="#">361659</a>	33180	730	6	203	pleckstrin-like (23.2 kD) (Plekha1) alternative variant bSep08, complete mRNA.
<a href="#">Plekha3</a>	<a href="#">Plekha3.cSep08</a>	<a href="#">295674</a>	6776	622	2	45	pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 3 (5.2 kD) (Plekha3) alternative variant cSep08, mRNA.
<a href="#">Plekha4</a>	<a href="#">Plekha4.aSep08</a>	<a href="#">308584</a>	18967	2348	17	715	pleckstrin-like (Plekha4) alternative variant aSep08, mRNA.
<a href="#">Plekha4</a>	<a href="#">Plekha4.cSep08</a>	<a href="#">308584</a>	5218	650	5	216	pleckstrin-like (Plekha4) alternative variant cSep08, mRNA.
<a href="#">Plekha4</a>	<a href="#">Plekha4.dSep08</a>	<a href="#">308584</a>	4969	777	4	126	putative protein of mammalian origin (Plekha4) alternative variant dSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.aSep08</a>	<a href="#">246237</a>	168773	4046	26	1077	WW/Rsp5/WWP and pleckstrin-like (Plekha5) alternative variant aSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.bSep08</a>	<a href="#">246237</a>	18881	1348	5	378	putative protein of vertebrate origin (42.8 kD) (Plekha5) alternative variant bSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.cSep08</a>	<a href="#">246237</a>	112574	851	8	283	pleckstrin-like (Plekha5) alternative variant cSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.dSep08</a>	<a href="#">246237</a>	28261	739	8	246	pleckstrin-like (Plekha5) alternative variant dSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.eSep08</a>	<a href="#">246237</a>	10178	919	6	237	putative protein of vertebrate origin (Plekha5) alternative variant eSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.fSep08</a>	<a href="#">246237</a>	99646	700	7	232	pleckstrin-like (Plekha5) alternative variant fSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.gSep08</a>	<a href="#">246237</a>	18458	876	6	215	pleckstrin-like (Plekha5) alternative variant gSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.hSep08</a>	<a href="#">246237</a>	105784	795	8	212	pleckstrin-like (Plekha5) alternative variant hSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.iSep08</a>	<a href="#">246237</a>	68273	727	8	184	pleckstrin-like (Plekha5) alternative variant iSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.jSep08</a>	<a href="#">246237</a>	11401	866	5	134	putative protein, with a coiled coil domain, of vertebrate origin (Plekha5) alternative variant jSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.kSep08</a>	<a href="#">246237</a>	15240	379	6	126	phosphoinositol 3-phosphate-binding like (Plekha5) alternative variant kSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.lSep08</a>	<a href="#">246237</a>	7847	427	5	90	putative protein, with a coiled coil domain, of vertebrate origin (Plekha5) alternative variant lSep08, mRNA.
<a href="#">Plekha5</a>	<a href="#">Plekha5.mSep08</a>	<a href="#">246237</a>	2703	737	2	118	putative protein (13.2 kD) (Plekha5) alternative variant mSep08, mRNA.
<a href="#">Plekha6</a>	<a href="#">Plekha6.aSep08</a>	<a href="#">360842</a>	109725	1746	10	434	WW/Rsp5/WWP and pleckstrin-like (Plekha6) alternative variant aSep08, mRNA.
<a href="#">Plekha6</a>	<a href="#">Plekha6.bSep08</a>	<a href="#">360842</a>	92627	943	8	195	pleckstrin-like (Plekha6) alternative variant bSep08, mRNA.
<a href="#">Plekha6</a>	<a href="#">Plekha6.cSep08</a>	<a href="#">360842</a>	86096	299	3	45	putative protein (Plekha6) alternative variant cSep08, mRNA.
<a href="#">Plekhb1</a>	<a href="#">Plekhb1.bSep08</a>	<a href="#">64471</a>	11839	824	6	188	pleckstrin-like (21.3 kD) (Plekhb1) alternative variant bSep08, mRNA.
<a href="#">Plekhb1</a>	<a href="#">Plekhb1.cSep08</a>	<a href="#">64471</a>	13777	1307	7	188	pleckstrin-like (21.3 kD) (Plekhb1) alternative variant cSep08, mRNA.
<a href="#">Plekhb1</a>	<a href="#">Plekhb1.dSep08</a>	<a href="#">64471</a>	7487	454	5	111	putative protein (Plekhb1) alternative variant dSep08, mRNA.
<a href="#">Plekhb1</a>	<a href="#">Plekhb1.fSep08</a>	<a href="#">64471</a>	5154	411	3	77	putative nuclear protein (8.3 kD) (Plekhb1) alternative variant fSep08, mRNA.
<a href="#">Plekhb2</a>	<a href="#">Plekhb2.bSep08</a>	<a href="#">301337</a>	21108	484	1	136	putative protein of metazoan origin (Plekhb2) alternative variant bSep08, mRNA.
<a href="#">Plekhg2</a>	<a href="#">Plekhg2.aSep08</a>	<a href="#">292750</a>	9993	3612	13	492	DH and pleckstrin-like (54.3 kD) (Plekhg2) alternative variant aSep08, mRNA.

<a href="#">Plekhg2</a>	<a href="#">Plekhg2.bSep08</a>	<a href="#">292750</a>	2446	867	3	289	putative protein of ancient origin (Plekhg2) alternative variant bSep08, mRNA.
<a href="#">Plekhg3</a>	<a href="#">Plekhg3.aSep08</a>	<a href="#">314249</a>	24948	396		131	putative protein of mammalian origin (Plekhg3) mRNA.
<a href="#">Plekhg4</a>	<a href="#">Plekhg4.aSep08</a>	<a href="#">307796</a>	1080	894		12	putative protein (Plekhg4) mRNA.
<a href="#">Plekhh1</a>	<a href="#">Plekhh1.bSep08</a>	<a href="#">314262</a>	4765	982	3	280	putative protein of bilaterial origin (Plekhh1) alternative variant bSep08, mRNA.
<a href="#">Plekhh1</a>	<a href="#">Plekhh1.cSep08</a>	<a href="#">314262</a>	6280	1476	7	280	putative protein of eukaryotic origin (Plekhh1) alternative variant cSep08, mRNA.
<a href="#">Plekhh1</a>	<a href="#">Plekhh1.dSep08</a>	<a href="#">314262</a>	2162	344	3	114	putative protein of eukaryotic origin (Plekhh1) alternative variant dSep08, mRNA.
<a href="#">Plekhh2</a>	<a href="#">Plekhh2.aSep08</a>	<a href="#">313866</a>	20637	1219		406	pleckstrin-like and unconventional myosin/plant kinesin-like protein/non-motor protein conserved region MyTH4 (Plekhh2) mRNA.
<a href="#">Plekhh3</a>	<a href="#">Plekhh3.bSep08</a>	<a href="#">360634</a>	1230	712	4	130	putative protein of eukaryotic origin (Plekhh3) alternative variant bSep08, mRNA.
<a href="#">Plekhh3</a>	<a href="#">Plekhh3.cSep08</a>	<a href="#">360634</a>	1477	407	3	104	putative protein of mammalian origin (Plekhh3) alternative variant cSep08, mRNA.
<a href="#">Plekhj1</a>	<a href="#">Plekhj1.bSep08</a>	<a href="#">314634</a>	746	644	2	125	putative protein of vertebrate origin (Plekhj1) alternative variant bSep08, mRNA.
<a href="#">Plekhj1</a>	<a href="#">Plekhj1.cSep08</a>	<a href="#">314634</a>	2139	1043	3	111	putative cytoplasmic protein of vertebrate origin (12.2 kD) (Plekhj1) alternative variant cSep08, complete mRNA.
<a href="#">Plekhj1</a>	<a href="#">Plekhj1.dSep08</a>	<a href="#">314634</a>	3499	1286	2	31	putative protein (3.6 kD) (Plekhj1) alternative variant dSep08, mRNA.
<a href="#">Plekhm2</a>	<a href="#">Plekhm2.aSep08</a>	<a href="#">313667</a>	5944	2478	12	571	pleckstrin-like (Plekhm2) alternative variant aSep08, mRNA.
<a href="#">Plekhm2</a>	<a href="#">Plekhm2.cSep08</a>	<a href="#">313667</a>	849	772	2	93	putative protein of vertebrate origin (Plekhm2) alternative variant cSep08, mRNA.
<a href="#">Plekhm2</a>	<a href="#">Plekhm2.dSep08</a>	<a href="#">313667</a>	995	376	3	77	putative protein of vertebrate origin (Plekhm2) alternative variant dSep08, mRNA.
<a href="#">Plekhm3</a>	<a href="#">Plekhm3.aSep08</a>	<a href="#">316455</a>	30689	1628		115	putative protein of eukaryotic origin (13.1 kD) (Plekhm3) alternative variant aSep08, mRNA.
<a href="#">Plekho1</a>	<a href="#">Plekho1.cSep08</a>	<a href="#">310674</a>	5891	653	4	160	pleckstrin-like (Plekho1) alternative variant cSep08, mRNA.
<a href="#">plerbor</a>	<a href="#">plerbor.aSep08</a>		14296	1226	4	154	CRA b (plerbor) alternative variant aSep08, mRNA.
<a href="#">plerchy</a>	<a href="#">plerchy.aSep08</a>		13110	479		159	putative protein of vertebrate origin (plerchy) mRNA.
<a href="#">plerdoy</a>	<a href="#">plerdoy.aSep08</a>		3537	1137	3	110	CRA b like (12.4 kD) (plerdoy) mRNA.
<a href="#">plerfee</a>	<a href="#">plerfee.aSep08</a>		3545	425		39	putative protein (4.6 kD) (plerfee) mRNA.
<a href="#">plerflu</a>	<a href="#">plerflu.aSep08</a>		5081	616		205	putative protein of eukaryotic origin (plerflu) mRNA.
<a href="#">plerfly</a>	<a href="#">plerfly.bSep08</a>		61321	766	5	110	putative protein (plerfly) alternative variant bSep08, mRNA.
<a href="#">plerfly</a>	<a href="#">plerfly.cSep08</a>		61289	330	2	39	wd repeat domain 27 CRA c (plerfly) alternative variant cSep08, mRNA.
<a href="#">plerja</a>	<a href="#">plerja.aSep08</a>		4611	632		57	putative protein (6.6 kD) (plerja) mRNA.
<a href="#">plerjey</a>	<a href="#">plerjey.aSep08</a>		16500	641		213	ubiquitin 34 (plerjey) mRNA.
<a href="#">plerlo</a>	<a href="#">plerlo.aSep08</a>		1058	387		19	putative protein (2.1 kD) (plerlo) mRNA.
<a href="#">plermee</a>	<a href="#">plermee.aSep08</a>		1050	322		107	CRA a like (plermee) mRNA.
<a href="#">plerpey</a>	<a href="#">plerpey.aSep08</a>		1458	336		32	putative protein (3.4 kD) (plerpey) mRNA.

<a href="#">plerpor</a>	<a href="#">plerpor.aSep08</a>		3462	741		98	deleted in lung esophageal cancer 1 like (plerpor) mRNA.
<a href="#">plerroy</a>	<a href="#">plerroy.aSep08</a>		3316	416		36	putative protein (plerroy) mRNA.
<a href="#">plershaw</a>	<a href="#">plershaw.aSep08</a>		6140	750		126	putative protein (plershaw) mRNA.
<a href="#">plershee</a>	<a href="#">plershee.aSep08</a>		1497	370		122	centromere protein E CRA a (plershee) mRNA.
<a href="#">plertu</a>	<a href="#">plertu.aSep08</a>		39272	726	3	63	putative protein (6.7 kD) (plertu) alternative variant aSep08, mRNA.
<a href="#">plertu</a>	<a href="#">plertu.bSep08</a>		29597	487	2	49	putative protein (5.7 kD) (plertu) alternative variant bSep08, mRNA.
<a href="#">plervo</a>	<a href="#">plervo.aSep08</a>		4272	737		177	putative protein, with at least 2 transmembrane domains (plervo) mRNA.
<a href="#">plerwer</a>	<a href="#">plerwer.aSep08</a>		2849	906		77	putative protein (9.1 kD) (plerwer) mRNA.
<a href="#">Plexin_cytopl.0</a>	<a href="#">Plexin_cytopl.0.aSep08</a>		2295	587	4	195	plexin A1 CRA c (Plexin_cytopl.0) alternative variant aSep08, mRNA.
<a href="#">pleybor</a>	<a href="#">pleybor.aSep08</a>		4383	920		55	putative protein (pleybor) mRNA.
<a href="#">pleychy</a>	<a href="#">pleychy.aSep08</a>		2764	573		35	putative protein (pleychy) mRNA.
<a href="#">pleydoy</a>	<a href="#">pleydoy.aSep08</a>		4145	499		52	putative protein (pleydoy) mRNA.
<a href="#">pleyfee</a>	<a href="#">pleyfee.aSep08</a>		823	285		33	putative protein (pleyfee) mRNA.
<a href="#">pleyflu</a>	<a href="#">pleyflu.aSep08</a>		4089	1555		54	putative protein (5.9 kD) (pleyflu) mRNA.
<a href="#">pleyfly</a>	<a href="#">pleyfly.aSep08</a>		7295	2150	2	288	putative cytoplasmic protein of eukaryotic origin (32.1 kD) (pleyfly) alternative variant aSep08, complete mRNA.
<a href="#">pleyfly</a>	<a href="#">pleyfly.bSep08</a>		5364	785	1	202	putative cytoplasmic protein of eukaryotic origin (22.4 kD) (pleyfly) alternative variant bSep08, mRNA.
<a href="#">pleyja</a>	<a href="#">pleyja.aSep08</a>		16267	604		201	tbc1 domain family member 4 (pleyja) mRNA.
<a href="#">pleyjej</a>	<a href="#">pleyjej.aSep08</a>		5603	670	6	223	ubiquitin 34 (pleyjej) alternative variant aSep08, mRNA.
<a href="#">pleylo</a>	<a href="#">pleylo.aSep08</a>		1823	327	3	63	putative protein (6.7 kD) (pleylo) alternative variant aSep08, mRNA.
<a href="#">pleymee</a>	<a href="#">pleymee.aSep08</a>		12006	435		115	putative protein (pleymee) mRNA.
<a href="#">pleypey</a>	<a href="#">pleypey.bSep08</a>		2923	713	3	68	CRA a like (pleypey) alternative variant bSep08, mRNA.
<a href="#">pleypey</a>	<a href="#">pleypey.cSep08</a>		2876	631	3	76	putative protein (pleypey) alternative variant cSep08, mRNA.
<a href="#">pleypor</a>	<a href="#">pleypor.aSep08</a>		2257	1940		73	deleted in lung esophageal cancer 1 like (8.3 kD) (pleypor) mRNA.
<a href="#">pleyro</a>	<a href="#">pleyro.aSep08</a>		7968	658		151	stabilin 2 (pleyro) mRNA.
<a href="#">pleyshaw</a>	<a href="#">pleyshaw.aSep08</a>		3384	571		66	putative protein (pleyshaw) mRNA.
<a href="#">pleyshee</a>	<a href="#">pleyshee.aSep08</a>		7634	1559	6	519	centromere protein E (pleyshee) alternative variant aSep08, mRNA.
<a href="#">pleyshee</a>	<a href="#">pleyshee.bSep08</a>		629	397	1	132	centromere protein e (pleyshee) alternative variant bSep08, mRNA.
<a href="#">pleytu</a>	<a href="#">pleytu.aSep08</a>		1137	647		59	putative protein (6.4 kD) (pleytu) mRNA.
<a href="#">pleyvo</a>	<a href="#">pleyvo.aSep08</a>		1046	789		53	putative protein (5.6 kD) (pleyvo) mRNA.
<a href="#">pleywer</a>	<a href="#">pleywer.aSep08</a>		3125	446		49	putative protein (5.7 kD) (pleywer) mRNA.
<a href="#">Plg</a>	<a href="#">Plg.aSep08</a>	<a href="#">85253</a>	42718	2749		812	plasminogen (90.5 kD) (Plg) mRNA.
<a href="#">Plk1</a>	<a href="#">Plk1.bSep08</a>	<a href="#">25515</a>	1928	684	3	227	polo-like kinase 1 (Drosophila) (Plk1) alternative variant bSep08, mRNA.

<a href="#">Plk1</a>	<a href="#">Plk1.cSep08</a>	<a href="#">25515</a>	1726	743	1	169	polo-like kinase 1 (Drosophila) (Plk1) alternative variant cSep08, mRNA.
<a href="#">Plk3</a>	<a href="#">Plk3.aSep08</a>	<a href="#">58936</a>	2667	1339		334	polo-like kinase 3 (Drosophila) (Plk3) mRNA.
<a href="#">Plk4</a>	<a href="#">Plk4.aSep08</a>	<a href="#">310344</a>	19088	3419	16	924	polo-like kinase 4 (Drosophila) (103.8 kD) (Plk4) alternative variant aSep08, complete mRNA.
<a href="#">Plk4</a>	<a href="#">Plk4.bSep08</a>	<a href="#">310344</a>	3089	557	3	119	polo-like kinase 4 (Drosophila) (Plk4) alternative variant bSep08, mRNA.
<a href="#">Plip</a>	<a href="#">Plip.bSep08</a>	<a href="#">64364</a>	6442	380	1	49	plasma membrane proteolipid (Plip) alternative variant bSep08, mRNA.
<a href="#">Pln</a>	<a href="#">Pln.bSep08</a>	<a href="#">64672</a>	2986	281	1	22	phospholamban (Pln) alternative variant bSep08, mRNA.
<a href="#">plobor</a>	<a href="#">plobor.aSep08</a>		3344	634		113	putative protein, with a coiled coil domain, of metazoan origin (plobor) mRNA.
<a href="#">plochy</a>	<a href="#">plochy.bSep08</a>		4164	371	3	63	putative protein (7.1 kD) (plochy) alternative variant bSep08, mRNA.
<a href="#">Plod1</a>	<a href="#">Plod1.aSep08</a>	<a href="#">116552</a>	19985	3257		574	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1 (Plod1) mRNA.
<a href="#">Plod2</a>	<a href="#">Plod2.bSep08</a>	<a href="#">300901</a>	33767	1782	2	184	procollagen lysine, 2-oxoglutarate 5-dioxygenase 2 (21.5 kD) (Plod2) alternative variant bSep08, mRNA.
<a href="#">Plod3</a>	<a href="#">Plod3.bSep08</a>	<a href="#">288583</a>	1517	744	6	248	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (Plod3) alternative variant bSep08, mRNA.
<a href="#">Plod3</a>	<a href="#">Plod3.cSep08</a>	<a href="#">288583</a>	1145	652	5	114	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (Plod3) alternative variant cSep08, mRNA.
<a href="#">Plod3</a>	<a href="#">Plod3.dSep08</a>	<a href="#">288583</a>	527	386	2	98	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (Plod3) alternative variant dSep08, mRNA.
<a href="#">plodoy</a>	<a href="#">plodoy.aSep08</a>		21188	697		61	putative protein (plodoy) mRNA.
<a href="#">plofee</a>	<a href="#">plofee.aSep08</a>		7137	844		242	putative protein of vertebrate origin (plofee) mRNA.
<a href="#">ploflu</a>	<a href="#">ploflu.aSep08</a>		4273	430		47	putative protein (5.2 kD) (ploflu) mRNA.
<a href="#">plofly</a>	<a href="#">plofly.aSep08</a>		14228	818		102	putative protein of mammalian origin (11.4 kD) (plofly) mRNA.
<a href="#">ploja</a>	<a href="#">ploja.bSep08</a>		745	222		28	putative protein (3.3 kD) (ploja) alternative variant bSep08, mRNA.
<a href="#">plojey</a>	<a href="#">plojey.aSep08</a>		1586	1051		37	putative protein (4.2 kD) (plojey) mRNA.
<a href="#">plolo</a>	<a href="#">plolo.aSep08</a>		474	367		116	putative protein (plolo) mRNA.
<a href="#">plomee</a>	<a href="#">plomee.aSep08</a>		24043	529		175	seizure related 6 (plomee) mRNA.
<a href="#">plopey</a>	<a href="#">plopey.aSep08</a>		27392	248		36	putative protein (plopey) mRNA.
<a href="#">plopor</a>	<a href="#">plopor.aSep08</a>		3728	356		118	golgi autoantigen golgin subfamily a 4 CRA a like (plopor) mRNA.
<a href="#">plorbor</a>	<a href="#">plorbor.bSep08</a>		2392	355	3	70	putative protein (plorbor) alternative variant bSep08, mRNA.
<a href="#">plorchy</a>	<a href="#">plorchy.aSep08</a>		756	662		71	putative secreted or extracellular protein precursor (7.9 kD) (plorchy) mRNA.
<a href="#">plordoy</a>	<a href="#">plordoy.aSep08</a>		8069	2195		114	guanine nucleotide binding protein alpha olfactory type like (plordoy) mRNA.
<a href="#">plorfee</a>	<a href="#">plorfee.aSep08</a>		6350	858		39	putative protein (plorfee) mRNA.
<a href="#">plorflu</a>	<a href="#">plorflu.aSep08</a>		30634	856	2	77	putative protein (8.3 kD) (plorflu) alternative variant aSep08, mRNA.

<a href="#">plorfly</a>	<a href="#">plorfly.aSep08</a>		16941	376		46	putative protein (plorfly) mRNA.
<a href="#">plorja</a>	<a href="#">plorja.aSep08</a>		2237	212		70	tbc1 domain family member 4 (plorja) mRNA.
<a href="#">plorjey</a>	<a href="#">plorjey.aSep08</a>		10902	1261	10	339	ubiquitin 34 (plorjey) alternative variant aSep08, mRNA.
<a href="#">plorjey</a>	<a href="#">plorjey.bSep08</a>		7599	1092	8	236	ubiquitin 34 (plorjey) alternative variant bSep08, mRNA.
<a href="#">plorlo</a>	<a href="#">plorlo.aSep08</a>		3031	473		157	trichoplein keratin filament binding CRA b like (plorlo) mRNA.
<a href="#">plormee</a>	<a href="#">plormee.aSep08</a>		2502	1108		368	suppressor of Ty 6 homolog (plormee) mRNA.
<a href="#">ploroy</a>	<a href="#">ploroy.aSep08</a>		13093	403	3	133	aldehyde dehydrogenase 1 family member L2 CRA a (ploroy) alternative variant aSep08, mRNA.
<a href="#">plorpey</a>	<a href="#">plorpey.aSep08</a>		17906	497		84	putative protein (9.0 kD) (plorpey) mRNA.
<a href="#">plorpor</a>	<a href="#">plorpor.aSep08</a>		1722	665		153	putative protein (plorpor) mRNA.
<a href="#">plorroy</a>	<a href="#">plorroy.aSep08</a>		1860	772		48	putative protein (plorroy) mRNA.
<a href="#">plorshaw</a>	<a href="#">plorshaw.aSep08</a>		15184	270		47	putative protein (plorshaw) mRNA.
<a href="#">plorshee</a>	<a href="#">plorshee.aSep08</a>		5786	552		49	putative protein (plorshee) mRNA.
<a href="#">plortu</a>	<a href="#">plortu.aSep08</a>		12043	577		79	vesicle transport through interaction with t-snares homolog 1b (9.0 kD) (plortu) alternative variant aSep08, mRNA.
<a href="#">plortu</a>	<a href="#">plortu.bSep08</a>		12055	672	1	62	vesicle transport through interaction with t-snares 1b (plortu) alternative variant bSep08, mRNA.
<a href="#">plorvo</a>	<a href="#">plorvo.aSep08</a>		1977	554		46	putative protein (plorvo) mRNA.
<a href="#">plorwer</a>	<a href="#">plorwer.aSep08</a>		630	263		87	CRA a (plorwer) mRNA.
<a href="#">ploshaw</a>	<a href="#">ploshaw.aSep08</a>		44137	318		26	putative protein (2.9 kD) (ploshaw) mRNA.
<a href="#">ploshee</a>	<a href="#">ploshee.aSep08</a>		3544	826		133	putative protein of eukaryotic origin (14.8 kD) (ploshee) mRNA.
<a href="#">plotu</a>	<a href="#">plotu.aSep08</a>		2955	767		38	putative protein (4.4 kD) (plotu) mRNA.
<a href="#">plovo</a>	<a href="#">plovo.aSep08</a>		1530	383	2	92	putative protein (plovo) alternative variant aSep08, mRNA.
<a href="#">plower</a>	<a href="#">plower.aSep08</a>		1175	723		67	putative protein (8.1 kD) (plower) mRNA.
<a href="#">ploybor</a>	<a href="#">ploybor.aSep08</a>		12612	815		65	alpha 1 type XIII collagen (6.4 kD) (ploybor) mRNA.
<a href="#">ploychy</a>	<a href="#">ploychy.aSep08</a>		7678	564		188	uncharacterized protein homolog (ploychy) mRNA.
<a href="#">ploydoy</a>	<a href="#">ploydoy.aSep08</a>		5051	1419	2	150	centrosomal protein 76 (ploydoy) alternative variant aSep08, mRNA.
<a href="#">ploydoy</a>	<a href="#">ploydoy.bSep08</a>		6516	360	2	57	centrosomal protein 76 (ploydoy) alternative variant bSep08, mRNA.
<a href="#">ployfee</a>	<a href="#">ployfee.aSep08</a>		8281	354		117	putative protein (ployfee) mRNA.
<a href="#">ployflu</a>	<a href="#">ployflu.aSep08</a>		3000	332		87	putative protein (ployflu) mRNA.
<a href="#">ployfly</a>	<a href="#">ployfly.aSep08</a>		33639	701	2	54	putative protein (ployfly) alternative variant aSep08, mRNA.
<a href="#">ployfly</a>	<a href="#">ployfly.bSep08</a>		10244	637	1	72	putative protein (ployfly) alternative variant bSep08, mRNA.
<a href="#">ployja</a>	<a href="#">ployja.aSep08</a>		28856	663		48	putative protein (ployja) mRNA.
<a href="#">ployjey</a>	<a href="#">ployjey.aSep08</a>		14371	2133	7	152	putative protein (ployjey) alternative variant aSep08, mRNA.
<a href="#">ployjey</a>	<a href="#">ployjey.bSep08</a>		5547	267	3	68	putative protein (ployjey) alternative variant bSep08, mRNA.
<a href="#">ploylo</a>	<a href="#">ploylo.aSep08</a>		7339	1024	2	341	trichoplein keratin filament binding like (ploylo) alternative variant aSep08, mRNA.



<a href="#">poylo</a>	<a href="#">poylo.bSep08</a>		6392	581	1	193	trichoplein keratin filament binding CRA a like (poylo) alternative variant bSep08, mRNA.
<a href="#">ploymee</a>	<a href="#">ploymee.aSep08</a>		2945	727	3	92	putative protein (ploymee) alternative variant aSep08, mRNA.
<a href="#">ploypey</a>	<a href="#">ploypey.aSep08</a>		699	423		85	putative protein (ploypey) mRNA.
<a href="#">ploypor</a>	<a href="#">ploypor.aSep08</a>		3199	701	3	190	putative protein (ploypor) alternative variant aSep08, mRNA.
<a href="#">poyroy</a>	<a href="#">poyroy.aSep08</a>		31274	642		214	CRA b (poyroy) mRNA.
<a href="#">poyshaw</a>	<a href="#">poyshaw.aSep08</a>		9427	845		47	single-stranded DNA binding protein 2 like (poyshaw) mRNA.
<a href="#">ployshee</a>	<a href="#">ployshee.aSep08</a>		4437	549		94	putative protein (10.4 kD) (ployshee) mRNA.
<a href="#">poytu</a>	<a href="#">poytu.aSep08</a>		3341	738		69	putative protein (poytu) mRNA.
<a href="#">poyvo</a>	<a href="#">poyvo.aSep08</a>		7589	565		154	putative protein (poyvo) mRNA.
<a href="#">poywer</a>	<a href="#">poywer.aSep08</a>		17888	738		80	putative cytoplasmic protein (8.8 kD) (poywer) mRNA.
<a href="#">Plp1</a>	<a href="#">Plp1.aSep08</a>	<a href="#">24943</a>	8202	2929	7	286	proteolipid protein Dm-20 (Plp1) alternative variant aSep08, mRNA.
<a href="#">Plp1</a>	<a href="#">Plp1.cSep08</a>	<a href="#">24943</a>	12041	828	6	219	proteolipid protein Dm-20 (Plp1) alternative variant cSep08, mRNA.
<a href="#">Plp1</a>	<a href="#">Plp1.dSep08</a>	<a href="#">24943</a>	12121	719	6	131	proteolipid protein Dm-20 (Plp1) alternative variant dSep08, mRNA.
<a href="#">Plp1</a>	<a href="#">Plp1.fSep08</a>	<a href="#">24943</a>	3327	1083	4	72	proteolipid protein Dm-20 (7.8 kD) (Plp1) alternative variant fSep08, mRNA.
<a href="#">Plp1</a>	<a href="#">Plp1.gSep08</a>	<a href="#">24943</a>	8379	376	2	64	proteolipid protein Dm-20 (7.0 kD) (Plp1) alternative variant gSep08, mRNA.
<a href="#">Plp2</a>	<a href="#">Plp2.bSep08</a>	<a href="#">302562</a>	1957	821	4	95	proteolipid protein 2 (10.6 kD) (Plp2) alternative variant bSep08, mRNA.
<a href="#">Plp2</a>	<a href="#">Plp2.cSep08</a>	<a href="#">302562</a>	3423	1033	5	95	proteolipid protein 2 (10.6 kD) (Plp2) alternative variant cSep08, complete mRNA.
<a href="#">Plrg1</a>	<a href="#">Plrg1.bSep08</a>	<a href="#">60376</a>	4832	610	1	171	pleiotropic regulator 1, PRL1 homolog (Arabidopsis) (Plrg1) alternative variant bSep08, mRNA.
<a href="#">Pls1</a>	<a href="#">Pls1.bSep08</a>	<a href="#">315926</a>	70005	3711	2	630	plastin 1 (I isoform) (70.4 kD) (Pls1) alternative variant bSep08, mRNA.
<a href="#">Pls3</a>	<a href="#">Pls3.aSep08</a>	<a href="#">81748</a>	95350	2246	13	630	plastin 3 (T-isoform) (70.7 kD) (Pls3) alternative variant aSep08, mRNA.
<a href="#">Pls3</a>	<a href="#">Pls3.bSep08</a>	<a href="#">81748</a>	7562	881	8	293	plastin 3 (T-isoform) (Pls3) alternative variant bSep08, mRNA.
<a href="#">Pls3</a>	<a href="#">Pls3.cSep08</a>	<a href="#">81748</a>	42462	383	1	127	plastin 3 (T-isoform) (Pls3) alternative variant cSep08, mRNA.
<a href="#">Plscr1</a>	<a href="#">Plscr1.bSep08</a>	<a href="#">117540</a>	15901	1399	3	294	phospholipid scramblase 1 (31.9 kD) (Plscr1) alternative variant bSep08, complete mRNA.
<a href="#">Plscr1</a>	<a href="#">Plscr1.cSep08</a>	<a href="#">117540</a>	15254	801	1	207	phospholipid scramblase 1 (Plscr1) alternative variant cSep08, mRNA.
<a href="#">Plscr2</a>	<a href="#">Plscr2.bSep08</a>	<a href="#">315883</a>	11915	1292		265	phospholipid scramblase 2 (Plscr2) alternative variant bSep08, mRNA.
<a href="#">Plscr3</a>	<a href="#">Plscr3.bSep08</a>	<a href="#">360549</a>	601	416	1	138	phospholipid scramblase 3 (Plscr3) alternative variant bSep08, mRNA.

<a href="#">Pltp</a>	<a href="#">Pltp.aSep08</a>	<a href="#">296371</a>	18118	2025	16	580	phospholipid transfer protein (Pltp) alternative variant aSep08, mRNA.
<a href="#">Pltp</a>	<a href="#">Pltp.bSep08</a>	<a href="#">296371</a>	22131	1520	13	292	phospholipid transfer protein (Pltp) alternative variant bSep08, mRNA.
<a href="#">Pltp</a>	<a href="#">Pltp.cSep08</a>	<a href="#">296371</a>	12768	897	3	62	phospholipid transfer protein (Pltp) alternative variant cSep08, mRNA.
<a href="#">plubor</a>	<a href="#">plubor.aSep08</a>	<a href="#">309745</a>	276442	630		210	catenin Alpha-3 (plubor) mRNA.
<a href="#">pluchy</a>	<a href="#">pluchy.aSep08</a>		7846	240		33	putative protein (pluchy) mRNA.
<a href="#">pludoy</a>	<a href="#">pludoy.aSep08</a>		22603	278		34	putative protein (4.1 kD) (pludoy) mRNA.
<a href="#">plufee</a>	<a href="#">plufee.aSep08</a>		7661	376		89	putative protein (10.1 kD) (plufee) mRNA.
<a href="#">plufu</a>	<a href="#">plufu.aSep08</a>		961	639	3	148	protease serine 36 (plufu) alternative variant aSep08, mRNA.
<a href="#">plufly</a>	<a href="#">plufly.aSep08</a>		28066	783		86	putative protein of metazoan origin (9.2 kD) (plufly) mRNA.
<a href="#">plugar</a>	<a href="#">plugar.aSep08</a>		6497	244		81	F-box protein 25 (plugar) mRNA.
<a href="#">pluja</a>	<a href="#">pluja.aSep08</a>		592	318	2	13	putative protein (pluja) mRNA.
<a href="#">plujey</a>	<a href="#">plujey.aSep08</a>		38789	1820		33	putative protein (plujey) mRNA.
<a href="#">plulo</a>	<a href="#">plulo.aSep08</a>		662	435	2	30	putative protein (3.4 kD) (plulo) alternative variant aSep08, mRNA.
<a href="#">plumee</a>	<a href="#">plumee.aSep08</a>		4576	422		44	putative protein (4.7 kD) (plumee) mRNA.
<a href="#">plupey</a>	<a href="#">plupey.aSep08</a>		39640	604	1	129	putative protein (14.4 kD) (plupey) alternative variant aSep08, mRNA.
<a href="#">plupey</a>	<a href="#">plupey.cSep08</a>		40342	1327	2	86	putative protein (plupey) alternative variant cSep08, mRNA.
<a href="#">plupor</a>	<a href="#">plupor.aSep08</a>		28645	484		160	golgi autoantigen golgin subfamily a 4 CRA a like (plupor) mRNA.
<a href="#">pluroy</a>	<a href="#">pluroy.aSep08</a>		12453	647		192	CRA b (pluroy) mRNA.
<a href="#">plushaw</a>	<a href="#">plushaw.aSep08</a>		624	368		113	putative protein (plushaw) mRNA.
<a href="#">plushee</a>	<a href="#">plushee.aSep08</a>		3897	707		108	polyprotein (plushee) mRNA.
<a href="#">plutu</a>	<a href="#">plutu.aSep08</a>		2803	2463		534	putative protein of vertebrate origin (plutu) alternative variant aSep08, mRNA.
<a href="#">plutu</a>	<a href="#">plutu.bSep08</a>		5814	1784		399	putative protein of mammalian origin (plutu) alternative variant bSep08, mRNA.
<a href="#">pluvo</a>	<a href="#">pluvo.bSep08</a>		4304	574	2	18	putative protein (2.2 kD) (pluvo) alternative variant bSep08, mRNA.
<a href="#">pluwer</a>	<a href="#">pluwer.aSep08</a>		25680	341		113	activator for secretion protein 2 (pluwer) mRNA.
<a href="#">Plvap</a>	<a href="#">Plvap.bSep08</a>	<a href="#">56765</a>	1459	828	3	274	plasmalemma vesicle associated protein (Plvap) alternative variant bSep08, mRNA.
<a href="#">Plxna3</a>	<a href="#">Plxna3.bSep08</a>	<a href="#">309280</a>	992	733	4	148	plexin A3 (Plxna3) alternative variant bSep08, mRNA.
<a href="#">Plxna3</a>	<a href="#">Plxna3.cSep08</a>	<a href="#">309280</a>	1607	547	2	74	plexin A3 (8.6 kD) (Plxna3) alternative variant cSep08, mRNA.
<a href="#">Plxna4</a>	<a href="#">Plxna4.bSep08</a>	<a href="#">312213</a>	17275	734	1	112	plexin A4 CRA b (Plxna4) alternative variant bSep08, mRNA.
<a href="#">Plxnb1</a>	<a href="#">Plxnb1.bSep08</a>	<a href="#">316009</a>	9675	431	4	137	plexin B1 (Plxnb1) alternative variant bSep08, mRNA.
<a href="#">Plxnb1</a>	<a href="#">Plxnb1.cSep08</a>	<a href="#">316009</a>	972	401	3	133	plexin B1 (Plxnb1) alternative variant cSep08, mRNA.
<a href="#">Plxnb1</a>	<a href="#">Plxnb1.dSep08</a>	<a href="#">316009</a>	1253	414	5	107	plexin B1 (Plxnb1) alternative variant dSep08, mRNA.

<a href="#">Plxnb2</a>	<a href="#">Plxnb2.bSep08</a>	<a href="#">315217</a>	3072	1101	2	335	plexin B2 (Plxnb2) alternative variant bSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.cSep08</a>	<a href="#">315217</a>	2201	749	2	233	plexin B2 (Plxnb2) alternative variant cSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.dSep08</a>	<a href="#">315217</a>	2834	1308	8	217	plexin B2 (Plxnb2) alternative variant dSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.eSep08</a>	<a href="#">315217</a>	2689	1253	7	201	plexin B2 (Plxnb2) alternative variant eSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.fSep08</a>	<a href="#">315217</a>	1391	607	5	181	plexin B2 (Plxnb2) alternative variant fSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.gSep08</a>	<a href="#">315217</a>	756	435	2	123	plexin B2 (Plxnb2) alternative variant gSep08, mRNA.
<a href="#">Plxnb2</a>	<a href="#">Plxnb2.hSep08</a>	<a href="#">315217</a>	642	542	2	65	putative protein (Plxnb2) alternative variant hSep08, mRNA.
<a href="#">Plxnb3</a>	<a href="#">Plxnb3.aSep08</a>	<a href="#">363517</a>	1840	791	5	182	plexin B3 (Plxnb3) alternative variant aSep08, mRNA.
<a href="#">Plxnb3</a>	<a href="#">Plxnb3.bSep08</a>	<a href="#">363517</a>	385	318	1	58	plexin B3 (Plxnb3) alternative variant bSep08, mRNA.
<a href="#">Plxnc1</a>	<a href="#">Plxnc1.bSep08</a>	<a href="#">362873</a>	7145	2806	5	162	plexin C1 (Plxnc1) alternative variant bSep08, mRNA.
<a href="#">Plxnc1</a>	<a href="#">Plxnc1.cSep08</a>	<a href="#">362873</a>	19404	256	3	84	plexin C1 (Plxnc1) alternative variant cSep08, mRNA.
<a href="#">Plxnd1</a>	<a href="#">Plxnd1.bSep08</a>	<a href="#">312652</a>	9040	3060	14	274	plexin D1 (32.2 kD) (Plxnd1) alternative variant bSep08, mRNA.
<a href="#">Plxnd1</a>	<a href="#">Plxnd1.cSep08</a>	<a href="#">312652</a>	1525	881	3	81	plexin D1 (9.6 kD) (Plxnd1) alternative variant cSep08, mRNA.
<a href="#">plybor</a>	<a href="#">plybor.aSep08</a>		21054	613		76	putative protein (plybor) mRNA.
<a href="#">plychy</a>	<a href="#">plychy.aSep08</a>		11255	290		56	putative protein (plychy) mRNA.
<a href="#">plydoy</a>	<a href="#">plydoy.aSep08</a>		72826	661		78	putative protein (plydoy) mRNA.
<a href="#">plyfee</a>	<a href="#">plyfee.aSep08</a>		66682	516	1	96	CRA a like (11.2 kD) (plyfee) alternative variant aSep08, mRNA.
<a href="#">plyfee</a>	<a href="#">plyfee.bSep08</a>		84286	249	2	31	putative protein (plyfee) alternative variant bSep08, mRNA.
<a href="#">plyflu</a>	<a href="#">plyflu.aSep08</a>		626	271		90	putative protein (plyflu) mRNA.
<a href="#">plyfly</a>	<a href="#">plyfly.aSep08</a>		2895	754		95	putative protein (10.7 kD) (plyfly) mRNA.
<a href="#">plygar</a>	<a href="#">plygar.aSep08</a>		707	347		115	mcf.2 transforming sequence-like CRA a (plygar) mRNA.
<a href="#">plyja</a>	<a href="#">plyja.aSep08</a>		751	542		11	putative protein (1.2 kD) (plyja) mRNA.
<a href="#">plyjey</a>	<a href="#">plyjey.aSep08</a>		744	488		67	putative protein (plyjey) mRNA.
<a href="#">plylo</a>	<a href="#">plylo.aSep08</a>		16378	731		52	putative protein (plylo) mRNA.
<a href="#">plymee</a>	<a href="#">plymee.aSep08</a>		3068	449		54	putative protein (plymee) mRNA.
<a href="#">plypey</a>	<a href="#">plypey.aSep08</a>		5877	725		50	putative protein (5.8 kD) (plypey) mRNA.
<a href="#">plypor</a>	<a href="#">plypor.aSep08</a>		11870	350		116	putative protein of mammalian origin (plypor) mRNA.
<a href="#">plyroy</a>	<a href="#">plyroy.aSep08</a>		13432	1534		421	CRA a (plyroy) mRNA.
<a href="#">plyshaw</a>	<a href="#">plyshaw.aSep08</a>		4155	1732	2	139	catalytic beta polypeptide (plyshaw) alternative variant aSep08, mRNA.
<a href="#">plyshaw</a>	<a href="#">plyshaw.bSep08</a>		1788	738	1	106	catalytic beta polypeptide (plyshaw) alternative variant bSep08, mRNA.
<a href="#">plyshee</a>	<a href="#">plyshee.aSep08</a>		714	554		41	putative protein (4.8 kD) (plyshee) mRNA.
<a href="#">plytu</a>	<a href="#">plytu.aSep08</a>		1317	567		188	putative protein of eukaryotic origin (plytu) mRNA.
<a href="#">plyvo</a>	<a href="#">plyvo.aSep08</a>		945	411		104	microprotein (plyvo) mRNA.
<a href="#">plywer</a>	<a href="#">plywer.aSep08</a>		1176	314		58	putative protein (plywer) mRNA.
<a href="#">Pm20d2</a>	<a href="#">Pm20d2.aSep08</a>	<a href="#">313130</a>	14148	1331	6	342	aminoacylase 1-like 2 (Pm20d2) alternative variant aSep08, mRNA.

<a href="#">Pmepa1</a>	<a href="#">Pmepa1.aSep08</a>	<a href="#">311676</a>	7634	1498		230	prostate transmembrane protein, androgen induced 1 (Pmepa1) mRNA.
<a href="#">Pmf1</a>	<a href="#">Pmf1.aSep08</a>	<a href="#">681050</a>	19736	788		198	polyamine-modulated factor 1 (Pmf1) mRNA.
<a href="#">Pmfbp1</a>	<a href="#">Pmfbp1.bSep08</a>	<a href="#">171414</a>	28572	980	7	303	polyamine modulated factor 1 binding protein 1 (Pmfbp1) alternative variant bSep08, mRNA.
<a href="#">Pmfbp1</a>	<a href="#">Pmfbp1.cSep08</a>	<a href="#">171414</a>	5244	606	5	202	polyamine modulated factor 1 binding protein 1 (Pmfbp1) alternative variant cSep08, mRNA.
<a href="#">Pmfbp1</a>	<a href="#">Pmfbp1.dSep08</a>	<a href="#">171414</a>	2619	576	4	192	polyamine modulated factor 1 binding protein 1 (Pmfbp1) alternative variant dSep08, mRNA.
<a href="#">Pmfbp1</a>	<a href="#">Pmfbp1.eSep08</a>	<a href="#">171414</a>	1014	503	2	92	polyamine modulated factor 1 binding protein 1 (Pmfbp1) alternative variant eSep08, mRNA.
<a href="#">Pml</a>	<a href="#">Pml.aSep08</a>	<a href="#">315713</a>	11719	604		201	promyelocytic leukemia (Pml) mRNA.
<a href="#">Pmm1</a>	<a href="#">Pmm1.bSep08</a>	<a href="#">300089</a>	9756	756	8	170	phosphomannomutase 1 CRA b (Pmm1) alternative variant bSep08, mRNA.
<a href="#">Pmm1</a>	<a href="#">Pmm1.cSep08</a>	<a href="#">300089</a>	5742	763	4	125	phosphomannomutase 1 CRA b (14.1 kD) (Pmm1) alternative variant cSep08, mRNA.
<a href="#">Pmm1</a>	<a href="#">Pmm1.dSep08</a>	<a href="#">300089</a>	3584	374	3	65	phosphomannomutase (Pmm1) alternative variant dSep08, mRNA.
<a href="#">Pmm2</a>	<a href="#">Pmm2.bSep08</a>	<a href="#">302915</a>	11087	693	5	126	phosphomannomutase 2 CRA b (14.3 kD) (Pmm2) alternative variant bSep08, complete mRNA.
<a href="#">Pmm2</a>	<a href="#">Pmm2.cSep08</a>	<a href="#">302915</a>	2027	712	2	101	putative protein (10.9 kD) (Pmm2) alternative variant cSep08, mRNA.
<a href="#">Pmm2</a>	<a href="#">Pmm2.dSep08</a>	<a href="#">302915</a>	9365	747	5	97	phosphomannomutase 2 CRA b (Pmm2) alternative variant dSep08, mRNA.
<a href="#">Pmm2</a>	<a href="#">Pmm2.eSep08</a>	<a href="#">302915</a>	15420	849	3	63	phosphomannomutase 2 CRA b (Pmm2) alternative variant eSep08, mRNA.
<a href="#">Pmm2</a>	<a href="#">Pmm2.fSep08</a>	<a href="#">302915</a>	6649	1156	3	35	phosphomannomutase 2 (4.0 kD) (Pmm2) alternative variant fSep08, mRNA.
<a href="#">Pmp2</a>	<a href="#">Pmp2.bSep08</a>	<a href="#">688790</a>	3257	682	4	110	peripheral myelin protein 2 (12.4 kD) (Pmp2) alternative variant bSep08, complete mRNA.
<a href="#">Pmp22</a>	<a href="#">Pmp22.bSep08</a>	<a href="#">24660</a>	19202	354	3	117	peripheral myelin protein 22 (Pmp22) alternative variant bSep08, mRNA.
<a href="#">Pmp22</a>	<a href="#">Pmp22.cSep08</a>	<a href="#">24660</a>	19622	617	4	97	peripheral myelin protein 22 (Pmp22) alternative variant cSep08, mRNA.
<a href="#">Pmp22</a>	<a href="#">Pmp22.dSep08</a>	<a href="#">24660</a>	20825	492	4	97	peripheral myelin protein 22 (Pmp22) alternative variant dSep08, mRNA.
<a href="#">Pmpca</a>	<a href="#">Pmpca.bSep08</a>	<a href="#">296588</a>	866	774	2	247	peptidase (mitochondrial processing) alpha (Pmpca) alternative variant bSep08, mRNA.
<a href="#">Pmpca</a>	<a href="#">Pmpca.cSep08</a>	<a href="#">296588</a>	4804	761	7	219	peptidase (mitochondrial processing) alpha (Pmpca) alternative variant cSep08, mRNA.
<a href="#">Pmpca</a>	<a href="#">Pmpca.dSep08</a>	<a href="#">296588</a>	1826	531	5	177	peptidase (mitochondrial processing) alpha (Pmpca) alternative variant dSep08, mRNA.
<a href="#">Pmpcb</a>	<a href="#">Pmpcb.bSep08</a>	<a href="#">64198</a>	6106	827	6	259	peptidase (mitochondrial processing) beta (Pmpcb) alternative variant bSep08, mRNA.
<a href="#">Pmpcb</a>	<a href="#">Pmpcb.dSep08</a>	<a href="#">64198</a>	561	341	3	84	peptidase (mitochondrial processing) beta (Pmpcb) alternative variant dSep08, mRNA.

<a href="#">Pmpcb</a>	<a href="#">Pmpcb.eSep08</a>	<a href="#">64198</a>	1721	780	3	84	peptidase (mitochondrial processing) beta (Pmpcb) alternative variant eSep08, mRNA.
<a href="#">Pms2</a>	<a href="#">Pms2.aSep08</a>	<a href="#">288479</a>	24197	2875	15	853	postmeiotic segregation increased 2 (S. cerevisiae) (94.5 kD) (Pms2) alternative variant aSep08, mRNA.
<a href="#">Pms2</a>	<a href="#">Pms2.cSep08</a>	<a href="#">288479</a>	4913	762	3	165	postmeiotic segregation increased 2 (S. cerevisiae) (Pms2) alternative variant cSep08, mRNA.
<a href="#">Pms2</a>	<a href="#">Pms2.dSep08</a>	<a href="#">288479</a>	4804	687	3	74	postmeiotic segregation increased 2 (S. cerevisiae) (8.3 kD) (Pms2) alternative variant dSep08, mRNA.
<a href="#">Pmvk</a>	<a href="#">Pmvk.bSep08</a>	<a href="#">310645</a>	9550	751	3	83	phosphomevalonate kinase (9.6 kD) (Pmvk) alternative variant bSep08, mRNA.
<a href="#">Pnkd</a>	<a href="#">Pnkd.bSep08</a>	<a href="#">100188944</a>	86084	3065	9	369	paroxysmal nonkinesinogenic dyskinesia (41.0 kD) (Pnkd) alternative variant bSep08, complete mRNA.
<a href="#">Pnkd</a>	<a href="#">Pnkd.fSep08</a>	<a href="#">100188944</a>	20834	1073	3	79	paroxysmal nonkinesinogenic dyskinesia (8.4 kD) (Pnkd) alternative variant fSep08, complete mRNA.
<a href="#">Pnkp</a>	<a href="#">Pnkp.bSep08</a>	<a href="#">308576</a>	3676	1105	12	255	polynucleotide kinase 3'-phosphatase (Pnkp) alternative variant bSep08, mRNA.
<a href="#">Pnkp</a>	<a href="#">Pnkp.cSep08</a>	<a href="#">308576</a>	3347	987	6	203	polynucleotide kinase 3'-phosphatase (Pnkp) alternative variant cSep08, mRNA.
<a href="#">Pnkp</a>	<a href="#">Pnkp.dSep08</a>	<a href="#">308576</a>	792	716	2	80	polynucleotide kinase 3'-phosphatase (Pnkp) alternative variant dSep08, mRNA.
<a href="#">Pnlhc1</a>	<a href="#">Pnlhc1.bSep08</a>	<a href="#">361478</a>	9714	877	6	292	ribonuclease CAF1 (Pnlhc1) alternative variant bSep08, mRNA.
<a href="#">Pnlhc1</a>	<a href="#">Pnlhc1.cSep08</a>	<a href="#">361478</a>	7573	759	3	252	putative protein of eukaryotic origin (Pnlhc1) alternative variant cSep08, mRNA.
<a href="#">Pnlhc1</a>	<a href="#">Pnlhc1.dSep08</a>	<a href="#">361478</a>	11091	761	7	158	putative protein of eukaryotic origin (Pnlhc1) alternative variant dSep08, mRNA.
<a href="#">Pnlipr2</a>	<a href="#">Pnlipr2.bSep08</a>	<a href="#">117554</a>	6312	444	4	129	pancreatic lipase-related protein 2 (Pnlipr2) alternative variant bSep08, mRNA.
<a href="#">Pnmal1</a>	<a href="#">Pnmal1.bSep08</a>	<a href="#">361515</a>	743	582	2	90	PNMA-like 1 (Pnmal1) alternative variant bSep08, mRNA.
<a href="#">Pnmal2</a>	<a href="#">Pnmal2.bSep08</a>	<a href="#">308393</a>	2705	1835	2	375	PNMA-like 2 (Pnmal2) alternative variant bSep08, mRNA.
<a href="#">Pno1</a>	<a href="#">Pno1.bSep08</a>	<a href="#">289809</a>	7551	614	2	202	partner of NOB1 homolog (S. cerevisiae) (Pno1) alternative variant bSep08, mRNA.
<a href="#">Pnpla1</a>	<a href="#">Pnpla1.aSep08</a>	<a href="#">361812</a>	8469	1781		465	putative protein (Pnpla1) alternative variant aSep08, mRNA.
<a href="#">Pnpla3</a>	<a href="#">Pnpla3.aSep08</a>	<a href="#">362972</a>	21685	3431	9	383	adiponutrin (42.4 kD) (Pnpla3) alternative variant aSep08, complete mRNA.
<a href="#">Pnpla3</a>	<a href="#">Pnpla3.bSep08</a>	<a href="#">362972</a>	10745	879	6	280	patatin (Pnpla3) alternative variant bSep08, mRNA.
<a href="#">Pnpla3</a>	<a href="#">Pnpla3.cSep08</a>	<a href="#">362972</a>	3430	638	2	132	putative protein of eukaryotic origin (Pnpla3) alternative variant cSep08, mRNA.
<a href="#">Pnpla6</a>	<a href="#">Pnpla6.aSep08</a>	<a href="#">360753</a>	1698	632	4	128	putative protein of vertebrate origin (Pnpla6) alternative variant aSep08, mRNA.
<a href="#">Pnpla6</a>	<a href="#">Pnpla6.bSep08</a>	<a href="#">360753</a>	29337	873	5	39	putative protein (Pnpla6) alternative variant bSep08, mRNA.
<a href="#">Pnpla6</a>	<a href="#">Pnpla6.cSep08</a>	<a href="#">360753</a>	880	750	2	59	putative protein (Pnpla6) alternative variant cSep08, mRNA.
<a href="#">Pnpla7</a>	<a href="#">Pnpla7.bSep08</a>	<a href="#">246246</a>	70095	4446	26	957	cyclic nucleotide-binding (106.3 kD) (Pnpla7) alternative variant bSep08, mRNA.

<a href="#">Pnpla7</a>	<a href="#">Pnpla7.cSep08</a>	<a href="#">246246</a>	37352	2373	18	682	patatin (75.8 kD) (Pnpla7) alternative variant cSep08, mRNA.
<a href="#">Pnpla7</a>	<a href="#">Pnpla7.dSep08</a>	<a href="#">246246</a>	4382	1262	4	225	putative protein of ancient origin (Pnpla7) alternative variant dSep08, mRNA.
<a href="#">Pnpla7</a>	<a href="#">Pnpla7.fSep08</a>	<a href="#">246246</a>	1425	388	2	70	neuropathy target esterase-related esterase splice like (7.7 kD) (Pnpla7) alternative variant fSep08, mRNA.
<a href="#">Pnpla8</a>	<a href="#">Pnpla8.aSep08</a>	<a href="#">314075</a>	34741	4414	8	776	membrane-associated calcium-independent phospholipase A2 gamma (88.0 kD) (Pnpla8) alternative variant aSep08, mRNA.
<a href="#">Pnpla8</a>	<a href="#">Pnpla8.bSep08</a>	<a href="#">314075</a>	59690	1786	5	595	putative protein of eukaryotic origin (Pnpla8) alternative variant bSep08, mRNA.
<a href="#">Pnpla8</a>	<a href="#">Pnpla8.dSep08</a>	<a href="#">314075</a>	3284	409	2	19	putative protein (2.3 kD) (Pnpla8) alternative variant dSep08, mRNA.
<a href="#">Pnpt1</a>	<a href="#">Pnpt1.aSep08</a>	<a href="#">360992</a>	23454	1801	20	532	polyribonucleotide nucleotidyltransferase 1 (Pnpt1) alternative variant aSep08, mRNA.
<a href="#">Pnpt1</a>	<a href="#">Pnpt1.bSep08</a>	<a href="#">360992</a>	1372	330	3	63	polyribonucleotide nucleotidyltransferase 1 (Pnpt1) alternative variant bSep08, mRNA.
<a href="#">Pnpt1</a>	<a href="#">Pnpt1.cSep08</a>	<a href="#">360992</a>	629	543	2	34	polyribonucleotide nucleotidyltransferase 1 (4.0 kD) (Pnpt1) alternative variant cSep08, mRNA.
<a href="#">Pnpt1</a>	<a href="#">Pnpt1.dSep08</a>	<a href="#">360992</a>	3898	917	2	40	polyribonucleotide nucleotidyltransferase 1 (4.3 kD) (Pnpt1) alternative variant dSep08, mRNA.
<a href="#">Pnrc1</a>	<a href="#">Pnrc1.bSep08</a>	<a href="#">286988</a>	45654	829	2	142	proline-rich nuclear receptor coactivator 1 (16.0 kD) (Pnrc1) alternative variant bSep08, mRNA.
<a href="#">poby</a>	<a href="#">poby.aSep08</a>		4048	384		23	putative protein (2.4 kD) (poby) mRNA.
<a href="#">pochy</a>	<a href="#">pochy.aSep08</a>		13305	926		138	putative protein (pochy) alternative variant aSep08, mRNA.
<a href="#">podar</a>	<a href="#">podar.aSep08</a>		9351	393		61	putative protein (podar) mRNA.
<a href="#">Podnl1</a>	<a href="#">Podnl1.bSep08</a>	<a href="#">288907</a>	2170	1396	3	218	podocan-like 1 (Podnl1) alternative variant bSep08, mRNA.
<a href="#">Podnl1</a>	<a href="#">Podnl1.cSep08</a>	<a href="#">288907</a>	730	658	2	132	podocan-like 1 (Podnl1) alternative variant cSep08, mRNA.
<a href="#">Podxl2</a>	<a href="#">Podxl2.cSep08</a>	<a href="#">297433</a>	2930	401	2	69	podocalyxin-like 2 (Podxl2) alternative variant cSep08, mRNA.
<a href="#">Pof1b</a>	<a href="#">Pof1b.bSep08</a>	<a href="#">302328</a>	34368	605	1	201	premature ovarian failure 1B (Pof1b) alternative variant bSep08, mRNA.
<a href="#">pofer</a>	<a href="#">pofer.aSep08</a>		7994	1732		78	putative cytoplasmic protein of eukaryotic origin (9.2 kD) (pofer) mRNA.
<a href="#">poflo</a>	<a href="#">poflo.aSep08</a>		88052	418	1	88	putative protein (poflo) alternative variant aSep08, mRNA.
<a href="#">poflo</a>	<a href="#">poflo.bSep08</a>		62176	537	1	71	golgi-specific brefeldin A-resistance guanine nucleotide exchange factor 1 (7.9 kD) (poflo) alternative variant bSep08, mRNA.
<a href="#">poflu</a>	<a href="#">poflu.aSep08</a>		3099	400		133	leucine-rich repeat kinase 1 CRA b (poflu) mRNA.
<a href="#">Pofut2</a>	<a href="#">Pofut2.bSep08</a>	<a href="#">309686</a>	1033	901	2	94	protein O-fucosyltransferase 2 (Pofut2) alternative variant bSep08, mRNA.
<a href="#">pogar</a>	<a href="#">pogar.aSep08</a>		2891	744		100	putative protein (pogar) mRNA.
<a href="#">poja</a>	<a href="#">poja.aSep08</a>		1766	971	2	82	putative protein (9.8 kD) (poja) alternative variant aSep08, mRNA.
<a href="#">poja</a>	<a href="#">poja.bSep08</a>		1784	802	3	45	putative protein (5.3 kD) (poja) alternative variant bSep08, mRNA.

<a href="#">pojey</a>	<a href="#">pojey.aSep08</a>		2646	497		60	putative protein (pojey) mRNA.
<a href="#">pokee</a>	<a href="#">pokee.bSep08</a>		15951	434		94	putative nuclear protein (9.7 kD) (pokee) alternative variant bSep08, mRNA.
<a href="#">pokler</a>	<a href="#">pokler.aSep08</a>		2318	356		26	putative protein (pokler) mRNA.
<a href="#">Pola1</a>	<a href="#">Pola1.aSep08</a>	<a href="#">85241</a>	189946	2297		468	polymerase (DNA directed), alpha 1 (Pola1) mRNA.
<a href="#">Pola2</a>	<a href="#">Pola2.bSep08</a>	<a href="#">85242</a>	10608	714	5	219	polymerase (DNA directed), alpha 2 (Pola2) alternative variant bSep08, mRNA.
<a href="#">Pola2</a>	<a href="#">Pola2.cSep08</a>	<a href="#">85242</a>	9029	760	7	148	polymerase (DNA directed), alpha 2 (Pola2) alternative variant cSep08, mRNA.
<a href="#">Pola2</a>	<a href="#">Pola2.dSep08</a>	<a href="#">85242</a>	1432	719	2	139	polymerase (DNA directed), alpha 2 (Pola2) alternative variant dSep08, mRNA.
<a href="#">Pola2</a>	<a href="#">Pola2.fSep08</a>	<a href="#">85242</a>	2174	1004	3	35	polymerase (DNA directed), alpha 2 (3.9 kD) (Pola2) alternative variant fSep08, mRNA.
<a href="#">Polb</a>	<a href="#">Polb.bSep08</a>	<a href="#">29240</a>	5634	671	5	92	polymerase (DNA directed), beta (Polb) alternative variant bSep08, mRNA.
<a href="#">Polb</a>	<a href="#">Polb.cSep08</a>	<a href="#">29240</a>	3024	300	3	91	polymerase (DNA directed), beta (10.1 kD) (Polb) alternative variant cSep08, mRNA.
<a href="#">Pold2</a>	<a href="#">Pold2.bSep08</a>	<a href="#">289758</a>	4792	799	5	265	polymerase (DNA directed), delta 2, regulatory subunit (Pold2) alternative variant bSep08, mRNA.
<a href="#">Pold2</a>	<a href="#">Pold2.dSep08</a>	<a href="#">289758</a>	560	316	2	94	polymerase (DNA directed), delta 2, regulatory subunit (Pold2) alternative variant dSep08, mRNA.
<a href="#">Pold2</a>	<a href="#">Pold2.eSep08</a>	<a href="#">289758</a>	823	494	2	68	polymerase (DNA directed), delta 2, regulatory subunit (Pold2) alternative variant eSep08, mRNA.
<a href="#">Pold4</a>	<a href="#">Pold4.bSep08</a>	<a href="#">361698</a>	1555	543	4	110	polymerase (DNA-directed), delta 4 (Pold4) alternative variant bSep08, mRNA.
<a href="#">Pold4</a>	<a href="#">Pold4.dSep08</a>	<a href="#">361698</a>	1320	767	3	32	polymerase (DNA-directed), delta 4 (3.8 kD) (Pold4) alternative variant dSep08, mRNA.
<a href="#">Pold4</a>	<a href="#">Pold4.eSep08</a>	<a href="#">361698</a>	1331	594	3	32	polymerase (DNA-directed), delta 4 (3.7 kD) (Pold4) alternative variant eSep08, mRNA.
<a href="#">Poldip2</a>	<a href="#">Poldip2.bSep08</a>	<a href="#">287544</a>	6505	858	8	286	polymerase (DNA-directed), delta interacting protein 2 (Poldip2) alternative variant bSep08, mRNA.
<a href="#">Poldip2</a>	<a href="#">Poldip2.cSep08</a>	<a href="#">287544</a>	3353	897	3	229	polymerase (DNA-directed), delta interacting protein 2 (26.2 kD) (Poldip2) alternative variant cSep08, mRNA.
<a href="#">Poldip2</a>	<a href="#">Poldip2.dSep08</a>	<a href="#">287544</a>	1518	922	2	109	polymerase (DNA-directed), delta interacting protein 2 (Poldip2) alternative variant dSep08, mRNA.
<a href="#">Pole</a>	<a href="#">Pole.bSep08</a>	<a href="#">304573</a>	7713	1010	4	336	polymerase (DNA directed), epsilon (Pole) alternative variant bSep08, mRNA.
<a href="#">Pole</a>	<a href="#">Pole.cSep08</a>	<a href="#">304573</a>	5952	849	2	278	polymerase (DNA directed), epsilon (Pole) alternative variant cSep08, mRNA.
<a href="#">Pole2</a>	<a href="#">Pole2.aSep08</a>	<a href="#">299112</a>	17587	1011	6	336	polymerase (DNA directed), epsilon 2 (p59 subunit) (Pole2) alternative variant aSep08, mRNA.
<a href="#">Pole2</a>	<a href="#">Pole2.bSep08</a>	<a href="#">299112</a>	12768	677	2	180	polymerase (DNA directed), epsilon 2 (p59 subunit) (Pole2) alternative variant bSep08, mRNA.
<a href="#">Pole4</a>	<a href="#">Pole4.bSep08</a>	<a href="#">362385</a>	5189	1980	2	107	polymerase (DNA-directed), epsilon 4 (p12 subunit) (10.9 kD) (Pole4) alternative variant bSep08, complete mRNA.
<a href="#">Pole4</a>	<a href="#">Pole4.dSep08</a>	<a href="#">362385</a>	4210	559	3	88	polymerase (DNA-directed), epsilon 4 (p12 subunit) (9.8 kD) (Pole4) alternative variant dSep08, mRNA.

<a href="#">Polg</a>	<a href="#">Polg.bSep08</a>	<a href="#">85472</a>	9090	2176	1	522	polymerase (DNA directed), gamma (Polg) alternative variant bSep08, mRNA.
<a href="#">Polg2</a>	<a href="#">Polg2.bSep08</a>	<a href="#">303612</a>	6088	849	5	174	polymerase (DNA directed), gamma 2, accessory subunit (Polg2) alternative variant bSep08, mRNA.
<a href="#">Poli</a>	<a href="#">Poli.bSep08</a>	<a href="#">291526</a>	8556	767	2	255	polymerase (DNA directed), iota (Poli) alternative variant bSep08, mRNA.
<a href="#">Poll</a>	<a href="#">Poll.bSep08</a>	<a href="#">361767</a>	3995	1321	4	261	polymerase (DNA directed), lambda (Poll) alternative variant bSep08, mRNA.
<a href="#">Poll</a>	<a href="#">Poll.cSep08</a>	<a href="#">361767</a>	920	632	2	172	polymerase (DNA directed), lambda (Poll) alternative variant cSep08, mRNA.
<a href="#">Poll</a>	<a href="#">Poll.dSep08</a>	<a href="#">361767</a>	2946	625	3	164	polymerase (DNA directed), lambda (Poll) alternative variant dSep08, mRNA.
<a href="#">Poll</a>	<a href="#">Poll.fSep08</a>	<a href="#">361767</a>	2050	549	2	113	polymerase (DNA directed), lambda (Poll) alternative variant fSep08, mRNA.
<a href="#">Poll</a>	<a href="#">Poll.gSep08</a>	<a href="#">361767</a>	1959	298	2	83	polymerase (DNA directed), lambda (Poll) alternative variant gSep08, mRNA.
<a href="#">Poln</a>	<a href="#">Poln.aSep08</a>	<a href="#">498396</a>	2431	441		62	polymerase (DNA directed) nu (Poln) mRNA.
<a href="#">poloy</a>	<a href="#">poloy.aSep08</a>		9611	2987		282	DiGeorge syndrome critical region gene 2 like (poloy) mRNA.
<a href="#">Polr1e</a>	<a href="#">Polr1e.bSep08</a>	<a href="#">313245</a>	8919	711	3	204	polymerase (RNA) I polypeptide E (22.6 kD) (Polr1e) alternative variant bSep08, mRNA.
<a href="#">Polr1e</a>	<a href="#">Polr1e.cSep08</a>	<a href="#">313245</a>	10241	1569	4	171	polymerase (RNA) I polypeptide E (Polr1e) alternative variant cSep08, mRNA.
<a href="#">Polr1e</a>	<a href="#">Polr1e.dSep08</a>	<a href="#">313245</a>	9326	769	5	162	polymerase (RNA) I polypeptide E (18.2 kD) (Polr1e) alternative variant dSep08, mRNA.
<a href="#">Polr2a</a>	<a href="#">Polr2a.aSep08</a>	<a href="#">363633</a>	2842	2040		542	polymerase (RNA) II (DNA directed) polypeptide A (Polr2a) alternative variant aSep08, mRNA.
<a href="#">Polr2a</a>	<a href="#">Polr2a.bSep08</a>	<a href="#">363633</a>	6692	1670		429	polymerase (RNA) II (DNA directed) polypeptide A (Polr2a) alternative variant bSep08, mRNA.
<a href="#">Polr2a</a>	<a href="#">Polr2a.cSep08</a>	<a href="#">363633</a>	3046	501		166	polymerase (RNA) II (DNA directed) polypeptide A (Polr2a) alternative variant cSep08, mRNA.
<a href="#">Polr2c</a>	<a href="#">Polr2c.bSep08</a>	<a href="#">361365</a>	1572	1003	2	87	polymerase (RNA) II (DNA directed) polypeptide C (Polr2c) alternative variant bSep08, mRNA.
<a href="#">Polr2d</a>	<a href="#">Polr2d.bSep08</a>	<a href="#">364834</a>	2277	749	1	29	polymerase (RNA) II (DNA directed) polypeptide D (3.3 kD) (Polr2d) alternative variant bSep08, mRNA.
<a href="#">Polr2e</a>	<a href="#">Polr2e.bSep08</a>	<a href="#">690966</a>	3331	828	6	202	polymerase (RNA) II (DNA directed) polypeptide E (23.1 kD) (Polr2e) alternative variant bSep08, mRNA.
<a href="#">Polr2e</a>	<a href="#">Polr2e.cSep08</a>	<a href="#">690966</a>	2785	700	6	166	polymerase (RNA) II (DNA directed) polypeptide E (Polr2e) alternative variant cSep08, mRNA.
<a href="#">Polr2e</a>	<a href="#">Polr2e.dSep08</a>	<a href="#">690966</a>	1736	655	6	132	polymerase (RNA) II (DNA directed) polypeptide E (Polr2e) alternative variant dSep08, mRNA.
<a href="#">Polr2e</a>	<a href="#">Polr2e.eSep08</a>	<a href="#">690966</a>	879	538	2	96	polymerase (RNA) II (DNA directed) polypeptide E (Polr2e) alternative variant eSep08, mRNA.
<a href="#">Polr2e</a>	<a href="#">Polr2e.fSep08</a>	<a href="#">690966</a>	1554	550	4	60	polymerase (RNA) II (DNA directed) polypeptide E (Polr2e) alternative variant fSep08, mRNA.
<a href="#">Polr2g</a>	<a href="#">Polr2g.aSep08</a>	<a href="#">117017</a>	6546	1187	8	174	polymerase (RNA) II (DNA directed) polypeptide G (19.5 kD) (Polr2g) alternative variant aSep08, mRNA.



<a href="#">Polr2g</a>	<a href="#">Polr2g.cSep08</a>	<a href="#">117017</a>	2622	616	7	153	polymerase (RNA) II (DNA directed) polypeptide G (17.1 kD) (Polr2g) alternative variant cSep08, mRNA.
<a href="#">Polr2g</a>	<a href="#">Polr2g.dSep08</a>	<a href="#">117017</a>	2135	416	4	138	polymerase (RNA) II (DNA directed) polypeptide G (Polr2g) alternative variant dSep08, mRNA.
<a href="#">Polr2h</a>	<a href="#">Polr2h.bSep08</a>	<a href="#">498109</a>	2702	451	4	112	polymerase (RNA) II (DNA directed) polypeptide H (12.9 kD) (Polr2h) alternative variant bSep08, mRNA.
<a href="#">Polr2i</a>	<a href="#">Polr2i.bSep08</a>	<a href="#">292778</a>	1055	493	5	103	polymerase (RNA) II (DNA directed) polypeptide I (12.0 kD) (Polr2i) alternative variant bSep08, mRNA.
<a href="#">Polr2i</a>	<a href="#">Polr2i.dSep08</a>	<a href="#">292778</a>	920	444	4	67	polymerase (RNA) II (DNA directed) polypeptide I (Polr2i) alternative variant dSep08, mRNA.
<a href="#">Polr3a</a>	<a href="#">Polr3a.aSep08</a>	<a href="#">361102</a>	14897	1498		499	polymerase (RNA) III (DNA directed) polypeptide A (Polr3a) mRNA.
<a href="#">Polr3b</a>	<a href="#">Polr3b.aSep08</a>	<a href="#">362858</a>	54465	2412	12	560	polymerase (RNA) III (DNA directed) polypeptide B (Polr3b) mRNA.
<a href="#">Polr3c</a>	<a href="#">Polr3c.bSep08</a>	<a href="#">310685</a>	3925	531	3	103	polymerase (RNA) III (DNA directed) polypeptide C (Polr3c) alternative variant bSep08, mRNA.
<a href="#">Polr3c</a>	<a href="#">Polr3c.cSep08</a>	<a href="#">310685</a>	1160	343	2	84	polymerase (RNA) III (DNA directed) polypeptide C (Polr3c) alternative variant cSep08, mRNA.
<a href="#">Polr3d</a>	<a href="#">Polr3d.bSep08</a>	<a href="#">306012</a>	2535	631	1	209	polymerase (RNA) III (DNA directed) polypeptide D (Polr3d) alternative variant bSep08, mRNA.
<a href="#">Polr3e</a>	<a href="#">Polr3e.bSep08</a>	<a href="#">361640</a>	760	258	3	85	polymerase (RNA) III (DNA directed) polypeptide E (Polr3e) alternative variant bSep08, mRNA.
<a href="#">Polr3gl</a>	<a href="#">Polr3gl.bSep08</a>	<a href="#">690254</a>	1076	326	2	100	putative protein (Polr3gl) alternative variant bSep08, mRNA.
<a href="#">Polr3gl</a>	<a href="#">Polr3gl.cSep08</a>	<a href="#">690254</a>	1370	733	3	71	putative protein (8.0 kD) (Polr3gl) alternative variant cSep08, mRNA.
<a href="#">Polr3gl</a>	<a href="#">Polr3gl.dSep08</a>	<a href="#">690254</a>	649	543	2	58	putative protein, with a coiled coil domain (Polr3gl) alternative variant dSep08, mRNA.
<a href="#">Polr3gl</a>	<a href="#">Polr3gl.eSep08</a>	<a href="#">690254</a>	1230	550	2	25	putative protein (Polr3gl) alternative variant eSep08, mRNA.
<a href="#">Polr3h</a>	<a href="#">Polr3h.aSep08</a>	<a href="#">300088</a>	10478	1797	7	476	polymerase (RNA) III (DNA directed) polypeptide H (Polr3h) alternative variant aSep08, complete mRNA.
<a href="#">Polr3h</a>	<a href="#">Polr3h.cSep08</a>	<a href="#">300088</a>	21236	632	7	128	polymerase (RNA) III (DNA directed) polypeptide H (Polr3h) alternative variant cSep08, mRNA.
<a href="#">Polrmt</a>	<a href="#">Polrmt.bSep08</a>	<a href="#">299604</a>	2623	1706	9	538	polymerase mitochondrial CRA a (Polrmt) alternative variant bSep08, mRNA.
<a href="#">Polrmt</a>	<a href="#">Polrmt.cSep08</a>	<a href="#">299604</a>	3442	2523	7	262	polymerase mitochondrial CRA b (29.3 kD) (Polrmt) alternative variant cSep08, mRNA.
<a href="#">Polrmt</a>	<a href="#">Polrmt.dSep08</a>	<a href="#">299604</a>	530	431	2	84	polymerase mitochondrial CRA b (Polrmt) alternative variant dSep08, mRNA.
<a href="#">Pomc</a>	<a href="#">Pomc.bSep08</a>	<a href="#">24664</a>	5529	702	2	173	proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-endorphin) (Pomc) alternative variant bSep08, mRNA.

<a href="#">Pomc</a>	<a href="#">Pomc.cSep08</a>	<a href="#">24664</a>	5246	387	1	84	proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-endorphin) (Pomc) alternative variant cSep08, mRNA.
<a href="#">pomee</a>	<a href="#">pomee.aSep08</a>		9213	745		52	putative protein (6.1 kD) (pomee) mRNA.
<a href="#">pomer</a>	<a href="#">pomer.aSep08</a>		11456	647		175	CRA b like (pomer) alternative variant aSep08, mRNA.
<a href="#">pomer</a>	<a href="#">pomer.bSep08</a>		9030	456		132	CRA b like (pomer) alternative variant bSep08, mRNA.
<a href="#">Pomgnt1</a>	<a href="#">Pomgnt1.bSep08</a>	<a href="#">362567</a>	3342	618	4	189	protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase (Pomgnt1) alternative variant bSep08, mRNA.
<a href="#">Pomgnt1</a>	<a href="#">Pomgnt1.cSep08</a>	<a href="#">362567</a>	1401	773	2	120	protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase (Pomgnt1) alternative variant cSep08, mRNA.
<a href="#">Pomgnt1</a>	<a href="#">Pomgnt1.dSep08</a>	<a href="#">362567</a>	2066	1351	3	92	protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase (10.5 kD) (Pomgnt1) alternative variant dSep08, mRNA.
<a href="#">Pomp</a>	<a href="#">Pomp.bSep08</a>	<a href="#">288455</a>	11041	429	1	120	proteasome maturation protein (Pomp) alternative variant bSep08, mRNA.
<a href="#">Pomt1</a>	<a href="#">Pomt1.aSep08</a>	<a href="#">84430</a>	15667	1956	18	651	protein-O-mannosyltransferase 1 (Pomt1) alternative variant aSep08, mRNA.
<a href="#">Pomt1</a>	<a href="#">Pomt1.bSep08</a>	<a href="#">84430</a>	6520	604	7	200	protein-O-mannosyltransferase 1 (Pomt1) alternative variant bSep08, mRNA.
<a href="#">Pomt1</a>	<a href="#">Pomt1.cSep08</a>	<a href="#">84430</a>	2676	303	4	100	protein-O-mannosyltransferase 1 (Pomt1) alternative variant cSep08, mRNA.
<a href="#">Pomt2</a>	<a href="#">Pomt2.aSep08</a>	<a href="#">366697</a>	76922	1663	11	554	protein-O-mannosyltransferase 2 (Pomt2) alternative variant aSep08, mRNA.
<a href="#">Pomt2</a>	<a href="#">Pomt2.bSep08</a>	<a href="#">366697</a>	22957	1518	3	142	protein-O-mannosyltransferase 2 (Pomt2) alternative variant bSep08, mRNA.
<a href="#">Pomt2</a>	<a href="#">Pomt2.cSep08</a>	<a href="#">366697</a>	40516	362	3	120	protein-O-mannosyltransferase 2 (Pomt2) alternative variant cSep08, mRNA.
<a href="#">Pon2</a>	<a href="#">Pon2.bSep08</a>	<a href="#">296851</a>	4269	771	4	196	paraoxonase 2 (22.0 kD) (Pon2) alternative variant bSep08, mRNA.
<a href="#">Pon2</a>	<a href="#">Pon2.cSep08</a>	<a href="#">296851</a>	30920	939	7	158	paraoxonase 2 (18.3 kD) (Pon2) alternative variant cSep08, complete mRNA.
<a href="#">Pon3</a>	<a href="#">Pon3.bSep08</a>	<a href="#">312086</a>	4749	989	2	67	paraoxonase 3 (7.5 kD) (Pon3) alternative variant bSep08, mRNA.
<a href="#">ponoy</a>	<a href="#">ponoy.aSep08</a>		1713	378		126	putative protein (ponoy) mRNA.
<a href="#">Pop4</a>	<a href="#">Pop4.bSep08</a>	<a href="#">292831</a>	4867	1076	5	219	processing of precursor 4, ribonuclease P/MRP family, (S. cerevisiae) (25.2 kD) (Pop4) alternative variant bSep08, mRNA.
<a href="#">Pop4</a>	<a href="#">Pop4.cSep08</a>	<a href="#">292831</a>	5713	1048	5	159	processing of precursor 4, ribonuclease P/MRP family, (S. cerevisiae) (18.7 kD) (Pop4) alternative variant cSep08, mRNA.
<a href="#">Pop4</a>	<a href="#">Pop4.dSep08</a>	<a href="#">292831</a>	2257	1239	2	169	processing of precursor 4, ribonuclease P/MRP family, (S. cerevisiae) (18.8 kD) (Pop4) alternative variant dSep08, mRNA.

<a href="#">Pop5</a>	<a href="#">Pop5.aSep08</a>	<a href="#">117241</a>	3420	2095	2	174	processing of precursor 5, ribonuclease P/MRP family (S. cerevisiae) (Pop5) alternative variant aSep08, mRNA.
<a href="#">popor</a>	<a href="#">popor.aSep08</a>		558	465		155	probable E3 ubiquitin-protein ligase herc1 (popor) mRNA.
<a href="#">Por</a>	<a href="#">Por.bSep08</a>	<a href="#">29441</a>	70870	642	6	169	p450 (cytochrome) oxidoreductase (Por) alternative variant bSep08, mRNA.
<a href="#">Por</a>	<a href="#">Por.cSep08</a>	<a href="#">29441</a>	5283	381	3	54	p450 (cytochrome) oxidoreductase (Por) alternative variant cSep08, mRNA.
<a href="#">porby</a>	<a href="#">porby.aSep08</a>		7030	630		209	taf1 RNA polymerase ii TATA box binding protein - associated factor like (porby) mRNA.
<a href="#">porchy</a>	<a href="#">porchy.aSep08</a>		4825	414	4	138	cd44 (porchy) alternative variant aSep08, mRNA.
<a href="#">Porcn</a>	<a href="#">Porcn.aSep08</a>	<a href="#">317368</a>	12771	1410	11	360	porcupine homolog (Drosophila) (Porcn) alternative variant aSep08, mRNA.
<a href="#">Porcn</a>	<a href="#">Porcn.bSep08</a>	<a href="#">317368</a>	5474	811	8	269	porcupine homolog (Drosophila) (Porcn) alternative variant bSep08, mRNA.
<a href="#">Porcn</a>	<a href="#">Porcn.cSep08</a>	<a href="#">317368</a>	2707	536	4	168	porcupine homolog (Drosophila) (Porcn) alternative variant cSep08, mRNA.
<a href="#">Porcn</a>	<a href="#">Porcn.eSep08</a>	<a href="#">317368</a>	747	652	2	95	porcupine homolog (Drosophila) (Porcn) alternative variant eSep08, mRNA.
<a href="#">Porcn</a>	<a href="#">Porcn.fSep08</a>	<a href="#">317368</a>	2286	326	3	85	porcupine homolog (Drosophila) (Porcn) alternative variant fSep08, mRNA.
<a href="#">pordar</a>	<a href="#">pordar.aSep08</a>		7414	481		50	putative protein (5.6 kD) (pordar) mRNA.
<a href="#">porfer</a>	<a href="#">porfer.aSep08</a>		3225	903		55	putative protein (6.1 kD) (porfer) mRNA.
<a href="#">porflo</a>	<a href="#">porflo.aSep08</a>		2694	736		38	putative protein (4.2 kD) (porflo) mRNA.
<a href="#">porflu</a>	<a href="#">porflu.aSep08</a>		23702	493	4	70	putative protein (8.0 kD) (porflu) alternative variant aSep08, mRNA.
<a href="#">porflu</a>	<a href="#">porflu.bSep08</a>		67215	1793	6	53	putative protein (porflu) alternative variant bSep08, mRNA.
<a href="#">porflu</a>	<a href="#">porflu.dSep08</a>		1459	371	3	59	CRA a like (porflu) alternative variant dSep08, mRNA.
<a href="#">porgar</a>	<a href="#">porgar.aSep08</a>		2033	908		302	family with sequence similarity 170 member B like (porgar) mRNA.
<a href="#">porja</a>	<a href="#">porja.aSep08</a>		596	414		35	putative protein (porja) mRNA.
<a href="#">porjey</a>	<a href="#">porjey.aSep08</a>		1024	566		64	cc2-5 like (7.2 kD) (porjey) mRNA.
<a href="#">porkee</a>	<a href="#">porkee.aSep08</a>		9433	396		69	pappalysin 2 (porkee) mRNA.
<a href="#">porkler</a>	<a href="#">porkler.aSep08</a>		20132	573	1	141	protection of telomeres 1 (porkler) alternative variant aSep08, mRNA.
<a href="#">porkler</a>	<a href="#">porkler.bSep08</a>		20251	678	1	91	protection of telomeres 1 (10.4 kD) (porkler) alternative variant bSep08, mRNA.
<a href="#">porlo</a>	<a href="#">porlo.aSep08</a>		5769	351		43	putative protein (porlo) mRNA.
<a href="#">porloy</a>	<a href="#">porloy.aSep08</a>		4008	403	2	56	apolipoprotein D CRA a (porloy) alternative variant aSep08, mRNA.
<a href="#">porloy</a>	<a href="#">porloy.bSep08</a>		4037	333	1	81	putative protein (9.6 kD) (porloy) alternative variant bSep08, mRNA.
<a href="#">porloy</a>	<a href="#">porloy.cSep08</a>		4037	179	1	59	putative protein (porloy) alternative variant cSep08, mRNA.
<a href="#">pormee</a>	<a href="#">pormee.aSep08</a>		16026	1099		366	putative protein of metazoan origin (pormee) mRNA.
<a href="#">pormer</a>	<a href="#">pormer.aSep08</a>		36077	377		110	putative protein (pormer) mRNA.
<a href="#">pornoy</a>	<a href="#">pornoy.aSep08</a>		1735	380		44	serine threonine kinase 36 (pornoy) mRNA.

<a href="#">porpor</a>	<a href="#">porpor.aSep08</a>		7312	453		150	talin 2 (porpor) mRNA.
<a href="#">porsa</a>	<a href="#">porsa.aSep08</a>		1381	1083		90	binding protein 1 like (porsa) mRNA.
<a href="#">porshee</a>	<a href="#">porshee.aSep08</a>		9706	725		207	mediator of rna polymerase ii transcription homolog (porshee) mRNA.
<a href="#">portu</a>	<a href="#">portu.aSep08</a>		946	477	2	79	putative protein (9.0 kD) (portu) alternative variant aSep08, mRNA.
<a href="#">porvar</a>	<a href="#">porvar.aSep08</a>		15308	629		71	putative protein (7.8 kD) (porvar) mRNA.
<a href="#">porwey</a>	<a href="#">porwey.aSep08</a>		1578	516		57	putative protein (porwey) mRNA.
<a href="#">posa</a>	<a href="#">posa.aSep08</a>		1745	525		88	putative protein (9.9 kD) (posa) mRNA.
<a href="#">poshee</a>	<a href="#">poshee.bSep08</a>		104360	393	2	31	putative protein (3.8 kD) (poshee) alternative variant bSep08, mRNA.
<a href="#">Postn</a>	<a href="#">Postn.bSep08</a>	<a href="#">361945</a>	16177	1452	11	300	periostin, osteoblast specific factor (Postn) alternative variant bSep08, mRNA.
<a href="#">Postn</a>	<a href="#">Postn.cSep08</a>	<a href="#">361945</a>	13013	810	8	177	periostin, osteoblast specific factor (Postn) alternative variant cSep08, mRNA.
<a href="#">Postn</a>	<a href="#">Postn.dSep08</a>	<a href="#">361945</a>	13266	982	7	151	periostin, osteoblast specific factor (Postn) alternative variant dSep08, mRNA.
<a href="#">Postn</a>	<a href="#">Postn.eSep08</a>	<a href="#">361945</a>	7787	412	6	137	periostin, osteoblast specific factor (Postn) alternative variant eSep08, mRNA.
<a href="#">Postn</a>	<a href="#">Postn.fSep08</a>	<a href="#">361945</a>	4967	540	5	125	periostin, osteoblast specific factor (Postn) alternative variant fSep08, mRNA.
<a href="#">Pot1b</a>	<a href="#">Pot1b.aSep08</a>	<a href="#">690237</a>	22441	923	2	307	protection of telomeres 1B (Pot1b) alternative variant aSep08, mRNA.
<a href="#">Pot1b</a>	<a href="#">Pot1b.bSep08</a>	<a href="#">690237</a>	35776	689	1	227	protection of telomeres 1B (Pot1b) alternative variant bSep08, mRNA.
<a href="#">potu</a>	<a href="#">potu.aSep08</a>		10760	781		46	putative protein (potu) mRNA.
<a href="#">Pou.0</a>	<a href="#">Pou.0.aSep08</a>		24431	780		260	pou domain transcription factor Oct-1B (Pou.0) mRNA.
<a href="#">Pou1f1</a>	<a href="#">Pou1f1.aSep08</a>	<a href="#">25517</a>	21153	2426	1	317	POU class 1 homeobox 1 (35.8 kD) (Pou1f1) alternative variant aSep08, mRNA.
<a href="#">Pou2f1</a>	<a href="#">Pou2f1.aSep08</a>	<a href="#">171068</a>	95289	643	6	213	POU domain, class 2, transcription factor 1 (Pou2f1) alternative variant aSep08, mRNA.
<a href="#">Pou2f1</a>	<a href="#">Pou2f1.bSep08</a>	<a href="#">171068</a>	9763	459	2	44	POU domain, class 2, transcription factor 1 (4.7 kD) (Pou2f1) alternative variant bSep08, mRNA.
<a href="#">Pou4f1</a>	<a href="#">Pou4f1.aSep08</a>	<a href="#">114503</a>	1674	478		110	POU domain, class 4, transcription factor 1 (Pou4f1) alternative variant aSep08, mRNA.
<a href="#">povar</a>	<a href="#">povar.aSep08</a>		1665	306		101	putative protein (povar) mRNA.
<a href="#">powey</a>	<a href="#">powey.aSep08</a>		4353	377		88	ribosome assembly protein CRA b (powey) mRNA.
<a href="#">poyby</a>	<a href="#">poyby.aSep08</a>		488	407		129	putative protein (poyby) mRNA.
<a href="#">poychy</a>	<a href="#">poychy.aSep08</a>		17818	736		245	CRA a (poychy) mRNA.
<a href="#">poydar</a>	<a href="#">poydar.aSep08</a>		843	422		33	putative protein (poydar) mRNA.
<a href="#">poyfer</a>	<a href="#">poyfer.aSep08</a>		22797	425		89	putative protein (9.8 kD) (poyfer) mRNA.
<a href="#">poyflo</a>	<a href="#">poyflo.aSep08</a>		16529	270		52	putative protein (poyflo) mRNA.
<a href="#">poyflu</a>	<a href="#">poyflu.aSep08</a>		26473	501	2	101	CRA a like (poyflu) alternative variant aSep08, mRNA.
<a href="#">poyflu</a>	<a href="#">poyflu.bSep08</a>		5538	786	2	93	putative protein (10.3 kD) (poyflu) alternative variant bSep08, mRNA.

<a href="#">poygar</a>	<a href="#">poygar.aSep08</a>		32831	704		234	putative protein, with a transmembrane domain, of vertebrate origin (poygar) mRNA.
<a href="#">poyja</a>	<a href="#">poyja.aSep08</a>		3475	1507		62	putative protein (6.9 kD) (poyja) mRNA.
<a href="#">poyjey</a>	<a href="#">poyjey.aSep08</a>		2319	530		44	putative protein (poyjey) mRNA.
<a href="#">poykee</a>	<a href="#">poykee.aSep08</a>		30824	406		135	pappalysin 2 (poykee) mRNA.
<a href="#">poykler</a>	<a href="#">poykler.aSep08</a>		1063	189		62	putative protein (poykler) mRNA.
<a href="#">poylo</a>	<a href="#">poylo.aSep08</a>		971	349		68	putative protein (poylo) mRNA.
<a href="#">poymee</a>	<a href="#">poymee.aSep08</a>		12527	651		216	putative protein, with 2 coiled coil domains, of bilateral origin (poymee) mRNA.
<a href="#">poymer</a>	<a href="#">poymer.aSep08</a>		10385	463		45	putative protein (4.8 kD) (poymer) mRNA.
<a href="#">poynoy</a>	<a href="#">poynoy.aSep08</a>		4085	417		139	putative protein of mammalian origin (poynoy) mRNA.
<a href="#">poypor</a>	<a href="#">poypor.aSep08</a>		11353	372		104	talin 2 (poypor) mRNA.
<a href="#">poysa</a>	<a href="#">poysa.aSep08</a>		610	499		64	putative protein (poysa) mRNA.
<a href="#">poyshee</a>	<a href="#">poyshee.aSep08</a>		1073	422		140	mediator of rna polymerase ii transcription (poyshee) mRNA.
<a href="#">poytu</a>	<a href="#">poytu.aSep08</a>		34196	519		114	putative protein (poytu) mRNA.
<a href="#">poyvar</a>	<a href="#">poyvar.aSep08</a>		2900	665	5	189	absent in melanoma 1-like (poyvar) alternative variant aSep08, mRNA.
<a href="#">poywey</a>	<a href="#">poywey.aSep08</a>		6915	566	2	75	putative nuclear protein (8.6 kD) (poywey) alternative variant aSep08, mRNA.
<a href="#">Pp11r</a>	<a href="#">Pp11r.aSep08</a>	<a href="#">680317</a>	12998	1777		495	placental protein 11 related (Pp11r) alternative variant aSep08, mRNA.
<a href="#">Ppa1</a>	<a href="#">Ppa1.aSep08</a>	<a href="#">294504</a>	26758	1318	11	333	pyrophosphatase (inorganic) 1 (Ppa1) alternative variant aSep08, mRNA.
<a href="#">Ppa1</a>	<a href="#">Ppa1.bSep08</a>	<a href="#">294504</a>	19139	881	8	198	pyrophosphatase (inorganic) 1 (22.6 kD) (Ppa1) alternative variant bSep08, mRNA.
<a href="#">Ppa2</a>	<a href="#">Ppa2.aSep08</a>	<a href="#">310856</a>	78357	1158	12	330	pyrophosphatase (inorganic) 2 (37.8 kD) (Ppa2) alternative variant aSep08, complete mRNA.
<a href="#">Ppa2</a>	<a href="#">Ppa2.bSep08</a>	<a href="#">310856</a>	73939	738	8	219	pyrophosphatase (inorganic) 2 (Ppa2) alternative variant bSep08, mRNA.
<a href="#">Ppa2</a>	<a href="#">Ppa2.cSep08</a>	<a href="#">310856</a>	1950	664	2	55	pyrophosphatase (inorganic) 2 (5.9 kD) (Ppa2) alternative variant cSep08, mRNA.
<a href="#">Ppa2</a>	<a href="#">Ppa2.dSep08</a>	<a href="#">310856</a>	9477	421	2	29	pyrophosphatase (inorganic) 2 (3.5 kD) (Ppa2) alternative variant dSep08, mRNA.
<a href="#">PPAK.0</a>	<a href="#">PPAK.0.aSep08</a>		2599	595		198	titin (PPAK.0) mRNA.
<a href="#">Ppan</a>	<a href="#">Ppan.aSep08</a>	<a href="#">298699</a>	3963	1797	4	553	peter pan homolog (Drosophila) (Ppan) alternative variant aSep08, mRNA.
<a href="#">Ppan</a>	<a href="#">Ppan.cSep08</a>	<a href="#">298699</a>	2490	834	4	277	peter pan homolog (Drosophila) (Ppan) alternative variant cSep08, mRNA.
<a href="#">Ppan</a>	<a href="#">Ppan.dSep08</a>	<a href="#">298699</a>	2446	730	2	243	peter pan homolog (Drosophila) (Ppan) alternative variant dSep08, mRNA.
<a href="#">Ppan</a>	<a href="#">Ppan.eSep08</a>	<a href="#">298699</a>	935	682	4	178	peter pan homolog (Drosophila) (Ppan) alternative variant eSep08, mRNA.
<a href="#">Ppap2a</a>	<a href="#">Ppap2a.bSep08</a>	<a href="#">64369</a>	64445	1225	6	326	phosphatidic acid phosphatase 2a (Ppap2a) alternative variant bSep08, mRNA.

<a href="#">Ppap2a</a>	<a href="#">Ppap2a.cSep08</a>	<a href="#">64369</a>	63588	764	4	136	phosphatidic acid phosphatase 2a (Ppap2a) alternative variant cSep08, mRNA.
<a href="#">Ppap2c</a>	<a href="#">Ppap2c.aSep08</a>	<a href="#">246115</a>	7654	1138	1	279	phosphatidic acid phosphatase type 2c (Ppap2c) alternative variant aSep08, mRNA.
<a href="#">Ppapdc1a</a>	<a href="#">Ppapdc1a.aSep08</a>	<a href="#">309014</a>	62715	527		175	putative protein, with at least 2 transmembrane domains, of eukaryotic origin (Ppapdc1a) mRNA.
<a href="#">Ppapdc1b</a>	<a href="#">Ppapdc1b.bSep08</a>	<a href="#">680466</a>	3523	826	6	251	phosphoesterase, PA-phosphatase related (Ppapdc1b) alternative variant bSep08, mRNA.
<a href="#">Ppapdc1b</a>	<a href="#">Ppapdc1b.cSep08</a>	<a href="#">680466</a>	3662	1777	4	212	putative protein, with a transmembrane domain, of eukaryotic origin (Ppapdc1b) alternative variant cSep08, mRNA.
<a href="#">Ppapdc1b</a>	<a href="#">Ppapdc1b.dSep08</a>	<a href="#">680466</a>	1734	753	4	161	putative protein, with at least 2 transmembrane domains, of eukaryotic origin (Ppapdc1b) alternative variant dSep08, mRNA.
<a href="#">Ppapdc1b</a>	<a href="#">Ppapdc1b.eSep08</a>	<a href="#">680466</a>	4097	1801	5	160	putative protein, with at least 2 transmembrane domains, of eukaryotic origin (Ppapdc1b) alternative variant eSep08, mRNA.
<a href="#">Ppapdc1b</a>	<a href="#">Ppapdc1b.fSep08</a>	<a href="#">680466</a>	1308	439	3	146	putative protein, with a transmembrane domain, of eukaryotic origin (Ppapdc1b) alternative variant fSep08, mRNA.
<a href="#">Ppard</a>	<a href="#">Ppard.bSep08</a>	<a href="#">25682</a>	15737	328	2	48	peroxisome proliferator-activated receptor delta (5.0 kD) (Ppard) alternative variant bSep08, mRNA.
<a href="#">Pparg</a>	<a href="#">Pparg.bSep08</a>	<a href="#">25664</a>	96675	778	6	215	peroxisome proliferator activated receptor gamma (Pparg) alternative variant bSep08, mRNA.
<a href="#">Pparg</a>	<a href="#">Pparg.cSep08</a>	<a href="#">25664</a>	97958	1012	6	205	peroxisome proliferator activated receptor gamma (Pparg) alternative variant cSep08, mRNA.
<a href="#">Pparg</a>	<a href="#">Pparg.dSep08</a>	<a href="#">25664</a>	47520	619	4	100	peroxisome proliferator activated receptor gamma (Pparg) alternative variant dSep08, mRNA.
<a href="#">Ppargc1a</a>	<a href="#">Ppargc1a.bSep08</a>	<a href="#">83516</a>	24686	504	3	85	peroxisome proliferative activated receptor, gamma, coactivator 1 alpha (9.5 kD) (Ppargc1a) alternative variant bSep08, mRNA.
<a href="#">Ppargc1a</a>	<a href="#">Ppargc1a.dSep08</a>	<a href="#">83516</a>	23694	525	3	70	peroxisome proliferative activated receptor, gamma, coactivator 1 alpha (7.7 kD) (Ppargc1a) alternative variant dSep08, mRNA.
<a href="#">Ppat</a>	<a href="#">Ppat.aSep08</a>	<a href="#">117544</a>	34397	4053	8	517	phosphoribosyl pyrophosphate amidotransferase (57.4 kD) (Ppat) alternative variant aSep08, mRNA.
<a href="#">Ppat</a>	<a href="#">Ppat.bSep08</a>	<a href="#">117544</a>	25226	857	4	285	phosphoribosyl pyrophosphate amidotransferase (Ppat) alternative variant bSep08, mRNA.
<a href="#">Ppat</a>	<a href="#">Ppat.cSep08</a>	<a href="#">117544</a>	7212	880	5	268	phosphoribosyl pyrophosphate amidotransferase (Ppat) alternative variant cSep08, mRNA.
<a href="#">Ppcdc</a>	<a href="#">Ppcdc.bSep08</a>	<a href="#">363069</a>	20690	807	5	216	phosphopantothenoylcysteine decarboxylase (Ppcdc) alternative variant bSep08, mRNA.
<a href="#">Ppcdc</a>	<a href="#">Ppcdc.cSep08</a>	<a href="#">363069</a>	5716	681	3	144	phosphopantothenoylcysteine decarboxylase (Ppcdc) alternative variant cSep08, mRNA.
<a href="#">Ppef1</a>	<a href="#">Ppef1.bSep08</a>	<a href="#">317498</a>	13207	548	3	182	protein phosphatase with EF hand calcium-binding domain 1 (Ppef1) alternative variant bSep08, mRNA.

<a href="#">Ppfia1</a>	<a href="#">Ppfia1.bSep08</a>	<a href="#">293645</a>	24351	1744	12	310	protein interacting alpha 1 (35.3 kD) (Ppfia1) alternative variant bSep08, mRNA.
<a href="#">Ppfia1</a>	<a href="#">Ppfia1.cSep08</a>	<a href="#">293645</a>	4908	763	4	149	putative protein (Ppfia1) alternative variant cSep08, mRNA.
<a href="#">Ppfia1</a>	<a href="#">Ppfia1.eSep08</a>	<a href="#">293645</a>	6571	379	4	126	protein tyrosine phosphatase receptor type f polypeptide interacting alpha (Ppfia1) alternative variant eSep08, mRNA.
<a href="#">Ppfia1</a>	<a href="#">Ppfia1.fSep08</a>	<a href="#">293645</a>	3836	373	3	123	protein tyrosine phosphatase receptor type f polypeptide interacting alpha 1 (Ppfia1) alternative variant fSep08, mRNA.
<a href="#">Ppfia1</a>	<a href="#">Ppfia1.gSep08</a>	<a href="#">293645</a>	558	391	2	122	protein interacting alpha 1 (Ppfia1) alternative variant gSep08, mRNA.
<a href="#">Ppfia2</a>	<a href="#">Ppfia2.bSep08</a>	<a href="#">362876</a>	21344	456	2	74	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2 (8.2 kD) (Ppfia2) alternative variant bSep08, mRNA.
<a href="#">Ppfia3</a>	<a href="#">Ppfia3.aSep08</a>	<a href="#">140591</a>	1344	744		156	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 3 (17.1 kD) (Ppfia3) mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.aSep08</a>	<a href="#">140592</a>	5041	1330	7	443	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant aSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.aSep08</a>	<a href="#">685423</a>	5041	1330	7	443	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant aSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.bSep08</a>	<a href="#">140592</a>	17666	2732	11	367	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant bSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.bSep08</a>	<a href="#">685423</a>	17666	2732	11	367	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant bSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.cSep08</a>	<a href="#">140592</a>	4057	594	5	197	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant cSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.cSep08</a>	<a href="#">685423</a>	4057	594	5	197	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant cSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.dSep08</a>	<a href="#">140592</a>	1100	640	2	167	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant dSep08, mRNA.

<a href="#">Ppfia4</a>	<a href="#">Ppfia4.dSep08</a>	<a href="#">685423</a>	1100	640	2	167	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant dSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.eSep08</a>	<a href="#">140592</a>	47724	1416	3	127	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant eSep08, mRNA.
<a href="#">Ppfia4</a>	<a href="#">Ppfia4.eSep08</a>	<a href="#">685423</a>	47724	1416	3	127	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4 and hypothetical protein LOC685423 (Ppfia4) alternative variant eSep08, mRNA.
<a href="#">Ppfibp1</a>	<a href="#">Ppfibp1.aSep08</a>	<a href="#">312855</a>	103251	684		157	PTPRF interacting protein, binding protein 1 (liprin beta 1) (Ppfibp1) mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.aSep08</a>	<a href="#">308918</a>	76852	2980	22	799	protein tyrosine phosphatase receptor-type F interacting binding 2 CRA b like (90.3 kD) (Ppfibp2) alternative variant aSep08, mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.bSep08</a>	<a href="#">308918</a>	100296	891	6	197	protein interacting binding 2 like (Ppfibp2) alternative variant bSep08, mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.cSep08</a>	<a href="#">308918</a>	2755	466	3	155	protein tyrosine phosphatase receptor-type F interacting binding 2 like (Ppfibp2) alternative variant cSep08, mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.dSep08</a>	<a href="#">308918</a>	8189	925	4	148	protein tyrosine phosphatase receptor-type F interacting binding 2 like (Ppfibp2) alternative variant dSep08, mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.eSep08</a>	<a href="#">308918</a>	17532	406	5	90	protein tyrosine phosphatase receptor-type F interacting binding 2 CRA a like (10.6 kD) (Ppfibp2) alternative variant eSep08, mRNA.
<a href="#">Ppfibp2</a>	<a href="#">Ppfibp2.fSep08</a>	<a href="#">308918</a>	6234	796	3	71	putative secreted or extracellular protein precursor (8.0 kD) (Ppfibp2) alternative variant fSep08, mRNA.
<a href="#">Ppgeb</a>	<a href="#">Ppgeb.bSep08</a>	<a href="#">296370</a>	6106	2216	15	493	protective protein for beta-galactosidase (55.8 kD) (Ppgeb) alternative variant bSep08, mRNA.
<a href="#">Ppgeb</a>	<a href="#">Ppgeb.cSep08</a>	<a href="#">296370</a>	3341	1059	9	253	protective protein for beta-galactosidase (Ppgeb) alternative variant cSep08, mRNA.
<a href="#">Ppgeb</a>	<a href="#">Ppgeb.dSep08</a>	<a href="#">296370</a>	3127	743	8	210	protective protein for beta-galactosidase (23.6 kD) (Ppgeb) alternative variant dSep08, mRNA.
<a href="#">Ppgeb</a>	<a href="#">Ppgeb.eSep08</a>	<a href="#">296370</a>	2222	703	4	116	protective protein for beta-galactosidase (Ppgeb) alternative variant eSep08, mRNA.
<a href="#">Pphln1</a>	<a href="#">Pphln1.bSep08</a>	<a href="#">366975</a>	82925	1787	1	359	periphilin 1 (Pphln1) alternative variant bSep08, complete mRNA.
<a href="#">Pphln1</a>	<a href="#">Pphln1.cSep08</a>	<a href="#">366975</a>	82925	3596	2	312	periphilin 1 (36.0 kD) (Pphln1) alternative variant cSep08, complete mRNA.
<a href="#">Ppia</a>	<a href="#">Ppia.bSep08</a>	<a href="#">25518</a>	3967	2640	4	104	peptidylprolyl isomerase A (11.3 kD) (Ppia) alternative variant bSep08, complete mRNA.
<a href="#">Ppib</a>	<a href="#">Ppib.bSep08</a>	<a href="#">64367</a>	3331	738	3	116	peptidylprolyl isomerase B (12.8 kD) (Ppib) alternative variant bSep08, mRNA.
<a href="#">Ppib</a>	<a href="#">Ppib.cSep08</a>	<a href="#">64367</a>	1396	730	2	77	peptidylprolyl isomerase B (8.5 kD) (Ppib) alternative variant cSep08, mRNA.



<a href="#">Ppic</a>	<a href="#">Ppic.bSep08</a>	<a href="#">291463</a>	6100	840	2	118	peptidylprolyl isomerase C (12.9 kD) (Ppic) alternative variant bSep08, mRNA.
<a href="#">Ppic</a>	<a href="#">Ppic.cSep08</a>	<a href="#">291463</a>	12134	878	4	118	peptidylprolyl isomerase C (12.9 kD) (Ppic) alternative variant cSep08, mRNA.
<a href="#">Ppid</a>	<a href="#">Ppid.bSep08</a>	<a href="#">361967</a>	2937	1889	3	132	peptidylprolyl isomerase D (14.5 kD) (Ppid) alternative variant bSep08, mRNA.
<a href="#">Ppid</a>	<a href="#">Ppid.cSep08</a>	<a href="#">361967</a>	4479	785	3	116	peptidylprolyl isomerase D (12.8 kD) (Ppid) alternative variant cSep08, mRNA.
<a href="#">Ppie</a>	<a href="#">Ppie.aSep08</a>	<a href="#">298508</a>	13288	1184	10	301	peptidylprolyl isomerase E (cyclophilin E) (33.4 kD) (Ppie) alternative variant aSep08, complete mRNA.
<a href="#">Ppie</a>	<a href="#">Ppie.cSep08</a>	<a href="#">298508</a>	3297	752	4	100	peptidylprolyl isomerase E (cyclophilin E) (Ppie) alternative variant cSep08, mRNA.
<a href="#">Ppig</a>	<a href="#">Ppig.bSep08</a>	<a href="#">83624</a>	26533	1441	14	418	peptidylprolyl isomerase G (Ppig) alternative variant bSep08, mRNA.
<a href="#">Ppig</a>	<a href="#">Ppig.cSep08</a>	<a href="#">83624</a>	6353	401	3	133	peptidylprolyl isomerase G (Ppig) alternative variant cSep08, mRNA.
<a href="#">Ppig</a>	<a href="#">Ppig.dSep08</a>	<a href="#">83624</a>	11060	745	7	119	peptidylprolyl isomerase G (Ppig) alternative variant dSep08, mRNA.
<a href="#">Ppig</a>	<a href="#">Ppig.fSep08</a>	<a href="#">83624</a>	810	360	2	33	peptidylprolyl isomerase G (Ppig) alternative variant fSep08, mRNA.
<a href="#">Ppil1</a>	<a href="#">Ppil1.bSep08</a>	<a href="#">309651</a>	9051	805	1	121	peptidylprolyl isomerase (cyclophilin)-like 1 (13.1 kD) (Ppil1) alternative variant bSep08, mRNA.
<a href="#">Ppil1</a>	<a href="#">Ppil1.cSep08</a>	<a href="#">309651</a>	11820	934	1	67	peptidylprolyl isomerase (cyclophilin)-like 1 (7.2 kD) (Ppil1) alternative variant cSep08, mRNA.
<a href="#">Ppil1</a>	<a href="#">Ppil1.dSep08</a>	<a href="#">309651</a>	7944	620	1	67	peptidylprolyl isomerase (cyclophilin)-like 1 (7.2 kD) (Ppil1) alternative variant dSep08, mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.aSep08</a>	<a href="#">360746</a>	22579	1778	20	526	peptidylprolyl isomerase (cyclophilin)-like 2 (59.5 kD) (Ppil2) alternative variant aSep08, complete mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.bSep08</a>	<a href="#">360746</a>	18360	858	12	267	peptidylprolyl isomerase (cyclophilin)-like 2 (Ppil2) alternative variant bSep08, mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.cSep08</a>	<a href="#">360746</a>	2841	529	6	175	peptidylprolyl isomerase (cyclophilin)-like 2 (Ppil2) alternative variant cSep08, mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.dSep08</a>	<a href="#">360746</a>	7840	502	5	161	peptidylprolyl isomerase (cyclophilin)-like 2 (Ppil2) alternative variant dSep08, mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.eSep08</a>	<a href="#">360746</a>	16716	775	11	118	peptidylprolyl isomerase (cyclophilin)-like 2 (13.5 kD) (Ppil2) alternative variant eSep08, mRNA.
<a href="#">Ppil2</a>	<a href="#">Ppil2.fSep08</a>	<a href="#">360746</a>	1308	736	2	42	peptidylprolyl isomerase (cyclophilin)-like 2 (Ppil2) alternative variant fSep08, mRNA.
<a href="#">Ppil3</a>	<a href="#">Ppil3.bSep08</a>	<a href="#">301432</a>	9868	439	1	121	peptidylprolyl isomerase (cyclophilin)-like 3 (Ppil3) alternative variant bSep08, mRNA.
<a href="#">Ppil4</a>	<a href="#">Ppil4.aSep08</a>	<a href="#">361449</a>	26366	1136	2	378	peptidylprolyl isomerase (cyclophilin)-like 4 (Ppil4) alternative variant aSep08, mRNA.
<a href="#">Ppil6</a>	<a href="#">Ppil6.aSep08</a>	<a href="#">685567</a>	19810	740		230	peptidylprolyl isomerase (cyclophilin)-like 6 (Ppil6) mRNA.
<a href="#">Ppm1a</a>	<a href="#">Ppm1a.bSep08</a>	<a href="#">24666</a>	5771	3456	2	280	protein phosphatase 1A, magnesium dependent, alpha isoform (Ppm1a) alternative variant bSep08, mRNA.

<a href="#">Ppm1a</a>	<a href="#">Ppm1a.cSep08</a>	<a href="#">24666</a>	41510	1496	6	99	protein phosphatase 1A, magnesium dependent, alpha isoform (11.0 kD) (Ppm1a) alternative variant cSep08, mRNA.
<a href="#">Ppm1a</a>	<a href="#">Ppm1a.dSep08</a>	<a href="#">24666</a>	2662	1791	2	87	protein phosphatase 1A, magnesium dependent, alpha isoform (9.5 kD) (Ppm1a) alternative variant dSep08, mRNA.
<a href="#">Ppm1b</a>	<a href="#">Ppm1b.bSep08</a>	<a href="#">24667</a>	21921	1897	5	375	protein phosphatase 1B, magnesium dependent, beta isoform (Ppm1b) alternative variant bSep08, mRNA.
<a href="#">Ppm1b</a>	<a href="#">Ppm1b.cSep08</a>	<a href="#">24667</a>	1125	756	2	71	protein phosphatase 1B, magnesium dependent, beta isoform (Ppm1b) alternative variant cSep08, mRNA.
<a href="#">Ppm1b</a>	<a href="#">Ppm1b.dSep08</a>	<a href="#">24667</a>	19689	443	3	63	protein phosphatase 1B, magnesium dependent, beta isoform (Ppm1b) alternative variant dSep08, mRNA.
<a href="#">Ppm1b</a>	<a href="#">Ppm1b.eSep08</a>	<a href="#">24667</a>	19670	472	4	59	protein phosphatase 1B, magnesium dependent, beta isoform (Ppm1b) alternative variant eSep08, mRNA.
<a href="#">Ppm1d</a>	<a href="#">Ppm1d.bSep08</a>	<a href="#">287585</a>	28849	808	1	256	protein phosphatase 1D magnesium-dependent, delta isoform (Ppm1d) alternative variant bSep08, mRNA.
<a href="#">Ppm1f</a>	<a href="#">Ppm1f.bSep08</a>	<a href="#">287931</a>	18873	811	5	206	protein phosphatase 1F (Ppm1f) alternative variant bSep08, mRNA.
<a href="#">Ppm1g</a>	<a href="#">Ppm1g.aSep08</a>	<a href="#">259229</a>	1658	1235		99	protein phosphatase 1G (formerly 2C), magnesium-dependent, gamma isoform (10.9 kD) (Ppm1g) mRNA.
<a href="#">Ppm1h</a>	<a href="#">Ppm1h.aSep08</a>	<a href="#">314897</a>	127516	806	7	249	protein phosphatase 1h (Ppm1h) alternative variant aSep08, mRNA.
<a href="#">Ppm1h</a>	<a href="#">Ppm1h.bSep08</a>	<a href="#">314897</a>	63616	1862	5	194	protein phosphatase 1H (21.8 kD) (Ppm1h) alternative variant bSep08, mRNA.
<a href="#">Ppm1k</a>	<a href="#">Ppm1k.bSep08</a>	<a href="#">312381</a>	24626	1800	3	268	protein phosphatase 1K CRA b (Ppm1k) alternative variant bSep08, mRNA.
<a href="#">Ppm1k</a>	<a href="#">Ppm1k.cSep08</a>	<a href="#">312381</a>	9287	765	1	142	protein phosphatase 1K CRA a (Ppm1k) alternative variant cSep08, mRNA.
<a href="#">Ppm2c</a>	<a href="#">Ppm2c.bSep08</a>	<a href="#">54705</a>	6778	2496	3	579	protein phosphatase 2C, magnesium dependent, catalytic subunit (Ppm2c) alternative variant bSep08, mRNA.
<a href="#">Ppm2c</a>	<a href="#">Ppm2c.cSep08</a>	<a href="#">54705</a>	4168	446	3	124	protein phosphatase 2C, magnesium dependent, catalytic subunit (Ppm2c) alternative variant cSep08, mRNA.
<a href="#">Ppme1</a>	<a href="#">Ppme1.aSep08</a>	<a href="#">361613</a>	48316	2461		386	protein phosphatase methylesterase 1 (42.3 kD) (Ppme1) mRNA.
<a href="#">Ppox</a>	<a href="#">Ppox.bSep08</a>	<a href="#">289219</a>	4133	2374	8	339	protoporphyrinogen oxidase (36.4 kD) (Ppox) alternative variant bSep08, mRNA.
<a href="#">Ppox</a>	<a href="#">Ppox.cSep08</a>	<a href="#">289219</a>	2155	1612	5	172	protoporphyrinogen oxidase (18.3 kD) (Ppox) alternative variant cSep08, mRNA.
<a href="#">Ppox</a>	<a href="#">Ppox.dSep08</a>	<a href="#">289219</a>	1600	710	6	148	protoporphyrinogen oxidase (Ppox) alternative variant dSep08, mRNA.
<a href="#">Ppox</a>	<a href="#">Ppox.eSep08</a>	<a href="#">289219</a>	446	291	2	61	protoporphyrinogen oxidase (Ppox) alternative variant eSep08, mRNA.
<a href="#">Ppp1cb</a>	<a href="#">Ppp1cb.bSep08</a>	<a href="#">25594</a>	24612	909	7	243	protein phosphatase 1, catalytic subunit, beta isoform (Ppp1cb) alternative variant bSep08, mRNA.
<a href="#">Ppp1cb</a>	<a href="#">Ppp1cb.cSep08</a>	<a href="#">25594</a>	30191	966	4		
<a href="#">Ppp1cc</a>	<a href="#">Ppp1cc.aSep08</a>	<a href="#">24669</a>	18169	1410	8	337	protein phosphatase 1 catalytic (38.5 kD) (Ppp1cc) alternative variant aSep08, complete mRNA.

<a href="#">Ppp1cc</a>	<a href="#">Ppp1cc.cSep08</a>	<a href="#">24669</a>	15819	879	5	263	protein phosphatase 1 (Ppp1cc) alternative variant cSep08, mRNA.
<a href="#">Ppp1cc</a>	<a href="#">Ppp1cc.dSep08</a>	<a href="#">24669</a>	15660	725	5	147	protein phosphatase 1 (17.0 kD) (Ppp1cc) alternative variant dSep08, mRNA.
<a href="#">Ppp1cc</a>	<a href="#">Ppp1cc.eSep08</a>	<a href="#">24669</a>	1480	737	2	118	protein phosphatase 1 catalytic (Ppp1cc) alternative variant eSep08, mRNA.
<a href="#">Ppp1r1a</a>	<a href="#">Ppp1r1a.bSep08</a>	<a href="#">58977</a>	3427	481	6	160	protein phosphatase 1, regulatory (inhibitor) subunit 1A (Ppp1r1a) alternative variant bSep08, mRNA.
<a href="#">Ppp1r1a</a>	<a href="#">Ppp1r1a.cSep08</a>	<a href="#">58977</a>	1839	641	3	97	protein phosphatase 1, regulatory (inhibitor) subunit 1A (10.3 kD) (Ppp1r1a) alternative variant cSep08, mRNA.
<a href="#">Ppp1r1a</a>	<a href="#">Ppp1r1a.dSep08</a>	<a href="#">58977</a>	998	601	2	96	protein phosphatase 1, regulatory (inhibitor) subunit 1A (Ppp1r1a) alternative variant dSep08, mRNA.
<a href="#">Ppp1r1b</a>	<a href="#">Ppp1r1b.aSep08</a>	<a href="#">360616</a>	9715	1761		205	protein phosphatase 1, regulatory (inhibitor) subunit 1B (22.9 kD) (Ppp1r1b) mRNA.
<a href="#">Ppp1r2</a>	<a href="#">Ppp1r2.bSep08</a>	<a href="#">192361</a>	4017	617	3	105	protein phosphatase 1, regulatory (inhibitor) subunit 2 (Ppp1r2) alternative variant bSep08, mRNA.
<a href="#">Ppp1r3b</a>	<a href="#">Ppp1r3b.bSep08</a>	<a href="#">192280</a>	7584	344	2	62	protein phosphatase 1, regulatory (inhibitor) subunit 3B (Ppp1r3b) alternative variant bSep08, mRNA.
<a href="#">Ppp1r3f_predicted</a>	<a href="#">Ppp1r3f_predicted.aSep08</a>	<a href="#">363453</a>	14239	316		105	protein phosphatase 1, regulatory (inhibitor) subunit 3F (predicted) (Ppp1r3f_predicted) mRNA.
<a href="#">Ppp1r8</a>	<a href="#">Ppp1r8.bSep08</a>	<a href="#">313030</a>	12384	668	5	218	protein phosphatase 1, regulatory (inhibitor) subunit 8 (Ppp1r8) alternative variant bSep08, mRNA.
<a href="#">Ppp1r9a</a>	<a href="#">Ppp1r9a.bSep08</a>	<a href="#">84685</a>	4312	1199	2	141	protein phosphatase 1, regulatory (inhibitor) subunit 9A (16.1 kD) (Ppp1r9a) alternative variant bSep08, mRNA.
<a href="#">Ppp1r9b</a>	<a href="#">Ppp1r9b.aSep08</a>	<a href="#">84686</a>	12613	1489	3	496	protein phosphatase 1, regulatory subunit 9B (Ppp1r9b) alternative variant aSep08, mRNA.
<a href="#">Ppp1r9b</a>	<a href="#">Ppp1r9b.bSep08</a>	<a href="#">84686</a>	4790	1783	1	433	protein phosphatase 1, regulatory subunit 9B (Ppp1r9b) alternative variant bSep08, mRNA.
<a href="#">Ppp1r9b</a>	<a href="#">Ppp1r9b.cSep08</a>	<a href="#">84686</a>	7146	1170	1	389	protein phosphatase 1, regulatory subunit 9B (Ppp1r9b) alternative variant cSep08, mRNA.
<a href="#">Ppp1r10</a>	<a href="#">Ppp1r10.bSep08</a>	<a href="#">65045</a>	12255	2683	17	654	protein phosphatase 1 regulatory (71.1 kD) (Ppp1r10) alternative variant bSep08, mRNA.
<a href="#">Ppp1r10</a>	<a href="#">Ppp1r10.cSep08</a>	<a href="#">65045</a>	1462	1202	2	128	protein phosphatase 1 regulatory (13.3 kD) (Ppp1r10) alternative variant cSep08, mRNA.
<a href="#">Ppp1r11</a>	<a href="#">Ppp1r11.bSep08</a>	<a href="#">294207</a>	2459	804	3	114	protein phosphatase 1, regulatory (inhibitor) subunit 11 (12.5 kD) (Ppp1r11) alternative variant bSep08, mRNA.
<a href="#">Ppp1r11</a>	<a href="#">Ppp1r11.cSep08</a>	<a href="#">294207</a>	1127	757	2	114	protein phosphatase 1, regulatory (inhibitor) subunit 11 (12.5 kD) (Ppp1r11) alternative variant cSep08, mRNA.
<a href="#">Ppp1r11</a>	<a href="#">Ppp1r11.dSep08</a>	<a href="#">294207</a>	2298	749	3	106	protein phosphatase 1, regulatory (inhibitor) subunit 11 (12.0 kD) (Ppp1r11) alternative variant dSep08, mRNA.
<a href="#">Ppp1r12a</a>	<a href="#">Ppp1r12a.bSep08</a>	<a href="#">116670</a>	12498	1711	8	413	protein phosphatase 1, regulatory (inhibitor) subunit 12A (Ppp1r12a) alternative variant bSep08, mRNA.
<a href="#">Ppp1r12a</a>	<a href="#">Ppp1r12a.cSep08</a>	<a href="#">116670</a>	16782	3150	7	152	protein phosphatase 1, regulatory (inhibitor) subunit 12A (17.9 kD) (Ppp1r12a) alternative variant cSep08, mRNA.
<a href="#">Ppp1r12a</a>	<a href="#">Ppp1r12a.dSep08</a>	<a href="#">116670</a>	17326	887	10	127	protein phosphatase 1, regulatory (inhibitor) subunit 12A (14.7 kD) (Ppp1r12a) alternative variant dSep08, mRNA.

<a href="#">Ppp1r12a</a>	<a href="#">Ppp1r12a.fSep08</a>	<a href="#">116670</a>	1470	436	2	51	protein phosphatase 1, regulatory (inhibitor) subunit 12A (Ppp1r12a) alternative variant fSep08, mRNA.
<a href="#">Ppp1r12b</a>	<a href="#">Ppp1r12b.aSep08</a>	<a href="#">304813</a>	31528	1104		367	protein phosphatase 1, regulatory (inhibitor) subunit 12B (Ppp1r12b) mRNA.
<a href="#">Ppp1r12c</a>	<a href="#">Ppp1r12c.aSep08</a>	<a href="#">499076</a>	23794	3857	22	318	protein phosphatase 1, regulatory (inhibitor) subunit 12C (35.8 kD) (Ppp1r12c) alternative variant aSep08, complete mRNA.
<a href="#">Ppp1r12c</a>	<a href="#">Ppp1r12c.bSep08</a>	<a href="#">499076</a>	1144	464	7	154	protein phosphatase 1, regulatory (inhibitor) subunit 12C (Ppp1r12c) alternative variant bSep08, mRNA.
<a href="#">Ppp1r12c</a>	<a href="#">Ppp1r12c.cSep08</a>	<a href="#">499076</a>	1452	734	8	143	protein phosphatase 1, regulatory (inhibitor) subunit 12C (Ppp1r12c) alternative variant cSep08, mRNA.
<a href="#">Ppp1r12c</a>	<a href="#">Ppp1r12c.dSep08</a>	<a href="#">499076</a>	1209	1120	2	50	protein phosphatase 1, regulatory (inhibitor) subunit 12C (Ppp1r12c) alternative variant dSep08, mRNA.
<a href="#">Ppp1r14a</a>	<a href="#">Ppp1r14a.aSep08</a>	<a href="#">114004</a>	5743	676	5	192	protein phosphatase 1, regulatory (inhibitor) subunit 14A (Ppp1r14a) alternative variant aSep08, mRNA.
<a href="#">Ppp1r15b</a>	<a href="#">Ppp1r15b.bSep08</a>	<a href="#">304799</a>	6129	2028	2	178	protein phosphatase 1, regulatory (inhibitor) subunit 15b (Ppp1r15b) alternative variant bSep08, mRNA.
<a href="#">Ppp1r16a</a>	<a href="#">Ppp1r16a.bSep08</a>	<a href="#">362944</a>	1084	781	4	171	protein phosphatase 1, regulatory (inhibitor) subunit 16A (Ppp1r16a) alternative variant bSep08, mRNA.
<a href="#">Ppp1r16a</a>	<a href="#">Ppp1r16a.bSep08</a>	<a href="#">681165</a>	1084	781	4	171	protein phosphatase 1, regulatory (inhibitor) subunit 16A (Ppp1r16a) alternative variant bSep08, mRNA.
<a href="#">Ppp1r16a</a>	<a href="#">Ppp1r16a.dSep08</a>	<a href="#">362944</a>	6100	498	3	32	protein phosphatase 1, regulatory (inhibitor) subunit 16A (3.5 kD) (Ppp1r16a) alternative variant dSep08, complete mRNA.
<a href="#">Ppp1r16a</a>	<a href="#">Ppp1r16a.dSep08</a>	<a href="#">681165</a>	6100	498	3	32	protein phosphatase 1, regulatory (inhibitor) subunit 16A (3.5 kD) (Ppp1r16a) alternative variant dSep08, complete mRNA.
<a href="#">Ppp1r16b</a>	<a href="#">Ppp1r16b.aSep08</a>	<a href="#">680616</a>	80828	750		190	protein phosphatase 1, regulatory (inhibitor) subunit 16B (Ppp1r16b) mRNA.
<a href="#">Ppp2cb</a>	<a href="#">Ppp2cb.bSep08</a>	<a href="#">24673</a>	5161	1574	2	115	protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform (13.1 kD) (Ppp2cb) alternative variant bSep08, mRNA.
<a href="#">Ppp2r1a</a>	<a href="#">Ppp2r1a.bSep08</a>	<a href="#">117281</a>	7874	752	1	98	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform (10.7 kD) (Ppp2r1a) alternative variant bSep08, mRNA.
<a href="#">Ppp2r1b</a>	<a href="#">Ppp2r1b.bSep08</a>	<a href="#">315648</a>	4037	345	3	115	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform (Ppp2r1b) alternative variant bSep08, mRNA.
<a href="#">Ppp2r1b</a>	<a href="#">Ppp2r1b.dSep08</a>	<a href="#">315648</a>	466	337	2	35	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform (4.3 kD) (Ppp2r1b) alternative variant dSep08, mRNA.
<a href="#">Ppp2r2a</a>	<a href="#">Ppp2r2a.bSep08</a>	<a href="#">117104</a>	8047	2112	4	240	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform (27.9 kD) (Ppp2r2a) alternative variant bSep08, mRNA.
<a href="#">Ppp2r2a</a>	<a href="#">Ppp2r2a.cSep08</a>	<a href="#">117104</a>	50591	789	6	113	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform (Ppp2r2a) alternative variant cSep08, mRNA.

<a href="#">Ppp2r2a</a>	<a href="#">Ppp2r2a.dSep08</a>	<a href="#">117104</a>	48503	339	4	112	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform (Ppp2r2a) alternative variant dSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.bSep08</a>	<a href="#">60660</a>	256027	754	6	166	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant bSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.cSep08</a>	<a href="#">60660</a>	256110	763	5	161	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant cSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.dSep08</a>	<a href="#">60660</a>	256120	883	6	153	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant dSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.eSep08</a>	<a href="#">60660</a>	217955	775	5	151	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant eSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.fSep08</a>	<a href="#">60660</a>	217968	745	5	149	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant fSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.gSep08</a>	<a href="#">60660</a>	179008	1081	2	135	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant gSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.hSep08</a>	<a href="#">60660</a>	25564	723	2	128	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant hSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.iSep08</a>	<a href="#">60660</a>	172289	629	2	111	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant iSep08, mRNA.
<a href="#">Ppp2r2b</a>	<a href="#">Ppp2r2b.jSep08</a>	<a href="#">60660</a>	211192	752	4	110	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform (Ppp2r2b) alternative variant jSep08, mRNA.
<a href="#">Ppp2r2c</a>	<a href="#">Ppp2r2c.bSep08</a>	<a href="#">117256</a>	2939	413	1	122	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), gamma isoform (Ppp2r2c) alternative variant bSep08, mRNA.
<a href="#">Ppp2r2d</a>	<a href="#">Ppp2r2d.bSep08</a>	<a href="#">246255</a>	24598	761	6	253	protein phosphatase 2, regulatory subunit B, delta isoform (Ppp2r2d) alternative variant bSep08, mRNA.
<a href="#">Ppp2r2d</a>	<a href="#">Ppp2r2d.cSep08</a>	<a href="#">246255</a>	24526	562	6	154	protein phosphatase 2, regulatory subunit B, delta isoform (Ppp2r2d) alternative variant cSep08, mRNA.
<a href="#">Ppp2r3a</a>	<a href="#">Ppp2r3a.aSep08</a>	<a href="#">363122</a>	74265	1677	10	558	protein phosphatase 2 (formerly 2A), regulatory subunit B", alpha (Ppp2r3a) alternative variant aSep08, mRNA.
<a href="#">Ppp2r3a</a>	<a href="#">Ppp2r3a.bSep08</a>	<a href="#">363122</a>	12914	739	3	167	protein phosphatase 2 (formerly 2A), regulatory subunit B", alpha (Ppp2r3a) alternative variant bSep08, mRNA.
<a href="#">Ppp2r3a</a>	<a href="#">Ppp2r3a.cSep08</a>	<a href="#">363122</a>	5021	807	1	94	protein phosphatase 2 (formerly 2A), regulatory subunit B", alpha (Ppp2r3a) alternative variant cSep08, mRNA.
<a href="#">Ppp2r3c</a>	<a href="#">Ppp2r3c.bSep08</a>	<a href="#">362739</a>	4496	377	1	120	protein phosphatase 2, regulatory subunit B", gamma (Ppp2r3c) alternative variant bSep08, mRNA.
<a href="#">Ppp2r4</a>	<a href="#">Ppp2r4.bSep08</a>	<a href="#">362102</a>	5212	580	4	95	protein phosphatase 2A, regulatory subunit B (PR 53) (Ppp2r4) alternative variant bSep08, mRNA.

<a href="#">Ppp2r4</a>	<a href="#">Ppp2r4.cSep08</a>	<a href="#">362102</a>	5724	852	3	72	protein phosphatase 2A, regulatory subunit B (PR 53) (Ppp2r4) alternative variant cSep08, mRNA.
<a href="#">Ppp2r5a</a>	<a href="#">Ppp2r5a.bSep08</a>	<a href="#">312754</a>	16010	688	8	228	protein phosphatase 2, regulatory subunit B (B56), alpha isoform and hypothetical protein LOC679797 (Ppp2r5a) alternative variant bSep08, mRNA.
<a href="#">Ppp2r5a</a>	<a href="#">Ppp2r5a.bSep08</a>	<a href="#">679797</a>	16010	688	8	228	protein phosphatase 2, regulatory subunit B (B56), alpha isoform and hypothetical protein LOC679797 (Ppp2r5a) alternative variant bSep08, mRNA.
<a href="#">Ppp2r5a</a>	<a href="#">Ppp2r5a.cSep08</a>	<a href="#">312754</a>	2348	1520	3	64	protein phosphatase 2, regulatory subunit B (B56), alpha isoform and hypothetical protein LOC679797 (7.7 kD) (Ppp2r5a) alternative variant cSep08, mRNA.
<a href="#">Ppp2r5a</a>	<a href="#">Ppp2r5a.cSep08</a>	<a href="#">679797</a>	2348	1520	3	64	protein phosphatase 2, regulatory subunit B (B56), alpha isoform and hypothetical protein LOC679797 (7.7 kD) (Ppp2r5a) alternative variant cSep08, mRNA.
<a href="#">Ppp2r5b</a>	<a href="#">Ppp2r5b.cSep08</a>	<a href="#">309179</a>	1751	1056	4	190	protein phosphatase 2, regulatory subunit B (B56), beta isoform (Ppp2r5b) alternative variant cSep08, mRNA.
<a href="#">Ppp2r5b</a>	<a href="#">Ppp2r5b.dSep08</a>	<a href="#">309179</a>	3816	1195	7	130	protein phosphatase 2, regulatory subunit B (B56), beta isoform (Ppp2r5b) alternative variant dSep08, mRNA.
<a href="#">Ppp2r5b</a>	<a href="#">Ppp2r5b.fSep08</a>	<a href="#">309179</a>	3649	1091	8	74	protein phosphatase 2, regulatory subunit B (B56), beta isoform (Ppp2r5b) alternative variant fSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.aSep08</a>	<a href="#">691318</a>	128772	1729	14	554	protein phosphatase 2, regulatory subunit B' gamma isoform (Ppp2r5c) alternative variant aSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.bSep08</a>	<a href="#">691318</a>	85186	2432	12	452	protein phosphatase 2, regulatory subunit B' gamma isoform (52.8 kD) (Ppp2r5c) alternative variant bSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.cSep08</a>	<a href="#">691318</a>	15984	2929	4	122	protein phosphatase 2, regulatory subunit B' gamma isoform (Ppp2r5c) alternative variant cSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.dSep08</a>	<a href="#">691318</a>	3547	471	3	116	protein phosphatase 2, regulatory subunit B' gamma isoform (Ppp2r5c) alternative variant dSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.eSep08</a>	<a href="#">691318</a>	7157	403	4	81	protein phosphatase 2, regulatory subunit B' gamma isoform (Ppp2r5c) alternative variant eSep08, mRNA.
<a href="#">Ppp2r5c</a>	<a href="#">Ppp2r5c.fSep08</a>	<a href="#">691318</a>	5868	1264	2	80	protein phosphatase 2, regulatory subunit B' gamma isoform (Ppp2r5c) alternative variant fSep08, mRNA.
<a href="#">Ppp2r5d</a>	<a href="#">Ppp2r5d.aSep08</a>	<a href="#">363193</a>	24517	2912	16	637	protein phosphatase 2, regulatory subunit B (B56), delta isoform (Ppp2r5d) alternative variant aSep08, mRNA.
<a href="#">Ppp2r5e</a>	<a href="#">Ppp2r5e.bSep08</a>	<a href="#">299147</a>	41967	2417	11	230	protein phosphatase 2, regulatory subunit B (B56), epsilon isoform (26.9 kD) (Ppp2r5e) alternative variant bSep08, mRNA.
<a href="#">Ppp3ca</a>	<a href="#">Ppp3ca.bSep08</a>	<a href="#">24674</a>	76579	1531	12	428	protein phosphatase 3, catalytic subunit, alpha isoform (Ppp3ca) alternative variant bSep08, mRNA.
<a href="#">Ppp3ca</a>	<a href="#">Ppp3ca.cSep08</a>	<a href="#">24674</a>	202492	441	3	68	protein phosphatase 3, catalytic subunit, alpha isoform (Ppp3ca) alternative variant cSep08, mRNA.
<a href="#">Ppp3cb</a>	<a href="#">Ppp3cb.bSep08</a>	<a href="#">24675</a>	14494	1432	6	211	protein phosphatase 3, catalytic subunit, beta isoform (Ppp3cb) alternative variant bSep08, mRNA.
<a href="#">Ppp3cb</a>	<a href="#">Ppp3cb.cSep08</a>	<a href="#">24675</a>	18819	2004	6	195	protein phosphatase 3, catalytic subunit, beta isoform (Ppp3cb) alternative variant cSep08, mRNA.

<a href="#">Ppp3cb</a>	<a href="#">Ppp3cb.dSep08</a>	<a href="#">24675</a>	2492	862	2	59	protein phosphatase 3, catalytic subunit, beta isoform (6.9 kD) (Ppp3cb) alternative variant dSep08, mRNA.
<a href="#">Ppp3cc</a>	<a href="#">Ppp3cc.aSep08</a>	<a href="#">171378</a>	49339	1583	11	332	protein phosphatase 3, catalytic subunit, gamma isoform (37.7 kD) (Ppp3cc) alternative variant aSep08, mRNA.
<a href="#">Ppp3cc</a>	<a href="#">Ppp3cc.bSep08</a>	<a href="#">171378</a>	24445	658	5	189	protein phosphatase 3, catalytic subunit, gamma isoform (Ppp3cc) alternative variant bSep08, mRNA.
<a href="#">Ppp3cc</a>	<a href="#">Ppp3cc.cSep08</a>	<a href="#">171378</a>	8424	2009	3	108	protein phosphatase 3, catalytic subunit, gamma isoform (12.0 kD) (Ppp3cc) alternative variant cSep08, mRNA.
<a href="#">Ppp4c</a>	<a href="#">Ppp4c.bSep08</a>	<a href="#">171366</a>	6769	1423	2	307	protein phosphatase 4, catalytic subunit (35.1 kD) (Ppp4c) alternative variant bSep08, mRNA.
<a href="#">Ppp4r1</a>	<a href="#">Ppp4r1.bSep08</a>	<a href="#">140943</a>	15838	901	4	300	protein phosphatase 4, regulatory subunit 1 (Ppp4r1) alternative variant bSep08, mRNA.
<a href="#">Ppp4r1</a>	<a href="#">Ppp4r1.cSep08</a>	<a href="#">140943</a>	13865	1100	8	252	protein phosphatase 4, regulatory subunit 1 (28.6 kD) (Ppp4r1) alternative variant cSep08, mRNA.
<a href="#">Ppp4r1</a>	<a href="#">Ppp4r1.dSep08</a>	<a href="#">140943</a>	12885	1352	6	198	protein phosphatase 4, regulatory subunit 1 (Ppp4r1) alternative variant dSep08, mRNA.
<a href="#">Ppp4r1</a>	<a href="#">Ppp4r1.eSep08</a>	<a href="#">140943</a>	2691	849	2	123	protein phosphatase 4, regulatory subunit 1 (Ppp4r1) alternative variant eSep08, mRNA.
<a href="#">Ppp4r1l</a>	<a href="#">Ppp4r1l.bSep08</a>	<a href="#">315549</a>	12680	482	4	160	protein phosphatase 4, regulatory subunit 1-like (Ppp4r1l) alternative variant bSep08, mRNA.
<a href="#">Ppp4r2</a>	<a href="#">Ppp4r2.bSep08</a>	<a href="#">297486</a>	37117	1517	2	375	protein phosphatase 4, regulatory subunit 2 (41.6 kD) (Ppp4r2) alternative variant bSep08, mRNA.
<a href="#">Ppp5c</a>	<a href="#">Ppp5c.bSep08</a>	<a href="#">65179</a>	4992	1113	4	314	protein phosphatase 5, catalytic subunit (Ppp5c) alternative variant bSep08, mRNA.
<a href="#">Ppp5c</a>	<a href="#">Ppp5c.cSep08</a>	<a href="#">65179</a>	2432	1317	3	220	protein phosphatase 5, catalytic subunit (Ppp5c) alternative variant cSep08, mRNA.
<a href="#">Pprc1</a>	<a href="#">Pprc1.bSep08</a>	<a href="#">294007</a>	10263	3800	9	1196	peroxisome proliferative activated receptor, gamma, coactivator-related 1 (127.3 kD) (Pprc1) alternative variant bSep08, mRNA.
<a href="#">Pprc1</a>	<a href="#">Pprc1.dSep08</a>	<a href="#">294007</a>	5810	414	3	138	peroxisome proliferative activated receptor, gamma, coactivator-related 1 (Pprc1) alternative variant dSep08, mRNA.
<a href="#">Pprc1</a>	<a href="#">Pprc1.eSep08</a>	<a href="#">294007</a>	3712	1264	5	111	peroxisome proliferative activated receptor, gamma, coactivator-related 1 (13.0 kD) (Pprc1) alternative variant eSep08, mRNA.
<a href="#">Pprc1</a>	<a href="#">Pprc1.fSep08</a>	<a href="#">294007</a>	672	457	2	104	peroxisome proliferative activated receptor, gamma, coactivator-related 1 (Pprc1) alternative variant fSep08, mRNA.
<a href="#">Ppt2</a>	<a href="#">Ppt2.bSep08</a>	<a href="#">54398</a>	3693	984	4	241	palmitoyl-protein thioesterase 2 (Ppt2) alternative variant bSep08, mRNA.
<a href="#">Ppt2</a>	<a href="#">Ppt2.cSep08</a>	<a href="#">54398</a>	4370	1082	6	232	palmitoyl-protein thioesterase 2 (Ppt2) alternative variant cSep08, mRNA.
<a href="#">Ppt2</a>	<a href="#">Ppt2.dSep08</a>	<a href="#">54398</a>	4444	1211	5	208	palmitoyl-protein thioesterase 2 (Ppt2) alternative variant dSep08, mRNA.
<a href="#">Ppt2</a>	<a href="#">Ppt2.eSep08</a>	<a href="#">54398</a>	1745	1119	3	180	palmitoyl-protein thioesterase 2 (Ppt2) alternative variant eSep08, mRNA.

<a href="#">Ppt2</a>	<a href="#">Ppt2.fSep08</a>	<a href="#">54398</a>	867	451	1	70	palmitoyl-protein thioesterase 2 (Ppt2) alternative variant fSep08, mRNA.
<a href="#">Ppwd1</a>	<a href="#">Ppwd1.bSep08</a>	<a href="#">294711</a>	8452	1077	4	254	peptidylprolyl isomerase domain and WD repeat containing 1 (Ppwd1) alternative variant bSep08, mRNA.
<a href="#">Ppwd1</a>	<a href="#">Ppwd1.cSep08</a>	<a href="#">294711</a>	2960	858	1	79	peptidylprolyl isomerase domain and WD repeat containing 1 (Ppwd1) alternative variant cSep08, mRNA.
<a href="#">Pqbp1</a>	<a href="#">Pqbp1.bSep08</a>	<a href="#">302557</a>	4077	1034	6	263	polyglutamine binding protein 1 (30.5 kD) (Pqbp1) alternative variant bSep08, mRNA.
<a href="#">Pqbp1</a>	<a href="#">Pqbp1.cSep08</a>	<a href="#">302557</a>	4418	994	6	263	polyglutamine binding protein 1 (30.5 kD) (Pqbp1) alternative variant cSep08, mRNA.
<a href="#">Pqbp1</a>	<a href="#">Pqbp1.dSep08</a>	<a href="#">302557</a>	3578	463	4	128	polyglutamine binding protein 1 (Pqbp1) alternative variant dSep08, mRNA.
<a href="#">Pqbp1</a>	<a href="#">Pqbp1.eSep08</a>	<a href="#">302557</a>	2737	299	2	74	polyglutamine binding protein 1 (Pqbp1) alternative variant eSep08, mRNA.
<a href="#">Pqlc1</a>	<a href="#">Pqlc1.aSep08</a>	<a href="#">361352</a>	36464	1459	4	271	PQ loop repeat containing 1 (30.6 kD) (Pqlc1) alternative variant aSep08, mRNA.
<a href="#">Pqlc1</a>	<a href="#">Pqlc1.bSep08</a>	<a href="#">361352</a>	36209	1602	2	253	PQ loop repeat containing 1 (28.4 kD) (Pqlc1) alternative variant bSep08, mRNA.
<a href="#">Pqlc1</a>	<a href="#">Pqlc1.dSep08</a>	<a href="#">361352</a>	27013	772	3	184	PQ loop repeat containing 1 (Pqlc1) alternative variant dSep08, mRNA.
<a href="#">Pqlc2</a>	<a href="#">Pqlc2.aSep08</a>	<a href="#">362642</a>	9905	1388	6	293	PQ loop repeat containing 2 (32.2 kD) (Pqlc2) alternative variant aSep08, mRNA.
<a href="#">Pqlc2</a>	<a href="#">Pqlc2.bSep08</a>	<a href="#">362642</a>	6163	1057	2	146	PQ loop repeat containing 2 (15.9 kD) (Pqlc2) alternative variant bSep08, mRNA.
<a href="#">Pqlc3</a>	<a href="#">Pqlc3.bSep08</a>	<a href="#">298906</a>	7105	626	1	153	PQ loop repeat containing 3 (17.2 kD) (Pqlc3) alternative variant bSep08, mRNA.
<a href="#">Praf2</a>	<a href="#">Praf2.cSep08</a>	<a href="#">367743</a>	1351	522	2	29	PRA1 domain family 2 (3.2 kD) (Praf2) alternative variant cSep08, mRNA.
<a href="#">Pram1</a>	<a href="#">Pram1.aSep08</a>	<a href="#">362848</a>	4272	1781	4	556	PML-RAR alpha-regulated adaptor molecule 1 (Pram1) alternative variant aSep08, mRNA.
<a href="#">Pram1</a>	<a href="#">Pram1.bSep08</a>	<a href="#">362848</a>	601	354	3	80	PML-RAR alpha-regulated adaptor molecule 1 (Pram1) alternative variant bSep08, mRNA.
<a href="#">Pram1</a>	<a href="#">Pram1.cSep08</a>	<a href="#">362848</a>	499	415	1	63	PML-RAR alpha-regulated adaptor molecule 1 (7.3 kD) (Pram1) alternative variant cSep08, mRNA.
<a href="#">Pramef12</a>	<a href="#">Pramef12.aSep08</a>	<a href="#">691157</a>	7779	678	3	189	PRAME family member 12 (Pramef12) alternative variant aSep08, mRNA.
<a href="#">Prc1</a>	<a href="#">Prc1.bSep08</a>	<a href="#">308761</a>	6108	1707	7	239	protein regulator of cytokinesis 1 (Prc1) alternative variant bSep08, mRNA.
<a href="#">Prc1</a>	<a href="#">Prc1.cSep08</a>	<a href="#">308761</a>	4177	1063	4	116	protein regulator of cytokinesis 1 (Prc1) alternative variant cSep08, mRNA.
<a href="#">Prpcp</a>	<a href="#">Prpcp.bSep08</a>	<a href="#">293118</a>	3543	679	2	187	prolylcarboxypeptidase (angiotensinase C) (Prpcp) alternative variant bSep08, mRNA.
<a href="#">Prpcp</a>	<a href="#">Prpcp.dSep08</a>	<a href="#">293118</a>	2656	518	1	125	prolylcarboxypeptidase (angiotensinase C) (14.2 kD) (Prpcp) alternative variant dSep08, mRNA.
<a href="#">Prdm2</a>	<a href="#">Prdm2.bSep08</a>	<a href="#">313678</a>	75830	512	5	124	putative protein of metazoan origin (Prdm2) alternative variant bSep08, mRNA.



<a href="#">Prdm4</a>	<a href="#">Prdm4.bSep08</a>	<a href="#">170820</a>	1288	737	1	141	zinc finger, C2H2-type (Prdm4) alternative variant bSep08, mRNA.
<a href="#">Prdm5</a>	<a href="#">Prdm5.aSep08</a>	<a href="#">689788</a>	102863	1130		376	domain 5 (Prdm5) mRNA.
<a href="#">Prdm6</a>	<a href="#">Prdm6.aSep08</a>	<a href="#">307305</a>	19715	426		77	PR domain 6 (Prdm6) mRNA.
<a href="#">Prdm8</a>	<a href="#">Prdm8.aSep08</a>	<a href="#">305198</a>	2172	840	1	279	pr domain 8 (Prdm8) alternative variant aSep08, mRNA.
<a href="#">Prdm10</a>	<a href="#">Prdm10.aSep08</a>	<a href="#">500964</a>	28329	2190		700	zinc finger, C2H2-type (Prdm10) mRNA.
<a href="#">Prdm12</a>	<a href="#">Prdm12.aSep08</a>	<a href="#">688699</a>	2839	486		161	zinc finger, C2H2-type (Prdm12) mRNA.
<a href="#">Prdx2</a>	<a href="#">Prdx2.bSep08</a>	<a href="#">29338</a>	5034	813	6	192	peroxiredoxin 2 (Prdx2) alternative variant bSep08, mRNA.
<a href="#">Prdx2</a>	<a href="#">Prdx2.cSep08</a>	<a href="#">29338</a>	5760	1286	5	181	peroxiredoxin 2 (19.8 kD) (Prdx2) alternative variant cSep08, mRNA.
<a href="#">Prdx2</a>	<a href="#">Prdx2.eSep08</a>	<a href="#">29338</a>	1756	419	3	88	peroxiredoxin 2 (Prdx2) alternative variant eSep08, mRNA.
<a href="#">Prdx2</a>	<a href="#">Prdx2.fSep08</a>	<a href="#">29338</a>	505	416	2	47	peroxiredoxin 2 (Prdx2) alternative variant fSep08, mRNA.
<a href="#">Prdx3</a>	<a href="#">Prdx3.bSep08</a>	<a href="#">64371</a>	1831	1369	2	55	peroxiredoxin 3 (6.2 kD) (Prdx3) alternative variant bSep08, mRNA.
<a href="#">Prdx5</a>	<a href="#">Prdx5.bSep08</a>	<a href="#">113898</a>	2947	789	4	217	peroxiredoxin 5 (Prdx5) alternative variant bSep08, complete mRNA.
<a href="#">Prdx5</a>	<a href="#">Prdx5.cSep08</a>	<a href="#">113898</a>	3344	687	6	158	peroxiredoxin 5 (16.7 kD) (Prdx5) alternative variant cSep08, mRNA.
<a href="#">Prdx5</a>	<a href="#">Prdx5.dSep08</a>	<a href="#">113898</a>	3454	430	6	142	peroxiredoxin 5 (Prdx5) alternative variant dSep08, mRNA.
<a href="#">Prdx5</a>	<a href="#">Prdx5.gSep08</a>	<a href="#">113898</a>	1113	777	2	78	peroxiredoxin 5 (9.6 kD) (Prdx5) alternative variant gSep08, mRNA.
<a href="#">Prdx5</a>	<a href="#">Prdx5.hSep08</a>	<a href="#">113898</a>	439	347	2	42	peroxiredoxin 5 (Prdx5) alternative variant hSep08, mRNA.
<a href="#">Pre-SET.0</a>	<a href="#">Pre-SET.0.aSep08</a>		20762	4460	16	1113	SET domain bifurcated 1 (Pre-SET.0) alternative variant aSep08, mRNA.
<a href="#">Pre-SET.0</a>	<a href="#">Pre-SET.0.bSep08</a>		8758	1011	2	197	CRA a (Pre-SET.0) alternative variant bSep08, mRNA.
<a href="#">Pre-SET.0</a>	<a href="#">Pre-SET.0.cSep08</a>		1253	1136	2	70	CRA b like (7.6 kD) (Pre-SET.0) alternative variant cSep08, mRNA.
<a href="#">Pre-SET.0</a>	<a href="#">Pre-SET.0.dSep08</a>		499	384	2	55	SET domain bifurcated 1 (Pre-SET.0) alternative variant dSep08, mRNA.
<a href="#">Prelid1</a>	<a href="#">Prelid1.bSep08</a>	<a href="#">290995</a>	2359	675	1	174	preli (Prelid1) alternative variant bSep08, mRNA.
<a href="#">Prelid2</a>	<a href="#">Prelid2.aSep08</a>	<a href="#">681037</a>	74752	598	2	108	putative protein (Prelid2) alternative variant aSep08, mRNA.
<a href="#">Prelid2</a>	<a href="#">Prelid2.bSep08</a>	<a href="#">681037</a>	94115	397	1	44	putative protein of mammalian origin (Prelid2) alternative variant bSep08, mRNA.
<a href="#">Prelp</a>	<a href="#">Prelp.bSep08</a>	<a href="#">84400</a>	12951	3581	2	377	proline arginine-rich end leucine-rich repeat protein (43.2 kD) (Prelp) alternative variant bSep08, mRNA.
<a href="#">Prenyltrans.0</a>	<a href="#">Prenyltrans.0.aSep08</a>		9219	2761	4	170	lanosterol synthase (Prenyltrans.0) alternative variant aSep08, mRNA.
<a href="#">Prenyltrans.0</a>	<a href="#">Prenyltrans.0.bSep08</a>		746	211	1	45	putative protein (Prenyltrans.0) alternative variant bSep08, mRNA.
<a href="#">Prep</a>	<a href="#">Prep.bSep08</a>	<a href="#">83471</a>	32885	2847	6	259	prolyl endopeptidase (28.7 kD) (Prep) alternative variant bSep08, mRNA.
<a href="#">Prep</a>	<a href="#">Prep.cSep08</a>	<a href="#">83471</a>	1888	458	2	99	prolyl endopeptidase (Prep) alternative variant cSep08, mRNA.

<a href="#">Prepl</a>	<a href="#">Prepl.aSep08</a>	<a href="#">298771</a>	29761	2387	8	640	prolyl endopeptidase-like (Prepl) alternative variant aSep08, mRNA.
<a href="#">Prepl</a>	<a href="#">Prepl.cSep08</a>	<a href="#">298771</a>	14124	679	5	140	prolyl endopeptidase-like (Prepl) alternative variant cSep08, mRNA.
<a href="#">Prepl</a>	<a href="#">Prepl.dSep08</a>	<a href="#">298771</a>	10181	734	2	121	prolyl endopeptidase-like (Prepl) alternative variant dSep08, mRNA.
<a href="#">Prg-2</a>	<a href="#">Prg-2.bSep08</a>	<a href="#">314614</a>	9633	1247	4	136	plasticity-related gene 2 (15.1 kD) (Prg-2) alternative variant bSep08, mRNA.
<a href="#">Prg-2</a>	<a href="#">Prg-2.cSep08</a>	<a href="#">314614</a>	11006	247	1	55	plasticity-related gene 2 (Prg-2) alternative variant cSep08, mRNA.
<a href="#">Prg-4</a>	<a href="#">Prg-4.aSep08</a>	<a href="#">300443</a>	11881	2364	10	452	plasticity-related gene 4 (48.4 kD) (Prg-4) alternative variant aSep08, mRNA.
<a href="#">Prg-4</a>	<a href="#">Prg-4.bSep08</a>	<a href="#">300443</a>	5447	1881	5	206	plasticity-related gene 4 (Prg-4) alternative variant bSep08, mRNA.
<a href="#">Prg-4</a>	<a href="#">Prg-4.cSep08</a>	<a href="#">300443</a>	797	407	2	62	plasticity-related gene 4 (Prg-4) alternative variant cSep08, mRNA.
<a href="#">Prg-4</a>	<a href="#">Prg-4.dSep08</a>	<a href="#">300443</a>	2769	343	3	53	plasticity-related gene 4 (Prg-4) alternative variant dSep08, mRNA.
<a href="#">Prg4</a>	<a href="#">Prg4.bSep08</a>	<a href="#">289104</a>	4776	536	4	172	proteoglycan 4, (megakaryocyte stimulating factor, articular superficial zone protein, camptodactyly, arthropathy, coxa vara, pericarditis syndrome) (Prg4) alternative variant bSep08, mRNA.
<a href="#">Pric285</a>	<a href="#">Pric285.aSep08</a>	<a href="#">296474</a>	3010	2064	5	555	peroxisomal proliferator-activated receptor A interacting complex 285 (Pric285) alternative variant aSep08, mRNA.
<a href="#">Pric285</a>	<a href="#">Pric285.bSep08</a>	<a href="#">296474</a>	1818	1529	4	266	peroxisomal proliferator-activated receptor A interacting complex 285 (29.9 kD) (Pric285) alternative variant bSep08, mRNA.
<a href="#">Prickle3</a>	<a href="#">Prickle3.bSep08</a>	<a href="#">317380</a>	3469	1158	5	315	prickle homolog 3 (Drosophila) (Prickle3) alternative variant bSep08, mRNA.
<a href="#">Prickle4</a>	<a href="#">Prickle4.aSep08</a>	<a href="#">681123</a>	1534	677	3	135	prickle homolog 4 (Drosophila) (Prickle4) alternative variant aSep08, mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.aSep08</a>	<a href="#">246327</a>	29441	1560	11	417	DNA primase, p49 subunit (Prim1) alternative variant aSep08, mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.bSep08</a>	<a href="#">246327</a>	29903	1451	12	333	DNA primase, p49 subunit (38.9 kD) (Prim1) alternative variant bSep08, complete mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.cSep08</a>	<a href="#">246327</a>	23720	877	5	277	DNA primase, p49 subunit (Prim1) alternative variant cSep08, mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.dSep08</a>	<a href="#">246327</a>	9692	727	8	242	DNA primase, p49 subunit (Prim1) alternative variant dSep08, mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.eSep08</a>	<a href="#">246327</a>	9281	726	8	219	DNA primase, p49 subunit (Prim1) alternative variant eSep08, mRNA.
<a href="#">Prim1</a>	<a href="#">Prim1.fSep08</a>	<a href="#">246327</a>	7863	1110	6	178	DNA primase, p49 subunit (Prim1) alternative variant fSep08, mRNA.
<a href="#">Prim2</a>	<a href="#">Prim2.bSep08</a>	<a href="#">301323</a>	40275	826	7	275	DNA primase, p58 subunit (Prim2) alternative variant bSep08, mRNA.
<a href="#">Prim2</a>	<a href="#">Prim2.cSep08</a>	<a href="#">301323</a>	50188	732	7	244	DNA primase, p58 subunit (Prim2) alternative variant cSep08, mRNA.

<a href="#">Prim2</a>	<a href="#">Prim2.dSep08</a>	<a href="#">301323</a>	167897	1269	8	220	DNA primase, p58 subunit (25.3 kD) (Prim2) alternative variant dSep08, mRNA.
<a href="#">Prima1</a>	<a href="#">Prima1.aSep08</a>	<a href="#">690195</a>	43598	441	4	146	proline rich membrane anchor 1 (Prima1) alternative variant aSep08, mRNA.
<a href="#">Prkaa1</a>	<a href="#">Prkaa1.aSep08</a>	<a href="#">65248</a>	4895	1783	4	558	protein kinase, AMP-activated, alpha 1 catalytic subunit (Prkaa1) alternative variant aSep08, mRNA.
<a href="#">Prkab1</a>	<a href="#">Prkab1.bSep08</a>	<a href="#">83803</a>	9101	806	7	241	5-AMP-activated protein kinase, beta subunit, complex-interacting region (Prkab1) alternative variant bSep08, mRNA.
<a href="#">Prkab1</a>	<a href="#">Prkab1.cSep08</a>	<a href="#">83803</a>	5153	2021	3	116	5-AMP-activated protein kinase, beta subunit, complex-interacting region (13.1 kD) (Prkab1) alternative variant cSep08, mRNA.
<a href="#">Prkab1</a>	<a href="#">Prkab1.dSep08</a>	<a href="#">83803</a>	7225	928	5	87	putative protein of ancient origin (Prkab1) alternative variant dSep08, mRNA.
<a href="#">Prkab1</a>	<a href="#">Prkab1.fSep08</a>	<a href="#">83803</a>	1122	917	2	61	AMP-activated protein kinase (Prkab1) alternative variant fSep08, mRNA.
<a href="#">Prkab1</a>	<a href="#">Prkab1.gSep08</a>	<a href="#">83803</a>	1296	348	2	52	putative protein of eukaryotic origin (Prkab1) alternative variant gSep08, mRNA.
<a href="#">Prkacb</a>	<a href="#">Prkacb.bSep08</a>	<a href="#">293508</a>	24329	1345	6	216	protein kinase, cAMP dependent, catalytic, beta (25.3 kD) (Prkacb) alternative variant bSep08, mRNA.
<a href="#">Prkacb</a>	<a href="#">Prkacb.cSep08</a>	<a href="#">293508</a>	25402	634	7	187	protein kinase, cAMP dependent, catalytic, beta (Prkacb) alternative variant cSep08, mRNA.
<a href="#">Prkag1</a>	<a href="#">Prkag1.bSep08</a>	<a href="#">25520</a>	15915	701	7	189	protein kinase, AMP-activated, gamma 1 non-catalytic subunit (21.6 kD) (Prkag1) alternative variant bSep08, mRNA.
<a href="#">Prkag1</a>	<a href="#">Prkag1.cSep08</a>	<a href="#">25520</a>	12250	1786	8	177	protein kinase, AMP-activated, gamma 1 non-catalytic subunit (Prkag1) alternative variant cSep08, mRNA.
<a href="#">Prkag1</a>	<a href="#">Prkag1.dSep08</a>	<a href="#">25520</a>	14846	402	2	68	protein kinase, AMP-activated, gamma 1 non-catalytic subunit (7.6 kD) (Prkag1) alternative variant dSep08, mRNA.
<a href="#">Prkag2</a>	<a href="#">Prkag2.aSep08</a>	<a href="#">373545</a>	233803	1657	11	407	protein kinase AMP-activated (Prkag2) alternative variant aSep08, mRNA.
<a href="#">Prkag2</a>	<a href="#">Prkag2.dSep08</a>	<a href="#">373545</a>	80879	1120	9	154	kinase -activated protein (Prkag2) alternative variant dSep08, mRNA.
<a href="#">Prkag2</a>	<a href="#">Prkag2.eSep08</a>	<a href="#">373545</a>	3571	498	2	102	protein kinase AMP-activated (11.4 kD) (Prkag2) alternative variant eSep08, mRNA.
<a href="#">Prkar1b</a>	<a href="#">Prkar1b.bSep08</a>	<a href="#">25521</a>	76062	729	6	243	protein kinase, cAMP dependent regulatory, type I, beta (Prkar1b) alternative variant bSep08, mRNA.
<a href="#">Prkar1b</a>	<a href="#">Prkar1b.cSep08</a>	<a href="#">25521</a>	40099	532	3	138	protein kinase, cAMP dependent regulatory, type I, beta (Prkar1b) alternative variant cSep08, mRNA.
<a href="#">Prkar1b</a>	<a href="#">Prkar1b.dSep08</a>	<a href="#">25521</a>	26919	1677	3	111	protein kinase, cAMP dependent regulatory, type I, beta (12.5 kD) (Prkar1b) alternative variant dSep08, mRNA.
<a href="#">Prkar2a</a>	<a href="#">Prkar2a.bSep08</a>	<a href="#">29699</a>	12888	4090	2	154	protein kinase, cAMP dependent regulatory, type II alpha (17.3 kD) (Prkar2a) alternative variant bSep08, mRNA.
<a href="#">Prkar2b</a>	<a href="#">Prkar2b.bSep08</a>	<a href="#">24679</a>	46358	675	8	203	protein kinase, cAMP dependent regulatory, type II beta (Prkar2b) alternative variant bSep08, mRNA.

<a href="#">Prkca</a>	<a href="#">Prkca.aSep08</a>	<a href="#">24680</a>	114646	3677	2	576	protein kinase C, alpha (Prkca) alternative variant aSep08, mRNA.
<a href="#">Prkcb1</a>	<a href="#">Prkcb1.bSep08</a>	<a href="#">25023</a>	109621	1887	12	459	protein kinase C, beta 1 (Prkcb1) alternative variant bSep08, mRNA.
<a href="#">Prkcb1</a>	<a href="#">Prkcb1.cSep08</a>	<a href="#">25023</a>	30175	1119	2	90	protein kinase C, beta 1 (Prkcb1) alternative variant cSep08, mRNA.
<a href="#">Prkcb1</a>	<a href="#">Prkcb1.eSep08</a>	<a href="#">25023</a>	6622	688	2	61	protein kinase C, beta 1 (Prkcb1) alternative variant eSep08, mRNA.
<a href="#">Prkcb1</a>	<a href="#">Prkcb1.fSep08</a>	<a href="#">25023</a>	176794	1406	4	51	protein kinase C, beta 1 (6.0 kD) (Prkcb1) alternative variant fSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.bSep08</a>	<a href="#">296374</a>	12741	1773	6	334	protein kinase C binding protein 1 (Prkcbp1) alternative variant bSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.cSep08</a>	<a href="#">296374</a>	40934	669	7	222	protein kinase C binding protein 1 (Prkcbp1) alternative variant cSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.dSep08</a>	<a href="#">296374</a>	44536	1010	5	220	protein kinase C binding protein 1 (Prkcbp1) alternative variant dSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.eSep08</a>	<a href="#">296374</a>	10644	2036	5	219	protein kinase C binding protein 1 (Prkcbp1) alternative variant eSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.gSep08</a>	<a href="#">296374</a>	10201	870	6	167	protein kinase C binding protein 1 (Prkcbp1) alternative variant gSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.hSep08</a>	<a href="#">296374</a>	15104	757	6	161	protein kinase C binding protein 1 (Prkcbp1) alternative variant hSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.iSep08</a>	<a href="#">296374</a>	40807	398	4	132	protein kinase C binding protein 1 (Prkcbp1) alternative variant iSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.jSep08</a>	<a href="#">296374</a>	40770	373	5	105	protein kinase C binding protein 1 (Prkcbp1) alternative variant jSep08, mRNA.
<a href="#">Prkcbp1</a>	<a href="#">Prkcbp1.kSep08</a>	<a href="#">296374</a>	7965	445	2	51	protein kinase C binding protein 1 (Prkcbp1) alternative variant kSep08, mRNA.
<a href="#">Prkcc</a>	<a href="#">Prkcc.bSep08</a>	<a href="#">24681</a>	1713	853	3	130	kinase C gamma (14.6 kD) (Prkcc) alternative variant bSep08, mRNA.
<a href="#">Prkcc</a>	<a href="#">Prkcc.cSep08</a>	<a href="#">24681</a>	2062	668	4	95	kinase C gamma (Prkcc) alternative variant cSep08, mRNA.
<a href="#">Prkcc</a>	<a href="#">Prkcc.dSep08</a>	<a href="#">24681</a>	1589	495	2	74	protein kinase C gamma (Prkcc) alternative variant dSep08, mRNA.
<a href="#">Prkcd</a>	<a href="#">Prkcd.bSep08</a>	<a href="#">170538</a>	4178	743	7	206	protein kinase C delta CRA a (Prkcd) alternative variant bSep08, mRNA.
<a href="#">Prkcd</a>	<a href="#">Prkcd.cSep08</a>	<a href="#">170538</a>	2115	726	3	116	protein kinase C delta CRA b (Prkcd) alternative variant cSep08, mRNA.
<a href="#">Prkcd</a>	<a href="#">Prkcd.dSep08</a>	<a href="#">170538</a>	2454	526	2	61	putative protein (6.5 kD) (Prkcd) alternative variant dSep08, mRNA.
<a href="#">Prkce</a>	<a href="#">Prkce.bSep08</a>	<a href="#">29340</a>	68109	484	4	161	protein kinase C, epsilon (Prkce) alternative variant bSep08, mRNA.
<a href="#">Prkce</a>	<a href="#">Prkce.cSep08</a>	<a href="#">29340</a>	9114	637	3	129	protein kinase C, epsilon (Prkce) alternative variant cSep08, mRNA.
<a href="#">Prkce</a>	<a href="#">Prkce.eSep08</a>	<a href="#">29340</a>	9017	300	3	74	protein kinase C, epsilon (8.8 kD) (Prkce) alternative variant eSep08, mRNA.
<a href="#">Prkcq</a>	<a href="#">Prkcq.aSep08</a>	<a href="#">85420</a>	58808	2727		464	protein kinase C, theta (53.9 kD) (Prkcq) mRNA.

<a href="#">Prkcsh</a>	<a href="#">Prkcsh.bSep08</a>	<a href="#">300445</a>	10178	1601	17	497	protein kinase C substrate 80K-H (Prkcsh) alternative variant bSep08, mRNA.
<a href="#">Prkcsh</a>	<a href="#">Prkcsh.cSep08</a>	<a href="#">300445</a>	733	561	3	130	protein kinase C substrate 80K-H (14.2 kD) (Prkcsh) alternative variant cSep08, mRNA.
<a href="#">Prkcsh</a>	<a href="#">Prkcsh.dSep08</a>	<a href="#">300445</a>	4093	664	3	119	protein kinase C substrate 80K-H (Prkcsh) alternative variant dSep08, mRNA.
<a href="#">Prkcz</a>	<a href="#">Prkcz.bSep08</a>	<a href="#">25522</a>	109433	2188	13	546	protein kinase C, zeta (62.5 kD) (Prkcz) alternative variant bSep08, mRNA.
<a href="#">Prkcz</a>	<a href="#">Prkcz.cSep08</a>	<a href="#">25522</a>	63277	1062	1	96	protein kinase C, zeta (Prkcz) alternative variant cSep08, mRNA.
<a href="#">Prkd1</a>	<a href="#">Prkd1.aSep08</a>	<a href="#">85421</a>	229432	456		152	protein kinase D1 (Prkd1) mRNA.
<a href="#">Prkd2</a>	<a href="#">Prkd2.bSep08</a>	<a href="#">292658</a>	8062	1556	7	231	protein kinase D2 (24.6 kD) (Prkd2) alternative variant bSep08, mRNA.
<a href="#">Prkd2</a>	<a href="#">Prkd2.cSep08</a>	<a href="#">292658</a>	1814	266	2	68	protein kinase D2 (Prkd2) alternative variant cSep08, mRNA.
<a href="#">Prkd3</a>	<a href="#">Prkd3.bSep08</a>	<a href="#">313834</a>	13211	1745	2	297	protein kinase D3 (33.6 kD) (Prkd3) alternative variant bSep08, mRNA.
<a href="#">Prkdc</a>	<a href="#">Prkdc.bSep08</a>	<a href="#">360748</a>	24474	3090	10	404	protein kinase catalytic (46.5 kD) (Prkdc) alternative variant bSep08, mRNA.
<a href="#">Prkdc</a>	<a href="#">Prkdc.cSep08</a>	<a href="#">360748</a>	8571	1161	9	289	protein kinase DNA-activated catalytic polypeptide CRA d (Prkdc) alternative variant cSep08, mRNA.
<a href="#">Prkdc</a>	<a href="#">Prkdc.dSep08</a>	<a href="#">360748</a>	8541	423	3	140	protein kinase catalytic (Prkdc) alternative variant dSep08, mRNA.
<a href="#">Prkdc</a>	<a href="#">Prkdc.eSep08</a>	<a href="#">360748</a>	2739	922	2	80	putative protein of eukaryotic origin (Prkdc) alternative variant eSep08, mRNA.
<a href="#">Prkg1</a>	<a href="#">Prkg1.aSep08</a>	<a href="#">54286</a>	5888	785		104	protein kinase, cGMP-dependent, type 1 (Prkg1) mRNA.
<a href="#">Prkg2</a>	<a href="#">Prkg2.bSep08</a>	<a href="#">25523</a>	12169	417	1	138	protein kinase, cGMP-dependent, type II (Prkg2) alternative variant bSep08, mRNA.
<a href="#">Prkra</a>	<a href="#">Prkra.bSep08</a>	<a href="#">311130</a>	5242	720	4	90	protein kinase, interferon inducible double stranded RNA dependent activator (Prkra) alternative variant bSep08, mRNA.
<a href="#">Prkrir</a>	<a href="#">Prkrir.aSep08</a>	<a href="#">308845</a>	13049	618		185	protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor) (Prkrir) mRNA.
<a href="#">Prkx</a>	<a href="#">Prkx.bSep08</a>	<a href="#">501563</a>	994	784	2	153	protein kinase, X-linked (Prkx) alternative variant bSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.bSep08</a>	<a href="#">24683</a>	10408	1345	5	225	prolactin (25.7 kD) (Prl) alternative variant bSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.cSep08</a>	<a href="#">24683</a>	9945	876	5	223	prolactin (25.4 kD) (Prl) alternative variant cSep08, complete mRNA.
<a href="#">Prl</a>	<a href="#">Prl.dSep08</a>	<a href="#">24683</a>	9776	704	5	216	prolactin (Prl) alternative variant dSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.eSep08</a>	<a href="#">24683</a>	9709	654	5	200	prolactin (Prl) alternative variant eSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.fSep08</a>	<a href="#">24683</a>	9935	767	4	190	prolactin (21.7 kD) (Prl) alternative variant fSep08, complete mRNA.
<a href="#">Prl</a>	<a href="#">Prl.gSep08</a>	<a href="#">24683</a>	9881	710	4	189	prolactin (21.6 kD) (Prl) alternative variant gSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.hSep08</a>	<a href="#">24683</a>	16150	739	5	174	prolactin (19.7 kD) (Prl) alternative variant hSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.iSep08</a>	<a href="#">24683</a>	9854	715	5	133	prolactin (15.2 kD) (Prl) alternative variant iSep08, mRNA.

<a href="#">Prl</a>	<a href="#">Prl.jSep08</a>	<a href="#">24683</a>	3700	361	3	119	prolactin (Prl) alternative variant jSep08, mRNA.
<a href="#">Prl</a>	<a href="#">Prl.kSep08</a>	<a href="#">24683</a>	4356	1020	3	114	prolactin (12.8 kD) (Prl) alternative variant kSep08, complete mRNA.
<a href="#">Prl3b1</a>	<a href="#">Prl3b1.cSep08</a>	<a href="#">24283</a>	25739	217	3	38	prolactin family 3, subfamily b, member 1 (Prl3b1) alternative variant cSep08, mRNA.
<a href="#">Prl3d4</a>	<a href="#">Prl3d4.aSep08</a>	<a href="#">24282</a>	6753	734	4	224	prolactin family 3, subfamily d, member 4 (25.9 kD) (Prl3d4) alternative variant aSep08, mRNA.
<a href="#">Prl3d4</a>	<a href="#">Prl3d4.cSep08</a>	<a href="#">24282</a>	3965	440	2	117	prolactin family 3, subfamily d, member 4 (13.7 kD) (Prl3d4) alternative variant cSep08, mRNA.
<a href="#">Prl3d4</a>	<a href="#">Prl3d4.dSep08</a>	<a href="#">24282</a>	1944	767	1	78	prolactin family 3, subfamily d, member 4 (8.9 kD) (Prl3d4) alternative variant dSep08, mRNA.
<a href="#">Prl4a1</a>	<a href="#">Prl4a1.cSep08</a>	<a href="#">24656</a>	2147	487	2		
<a href="#">Prl6a1</a>	<a href="#">Prl6a1.bSep08</a>	<a href="#">24657</a>	7367	888	2	216	prolactin family 6, subfamily a, member 1 (25.1 kD) (Prl6a1) alternative variant bSep08, mRNA.
<a href="#">Prl7d1</a>	<a href="#">Prl7d1.bSep08</a>	<a href="#">84377</a>	6464	1056	2	217	prolactin family 7, subfamily d, member 1 (24.3 kD) (Prl7d1) alternative variant bSep08, mRNA.
<a href="#">Prl8a2</a>	<a href="#">Prl8a2.aSep08</a>	<a href="#">24315</a>	10175	928	1	240	prolactin family 8, subfamily a, member 2 (27.9 kD) (Prl8a2) alternative variant aSep08, mRNA.
<a href="#">Prl8a4</a>	<a href="#">Prl8a4.bSep08</a>	<a href="#">59088</a>	6339	1125	1	226	prolactin family 8, subfamily a, member 4 (25.8 kD) (Prl8a4) alternative variant bSep08, complete mRNA.
<a href="#">Prl8a7</a>	<a href="#">Prl8a7.aSep08</a>	<a href="#">64368</a>	6409	1577	2	241	prolactin family 8, subfamily a, member 7 (27.4 kD) (Prl8a7) alternative variant aSep08, complete mRNA.
<a href="#">Prl8a9</a>	<a href="#">Prl8a9.bSep08</a>	<a href="#">171406</a>	4481	731	4	184	prolactin family 8, subfamily a, member 9 (20.9 kD) (Prl8a9) alternative variant bSep08, complete mRNA.
<a href="#">Prl8a9</a>	<a href="#">Prl8a9.cSep08</a>	<a href="#">171406</a>	4216	427	3	142	prolactin family 8, subfamily a, member 9 (Prl8a9) alternative variant cSep08, mRNA.
<a href="#">Prl8a9</a>	<a href="#">Prl8a9.dSep08</a>	<a href="#">171406</a>	4198	445	4	139	prolactin family 8, subfamily a, member 9 (Prl8a9) alternative variant dSep08, mRNA.
<a href="#">Prl8a9</a>	<a href="#">Prl8a9.eSep08</a>	<a href="#">171406</a>	2737	751	2	92	prolactin family 8, subfamily a, member 9 (Prl8a9) alternative variant eSep08, mRNA.
<a href="#">Prlr</a>	<a href="#">Prlr.cSep08</a>	<a href="#">24684</a>	23329	555	4	134	prolactin receptor (Prlr) alternative variant cSep08, mRNA.
<a href="#">Prlr</a>	<a href="#">Prlr.dSep08</a>	<a href="#">24684</a>	158504	548	3	84	putative protein (Prlr) alternative variant dSep08, mRNA.
<a href="#">Prlr</a>	<a href="#">Prlr.eSep08</a>	<a href="#">24684</a>	34265	627	5	17	putative protein (2.2 kD) (Prlr) alternative variant eSep08, mRNA.
<a href="#">Prlr</a>	<a href="#">Prlr.fSep08</a>	<a href="#">24684</a>	120132	567	3	35	putative protein (Prlr) alternative variant fSep08, mRNA.
<a href="#">Prlr</a>	<a href="#">Prlr.gSep08</a>	<a href="#">24684</a>	58263	484	2	34	putative protein (3.7 kD) (Prlr) alternative variant gSep08, mRNA.
<a href="#">Prmt1</a>	<a href="#">Prmt1.bSep08</a>	<a href="#">60421</a>	6945	784	8	253	protein arginine N-methyltransferase 1 (Prmt1) alternative variant bSep08, mRNA.
<a href="#">Prmt1</a>	<a href="#">Prmt1.cSep08</a>	<a href="#">60421</a>	4232	661	5	200	protein arginine N-methyltransferase 1 (Prmt1) alternative variant cSep08, mRNA.
<a href="#">Prmt2</a>	<a href="#">Prmt2.bSep08</a>	<a href="#">499420</a>	7373	562	1	187	protein arginine N-methyltransferase 2 (Prmt2) alternative variant bSep08, mRNA.
<a href="#">Prmt5</a>	<a href="#">Prmt5.bSep08</a>	<a href="#">364382</a>	1102	428	2	64	protein arginine N-methyltransferase 5 (Prmt5) alternative variant bSep08, mRNA.

<a href="#">Prmt7</a>	<a href="#">Prmt7.bSep08</a>	<a href="#">361402</a>	35122	865	7	284	protein arginine N-methyltransferase 7 (Prmt7) alternative variant bSep08, mRNA.
<a href="#">Prmt7</a>	<a href="#">Prmt7.cSep08</a>	<a href="#">361402</a>	2273	389	3	114	protein arginine N-methyltransferase 7 (Prmt7) alternative variant cSep08, mRNA.
<a href="#">Prnd</a>	<a href="#">Prnd.bSep08</a>	<a href="#">113910</a>	3193	2329	2	181	prion protein dublet (Prnd) alternative variant bSep08, mRNA.
<a href="#">Prnp</a>	<a href="#">Prnp.bSep08</a>	<a href="#">24686</a>	13954	821	1	194	prion protein (Prnp) alternative variant bSep08, mRNA.
<a href="#">Prnpip1</a>	<a href="#">Prnpip1.aSep08</a>	<a href="#">313535</a>	42599	986	6	328	prion protein interacting protein 1 (Prnpip1) alternative variant aSep08, mRNA.
<a href="#">Prnpip1</a>	<a href="#">Prnpip1.bSep08</a>	<a href="#">313535</a>	38986	807	5	268	prion protein interacting protein 1 (Prnpip1) alternative variant bSep08, mRNA.
<a href="#">Prnpip1</a>	<a href="#">Prnpip1.cSep08</a>	<a href="#">313535</a>	24300	793	3	144	prion protein interacting protein 1 (Prnpip1) alternative variant cSep08, mRNA.
<a href="#">Prnpip1</a>	<a href="#">Prnpip1.eSep08</a>	<a href="#">313535</a>	40180	346	4	76	prion protein interacting protein 1 (Prnpip1) alternative variant eSep08, mRNA.
<a href="#">PRO8NT.0</a>	<a href="#">PRO8NT.0.aSep08</a>		4934	1869		533	pre-mRNA processing factor 8 (PRO8NT.0) mRNA.
<a href="#">Proc</a>	<a href="#">Proc.bSep08</a>	<a href="#">25268</a>	8064	836	8	259	protein C (Proc) alternative variant bSep08, mRNA.
<a href="#">PROCN.0</a>	<a href="#">PROCN.0.aSep08</a>		2713	1177		391	pre-mRNA processing factor 8 (PROCN.0) mRNA.
<a href="#">Prodh2</a>	<a href="#">Prodh2.bSep08</a>	<a href="#">361538</a>	5998	750	4	109	proline dehydrogenase 2 CRA a (11.4 kD) (Prodh2) alternative variant bSep08, mRNA.
<a href="#">Prodh2</a>	<a href="#">Prodh2.cSep08</a>	<a href="#">361538</a>	3796	425	3	85	proline dehydrogenase 2 CRA a (9.6 kD) (Prodh2) alternative variant cSep08, mRNA.
<a href="#">Prok2</a>	<a href="#">Prok2.cSep08</a>	<a href="#">192206</a>	4185	832	2	90	prokineticin 2 (Prok2) alternative variant cSep08, mRNA.
<a href="#">Prom1</a>	<a href="#">Prom1.cSep08</a>	<a href="#">60357</a>	7932	521	4	129	prominin 1 (Prom1) alternative variant cSep08, mRNA.
<a href="#">Prom1</a>	<a href="#">Prom1.dSep08</a>	<a href="#">60357</a>	38223	1374	3	85	prominin 1 (9.3 kD) (Prom1) alternative variant dSep08, mRNA.
<a href="#">Prom2</a>	<a href="#">Prom2.bSep08</a>	<a href="#">192211</a>	1645	459	5	90	prominin 2 (Prom2) alternative variant bSep08, mRNA.
<a href="#">Prom2</a>	<a href="#">Prom2.cSep08</a>	<a href="#">192211</a>	528	416	2	70	prominin 2 (Prom2) alternative variant cSep08, mRNA.
<a href="#">Prosc</a>	<a href="#">Prosc.bSep08</a>	<a href="#">306544</a>	8033	852	3	108	proline synthetase co-transcribed (Prosc) alternative variant bSep08, mRNA.
<a href="#">Prosc</a>	<a href="#">Prosc.cSep08</a>	<a href="#">306544</a>	4018	466	5	85	proline synthetase co-transcribed (9.4 kD) (Prosc) alternative variant cSep08, mRNA.
<a href="#">Proz</a>	<a href="#">Proz.aSep08</a>	<a href="#">306608</a>	13125	3221	1	410	protein Z, vitamin K-dependent plasma glycoprotein (Proz) alternative variant aSep08, mRNA.
<a href="#">Pro_isomerase.0</a>	<a href="#">Pro_isomerase.0.cSep08</a>		10090	718	2	162	fusion protein (Pro_isomerase.0) alternative variant cSep08, mRNA.
<a href="#">PRP-2</a>	<a href="#">PRP-2.aSep08</a>	<a href="#">287750</a>	54922	3592	14	1153	leucine-rich repeat containing protein (PRP-2) alternative variant aSep08, mRNA.
<a href="#">PRP-2</a>	<a href="#">PRP-2.bSep08</a>	<a href="#">287750</a>	25279	759	4	252	putative protein of mammalian origin (PRP-2) alternative variant bSep08, mRNA.
<a href="#">PRP-2</a>	<a href="#">PRP-2.cSep08</a>	<a href="#">287750</a>	5680	2441	4	238	CRA a (PRP-2) alternative variant cSep08, mRNA.
<a href="#">PRP-2</a>	<a href="#">PRP-2.dSep08</a>	<a href="#">287750</a>	1208	1056	3	124	CRA a like (14.5 kD) (PRP-2) alternative variant dSep08, mRNA.
<a href="#">Prpf3</a>	<a href="#">Prpf3.bSep08</a>	<a href="#">361995</a>	6267	693	5	207	prp3 pre-mRNA processing factor 3 homolog (Prpf3) alternative variant bSep08, mRNA.

<a href="#">Prpf3</a>	<a href="#">Prpf3.cSep08</a>	<a href="#">361995</a>	6327	1078	7	202	prp3 Pre-mRNA processing factor 3 homolog (23.0 kD) (Prpf3) alternative variant cSep08, mRNA.
<a href="#">Prpf3</a>	<a href="#">Prpf3.dSep08</a>	<a href="#">361995</a>	9491	1069	7	177	prp3 pre-mRNA processing factor 3 homolog (Prpf3) alternative variant dSep08, mRNA.
<a href="#">Prpf3</a>	<a href="#">Prpf3.eSep08</a>	<a href="#">361995</a>	3331	459	4	93	prp3 pre-mRNA processing factor 3 homolog (Prpf3) alternative variant eSep08, mRNA.
<a href="#">Prpf3</a>	<a href="#">Prpf3.fSep08</a>	<a href="#">361995</a>	2850	1252	2	38	prp3 pre-mRNA processing factor 3 homolog (4.5 kD) (Prpf3) alternative variant fSep08, mRNA.
<a href="#">Prpf3</a>	<a href="#">Prpf3.gSep08</a>	<a href="#">361995</a>	8162	664	6	99	prp3 pre-mRNA processing factor 3 homolog (11.3 kD) (Prpf3) alternative variant gSep08, mRNA.
<a href="#">Prpf4b</a>	<a href="#">Prpf4b.bSep08</a>	<a href="#">291078</a>	2186	686	3	136	PRP4 pre-mRNA processing factor 4 homolog B (yeast) (15.8 kD) (Prpf4b) alternative variant bSep08, mRNA.
<a href="#">Prpf4b</a>	<a href="#">Prpf4b.dSep08</a>	<a href="#">291078</a>	5089	1369	8	136	PRP4 pre-mRNA processing factor 4 homolog B (yeast) (15.8 kD) (Prpf4b) alternative variant dSep08, mRNA.
<a href="#">Prpf6</a>	<a href="#">Prpf6.bSep08</a>	<a href="#">366276</a>	8292	1778	7	558	PRP6 pre-mRNA splicing factor 6 homolog ( <i>S. cerevisiae</i> ) (Prpf6) alternative variant bSep08, mRNA.
<a href="#">Prpf8</a>	<a href="#">Prpf8.aSep08</a>	<a href="#">287530</a>	14514	3986	23	1275	pre-mrna processing factor 8 (Prpf8) alternative variant aSep08, mRNA.
<a href="#">Prpf8</a>	<a href="#">Prpf8.bSep08</a>	<a href="#">287530</a>	2418	769	5	197	pre-mRNA processing factor 8 (Prpf8) alternative variant bSep08, mRNA.
<a href="#">Prpf8</a>	<a href="#">Prpf8.cSep08</a>	<a href="#">287530</a>	577	398	2	107	pre-mRNA processing factor 8 (Prpf8) alternative variant cSep08, mRNA.
<a href="#">Prpf8</a>	<a href="#">Prpf8.dSep08</a>	<a href="#">287530</a>	2523	761	4	87	splicing factor Prp8 (Prpf8) alternative variant dSep08, mRNA.
<a href="#">Prpf8</a>	<a href="#">Prpf8.eSep08</a>	<a href="#">287530</a>	558	428	2	67	pre-mRNA processing factor 8 (Prpf8) alternative variant eSep08, mRNA.
<a href="#">Prpf18</a>	<a href="#">Prpf18.bSep08</a>	<a href="#">171552</a>	13589	1789	4	506	PRP18 pre-mRNA processing factor 18 homolog (Prpf18) alternative variant bSep08, mRNA.
<a href="#">Prpf18</a>	<a href="#">Prpf18.cSep08</a>	<a href="#">171552</a>	12840	663	3	78	PRP18 pre-mRNA processing factor 18 homolog (Prpf18) alternative variant cSep08, mRNA.
<a href="#">Prpf18</a>	<a href="#">Prpf18.eSep08</a>	<a href="#">171552</a>	9733	468	2	72	PRP18 pre-mRNA processing factor 18 homolog (Prpf18) alternative variant eSep08, mRNA.
<a href="#">Prpf18</a>	<a href="#">Prpf18.fSep08</a>	<a href="#">171552</a>	4034	469	3	51	PRP18 pre-mRNA processing factor 18 homolog (6.2 kD) (Prpf18) alternative variant fSep08, mRNA.
<a href="#">Prpf18</a>	<a href="#">Prpf18.gSep08</a>	<a href="#">171552</a>	11232	360	3		
<a href="#">Prpf19</a>	<a href="#">Prpf19.bSep08</a>	<a href="#">246216</a>	4266	782	5	194	PRP19/PSO4 pre-mRNA processing factor 19 homolog ( <i>S. cerevisiae</i> ) (Prpf19) alternative variant bSep08, mRNA.
<a href="#">Prpf19</a>	<a href="#">Prpf19.cSep08</a>	<a href="#">246216</a>	5085	912	5	154	PRP19/PSO4 pre-mRNA processing factor 19 homolog ( <i>S. cerevisiae</i> ) (17.5 kD) (Prpf19) alternative variant cSep08, mRNA.
<a href="#">Prpf19</a>	<a href="#">Prpf19.dSep08</a>	<a href="#">246216</a>	8823	249	2	69	PRP19/PSO4 pre-mRNA processing factor 19 homolog ( <i>S. cerevisiae</i> ) (Prpf19) alternative variant dSep08, mRNA.
<a href="#">Prpf38a</a>	<a href="#">Prpf38a.aSep08</a>	<a href="#">298374</a>	11539	1495	9	312	PRP38 (37.4 kD) (Prpf38a) alternative variant aSep08, mRNA.
<a href="#">Prpf38a</a>	<a href="#">Prpf38a.bSep08</a>	<a href="#">298374</a>	1404	423	1	33	putative protein of eukaryotic origin (Prpf38a) alternative variant bSep08, mRNA.



<a href="#">Prpf38b</a>	<a href="#">Prpf38b.bSep08</a>	<a href="#">499691</a>	3970	750	3	168	putative protein of eukaryotic origin (Prpf38b) alternative variant bSep08, mRNA.
<a href="#">Prpf38b</a>	<a href="#">Prpf38b.cSep08</a>	<a href="#">499691</a>	8599	2139	8	139	PRP38 pre-mRNA processing factor 38 domain (Prpf38b) alternative variant cSep08, mRNA.
<a href="#">Prpf38b</a>	<a href="#">Prpf38b.dSep08</a>	<a href="#">499691</a>	3539	372	4	52	putative protein (Prpf38b) alternative variant dSep08, mRNA.
<a href="#">Prpf38b</a>	<a href="#">Prpf38b.fSep08</a>	<a href="#">499691</a>	1104	522	2	37	putative protein of metazoan origin (Prpf38b) alternative variant fSep08, mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.bSep08</a>	<a href="#">314171</a>	6064	2588	6	278	prp39 pre-mRNA processing factor 39 homolog CRA b (33.2 kD) (Prpf39) alternative variant bSep08, mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.cSep08</a>	<a href="#">314171</a>	24854	3645	14	271	prp39 pre-mRNA processing factor 39 homolog CRA a (31.6 kD) (Prpf39) alternative variant cSep08, complete mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.dSep08</a>	<a href="#">314171</a>	8412	742	6	247	prp39 pre-mRNA processing factor 39 homolog CRA b (Prpf39) alternative variant dSep08, mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.eSep08</a>	<a href="#">314171</a>	2587	876	3	111	prp39 pre-mRNA processing factor 39 homolog CRA a (Prpf39) alternative variant eSep08, mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.fSep08</a>	<a href="#">314171</a>	7739	698	3	53	prp39 pre-mRNA processing factor 39 homolog (6.6 kD) (Prpf39) alternative variant fSep08, complete mRNA.
<a href="#">Prpf39</a>	<a href="#">Prpf39.iSep08</a>	<a href="#">314171</a>	1583	412	3	9	putative protein (1.2 kD) (Prpf39) alternative variant iSep08, mRNA.
<a href="#">Prpf40a</a>	<a href="#">Prpf40a.bSep08</a>	<a href="#">295607</a>	5262	684	7	146	PRP40 pre-mRNA processing factor 40 homolog A ( <i>S. cerevisiae</i> ) (Prpf40a) alternative variant bSep08, mRNA.
<a href="#">Prph</a>	<a href="#">Prph.bSep08</a>	<a href="#">24688</a>	3196	1517	7	311	peripherin (35.8 kD) (Prph) alternative variant bSep08, mRNA.
<a href="#">Prph</a>	<a href="#">Prph.cSep08</a>	<a href="#">24688</a>	2149	995	4	88	peripherin (10.3 kD) (Prph) alternative variant cSep08, mRNA.
<a href="#">Prph</a>	<a href="#">Prph.dSep08</a>	<a href="#">24688</a>	793	527	2	53	peripherin (Prph) alternative variant dSep08, mRNA.
<a href="#">Prps1</a>	<a href="#">Prps1.bSep08</a>	<a href="#">29562</a>	21112	540	1	152	phosphoribosyl pyrophosphate synthetase 1 (16.8 kD) (Prps1) alternative variant bSep08, mRNA.
<a href="#">Prps2</a>	<a href="#">Prps2.bSep08</a>	<a href="#">24689</a>	6692	322	2	64	phosphoribosyl pyrophosphate synthetase 2 (Prps2) alternative variant bSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.bSep08</a>	<a href="#">64390</a>	14558	1486	4	210	synthetase-associated protein (Prpsap1) alternative variant bSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.cSep08</a>	<a href="#">64390</a>	49745	1041	9	150	synthetase-associated protein (16.5 kD) (Prpsap1) alternative variant cSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.dSep08</a>	<a href="#">64390</a>	6778	1265	5	148	synthetase-associated protein (16.3 kD) (Prpsap1) alternative variant dSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.gSep08</a>	<a href="#">64390</a>	2121	1553	2	74	synthetase-associated protein (8.3 kD) (Prpsap1) alternative variant gSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.hSep08</a>	<a href="#">64390</a>	23652	1535	4	34	putative protein (3.8 kD) (Prpsap1) alternative variant hSep08, mRNA.
<a href="#">Prpsap1</a>	<a href="#">Prpsap1.iSep08</a>	<a href="#">64390</a>	22834	671	3	54	putative protein (6.0 kD) (Prpsap1) alternative variant iSep08, mRNA.
<a href="#">Prpsap2</a>	<a href="#">Prpsap2.bSep08</a>	<a href="#">117272</a>	33102	1790	12	369	phosphoribosyl pyrophosphate synthetase-associated protein 2 (40.9 kD) (Prpsap2) alternative variant bSep08, mRNA.

<a href="#">Prpsap2</a>	<a href="#">Prpsap2.cSep08</a>	<a href="#">117272</a>	35045	1634	11	320	phosphoribosyl pyrophosphate synthetase-associated protein 2 (35.5 kD) (Prpsap2) alternative variant cSep08, complete mRNA.
<a href="#">Prpsap2</a>	<a href="#">Prpsap2.dSep08</a>	<a href="#">117272</a>	33166	915	9	208	phosphoribosyl pyrophosphate synthetase-associated protein 2 (Prpsap2) alternative variant dSep08, mRNA.
<a href="#">Prpsap2</a>	<a href="#">Prpsap2.eSep08</a>	<a href="#">117272</a>	16756	786	8	172	phosphoribosyl pyrophosphate synthetase-associated protein 2 (Prpsap2) alternative variant eSep08, mRNA.
<a href="#">Prpsap2</a>	<a href="#">Prpsap2.fSep08</a>	<a href="#">117272</a>	28297	887	8	152	phosphoribosyl pyrophosphate synthetase-associated protein 2 (Prpsap2) alternative variant fSep08, mRNA.
<a href="#">Prpsap2</a>	<a href="#">Prpsap2.gSep08</a>	<a href="#">117272</a>	12903	741	7	126	phosphoribosyl pyrophosphate synthetase-associated protein 2 (Prpsap2) alternative variant gSep08, mRNA.
<a href="#">Prr3</a>	<a href="#">Prr3.bSep08</a>	<a href="#">361788</a>	4725	2121	4	143	proline-rich polypeptide 3 (15.8 kD) (Prr3) alternative variant bSep08, mRNA.
<a href="#">Prr3</a>	<a href="#">Prr3.cSep08</a>	<a href="#">361788</a>	4246	1140	5	116	proline-rich polypeptide 3 (Prr3) alternative variant cSep08, mRNA.
<a href="#">Prr3</a>	<a href="#">Prr3.dSep08</a>	<a href="#">361788</a>	3679	744	5	101	proline-rich polypeptide 3 (Prr3) alternative variant dSep08, mRNA.
<a href="#">Prr5</a>	<a href="#">Prr5.bSep08</a>	<a href="#">315189</a>	30638	761	8	253	proline rich 5 (renal) (Prr5) alternative variant bSep08, mRNA.
<a href="#">Prr5</a>	<a href="#">Prr5.cSep08</a>	<a href="#">315189</a>	30677	680	7	213	proline rich 5 (renal) (Prr5) alternative variant cSep08, mRNA.
<a href="#">Prr6</a>	<a href="#">Prr6.aSep08</a>	<a href="#">501702</a>	8522	1780	3	513	proline-rich polypeptide 6 (Prr6) alternative variant aSep08, mRNA.
<a href="#">Prr6</a>	<a href="#">Prr6.bSep08</a>	<a href="#">501702</a>	13347	773	5	185	proline-rich polypeptide 6 (Prr6) alternative variant bSep08, mRNA.
<a href="#">Prr6</a>	<a href="#">Prr6.cSep08</a>	<a href="#">501702</a>	13341	652	4	133	proline-rich polypeptide 6 (Prr6) alternative variant cSep08, mRNA.
<a href="#">Prr11</a>	<a href="#">Prr11.bSep08</a>	<a href="#">360591</a>	10548	1782	1	139	proline rich 11 (Prr11) alternative variant bSep08, mRNA.
<a href="#">Prr12</a>	<a href="#">Prr12.aSep08</a>	<a href="#">361569</a>	16415	2875		651	proline rich 12 (Prr12) mRNA.
<a href="#">Prr13</a>	<a href="#">Prr13.aSep08</a>	<a href="#">363004</a>	1323	983	2	177	proline rich 13 (18.6 kD) (Prr13) alternative variant aSep08, mRNA.
<a href="#">Prr14</a>	<a href="#">Prr14.aSep08</a>	<a href="#">691898</a>	5422	2740	11	649	proline rich 14 (Prr14) alternative variant aSep08, mRNA.
<a href="#">Prr14</a>	<a href="#">Prr14.bSep08</a>	<a href="#">691898</a>	2626	1323	7	359	proline rich 14 (Prr14) alternative variant bSep08, mRNA.
<a href="#">Prr14</a>	<a href="#">Prr14.cSep08</a>	<a href="#">691898</a>	2921	856	5	169	proline rich 14 (Prr14) alternative variant cSep08, mRNA.
<a href="#">Prr14</a>	<a href="#">Prr14.dSep08</a>	<a href="#">691898</a>	3339	692	6	159	proline rich 14 (Prr14) alternative variant dSep08, mRNA.
<a href="#">Prr19</a>	<a href="#">Prr19.aSep08</a>	<a href="#">681408</a>	3017	1286		363	proline rich 19 (39.9 kD) (Prr19) mRNA.
<a href="#">Prrg1</a>	<a href="#">Prrg1.aSep08</a>	<a href="#">363472</a>	113616	1310		217	proline rich Gla (G-carboxyglutamic acid) 1 (Prrg1) mRNA.
<a href="#">Prrg2</a>	<a href="#">Prrg2.aSep08</a>	<a href="#">361570</a>	8236	1364	2	198	proline-rich Gla (G-carboxyglutamic acid) polypeptide 2 (22.3 kD) (Prrg2) alternative variant aSep08, complete mRNA.
<a href="#">Prrg4</a>	<a href="#">Prrg4.bSep08</a>	<a href="#">499847</a>	14554	1788	2	112	proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane) (12.2 kD) (Prrg4) alternative variant bSep08, mRNA.
<a href="#">Prrg4</a>	<a href="#">Prrg4.cSep08</a>	<a href="#">499847</a>	12959	390	2	97	proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane) (Prrg4) alternative variant cSep08, mRNA.

<a href="#">Prrt1</a>	<a href="#">Prrt1.aSep08</a>	<a href="#">406167</a>	3642	2544	1	357	proline-rich transmembrane protein 1 (Prrt1) alternative variant aSep08, mRNA.
<a href="#">Prrt1</a>	<a href="#">Prrt1.bSep08</a>	<a href="#">406167</a>	1489	753	2	179	proline-rich transmembrane protein 1 (Prrt1) alternative variant bSep08, mRNA.
<a href="#">Prrt3</a>	<a href="#">Prrt3.bSep08</a>	<a href="#">502873</a>	5875	1726	3	386	proline-rich transmembrane protein 3 (40.1 kD) (Prrt3) alternative variant bSep08, mRNA.
<a href="#">Prrx2</a>	<a href="#">Prrx2.bSep08</a>	<a href="#">113931</a>	35219	511	3	169	paired related homeobox 2 (Prrx2) alternative variant bSep08, mRNA.
<a href="#">Prrx2</a>	<a href="#">Prrx2.cSep08</a>	<a href="#">113931</a>	3422	695	3	123	paired related homeobox 2 (Prrx2) alternative variant cSep08, mRNA.
<a href="#">Prss7</a>	<a href="#">Prss7.bSep08</a>	<a href="#">288291</a>	3050	366	1	70	protease, serine, 7 (enterokinase) (Prss7) alternative variant bSep08, mRNA.
<a href="#">Prss12</a>	<a href="#">Prss12.bSep08</a>	<a href="#">85266</a>	8588	783	1	207	protease, serine, 12 neurotrypsin (motopsin) (Prss12) alternative variant bSep08, mRNA.
<a href="#">Prss16</a>	<a href="#">Prss16.bSep08</a>	<a href="#">364719</a>	1457	656	2	166	protease, serine, 16 (thymus) (18.2 kD) (Prss16) alternative variant bSep08, mRNA.
<a href="#">Prss36</a>	<a href="#">Prss36.aSep08</a>	<a href="#">497040</a>	1015	599		199	protease, serine, 36 (Prss36) mRNA.
<a href="#">Prtfdc1</a>	<a href="#">Prtfdc1.bSep08</a>	<a href="#">291355</a>	119563	1965	10	215	phosphoribosyltransferase (24.6 kD) (Prtfdc1) alternative variant bSep08, complete mRNA.
<a href="#">Prtfdc1</a>	<a href="#">Prtfdc1.cSep08</a>	<a href="#">291355</a>	102906	647	6	136	phosphoribosyl transferase 1 (Prtfdc1) alternative variant cSep08, mRNA.
<a href="#">Prtfdc1</a>	<a href="#">Prtfdc1.dSep08</a>	<a href="#">291355</a>	15783	338	3	96	putative protein of ancient origin (Prtfdc1) alternative variant dSep08, mRNA.
<a href="#">Prtfdc1</a>	<a href="#">Prtfdc1.eSep08</a>	<a href="#">291355</a>	35563	408	2	34	putative protein (Prtfdc1) alternative variant eSep08, mRNA.
<a href="#">Prune</a>	<a href="#">Prune.bSep08</a>	<a href="#">310664</a>	16016	421	3	55	prune homolog (Drosophila) (Prune) alternative variant bSep08, mRNA.
<a href="#">Psap</a>	<a href="#">Psap.bSep08</a>	<a href="#">25524</a>	26906	2554	14	553	prosaposin (61.0 kD) (Psap) alternative variant bSep08, mRNA.
<a href="#">Psap</a>	<a href="#">Psap.cSep08</a>	<a href="#">25524</a>	1484	738	2	71	prosaposin (7.6 kD) (Psap) alternative variant cSep08, mRNA.
<a href="#">Psbpc1</a>	<a href="#">Psbpc1.bSep08</a>	<a href="#">309203</a>	3004	461	2	73	prostatic steroid binding protein C1 (8.6 kD) (Psbpc1) alternative variant bSep08, complete mRNA.
<a href="#">Pscd1</a>	<a href="#">Pscd1.bSep08</a>	<a href="#">116691</a>	16801	797	7	161	pleckstrin homology, Sec7 and coiled-coil domains 1 (Pscd1) alternative variant bSep08, mRNA.
<a href="#">Pscd2</a>	<a href="#">Pscd2.bSep08</a>	<a href="#">116692</a>	2748	688	4	145	pleckstrin homology, Sec7 and coiled-coil domains 2 (Pscd2) alternative variant bSep08, mRNA.
<a href="#">Pscdbp</a>	<a href="#">Pscdbp.bSep08</a>	<a href="#">311047</a>	27878	2521	7	152	pleckstrin homology, Sec7 and coiled-coil domains, binding protein (Pscdbp) alternative variant bSep08, mRNA.
<a href="#">Psd</a>	<a href="#">Psd.aSep08</a>	<a href="#">171381</a>	2571	1118	2	294	pleckstrin-like (Psd) alternative variant aSep08, mRNA.
<a href="#">Psd3</a>	<a href="#">Psd3.aSep08</a>	<a href="#">306380</a>	28868	623		166	putative protein, with a coiled coil domain, of bilateral origin (Psd3) mRNA.
<a href="#">Psd4</a>	<a href="#">Psd4.bSep08</a>	<a href="#">311785</a>	22582	773	2	89	putative protein (Psd4) alternative variant bSep08, mRNA.
<a href="#">Psen1</a>	<a href="#">Psen1.bSep08</a>	<a href="#">29192</a>	39325	730	6	183	presenilin 1 CRA a (21.1 kD) (Psen1) alternative variant bSep08, mRNA.
<a href="#">Psen1</a>	<a href="#">Psen1.cSep08</a>	<a href="#">29192</a>	2332	798	2	100	presenilin 1 I-467 precursor (11.0 kD) (Psen1) alternative variant cSep08, mRNA.

<a href="#">Psen2</a>	<a href="#">Psen2.bSep08</a>	<a href="#">81751</a>	2807	1317	3	107	presenilin 2 (11.8 kD) (Psen2) alternative variant bSep08, mRNA.
<a href="#">Psen2</a>	<a href="#">Psen2.cSep08</a>	<a href="#">81751</a>	2083	512	2	67	presenilin 2 (11.4 kD) (Psen2) alternative variant cSep08, mRNA.
<a href="#">PseudoU_synth_2.0</a>	<a href="#">PseudoU_synth_2.0.aSep08</a>		1260	398		132	pseudouridine synthase (PseudoU_synth_2.0) mRNA.
<a href="#">Psg-ps1_predicted</a>	<a href="#">Psg-ps1_predicted.aSep08</a>	<a href="#">308396</a>	6289	1203		400	pregnancy specific glycoprotein pseudogene 1 (predicted) (Psg-ps1_predicted) mRNA.
<a href="#">Psg19</a>	<a href="#">Psg19.bSep08</a>	<a href="#">24256</a>	10322	2470	4	355	pregnancy specific glycoprotein 19 (39.9 kD) (Psg19) alternative variant bSep08, complete mRNA.
<a href="#">Psg19</a>	<a href="#">Psg19.cSep08</a>	<a href="#">24256</a>	971	350	2	116	pregnancy specific glycoprotein 19 (Psg19) alternative variant cSep08, mRNA.
<a href="#">Psg19</a>	<a href="#">Psg19.dSep08</a>	<a href="#">24256</a>	1150	378	2	114	pregnancy specific glycoprotein 19 (Psg19) alternative variant dSep08, mRNA.
<a href="#">Psgb1</a>	<a href="#">Psgb1.bSep08</a>	<a href="#">59313</a>	2191	542	2	125	pregnancy-specific beta 1-glycoprotein (Psgb1) alternative variant bSep08, mRNA.
<a href="#">Psgb1</a>	<a href="#">Psgb1.cSep08</a>	<a href="#">59313</a>	5157	378	2	104	pregnancy-specific beta 1-glycoprotein (Psgb1) alternative variant cSep08, mRNA.
<a href="#">PSI.0</a>	<a href="#">PSI.0.aSep08</a>		1756	495		164	plexin A1 CRA c (PSI.0) mRNA.
<a href="#">PSI.1</a>	<a href="#">PSI.1.aSep08</a>		2665	181		60	sema 4G (PSI.1) mRNA.
<a href="#">PSI.2</a>	<a href="#">PSI.2.aSep08</a>		51047	941		313	attractin-like 1 CRA a (PSI.2) mRNA.
<a href="#">Psp1</a>	<a href="#">Psp1.bSep08</a>	<a href="#">313323</a>	28512	1783	9	357	PC4 and SFRS1 interacting protein 1 (Psp1) alternative variant bSep08, mRNA.
<a href="#">Psp1</a>	<a href="#">Psp1.cSep08</a>	<a href="#">313323</a>	22493	1521	9	285	PC4 and SFRS1 interacting protein 1 (Psp1) alternative variant cSep08, mRNA.
<a href="#">Psp1</a>	<a href="#">Psp1.dSep08</a>	<a href="#">313323</a>	11908	1101	6	205	PC4 and SFRS1 interacting protein 1 (Psp1) alternative variant dSep08, mRNA.
<a href="#">Psp1</a>	<a href="#">Psp1.eSep08</a>	<a href="#">313323</a>	6430	1494	5	141	PC4 and SFRS1 interacting protein 1 (16.5 kD) (Psp1) alternative variant eSep08, mRNA.
<a href="#">Pskh1</a>	<a href="#">Pskh1.aSep08</a>	<a href="#">364993</a>	12510	1065	2	294	protein serine kinase H1 (Pskh1) alternative variant aSep08, mRNA.
<a href="#">Psm2</a>	<a href="#">Psm2.bSep08</a>	<a href="#">29669</a>	4238	495		111	proteasome (prosome, macropain) subunit, alpha type 2 (Psm2) alternative variant bSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.bSep08</a>	<a href="#">29670</a>	11917	526	7	174	proteasome alpha (Psm3l) alternative variant bSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.bSep08</a>	<a href="#">408248</a>	11917	526	7	174	proteasome alpha (Psm3l) alternative variant bSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.cSep08</a>	<a href="#">29670</a>	4491	620	3	56	proteasome alpha type 3 (6.7 kD) (Psm3l) alternative variant cSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.cSep08</a>	<a href="#">408248</a>	4491	620	3	56	proteasome alpha type 3 (6.7 kD) (Psm3l) alternative variant cSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.dSep08</a>	<a href="#">29670</a>	1151	375	2	22	putative protein (Psm3l) alternative variant dSep08, mRNA.
<a href="#">Psm3l</a>	<a href="#">Psm3l.dSep08</a>	<a href="#">408248</a>	1151	375	2	22	putative protein (Psm3l) alternative variant dSep08, mRNA.
<a href="#">Psm4</a>	<a href="#">Psm4.bSep08</a>	<a href="#">29671</a>	6474	818	8	219	proteasome (prosome, macropain) subunit, alpha type 4 (24.3 kD) (Psm4) alternative variant bSep08, mRNA.

<a href="#">Psm4</a>	<a href="#">Psm4.cSep08</a>	<a href="#">29671</a>	4999	658	7	208	proteasome (prosome, macropain) subunit, alpha type 4 (Psm4) alternative variant cSep08, mRNA.
<a href="#">Psm4</a>	<a href="#">Psm4.eSep08</a>	<a href="#">29671</a>	1796	481	2	57	proteasome (prosome, macropain) subunit, alpha type 4 (Psm4) alternative variant eSep08, mRNA.
<a href="#">Psm5</a>	<a href="#">Psm5.aSep08</a>	<a href="#">29672</a>	22984	976	9	244	proteasome (prosome, macropain) subunit, alpha type 5 (26.5 kD) (Psm5) alternative variant aSep08, mRNA.
<a href="#">Psm6</a>	<a href="#">Psm6.bSep08</a>	<a href="#">29673</a>	30996	794	1	243	proteasome (prosome, macropain) subunit, alpha type 6 (Psm6) alternative variant bSep08, mRNA.
<a href="#">Psm7</a>	<a href="#">Psm7.bSep08</a>	<a href="#">29674</a>	6370	1717	5	174	proteasome (prosome, macropain) subunit, alpha type 7 (19.5 kD) (Psm7) alternative variant bSep08, mRNA.
<a href="#">Psm7</a>	<a href="#">Psm7.cSep08</a>	<a href="#">29674</a>	4631	406	4	130	proteasome (prosome, macropain) subunit, alpha type 7 (Psm7) alternative variant cSep08, mRNA.
<a href="#">Psm7</a>	<a href="#">Psm7.eSep08</a>	<a href="#">29674</a>	2353	813	3	45	proteasome (prosome, macropain) subunit, alpha type 7 (Psm7) alternative variant eSep08, mRNA.
<a href="#">Psm7</a>	<a href="#">Psm7.fSep08</a>	<a href="#">29674</a>	615	499	2	55	proteasome (prosome, macropain) subunit, alpha type 7 (Psm7) alternative variant fSep08, mRNA.
<a href="#">Psm1</a>	<a href="#">Psm1.bSep08</a>	<a href="#">94198</a>	17200	783	6	201	proteasome (prosome, macropain) subunit, beta type 1 (22.3 kD) (Psm1) alternative variant bSep08, mRNA.
<a href="#">Psm1</a>	<a href="#">Psm1.cSep08</a>	<a href="#">94198</a>	19306	995	6	201	proteasome (prosome, macropain) subunit, beta type 1 (22.3 kD) (Psm1) alternative variant cSep08, complete mRNA.
<a href="#">Psm3</a>	<a href="#">Psm3.aSep08</a>	<a href="#">29676</a>	4596	794	3	197	proteasome (prosome, macropain) subunit, beta type 3 (22.0 kD) (Psm3) alternative variant aSep08, mRNA.
<a href="#">Psm3</a>	<a href="#">Psm3.bSep08</a>	<a href="#">29676</a>	7012	744	6	173	proteasome (prosome, macropain) subunit, beta type 3 (19.4 kD) (Psm3) alternative variant bSep08, complete mRNA.
<a href="#">Psm7</a>	<a href="#">Psm7.bSep08</a>	<a href="#">85492</a>	22537	742	1	86	proteasome (prosome, macropain) subunit, beta type 7 (Psm7) alternative variant bSep08, mRNA.
<a href="#">Psm8</a>	<a href="#">Psm8.bSep08</a>	<a href="#">24968</a>	2916	1457	5	189	proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional peptidase 7) (20.6 kD) (Psm8) alternative variant bSep08, complete mRNA.
<a href="#">Psm8</a>	<a href="#">Psm8.cSep08</a>	<a href="#">24968</a>	3157	2066	4	160	proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional peptidase 7) (17.5 kD) (Psm8) alternative variant cSep08, mRNA.
<a href="#">Psm8</a>	<a href="#">Psm8.dSep08</a>	<a href="#">24968</a>	463	276	2	83	proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional peptidase 7) (Psm8) alternative variant dSep08, mRNA.
<a href="#">Psm1</a>	<a href="#">Psm1.bSep08</a>	<a href="#">117263</a>	6687	758	1	252	protease (prosome, macropain) 26S subunit, ATPase 1 (Psm1) alternative variant bSep08, mRNA.
<a href="#">Psm3</a>	<a href="#">Psm3.bSep08</a>	<a href="#">29677</a>	854	748	1	63	proteasome (prosome, macropain) 26S subunit, ATPase 3 (Psm3) alternative variant bSep08, mRNA.
<a href="#">Psm3ip</a>	<a href="#">Psm3ip.bSep08</a>	<a href="#">140938</a>	774	551	3	103	psmc3 interacting protein (Psm3ip) alternative variant bSep08, mRNA.
<a href="#">Psm3ip</a>	<a href="#">Psm3ip.cSep08</a>	<a href="#">140938</a>	858	777	2	86	psmc3 interacting protein CRA c (9.9 kD) (Psm3ip) alternative variant cSep08, mRNA.
<a href="#">Psm3ip</a>	<a href="#">Psm3ip.dSep08</a>	<a href="#">140938</a>	438	355	2	71	proteasome ATPase 3 interacting protein CRA b (Psm3ip) alternative variant dSep08, mRNA.

<a href="#">Psmc4</a>	<a href="#">Psmc4.bSep08</a>	<a href="#">117262</a>	1102	670	4	110	proteasome (prosome, macropain) 26S subunit, ATPase, 4 (12.8 kD) (Psmc4) alternative variant bSep08, mRNA.
<a href="#">Psmc5</a>	<a href="#">Psmc5.bSep08</a>	<a href="#">81827</a>	5860	794	9	139	protease (prosome, macropain) 26S subunit, ATPase 5 (15.9 kD) (Psmc5) alternative variant bSep08, mRNA.
<a href="#">Psmc5</a>	<a href="#">Psmc5.cSep08</a>	<a href="#">81827</a>	4303	409	5	126	protease (prosome, macropain) 26S subunit, ATPase 5 (Psmc5) alternative variant cSep08, mRNA.
<a href="#">Psmc5</a>	<a href="#">Psmc5.dSep08</a>	<a href="#">81827</a>	967	664	4	71	protease (prosome, macropain) 26S subunit, ATPase 5 (8.0 kD) (Psmc5) alternative variant dSep08, mRNA.
<a href="#">Psmc6</a>	<a href="#">Psmc6.aSep08</a>	<a href="#">289990</a>	21299	1726	14	407	proteasome (prosome, macropain) 26S subunit, ATPase, 6 (Psmc6) alternative variant aSep08, mRNA.
<a href="#">Psmc6</a>	<a href="#">Psmc6.bSep08</a>	<a href="#">289990</a>	11353	812	9	200	proteasome (prosome, macropain) 26S subunit, ATPase, 6 (Psmc6) alternative variant bSep08, mRNA.
<a href="#">Psmc1</a>	<a href="#">Psmc1.bSep08</a>	<a href="#">83806</a>	16391	825	7	275	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (Psmc1) alternative variant bSep08, mRNA.
<a href="#">Psmc1</a>	<a href="#">Psmc1.cSep08</a>	<a href="#">83806</a>	38475	727	6	183	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (Psmc1) alternative variant cSep08, mRNA.
<a href="#">Psmc1</a>	<a href="#">Psmc1.dSep08</a>	<a href="#">83806</a>	19460	653	5	142	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (Psmc1) alternative variant dSep08, mRNA.
<a href="#">Psmc1</a>	<a href="#">Psmc1.fSep08</a>	<a href="#">83806</a>	2166	614	2	48	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (Psmc1) alternative variant fSep08, mRNA.
<a href="#">Psmc1</a>	<a href="#">Psmc1.gSep08</a>	<a href="#">83806</a>	552	367	2	34	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (Psmc1) alternative variant gSep08, mRNA.
<a href="#">Psmc2</a>	<a href="#">Psmc2.bSep08</a>	<a href="#">287984</a>	2480	735	1	220	proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 (Psmc2) alternative variant bSep08, mRNA.
<a href="#">Psmc3</a>	<a href="#">Psmc3.bSep08</a>	<a href="#">287670</a>	2216	750	1	121	proteasome (prosome, macropain) 26S subunit, non-ATPase, 3 (Psmc3) alternative variant bSep08, mRNA.
<a href="#">Psmc4</a>	<a href="#">Psmc4.bSep08</a>	<a href="#">83499</a>	9217	1281	9	377	proteasome (prosome, macropain) 26S subunit, non-ATPase, 4 (40.7 kD) (Psmc4) alternative variant bSep08, complete mRNA.
<a href="#">Psmc4</a>	<a href="#">Psmc4.cSep08</a>	<a href="#">83499</a>	9179	1397	8	273	proteasome (prosome, macropain) 26S subunit, non-ATPase, 4 (29.2 kD) (Psmc4) alternative variant cSep08, complete mRNA.
<a href="#">Psmc5</a>	<a href="#">Psmc5.bSep08</a>	<a href="#">296651</a>	10739	768	6	191	proteasome (prosome, macropain) 26S subunit, non-ATPase, 5 (Psmc5) alternative variant bSep08, mRNA.
<a href="#">Psmc5</a>	<a href="#">Psmc5.cSep08</a>	<a href="#">296651</a>	3828	354	2	110	proteasome (prosome, macropain) 26S subunit, non-ATPase, 5 (Psmc5) alternative variant cSep08, mRNA.
<a href="#">Psmc6</a>	<a href="#">Psmc6.bSep08</a>	<a href="#">289924</a>	6975	264	3	88	proteasome (prosome, macropain) 26S subunit, non-ATPase, 6 (Psmc6) alternative variant bSep08, mRNA.
<a href="#">Psmc7</a>	<a href="#">Psmc7.bSep08</a>	<a href="#">307821</a>	6357	746	3	158	proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Psmc7) alternative variant bSep08, mRNA.
<a href="#">Psmc8</a>	<a href="#">Psmc8.aSep08</a>	<a href="#">292766</a>	6833	1520	7	353	proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (39.9 kD) (Psmc8) alternative variant aSep08, mRNA.
<a href="#">Psmc8</a>	<a href="#">Psmc8.bSep08</a>	<a href="#">292766</a>	5389	709	6	215	proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (Psmc8) alternative variant bSep08, mRNA.
<a href="#">Psmc8</a>	<a href="#">Psmc8.cSep08</a>	<a href="#">292766</a>	5410	686	6	136	proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (Psmc8) alternative variant cSep08, mRNA.

<a href="#">Psm8</a>	<a href="#">Psm8.dSep08</a>	<a href="#">292766</a>	1508	875	2	90	proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (10.5 kD) (Psm8) alternative variant dSep08, mRNA.
<a href="#">Psm9</a>	<a href="#">Psm9.bSep08</a>	<a href="#">161475</a>	20047	747	3	167	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 (Psm9) alternative variant bSep08, mRNA.
<a href="#">Psm9</a>	<a href="#">Psm9.cSep08</a>	<a href="#">161475</a>	10474	775	5	120	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 (Psm9) alternative variant cSep08, mRNA.
<a href="#">Psm9</a>	<a href="#">Psm9.eSep08</a>	<a href="#">161475</a>	8803	699	5	91	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 (Psm9) alternative variant eSep08, mRNA.
<a href="#">Psm11</a>	<a href="#">Psm11.bSep08</a>	<a href="#">303353</a>	45221	2737	12	343	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 (39.2 kD) (Psm11) alternative variant bSep08, complete mRNA.
<a href="#">Psm11</a>	<a href="#">Psm11.cSep08</a>	<a href="#">303353</a>	13378	888	8	207	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 (Psm11) alternative variant cSep08, mRNA.
<a href="#">Psm11</a>	<a href="#">Psm11.dSep08</a>	<a href="#">303353</a>	12376	637	7	159	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 (Psm11) alternative variant dSep08, mRNA.
<a href="#">Psm12</a>	<a href="#">Psm12.bSep08</a>	<a href="#">287772</a>	12629	758	6	235	proteasome (prosome, macropain) 26S subunit, non-ATPase, 12 (Psm12) alternative variant bSep08, mRNA.
<a href="#">Psm13</a>	<a href="#">Psm13.bSep08</a>	<a href="#">365388</a>	6218	771	7	204	proteasome non-ATPase (23.0 kD) (Psm13) alternative variant bSep08, mRNA.
<a href="#">Psm13</a>	<a href="#">Psm13.cSep08</a>	<a href="#">365388</a>	3480	698	5	117	proteasome non-ATPase (Psm13) alternative variant cSep08, mRNA.
<a href="#">Psm13</a>	<a href="#">Psm13.dSep08</a>	<a href="#">365388</a>	1722	1612	2	92	proteasome non-ATPase (10.6 kD) (Psm13) alternative variant dSep08, mRNA.
<a href="#">Psme1</a>	<a href="#">Psme1.aSep08</a>	<a href="#">29630</a>	2883	1005	10	286	proteasome (prosome, macropain) 28 subunit, alpha (Psme1) alternative variant aSep08, mRNA.
<a href="#">Psme1</a>	<a href="#">Psme1.bSep08</a>	<a href="#">29630</a>	2830	1356	8	178	proteasome (prosome, macropain) 28 subunit, alpha (19.6 kD) (Psme1) alternative variant bSep08, complete mRNA.
<a href="#">Psme1</a>	<a href="#">Psme1.cSep08</a>	<a href="#">29630</a>	1830	712	5	155	proteasome (prosome, macropain) 28 subunit, alpha (17.0 kD) (Psme1) alternative variant cSep08, mRNA.
<a href="#">Psme1</a>	<a href="#">Psme1.dSep08</a>	<a href="#">29630</a>	1146	461	1	18	proteasome (prosome, macropain) 28 subunit, alpha (2.1 kD) (Psme1) alternative variant dSep08, mRNA.
<a href="#">Psme2.1</a>	<a href="#">Psme2.1.aSep08</a>	<a href="#">29614</a>	4302	1345	11	250	proteasome (prosome, macropain) 28 subunit, beta (Psme2.1) alternative variant aSep08, mRNA.
<a href="#">Psme2.1</a>	<a href="#">Psme2.1.cSep08</a>	<a href="#">29614</a>	629	539	2	55	proteasome (prosome, macropain) 28 subunit, beta (6.4 kD) (Psme2.1) alternative variant cSep08, mRNA.
<a href="#">Psme3</a>	<a href="#">Psme3.aSep08</a>	<a href="#">287716</a>	9046	3820		325	proteasome (prosome, macropain) 28 subunit, 3 (Psme3) mRNA.
<a href="#">Psme4</a>	<a href="#">Psme4.aSep08</a>	<a href="#">498433</a>	44846	4305	23	929	proteasome (prosome, macropain) activator subunit 4 (Psme4) alternative variant aSep08, mRNA.
<a href="#">Psme4</a>	<a href="#">Psme4.bSep08</a>	<a href="#">498433</a>	24979	1475	16	491	proteasome (prosome, macropain) activator subunit 4 (Psme4) alternative variant bSep08, mRNA.
<a href="#">Psme4</a>	<a href="#">Psme4.cSep08</a>	<a href="#">498433</a>	5088	635	7	156	proteasome (prosome, macropain) activator subunit 4 (Psme4) alternative variant cSep08, mRNA.
<a href="#">Psme4</a>	<a href="#">Psme4.dSep08</a>	<a href="#">498433</a>	5566	794	4	140	proteasome (prosome, macropain) activator subunit 4 (15.8 kD) (Psme4) alternative variant dSep08, mRNA.

<a href="#">Psmg1</a>	<a href="#">Psmg1.bSep08</a>	<a href="#">288236</a>	9455	1002	4	289	proteasome (prosome, macropain) assembly chaperone 1 (33.2 kD) (Psmg1) alternative variant bSep08, mRNA.
<a href="#">Psmg1</a>	<a href="#">Psmg1.cSep08</a>	<a href="#">288236</a>	6922	589	3	165	proteasome (prosome, macropain) assembly chaperone 1 (Psmg1) alternative variant cSep08, mRNA.
<a href="#">Psp</a>	<a href="#">Psp.bSep08</a>	<a href="#">50585</a>	2411	361	1	51	parotid secretory protein (Psp) alternative variant bSep08, mRNA.
<a href="#">Pspc1</a>	<a href="#">Pspc1.bSep08</a>	<a href="#">305910</a>	6844	1422	3	133	paraspeckle protein 1 (13.4 kD) (Pspc1) alternative variant bSep08, mRNA.
<a href="#">Pspc1</a>	<a href="#">Pspc1.cSep08</a>	<a href="#">305910</a>	29365	876	5	77	paraspeckle protein 1 (Pspc1) alternative variant cSep08, mRNA.
<a href="#">Psph</a>	<a href="#">Psph.bSep08</a>	<a href="#">304429</a>	16403	584	3	102	phosphoserine phosphatase (11.5 kD) (Psph) alternative variant bSep08, mRNA.
<a href="#">Psph</a>	<a href="#">Psph.cSep08</a>	<a href="#">304429</a>	17352	785	3	102	phosphoserine phosphatase (11.5 kD) (Psph) alternative variant cSep08, mRNA.
<a href="#">Psrc1</a>	<a href="#">Psrc1.bSep08</a>	<a href="#">691380</a>	2955	956	1	282	proline/serine-rich coiled-coil 1 (Psrc1) alternative variant bSep08, mRNA.
<a href="#">Psrc2</a>	<a href="#">Psrc2.aSep08</a>	<a href="#">314836</a>	24498	3400	20	859	proline serine-rich coiled-coil 2 (Psrc2) alternative variant aSep08, mRNA.
<a href="#">Psrc2</a>	<a href="#">Psrc2.bSep08</a>	<a href="#">314836</a>	5866	862	5	287	proline serine-rich coiled-coil 2 CRA b (Psrc2) alternative variant bSep08, mRNA.
<a href="#">Psrc2</a>	<a href="#">Psrc2.cSep08</a>	<a href="#">314836</a>	5594	713	4	165	proline serine-rich coiled-coil 2 CRA b (Psrc2) alternative variant cSep08, mRNA.
<a href="#">Psrc2</a>	<a href="#">Psrc2.dSep08</a>	<a href="#">314836</a>	2352	839	2	44	proline serine-rich coiled-coil 2 CRA d (5.2 kD) (Psrc2) alternative variant dSep08, mRNA.
<a href="#">Pstpip1</a>	<a href="#">Pstpip1.aSep08</a>	<a href="#">300732</a>	14824	1802	10	525	proline-serine-threonine phosphatase-interacting protein 1 (Pstpip1) alternative variant aSep08, mRNA.
<a href="#">Pstpip1</a>	<a href="#">Pstpip1.cSep08</a>	<a href="#">300732</a>	3088	586	2	65	proline-serine-threonine phosphatase-interacting protein 1 (Pstpip1) alternative variant cSep08, mRNA.
<a href="#">Pstpip2</a>	<a href="#">Pstpip2.aSep08</a>	<a href="#">307248</a>	20444	2745	10	216	proline-serine-threonine phosphatase-interacting protein 2 (Pstpip2) alternative variant aSep08, mRNA.
<a href="#">Ptar1</a>	<a href="#">Ptar1.bSep08</a>	<a href="#">286972</a>	9275	524	2	151	protein prenyltransferase alpha subunit repeat containing 1 (17.0 kD) (Ptar1) alternative variant bSep08, mRNA.
<a href="#">PTB.0</a>	<a href="#">PTB.0.aSep08</a>	<a href="#">498410</a>	13842	3335	5	218	tensin (PTB.0) alternative variant aSep08, mRNA.
<a href="#">PTB.0</a>	<a href="#">PTB.0.bSep08</a>	<a href="#">498410</a>	2024	740	1	111	tensin (PTB.0) alternative variant bSep08, mRNA.
<a href="#">PTB.1</a>	<a href="#">PTB.1.aSep08</a>		9338	915		305	tensin (PTB.1) mRNA.
<a href="#">Ptbp1</a>	<a href="#">Ptbp1.bSep08</a>	<a href="#">29497</a>	8286	3285	11	518	polypyrimidine tract binding protein 1 (55.2 kD) (Ptbp1) alternative variant bSep08, mRNA.
<a href="#">Ptbp1</a>	<a href="#">Ptbp1.cSep08</a>	<a href="#">29497</a>	3276	868	6	151	polypyrimidine tract binding protein 1 (16.7 kD) (Ptbp1) alternative variant cSep08, mRNA.
<a href="#">Ptbp1</a>	<a href="#">Ptbp1.dSep08</a>	<a href="#">29497</a>	2947	1049	5	148	polypyrimidine tract binding protein 1 (Ptbp1) alternative variant dSep08, mRNA.
<a href="#">Ptbp1</a>	<a href="#">Ptbp1.eSep08</a>	<a href="#">29497</a>	3571	856	7	140	polypyrimidine tract binding protein 1 (Ptbp1) alternative variant eSep08, mRNA.
<a href="#">Ptbp2</a>	<a href="#">Ptbp2.bSep08</a>	<a href="#">310820</a>	24101	2386	7	265	polypyrimidine tract binding protein 2 like (Ptbp2) alternative variant bSep08, mRNA.
<a href="#">Ptbp2</a>	<a href="#">Ptbp2.cSep08</a>	<a href="#">310820</a>	6277	1138	6	197	polypyrimidine tract binding protein 2 CRA b like (Ptbp2) alternative variant cSep08, mRNA.



<a href="#">Ptbp2</a>	<a href="#">Ptbp2.dSep08</a>	<a href="#">310820</a>	22475	726	6	150	polypyrimidine tract binding protein 2 like (16.8 kD) (Ptbp2) alternative variant dSep08, mRNA.
<a href="#">Ptbp2</a>	<a href="#">Ptbp2.eSep08</a>	<a href="#">310820</a>	22547	933	5	147	polypyrimidine tract binding protein 2 CRA e like (15.4 kD) (Ptbp2) alternative variant eSep08, mRNA.
<a href="#">Ptbp2</a>	<a href="#">Ptbp2.fSep08</a>	<a href="#">310820</a>	3676	628	2	139	polypyrimidine tract binding protein 2 CRA a like (Ptbp2) alternative variant fSep08, mRNA.
<a href="#">Ptcd1</a>	<a href="#">Ptcd1.aSep08</a>	<a href="#">304278</a>	17676	2946	9	686	pentatricopeptide repeat domain 1 (77.1 kD) (Ptcd1) alternative variant aSep08, complete mRNA.
<a href="#">Ptcd3</a>	<a href="#">Ptcd3.bSep08</a>	<a href="#">500199</a>	14610	1965	14	230	pentatricopeptide repeat domain 3 (26.4 kD) (Ptcd3) alternative variant bSep08, mRNA.
<a href="#">Ptch1</a>	<a href="#">Ptch1.bSep08</a>	<a href="#">89830</a>	29287	857	3	183	patched homolog 1 (Drosophila) (20.8 kD) (Ptch1) alternative variant bSep08, mRNA.
<a href="#">Ptch1</a>	<a href="#">Ptch1.cSep08</a>	<a href="#">89830</a>	18439	475	2	140	patched homolog 1 (Drosophila) (Ptch1) alternative variant cSep08, mRNA.
<a href="#">Ptch2</a>	<a href="#">Ptch2.bSep08</a>	<a href="#">366452</a>	5155	1433		41	patched homolog 2 (Ptch2) alternative variant bSep08, mRNA.
<a href="#">Ptchd2</a>	<a href="#">Ptchd2.bSep08</a>	<a href="#">313705</a>	12757	1802	6	600	putative protein of metazoan origin (Ptchd2) alternative variant bSep08, mRNA.
<a href="#">Ptchd2</a>	<a href="#">Ptchd2.cSep08</a>	<a href="#">313705</a>	1930	584	3	154	putative protein of metazoan origin (Ptchd2) alternative variant cSep08, mRNA.
<a href="#">Ptchd2</a>	<a href="#">Ptchd2.dSep08</a>	<a href="#">313705</a>	2403	418	4	139	putative protein, with a transmembrane domain, of metazoan origin (Ptchd2) alternative variant dSep08, mRNA.
<a href="#">Ptcrs</a>	<a href="#">Ptcrs.aSep08</a>	<a href="#">116462</a>	1240	602		93	pre T-cell antigen receptor alpha (Ptcrs) mRNA.
<a href="#">Ptdss1</a>	<a href="#">Ptdss1.bSep08</a>	<a href="#">314553</a>	22904	591	5	197	phosphatidylserine synthase 1 (Ptdss1) alternative variant bSep08, mRNA.
<a href="#">Ptdss2</a>	<a href="#">Ptdss2.bSep08</a>	<a href="#">293620</a>	22633	2006	1	168	phosphatidylserine synthase 2 (20.1 kD) (Ptdss2) alternative variant bSep08, complete mRNA.
<a href="#">Pten</a>	<a href="#">Pten.bSep08</a>	<a href="#">50557</a>	18749	423	2	61	phosphatase and tensin homolog (Pten) alternative variant bSep08, mRNA.
<a href="#">Pter</a>	<a href="#">Pter.bSep08</a>	<a href="#">63852</a>	41368	750	1	250	phosphotriesterase related (Pter) alternative variant bSep08, mRNA.
<a href="#">Pter</a>	<a href="#">Pter.cSep08</a>	<a href="#">63852</a>	41341	752	1	250	phosphotriesterase related (Pter) alternative variant cSep08, mRNA.
<a href="#">Ptgds</a>	<a href="#">Ptgds.bSep08</a>	<a href="#">25526</a>	2853	633	7	195	prostaglandin D2 synthase (brain) (Ptgds) alternative variant bSep08, mRNA.
<a href="#">Ptgds</a>	<a href="#">Ptgds.cSep08</a>	<a href="#">25526</a>	2530	758	6	126	prostaglandin D2 synthase (brain) (14.2 kD) (Ptgds) alternative variant cSep08, mRNA.
<a href="#">Ptgds2</a>	<a href="#">Ptgds2.bSep08</a>	<a href="#">58962</a>	11201	609	1	103	prostaglandin D2 synthase 2, hematopoietic (Ptgds2) alternative variant bSep08, mRNA.
<a href="#">Ptger1</a>	<a href="#">Ptger1.cSep08</a>	<a href="#">25637</a>	1716	763	2	219	prostaglandin E receptor 1 (subtype EP1) (Ptger1) alternative variant cSep08, mRNA.
<a href="#">Ptger3</a>	<a href="#">Ptger3.aSep08</a>	<a href="#">24929</a>	79248	2078	3	366	prostaglandin E receptor 3 (subtype EP3) (40.1 kD) (Ptger3) alternative variant aSep08, mRNA.
<a href="#">Ptger3</a>	<a href="#">Ptger3.bSep08</a>	<a href="#">24929</a>	39240	1188	1	365	prostaglandin E receptor 3 (subtype EP3) (39.6 kD) (Ptger3) alternative variant bSep08, mRNA.

<a href="#">Ptger4</a>	<a href="#">Ptger4.bSep08</a>	<a href="#">84023</a>	2373	755	4	122	putative mitochondrial protein (13.9 kD) (Ptger4) alternative variant bSep08, mRNA.
<a href="#">Ptges</a>	<a href="#">Ptges.bSep08</a>	<a href="#">59103</a>	895	493	2	38	prostaglandin E synthase (4.2 kD) (Ptges) alternative variant bSep08, mRNA.
<a href="#">Ptges2</a>	<a href="#">Ptges2.bSep08</a>	<a href="#">311865</a>	5200	1311	4	189	prostaglandin E synthase 2 (20.8 kD) (Ptges2) alternative variant bSep08, mRNA.
<a href="#">Ptgir</a>	<a href="#">Ptgir.bSep08</a>	<a href="#">292661</a>	2226	593	2	197	prostaglandin I receptor (IP) (Ptgir) alternative variant bSep08, mRNA.
<a href="#">Ptgs1</a>	<a href="#">Ptgs1.bSep08</a>	<a href="#">24693</a>	6223	324	3	92	prostaglandin-endoperoxide synthase 1 (Ptgs1) alternative variant bSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.bSep08</a>	<a href="#">56813</a>	23300	2250	12	363	parathyroid hormone receptor (40.3 kD) (Pthr1) alternative variant bSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.cSep08</a>	<a href="#">56813</a>	4107	2604	5	269	parathyroid hormone receptor (29.9 kD) (Pthr1) alternative variant cSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.dSep08</a>	<a href="#">56813</a>	17113	736	7	212	parathyroid hormone receptor (Pthr1) alternative variant dSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.eSep08</a>	<a href="#">56813</a>	19479	655	7	198	parathyroid hormone receptor (Pthr1) alternative variant eSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.fSep08</a>	<a href="#">56813</a>	1954	713	4	194	parathyroid hormone receptor (Pthr1) alternative variant fSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.gSep08</a>	<a href="#">56813</a>	12664	458	3	152	putative protein (Pthr1) alternative variant gSep08, mRNA.
<a href="#">Pthr1</a>	<a href="#">Pthr1.iSep08</a>	<a href="#">56813</a>	1994	617	3	75	parathyroid hormone receptor (Pthr1) alternative variant iSep08, mRNA.
<a href="#">Ptk2</a>	<a href="#">Ptk2.bSep08</a>	<a href="#">25614</a>	22578	1784	5	449	PTK2 protein tyrosine kinase 2 (Ptk2) alternative variant bSep08, mRNA.
<a href="#">Ptk2</a>	<a href="#">Ptk2.cSep08</a>	<a href="#">25614</a>	119169	1907	21	248	PTK2 protein tyrosine kinase 2 (Ptk2) alternative variant cSep08, mRNA.
<a href="#">Ptk2</a>	<a href="#">Ptk2.dSep08</a>	<a href="#">25614</a>	10824	398	5	113	PTK2 protein tyrosine kinase 2 (Ptk2) alternative variant dSep08, mRNA.
<a href="#">Ptk2</a>	<a href="#">Ptk2.fSep08</a>	<a href="#">25614</a>	8443	436	3	38	PTK2 protein tyrosine kinase 2 (Ptk2) alternative variant fSep08, mRNA.
<a href="#">Ptk7</a>	<a href="#">Ptk7.bSep08</a>	<a href="#">301242</a>	11957	1105	5	214	PTK7 protein tyrosine kinase 7 (Ptk7) alternative variant bSep08, mRNA.
<a href="#">Ptk7</a>	<a href="#">Ptk7.cSep08</a>	<a href="#">301242</a>	3601	459	4	152	PTK7 protein tyrosine kinase 7 (Ptk7) alternative variant cSep08, mRNA.
<a href="#">Ptma</a>	<a href="#">Ptma.bSep08</a>	<a href="#">29222</a>	3656	975	4	107	prothymosin alpha (11.7 kD) (Ptma) alternative variant bSep08, mRNA.
<a href="#">Ptma</a>	<a href="#">Ptma.cSep08</a>	<a href="#">29222</a>	3966	1041	5	96	prothymosin alpha (10.6 kD) (Ptma) alternative variant cSep08, mRNA.
<a href="#">Ptma</a>	<a href="#">Ptma.dSep08</a>	<a href="#">29222</a>	3835	1179	4	73	prothymosin alpha (8.0 kD) (Ptma) alternative variant dSep08, complete mRNA.
<a href="#">Ptma</a>	<a href="#">Ptma.eSep08</a>	<a href="#">29222</a>	3500	401	3	38	prothymosin alpha (4.3 kD) (Ptma) alternative variant eSep08, mRNA.
<a href="#">Ptn</a>	<a href="#">Ptn.aSep08</a>	<a href="#">24924</a>	50563	1294	5	180	pleiotrophin (Ptn) alternative variant aSep08, mRNA.
<a href="#">Ptn</a>	<a href="#">Ptn.bSep08</a>	<a href="#">24924</a>	80888	916	6	168	pleiotrophin (18.9 kD) (Ptn) alternative variant bSep08, complete mRNA.

<a href="#">Ptn</a>	<a href="#">Ptn.dSep08</a>	<a href="#">24924</a>	65113	1094	4	156	pleiotrophin (17.4 kD) (Ptn) alternative variant dSep08, mRNA.
<a href="#">Ptn</a>	<a href="#">Ptn.eSep08</a>	<a href="#">24924</a>	64607	621	4	152	pleiotrophin (Ptn) alternative variant eSep08, mRNA.
<a href="#">Ptn</a>	<a href="#">Ptn.fSep08</a>	<a href="#">24924</a>	40805	539	4	146	pleiotrophin (Ptn) alternative variant fSep08, mRNA.
<a href="#">Ptov1</a>	<a href="#">Ptov1.bSep08</a>	<a href="#">292888</a>	2745	1054	8	227	prostate tumor overexpressed 1 like (26.2 kD) (Ptov1) alternative variant bSep08, mRNA.
<a href="#">Ptov1</a>	<a href="#">Ptov1.cSep08</a>	<a href="#">292888</a>	4625	1296	9	224	prostate tumor overexpressed 1 like (Ptov1) alternative variant cSep08, mRNA.
<a href="#">Ptp4a1</a>	<a href="#">Ptp4a1.cSep08</a>	<a href="#">29463</a>	3232	581	2	46	putative protein (Ptp4a1) alternative variant cSep08, mRNA.
<a href="#">Ptp4a3</a>	<a href="#">Ptp4a3.bSep08</a>	<a href="#">362930</a>	2980	365	3	37	putative protein (Ptp4a3) alternative variant bSep08, mRNA.
<a href="#">Ptp4a3</a>	<a href="#">Ptp4a3.dSep08</a>	<a href="#">362930</a>	3109	416	2	39	putative protein (Ptp4a3) alternative variant dSep08, mRNA.
<a href="#">Ptpdc1</a>	<a href="#">Ptpdc1.bSep08</a>	<a href="#">291022</a>	7372	1521	4	129	protein tyrosine phosphatase (Ptpdc1) alternative variant bSep08, mRNA.
<a href="#">Ptpdc1</a>	<a href="#">Ptpdc1.cSep08</a>	<a href="#">291022</a>	777	368	2	122	putative protein of vertebrate origin (Ptpdc1) alternative variant cSep08, mRNA.
<a href="#">Ptpdc1</a>	<a href="#">Ptpdc1.dSep08</a>	<a href="#">291022</a>	34699	422	3	52	putative protein of vertebrate origin (Ptpdc1) alternative variant dSep08, mRNA.
<a href="#">Ptpla</a>	<a href="#">Ptpla.bSep08</a>	<a href="#">680115</a>	14016	683	5	196	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (Ptpla) alternative variant bSep08, mRNA.
<a href="#">Ptpla</a>	<a href="#">Ptpla.cSep08</a>	<a href="#">680115</a>	13952	896	3	175	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (20.3 kD) (Ptpla) alternative variant cSep08, mRNA.
<a href="#">Ptpla</a>	<a href="#">Ptpla.dSep08</a>	<a href="#">680115</a>	2242	373	1	51	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (Ptpla) alternative variant dSep08, mRNA.
<a href="#">Ptplb</a>	<a href="#">Ptplb.aSep08</a>	<a href="#">288058</a>	91751	994	7	254	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b (28.4 kD) (Ptplb) alternative variant aSep08, complete mRNA.
<a href="#">Ptplb</a>	<a href="#">Ptplb.bSep08</a>	<a href="#">288058</a>	18246	679	5	183	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b (Ptplb) alternative variant bSep08, mRNA.
<a href="#">Ptpmt1</a>	<a href="#">Ptpmt1.bSep08</a>	<a href="#">29390</a>	9074	580	3	123	protein tyrosine phosphatase, mitochondrial 1 (14.0 kD) (Ptpmt1) alternative variant bSep08, mRNA.
<a href="#">Ptpmt1</a>	<a href="#">Ptpmt1.cSep08</a>	<a href="#">29390</a>	9715	915	5	46	protein tyrosine phosphatase, mitochondrial 1 (Ptpmt1) alternative variant cSep08, mRNA.
<a href="#">Ptpn1</a>	<a href="#">Ptpn1.bSep08</a>	<a href="#">24697</a>	31509	581	2	67	protein tyrosine phosphatase, non-receptor type 1 (8.0 kD) (Ptpn1) alternative variant bSep08, mRNA.
<a href="#">Ptpn2</a>	<a href="#">Ptpn2.bSep08</a>	<a href="#">117063</a>	24843	1210	7	289	protein tyrosine phosphatase, non-receptor type 2 (Ptpn2) alternative variant bSep08, mRNA.
<a href="#">Ptpn3</a>	<a href="#">Ptpn3.aSep08</a>	<a href="#">362524</a>	29759	1157		384	protein tyrosine phosphatase, non-receptor type 3 (Ptpn3) mRNA.

<a href="#">Ptpn4</a>	<a href="#">Ptpn4.aSep08</a>	<a href="#">246116</a>	144629	1757	6	525	protein tyrosine phosphatase, non-receptor type 4 and hypothetical protein LOC680095 (Ptpn4) alternative variant aSep08, mRNA.
<a href="#">Ptpn4</a>	<a href="#">Ptpn4.aSep08</a>	<a href="#">680095</a>	144629	1757	6	525	protein tyrosine phosphatase, non-receptor type 4 and hypothetical protein LOC680095 (Ptpn4) alternative variant aSep08, mRNA.
<a href="#">Ptpn4</a>	<a href="#">Ptpn4.bSep08</a>	<a href="#">246116</a>	69125	494	2	104	protein tyrosine phosphatase, non-receptor type 4 and hypothetical protein LOC680095 (Ptpn4) alternative variant bSep08, mRNA.
<a href="#">Ptpn4</a>	<a href="#">Ptpn4.bSep08</a>	<a href="#">680095</a>	69125	494	2	104	protein tyrosine phosphatase, non-receptor type 4 and hypothetical protein LOC680095 (Ptpn4) alternative variant bSep08, mRNA.
<a href="#">Ptpn5</a>	<a href="#">Ptpn5.bSep08</a>	<a href="#">29644</a>	3408	700	1	140	protein tyrosine phosphatase, non-receptor type 5 (16.1 kD) (Ptpn5) alternative variant bSep08, mRNA.
<a href="#">Ptpn6</a>	<a href="#">Ptpn6.bSep08</a>	<a href="#">116689</a>	5139	846	6	231	protein tyrosine phosphatase, non-receptor type 6 (Ptpn6) alternative variant bSep08, mRNA.
<a href="#">Ptpn6</a>	<a href="#">Ptpn6.cSep08</a>	<a href="#">116689</a>	5197	739	6	176	protein tyrosine phosphatase, non-receptor type 6 (Ptpn6) alternative variant cSep08, mRNA.
<a href="#">Ptpn6</a>	<a href="#">Ptpn6.dSep08</a>	<a href="#">116689</a>	1351	808	4	141	protein tyrosine phosphatase, non-receptor type 6 (Ptpn6) alternative variant dSep08, mRNA.
<a href="#">Ptpn9</a>	<a href="#">Ptpn9.bSep08</a>	<a href="#">266611</a>	29990	1582	8	367	protein tyrosine phosphatase, non-receptor type 9 (42.1 kD) (Ptpn9) alternative variant bSep08, mRNA.
<a href="#">Ptpn9</a>	<a href="#">Ptpn9.cSep08</a>	<a href="#">266611</a>	12680	621	6	144	protein tyrosine phosphatase, non-receptor type 9 (Ptpn9) alternative variant cSep08, mRNA.
<a href="#">Ptpn9</a>	<a href="#">Ptpn9.dSep08</a>	<a href="#">266611</a>	14031	414	5	104	protein tyrosine phosphatase, non-receptor type 9 (Ptpn9) alternative variant dSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.aSep08</a>	<a href="#">498331</a>	44073	1897	7	570	tyrosine phosphatase (Ptpn13) alternative variant aSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.bSep08</a>	<a href="#">498331</a>	20412	1822	10	413	protein Tyrosine phosphatase (Ptpn13) alternative variant bSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.cSep08</a>	<a href="#">498331</a>	35649	1788	8	404	protein Tyrosine phosphatase (Ptpn13) alternative variant cSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.dSep08</a>	<a href="#">498331</a>	5992	924	6	307	protein Tyrosine phosphatase (Ptpn13) alternative variant dSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.eSep08</a>	<a href="#">498331</a>	9945	734	6	244	protein Tyrosine phosphatase (Ptpn13) alternative variant eSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.fSep08</a>	<a href="#">498331</a>	6730	318	3	83	protein Tyrosine phosphatase (Ptpn13) alternative variant fSep08, mRNA.
<a href="#">Ptpn13</a>	<a href="#">Ptpn13.gSep08</a>	<a href="#">498331</a>	1587	529	2	67	protein Tyrosine phosphatase (Ptpn13) alternative variant gSep08, mRNA.
<a href="#">Ptpn18</a>	<a href="#">Ptpn18.bSep08</a>	<a href="#">301333</a>	6156	1866	10	333	protein tyrosine phosphatase, non-receptor type 18 (36.5 kD) (Ptpn18) alternative variant bSep08, mRNA.
<a href="#">Ptpn18</a>	<a href="#">Ptpn18.cSep08</a>	<a href="#">301333</a>	2187	1184	5	133	protein tyrosine phosphatase, non-receptor type 18 (Ptpn18) alternative variant cSep08, mRNA.
<a href="#">Ptpn18</a>	<a href="#">Ptpn18.dSep08</a>	<a href="#">301333</a>	13535	697	7	71	protein tyrosine phosphatase, non-receptor type 18 (Ptpn18) alternative variant dSep08, mRNA.

<a href="#">Ptpn21</a>	<a href="#">Ptpn21.bSep08</a>	<a href="#">171070</a>	12605	3818	6	687	protein tyrosine phosphatase, non-receptor type 21 (Ptpn21) alternative variant bSep08, mRNA.
<a href="#">Ptpn21</a>	<a href="#">Ptpn21.cSep08</a>	<a href="#">171070</a>	2567	539	1	84	protein tyrosine phosphatase, non-receptor type 21 (Ptpn21) alternative variant cSep08, mRNA.
<a href="#">Ptpn23</a>	<a href="#">Ptpn23.bSep08</a>	<a href="#">117552</a>	754	388	1	57	protein tyrosine phosphatase, non-receptor type 23 (Ptpn23) alternative variant bSep08, mRNA.
<a href="#">Ptpra</a>	<a href="#">Ptpra.bSep08</a>	<a href="#">25167</a>	109886	1481	7	327	protein tyrosine phosphatase, receptor type, A (35.1 kD) (Ptpra) alternative variant bSep08, complete mRNA.
<a href="#">Ptpra</a>	<a href="#">Ptpra.cSep08</a>	<a href="#">25167</a>	27313	528	6	175	protein tyrosine phosphatase, receptor type, A (Ptpra) alternative variant cSep08, mRNA.
<a href="#">Ptpra</a>	<a href="#">Ptpra.dSep08</a>	<a href="#">25167</a>	19644	738	7	167	protein tyrosine phosphatase, receptor type, A (Ptpra) alternative variant dSep08, mRNA.
<a href="#">Ptprb</a>	<a href="#">Ptprb.bSep08</a>	<a href="#">314843</a>	8882	501	3	166	protein tyrosine phosphatase receptor type B CRA b (Ptprb) alternative variant bSep08, mRNA.
<a href="#">Ptprb</a>	<a href="#">Ptprb.bSep08</a>	<a href="#">688895</a>	8882	501	3	166	protein tyrosine phosphatase receptor type B CRA b (Ptprb) alternative variant bSep08, mRNA.
<a href="#">Ptprb</a>	<a href="#">Ptprb.cSep08</a>	<a href="#">314843</a>	791	377	2	77	protein tyrosine phosphatase receptor type B (Ptprb) alternative variant cSep08, mRNA.
<a href="#">Ptprb</a>	<a href="#">Ptprb.cSep08</a>	<a href="#">688895</a>	791	377	2	77	protein tyrosine phosphatase receptor type B (Ptprb) alternative variant cSep08, mRNA.
<a href="#">Ptprc</a>	<a href="#">Ptprc.fSep08</a>	<a href="#">24699</a>	58176	443	5	108	protein tyrosine phosphatase, receptor type, C (Ptprc) alternative variant fSep08, mRNA.
<a href="#">Ptprc</a>	<a href="#">Ptprc.gSep08</a>	<a href="#">24699</a>	7596	755	4	101	protein tyrosine phosphatase, receptor type, C (Ptprc) alternative variant gSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.bSep08</a>	<a href="#">25529</a>	32824	2145	8	559	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant bSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.dSep08</a>	<a href="#">25529</a>	3633	851	5	283	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant dSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.eSep08</a>	<a href="#">25529</a>	15535	1171	7	248	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant eSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.fSep08</a>	<a href="#">25529</a>	1705	618	3	198	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant fSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.gSep08</a>	<a href="#">25529</a>	9390	716	5	172	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant gSep08, mRNA.
<a href="#">Ptprd</a>	<a href="#">Ptprd.hSep08</a>	<a href="#">25529</a>	1580	881	3	124	protein tyrosine phosphatase, receptor type, D (Ptprd) alternative variant hSep08, mRNA.
<a href="#">Ptpre</a>	<a href="#">Ptpre.bSep08</a>	<a href="#">114767</a>	53644	678	6	95	protein tyrosine phosphatase, receptor type, E (10.5 kD) (Ptpre) alternative variant bSep08, mRNA.
<a href="#">Ptprf</a>	<a href="#">Ptprf.bSep08</a>	<a href="#">360406</a>	6443	711	6	236	protein tyrosine phosphatase, receptor type, F (Ptprf) alternative variant bSep08, mRNA.
<a href="#">Ptprf</a>	<a href="#">Ptprf.cSep08</a>	<a href="#">360406</a>	9315	563	5	187	protein tyrosine phosphatase, receptor type, F (Ptprf) alternative variant cSep08, mRNA.
<a href="#">Ptprf</a>	<a href="#">Ptprf.eSep08</a>	<a href="#">360406</a>	618	389	2	92	protein tyrosine phosphatase, receptor type, F (Ptprf) alternative variant eSep08, mRNA.
<a href="#">Ptprf</a>	<a href="#">Ptprf.fSep08</a>	<a href="#">360406</a>	4304	274	3	90	protein tyrosine phosphatase, receptor type, F (Ptprf) alternative variant fSep08, mRNA.

<a href="#">Ptprg</a>	<a href="#">Ptprg.bSep08</a>	<a href="#">171357</a>	55964	3014	17	668	protein tyrosine phosphatase, receptor type, G (Ptprg) alternative variant bSep08, mRNA.
<a href="#">Ptprg</a>	<a href="#">Ptprg.cSep08</a>	<a href="#">171357</a>	39170	417	3	138	protein tyrosine phosphatase, receptor type, G (Ptprg) alternative variant cSep08, mRNA.
<a href="#">Ptprh</a>	<a href="#">Ptprh.aSep08</a>	<a href="#">171125</a>	29203	1714	7	571	protein tyrosine phosphatase, receptor type, H (Ptprh) alternative variant aSep08, mRNA.
<a href="#">Ptprh</a>	<a href="#">Ptprh.bSep08</a>	<a href="#">171125</a>	10708	391	3	130	protein tyrosine phosphatase, receptor type, H (Ptprh) alternative variant bSep08, mRNA.
<a href="#">Ptprh</a>	<a href="#">Ptprh.cSep08</a>	<a href="#">171125</a>	2392	291	2	96	protein tyrosine phosphatase, receptor type, H (Ptprh) alternative variant cSep08, mRNA.
<a href="#">Ptprj</a>	<a href="#">Ptprj.aSep08</a>	<a href="#">29645</a>	111816	1795		598	protein tyrosine phosphatase, receptor type, J (Ptprj) alternative variant aSep08, mRNA.
<a href="#">Ptprj</a>	<a href="#">Ptprj.bSep08</a>	<a href="#">29645</a>	3240	346		115	protein tyrosine phosphatase, receptor type, J (Ptprj) alternative variant bSep08, mRNA.
<a href="#">Ptprk</a>	<a href="#">Ptprk.bSep08</a>	<a href="#">360302</a>	299803	1795	1	597	protein tyrosine phosphatase, receptor type, K, extracellular region (Ptprk) alternative variant bSep08, mRNA.
<a href="#">Ptprm</a>	<a href="#">Ptprm.aSep08</a>	<a href="#">29616</a>	307975	3967	24	1059	protein tyrosine phosphatase, receptor type, M (Ptprm) alternative variant aSep08, mRNA.
<a href="#">Ptprm</a>	<a href="#">Ptprm.bSep08</a>	<a href="#">29616</a>	20555	792	6	227	protein tyrosine phosphatase, receptor type, M (Ptprm) alternative variant bSep08, mRNA.
<a href="#">Ptprm</a>	<a href="#">Ptprm.cSep08</a>	<a href="#">29616</a>	142898	704	7	187	protein tyrosine phosphatase, receptor type, M (Ptprm) alternative variant cSep08, mRNA.
<a href="#">Ptprm</a>	<a href="#">Ptprm.dSep08</a>	<a href="#">29616</a>	8831	350	4	116	protein tyrosine phosphatase, receptor type, M (Ptprm) alternative variant dSep08, mRNA.
<a href="#">Ptprm</a>	<a href="#">Ptprm.gSep08</a>	<a href="#">29616</a>	1428	309	2	56	protein tyrosine phosphatase, receptor type, M (Ptprm) alternative variant gSep08, mRNA.
<a href="#">Ptprn</a>	<a href="#">Ptprn.aSep08</a>	<a href="#">116660</a>	15082	3372	23	952	protein tyrosine phosphatase, receptor type, N (Ptprn) alternative variant aSep08, mRNA.
<a href="#">Ptprn</a>	<a href="#">Ptprn.bSep08</a>	<a href="#">116660</a>	2262	587	6	180	protein tyrosine phosphatase, receptor type, N (Ptprn) alternative variant bSep08, mRNA.
<a href="#">Ptprn</a>	<a href="#">Ptprn.cSep08</a>	<a href="#">116660</a>	9093	1831	7	172	protein tyrosine phosphatase, receptor type, N (Ptprn) alternative variant cSep08, mRNA.
<a href="#">Ptprn</a>	<a href="#">Ptprn.dSep08</a>	<a href="#">116660</a>	465	383	2	127	protein tyrosine phosphatase, receptor type, N (Ptprn) alternative variant dSep08, mRNA.
<a href="#">Ptprn</a>	<a href="#">Ptprn.eSep08</a>	<a href="#">116660</a>	1141	428	5	60	protein tyrosine phosphatase, receptor type, N (Ptprn) alternative variant eSep08, mRNA.
<a href="#">Ptpro</a>	<a href="#">Ptpro.bSep08</a>	<a href="#">50677</a>	34514	809	1	217	protein tyrosine phosphatase, receptor type, O (Ptpro) alternative variant bSep08, mRNA.
<a href="#">Ptpru</a>	<a href="#">Ptpru.aSep08</a>	<a href="#">116680</a>	8623	1113	8	370	protein tyrosine phosphatase, receptor type, U (Ptpru) alternative variant aSep08, mRNA.
<a href="#">Ptpru</a>	<a href="#">Ptpru.bSep08</a>	<a href="#">116680</a>	3609	2339	3	175	protein tyrosine phosphatase, receptor type, U (20.0 kD) (Ptpru) alternative variant bSep08, mRNA.
<a href="#">Ptprv</a>	<a href="#">Ptprv.aSep08</a>	<a href="#">64576</a>	2595	786		261	protein tyrosine phosphatase, receptor type, V (Ptprv) mRNA.
<a href="#">Ptprz1</a>	<a href="#">Ptprz1.cSep08</a>	<a href="#">25613</a>	11078	625	6	207	protein tyrosine phosphatase, receptor-type, Z polypeptide 1 (Ptprz1) alternative variant cSep08, mRNA.

<a href="#">Pts</a>	<a href="#">Pts.bSep08</a>	<a href="#">29498</a>	6581	724		91	6-pyruvoyl-tetrahydropterin synthase (10.4 kD) (Pts) alternative variant bSep08, mRNA.
<a href="#">Pttg1</a>	<a href="#">Pttg1.bSep08</a>	<a href="#">64193</a>	5671	1072	2	199	pituitary tumor-transforming 1 (21.6 kD) (Pttg1) alternative variant bSep08, mRNA.
<a href="#">Pttg1</a>	<a href="#">Pttg1.cSep08</a>	<a href="#">64193</a>	5147	715	2	108	pituitary tumor-transforming 1 (Pttg1) alternative variant cSep08, mRNA.
<a href="#">Pttg1ip</a>	<a href="#">Pttg1ip.bSep08</a>	<a href="#">365548</a>	12542	746	6	162	pituitary tumor-transforming 1 interacting protein (Pttg1ip) alternative variant bSep08, mRNA.
<a href="#">puby</a>	<a href="#">puby.aSep08</a>		2858	1967		160	mediator of RNA polymerase II transcription homolog (19.1 kD) (puby) mRNA.
<a href="#">puchy</a>	<a href="#">puchy.aSep08</a>		2132	881		83	putative protein (9.1 kD) (puchy) mRNA.
<a href="#">pudar</a>	<a href="#">pudar.aSep08</a>		2288	757		77	putative mitochondrial protein (9.2 kD) (pudar) mRNA.
<a href="#">pufer</a>	<a href="#">pufer.aSep08</a>		1075	307		85	putative protein (pufer) mRNA.
<a href="#">puflo</a>	<a href="#">puflo.aSep08</a>		34318	842	9	238	CRA a (puflo) alternative variant aSep08, mRNA.
<a href="#">puflo</a>	<a href="#">puflo.cSep08</a>		16453	282	2	74	CRA a like (puflo) alternative variant cSep08, mRNA.
<a href="#">puflu</a>	<a href="#">puflu.aSep08</a>		3083	400		42	putative protein (4.7 kD) (puflu) mRNA.
<a href="#">pugar</a>	<a href="#">pugar.aSep08</a>		2021	384		127	stabilin 1 (pugar) mRNA.
<a href="#">puja</a>	<a href="#">puja.aSep08</a>		7631	696		105	putative protein (12.0 kD) (puja) mRNA.
<a href="#">pujey</a>	<a href="#">pujey.aSep08</a>		764	509		35	putative protein (3.9 kD) (pujey) mRNA.
<a href="#">pukee</a>	<a href="#">pukee.bSep08</a>		1225	440		146	centrosomal protein 350kDa CRA a (pukee) alternative variant bSep08, mRNA.
<a href="#">pukler</a>	<a href="#">pukler.aSep08</a>		2315	739		45	putative protein (5.2 kD) (pukler) mRNA.
<a href="#">puloy</a>	<a href="#">puloy.aSep08</a>		1594	1421	2	80	putative mitochondrial protein (9.0 kD) (puloy) alternative variant aSep08, mRNA.
<a href="#">Pum2</a>	<a href="#">Pum2.bSep08</a>	<a href="#">298874</a>	38452	1972	13	656	pumilio 2 (Drosophila) (Pum2) alternative variant bSep08, mRNA.
<a href="#">Pum2</a>	<a href="#">Pum2.cSep08</a>	<a href="#">298874</a>	35808	684	5	165	pumilio 2 (Drosophila) (Pum2) alternative variant cSep08, mRNA.
<a href="#">Pum2</a>	<a href="#">Pum2.dSep08</a>	<a href="#">298874</a>	11361	310	2	103	pumilio 2 (Drosophila) (Pum2) alternative variant dSep08, mRNA.
<a href="#">pumee</a>	<a href="#">pumee.aSep08</a>		8755	811		97	putative protein (10.6 kD) (pumee) mRNA.
<a href="#">pumer</a>	<a href="#">pumer.aSep08</a>		3895	1346	4	448	ubiquitin-conjugating enzyme E2O CRA d (pumer) alternative variant aSep08, mRNA.
<a href="#">Punc</a>	<a href="#">Punc.aSep08</a>	<a href="#">315759</a>	47801	4756	12	923	putative neuronal cell adhesion molecule (Punc) alternative variant aSep08, mRNA.
<a href="#">Punc</a>	<a href="#">Punc.bSep08</a>	<a href="#">315759</a>	8239	2127	10	528	putative neuronal cell adhesion molecule (Punc) alternative variant bSep08, mRNA.
<a href="#">Punc</a>	<a href="#">Punc.dSep08</a>	<a href="#">315759</a>	2823	2093	2	108	putative neuronal cell adhesion molecule (Punc) alternative variant dSep08, mRNA.
<a href="#">punoy</a>	<a href="#">punoy.aSep08</a>		3311	555		37	putative protein (punoy) mRNA.
<a href="#">pupor</a>	<a href="#">pupor.aSep08</a>		3132	906	1	302	probable E3 ubiquitin-protein ligase herc1 (pupor) alternative variant aSep08, mRNA.
<a href="#">pupor</a>	<a href="#">pupor.bSep08</a>		2850	289	1	96	probable E3 ubiquitin-protein ligase herc1 like (pupor) alternative variant bSep08, mRNA.
<a href="#">Purg</a>	<a href="#">Purg.aSep08</a>	<a href="#">361162</a>	29855	1287		162	purine-rich element binding protein G (Purg) mRNA.

<a href="#">Pus1</a>	<a href="#">Pus1.bSep08</a>	<a href="#">304567</a>	7910	1032	5	336	pseudouridine synthase 1 (Pus1) alternative variant bSep08, mRNA.
<a href="#">Pus1</a>	<a href="#">Pus1.cSep08</a>	<a href="#">304567</a>	7586	807	4	268	pseudouridine synthase 1 (Pus1) alternative variant cSep08, mRNA.
<a href="#">Pus1</a>	<a href="#">Pus1.dSep08</a>	<a href="#">304567</a>	6975	742	4	247	pseudouridine synthase 1 (Pus1) alternative variant dSep08, mRNA.
<a href="#">Pus3</a>	<a href="#">Pus3.bSep08</a>	<a href="#">315554</a>	709	624	1	115	pseudouridine synthase 3 (Pus3) alternative variant bSep08, mRNA.
<a href="#">Pus7</a>	<a href="#">Pus7.aSep08</a>	<a href="#">296751</a>	14182	2401	9	341	pseudouridylate synthase 7 homolog CRA a (38.9 kD) (Pus7) alternative variant aSep08, mRNA.
<a href="#">Pus7</a>	<a href="#">Pus7.bSep08</a>	<a href="#">296751</a>	8154	650	3	200	putative protein (Pus7) alternative variant bSep08, mRNA.
<a href="#">Pus7</a>	<a href="#">Pus7.cSep08</a>	<a href="#">296751</a>	15774	757	8	182	pseudouridylate synthase 7 homolog CRA a (Pus7) alternative variant cSep08, mRNA.
<a href="#">Pus7</a>	<a href="#">Pus7.dSep08</a>	<a href="#">296751</a>	934	557	2	44	putative protein (Pus7) alternative variant dSep08, mRNA.
<a href="#">pusa</a>	<a href="#">pusa.aSep08</a>		1203	1045			
<a href="#">pushee</a>	<a href="#">pushee.aSep08</a>		31039	834		277	neurobeachin (pushee) mRNA.
<a href="#">Pusl1</a>	<a href="#">Pusl1.aSep08</a>	<a href="#">362681</a>	5102	3340	8	291	pseudouridylate synthase-like 1 (32.1 kD) (Pusl1) alternative variant aSep08, complete mRNA.
<a href="#">Pusl1</a>	<a href="#">Pusl1.bSep08</a>	<a href="#">362681</a>	1357	760	5	211	pseudouridylate synthase-like 1 (Pusl1) alternative variant bSep08, mRNA.
<a href="#">Pusl1</a>	<a href="#">Pusl1.cSep08</a>	<a href="#">362681</a>	2223	806	6	167	pseudouridylate synthase-like 1 (18.4 kD) (Pusl1) alternative variant cSep08, mRNA.
<a href="#">Pusl1</a>	<a href="#">Pusl1.dSep08</a>	<a href="#">362681</a>	1876	1316	5	129	pseudouridylate synthase-like 1 (13.9 kD) (Pusl1) alternative variant dSep08, mRNA.
<a href="#">Pusl1</a>	<a href="#">Pusl1.eSep08</a>	<a href="#">362681</a>	977	748	2	114	pseudouridylate synthase-like 1 (Pusl1) alternative variant eSep08, mRNA.
<a href="#">putu</a>	<a href="#">putu.aSep08</a>		2234	430		43	putative protein (5.0 kD) (putu) mRNA.
<a href="#">puvar</a>	<a href="#">puvar.aSep08</a>		1464	811		27	putative protein (2.9 kD) (puvar) mRNA.
<a href="#">puwey</a>	<a href="#">puwey.aSep08</a>		1691	813		44	putative protein (puwey) mRNA.
<a href="#">Pvalb</a>	<a href="#">Pvalb.aSep08</a>	<a href="#">25269</a>	14998	965	3	110	parvalbumin (11.9 kD) (Pvalb) alternative variant aSep08, complete mRNA.
<a href="#">Pvrl1</a>	<a href="#">Pvrl1.aSep08</a>	<a href="#">192183</a>	1345	571		190	poliovirus receptor-related 1 (Pvrl1) mRNA.
<a href="#">Pvrl3</a>	<a href="#">Pvrl3.bSep08</a>	<a href="#">288124</a>	55420	1364	7	377	poliovirus receptor-related 3 (41.6 kD) (Pvrl3) alternative variant bSep08, mRNA.
<a href="#">Pvrl3</a>	<a href="#">Pvrl3.cSep08</a>	<a href="#">288124</a>	8890	3308	2	216	poliovirus receptor-related 3 (Pvrl3) alternative variant cSep08, mRNA.
<a href="#">Pvrl3</a>	<a href="#">Pvrl3.dSep08</a>	<a href="#">288124</a>	27679	761	2	105	poliovirus receptor-related 3 (Pvrl3) alternative variant dSep08, mRNA.
<a href="#">Pvrl3</a>	<a href="#">Pvrl3.eSep08</a>	<a href="#">288124</a>	11904	678	3	93	poliovirus receptor-related 3 (10.3 kD) (Pvrl3) alternative variant eSep08, mRNA.
<a href="#">Pvrl4</a>	<a href="#">Pvrl4.aSep08</a>	<a href="#">498281</a>	3404	1787	2	517	poliovirus receptor-related 4 (Pvrl4) alternative variant aSep08, mRNA.
<a href="#">Pvrl4</a>	<a href="#">Pvrl4.cSep08</a>	<a href="#">498281</a>	3227	779	2	208	poliovirus receptor-related 4 (Pvrl4) alternative variant cSep08, mRNA.
<a href="#">Pwp1</a>	<a href="#">Pwp1.aSep08</a>	<a href="#">362856</a>	14648	1427	13	456	PWP1 homolog ( <i>S. cerevisiae</i> ) (Pwp1) alternative variant aSep08, mRNA.



<a href="#">Pwp1</a>	<a href="#">Pwp1.bSep08</a>	<a href="#">362856</a>	4045	1325	5	184	PWP1 homolog ( <i>S. cerevisiae</i> ) (20.5 kD) (Pwp1) alternative variant bSep08, mRNA.
<a href="#">Pwp2</a>	<a href="#">Pwp2.aSep08</a>	<a href="#">690297</a>	6771	1767	10	470	PWP2 periodic tryptophan protein homolog (yeast) (Pwp2) alternative variant aSep08, mRNA.
<a href="#">Pwp2</a>	<a href="#">Pwp2.cSep08</a>	<a href="#">690297</a>	1117	1001	2	76	PWP2 periodic tryptophan protein homolog (yeast) (Pwp2) alternative variant cSep08, mRNA.
<a href="#">Pwwp2a</a>	<a href="#">Pwwp2a.bSep08</a>	<a href="#">303060</a>	14230	817	2	97	putative protein of vertebrate origin (Pwwp2a) alternative variant bSep08, mRNA.
<a href="#">Pwwp2b</a>	<a href="#">Pwwp2b.bSep08</a>	<a href="#">361671</a>	11056	1177	2	147	PWWP (Pwwp2b) alternative variant bSep08, mRNA.
<a href="#">PX.0</a>	<a href="#">PX.0.aSep08</a>		42870	986	1	303	phox-like and MIT containing protein (33.8 kD) (PX.0) alternative variant aSep08, mRNA.
<a href="#">PX.0</a>	<a href="#">PX.0.bSep08</a>		26609	658	1	98	putative protein of metazoan origin (PX.0) alternative variant bSep08, mRNA.
<a href="#">PX.1</a>	<a href="#">PX.1.aSep08</a>		7816	415		138	protein 3 CRA d (PX.1) mRNA.
<a href="#">PX.2</a>	<a href="#">PX.2.aSep08</a>		9200	2230	1	278	sorting nexin 21 (PX.2) alternative variant aSep08, mRNA.
<a href="#">PX.2</a>	<a href="#">PX.2.bSep08</a>		518	351	1	116	sorting nexin 21 (PX.2) alternative variant bSep08, mRNA.
<a href="#">PX.3</a>	<a href="#">PX.3.aSep08</a>		21888	820		273	phospholipase D1 (PX.3) mRNA.
<a href="#">Pxdn</a>	<a href="#">Pxdn.aSep08</a>	<a href="#">554172</a>	15295	4278		192	peroxidasin homolog ( <i>Drosophila</i> ) (21.6 kD) (Pxdn) mRNA.
<a href="#">Pxm</a>	<a href="#">Pxm.dSep08</a>	<a href="#">306203</a>	3208	646	2	46	putative protein of metazoan origin (Pxm) alternative variant dSep08, mRNA.
<a href="#">Pxmp2</a>	<a href="#">Pxmp2.bSep08</a>	<a href="#">29533</a>	2615	624	2	74	peroxisomal membrane protein 2 (Pxmp2) alternative variant bSep08, mRNA.
<a href="#">Pxmp3</a>	<a href="#">Pxmp3.aSep08</a>	<a href="#">29534</a>	15954	931	2	310	peroxisomal membrane protein 3 (Pxmp3) alternative variant aSep08, mRNA.
<a href="#">Pxmp3</a>	<a href="#">Pxmp3.cSep08</a>	<a href="#">29534</a>	16663	1750	3	305	peroxisomal membrane protein 3 (34.8 kD) (Pxmp3) alternative variant cSep08, complete mRNA.
<a href="#">Pxmp3</a>	<a href="#">Pxmp3.dSep08</a>	<a href="#">29534</a>	15956	750	3	209	peroxisomal membrane protein 3 (Pxmp3) alternative variant dSep08, mRNA.
<a href="#">Pxn</a>	<a href="#">Pxn.bSep08</a>	<a href="#">360820</a>	38036	767	2	207	paxillin (Pxn) alternative variant bSep08, mRNA.
<a href="#">pyby</a>	<a href="#">pyby.aSep08</a>		2675	622	5	207	mediator of RNA polymerase II transcription homolog (pyby) alternative variant aSep08, mRNA.
<a href="#">pychy</a>	<a href="#">pychy.aSep08</a>		68454	367	1	64	putative protein (pychy) alternative variant aSep08, mRNA.
<a href="#">pychy</a>	<a href="#">pychy.bSep08</a>		858	395	1	49	putative protein (pychy) alternative variant bSep08, mRNA.
<a href="#">Pycr1</a>	<a href="#">Pycr1.bSep08</a>	<a href="#">287877</a>	948	337	1	93	pyrroline-5-carboxylate reductase 1 (Pycr1) alternative variant bSep08, mRNA.
<a href="#">Pycr2</a>	<a href="#">Pycr2.bSep08</a>	<a href="#">364064</a>	2284	386	3	84	pyrroline-5-carboxylate reductase family, member 2 (8.7 kD) (Pycr2) alternative variant bSep08, mRNA.
<a href="#">Pycr2</a>	<a href="#">Pycr2.cSep08</a>	<a href="#">364064</a>	1682	373	2	75	pyrroline-5-carboxylate reductase family, member 2 (Pycr2) alternative variant cSep08, mRNA.
<a href="#">Pycl</a>	<a href="#">Pycl.bSep08</a>	<a href="#">300035</a>	4547	789	1	260	pyrroline-5-carboxylate reductase-like (Pycl) alternative variant bSep08, mRNA.
<a href="#">Pycs</a>	<a href="#">Pycs.bSep08</a>	<a href="#">361755</a>	20258	1357	11	413	pyrroline-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase) (Pycs) alternative variant bSep08, mRNA.

<a href="#">Pycs</a>	<a href="#">Pycs.dSep08</a>	<a href="#">361755</a>	1946	677	3	52	pyrroline-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase) (5.6 kD) (Pycs) alternative variant dSep08, mRNA.
<a href="#">pydar</a>	<a href="#">pydar.aSep08</a>		5910	735		25	putative protein (2.9 kD) (pydar) mRNA.
<a href="#">pyfer</a>	<a href="#">pyfer.aSep08</a>		607	375		64	putative protein (pyfer) mRNA.
<a href="#">pyflo</a>	<a href="#">pyflo.aSep08</a>		6349	382		101	CRA b like (pyflo) mRNA.
<a href="#">pyflu</a>	<a href="#">pyflu.aSep08</a>		13054	1029	9	255	small nuclear ribonucleoprotein polypeptide a' (28.3 kD) (pyflu) alternative variant aSep08, mRNA.
<a href="#">pyflu</a>	<a href="#">pyflu.bSep08</a>		602	388	2	74	CRA c (pyflu) alternative variant bSep08, mRNA.
<a href="#">pyflu</a>	<a href="#">pyflu.cSep08</a>		9595	611	6	62	small nuclear ribonucleoprotein polypeptide A' (pyflu) alternative variant cSep08, mRNA.
<a href="#">pyflu</a>	<a href="#">pyflu.dSep08</a>		2905	532	2	31	putative protein (pyflu) alternative variant dSep08, mRNA.
<a href="#">pygar</a>	<a href="#">pygar.aSep08</a>		2370	753		250	stabilin 1 (pygar) mRNA.
<a href="#">Pygb</a>	<a href="#">Pygb.aSep08</a>	<a href="#">25739</a>	51705	4191	21	846	brain glycogen phosphorylase (96.8 kD) (Pygb) alternative variant aSep08, mRNA.
<a href="#">Pygl</a>	<a href="#">Pygl.aSep08</a>	<a href="#">64035</a>	43713	3416		910	liver glycogen phosphorylase (Pygl) mRNA.
<a href="#">Pygm</a>	<a href="#">Pygm.bSep08</a>	<a href="#">24701</a>	949	378	4	103	muscle glycogen phosphorylase (Pygm) alternative variant bSep08, mRNA.
<a href="#">Pygo2</a>	<a href="#">Pygo2.bSep08</a>	<a href="#">295251</a>	2865	757	2	252	pygopus 2 (Pygo2) alternative variant bSep08, mRNA.
<a href="#">Pygo2</a>	<a href="#">Pygo2.cSep08</a>	<a href="#">295251</a>	2853	1418	1	245	pygopus 2 (Pygo2) alternative variant cSep08, mRNA.
<a href="#">pyja</a>	<a href="#">pyja.aSep08</a>		5667	509		64	putative protein (7.2 kD) (pyja) mRNA.
<a href="#">pyjey</a>	<a href="#">pyjey.aSep08</a>		1174	670		70	putative protein (7.9 kD) (pyjey) mRNA.
<a href="#">pykee</a>	<a href="#">pykee.aSep08</a>		10720	865		150	centrosomal protein 350kDa CRA a (pykee) mRNA.
<a href="#">pykler</a>	<a href="#">pykler.aSep08</a>		25794	800		51	putative protein (5.6 kD) (pykler) mRNA.
<a href="#">pyloy</a>	<a href="#">pyloy.aSep08</a>		1799	226		60	putative protein (pyloy) mRNA.
<a href="#">pymee</a>	<a href="#">pymee.aSep08</a>		7947	592		196	serine threonine kinase 10 (pymee) mRNA.
<a href="#">pymer</a>	<a href="#">pymer.aSep08</a>		826	687		133	ubiquitin-conjugating enzyme E2O CRA c (pymer) mRNA.
<a href="#">pynoy</a>	<a href="#">pynoy.aSep08</a>		25017	303		100	tensin (pynoy) mRNA.
<a href="#">pypor</a>	<a href="#">pypor.aSep08</a>		8115	640		213	probable e3 ubiquitin-protein ligase herc1 (pypor) mRNA.
<a href="#">Pyroxd1</a>	<a href="#">Pyroxd1.bSep08</a>	<a href="#">297708</a>	1454	817	2	140	pyridine nucleotide-disulphide oxidoreductase domain 1 (Pyroxd1) alternative variant bSep08, mRNA.
<a href="#">Pyroxd1</a>	<a href="#">Pyroxd1.cSep08</a>	<a href="#">297708</a>	1676	919	2	117	pyridine nucleotide-disulphide oxidoreductase domain 1 (13.2 kD) (Pyroxd1) alternative variant cSep08, mRNA.
<a href="#">Pyroxd1</a>	<a href="#">Pyroxd1.dSep08</a>	<a href="#">297708</a>	1811	343	2	49	pyridine nucleotide-disulphide oxidoreductase domain 1 (Pyroxd1) alternative variant dSep08, mRNA.
<a href="#">pysa</a>	<a href="#">pysa.aSep08</a>		21599	505		74	putative protein (pysa) mRNA.
<a href="#">pyshee</a>	<a href="#">pyshee.aSep08</a>		21705	605		201	neurobeachin CRA e (pyshee) mRNA.
<a href="#">pytu</a>	<a href="#">pytu.aSep08</a>		2123	732		100	putative protein (pytu) mRNA.
<a href="#">pyvar</a>	<a href="#">pyvar.aSep08</a>		3126	858		63	putative protein (pyvar) mRNA.
<a href="#">pywey</a>	<a href="#">pywey.bSep08</a>		626	446	2	59	putative protein (pywey) alternative variant bSep08, mRNA.
<a href="#">Pzp</a>	<a href="#">Pzp.bSep08</a>	<a href="#">252922</a>	4832	1204	8	265	zone protein (Pzp) alternative variant bSep08, mRNA.
<a href="#">Pzp</a>	<a href="#">Pzp.cSep08</a>	<a href="#">252922</a>	2137	771	4	221	zone protein (Pzp) alternative variant cSep08, mRNA.

<a href="#">Pzp</a>	<a href="#">Pzp.eSep08</a>	<a href="#">252922</a>	4595	253	3	84	pregnancy-zone protein like (Pzp) alternative variant eSep08, mRNA.
<a href="#">Pzp</a>	<a href="#">Pzp.fSep08</a>	<a href="#">252922</a>	2087	1283	2	64	zone protein (7.2 kD) (Pzp) alternative variant fSep08, mRNA.
<a href="#">Qars</a>	<a href="#">Qars.bSep08</a>	<a href="#">290868</a>	2249	639	7	212	glutaminyl-tRNA synthetase (Qars) alternative variant bSep08, mRNA.
<a href="#">Qars</a>	<a href="#">Qars.cSep08</a>	<a href="#">290868</a>	1756	691	6	175	glutaminyl-tRNA synthetase (19.7 kD) (Qars) alternative variant cSep08, mRNA.
<a href="#">Qars</a>	<a href="#">Qars.dSep08</a>	<a href="#">290868</a>	761	396	3	131	glutaminyl-tRNA synthetase (Qars) alternative variant dSep08, mRNA.
<a href="#">Qars</a>	<a href="#">Qars.eSep08</a>	<a href="#">290868</a>	1847	672	7	117	glutaminyl-tRNA synthetase (Qars) alternative variant eSep08, mRNA.
<a href="#">Qki</a>	<a href="#">Qki.bSep08</a>	<a href="#">499022</a>	61970	3747	6	229	quaking homolog, KH domain RNA binding (mouse) (Qki) alternative variant bSep08, mRNA.
<a href="#">Qki</a>	<a href="#">Qki.cSep08</a>	<a href="#">499022</a>	59043	796	6	214	quaking homolog, KH domain RNA binding (mouse) (Qki) alternative variant cSep08, mRNA.
<a href="#">QLQ.0</a>	<a href="#">QLQ.0.aSep08</a>		25426	714		214	SWI snf-related matrix-associated actin-dependent regulator of chromatin a2 (QLQ.0) mRNA.
<a href="#">Qpctl</a>	<a href="#">Qpctl.bSep08</a>	<a href="#">292687</a>	8335	1349	6	221	glutaminyl-peptide cyclotransferase-like (24.2 kD) (Qpctl) alternative variant bSep08, complete mRNA.
<a href="#">Qpctl</a>	<a href="#">Qpctl.cSep08</a>	<a href="#">292687</a>	2537	661	2	127	glutaminyl-peptide cyclotransferase-like (Qpctl) alternative variant cSep08, mRNA.
<a href="#">Qprt</a>	<a href="#">Qprt.aSep08</a>	<a href="#">293504</a>	15504	1218	3	305	quinolinate phosphoribosyltransferase (Qprt) alternative variant aSep08, mRNA.
<a href="#">Qprt</a>	<a href="#">Qprt.cSep08</a>	<a href="#">293504</a>	14414	635	3	131	quinolinate phosphoribosyltransferase (Qprt) alternative variant cSep08, mRNA.
<a href="#">Qprt</a>	<a href="#">Qprt.dSep08</a>	<a href="#">293504</a>	4943	607	3	27	quinolinate phosphoribosyltransferase (3.0 kD) (Qprt) alternative variant dSep08, complete mRNA.
<a href="#">Qrich2</a>	<a href="#">Qrich2.aSep08</a>	<a href="#">501747</a>	1816	573	4	157	glutamine rich 2 (Qrich2) alternative variant aSep08, mRNA.
<a href="#">Qrich2</a>	<a href="#">Qrich2.bSep08</a>	<a href="#">501747</a>	4588	716	6	156	glutamine rich 2 (17.2 kD) (Qrich2) alternative variant bSep08, mRNA.
<a href="#">Qrs1</a>	<a href="#">Qrs1.bSep08</a>	<a href="#">309911</a>	14359	1040	6	183	glutaminyl-tRNA synthase (glutamine-hydrolyzing)-like 1 (19.8 kD) (Qrs1) alternative variant bSep08, complete mRNA.
<a href="#">Qrs1</a>	<a href="#">Qrs1.cSep08</a>	<a href="#">309911</a>	2738	448	3	142	glutaminyl-tRNA synthase (glutamine-hydrolyzing)-like 1 (Qrs1) alternative variant cSep08, mRNA.
<a href="#">Qser1</a>	<a href="#">Qser1.aSep08</a>	<a href="#">311266</a>	9001	708		200	glutamine and serine rich 1 (Qser1) mRNA.
<a href="#">Qtrt1</a>	<a href="#">Qtrt1.bSep08</a>	<a href="#">64016</a>	2367	750	5	141	queuine tRNA-ribosyltransferase 1 (Qtrt1) alternative variant bSep08, mRNA.
<a href="#">Qtrt1</a>	<a href="#">Qtrt1.cSep08</a>	<a href="#">64016</a>	801	360	1	97	queuine tRNA-ribosyltransferase 1 (10.9 kD) (Qtrt1) alternative variant cSep08, mRNA.
<a href="#">Qtrtd1</a>	<a href="#">Qtrtd1.aSep08</a>	<a href="#">288364</a>	6360	909		230	queuine/other tRNA-ribosyltransferase (Qtrtd1) mRNA.
<a href="#">R3hcc1</a>	<a href="#">R3hcc1.aSep08</a>	<a href="#">361064</a>	7395	1714	8	515	R3H domain and coiled-coil containing 1 (R3hcc1) alternative variant aSep08, mRNA.
<a href="#">R3hcc1</a>	<a href="#">R3hcc1.bSep08</a>	<a href="#">361064</a>	2147	825	4	247	R3H domain and coiled-coil containing 1 (R3hcc1) alternative variant bSep08, mRNA.

<a href="#">R3hcc1</a>	<a href="#">R3hcc1.cSep08</a>	<a href="#">361064</a>	10681	745	4	247	R3H domain and coiled-coil containing 1 (R3hcc1) alternative variant cSep08, mRNA.
<a href="#">R3hdm1</a>	<a href="#">R3hdm1.bSep08</a>	<a href="#">304763</a>	29853	1775	13	455	R3H domain 1 (52.3 kD) (R3hdm1) alternative variant bSep08, mRNA.
<a href="#">R3hdm1</a>	<a href="#">R3hdm1.cSep08</a>	<a href="#">304763</a>	16083	463	3	81	R3H domain 1 (R3hdm1) alternative variant cSep08, mRNA.
<a href="#">R3hdm1</a>	<a href="#">R3hdm1.dSep08</a>	<a href="#">304763</a>	5422	770	4	79	putative protein (9.1 kD) (R3hdm1) alternative variant dSep08, mRNA.
<a href="#">R3hdm1</a>	<a href="#">R3hdm1.eSep08</a>	<a href="#">304763</a>	827	727	2	43	putative protein (R3hdm1) alternative variant eSep08, mRNA.
<a href="#">R3hdm2</a>	<a href="#">R3hdm2.bSep08</a>	<a href="#">362894</a>	14877	1793	8	421	putative protein of vertebrate origin (R3hdm2) alternative variant bSep08, mRNA.
<a href="#">R3hdm2</a>	<a href="#">R3hdm2.cSep08</a>	<a href="#">362894</a>	5856	1585	5	312	putative protein of vertebrate origin (R3hdm2) alternative variant cSep08, mRNA.
<a href="#">R3hdm2</a>	<a href="#">R3hdm2.dSep08</a>	<a href="#">362894</a>	11089	675	5	225	putative protein of vertebrate origin (R3hdm2) alternative variant dSep08, mRNA.
<a href="#">RA.0</a>	<a href="#">RA.0.aSep08</a>		7636	539	3	179	guanine nucleotide exchange factor (RA.0) alternative variant aSep08, mRNA.
<a href="#">RA.0</a>	<a href="#">RA.0.bSep08</a>		5379	418	1	123	guanine nucleotide exchange factor (RA.0) alternative variant bSep08, mRNA.
<a href="#">Rab1</a>	<a href="#">Rab1.bSep08</a>	<a href="#">81754</a>	47195	331	4	58	RAB1, member RAS oncogene family (Rab1) alternative variant bSep08, mRNA.
<a href="#">Rab2b</a>	<a href="#">Rab2b.bSep08</a>	<a href="#">305853</a>	17926	1116	6	169	RAB2B, member RAS oncogene family (19.3 kD) (Rab2b) alternative variant bSep08, mRNA.
<a href="#">Rab2b</a>	<a href="#">Rab2b.cSep08</a>	<a href="#">305853</a>	17455	675	7	168	RAB2B, member RAS oncogene family (Rab2b) alternative variant cSep08, mRNA.
<a href="#">Rab2b</a>	<a href="#">Rab2b.dSep08</a>	<a href="#">305853</a>	21556	2601	7	146	RAB2B, member RAS oncogene family (16.6 kD) (Rab2b) alternative variant dSep08, mRNA.
<a href="#">Rab2b</a>	<a href="#">Rab2b.eSep08</a>	<a href="#">305853</a>	12778	534	6	142	RAB2B, member RAS oncogene family (Rab2b) alternative variant eSep08, mRNA.
<a href="#">Rab2b</a>	<a href="#">Rab2b.fSep08</a>	<a href="#">305853</a>	4853	524	2	76	RAB2B, member RAS oncogene family (Rab2b) alternative variant fSep08, mRNA.
<a href="#">Rab2l</a>	<a href="#">Rab2l.bSep08</a>	<a href="#">294283</a>	1144	705	5	201	RAB2, member RAS oncogene family-like (Rab2l) alternative variant bSep08, mRNA.
<a href="#">Rab2l</a>	<a href="#">Rab2l.cSep08</a>	<a href="#">294283</a>	934	740	3	159	RAB2, member RAS oncogene family-like (Rab2l) alternative variant cSep08, mRNA.
<a href="#">Rab2l</a>	<a href="#">Rab2l.dSep08</a>	<a href="#">294283</a>	1214	872	3	119	RAB2, member RAS oncogene family-like (Rab2l) alternative variant dSep08, mRNA.
<a href="#">Rab2l</a>	<a href="#">Rab2l.eSep08</a>	<a href="#">294283</a>	775	688	2	115	RAB2, member RAS oncogene family-like (Rab2l) alternative variant eSep08, mRNA.
<a href="#">Rab3b</a>	<a href="#">Rab3b.aSep08</a>	<a href="#">81755</a>	98895	1153	5	259	RAB3B, member RAS oncogene family (Rab3b) alternative variant aSep08, mRNA.
<a href="#">Rab3b</a>	<a href="#">Rab3b.bSep08</a>	<a href="#">81755</a>	98325	610	5	202	RAB3B, member RAS oncogene family (Rab3b) alternative variant bSep08, mRNA.
<a href="#">Rab3b</a>	<a href="#">Rab3b.cSep08</a>	<a href="#">81755</a>	17764	588	3	121	RAB3B, member RAS oncogene family (Rab3b) alternative variant cSep08, mRNA.

<a href="#">Rab3b</a>	<a href="#">Rab3b.dSep08</a>	<a href="#">81755</a>	8808	267	2	89	RAB3B, member RAS oncogene family (Rab3b) alternative variant dSep08, mRNA.
<a href="#">Rab3c</a>	<a href="#">Rab3c.bSep08</a>	<a href="#">171058</a>	19396	315	2	34	RAB3C, member RAS oncogene family (Rab3c) alternative variant bSep08, mRNA.
<a href="#">Rab3c</a>	<a href="#">Rab3c.cSep08</a>	<a href="#">171058</a>	1419	299	2	33	RAB3C, member RAS oncogene family (3.5 kD) (Rab3c) alternative variant cSep08, mRNA.
<a href="#">Rab3gap2</a>	<a href="#">Rab3gap2.aSep08</a>	<a href="#">289350</a>	14034	1655	10	551	RAB3 GTPase activating protein subunit 2 (Rab3gap2) alternative variant aSep08, mRNA.
<a href="#">Rab3gap2</a>	<a href="#">Rab3gap2.bSep08</a>	<a href="#">289350</a>	6143	731	3	208	RAB3 GTPase activating protein subunit 2 (Rab3gap2) alternative variant bSep08, mRNA.
<a href="#">Rab3il1</a>	<a href="#">Rab3il1.bSep08</a>	<a href="#">171452</a>	25971	707	4	40	RAB3A interacting protein (rabin3)-like 1 (4.1 kD) (Rab3il1) alternative variant bSep08, mRNA.
<a href="#">Rab3il1</a>	<a href="#">Rab3il1.cSep08</a>	<a href="#">171452</a>	13551	855	4	50	RAB3A interacting protein (rabin3)-like 1 (Rab3il1) alternative variant cSep08, mRNA.
<a href="#">Rab3il1</a>	<a href="#">Rab3il1.dSep08</a>	<a href="#">171452</a>	13218	634	5	93	RAB3A interacting protein (rabin3)-like 1 (Rab3il1) alternative variant dSep08, mRNA.
<a href="#">Rab3ip</a>	<a href="#">Rab3ip.bSep08</a>	<a href="#">29885</a>	3061	548	3	104	RAB3A interacting protein (Rab3ip) alternative variant bSep08, mRNA.
<a href="#">Rab4a</a>	<a href="#">Rab4a.bSep08</a>	<a href="#">25532</a>	26747	752	7	164	RAB4A, member RAS oncogene family (Rab4a) alternative variant bSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.bSep08</a>	<a href="#">50866</a>	5267	753	4	250	EGL nine homolog 2 (Rab4bandEgln2) alternative variant bSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.bSep08</a>	<a href="#">308457</a>	5267	753	4	250	EGL nine homolog 2 (Rab4bandEgln2) alternative variant bSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.dSep08</a>	<a href="#">50866</a>	14616	1178	7	186	GTP-binding protein like (20.6 kD) (Rab4bandEgln2) alternative variant dSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.dSep08</a>	<a href="#">308457</a>	14616	1178	7	186	GTP-binding protein like (20.6 kD) (Rab4bandEgln2) alternative variant dSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.eSep08</a>	<a href="#">50866</a>	7436	518	5	154	member ras oncogene family (Rab4bandEgln2) alternative variant eSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.eSep08</a>	<a href="#">308457</a>	7436	518	5	154	member ras oncogene family (Rab4bandEgln2) alternative variant eSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.fSep08</a>	<a href="#">50866</a>	2399	740	4	138	protein rab-4a (Rab4bandEgln2) alternative variant fSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.fSep08</a>	<a href="#">308457</a>	2399	740	4	138	protein rab-4a (Rab4bandEgln2) alternative variant fSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.gSep08</a>	<a href="#">50866</a>	3004	2156	4	124	EGL nine homolog 2 (Rab4bandEgln2) alternative variant gSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.gSep08</a>	<a href="#">308457</a>	3004	2156	4	124	EGL nine homolog 2 (Rab4bandEgln2) alternative variant gSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.hSep08</a>	<a href="#">50866</a>	1128	769	2	115	EGL nine homolog 2 (13.3 kD) (Rab4bandEgln2) alternative variant hSep08, mRNA.
<a href="#">Rab4bandEgln2</a>	<a href="#">Rab4bandEgln2.hSep08</a>	<a href="#">308457</a>	1128	769	2	115	EGL nine homolog 2 (13.3 kD) (Rab4bandEgln2) alternative variant hSep08, mRNA.
<a href="#">Rab5a</a>	<a href="#">Rab5a.bSep08</a>	<a href="#">64633</a>	491	396	2	111	putative protein (Rab5a) alternative variant bSep08, mRNA.

<a href="#">Rab5a</a>	<a href="#">Rab5a.cSep08</a>	<a href="#">64633</a>	3317	291	2	94	putative protein (Rab5a) alternative variant cSep08, mRNA.
<a href="#">Rab5a</a>	<a href="#">Rab5a.dSep08</a>	<a href="#">64633</a>	28645	1674	4	78	RAB5C member RAS oncogene family (8.5 kD) (Rab5a) alternative variant dSep08, complete mRNA.
<a href="#">Rab5b</a>	<a href="#">Rab5b.bSep08</a>	<a href="#">288779</a>	13639	675	4	205	RAB5B, member RAS oncogene family (Rab5b) alternative variant bSep08, mRNA.
<a href="#">Rab5b</a>	<a href="#">Rab5b.cSep08</a>	<a href="#">288779</a>	10067	919	3	113	RAB5B, member RAS oncogene family (Rab5b) alternative variant cSep08, mRNA.
<a href="#">Rab6a</a>	<a href="#">Rab6a.aSep08</a>	<a href="#">84379</a>	38986	3231	8	208	RAB6A, member RAS oncogene family (23.6 kD) (Rab6a) alternative variant aSep08, mRNA.
<a href="#">Rab6a</a>	<a href="#">Rab6a.bSep08</a>	<a href="#">84379</a>	13403	1069	5	132	RAB6A, member RAS oncogene family (Rab6a) alternative variant bSep08, mRNA.
<a href="#">Rab6a</a>	<a href="#">Rab6a.cSep08</a>	<a href="#">84379</a>	24252	831	4	103	RAB6A, member RAS oncogene family (11.7 kD) (Rab6a) alternative variant cSep08, mRNA.
<a href="#">Rab6ip1.1</a>	<a href="#">Rab6ip1.1.aSep08</a>	<a href="#">308942</a>	26919	3131	15	708	rab6 interacting protein 1 (81.2 kD) (Rab6ip1.1) alternative variant aSep08, mRNA.
<a href="#">Rab6ip1.1</a>	<a href="#">Rab6ip1.1.bSep08</a>	<a href="#">308942</a>	3404	516	3	148	rab6 interacting protein 1 (Rab6ip1.1) alternative variant bSep08, mRNA.
<a href="#">Rab7</a>	<a href="#">Rab7.bSep08</a>	<a href="#">29448</a>	11609	633	4	183	RAB7, member RAS oncogene family (Rab7) alternative variant bSep08, mRNA.
<a href="#">Rab7</a>	<a href="#">Rab7.cSep08</a>	<a href="#">29448</a>	6214	459	4	82	RAB7, member RAS oncogene family (Rab7) alternative variant cSep08, mRNA.
<a href="#">Rab8a</a>	<a href="#">Rab8a.bSep08</a>	<a href="#">117103</a>	14904	685	1	134	RAB8A, member RAS oncogene family (Rab8a) alternative variant bSep08, mRNA.
<a href="#">Rab11a</a>	<a href="#">Rab11a.bSep08</a>	<a href="#">81830</a>	35539	784	2	101	RAB11a, member RAS oncogene family (11.7 kD) (Rab11a) alternative variant bSep08, mRNA.
<a href="#">Rab11b</a>	<a href="#">Rab11b.bSep08</a>	<a href="#">79434</a>	11939	959	1	178	RAB11B, member RAS oncogene family (19.9 kD) (Rab11b) alternative variant bSep08, mRNA.
<a href="#">Rab11fip2</a>	<a href="#">Rab11fip2.bSep08</a>	<a href="#">308003</a>	6396	1100	2	135	RAB11 family interacting protein 2 (class I) (Rab11fip2) alternative variant bSep08, mRNA.
<a href="#">Rab11fip2</a>	<a href="#">Rab11fip2.cSep08</a>	<a href="#">308003</a>	28526	429	2	84	RAB11 family interacting protein 2 (class I) (Rab11fip2) alternative variant cSep08, mRNA.
<a href="#">Rab11fip4</a>	<a href="#">Rab11fip4.cSep08</a>	<a href="#">303337</a>	28304	380	3	95	RAB11 family interacting protein 4 (class II) (Rab11fip4) alternative variant cSep08, mRNA.
<a href="#">Rab12</a>	<a href="#">Rab12.aSep08</a>	<a href="#">25530</a>	19693	566		188	RAB12, member RAS oncogene family (Rab12) mRNA.
<a href="#">Rab13</a>	<a href="#">Rab13.bSep08</a>	<a href="#">81756</a>	2036	806	6	96	RAB13, member RAS oncogene family (Rab13) alternative variant bSep08, mRNA.
<a href="#">Rab13</a>	<a href="#">Rab13.cSep08</a>	<a href="#">81756</a>	1260	746	3	86	RAB13, member RAS oncogene family (9.3 kD) (Rab13) alternative variant cSep08, mRNA.
<a href="#">Rab14</a>	<a href="#">Rab14.bSep08</a>	<a href="#">94197</a>	1758	534	3	88	RAB14, member RAS oncogene family (9.5 kD) (Rab14) alternative variant bSep08, mRNA.
<a href="#">Rab18</a>	<a href="#">Rab18.bSep08</a>	<a href="#">307039</a>	31799	2116	1	89	RAB18, member RAS oncogene family (9.7 kD) (Rab18) alternative variant bSep08, complete mRNA.
<a href="#">Rab21</a>	<a href="#">Rab21.bSep08</a>	<a href="#">299799</a>	8006	410	5	89	RAB21, member RAS oncogene family (Rab21) alternative variant bSep08, mRNA.
<a href="#">Rab22a</a>	<a href="#">Rab22a.aSep08</a>	<a href="#">366265</a>	41865	1726	5	194	RAB22A, member RAS oncogene family (21.8 kD) (Rab22a) alternative variant aSep08, complete mRNA.

<a href="#">Rab22a</a>	<a href="#">Rab22a.cSep08</a>	<a href="#">366265</a>	36674	682	3	186	RAB22A, member RAS oncogene family (Rab22a) alternative variant cSep08, mRNA.
<a href="#">Rab24</a>	<a href="#">Rab24.aSep08</a>	<a href="#">361208</a>	2307	1418	8	203	RAB24, member RAS oncogene family (23.1 kD) (Rab24) alternative variant aSep08, mRNA.
<a href="#">Rab24</a>	<a href="#">Rab24.cSep08</a>	<a href="#">361208</a>	1182	887	3	107	RAB24, member RAS oncogene family (11.9 kD) (Rab24) alternative variant cSep08, mRNA.
<a href="#">Rab24</a>	<a href="#">Rab24.dSep08</a>	<a href="#">361208</a>	2076	1889	3	76	RAB24, member RAS oncogene family (9.0 kD) (Rab24) alternative variant dSep08, mRNA.
<a href="#">Rab24</a>	<a href="#">Rab24.eSep08</a>	<a href="#">361208</a>	770	396	4	60	RAB24, member RAS oncogene family (Rab24) alternative variant eSep08, mRNA.
<a href="#">Rab26</a>	<a href="#">Rab26.bSep08</a>	<a href="#">171111</a>	3576	847	5	212	RAB26, member RAS oncogene family (Rab26) alternative variant bSep08, mRNA.
<a href="#">Rab26</a>	<a href="#">Rab26.cSep08</a>	<a href="#">171111</a>	814	715	2	78	RAB26, member RAS oncogene family (Rab26) alternative variant cSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.bSep08</a>	<a href="#">117049</a>	79248	884	7	220	RAB28, member RAS oncogene family (24.7 kD) (Rab28) alternative variant bSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.cSep08</a>	<a href="#">117049</a>	79890	662	6	184	RAB28, member RAS oncogene family (Rab28) alternative variant cSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.dSep08</a>	<a href="#">117049</a>	79408	897	6	83	RAB28, member RAS oncogene family (9.4 kD) (Rab28) alternative variant dSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.eSep08</a>	<a href="#">117049</a>	5613	233	2	77	RAB28, member RAS oncogene family (Rab28) alternative variant eSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.fSep08</a>	<a href="#">117049</a>	79403	855	5	86	RAB28, member RAS oncogene family (9.7 kD) (Rab28) alternative variant fSep08, mRNA.
<a href="#">Rab28</a>	<a href="#">Rab28.gSep08</a>	<a href="#">117049</a>	75573	469	4	58	RAB28, member RAS oncogene family (6.5 kD) (Rab28) alternative variant gSep08, mRNA.
<a href="#">Rab30</a>	<a href="#">Rab30.bSep08</a>	<a href="#">308821</a>	85854	735	4	173	RAB30, member RAS oncogene family (Rab30) alternative variant bSep08, mRNA.
<a href="#">Rab30</a>	<a href="#">Rab30.cSep08</a>	<a href="#">308821</a>	81417	410	3	53	RAB30, member RAS oncogene family (Rab30) alternative variant cSep08, mRNA.
<a href="#">Rab31</a>	<a href="#">Rab31.bSep08</a>	<a href="#">246324</a>	23344	714	1	91	RAB31, member RAS oncogene family (9.7 kD) (Rab31) alternative variant bSep08, mRNA.
<a href="#">Rab32</a>	<a href="#">Rab32.bSep08</a>	<a href="#">365042</a>	15628	2910	1	163	RAB32, member RAS oncogene family (Rab32) alternative variant bSep08, mRNA.
<a href="#">Rab34</a>	<a href="#">Rab34.bSep08</a>	<a href="#">360571</a>	3699	1498	8	215	RAB34, member of RAS oncogene family (Rab34) alternative variant bSep08, mRNA.
<a href="#">Rab34</a>	<a href="#">Rab34.cSep08</a>	<a href="#">360571</a>	965	799	3	146	RAB34, member of RAS oncogene family (Rab34) alternative variant cSep08, mRNA.
<a href="#">Rab34</a>	<a href="#">Rab34.dSep08</a>	<a href="#">360571</a>	911	699	3	81	RAB34, member of RAS oncogene family (9.1 kD) (Rab34) alternative variant dSep08, mRNA.
<a href="#">Rab35</a>	<a href="#">Rab35.bSep08</a>	<a href="#">288700</a>	15232	3235	3	235	RAB35, member RAS oncogene family (Rab35) alternative variant bSep08, mRNA.
<a href="#">Rab35</a>	<a href="#">Rab35.cSep08</a>	<a href="#">288700</a>	13915	755	3	183	RAB35, member RAS oncogene family (21.0 kD) (Rab35) alternative variant cSep08, mRNA.
<a href="#">Rab36</a>	<a href="#">Rab36.bSep08</a>	<a href="#">690407</a>	11890	750	6	236	RAB36, member RAS oncogene family (Rab36) alternative variant bSep08, mRNA.

<a href="#">Rab40b</a>	<a href="#">Rab40b.bSep08</a>	<a href="#">303754</a>	25611	685	6	106	rab40b, member RAS oncogene family (Rab40b) alternative variant bSep08, mRNA.
<a href="#">Rab43</a>	<a href="#">Rab43.bSep08</a>	<a href="#">500249</a>	17504	887	2	122	RAB43, member RAS oncogene family (13.5 kD) (Rab43) alternative variant bSep08, mRNA.
<a href="#">Rabac1</a>	<a href="#">Rabac1.bSep08</a>	<a href="#">83583</a>	1622	1377	2	161	rab acceptor 1 (18.4 kD) (Rabac1) alternative variant bSep08, mRNA.
<a href="#">Rabac1</a>	<a href="#">Rabac1.cSep08</a>	<a href="#">83583</a>	3020	705	5	91	putative protein (Rabac1) alternative variant cSep08, mRNA.
<a href="#">Rabac1</a>	<a href="#">Rabac1.dSep08</a>	<a href="#">83583</a>	922	679	2	90	rab acceptor 1 (Rabac1) alternative variant dSep08, mRNA.
<a href="#">Rabac1</a>	<a href="#">Rabac1.eSep08</a>	<a href="#">83583</a>	682	598	2	75	putative protein (Rabac1) alternative variant eSep08, mRNA.
<a href="#">Rabep1</a>	<a href="#">Rabep1.bSep08</a>	<a href="#">54190</a>	5900	407	4	109	rabaptin, RAB GTPase binding effector protein 1 (Rabep1) alternative variant bSep08, mRNA.
<a href="#">Rabep2</a>	<a href="#">Rabep2.aSep08</a>	<a href="#">80754</a>	10482	1089	5	348	rabaptin, RAB GTPase binding effector protein 2 (Rabep2) alternative variant aSep08, mRNA.
<a href="#">Rabep2</a>	<a href="#">Rabep2.bSep08</a>	<a href="#">80754</a>	6297	541	1	151	rabaptin, RAB GTPase binding effector protein 2 (16.5 kD) (Rabep2) alternative variant bSep08, mRNA.
<a href="#">Rabepk</a>	<a href="#">Rabepk.bSep08</a>	<a href="#">296649</a>	19237	1019	7	331	rab9 effector protein with kelch motifs (Rabepk) alternative variant bSep08, mRNA.
<a href="#">Rabepk</a>	<a href="#">Rabepk.cSep08</a>	<a href="#">296649</a>	17384	987	6	175	rab9 effector protein with kelch motifs (Rabepk) alternative variant cSep08, mRNA.
<a href="#">Rabepk</a>	<a href="#">Rabepk.dSep08</a>	<a href="#">296649</a>	11072	410	4	136	rab9 effector protein with kelch motifs (Rabepk) alternative variant dSep08, mRNA.
<a href="#">Rabgap1</a>	<a href="#">Rabgap1.bSep08</a>	<a href="#">311911</a>	8205	882	6	213	RAB GTPase activating protein 1 (Rabgap1) alternative variant bSep08, mRNA.
<a href="#">Rabgap1</a>	<a href="#">Rabgap1.cSep08</a>	<a href="#">311911</a>	18675	457	3	121	RAB GTPase activating protein 1 (Rabgap1) alternative variant cSep08, mRNA.
<a href="#">Rabgap1l</a>	<a href="#">Rabgap1l.bSep08</a>	<a href="#">304914</a>	51722	940	2	72	RAB GTPase activating protein 1-like (8.7 kD) (Rabgap1l) alternative variant bSep08, mRNA.
<a href="#">Rabgef1</a>	<a href="#">Rabgef1.bSep08</a>	<a href="#">360797</a>	40931	759	2	252	RAB guanine nucleotide exchange factor (GEF) 1 (Rabgef1) alternative variant bSep08, mRNA.
<a href="#">Rabggta</a>	<a href="#">Rabggta.bSep08</a>	<a href="#">58983</a>	4023	1668	6	252	rab geranylgeranyl transferase, a subunit (Rabggta) alternative variant bSep08, mRNA.
<a href="#">Rabggta</a>	<a href="#">Rabggta.cSep08</a>	<a href="#">58983</a>	1511	691	6	230	rab geranylgeranyl transferase, a subunit (Rabggta) alternative variant cSep08, mRNA.
<a href="#">Rabggta</a>	<a href="#">Rabggta.dSep08</a>	<a href="#">58983</a>	1847	815	6	210	rab geranylgeranyl transferase, a subunit (Rabggta) alternative variant dSep08, mRNA.
<a href="#">Rabggta</a>	<a href="#">Rabggta.eSep08</a>	<a href="#">58983</a>	1718	632	2	96	rab geranylgeranyl transferase, a subunit (10.4 kD) (Rabggta) alternative variant eSep08, mRNA.
<a href="#">Rabggta</a>	<a href="#">Rabggta.iSep08</a>	<a href="#">58983</a>	488	293	3	49	rab geranylgeranyl transferase, a subunit (Rabggta) alternative variant iSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.bSep08</a>	<a href="#">25533</a>	6193	1235	8	208	RAB geranylgeranyl transferase CRA b (22.8 kD) (Rabggtb) alternative variant bSep08, complete mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.cSep08</a>	<a href="#">25533</a>	4887	929	8	116	rab geranylgeranyl transferase CRA b (12.4 kD) (Rabggtb) alternative variant cSep08, mRNA.



<a href="#">Rabggtb</a>	<a href="#">Rabggtb.dSep08</a>	<a href="#">25533</a>	5240	947	9	113	rabggtb protein (12.8 kD) (Rabggtb) alternative variant dSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.eSep08</a>	<a href="#">25533</a>	4075	638	3	87	RAB geranylgeranyl transferase (Rabggtb) alternative variant eSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.fSep08</a>	<a href="#">25533</a>	3305	1343	5	70	rab geranylgeranyl transferase (7.5 kD) (Rabggtb) alternative variant fSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.gSep08</a>	<a href="#">25533</a>	1629	1092	3	60	rab geranylgeranyl transferase (Rabggtb) alternative variant gSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.iSep08</a>	<a href="#">25533</a>	3353	1412	4	73	rabggtb protein (8.3 kD) (Rabggtb) alternative variant iSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.jSep08</a>	<a href="#">25533</a>	2850	1340	4	88	rab geranylgeranyl transferase (Rabggtb) alternative variant jSep08, mRNA.
<a href="#">Rabggtb</a>	<a href="#">Rabggtb.kSep08</a>	<a href="#">25533</a>	4140	986	9	73	rabggtb protein (8.3 kD) (Rabggtb) alternative variant kSep08, mRNA.
<a href="#">Rabl2a</a>	<a href="#">Rabl2a.aSep08</a>	<a href="#">362987</a>	8191	1388	8	250	RAB, member of RAS oncogene family-like 2A (28.6 kD) (Rabl2a) alternative variant aSep08, mRNA.
<a href="#">Rabl2a</a>	<a href="#">Rabl2a.cSep08</a>	<a href="#">362987</a>	2805	601	2	144	RAB, member of RAS oncogene family-like 2A (Rabl2a) alternative variant cSep08, mRNA.
<a href="#">Rabl2a</a>	<a href="#">Rabl2a.dSep08</a>	<a href="#">362987</a>	6871	366	3	97	RAB, member of RAS oncogene family-like 2A (Rabl2a) alternative variant dSep08, mRNA.
<a href="#">Rabl3</a>	<a href="#">Rabl3.bSep08</a>	<a href="#">360720</a>	6752	1757	4	80	RAB, member of RAS oncogene family-like 3 (Rabl3) alternative variant bSep08, mRNA.
<a href="#">Rabl4</a>	<a href="#">Rabl4.aSep08</a>	<a href="#">300062</a>	7615	545		128	RAB, member of RAS oncogene family-like 4 (Rabl4) alternative variant aSep08, mRNA.
<a href="#">Rabl5</a>	<a href="#">Rabl5.bSep08</a>	<a href="#">288585</a>	7015	1270	2	100	RAB, member RAS oncogene family-like 5 (11.0 kD) (Rabl5) alternative variant bSep08, mRNA.
<a href="#">raby</a>	<a href="#">raby.aSep08</a>		5002	390		124	putative protein (raby) mRNA.
<a href="#">Rac1</a>	<a href="#">Rac1.aSep08</a>	<a href="#">363875</a>	21681	2312	7	211	ras-related C3 botulinum toxin substrate 1 (23.4 kD) (Rac1) alternative variant aSep08, complete mRNA.
<a href="#">Rac2</a>	<a href="#">Rac2.aSep08</a>	<a href="#">366957</a>	12381	1846	6	230	RAS-related C3 botulinum substrate 2 (25.9 kD) (Rac2) alternative variant aSep08, mRNA.
<a href="#">Rac2</a>	<a href="#">Rac2.cSep08</a>	<a href="#">366957</a>	11805	670	7	185	RAS-related C3 botulinum substrate 2 (20.8 kD) (Rac2) alternative variant cSep08, mRNA.
<a href="#">Rac2</a>	<a href="#">Rac2.dSep08</a>	<a href="#">366957</a>	11894	717	6	171	RAS-related C3 botulinum substrate 2 (19.1 kD) (Rac2) alternative variant dSep08, mRNA.
<a href="#">Racgap1</a>	<a href="#">Racgap1.bSep08</a>	<a href="#">315298</a>	29804	2913	2	626	rac GTPase-activating protein 1 (69.9 kD) (Racgap1) alternative variant bSep08, mRNA.
<a href="#">rachy</a>	<a href="#">rachy.aSep08</a>		1103	513		61	putative protein (6.9 kD) (rachy) mRNA.
<a href="#">Rad1</a>	<a href="#">Rad1.bSep08</a>	<a href="#">294800</a>	3464	676	3	98	RAD1 homolog (S. pombe) (11.2 kD) (Rad1) alternative variant bSep08, mRNA.
<a href="#">Rad9.0</a>	<a href="#">Rad9.0.aSep08</a>		2746	1775	5	205	CRA a (Rad9.0) alternative variant aSep08, mRNA.
<a href="#">Rad9.0</a>	<a href="#">Rad9.0.bSep08</a>		5091	811	6	171	CRA a (19.0 kD) (Rad9.0) alternative variant bSep08, mRNA.
<a href="#">Rad9b</a>	<a href="#">Rad9b.bSep08</a>	<a href="#">363924</a>	2751	388	3	116	RAD9 homolog B (S. cerevisiae) (Rad9b) alternative variant bSep08, mRNA.
<a href="#">Rad18</a>	<a href="#">Rad18.bSep08</a>	<a href="#">362412</a>	66980	712	5	175	RAD18 homolog (S. cerevisiae) (Rad18) alternative variant bSep08, mRNA.

<a href="#">Rad18</a>	<a href="#">Rad18.cSep08</a>	<a href="#">362412</a>	59855	440	3	63	RAD18 homolog (S. cerevisiae) (Rad18) alternative variant cSep08, mRNA.
<a href="#">Rad18</a>	<a href="#">Rad18.dSep08</a>	<a href="#">362412</a>	7074	381	3	23	RAD18 homolog (S. cerevisiae) (2.5 kD) (Rad18) alternative variant dSep08, mRNA.
<a href="#">Rad18</a>	<a href="#">Rad18.eSep08</a>	<a href="#">362412</a>	21234	330	2	46	RAD18 homolog (S. cerevisiae) (Rad18) alternative variant eSep08, mRNA.
<a href="#">Rad21</a>	<a href="#">Rad21.bSep08</a>	<a href="#">314949</a>	763	382	2	117	RAD21 homolog (S. pombe) (Rad21) alternative variant bSep08, mRNA.
<a href="#">Rad23a</a>	<a href="#">Rad23a.bSep08</a>	<a href="#">361381</a>	3993	1668	7	255	RAD23a homolog (S. cerevisiae) (Rad23a) alternative variant bSep08, mRNA.
<a href="#">Rad23a</a>	<a href="#">Rad23a.cSep08</a>	<a href="#">361381</a>	2767	776	5	244	RAD23a homolog (S. cerevisiae) (Rad23a) alternative variant cSep08, mRNA.
<a href="#">Rad23a</a>	<a href="#">Rad23a.dSep08</a>	<a href="#">361381</a>	2966	723	6	241	RAD23a homolog (S. cerevisiae) (Rad23a) alternative variant dSep08, mRNA.
<a href="#">Rad23b</a>	<a href="#">Rad23b.bSep08</a>	<a href="#">298012</a>	16672	421	3	94	RAD23b homolog (S. cerevisiae) (Rad23b) alternative variant bSep08, mRNA.
<a href="#">Rad23b</a>	<a href="#">Rad23b.cSep08</a>	<a href="#">298012</a>	16407	891	2	74	RAD23b homolog (S. cerevisiae) (8.3 kD) (Rad23b) alternative variant cSep08, mRNA.
<a href="#">Rad51ap1</a>	<a href="#">Rad51ap1.aSep08</a>	<a href="#">689055</a>	12309	1225	9	349	RAD51 associated protein 1 (Rad51ap1) alternative variant aSep08, mRNA.
<a href="#">Rad51ap1</a>	<a href="#">Rad51ap1.cSep08</a>	<a href="#">689055</a>	4318	856	5	202	RAD51 associated protein 1 (Rad51ap1) alternative variant cSep08, mRNA.
<a href="#">Rad51c</a>	<a href="#">Rad51c.aSep08</a>	<a href="#">497976</a>	26070	1832		345	rad51 homolog c (S. cerevisiae) (Rad51c) mRNA.
<a href="#">Rad5111</a>	<a href="#">Rad5111.aSep08</a>	<a href="#">500679</a>	41992	768		237	RAD51-like 1 (S. cerevisiae) (Rad5111) alternative variant aSep08, mRNA.
<a href="#">Rad5111</a>	<a href="#">Rad5111.bSep08</a>	<a href="#">500679</a>	41960	839	1	223	RAD51-like 1 (S. cerevisiae) (Rad5111) alternative variant bSep08, mRNA.
<a href="#">Rad52</a>	<a href="#">Rad52.bSep08</a>	<a href="#">297561</a>	2366	735	3	133	RAD52 homolog (S. cerevisiae) (Rad52) alternative variant bSep08, mRNA.
<a href="#">Rad52</a>	<a href="#">Rad52.cSep08</a>	<a href="#">297561</a>	2512	1464	2	111	RAD52 homolog (S. cerevisiae) (12.6 kD) (Rad52) alternative variant cSep08, mRNA.
<a href="#">Rad52</a>	<a href="#">Rad52.dSep08</a>	<a href="#">297561</a>	1484	427	2	39	RAD52 homolog (S. cerevisiae) (Rad52) alternative variant dSep08, mRNA.
<a href="#">radar</a>	<a href="#">radar.aSep08</a>		67291	1429	1	46	putative protein (4.8 kD) (radar) alternative variant aSep08, mRNA.
<a href="#">radar</a>	<a href="#">radar.bSep08</a>		40118	313	1	43	putative protein (radar) alternative variant bSep08, mRNA.
<a href="#">Radical_SAM.0</a>	<a href="#">Radical_SAM.0.aSep08</a>		62107	2812	15	591	elongation protein 3 homolog (Radical_SAM.0) alternative variant aSep08, mRNA.
<a href="#">Radical_SAM.0</a>	<a href="#">Radical_SAM.0.bSep08</a>		27137	784	8	261	elongation protein 3 homolog CRA c (Radical_SAM.0) alternative variant bSep08, mRNA.
<a href="#">Radical_SAM.0</a>	<a href="#">Radical_SAM.0.cSep08</a>		16959	674	7	212	elongation protein 3 homolog (Radical_SAM.0) alternative variant cSep08, mRNA.
<a href="#">Radil</a>	<a href="#">Radil.bSep08</a>	<a href="#">304299</a>	1855	740	3	203	rap GTPase interactor (Radil) alternative variant bSep08, mRNA.
<a href="#">Radil</a>	<a href="#">Radil.cSep08</a>	<a href="#">304299</a>	2664	538	3	179	rap GTPase interactor (Radil) alternative variant cSep08, mRNA.

<a href="#">Rae1</a>	<a href="#">Rae1.aSep08</a>	<a href="#">362281</a>	9210	1791	7	500	RAE1 RNA export 1 homolog (S. pombe) (Rae1) alternative variant aSep08, mRNA.
<a href="#">Rae1</a>	<a href="#">Rae1.cSep08</a>	<a href="#">362281</a>	696	621	2	116	RAE1 RNA export 1 homolog (S. pombe) (Rae1) alternative variant cSep08, mRNA.
<a href="#">Rae1</a>	<a href="#">Rae1.dSep08</a>	<a href="#">362281</a>	3168	400	3	103	RAE1 RNA export 1 homolog (S. pombe) (Rae1) alternative variant dSep08, mRNA.
<a href="#">Raf1</a>	<a href="#">Raf1.bSep08</a>	<a href="#">24703</a>	13099	1133	11	377	v-raf-leukemia viral oncogene 1 (Raf1) alternative variant bSep08, mRNA.
<a href="#">Raf1</a>	<a href="#">Raf1.cSep08</a>	<a href="#">24703</a>	5262	2121	6	257	v-raf-leukemia viral oncogene 1 (Raf1) alternative variant cSep08, mRNA.
<a href="#">Raf1</a>	<a href="#">Raf1.dSep08</a>	<a href="#">24703</a>	2783	719	5	164	v-raf-leukemia viral oncogene 1 (Raf1) alternative variant dSep08, mRNA.
<a href="#">Raf1</a>	<a href="#">Raf1.eSep08</a>	<a href="#">24703</a>	1237	737	4	122	v-raf-leukemia viral oncogene 1 (Raf1) alternative variant eSep08, mRNA.
<a href="#">Raf1</a>	<a href="#">Raf1.fSep08</a>	<a href="#">24703</a>	1309	1152	2	89	v-raf-leukemia viral oncogene 1 (10.0 kD) (Raf1) alternative variant fSep08, mRNA.
<a href="#">rafer</a>	<a href="#">rafer.aSep08</a>		7786	988		47	putative protein (rafer) mRNA.
<a href="#">raflo</a>	<a href="#">raflo.aSep08</a>		738	236	2	64	putative protein (raflo) alternative variant aSep08, mRNA.
<a href="#">raflu</a>	<a href="#">raflu.aSep08</a>		50513	706	2	58	putative protein (raflu) alternative variant aSep08, mRNA.
<a href="#">raflu</a>	<a href="#">raflu.bSep08</a>		1520	593	1	99	CRA a like (11.1 kD) (raflu) alternative variant bSep08, mRNA.
<a href="#">Rag1ap1</a>	<a href="#">Rag1ap1.bSep08</a>	<a href="#">295245</a>	1577	755	1	116	recombination activating gene 1 activating protein 1 (Rag1ap1) alternative variant bSep08, mRNA.
<a href="#">ragar</a>	<a href="#">ragar.aSep08</a>		5116	593		87	CRA b like (ragar) mRNA.
<a href="#">Rage</a>	<a href="#">Rage.aSep08</a>	<a href="#">362787</a>	20734	1485	7	301	renal tumor antigen (Rage) alternative variant aSep08, mRNA.
<a href="#">Rage</a>	<a href="#">Rage.cSep08</a>	<a href="#">362787</a>	5719	743	2	75	renal tumor antigen (Rage) alternative variant cSep08, mRNA.
<a href="#">Rai1</a>	<a href="#">Rai1.aSep08</a>	<a href="#">303188</a>	10480	2870	3	415	retinoic acid induced 1 (Rai1) alternative variant aSep08, mRNA.
<a href="#">Rai1</a>	<a href="#">Rai1.bSep08</a>	<a href="#">303188</a>	2069	1870	1	51	retinoic acid induced 1 (6.1 kD) (Rai1) alternative variant bSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.aSep08</a>	<a href="#">287446</a>	11626	1457	8	320	retinoic acid induced 12 like (35.6 kD) (Rai12) alternative variant aSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.cSep08</a>	<a href="#">287446</a>	6399	1530	4	183	retinoic acid induced 12 like (20.2 kD) (Rai12) alternative variant cSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.dSep08</a>	<a href="#">287446</a>	3714	1276	4	173	retinoic acid induced 12 CRA b like (19.5 kD) (Rai12) alternative variant dSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.eSep08</a>	<a href="#">287446</a>	6008	612	4	153	retinoic acid induced 12 CRA e like (Rai12) alternative variant eSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.fSep08</a>	<a href="#">287446</a>	1415	684	2	98	S-phase 2 protein (Rai12) alternative variant fSep08, mRNA.
<a href="#">Rai12</a>	<a href="#">Rai12.gSep08</a>	<a href="#">287446</a>	3023	484	3	88	dermal papilla derived protein 6 CRA a like (Rai12) alternative variant gSep08, mRNA.
<a href="#">Rai14</a>	<a href="#">Rai14.bSep08</a>	<a href="#">294804</a>	10806	772	5	257	retinoic acid induced 14 (Rai14) alternative variant bSep08, mRNA.

<a href="#">Rai14</a>	<a href="#">Rai14.cSep08</a>	<a href="#">294804</a>	117597	698	8	176	retinoic acid induced 14 (Rai14) alternative variant cSep08, mRNA.
<a href="#">Rai16</a>	<a href="#">Rai16.aSep08</a>	<a href="#">306015</a>	17467	3814	17	756	retinoic acid induced 16 (Rai16) alternative variant aSep08, mRNA.
<a href="#">raja</a>	<a href="#">raja.aSep08</a>		60013	330		36	putative protein (raja) mRNA.
<a href="#">rajey</a>	<a href="#">rajey.aSep08</a>		11565	1331	4	443	shroom (rajey) alternative variant aSep08, mRNA.
<a href="#">rakee</a>	<a href="#">rakee.aSep08</a>		2564	687	2	63	putative protein (7.5 kD) (rakee) alternative variant aSep08, mRNA.
<a href="#">rakler</a>	<a href="#">rakler.aSep08</a>		16211	537	2	143	putative protein of mammalian origin (15.7 kD) (rakler) alternative variant aSep08, mRNA.
<a href="#">rakler</a>	<a href="#">rakler.bSep08</a>		14571	824	1	101	putative protein of mammalian origin (rakler) alternative variant bSep08, mRNA.
<a href="#">Rala</a>	<a href="#">Rala.bSep08</a>	<a href="#">81757</a>	6211	584	1	120	v-ral simian leukemia viral oncogene homolog A (ras related) (Rala) alternative variant bSep08, mRNA.
<a href="#">Ralbp1</a>	<a href="#">Ralbp1.bSep08</a>	<a href="#">84014</a>	20670	625	3	156	ralA binding protein 1 (Ralbp1) alternative variant bSep08, mRNA.
<a href="#">Ralbp1</a>	<a href="#">Ralbp1.cSep08</a>	<a href="#">84014</a>	20520	378	2	125	ralA binding protein 1 (Ralbp1) alternative variant cSep08, mRNA.
<a href="#">Ralgds</a>	<a href="#">Ralgds.aSep08</a>	<a href="#">29622</a>	33574	1664	4	422	ral guanine nucleotide dissociation stimulator (Ralgds) alternative variant aSep08, mRNA.
<a href="#">Ralgds</a>	<a href="#">Ralgds.bSep08</a>	<a href="#">29622</a>	12145	1697	3	407	ral guanine nucleotide dissociation stimulator (Ralgds) alternative variant bSep08, mRNA.
<a href="#">Ralgds</a>	<a href="#">Ralgds.cSep08</a>	<a href="#">29622</a>	7149	1719	10	370	ral guanine nucleotide dissociation stimulator (41.0 kD) (Ralgds) alternative variant cSep08, mRNA.
<a href="#">Ralgds</a>	<a href="#">Ralgds.dSep08</a>	<a href="#">29622</a>	2405	570	2	189	ral guanine nucleotide dissociation stimulator (Ralgds) alternative variant dSep08, mRNA.
<a href="#">Ralgps2</a>	<a href="#">Ralgps2.aSep08</a>	<a href="#">304887</a>	68984	2424	14	454	GEF with PH domain motif 2 (Ralgps2) alternative variant aSep08, mRNA.
<a href="#">Ralgps2</a>	<a href="#">Ralgps2.cSep08</a>	<a href="#">304887</a>	14826	886	6	172	GEF with PH domain motif 2 (19.6 kD) (Ralgps2) alternative variant cSep08, mRNA.
<a href="#">ralo</a>	<a href="#">ralo.aSep08</a>		18387	535		107	ecotropic viral integration site 5-like (ralo) mRNA.
<a href="#">Raly</a>	<a href="#">Raly.bSep08</a>	<a href="#">296301</a>	61273	1359	6	244	hnRNP-associated with lethal yellow (26.5 kD) (Raly) alternative variant bSep08, mRNA.
<a href="#">Raly</a>	<a href="#">Raly.cSep08</a>	<a href="#">296301</a>	61572	692	5	230	hnRNP-associated with lethal yellow (Raly) alternative variant cSep08, mRNA.
<a href="#">Raly</a>	<a href="#">Raly.dSep08</a>	<a href="#">296301</a>	60051	842	6	165	hnRNP-associated with lethal yellow (Raly) alternative variant dSep08, mRNA.
<a href="#">Raly</a>	<a href="#">Raly.eSep08</a>	<a href="#">296301</a>	58004	652	5	126	hnRNP-associated with lethal yellow (Raly) alternative variant eSep08, mRNA.
<a href="#">Raly</a>	<a href="#">Raly.fSep08</a>	<a href="#">296301</a>	55785	441	3	68	hnRNP-associated with lethal yellow (Raly) alternative variant fSep08, mRNA.
<a href="#">Raly</a>	<a href="#">Raly.gSep08</a>	<a href="#">296301</a>	33389	294	3	43	hnRNP-associated with lethal yellow (Raly) alternative variant gSep08, mRNA.
<a href="#">ramee</a>	<a href="#">ramee.aSep08</a>		4061	345		114	odd Oz ten-m homolog 2 (ramee) mRNA.
<a href="#">ramer</a>	<a href="#">ramer.aSep08</a>		879	671		55	putative protein (ramer) mRNA.

<a href="#">Ramp2</a>	<a href="#">Ramp2.bSep08</a>	<a href="#">58966</a>	1746	503	4	105	receptor (calcitonin) activity modifying protein 2 (Ramp2) alternative variant bSep08, mRNA.
<a href="#">Ramp2</a>	<a href="#">Ramp2.cSep08</a>	<a href="#">58966</a>	996	866	2	76	receptor (calcitonin) activity modifying protein 2 (Ramp2) alternative variant cSep08, mRNA.
<a href="#">Ramp3</a>	<a href="#">Ramp3.bSep08</a>	<a href="#">56820</a>	14325	558	3	116	receptor (calcitonin) activity modifying protein 3 (13.5 kD) (Ramp3) alternative variant bSep08, mRNA.
<a href="#">Ramp3</a>	<a href="#">Ramp3.cSep08</a>	<a href="#">56820</a>	15299	608	2	63	receptor (calcitonin) activity modifying protein 3 (7.1 kD) (Ramp3) alternative variant cSep08, mRNA.
<a href="#">Ran</a>	<a href="#">Ran.aSep08</a>	<a href="#">84509</a>	2343	777	6	259	RAN, member RAS oncogene family (Ran) alternative variant aSep08, mRNA.
<a href="#">Ran</a>	<a href="#">Ran.bSep08</a>	<a href="#">84509</a>	2954	1104	6	216	RAN, member RAS oncogene family (24.4 kD) (Ran) alternative variant bSep08, mRNA.
<a href="#">Ran</a>	<a href="#">Ran.cSep08</a>	<a href="#">84509</a>	3184	1101	7	216	RAN, member RAS oncogene family (24.4 kD) (Ran) alternative variant cSep08, mRNA.
<a href="#">Ran</a>	<a href="#">Ran.dSep08</a>	<a href="#">84509</a>	4555	2488	7	216	RAN, member RAS oncogene family (24.4 kD) (Ran) alternative variant dSep08, mRNA.
<a href="#">Ran</a>	<a href="#">Ran.eSep08</a>	<a href="#">84509</a>	1934	775	3	115	RAN, member RAS oncogene family (Ran) alternative variant eSep08, mRNA.
<a href="#">Ranbp1</a>	<a href="#">Ranbp1.aSep08</a>	<a href="#">360739</a>	6968	952	1	216	RAN binding protein 1 (Ranbp1) alternative variant aSep08, mRNA.
<a href="#">Ranbp2</a>	<a href="#">Ranbp2.aSep08</a>	<a href="#">294429</a>	18805	5654	10	1529	RAN binding protein 2 (Ranbp2) alternative variant aSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.aSep08</a>	<a href="#">501281</a>	37125	2532	17	558	RAN binding protein 3 like (59.2 kD) (Ranbp3) alternative variant aSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.bSep08</a>	<a href="#">501281</a>	36104	1392	12	369	RAN binding protein 3 like (Ranbp3) alternative variant bSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.cSep08</a>	<a href="#">501281</a>	32630	779	10	255	RAN binding protein 3 like (Ranbp3) alternative variant cSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.dSep08</a>	<a href="#">501281</a>	2915	664	5	220	RAN binding protein 3 like (Ranbp3) alternative variant dSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.eSep08</a>	<a href="#">501281</a>	28176	803	9	211	RAN binding protein 3 like (Ranbp3) alternative variant eSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.fSep08</a>	<a href="#">501281</a>	2749	862	3	177	RAN binding protein 3 like (19.5 kD) (Ranbp3) alternative variant fSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.gSep08</a>	<a href="#">501281</a>	28174	489	8	162	RAN binding protein 3 like (Ranbp3) alternative variant gSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.hSep08</a>	<a href="#">501281</a>	902	810	2	95	putative protein (Ranbp3) alternative variant hSep08, mRNA.
<a href="#">Ranbp3</a>	<a href="#">Ranbp3.iSep08</a>	<a href="#">501281</a>	26086	722	8	87	RAN binding protein 3 like (Ranbp3) alternative variant iSep08, mRNA.
<a href="#">Ranbp3l</a>	<a href="#">Ranbp3l.aSep08</a>	<a href="#">294789</a>	9131	1762	6	425	RAN binding protein 3-like (Ranbp3l) alternative variant aSep08, mRNA.
<a href="#">Ranbp3l</a>	<a href="#">Ranbp3l.bSep08</a>	<a href="#">294789</a>	9205	1683	6	226	RAN binding protein 3-like (24.9 kD) (Ranbp3l) alternative variant bSep08, mRNA.
<a href="#">Ranbp3l</a>	<a href="#">Ranbp3l.cSep08</a>	<a href="#">294789</a>	6657	1458	4	181	RAN binding protein 3-like (Ranbp3l) alternative variant cSep08, mRNA.

<a href="#">Ranbp3l</a>	<a href="#">Ranbp3l.dSep08</a>	<a href="#">294789</a>	6277	527	5	175	RAN binding protein 3-like (Ranbp3l) alternative variant dSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.aSep08</a>	<a href="#">361396</a>	54755	1782	8	594	ran binding protein 10 like (Ranbp10) alternative variant aSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.bSep08</a>	<a href="#">361396</a>	20794	773	7	225	ran binding protein like (Ranbp10) alternative variant bSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.cSep08</a>	<a href="#">361396</a>	48420	597	6	198	ran binding protein like (Ranbp10) alternative variant cSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.dSep08</a>	<a href="#">361396</a>	25445	759	6	172	ran binding protein like (Ranbp10) alternative variant dSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.eSep08</a>	<a href="#">361396</a>	495	398	2	76	ran binding protein 10 like (Ranbp10) alternative variant eSep08, mRNA.
<a href="#">Ranbp10</a>	<a href="#">Ranbp10.fSep08</a>	<a href="#">361396</a>	20377	523	2	68	protein RAN binding like (Ranbp10) alternative variant fSep08, mRNA.
<a href="#">Ranbp17</a>	<a href="#">Ranbp17.bSep08</a>	<a href="#">303029</a>	49554	1801	4	530	RAN binding protein 17 (Ranbp17) alternative variant bSep08, mRNA.
<a href="#">RanBPM_CRA.0</a>	<a href="#">RanBPM_CRA.0.aSep08</a>		2005	571		190	ran binding protein 10 like (RanBPM_CRA.0) mRNA.
<a href="#">Rangap1</a>	<a href="#">Rangap1.bSep08</a>	<a href="#">362965</a>	29421	3982	17	418	RAN GTPase activating protein 1 (45.5 kD) (Rangap1) alternative variant bSep08, mRNA.
<a href="#">Rangap1</a>	<a href="#">Rangap1.cSep08</a>	<a href="#">362965</a>	1115	845	3	156	RAN GTPase activating protein 1 (17.6 kD) (Rangap1) alternative variant cSep08, mRNA.
<a href="#">Rangap1</a>	<a href="#">Rangap1.dSep08</a>	<a href="#">362965</a>	719	537	2	71	RAN GTPase activating protein 1 (Rangap1) alternative variant dSep08, mRNA.
<a href="#">Rangap1</a>	<a href="#">Rangap1.eSep08</a>	<a href="#">362965</a>	7971	707	3	42	RAN GTPase activating protein 1 (Rangap1) alternative variant eSep08, mRNA.
<a href="#">Rangap1</a>	<a href="#">Rangap1.gSep08</a>	<a href="#">362965</a>	9692	346	3	32	putative protein (Rangap1) alternative variant gSep08, mRNA.
<a href="#">Rangrf</a>	<a href="#">Rangrf.bSep08</a>	<a href="#">287419</a>	1256	468	2	110	RAN guanine nucleotide release factor (Rangrf) alternative variant bSep08, mRNA.
<a href="#">Rangrf</a>	<a href="#">Rangrf.cSep08</a>	<a href="#">287419</a>	884	801	1	65	RAN guanine nucleotide release factor (7.0 kD) (Rangrf) alternative variant cSep08, mRNA.
<a href="#">ranoy</a>	<a href="#">ranoy.aSep08</a>		2675	739		63	putative protein (ranoy) mRNA.
<a href="#">RAP.0</a>	<a href="#">RAP.0.aSep08</a>		7885	540		150	protein CRA a (17.2 kD) (RAP.0) mRNA.
<a href="#">Rap1a</a>	<a href="#">Rap1a.aSep08</a>	<a href="#">295347</a>	76317	681	1	193	RAS-related protein 1a (21.8 kD) (Rap1a) alternative variant aSep08, mRNA.
<a href="#">Rap1a</a>	<a href="#">Rap1a.cSep08</a>	<a href="#">295347</a>	76280	760	2	161	RAS-related protein 1a (18.2 kD) (Rap1a) alternative variant cSep08, mRNA.
<a href="#">Rap1b</a>	<a href="#">Rap1b.bSep08</a>	<a href="#">171337</a>	27312	2599	6	184	RAS related protein 1b (20.8 kD) (Rap1b) alternative variant bSep08, mRNA.
<a href="#">Rap1b</a>	<a href="#">Rap1b.cSep08</a>	<a href="#">171337</a>	33046	2685	6	184	RAS related protein 1b (20.8 kD) (Rap1b) alternative variant cSep08, mRNA.
<a href="#">Rap1b</a>	<a href="#">Rap1b.dSep08</a>	<a href="#">171337</a>	5688	713	3	133	RAS related protein 1b (Rap1b) alternative variant dSep08, mRNA.
<a href="#">Rap1b</a>	<a href="#">Rap1b.eSep08</a>	<a href="#">171337</a>	4335	725	2	130	RAS related protein 1b (Rap1b) alternative variant eSep08, mRNA.

<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.aSep08</a>	<a href="#">313644</a>	47130	3296	26	761	rap1 GTPase-activating protein (Rap1GAP) alternative variant aSep08, mRNA.
<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.bSep08</a>	<a href="#">313644</a>	9056	745	9	248	rap1 GTPase-activating protein (Rap1GAP) alternative variant bSep08, mRNA.
<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.cSep08</a>	<a href="#">313644</a>	51680	644	8	214	rap1 GTPase-activating protein (Rap1GAP) alternative variant cSep08, mRNA.
<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.dSep08</a>	<a href="#">313644</a>	7433	604	3	201	rap1 GTPase-activating protein (Rap1GAP) alternative variant dSep08, mRNA.
<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.fSep08</a>	<a href="#">313644</a>	882	562	2	87	rap1 GTPase-activating protein (10.1 kD) (Rap1GAP) alternative variant fSep08, mRNA.
<a href="#">Rap1GAP</a>	<a href="#">Rap1GAP.gSep08</a>	<a href="#">313644</a>	1217	531	2	103	rap1 GTPase-activating protein (Rap1GAP) alternative variant gSep08, mRNA.
<a href="#">Rap1gds1</a>	<a href="#">Rap1gds1.bSep08</a>	<a href="#">310909</a>	119516	528	2	176	RAP1, GTP-GDP dissociation stimulator 1 (Rap1gds1) alternative variant bSep08, mRNA.
<a href="#">Rap2a</a>	<a href="#">Rap2a.aSep08</a>	<a href="#">114560</a>	24401	1553	2	250	RAS related protein 2a? (26.9 kD) (Rap2a) alternative variant aSep08, mRNA.
<a href="#">Rap2ip</a>	<a href="#">Rap2ip.bSep08</a>	<a href="#">303569</a>	9056	1935	11	441	rap2 interacting protein (49.5 kD) (Rap2ip) alternative variant bSep08, mRNA.
<a href="#">Rap2ip</a>	<a href="#">Rap2ip.cSep08</a>	<a href="#">303569</a>	1571	404	3	104	rap2 interacting protein (Rap2ip) alternative variant cSep08, mRNA.
<a href="#">Rap2ip</a>	<a href="#">Rap2ip.eSep08</a>	<a href="#">303569</a>	892	393	3	72	rap2 interacting protein (Rap2ip) alternative variant eSep08, mRNA.
<a href="#">Rapgef1</a>	<a href="#">Rapgef1.aSep08</a>	<a href="#">63881</a>	6246	981	6	249	rap guanine nucleotide exchange factor (GEF) 1 (Rapgef1) alternative variant aSep08, mRNA.
<a href="#">Rapgef1</a>	<a href="#">Rapgef1.bSep08</a>	<a href="#">63881</a>	10884	396	4	131	rap guanine nucleotide exchange factor (GEF) 1 (Rapgef1) alternative variant bSep08, mRNA.
<a href="#">Rapgef2</a>	<a href="#">Rapgef2.bSep08</a>	<a href="#">310533</a>	10516	1147	9	381	rap guanine nucleotide exchange factor (GEF) 2 (Rapgef2) alternative variant bSep08, mRNA.
<a href="#">Rapgef2</a>	<a href="#">Rapgef2.cSep08</a>	<a href="#">310533</a>	8759	645	5	214	rap guanine nucleotide exchange factor (GEF) 2 (Rapgef2) alternative variant cSep08, mRNA.
<a href="#">Rapgef3</a>	<a href="#">Rapgef3.bSep08</a>	<a href="#">59326</a>	6268	518	4	118	rap guanine nucleotide exchange factor (GEF) 3 (Rapgef3) alternative variant bSep08, mRNA.
<a href="#">Rapgef4</a>	<a href="#">Rapgef4.aSep08</a>	<a href="#">252857</a>	60892	705	7	234	rap guanine nucleotide exchange factor (GEF) 4 (Rapgef4) alternative variant aSep08, mRNA.
<a href="#">Rapgef4</a>	<a href="#">Rapgef4.bSep08</a>	<a href="#">252857</a>	203167	671	5	223	rap guanine nucleotide exchange factor (GEF) 4 (Rapgef4) alternative variant bSep08, mRNA.
<a href="#">Rapgef4</a>	<a href="#">Rapgef4.cSep08</a>	<a href="#">252857</a>	88229	538	5	132	rap guanine nucleotide exchange factor (GEF) 4 (Rapgef4) alternative variant cSep08, mRNA.
<a href="#">Rapgef5</a>	<a href="#">Rapgef5.bSep08</a>	<a href="#">362799</a>	13481	527	3	153	rap guanine nucleotide exchange factor 5 (Rapgef5) alternative variant bSep08, mRNA.
<a href="#">Rapgef5</a>	<a href="#">Rapgef5.cSep08</a>	<a href="#">362799</a>	16269	441	4	147	rap guanine nucleotide exchange factor 5 (Rapgef5) alternative variant cSep08, mRNA.
<a href="#">Rapgef5</a>	<a href="#">Rapgef5.eSep08</a>	<a href="#">362799</a>	3044	1193	2	96	putative nuclear protein (10.8 kD) (Rapgef5) alternative variant eSep08, mRNA.
<a href="#">Rapgef5</a>	<a href="#">Rapgef5.fSep08</a>	<a href="#">362799</a>	7978	393	5	85	rap guanine nucleotide exchange factor 5 (Rapgef5) alternative variant fSep08, mRNA.

<a href="#">Rapgef6</a>	<a href="#">Rapgef6.bSep08</a>	<a href="#">303141</a>	10362	345	1	114	rap guanine nucleotide exchange factor (GEF) 6 (Rapgef6) alternative variant bSep08, mRNA.
<a href="#">Rapgef1</a>	<a href="#">Rapgef1.aSep08</a>	<a href="#">303515</a>	2399	916	7	217	rap guanine nucleotide exchange factor -like 1 (Rapgef1) alternative variant aSep08, mRNA.
<a href="#">Rapgef1</a>	<a href="#">Rapgef1.bSep08</a>	<a href="#">303515</a>	2563	2312	2	124	putative secreted or extracellular protein precursor (13.2 kD) (Rapgef1) alternative variant bSep08, mRNA.
<a href="#">Raph1</a>	<a href="#">Raph1.bSep08</a>	<a href="#">363239</a>	5072	468	2	75	ras association (RalGDS/AF-6) and pleckstrin homology domains 1 (8.4 kD) (Raph1) alternative variant bSep08, mRNA.
<a href="#">rapor</a>	<a href="#">rapor.aSep08</a>		24333	2096	9	698	talin 2 (rapor) alternative variant aSep08, mRNA.
<a href="#">rapor</a>	<a href="#">rapor.bSep08</a>		10535	560	4	186	talin 2 (rapor) alternative variant bSep08, mRNA.
<a href="#">rapor</a>	<a href="#">rapor.cSep08</a>		11630	581	5	119	talin 2 (rapor) alternative variant cSep08, mRNA.
<a href="#">rapor</a>	<a href="#">rapor.dSep08</a>		1495	407	2	104	talin 2 (rapor) alternative variant dSep08, mRNA.
<a href="#">Rapsn</a>	<a href="#">Rapsn.bSep08</a>	<a href="#">362161</a>	1258	711	2	208	receptor-associated protein of the synapse (23.1 kD) (Rapsn) alternative variant bSep08, mRNA.
<a href="#">Rapsn</a>	<a href="#">Rapsn.cSep08</a>	<a href="#">362161</a>	710	398	2	69	receptor-associated protein of the synapse (Rapsn) alternative variant cSep08, mRNA.
<a href="#">Rara</a>	<a href="#">Rara.bSep08</a>	<a href="#">24705</a>	3471	1784	2	165	retinoic acid receptor, alpha (Rara) alternative variant bSep08, mRNA.
<a href="#">Rarb</a>	<a href="#">Rarb.aSep08</a>	<a href="#">24706</a>	15876	1728		83	retinoic acid receptor, beta (8.9 kD) (Rarb) mRNA.
<a href="#">rarby</a>	<a href="#">rarby.aSep08</a>		34681	1245	4	48	putative protein (5.7 kD) (rarby) alternative variant aSep08, mRNA.
<a href="#">rarby</a>	<a href="#">rarby.bSep08</a>		51399	669	5	36	putative protein (4.1 kD) (rarby) alternative variant bSep08, mRNA.
<a href="#">rarchy</a>	<a href="#">rarchy.aSep08</a>		1219	244		81	glutamine serine rich 1 (rarchy) mRNA.
<a href="#">rardar</a>	<a href="#">rardar.bSep08</a>		21364	464	2	31	CRA b like (3.7 kD) (rardar) alternative variant bSep08, mRNA.
<a href="#">rarfer</a>	<a href="#">rarfer.aSep08</a>		14302	263		87	armadillo repeat containing 3 (rarfer) mRNA.
<a href="#">rarflo</a>	<a href="#">rarflo.aSep08</a>		6432	686		43	putative protein (4.6 kD) (rarflo) mRNA.
<a href="#">rarflu</a>	<a href="#">rarflu.aSep08</a>		10723	745		63	putative protein (rarflu) mRNA.
<a href="#">rargar</a>	<a href="#">rargar.aSep08</a>		2727	663		139	putative protein (rargar) mRNA.
<a href="#">rarja</a>	<a href="#">rarja.aSep08</a>		512	312		51	putative protein (rarja) mRNA.
<a href="#">rarjey</a>	<a href="#">rarjey.aSep08</a>		4428	439	2	62	putative protein (7.0 kD) (rarjey) alternative variant aSep08, mRNA.
<a href="#">rarjey</a>	<a href="#">rarjey.bSep08</a>		1475	261	2	49	putative protein (rarjey) alternative variant bSep08, mRNA.
<a href="#">rarkee</a>	<a href="#">rarkee.aSep08</a>		8178	987		130	RAB GTPase activating protein 1-like (rarkee) mRNA.
<a href="#">rarkler</a>	<a href="#">rarkler.aSep08</a>		4775	331		30	putative protein (3.6 kD) (rarkler) mRNA.
<a href="#">rarlo</a>	<a href="#">rarlo.aSep08</a>		4333	1681		560	CRA b (rarlo) mRNA.
<a href="#">rarmee</a>	<a href="#">rarmee.aSep08</a>		61252	284		31	putative protein (rarmee) mRNA.
<a href="#">rarmer</a>	<a href="#">rarmer.aSep08</a>		1954	755		145	ring finger protein 213 (rarmer) mRNA.
<a href="#">rarnoy</a>	<a href="#">rarnoy.aSep08</a>		885	454		130	putative protein of metazoan origin (rarnoy) mRNA.
<a href="#">rarpor</a>	<a href="#">rarpor.aSep08</a>		3099	646		215	vacuolar protein sorting 13C (rarpor) mRNA.
<a href="#">Rarres2</a>	<a href="#">Rarres2.bSep08</a>	<a href="#">297073</a>	3048	1046	5	162	retinoic acid receptor responder (tazarotene induced) 2 (18.4 kD) (Rarres2) alternative variant bSep08, mRNA.



<a href="#">Rarres2</a>	<a href="#">Rarres2.cSep08</a>	<a href="#">297073</a>	2998	717	6	162	retinoic acid receptor responder (tazarotene induced) 2 (18.4 kD) (Rarres2) alternative variant cSep08, mRNA.
<a href="#">Rarres2</a>	<a href="#">Rarres2.eSep08</a>	<a href="#">297073</a>	2228	745	2	72	retinoic acid receptor responder (tazarotene induced) 2 (Rarres2) alternative variant eSep08, mRNA.
<a href="#">Rars</a>	<a href="#">Rars.bSep08</a>	<a href="#">287191</a>	6063	605	3	153	arginyl-tRNA synthetase (17.6 kD) (Rars) alternative variant bSep08, complete mRNA.
<a href="#">Rars2</a>	<a href="#">Rars2.bSep08</a>	<a href="#">297969</a>	22369	743	9	160	arginyl-tRNA synthetase 2 mitochondrial (Rars2) alternative variant bSep08, mRNA.
<a href="#">Rars2</a>	<a href="#">Rars2.cSep08</a>	<a href="#">297969</a>	8971	1147	4	126	arginyl-tRNA synthetase-like (14.6 kD) (Rars2) alternative variant cSep08, mRNA.
<a href="#">Rars2</a>	<a href="#">Rars2.dSep08</a>	<a href="#">297969</a>	703	368	2	74	arginyl-tRNA synthetase-like (Rars2) alternative variant dSep08, mRNA.
<a href="#">rarsa</a>	<a href="#">rarsa.aSep08</a>		5298	705		48	putative protein (5.7 kD) (rarsa) mRNA.
<a href="#">rarshee</a>	<a href="#">rarshee.aSep08</a>		14873	470		32	putative protein (rarshee) mRNA.
<a href="#">rartu</a>	<a href="#">rartu.aSep08</a>		3365	747		81	putative protein (9.1 kD) (rartu) mRNA.
<a href="#">rarvar</a>	<a href="#">rarvar.aSep08</a>		944	282		20	putative protein (rarvar) mRNA.
<a href="#">rarwey</a>	<a href="#">rarwey.aSep08</a>		3579	1225		408	uncharacterized protein (rarwey) mRNA.
<a href="#">Ras.1</a>	<a href="#">Ras.1.aSep08</a>		2975	845		162	GTPase Rab37 (Ras.1) mRNA.
<a href="#">rasa</a>	<a href="#">rasa.aSep08</a>		838	732		69	putative protein (7.0 kD) (rasa) mRNA.
<a href="#">Rasa1</a>	<a href="#">Rasa1.bSep08</a>	<a href="#">25676</a>	675	419	2	127	RAS p21 protein activator 1 like (Rasa1) alternative variant bSep08, mRNA.
<a href="#">Rasa1</a>	<a href="#">Rasa1.cSep08</a>	<a href="#">25676</a>	4185	894	4	95	ras p21 protein activator 1 CRA b (10.9 kD) (Rasa1) alternative variant cSep08, mRNA.
<a href="#">Rasa2</a>	<a href="#">Rasa2.bSep08</a>	<a href="#">25597</a>	65170	1802	7	570	RAS p21 protein activator 2 (Rasa2) alternative variant bSep08, mRNA.
<a href="#">Rasa2</a>	<a href="#">Rasa2.cSep08</a>	<a href="#">25597</a>	11794	316	4	56	RAS p21 protein activator 2 (Rasa2) alternative variant cSep08, mRNA.
<a href="#">Rasa3</a>	<a href="#">Rasa3.bSep08</a>	<a href="#">29372</a>	26753	3458	17	624	RAS p21 protein activator 3 (71.6 kD) (Rasa3) alternative variant bSep08, mRNA.
<a href="#">Rasa3</a>	<a href="#">Rasa3.cSep08</a>	<a href="#">29372</a>	8980	1802	8	564	RAS p21 protein activator 3 (Rasa3) alternative variant cSep08, mRNA.
<a href="#">Rasa3</a>	<a href="#">Rasa3.dSep08</a>	<a href="#">29372</a>	21682	1776	9	331	RAS p21 protein activator 3 (Rasa3) alternative variant dSep08, mRNA.
<a href="#">Rasal1</a>	<a href="#">Rasal1.cSep08</a>	<a href="#">360814</a>	6514	441			
<a href="#">Rasal2</a>	<a href="#">Rasal2.bSep08</a>	<a href="#">304893</a>	7827	1799	4	599	RAS protein activator like 2 (Rasal2) alternative variant bSep08, mRNA.
<a href="#">Rasal2</a>	<a href="#">Rasal2.cSep08</a>	<a href="#">304893</a>	76176	828	6	275	RAS protein activator like 2 (Rasal2) alternative variant cSep08, mRNA.
<a href="#">Rasd1</a>	<a href="#">Rasd1.aSep08</a>	<a href="#">64455</a>	1771	1634		280	RAS, dexamethasone-induced 1 (31.7 kD) (Rasd1) mRNA.
<a href="#">RasGAP_C.0</a>	<a href="#">RasGAP_C.0.aSep08</a>		2284	728	2	186	IQ motif containing GTPase activating protein 3 (RasGAP_C.0) alternative variant aSep08, mRNA.
<a href="#">RasGAP_C.0</a>	<a href="#">RasGAP_C.0.bSep08</a>		2652	502	2	166	IQ motif containing GTPase activating protein 3 (RasGAP_C.0) alternative variant bSep08, mRNA.
<a href="#">RasGEF.0</a>	<a href="#">RasGEF.0.aSep08</a>		20365	762		253	ral guanine nucleotide dissociation stimulator-like 1 CRA d (RasGEF.0) mRNA.

<a href="#">RasGEF.1</a>	<a href="#">RasGEF.1.aSep08</a>		4053	897		243	domain-containing family member ras-gef 1 (RasGEF.1) mRNA.
<a href="#">Rasgef1a</a>	<a href="#">Rasgef1a.aSep08</a>	<a href="#">312664</a>	16502	388		129	RasGEF domain family, member 1A (Rasgef1a) mRNA.
<a href="#">Rasgef1c</a>	<a href="#">Rasgef1c.bSep08</a>	<a href="#">360519</a>	1520	763	1	39	RasGEF domain family, member 1C (Rasgef1c) alternative variant bSep08, mRNA.
<a href="#">RasGEF_N.0</a>	<a href="#">RasGEF_N.0.aSep08</a>		10844	425		100	RasGEF domain family member 1B (RasGEF_N.0) mRNA.
<a href="#">Rasgrf1</a>	<a href="#">Rasgrf1.aSep08</a>	<a href="#">192213</a>	35033	1581	12	418	RAS protein-specific guanine nucleotide-releasing factor 1 (Rasgrf1) alternative variant aSep08, mRNA.
<a href="#">Rasgrf2</a>	<a href="#">Rasgrf2.bSep08</a>	<a href="#">114513</a>	1000	426	2	141	RAS protein-specific guanine nucleotide-releasing factor 2 (Rasgrf2) alternative variant bSep08, mRNA.
<a href="#">Rasgrf2</a>	<a href="#">Rasgrf2.cSep08</a>	<a href="#">114513</a>	13304	642	2	127	RAS protein-specific guanine nucleotide-releasing factor 2 (Rasgrf2) alternative variant cSep08, mRNA.
<a href="#">Rasgrf2</a>	<a href="#">Rasgrf2.dSep08</a>	<a href="#">114513</a>	4782	1366	3	95	RAS protein-specific guanine nucleotide-releasing factor 2 (Rasgrf2) alternative variant dSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.bSep08</a>	<a href="#">361714</a>	9955	1223	8	273	RAS guanyl releasing protein 2 (Rasgrp2) alternative variant bSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.cSep08</a>	<a href="#">361714</a>	2554	888	4	145	RAS guanyl releasing protein 2 (Rasgrp2) alternative variant cSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.dSep08</a>	<a href="#">361714</a>	3799	587	5	141	RAS guanyl releasing protein 2 (16.1 kD) (Rasgrp2) alternative variant dSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.eSep08</a>	<a href="#">361714</a>	4705	623	5	141	RAS guanyl releasing protein 2 (16.1 kD) (Rasgrp2) alternative variant eSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.fSep08</a>	<a href="#">361714</a>	2010	572	4	141	RAS guanyl releasing protein 2 (16.1 kD) (Rasgrp2) alternative variant fSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.gSep08</a>	<a href="#">361714</a>	1543	510	4	110	RAS guanyl releasing protein 2 (Rasgrp2) alternative variant gSep08, mRNA.
<a href="#">Rasgrp2</a>	<a href="#">Rasgrp2.hSep08</a>	<a href="#">361714</a>	953	748	2	99	putative protein (Rasgrp2) alternative variant hSep08, mRNA.
<a href="#">Rasgrp3</a>	<a href="#">Rasgrp3.bSep08</a>	<a href="#">313874</a>	12348	707	5	192	RAS, guanyl releasing protein 3 (Rasgrp3) alternative variant bSep08, mRNA.
<a href="#">Rasgrp3</a>	<a href="#">Rasgrp3.cSep08</a>	<a href="#">313874</a>	10409	407	4	135	RAS, guanyl releasing protein 3 (Rasgrp3) alternative variant cSep08, mRNA.
<a href="#">Rasgrp3</a>	<a href="#">Rasgrp3.dSep08</a>	<a href="#">313874</a>	65923	729	5	120	RAS, guanyl releasing protein 3 (Rasgrp3) alternative variant dSep08, mRNA.
<a href="#">Rasgrp4</a>	<a href="#">Rasgrp4.cSep08</a>	<a href="#">170668</a>	420	348	2	58	RAS guanyl releasing protein 4 (6.2 kD) (Rasgrp4) alternative variant cSep08, mRNA.
<a href="#">rashee</a>	<a href="#">rashee.aSep08</a>		3552	324		108	mediator of rna polymerase ii transcription (rashee) mRNA.
<a href="#">Rasip1</a>	<a href="#">Rasip1.bSep08</a>	<a href="#">292912</a>	904	739	2	132	ras interacting protein 1 (Rasip1) alternative variant bSep08, mRNA.
<a href="#">Rasl10b</a>	<a href="#">Rasl10b.aSep08</a>	<a href="#">303382</a>	9931	1251		266	RAS-like, family 10, member B (Rasl10b) mRNA.
<a href="#">Rasl11a</a>	<a href="#">Rasl11a.bSep08</a>	<a href="#">304268</a>	2315	740	1	246	RAS-like family 11 member A (Rasl11a) alternative variant bSep08, mRNA.
<a href="#">Rasl12</a>	<a href="#">Rasl12.bSep08</a>	<a href="#">315762</a>	662	402	1	87	RAS-like, family 12 (Rasl12) alternative variant bSep08, mRNA.
<a href="#">Rasl12</a>	<a href="#">Rasl12.cSep08</a>	<a href="#">315762</a>	3130	313	2	86	RAS-like, family 12 (Rasl12) alternative variant cSep08, mRNA.

<a href="#">Rassf1</a>	<a href="#">Rassf1.cSep08</a>	<a href="#">363140</a>	6470	966	3	193	ras association (RalGDS/AF-6) domain family member 1 (22.1 kD) (Rassf1) alternative variant cSep08, mRNA.
<a href="#">Rassf5</a>	<a href="#">Rassf5.bSep08</a>	<a href="#">54355</a>	18842	709	3	167	ras association (RalGDS/AF-6) domain family member 5 (Rassf5) alternative variant bSep08, mRNA.
<a href="#">Rassf7</a>	<a href="#">Rassf7.bSep08</a>	<a href="#">293623</a>	1753	1116	3	279	ras association (RalGDS/AF-6) domain family (N-terminal) member 7 (Rassf7) alternative variant bSep08, mRNA.
<a href="#">Rassf8</a>	<a href="#">Rassf8.aSep08</a>	<a href="#">312846</a>	26682	940	5	199	CRA b like (Rassf8) alternative variant aSep08, mRNA.
<a href="#">Rassf8</a>	<a href="#">Rassf8.bSep08</a>	<a href="#">312846</a>	73704	673	3	132	CRA a (Rassf8) alternative variant bSep08, mRNA.
<a href="#">Rassf8</a>	<a href="#">Rassf8.cSep08</a>	<a href="#">312846</a>	69822	432	3	94	RA (Rassf8) alternative variant cSep08, mRNA.
<a href="#">ratu</a>	<a href="#">ratu.aSep08</a>		98638	720		122	putative nuclear protein (14.0 kD) (ratu) mRNA.
<a href="#">ravar</a>	<a href="#">ravar.aSep08</a>		5942	1759		408	uromodulin-like (ravar) mRNA.
<a href="#">Raver1</a>	<a href="#">Raver1.bSep08</a>	<a href="#">298705</a>	1948	748	6	249	ribonucleoprotein, PTB-binding 1 (Raver1) alternative variant bSep08, mRNA.
<a href="#">Raver1</a>	<a href="#">Raver1.dSep08</a>	<a href="#">298705</a>	1913	732	2	91	ribonucleoprotein, PTB-binding 1 (9.8 kD) (Raver1) alternative variant dSep08, mRNA.
<a href="#">Raver2</a>	<a href="#">Raver2.aSep08</a>	<a href="#">362551</a>	23418	1385	6	420	ribonucleoprotein, PTB-binding 2 (Raver2) alternative variant aSep08, mRNA.
<a href="#">Raver2</a>	<a href="#">Raver2.bSep08</a>	<a href="#">362551</a>	18494	1303	7	315	ribonucleoprotein, PTB-binding 2 (Raver2) alternative variant bSep08, mRNA.
<a href="#">rawby</a>	<a href="#">rawby.aSep08</a>		108128	386		79	putative protein (rawby) alternative variant aSep08, mRNA.
<a href="#">rawchy</a>	<a href="#">rawchy.aSep08</a>		5544	383		76	putative protein (8.8 kD) (rawchy) mRNA.
<a href="#">rawdardar</a>	<a href="#">rawdardar.aSep08</a>		1426	639		198	repeat-containing protein 16c (rawdardar) mRNA.
<a href="#">rawey</a>	<a href="#">rawey.aSep08</a>		1180	863	2	45	putative protein (5.2 kD) (rawey) alternative variant aSep08, mRNA.
<a href="#">rawfer</a>	<a href="#">rawfer.aSep08</a>		13842	826		227	armadillo repeat containing 3 CRA b (rawfer) alternative variant aSep08, mRNA.
<a href="#">rawfer</a>	<a href="#">rawfer.bSep08</a>		6768	494		124	armadillo repeat containing 3 CRA b (rawfer) alternative variant bSep08, mRNA.
<a href="#">rawflo</a>	<a href="#">rawflo.aSep08</a>		450	332		51	putative protein (rawflo) mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.aSep08</a>		24567	729	7	242	protein kinase A anchoring Rt31 (rawflu) alternative variant aSep08, mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.bSep08</a>		17661	688	6	229	protein kinase A anchoring Rt31 (rawflu) alternative variant bSep08, mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.cSep08</a>		17576	669	7	223	protein kinase A anchoring Rt31 (rawflu) alternative variant cSep08, mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.dSep08</a>		11338	645	2	177	protein kinase A anchoring Rt31 (rawflu) alternative variant dSep08, mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.eSep08</a>		9180	864	1	155	protein kinase A anchoring Rt31 (rawflu) alternative variant eSep08, mRNA.
<a href="#">rawflu</a>	<a href="#">rawflu.fSep08</a>		32096	789	1	110	protein kinase A anchoring Rt31 (rawflu) alternative variant fSep08, mRNA.
<a href="#">rawgar</a>	<a href="#">rawgar.aSep08</a>		3004	794		210	putative protein (rawgar) mRNA.
<a href="#">rawja</a>	<a href="#">rawja.aSep08</a>		13563	506		50	putative protein (5.7 kD) (rawja) mRNA.
<a href="#">rawjey</a>	<a href="#">rawjey.aSep08</a>		83063	246		55	putative protein (rawjey) mRNA.
<a href="#">rawkee</a>	<a href="#">rawkee.aSep08</a>		2325	727		31	putative protein (3.5 kD) (rawkee) mRNA.

<a href="#">rawkler</a>	<a href="#">rawkler.aSep08</a>		3659	404		62	putative protein (rawkler) mRNA.
<a href="#">rawlo</a>	<a href="#">rawlo.aSep08</a>		1999	507		168	putative protein of ancient origin (rawlo) mRNA.
<a href="#">rawmee</a>	<a href="#">rawmee.aSep08</a>		9552	995	5	77	putative mitochondrial protein (8.7 kD) (rawmee) alternative variant aSep08, mRNA.
<a href="#">rawmer</a>	<a href="#">rawmer.aSep08</a>		10311	1283		427	ring finger protein 213 (rawmer) mRNA.
<a href="#">rawnoy</a>	<a href="#">rawnoy.aSep08</a>		1622	397		67	putative protein (rawnoy) mRNA.
<a href="#">rawpor</a>	<a href="#">rawpor.aSep08</a>		12884	687		139	vacuolar protein sorting 13C CRA a (rawpor) mRNA.
<a href="#">rawsa</a>	<a href="#">rawsa.aSep08</a>		670	393			
<a href="#">rawshee</a>	<a href="#">rawshee.aSep08</a>		2847	232		44	putative protein (5.2 kD) (rawshee) mRNA.
<a href="#">rawtu</a>	<a href="#">rawtu.aSep08</a>		1303	875	1	95	phosphatase 1G (rawtu) alternative variant aSep08, mRNA.
<a href="#">rawtu</a>	<a href="#">rawtu.bSep08</a>		1340	475	2	90	phosphatase 1G (rawtu) alternative variant bSep08, mRNA.
<a href="#">rawvar</a>	<a href="#">rawvar.aSep08</a>		1479	709		118	putative protein (rawvar) mRNA.
<a href="#">rawwey</a>	<a href="#">rawwey.aSep08</a>		2138	355	3	61	putative protein of vertebrate origin (rawwey) mRNA.
<a href="#">Rax</a>	<a href="#">Rax.aSep08</a>	<a href="#">114213</a>	2929	504		167	retina and anterior neural fold homeobox (Rax) mRNA.
<a href="#">Rb1</a>	<a href="#">Rb1.aSep08</a>	<a href="#">24708</a>	133015	4490	27	910	retinoblastoma 1 (Rb1) alternative variant aSep08, mRNA.
<a href="#">Rb1</a>	<a href="#">Rb1.bSep08</a>	<a href="#">24708</a>	27260	512	5	170	retinoblastoma-associated protein (Rb1) alternative variant bSep08, mRNA.
<a href="#">Rb1</a>	<a href="#">Rb1.cSep08</a>	<a href="#">24708</a>	24207	494	4	106	retinoblastoma-associated protein (Rb1) alternative variant cSep08, mRNA.
<a href="#">Rb1</a>	<a href="#">Rb1.dSep08</a>	<a href="#">24708</a>	4124	735	3	92	retinoblastoma 1 (Rb1) alternative variant dSep08, mRNA.
<a href="#">Rb1cc1</a>	<a href="#">Rb1cc1.cSep08</a>	<a href="#">312927</a>	3087	427	3	80	RB1-inducible coiled-coil 1 (Rb1cc1) alternative variant cSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.bSep08</a>	<a href="#">313658</a>	15502	1785	4	522	ZUBR1 (Rbaf600) alternative variant bSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.cSep08</a>	<a href="#">313658</a>	21377	1791	12	478	ZUBR1 (Rbaf600) alternative variant cSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.dSep08</a>	<a href="#">313658</a>	3156	676	5	225	ZUBR1 (Rbaf600) alternative variant dSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.eSep08</a>	<a href="#">313658</a>	5048	617	5	205	ZUBR1 (Rbaf600) alternative variant eSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.fSep08</a>	<a href="#">313658</a>	3523	772	2	135	ZUBR1 (Rbaf600) alternative variant fSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.gSep08</a>	<a href="#">313658</a>	1053	423	3	123	ZUBR1 (Rbaf600) alternative variant gSep08, mRNA.
<a href="#">Rbaf600</a>	<a href="#">Rbaf600.hSep08</a>	<a href="#">313658</a>	1357	543	2	81	ZUBR1 (Rbaf600) alternative variant hSep08, mRNA.
<a href="#">Rbak</a>	<a href="#">Rbak.aSep08</a>	<a href="#">288489</a>	11419	1026		341	RB-associated KRAB repressor (Rbak) mRNA.
<a href="#">Rbbp5</a>	<a href="#">Rbbp5.bSep08</a>	<a href="#">304794</a>	8871	703	5	174	retinoblastoma binding protein 5 (Rbbp5) alternative variant bSep08, mRNA.
<a href="#">Rbbp5</a>	<a href="#">Rbbp5.cSep08</a>	<a href="#">304794</a>	500	414	2	71	retinoblastoma binding protein 5 (Rbbp5) alternative variant cSep08, mRNA.
<a href="#">Rbbp5</a>	<a href="#">Rbbp5.dSep08</a>	<a href="#">304794</a>	7545	416	2	49	retinoblastoma binding protein 5 (Rbbp5) alternative variant dSep08, mRNA.
<a href="#">Rbbp6</a>	<a href="#">Rbbp6.aSep08</a>	<a href="#">308968</a>	3685	3357		977	retinoblastoma binding protein 6 (Rbbp6) mRNA.
<a href="#">Rbbp7</a>	<a href="#">Rbbp7.bSep08</a>	<a href="#">83712</a>	13124	965	6	208	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (Rbbp7) alternative variant bSep08, mRNA.
<a href="#">Rbbp7</a>	<a href="#">Rbbp7.bSep08</a>	<a href="#">688840</a>	13124	965	6	208	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (Rbbp7) alternative variant bSep08, mRNA.

<a href="#">Rbbp7</a>	<a href="#">Rbbp7.cSep08</a>	<a href="#">83712</a>	2853	694	3	89	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (10.4 kD) (Rbbp7) alternative variant cSep08, mRNA.
<a href="#">Rbbp7</a>	<a href="#">Rbbp7.cSep08</a>	<a href="#">688840</a>	2853	694	3	89	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (10.4 kD) (Rbbp7) alternative variant cSep08, mRNA.
<a href="#">Rbbp7</a>	<a href="#">Rbbp7.dSep08</a>	<a href="#">83712</a>	3073	861	3	71	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (Rbbp7) alternative variant dSep08, mRNA.
<a href="#">Rbbp7</a>	<a href="#">Rbbp7.dSep08</a>	<a href="#">688840</a>	3073	861	3	71	retinoblastoma binding protein 7 and hypothetical protein LOC688840 (Rbbp7) alternative variant dSep08, mRNA.
<a href="#">Rbbp9</a>	<a href="#">Rbbp9.cSep08</a>	<a href="#">29459</a>	1380	1029	2	90	retinoblastoma binding protein 9 (10.3 kD) (Rbbp9) alternative variant cSep08, mRNA.
<a href="#">Rbck1</a>	<a href="#">Rbck1.bSep08</a>	<a href="#">60383</a>	13961	1151	9	329	zinc finger, RanBP2-type (Rbck1) alternative variant bSep08, mRNA.
<a href="#">Rbck1</a>	<a href="#">Rbck1.cSep08</a>	<a href="#">60383</a>	3357	1334	4	236	putative protein of eukaryotic origin (26.4 kD) (Rbck1) alternative variant cSep08, mRNA.
<a href="#">Rbck1</a>	<a href="#">Rbck1.dSep08</a>	<a href="#">60383</a>	7306	633	4	181	putative protein of metazoan origin (Rbck1) alternative variant dSep08, mRNA.
<a href="#">RBD-FIP.0</a>	<a href="#">RBD-FIP.0.aSep08</a>		6182	3141		376	rab11 family interacting protein 5 (RBD-FIP.0) mRNA.
<a href="#">RBD.0</a>	<a href="#">RBD.0.aSep08</a>		22344	646		215	T-cell lymphoma invasion metastasis 1 like (RBD.0) alternative variant aSep08, mRNA.
<a href="#">RBD.0</a>	<a href="#">RBD.0.bSep08</a>		15975	513		115	T-cell lymphoma invasion metastasis 1 like (RBD.0) alternative variant bSep08, mRNA.
<a href="#">Rbed1</a>	<a href="#">Rbed1.aSep08</a>	<a href="#">297342</a>	25762	1037	9	246	RNA binding motif and ELMO domain 1 (Rbed1) alternative variant aSep08, mRNA.
<a href="#">Rbed1</a>	<a href="#">Rbed1.bSep08</a>	<a href="#">297342</a>	22639	762	7	137	RNA binding motif and ELMO domain 1 (Rbed1) alternative variant bSep08, mRNA.
<a href="#">Rbed1</a>	<a href="#">Rbed1.cSep08</a>	<a href="#">297342</a>	19848	578	6	74	RNA binding motif and ELMO domain 1 (8.0 kD) (Rbed1) alternative variant cSep08, mRNA.
<a href="#">Rbj</a>	<a href="#">Rbj.bSep08</a>	<a href="#">298859</a>	16241	684	5	169	dnaj homolog subfamily C member 27 (Rbj) alternative variant bSep08, mRNA.
<a href="#">Rbks</a>	<a href="#">Rbks.bSep08</a>	<a href="#">362706</a>	31207	469	4	134	ribokinase (Rbks) alternative variant bSep08, mRNA.
<a href="#">Rbks</a>	<a href="#">Rbks.cSep08</a>	<a href="#">362706</a>	23499	818	2	121	ribokinase (Rbks) alternative variant cSep08, mRNA.
<a href="#">Rbks</a>	<a href="#">Rbks.dSep08</a>	<a href="#">362706</a>	8411	290	2	74	ribokinase (Rbks) alternative variant dSep08, mRNA.
<a href="#">Rbl1</a>	<a href="#">Rbl1.aSep08</a>	<a href="#">680111</a>	23492	1174		338	retinoblastoma-like 1 (p107) (Rbl1) mRNA.
<a href="#">Rbl2</a>	<a href="#">Rbl2.bSep08</a>	<a href="#">81758</a>	14922	2787	8	401	retinoblastoma-like 2 (Rbl2) alternative variant bSep08, mRNA.
<a href="#">Rbl2</a>	<a href="#">Rbl2.cSep08</a>	<a href="#">81758</a>	6849	719	5	239	retinoblastoma-like 2 (Rbl2) alternative variant cSep08, mRNA.
<a href="#">Rbl2</a>	<a href="#">Rbl2.dSep08</a>	<a href="#">81758</a>	8206	627	4	172	retinoblastoma-like 2 (Rbl2) alternative variant dSep08, mRNA.
<a href="#">Rbl2</a>	<a href="#">Rbl2.eSep08</a>	<a href="#">81758</a>	1173	727	2	117	retinoblastoma-like 2 (Rbl2) alternative variant eSep08, mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.aSep08</a>	<a href="#">114488</a>	6521	1799	7	456	RNA binding motif protein 3 (Rbm3) alternative variant aSep08, complete mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.bSep08</a>	<a href="#">114488</a>	3983	1735	7	189	RNA binding motif protein 3 (Rbm3) alternative variant bSep08, mRNA.

<a href="#">Rbm3</a>	<a href="#">Rbm3.dSep08</a>	<a href="#">114488</a>	3102	847	7	185	RNA binding motif protein 3 (Rbm3) alternative variant dSep08, mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.eSep08</a>	<a href="#">114488</a>	3023	772	7	155	RNA binding motif protein 3 (16.8 kD) (Rbm3) alternative variant eSep08, mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.fSep08</a>	<a href="#">114488</a>	3278	1292	8	118	RNA binding motif protein 3 (12.8 kD) (Rbm3) alternative variant fSep08, complete mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.gSep08</a>	<a href="#">114488</a>	1324	561	4	111	RNA binding motif protein 3 (11.9 kD) (Rbm3) alternative variant gSep08, mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.hSep08</a>	<a href="#">114488</a>	3431	1815	6	94	RNA binding motif protein 3 (10.0 kD) (Rbm3) alternative variant hSep08, complete mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.iSep08</a>	<a href="#">114488</a>	1375	783	4	88	RNA binding motif protein 3 (Rbm3) alternative variant iSep08, mRNA.
<a href="#">Rbm3</a>	<a href="#">Rbm3.kSep08</a>	<a href="#">114488</a>	1356	787	2	94	RNA binding motif protein 3 (10.6 kD) (Rbm3) alternative variant kSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.aSep08</a>	<a href="#">293663</a>	27506	2049	5	365	RNA binding motif protein 4B like (40.4 kD) (Rbm4) alternative variant aSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.bSep08</a>	<a href="#">293663</a>	6458	908	3	228	CRA b (Rbm4) alternative variant bSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.cSep08</a>	<a href="#">293663</a>	3842	3075	2	213	RNA binding motif protein 14 like (22.7 kD) (Rbm4) alternative variant cSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.eSep08</a>	<a href="#">293663</a>	9043	872	3	143	RNA binding motif protein 4B like (16.1 kD) (Rbm4) alternative variant eSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.fSep08</a>	<a href="#">293663</a>	27346	578	2	112	RNA binding motif protein 14 like (Rbm4) alternative variant fSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.gSep08</a>	<a href="#">293663</a>	58655	893	3	109	RNA binding motif protein 14 like (Rbm4) alternative variant gSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.hSep08</a>	<a href="#">293663</a>	18579	1306	4	100	putative membrane protein (10.7 kD) (Rbm4) alternative variant hSep08, mRNA.
<a href="#">Rbm4</a>	<a href="#">Rbm4.jSep08</a>	<a href="#">293663</a>	1400	915	2	28	putative protein (Rbm4) alternative variant jSep08, mRNA.
<a href="#">Rbm4b</a>	<a href="#">Rbm4b.bSep08</a>	<a href="#">474154</a>	1486	871	2	172	RNA binding motif protein 4B (19.7 kD) (Rbm4b) alternative variant bSep08, mRNA.
<a href="#">Rbm4b</a>	<a href="#">Rbm4b.cSep08</a>	<a href="#">474154</a>	2601	1833	3	7	RNA binding motif protein 4B (0.8 kD) (Rbm4b) alternative variant cSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.bSep08</a>	<a href="#">300996</a>	23193	1752	18	537	RNA binding motif protein 5 like (Rbm5) alternative variant bSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.cSep08</a>	<a href="#">300996</a>	10004	1692	5	146	RNA binding motif protein 5 like (17.4 kD) (Rbm5) alternative variant cSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.dSep08</a>	<a href="#">300996</a>	3001	374	4	124	RNA binding motif protein 5 like (Rbm5) alternative variant dSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.eSep08</a>	<a href="#">300996</a>	5047	846	4	122	RNA binding motif protein 5 like (14.6 kD) (Rbm5) alternative variant eSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.fSep08</a>	<a href="#">300996</a>	5620	1417	4	100	RNA binding motif protein 5 like (11.4 kD) (Rbm5) alternative variant fSep08, mRNA.
<a href="#">Rbm5</a>	<a href="#">Rbm5.hSep08</a>	<a href="#">300996</a>	3055	439	3	59	RNA binding motif protein 5 like (Rbm5) alternative variant hSep08, mRNA.
<a href="#">Rbm6</a>	<a href="#">Rbm6.bSep08</a>	<a href="#">315997</a>	82144	724	6	185	RNA binding motif protein 6 (Rbm6) alternative variant bSep08, mRNA.

<a href="#">Rbm6</a>	<a href="#">Rbm6.eSep08</a>	<a href="#">315997</a>	19671	460	4	79	RNA binding motif protein 6 (Rbm6) alternative variant eSep08, mRNA.
<a href="#">Rbm7</a>	<a href="#">Rbm7.aSep08</a>	<a href="#">315634</a>	6350	1736	3	264	RNA binding motif protein 7 like (Rbm7) alternative variant aSep08, mRNA.
<a href="#">Rbm7</a>	<a href="#">Rbm7.cSep08</a>	<a href="#">315634</a>	1822	414	2	79	RNA binding motif protein 7 like (8.8 kD) (Rbm7) alternative variant cSep08, mRNA.
<a href="#">Rbm8</a>	<a href="#">Rbm8.aSep08</a>	<a href="#">295284</a>	2749	1468	3	174	RNA binding motif protein 8 (19.9 kD) (Rbm8) alternative variant aSep08, complete mRNA.
<a href="#">Rbm8</a>	<a href="#">Rbm8.bSep08</a>	<a href="#">295284</a>	2177	893	3	173	RNA binding motif protein 8 (19.8 kD) (Rbm8) alternative variant bSep08, mRNA.
<a href="#">Rbm8</a>	<a href="#">Rbm8.cSep08</a>	<a href="#">295284</a>	1985	567	2	142	RNA binding motif protein 8 (16.4 kD) (Rbm8) alternative variant cSep08, complete mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.aSep08</a>	<a href="#">362950</a>	241334	2498	14	506	RNA binding motif protein 9 (53.3 kD) (Rbm9) alternative variant aSep08, complete mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.cSep08</a>	<a href="#">362950</a>	70427	1143	10	380	RNA binding motif protein 9 (Rbm9) alternative variant cSep08, mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.dSep08</a>	<a href="#">362950</a>	103955	729	5	178	RNA binding motif protein 9 (Rbm9) alternative variant dSep08, mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.eSep08</a>	<a href="#">362950</a>	142985	570	4	169	RNA binding motif protein 9 (18.6 kD) (Rbm9) alternative variant eSep08, mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.fSep08</a>	<a href="#">362950</a>	48627	499	3	165	RNA binding motif protein 9 (Rbm9) alternative variant fSep08, mRNA.
<a href="#">Rbm9</a>	<a href="#">Rbm9.gSep08</a>	<a href="#">362950</a>	154741	714	7	162	RNA binding motif protein 9 (Rbm9) alternative variant gSep08, mRNA.
<a href="#">Rbm10</a>	<a href="#">Rbm10.aSep08</a>	<a href="#">64510</a>	4866	1957	14	541	RNA binding motif protein 10 like (59.5 kD) (Rbm10) alternative variant aSep08, mRNA.
<a href="#">Rbm10</a>	<a href="#">Rbm10.cSep08</a>	<a href="#">64510</a>	26973	1468	10	263	RNA binding motif protein 10 like (Rbm10) alternative variant cSep08, mRNA.
<a href="#">Rbm10</a>	<a href="#">Rbm10.dSep08</a>	<a href="#">64510</a>	380	265	2	86	putative protein (Rbm10) alternative variant dSep08, mRNA.
<a href="#">Rbm11</a>	<a href="#">Rbm11.aSep08</a>	<a href="#">288321</a>	8089	711	1	214	RNA binding motif protein 11 (Rbm11) alternative variant aSep08, mRNA.
<a href="#">Rbm12</a>	<a href="#">Rbm12.bSep08</a>	<a href="#">652928</a>	33705	591	5	196	RNA binding motif protein 12 (Rbm12) alternative variant bSep08, mRNA.
<a href="#">Rbm12</a>	<a href="#">Rbm12.cSep08</a>	<a href="#">652928</a>	33226	385	4	61	RNA binding motif protein 12 (Rbm12) alternative variant cSep08, mRNA.
<a href="#">Rbm13</a>	<a href="#">Rbm13.bSep08</a>	<a href="#">306526</a>	5772	1251	6	229	RNA binding motif protein 13 (27.5 kD) (Rbm13) alternative variant bSep08, mRNA.
<a href="#">Rbm13</a>	<a href="#">Rbm13.cSep08</a>	<a href="#">306526</a>	1611	359	3	14	RNA binding motif protein 13 (1.5 kD) (Rbm13) alternative variant cSep08, mRNA.
<a href="#">Rbm16</a>	<a href="#">Rbm16.bSep08</a>	<a href="#">245926</a>	53178	2546	2	479	RNA binding motif protein 16 (Rbm16) alternative variant bSep08, mRNA.
<a href="#">Rbm17</a>	<a href="#">Rbm17.bSep08</a>	<a href="#">291295</a>	1900	363	2	120	RNA binding motif protein 17 like (Rbm17) alternative variant bSep08, mRNA.
<a href="#">Rbm17</a>	<a href="#">Rbm17.cSep08</a>	<a href="#">291295</a>	4957	461	4	97	RNA binding motif protein 17 like (Rbm17) alternative variant cSep08, mRNA.

<a href="#">Rbm17</a>	<a href="#">Rbm17.dSep08</a>	<a href="#">291295</a>	2421	365	2	80	RNA binding motif protein 17 like (Rbm17) alternative variant dSep08, mRNA.
<a href="#">Rbm17</a>	<a href="#">Rbm17.eSep08</a>	<a href="#">291295</a>	2147	996	3	69	RNA binding motif protein 17 like (Rbm17) alternative variant eSep08, mRNA.
<a href="#">Rbm18</a>	<a href="#">Rbm18.bSep08</a>	<a href="#">311902</a>	18243	747	5	180	RNA binding motif protein 18 (Rbm18) alternative variant bSep08, mRNA.
<a href="#">Rbm18</a>	<a href="#">Rbm18.cSep08</a>	<a href="#">311902</a>	10666	726	3	100	RNA binding motif protein 18 (11.6 kD) (Rbm18) alternative variant cSep08, mRNA.
<a href="#">Rbm18</a>	<a href="#">Rbm18.dSep08</a>	<a href="#">311902</a>	4863	540	3	62	RNA binding motif protein 18 (6.7 kD) (Rbm18) alternative variant dSep08, mRNA.
<a href="#">Rbm19</a>	<a href="#">Rbm19.bSep08</a>	<a href="#">304512</a>	4936	531	3	177	RNA binding motif protein 19 (Rbm19) alternative variant bSep08, mRNA.
<a href="#">Rbm20</a>	<a href="#">Rbm20.bSep08</a>	<a href="#">309544</a>	14470	816	2	171	RNA binding motif protein 20 (Rbm20) alternative variant bSep08, mRNA.
<a href="#">Rbm20</a>	<a href="#">Rbm20.cSep08</a>	<a href="#">309544</a>	36545	425	1	86	RNA binding motif protein 20 (Rbm20) alternative variant cSep08, mRNA.
<a href="#">Rbm22</a>	<a href="#">Rbm22.bSep08</a>	<a href="#">307410</a>	423	283	2	94	RNA binding motif protein 22 (Rbm22) alternative variant bSep08, mRNA.
<a href="#">Rbm25</a>	<a href="#">Rbm25.bSep08</a>	<a href="#">366693</a>	6795	1383	5	273	RNA binding motif protein 25 (Rbm25) alternative variant bSep08, mRNA.
<a href="#">Rbm25</a>	<a href="#">Rbm25.cSep08</a>	<a href="#">366693</a>	6170	1136	4	178	RNA binding motif protein 25 (20.4 kD) (Rbm25) alternative variant cSep08, mRNA.
<a href="#">Rbm25</a>	<a href="#">Rbm25.fSep08</a>	<a href="#">366693</a>	3479	584	4	45	RNA binding motif protein 25 (Rbm25) alternative variant fSep08, mRNA.
<a href="#">Rbm27</a>	<a href="#">Rbm27.bSep08</a>	<a href="#">361317</a>	25916	1126	7	323	RNA binding motif protein 27 (Rbm27) alternative variant bSep08, mRNA.
<a href="#">Rbm28</a>	<a href="#">Rbm28.bSep08</a>	<a href="#">312182</a>	22730	1020	9	325	RNA binding motif protein 28 (Rbm28) alternative variant bSep08, mRNA.
<a href="#">Rbm34</a>	<a href="#">Rbm34.bSep08</a>	<a href="#">307956</a>	14130	786	9	193	RNA binding motif protein 34 (Rbm34) alternative variant bSep08, mRNA.
<a href="#">Rbm35b</a>	<a href="#">Rbm35b.aSep08</a>	<a href="#">307810</a>	6662	3467	15	746	RNA binding motif protein 35b (Rbm35b) alternative variant aSep08, mRNA.
<a href="#">Rbm38</a>	<a href="#">Rbm38.aSep08</a>	<a href="#">366262</a>	12250	1193	2	271	RNA binding motif protein 38 (Rbm38) alternative variant aSep08, mRNA.
<a href="#">Rbm38</a>	<a href="#">Rbm38.cSep08</a>	<a href="#">366262</a>	12072	992	2	158	RNA binding motif protein 38 (16.9 kD) (Rbm38) alternative variant cSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.aSep08</a>	<a href="#">362251</a>	33169	2738	17	530	RNA binding motif protein 39 like (59.4 kD) (Rbm39) alternative variant aSep08, complete mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.cSep08</a>	<a href="#">362251</a>	15397	869	7	255	RNA binding motif protein 39 like (Rbm39) alternative variant cSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.dSep08</a>	<a href="#">362251</a>	18000	1144	10	235	RNA binding motif protein 39 like (Rbm39) alternative variant dSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.eSep08</a>	<a href="#">362251</a>	16579	1054	9	205	RNA binding motif protein 39 like (Rbm39) alternative variant eSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.fSep08</a>	<a href="#">362251</a>	7151	587	6	195	RNA binding motif protein 39 like (Rbm39) alternative variant fSep08, mRNA.



<a href="#">Rbm39</a>	<a href="#">Rbm39.gSep08</a>	<a href="#">362251</a>	14796	1688	7	169	RNA binding motif protein 39 like (Rbm39) alternative variant gSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.hSep08</a>	<a href="#">362251</a>	12414	795	7	152	RNA binding motif protein 39 like (Rbm39) alternative variant hSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.iSep08</a>	<a href="#">362251</a>	1373	763	2	49	binding CRA a like (Rbm39) alternative variant iSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.jSep08</a>	<a href="#">362251</a>	2316	1242	2	60	putative protein (6.9 kD) (Rbm39) alternative variant jSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.lSep08</a>	<a href="#">362251</a>	1409	778	2	46	putative protein (Rbm39) alternative variant lSep08, mRNA.
<a href="#">Rbm39</a>	<a href="#">Rbm39.mSep08</a>	<a href="#">362251</a>	1608	753	3	40	putative protein of mammalian origin (4.6 kD) (Rbm39) alternative variant mSep08, complete mRNA.
<a href="#">Rbm41</a>	<a href="#">Rbm41.bSep08</a>	<a href="#">680581</a>	8357	612	1	156	RNA binding motif protein 41 (Rbm41) alternative variant bSep08, mRNA.
<a href="#">Rbm42</a>	<a href="#">Rbm42.bSep08</a>	<a href="#">361545</a>	9548	1275	4	391	RNA binding motif protein 42 (Rbm42) alternative variant bSep08, mRNA.
<a href="#">Rbm43</a>	<a href="#">Rbm43.aSep08</a>	<a href="#">311020</a>	2819	2114	2	276	RNA binding motif protein 43 (Rbm43) alternative variant aSep08, mRNA.
<a href="#">Rbm43</a>	<a href="#">Rbm43.bSep08</a>	<a href="#">311020</a>	19421	737	4	206	RNA binding motif protein 43 (Rbm43) alternative variant bSep08, mRNA.
<a href="#">Rbm44</a>	<a href="#">Rbm44.aSep08</a>	<a href="#">501183</a>	3481	735		163	RNA binding motif protein 44 (Rbm44) mRNA.
<a href="#">Rbm45</a>	<a href="#">Rbm45.bSep08</a>	<a href="#">266631</a>	5377	1422	3	263	RNA binding motif protein 45 (Rbm45) alternative variant bSep08, mRNA.
<a href="#">Rbm45</a>	<a href="#">Rbm45.cSep08</a>	<a href="#">266631</a>	9490	746	4	223	RNA binding motif protein 45 (Rbm45) alternative variant cSep08, mRNA.
<a href="#">Rbms1</a>	<a href="#">Rbms1.bSep08</a>	<a href="#">362138</a>	134369	2513	15	498	RNA binding motif, single stranded interacting protein 1 (Rbms1) alternative variant bSep08, mRNA.
<a href="#">Rbms1</a>	<a href="#">Rbms1.cSep08</a>	<a href="#">362138</a>	113308	698	7	170	RNA binding motif, single stranded interacting protein 1 (Rbms1) alternative variant cSep08, mRNA.
<a href="#">Rbms1</a>	<a href="#">Rbms1.dSep08</a>	<a href="#">362138</a>	113296	693	7	169	RNA binding motif, single stranded interacting protein 1 (Rbms1) alternative variant dSep08, mRNA.
<a href="#">Rbms1</a>	<a href="#">Rbms1.eSep08</a>	<a href="#">362138</a>	7149	524	5	89	RNA binding motif, single stranded interacting protein 1 (Rbms1) alternative variant eSep08, mRNA.
<a href="#">Rbms2</a>	<a href="#">Rbms2.aSep08</a>	<a href="#">288771</a>	36090	1798	9	495	RNA binding motif, single stranded interacting protein 2 (Rbms2) alternative variant aSep08, mRNA.
<a href="#">Rbms2</a>	<a href="#">Rbms2.cSep08</a>	<a href="#">288771</a>	696	436	2	66	RNA binding motif, single stranded interacting protein 2 (Rbms2) alternative variant cSep08, mRNA.
<a href="#">Rbmx</a>	<a href="#">Rbmx.bSep08</a>	<a href="#">302855</a>	9089	2127	10	306	RBM1 (Rbmx) alternative variant bSep08, mRNA.
<a href="#">Rbmx</a>	<a href="#">Rbmx.cSep08</a>	<a href="#">302855</a>	3665	669	6	187	RBM1 (Rbmx) alternative variant cSep08, mRNA.
<a href="#">Rbmx</a>	<a href="#">Rbmx.dSep08</a>	<a href="#">302855</a>	3882	2282	3	126	putative cytoplasmic protein of vertebrate origin (13.9 kD) (Rbmx) alternative variant dSep08, mRNA.
<a href="#">Rbmx2</a>	<a href="#">Rbmx2.bSep08</a>	<a href="#">367930</a>	6525	450	1	91	RNA binding motif protein, X-linked 2 (Rbmx2) alternative variant bSep08, mRNA.
<a href="#">Rbmxrtl</a>	<a href="#">Rbmxrtl.aSep08</a>	<a href="#">307779</a>	3571	1676	1	388	RBM1 (42.2 kD) (Rbmxrtl) alternative variant aSep08, complete mRNA.
<a href="#">Rbp3</a>	<a href="#">Rbp3.aSep08</a>	<a href="#">24711</a>	3727	343		114	retinol binding protein 3, interstitial (Rbp3) mRNA.

<a href="#">Rbpjl</a>	<a href="#">Rbpjl.bSep08</a>	<a href="#">362268</a>	6266	767	1	255	recombination signal binding protein for immunoglobulin kappa J region-like (Rbpjl) alternative variant bSep08, mRNA.
<a href="#">Rbpms2</a>	<a href="#">Rbpms2.aSep08</a>	<a href="#">503214</a>	11824	1643		181	RNA binding protein with multiple splicing 2 (Rbpms2) alternative variant aSep08, mRNA.
<a href="#">Rbpms2</a>	<a href="#">Rbpms2.bSep08</a>	<a href="#">503214</a>	32428	1874		178	RNA binding protein with multiple splicing 2 (19.5 kD) (Rbpms2) alternative variant bSep08, mRNA.
<a href="#">Rbpms2</a>	<a href="#">Rbpms2.cSep08</a>	<a href="#">503214</a>	22937	687		89	RNA binding protein with multiple splicing 2 (Rbpms2) alternative variant cSep08, mRNA.
<a href="#">Rbx1</a>	<a href="#">Rbx1.bSep08</a>	<a href="#">300084</a>	10114	2248	4	86	ring-box 1 (9.4 kD) (Rbx1) alternative variant bSep08, complete mRNA.
<a href="#">Rbx1</a>	<a href="#">Rbx1.cSep08</a>	<a href="#">300084</a>	2393	1217	2	80	ring-box 1 (8.2 kD) (Rbx1) alternative variant cSep08, mRNA.
<a href="#">Rc3h2</a>	<a href="#">Rc3h2.bSep08</a>	<a href="#">311909</a>	2791	1742	6	454	ring finger CCCH-type zinc domains 2 (Rc3h2) alternative variant bSep08, mRNA.
<a href="#">Rc3h2</a>	<a href="#">Rc3h2.cSep08</a>	<a href="#">311909</a>	727	628	2	86	ring finger CCCH-type zinc domains 2 (Rc3h2) alternative variant cSep08, mRNA.
<a href="#">Rcan1</a>	<a href="#">Rcan1.aSep08</a>	<a href="#">266766</a>	92185	710	3	208	regulator of calcineurin 1 (Rcan1) alternative variant aSep08, mRNA.
<a href="#">Rcan1</a>	<a href="#">Rcan1.cSep08</a>	<a href="#">266766</a>	4753	685	2	119	regulator of calcineurin 1 (13.5 kD) (Rcan1) alternative variant cSep08, mRNA.
<a href="#">Rcan2</a>	<a href="#">Rcan2.bSep08</a>	<a href="#">140666</a>	72660	425	2	80	regulator of calcineurin 2 (Rcan2) alternative variant bSep08, mRNA.
<a href="#">Rcan2</a>	<a href="#">Rcan2.cSep08</a>	<a href="#">140666</a>	11959	537	2	83	regulator of calcineurin 2 (9.5 kD) (Rcan2) alternative variant cSep08, mRNA.
<a href="#">Rcbtb1</a>	<a href="#">Rcbtb1.bSep08</a>	<a href="#">361050</a>	66906	1359	9	159	regulator of chromosome condensation, RCC1 (16.9 kD) (Rcbtb1) alternative variant bSep08, mRNA.
<a href="#">Rcbtb1</a>	<a href="#">Rcbtb1.cSep08</a>	<a href="#">361050</a>	2458	610	3	110	putative protein of eukaryotic origin (Rcbtb1) alternative variant cSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.bSep08</a>	<a href="#">290363</a>	15441	1028	6	303	regulator of chromosome condensation, RCC1 (Rcbtb2) alternative variant bSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.cSep08</a>	<a href="#">290363</a>	14994	747	6	192	regulator of chromosome condensation, RCC1 (Rcbtb2) alternative variant cSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.dSep08</a>	<a href="#">290363</a>	3716	825	4	154	regulator of chromosome condensation, RCC1 (Rcbtb2) alternative variant dSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.eSep08</a>	<a href="#">290363</a>	13242	492	5	128	putative protein of eukaryotic origin (Rcbtb2) alternative variant eSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.fSep08</a>	<a href="#">290363</a>	11614	1337	5	63	putative protein (Rcbtb2) alternative variant fSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.gSep08</a>	<a href="#">290363</a>	20875	758	8	126	putative protein of ancient origin (Rcbtb2) alternative variant gSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.hSep08</a>	<a href="#">290363</a>	20824	752	8	120	putative protein of ancient origin (Rcbtb2) alternative variant hSep08, mRNA.
<a href="#">Rcbtb2</a>	<a href="#">Rcbtb2.iSep08</a>	<a href="#">290363</a>	20072	700	7	94	putative protein (Rcbtb2) alternative variant iSep08, mRNA.
<a href="#">RCC1.0</a>	<a href="#">RCC1.0.aSep08</a>		11325	890		296	retinitis pigmentosa GTPase regulator (RCC1.0) mRNA.
<a href="#">RCC1.1</a>	<a href="#">RCC1.1.aSep08</a>		5734	706		235	hect domain RLD 6 CRA b (RCC1.1) mRNA.

<a href="#">Rcc2</a>	<a href="#">Rcc2.aSep08</a>	<a href="#">298594</a>	15336	3278		427	regulator of chromosome condensation, RCC1 (Rcc2) mRNA.
<a href="#">Rce1</a>	<a href="#">Rce1.bSep08</a>	<a href="#">309153</a>	3051	1361	8	314	RCE1 homolog, prenyl protein peptidase ( <i>S. cerevisiae</i> ) (Rce1) alternative variant bSep08, mRNA.
<a href="#">Rce1</a>	<a href="#">Rce1.cSep08</a>	<a href="#">309153</a>	3100	1351	7	308	RCE1 homolog, prenyl protein peptidase ( <i>S. cerevisiae</i> ) (Rce1) alternative variant cSep08, complete mRNA.
<a href="#">Rce1</a>	<a href="#">Rce1.dSep08</a>	<a href="#">309153</a>	3075	1742	4	214	RCE1 homolog, prenyl protein peptidase ( <i>S. cerevisiae</i> ) (Rce1) alternative variant dSep08, mRNA.
<a href="#">Rce1</a>	<a href="#">Rce1.eSep08</a>	<a href="#">309153</a>	2451	723	7	211	RCE1 homolog, prenyl protein peptidase ( <i>S. cerevisiae</i> ) (Rce1) alternative variant eSep08, mRNA.
<a href="#">Rce1</a>	<a href="#">Rce1.fSep08</a>	<a href="#">309153</a>	1552	664	4	150	RCE1 homolog, prenyl protein peptidase ( <i>S. cerevisiae</i> ) (Rce1) alternative variant fSep08, mRNA.
<a href="#">rCG_22919</a>	<a href="#">rCG_22919.cSep08</a>	<a href="#">100125364</a>	880	628	2	209	hypothetical protein LOC100125364 (rCG_22919) alternative variant cSep08, mRNA.
<a href="#">rCG_22919</a>	<a href="#">rCG_22919.dSep08</a>	<a href="#">100125364</a>	905	658	2	185	hypothetical protein LOC100125364 (20.9 kD) (rCG_22919) alternative variant dSep08, mRNA.
<a href="#">rCG_22919</a>	<a href="#">rCG_22919.eSep08</a>	<a href="#">100125364</a>	9601	378	2	48	hypothetical protein LOC100125364 (5.2 kD) (rCG_22919) alternative variant eSep08, mRNA.
<a href="#">rCG_28701</a>	<a href="#">rCG_28701.aSep08</a>	<a href="#">691807</a>	1765	1085		65	hypothetical protein LOC691807 (7.4 kD) (rCG_28701) complete mRNA.
<a href="#">rCG_34148</a>	<a href="#">rCG_34148.bSep08</a>	<a href="#">100169747</a>	7514	574	5	113	alanine-glyoxylate aminotransferase 2-like 2 (rCG_34148) alternative variant bSep08, mRNA.
<a href="#">rCG_34148</a>	<a href="#">rCG_34148.cSep08</a>	<a href="#">100169747</a>	1045	387	2	77	alanine-glyoxylate aminotransferase 2-like 2 (9.0 kD) (rCG_34148) alternative variant cSep08, mRNA.
<a href="#">rCG_34148</a>	<a href="#">rCG_34148.dSep08</a>	<a href="#">100169747</a>	3820	772	3	67	alanine-glyoxylate aminotransferase 2-like 2 (rCG_34148) alternative variant dSep08, mRNA.
<a href="#">rCG_43687</a>	<a href="#">rCG_43687.bSep08</a>	<a href="#">688459</a>	10855	1206	2	181	hypothetical protein LOC688459 (rCG_43687) alternative variant bSep08, mRNA.
<a href="#">rCG_48149</a>	<a href="#">rCG_48149.bSep08</a>	<a href="#">100126191</a>	1063	467	1	127	RAB1B, member RAS oncogene family (rCG_48149) alternative variant bSep08, mRNA.
<a href="#">rCG_48647</a>	<a href="#">rCG_48647.aSep08</a>	<a href="#">688047</a>	5064	915		151	similar to Lysozyme C type 2 precursor (1,4-beta-N-acetylmuramidase C) (16.8 kD) (rCG_48647) alternative variant aSep08, mRNA.
<a href="#">rCG_49031</a>	<a href="#">rCG_49031.aSep08</a>	<a href="#">362863</a>	21741	1900		399	grp94 neighboring nucleotidase (46.2 kD) (rCG_49031) mRNA.
<a href="#">rCG_54677</a>	<a href="#">rCG_54677.bSep08</a>	<a href="#">678728</a>	16591	760	3	152	bone morphogenetic protein 1 (rCG_54677) alternative variant bSep08, mRNA.
<a href="#">rCG_54677</a>	<a href="#">rCG_54677.bSep08</a>	<a href="#">678743</a>	16591	760	3	152	bone morphogenetic protein 1 (rCG_54677) alternative variant bSep08, mRNA.
<a href="#">rCG_54677</a>	<a href="#">rCG_54677.cSep08</a>	<a href="#">678728</a>	12576	683	1	104	tolloid-like (rCG_54677) alternative variant cSep08, mRNA.
<a href="#">rCG_54677</a>	<a href="#">rCG_54677.cSep08</a>	<a href="#">678743</a>	12576	683	1	104	tolloid-like (rCG_54677) alternative variant cSep08, mRNA.
<a href="#">rCG_59505</a>	<a href="#">rCG_59505.bSep08</a>	<a href="#">100125385</a>	10808	500	2	121	hypothetical protein LOC100125385 (rCG_59505) alternative variant bSep08, mRNA.
<a href="#">rCG_60321</a>	<a href="#">rCG_60321.bSep08</a>	<a href="#">688613</a>	4160	735	2	55	hypothetical protein LOC688613 (5.7 kD) (rCG_60321) alternative variant bSep08, mRNA.
<a href="#">Rchy1</a>	<a href="#">Rchy1.bSep08</a>	<a href="#">289508</a>	10209	1127	8	204	zinc finger, RING-type (23.5 kD) (Rchy1) alternative variant bSep08, mRNA.

<a href="#">Rchy1</a>	<a href="#">Rchy1.cSep08</a>	<a href="#">289508</a>	1901	1100	2	75	putative protein of eukaryotic origin (8.5 kD) (Rchy1) alternative variant cSep08, mRNA.
<a href="#">Rcl1</a>	<a href="#">Rcl1.aSep08</a>	<a href="#">309301</a>	45305	1827	9	398	RNA terminal phosphate cyclase-like 1 (Rcl1) alternative variant aSep08, mRNA.
<a href="#">Rcl1</a>	<a href="#">Rcl1.cSep08</a>	<a href="#">309301</a>	16308	860	5	170	RNA terminal phosphate cyclase-like 1 (18.4 kD) (Rcl1) alternative variant cSep08, mRNA.
<a href="#">Rcn1</a>	<a href="#">Rcn1.bSep08</a>	<a href="#">362182</a>	5051	983	1	85	reticulocalbin 1 (9.6 kD) (Rcn1) alternative variant bSep08, complete mRNA.
<a href="#">Rcn2</a>	<a href="#">Rcn2.bSep08</a>	<a href="#">29218</a>	5280	741	2	120	reticulocalbin 2 (Rcn2) alternative variant bSep08, mRNA.
<a href="#">Rcor1</a>	<a href="#">Rcor1.aSep08</a>	<a href="#">314458</a>	29977	772		240	REST corepressor 1 (Rcor1) mRNA.
<a href="#">Rcor2</a>	<a href="#">Rcor2.bSep08</a>	<a href="#">305811</a>	2812	783	7	260	REST corepressor 2 (Rcor2) alternative variant bSep08, mRNA.
<a href="#">Rcor2</a>	<a href="#">Rcor2.cSep08</a>	<a href="#">305811</a>	2309	1472	4	223	REST corepressor 2 (24.2 kD) (Rcor2) alternative variant cSep08, mRNA.
<a href="#">Rcsd1</a>	<a href="#">Rcsd1.bSep08</a>	<a href="#">360872</a>	7953	384	4	127	putative protein of vertebrate origin (Rcsd1) alternative variant bSep08, mRNA.
<a href="#">Rcsd1</a>	<a href="#">Rcsd1.cSep08</a>	<a href="#">360872</a>	3491	753	3	106	putative protein of vertebrate origin (Rcsd1) alternative variant cSep08, mRNA.
<a href="#">Rcsd1</a>	<a href="#">Rcsd1.dSep08</a>	<a href="#">360872</a>	5846	486	4	86	putative protein of vertebrate origin (Rcsd1) alternative variant dSep08, mRNA.
<a href="#">Rdbp</a>	<a href="#">Rdbp.bSep08</a>	<a href="#">294258</a>	2844	1357	5	255	RD RNA-binding protein (29.2 kD) (Rdbp) alternative variant bSep08, mRNA.
<a href="#">Rdh2</a>	<a href="#">Rdh2.bSep08</a>	<a href="#">299511</a>	16938	3800	2	317	retinol dehydrogenase 2 (35.7 kD) (Rdh2) alternative variant bSep08, mRNA.
<a href="#">Rdh5</a>	<a href="#">Rdh5.aSep08</a>	<a href="#">366791</a>	1111	430		75	retinol dehydrogenase 5 (Rdh5) alternative variant aSep08, mRNA.
<a href="#">Rdh5</a>	<a href="#">Rdh5.bSep08</a>	<a href="#">366791</a>	1622	940		64	retinol dehydrogenase 5 (Rdh5) alternative variant bSep08, mRNA.
<a href="#">Rdh12</a>	<a href="#">Rdh12.bSep08</a>	<a href="#">314264</a>	5325	492	2	121	retinol dehydrogenase 12 (Rdh12) alternative variant bSep08, mRNA.
<a href="#">Rdh14</a>	<a href="#">Rdh14.bSep08</a>	<a href="#">298881</a>	4484	782	1	229	retinol dehydrogenase 14 (all-trans and 9-cis) (Rdh14) alternative variant bSep08, mRNA.
<a href="#">Rdm1</a>	<a href="#">Rdm1.bSep08</a>	<a href="#">287726</a>	3073	767	4	143	RAD52 motif 1 (Rdm1) alternative variant bSep08, mRNA.
<a href="#">Rdm1</a>	<a href="#">Rdm1.cSep08</a>	<a href="#">287726</a>	5495	668	3	88	RAD52 motif 1 (9.9 kD) (Rdm1) alternative variant cSep08, mRNA.
<a href="#">Rdx</a>	<a href="#">Rdx.aSep08</a>	<a href="#">315655</a>	20542	3201	6	318	radixin (Rdx) alternative variant aSep08, mRNA.
<a href="#">Rdx</a>	<a href="#">Rdx.cSep08</a>	<a href="#">315655</a>	15469	403	3	75	radixin (Rdx) alternative variant cSep08, mRNA.
<a href="#">Rdx</a>	<a href="#">Rdx.dSep08</a>	<a href="#">315655</a>	8868	753	5	58	radixin (Rdx) alternative variant dSep08, mRNA.
<a href="#">Rdx</a>	<a href="#">Rdx.eSep08</a>	<a href="#">315655</a>	17863	732	3	89	radixin (9.6 kD) (Rdx) alternative variant eSep08, mRNA.
<a href="#">Rdx</a>	<a href="#">Rdx.gSep08</a>	<a href="#">315655</a>	15580	620	2	30	radixin (Rdx) alternative variant gSep08, mRNA.
<a href="#">Reck</a>	<a href="#">Reck.bSep08</a>	<a href="#">313488</a>	12925	2608	2	430	reversion-inducing-cysteine-rich protein with kazal motifs (Reck) alternative variant bSep08, mRNA.
<a href="#">Recql</a>	<a href="#">Recql.bSep08</a>	<a href="#">312824</a>	18390	1027	8	342	RecQ protein-like (Recql) alternative variant bSep08, mRNA.
<a href="#">Recql</a>	<a href="#">Recql.cSep08</a>	<a href="#">312824</a>	13895	816	6	271	RecQ protein-like (Recql) alternative variant cSep08, mRNA.

<a href="#">Recql</a>	<a href="#">Recql.dSep08</a>	<a href="#">312824</a>	13940	763	6	209	RecQ protein-like (Recql) alternative variant dSep08, mRNA.
<a href="#">Recql</a>	<a href="#">Recql.eSep08</a>	<a href="#">312824</a>	2729	1745	3	168	RecQ protein-like (18.4 kD) (Recql) alternative variant eSep08, mRNA.
<a href="#">Recql</a>	<a href="#">Recql.fSep08</a>	<a href="#">312824</a>	1857	419	4	87	RecQ protein-like (Recql) alternative variant fSep08, mRNA.
<a href="#">Recql</a>	<a href="#">Recql.gSep08</a>	<a href="#">312824</a>	488	401	2	62	putative protein (7.3 kD) (Recql) alternative variant gSep08, mRNA.
<a href="#">Recql4</a>	<a href="#">Recql4.bSep08</a>	<a href="#">300057</a>	5349	892	4	286	RecQ protein-like 4 (Recql4) alternative variant bSep08, mRNA.
<a href="#">Recql4</a>	<a href="#">Recql4.cSep08</a>	<a href="#">300057</a>	5541	1084	4	272	RecQ protein-like 4 (Recql4) alternative variant cSep08, mRNA.
<a href="#">Recql5</a>	<a href="#">Recql5.bSep08</a>	<a href="#">287834</a>	11288	738	1	246	RecQ protein-like 5 (Recql5) alternative variant bSep08, mRNA.
<a href="#">reeby</a>	<a href="#">reeby.aSep08</a>		9583	1029	2	109	putative secreted or extracellular protein precursor (12.0 kD) (reeby) alternative variant aSep08, mRNA.
<a href="#">reechy</a>	<a href="#">reechy.aSep08</a>		996	645	1	52	putative protein of mammalian origin (5.8 kD) (reechy) alternative variant aSep08, mRNA.
<a href="#">reechy</a>	<a href="#">reechy.bSep08</a>		10141	519	2	52	putative protein of mammalian origin (5.8 kD) (reechy) alternative variant bSep08, mRNA.
<a href="#">reedar</a>	<a href="#">reedar.aSep08</a>		875	414	2	120	RGD motif leucine rich repeats tropomodulin domain proline-rich containing like (reedar) alternative variant aSep08, mRNA.
<a href="#">reedar</a>	<a href="#">reedar.bSep08</a>		797	600	1	81	RGD motif leucine rich repeats tropomodulin domain proline-rich containing like (reedar) alternative variant bSep08, mRNA.
<a href="#">reefer</a>	<a href="#">reefer.aSep08</a>		833	743		37	putative protein (reefer) mRNA.
<a href="#">reeflo</a>	<a href="#">reeflo.aSep08</a>		6808	711		147	putative nuclear protein, with 2 coiled coil domains, of eukaryotic origin (16.8 kD) (reeflo) mRNA.
<a href="#">reeflu</a>	<a href="#">reeflu.aSep08</a>		18052	795		82	putative protein (9.2 kD) (reeflu) mRNA.
<a href="#">reegar</a>	<a href="#">reegar.aSep08</a>		13062	1735		202	putative protein of metazoan origin (reegar) mRNA.
<a href="#">reeja</a>	<a href="#">reeja.aSep08</a>		3449	1310		55	putative protein (reeja) mRNA.
<a href="#">reejey</a>	<a href="#">reejey.aSep08</a>		12083	625		41	putative protein (4.8 kD) (reejey) mRNA.
<a href="#">reekee</a>	<a href="#">reekee.aSep08</a>		9952	1055		351	roquin (reekee) mRNA.
<a href="#">reekler</a>	<a href="#">reekler.aSep08</a>		1638	465	2	126	putative protein (reekler) alternative variant aSep08, mRNA.
<a href="#">reelo</a>	<a href="#">reelo.aSep08</a>		5650	737	3	82	putative protein (reelo) alternative variant aSep08, mRNA.
<a href="#">reemee</a>	<a href="#">reemee.aSep08</a>		8560	757		43	putative protein (reemee) mRNA.
<a href="#">reemer</a>	<a href="#">reemer.aSep08</a>		4907	2303		205	CRA a (reemer) mRNA.
<a href="#">reenoy</a>	<a href="#">reenoy.aSep08</a>		13718	575		34	putative protein (3.7 kD) (reenoy) mRNA.
<a href="#">Reep1</a>	<a href="#">Reep1.bSep08</a>	<a href="#">362384</a>	2986	629	2	81	receptor accessory protein 1 (Reep1) alternative variant bSep08, mRNA.
<a href="#">Reep5</a>	<a href="#">Reep5.aSep08</a>	<a href="#">364838</a>	30868	2480	5	220	receptor accessory protein 5 (Reep5) alternative variant aSep08, mRNA.
<a href="#">Reep6</a>	<a href="#">Reep6.bSep08</a>	<a href="#">362835</a>	2339	995		331	receptor accessory protein 6 (Reep6) alternative variant bSep08, mRNA.

<a href="#">reepor</a>	<a href="#">reepor.aSep08</a>		350	264		68	putative protein (reepor) mRNA.
<a href="#">reesa</a>	<a href="#">reesa.aSep08</a>		9248	404		134	E1A binding protein p300 like (reesa) mRNA.
<a href="#">reeshee</a>	<a href="#">reeshee.aSep08</a>		8737	1725	5	268	acetyl-CoA transporter (reeshee) alternative variant aSep08, mRNA.
<a href="#">reeshee</a>	<a href="#">reeshee.bSep08</a>		2076	931	2	122	acetyl-CoA transporter (13.4 kD) (reeshee) alternative variant bSep08, mRNA.
<a href="#">reeshee</a>	<a href="#">reeshee.cSep08</a>		2178	345	3	90	putative protein (reeshee) alternative variant cSep08, mRNA.
<a href="#">reetu</a>	<a href="#">reetu.aSep08</a>		1357	600	2	138	zinc finger protein 513 (reetu) alternative variant aSep08, mRNA.
<a href="#">reetu</a>	<a href="#">reetu.bSep08</a>		1613	608	3	133	zinc finger protein 513 (reetu) alternative variant bSep08, mRNA.
<a href="#">reetu</a>	<a href="#">reetu.cSep08</a>		844	596	2	77	putative protein (reetu) alternative variant cSep08, mRNA.
<a href="#">reevar</a>	<a href="#">reevar.aSep08</a>		7958	716		132	CRA b like (reevar) alternative variant aSep08, mRNA.
<a href="#">reevar</a>	<a href="#">reevar.bSep08</a>		6289	494		126	CRA b like (reevar) alternative variant bSep08, mRNA.
<a href="#">reewey</a>	<a href="#">reewey.aSep08</a>		3486	368		122	putative protein of eukaryotic origin (reewey) mRNA.
<a href="#">Reg3a</a>	<a href="#">Reg3a.aSep08</a>	<a href="#">171162</a>	2769	790	1	192	regenerating islet-derived 3 alpha (Reg3a) alternative variant aSep08, mRNA.
<a href="#">Reg3b</a>	<a href="#">Reg3b.cSep08</a>	<a href="#">24618</a>	1592	669	2	48	regenerating islet-derived 3 beta (5.5 kD) (Reg3b) alternative variant cSep08, mRNA.
<a href="#">Rela</a>	<a href="#">Rela.bSep08</a>	<a href="#">309165</a>	4506	467	3	65	v-rel reticuloendotheliosis viral oncogene homolog A (avian) (Rela) alternative variant bSep08, mRNA.
<a href="#">Rell1</a>	<a href="#">Rell1.bSep08</a>	<a href="#">289635</a>	30683	2762	1	229	RELT-like 1 (Rell1) alternative variant bSep08, mRNA.
<a href="#">Rell2</a>	<a href="#">Rell2.bSep08</a>	<a href="#">361313</a>	2537	763	5	253	RELT-like 2 (Rell2) alternative variant bSep08, mRNA.
<a href="#">Reln</a>	<a href="#">Reln.bSep08</a>	<a href="#">24718</a>	7735	622	5	206	reelin (Reln) alternative variant bSep08, mRNA.
<a href="#">Reln</a>	<a href="#">Reln.cSep08</a>	<a href="#">24718</a>	2880	518	2	106	reelin (Reln) alternative variant cSep08, mRNA.
<a href="#">Reln</a>	<a href="#">Reln.dSep08</a>	<a href="#">24718</a>	6396	1470	2	64	reelin (7.2 kD) (Reln) alternative variant dSep08, mRNA.
<a href="#">Rem1</a>	<a href="#">Rem1.bSep08</a>	<a href="#">366232</a>	7335	1384	4	213	rad and gem related GTP binding protein 1 (23.2 kD) (Rem1) alternative variant bSep08, mRNA.
<a href="#">Ren1</a>	<a href="#">Ren1.bSep08</a>	<a href="#">24715</a>	6240	692	1	207	renin 1 structural (Ren1) alternative variant bSep08, mRNA.
<a href="#">Renbp</a>	<a href="#">Renbp.bSep08</a>	<a href="#">81759</a>	3520	549	5	150	renin binding protein (Renbp) alternative variant bSep08, mRNA.
<a href="#">Reps1</a>	<a href="#">Reps1.bSep08</a>	<a href="#">292944</a>	33312	1808	15	537	calcium-binding EF-hand containing protein (Reps1) alternative variant bSep08, mRNA.
<a href="#">Reps1</a>	<a href="#">Reps1.cSep08</a>	<a href="#">292944</a>	28049	816	9	272	calcium-binding EF-hand containing protein (Reps1) alternative variant cSep08, mRNA.
<a href="#">Reps1</a>	<a href="#">Reps1.dSep08</a>	<a href="#">292944</a>	27871	811	8	270	putative protein of vertebrate origin (Reps1) alternative variant dSep08, mRNA.
<a href="#">Reps1</a>	<a href="#">Reps1.eSep08</a>	<a href="#">292944</a>	11095	535	2	127	putative protein of vertebrate origin (Reps1) alternative variant eSep08, mRNA.
<a href="#">Reps1</a>	<a href="#">Reps1.fSep08</a>	<a href="#">292944</a>	2724	463	4	80	putative protein of vertebrate origin (Reps1) alternative variant fSep08, mRNA.
<a href="#">Reps2</a>	<a href="#">Reps2.aSep08</a>	<a href="#">363466</a>	113951	483		160	putative protein of eukaryotic origin (Reps2) mRNA.

<a href="#">Rer1</a>	<a href="#">Rer1.aSep08</a>	<a href="#">298675</a>	12393	1797	6	327	RER1 retention in endoplasmic reticulum 1 homolog (S. cerevisiae) (Rer1) alternative variant aSep08, mRNA.
<a href="#">Rer1</a>	<a href="#">Rer1.cSep08</a>	<a href="#">298675</a>	4712	1085	2	68	RER1 retention in endoplasmic reticulum 1 homolog (S. cerevisiae) (8.1 kD) (Rer1) alternative variant cSep08, mRNA.
<a href="#">rerby</a>	<a href="#">rerby.aSep08</a>		23846	582		42	CRA b like (rerby) mRNA.
<a href="#">rerchy</a>	<a href="#">rerchy.aSep08</a>		1095	323		59	putative protein (6.6 kD) (rerchy) mRNA.
<a href="#">rerdar</a>	<a href="#">rerdar.aSep08</a>		17453	427		66	putative protein (rerdar) mRNA.
<a href="#">Rere</a>	<a href="#">Rere.bSep08</a>	<a href="#">116665</a>	2975	2206	3	685	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant bSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.bSep08</a>	<a href="#">691475</a>	2975	2206	3	685	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant bSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.dSep08</a>	<a href="#">116665</a>	39886	575	3	107	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant dSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.dSep08</a>	<a href="#">691475</a>	39886	575	3	107	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant dSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.eSep08</a>	<a href="#">116665</a>	21459	331	4	69	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant eSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.eSep08</a>	<a href="#">691475</a>	21459	331	4	69	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant eSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.fSep08</a>	<a href="#">116665</a>	35669	233	3	52	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant fSep08, mRNA.
<a href="#">Rere</a>	<a href="#">Rere.fSep08</a>	<a href="#">691475</a>	35669	233	3	52	arginine-glutamic acid dipeptide (RE) repeats and hypothetical protein LOC691475 (Rere) alternative variant fSep08, mRNA.
<a href="#">rerfer</a>	<a href="#">rerfer.aSep08</a>		4603	392		102	putative protein (rerfer) mRNA.
<a href="#">rerflo</a>	<a href="#">rerflo.aSep08</a>		8932	756		84	putative protein (rerflo) mRNA.
<a href="#">rerflu</a>	<a href="#">rerflu.aSep08</a>		418224	1728		541	protein CRA b (rerflu) alternative variant aSep08, mRNA.
<a href="#">Rerg</a>	<a href="#">Rerg.aSep08</a>	<a href="#">502916</a>	112501	1801			
<a href="#">Rerg</a>	<a href="#">Rerg.cSep08</a>	<a href="#">502916</a>	98795	416		76	RAS-like, estrogen-regulated, growth-inhibitor (Rerg) alternative variant cSep08, mRNA.
<a href="#">rergar</a>	<a href="#">rergar.aSep08</a>		1088	417		100	putative protein (rergar) mRNA.
<a href="#">Rergl</a>	<a href="#">Rergl.aSep08</a>	<a href="#">500356</a>	8835	723		214	RERG/RAS-like (Rergl) mRNA.
<a href="#">rerja</a>	<a href="#">rerja.aSep08</a>		15422	747	2	99	putative nuclear protein (11.1 kD) (rerja) alternative variant aSep08, mRNA.
<a href="#">rerja</a>	<a href="#">rerja.bSep08</a>		5781	737	1	44	putative protein, with a coiled coil domain (5.1 kD) (rerja) alternative variant bSep08, mRNA.
<a href="#">rerjey</a>	<a href="#">rerjey.aSep08</a>		4405	404		71	putative protein (8.3 kD) (rerjey) mRNA.

<a href="#">rerkee</a>	<a href="#">rerkee.aSep08</a>		3186	589	2	111	ring CCCH domains 1 (rerkee) alternative variant aSep08, mRNA.
<a href="#">rerkee</a>	<a href="#">rerkee.bSep08</a>		916	321	1	79	ring CCCH domains 1 (rerkee) alternative variant bSep08, mRNA.
<a href="#">rerkler</a>	<a href="#">rerkler.aSep08</a>		664	413		20	putative protein (2.1 kD) (rerkler) mRNA.
<a href="#">rerlo</a>	<a href="#">rerlo.aSep08</a>		1624	753		36	putative protein (4.0 kD) (rerlo) mRNA.
<a href="#">rermee</a>	<a href="#">rermee.aSep08</a>		1166	660		34	putative protein (rermee) mRNA.
<a href="#">rermer</a>	<a href="#">rermer.aSep08</a>		1723	441		102	putative protein (rermer) mRNA.
<a href="#">rernoy</a>	<a href="#">rernoy.aSep08</a>		2776	915		128	putative secreted or extracellular protein precursor (13.7 kD) (rernoy) mRNA.
<a href="#">rerpor</a>	<a href="#">rerpor.aSep08</a>		51702	647		41	putative protein (rerpor) mRNA.
<a href="#">rersa</a>	<a href="#">rersa.aSep08</a>		3635	327		108	E1A binding protein p300 like (rersa) mRNA.
<a href="#">rershee</a>	<a href="#">rershee.aSep08</a>		2304	509		42	putative protein (rershee) mRNA.
<a href="#">rertu</a>	<a href="#">rertu.cSep08</a>		5920	289	2	58	putative protein (rertu) alternative variant cSep08, mRNA.
<a href="#">rervar</a>	<a href="#">rervar.bSep08</a>		6099	492	2	73	putative secreted or extracellular protein precursor (8.0 kD) (rervar) alternative variant bSep08, mRNA.
<a href="#">rerwey</a>	<a href="#">rerwey.aSep08</a>		851	406	2	110	putative mitochondrial protein (12.5 kD) (rerwey) alternative variant aSep08, mRNA.
<a href="#">Resp18</a>	<a href="#">Resp18.bSep08</a>	<a href="#">50561</a>	4390	532	2	141	regulated endocrine-specific protein 18 (15.9 kD) (Resp18) alternative variant bSep08, mRNA.
<a href="#">Resp18</a>	<a href="#">Resp18.cSep08</a>	<a href="#">50561</a>	3917	417	2	103	regulated endocrine-specific protein 18 (Resp18) alternative variant cSep08, mRNA.
<a href="#">Ret</a>	<a href="#">Ret.dSep08</a>	<a href="#">24716</a>	5378	410	3	31	ret proto-oncogene (Ret) alternative variant dSep08, mRNA.
<a href="#">Retn</a>	<a href="#">Retn.bSep08</a>	<a href="#">246250</a>	889	728	1	25	resistin (Retn) alternative variant bSep08, mRNA.
<a href="#">Rev1</a>	<a href="#">Rev1.bSep08</a>	<a href="#">316344</a>	2477	1133	6	280	REV1 homolog (S. cerevisiae) (Rev1) alternative variant bSep08, mRNA.
<a href="#">Rev1</a>	<a href="#">Rev1.cSep08</a>	<a href="#">316344</a>	1530	723	3	211	REV1 homolog (S. cerevisiae) (Rev1) alternative variant cSep08, mRNA.
<a href="#">Rev1</a>	<a href="#">Rev1.dSep08</a>	<a href="#">316344</a>	6395	359	3	119	REV1 homolog (S. cerevisiae) (Rev1) alternative variant dSep08, mRNA.
<a href="#">Rev1</a>	<a href="#">Rev1.fSep08</a>	<a href="#">316344</a>	700	386	2	65	REV1 homolog (S. cerevisiae) (7.3 kD) (Rev1) alternative variant fSep08, mRNA.
<a href="#">Rev3l</a>	<a href="#">Rev3l.bSep08</a>	<a href="#">309812</a>	4624	383	3	127	REV3-like, catalytic subunit of DNA polymerase zeta RAD54 like (S. cerevisiae) (Rev3l) alternative variant bSep08, mRNA.
<a href="#">Rexo1</a>	<a href="#">Rexo1.bSep08</a>	<a href="#">314630</a>	2010	845	4	229	REX1, RNA exonuclease 1 homolog (S. cerevisiae) (Rexo1) alternative variant bSep08, mRNA.
<a href="#">Rexo1</a>	<a href="#">Rexo1.cSep08</a>	<a href="#">314630</a>	1882	1047	6	191	REX1, RNA exonuclease 1 homolog (S. cerevisiae) (Rexo1) alternative variant cSep08, mRNA.
<a href="#">Rexo1</a>	<a href="#">Rexo1.dSep08</a>	<a href="#">314630</a>	1521	404	3	62	REX1, RNA exonuclease 1 homolog (S. cerevisiae) (Rexo1) alternative variant dSep08, mRNA.
<a href="#">Rexo2</a>	<a href="#">Rexo2.bSep08</a>	<a href="#">300689</a>	7128	1882	3	153	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (Rexo2) alternative variant bSep08, mRNA.
<a href="#">Rexo2</a>	<a href="#">Rexo2.cSep08</a>	<a href="#">300689</a>	8160	667	6	145	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (17.0 kD) (Rexo2) alternative variant cSep08, mRNA.



<a href="#">Rexo2</a>	<a href="#">Rexo2.dSep08</a>	<a href="#">300689</a>	6627	615	4	117	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (Rexo2) alternative variant dSep08, mRNA.
<a href="#">Rexo2</a>	<a href="#">Rexo2.eSep08</a>	<a href="#">300689</a>	6430	534	4	82	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (Rexo2) alternative variant eSep08, mRNA.
<a href="#">Rexo2</a>	<a href="#">Rexo2.fSep08</a>	<a href="#">300689</a>	1054	430	2	74	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (Rexo2) alternative variant fSep08, mRNA.
<a href="#">reyby</a>	<a href="#">reyby.aSep08</a>		1368	616		30	putative protein (reyby) mRNA.
<a href="#">reychy</a>	<a href="#">reychy.aSep08</a>		32747	1778	5	552	CRA a like (reychy) alternative variant aSep08, mRNA.
<a href="#">reychy</a>	<a href="#">reychy.cSep08</a>		1302	364	2	115	CRA a (reychy) alternative variant cSep08, mRNA.
<a href="#">reychy</a>	<a href="#">reychy.dSep08</a>		30010	474	4	44	putative protein (reychy) alternative variant dSep08, mRNA.
<a href="#">reydar</a>	<a href="#">reydar.bSep08</a>		12459	643	2	42	putative protein (5.0 kD) (reydar) alternative variant bSep08, mRNA.
<a href="#">reyfer</a>	<a href="#">reyfer.aSep08</a>		3743	616		55	putative protein (5.8 kD) (reyfer) mRNA.
<a href="#">reyflo</a>	<a href="#">reyflo.aSep08</a>		69617	244		81	VPS10 domain receptor (reyflo) mRNA.
<a href="#">reyflu</a>	<a href="#">reyflu.aSep08</a>		2099	606		84	tyrosine kinase receptor (reyflu) mRNA.
<a href="#">reygar</a>	<a href="#">reygar.aSep08</a>		1181	387		128	elastin microfibril interfacier 3 (reygar) mRNA.
<a href="#">reyja</a>	<a href="#">reyja.aSep08</a>		3537	737	2	86	putative cytoplasmic protein (9.8 kD) (reyja) alternative variant aSep08, mRNA.
<a href="#">reyja</a>	<a href="#">reyja.bSep08</a>		1028	244	1	25	putative protein (reyja) alternative variant bSep08, mRNA.
<a href="#">reyjey</a>	<a href="#">reyjey.aSep08</a>		652	550		52	CRA a like (5.9 kD) (reyjey) mRNA.
<a href="#">reykee</a>	<a href="#">reykee.aSep08</a>		6144	481		37	putative protein (4.2 kD) (reykee) mRNA.
<a href="#">reykler</a>	<a href="#">reykler.bSep08</a>		1987	257		35	putative protein (reykler) alternative variant bSep08, mRNA.
<a href="#">reylo</a>	<a href="#">reylo.aSep08</a>		1482	807		64	putative protein (reylo) mRNA.
<a href="#">reymee</a>	<a href="#">reymee.aSep08</a>		2498	402		134	domain 19 (reymee) mRNA.
<a href="#">reymmer</a>	<a href="#">reymmer.aSep08</a>		1468	479	2	123	putative protein (reymmer) alternative variant aSep08, mRNA.
<a href="#">reynoy</a>	<a href="#">reynoy.aSep08</a>		7615	753		48	putative protein (reynoy) mRNA.
<a href="#">reypor</a>	<a href="#">reypor.aSep08</a>		4319	340	3	112	family with sequence similarity 63 member B CRA b like (reypor) alternative variant aSep08, mRNA.
<a href="#">reysa</a>	<a href="#">reysa.aSep08</a>		1153	338		112	putative protein (reysa) mRNA.
<a href="#">reyshee</a>	<a href="#">reyshee.aSep08</a>		2580	329		54	putative protein (reyshee) mRNA.
<a href="#">reytu</a>	<a href="#">reytu.aSep08</a>		918	613		154	putative mitochondrial protein (17.4 kD) (reytu) mRNA.
<a href="#">reyvar</a>	<a href="#">reyvar.aSep08</a>		2516	274		91	putative protein, with a transmembrane domain (reyvar) mRNA.
<a href="#">reywey</a>	<a href="#">reywey.aSep08</a>		2659	1125	2	60	ubiquitin specific peptidase 18 CRA a (7.2 kD) (reywey) alternative variant aSep08, mRNA.
<a href="#">reywey</a>	<a href="#">reywey.bSep08</a>		1874	665	1	47	putative protein (reywey) alternative variant bSep08, mRNA.
<a href="#">Rfc1</a>	<a href="#">Rfc1.bSep08</a>	<a href="#">89809</a>	9194	1051	6	296	replication factor C (activator 1) 1 (Rfc1) alternative variant bSep08, mRNA.
<a href="#">Rfc1</a>	<a href="#">Rfc1.cSep08</a>	<a href="#">89809</a>	3228	342	2	86	replication factor C (activator 1) 1 (Rfc1) alternative variant cSep08, mRNA.

<a href="#">Rfc1</a>	<a href="#">Rfc1.eSep08</a>	<a href="#">89809</a>	30060	379	4	57	replication factor C (activator 1) 1 (Rfc1) alternative variant eSep08, mRNA.
<a href="#">Rfc2</a>	<a href="#">Rfc2.bSep08</a>	<a href="#">116468</a>	5595	1048	1	152	replication factor C (activator 1) 2 (17.1 kD) (Rfc2) alternative variant bSep08, mRNA.
<a href="#">Rfc4</a>	<a href="#">Rfc4.bSep08</a>	<a href="#">288003</a>	15301	1347	10	223	replication factor C (activator 1) 4 (24.6 kD) (Rfc4) alternative variant bSep08, complete mRNA.
<a href="#">Rfc4</a>	<a href="#">Rfc4.cSep08</a>	<a href="#">288003</a>	13951	534	5	178	replication factor C (activator 1) 4 (Rfc4) alternative variant cSep08, mRNA.
<a href="#">Rfc4</a>	<a href="#">Rfc4.eSep08</a>	<a href="#">288003</a>	800	611	3	22	replication factor C (activator 1) 4 (2.6 kD) (Rfc4) alternative variant eSep08, mRNA.
<a href="#">Rfc5</a>	<a href="#">Rfc5.bSep08</a>	<a href="#">304528</a>	7054	783	8	205	replication factor C (activator 1) 5 (Rfc5) alternative variant bSep08, mRNA.
<a href="#">Rfc5</a>	<a href="#">Rfc5.cSep08</a>	<a href="#">304528</a>	1337	358	2	25	replication factor C (activator 1) 5 (Rfc5) alternative variant cSep08, mRNA.
<a href="#">Rffl</a>	<a href="#">Rffl.aSep08</a>	<a href="#">282844</a>	60616	1102	2	367	putative protein of metazoan origin (Rffl) alternative variant aSep08, mRNA.
<a href="#">Rft1</a>	<a href="#">Rft1.aSep08</a>	<a href="#">290552</a>	13108	1419		308	RFT1 homolog ( <i>S. cerevisiae</i> ) (Rft1) mRNA.
<a href="#">Rftn2</a>	<a href="#">Rftn2.aSep08</a>	<a href="#">363231</a>	35628	1264	1	348	raftlin family member 2 (Rftn2) alternative variant aSep08, mRNA.
<a href="#">Rftn2</a>	<a href="#">Rftn2.bSep08</a>	<a href="#">363231</a>	811	386	1	47	raftlin family member 2 (Rftn2) alternative variant bSep08, mRNA.
<a href="#">Rfwd3</a>	<a href="#">Rfwd3.aSep08</a>	<a href="#">361409</a>	7212	2719		252	ring finger and WD repeat domain 3 (Rfwd3) mRNA.
<a href="#">Rfx2</a>	<a href="#">Rfx2.bSep08</a>	<a href="#">301121</a>	8162	2159	9	361	regulatory factor X, 2 (influences HLA class II expression) (Rfx2) alternative variant bSep08, mRNA.
<a href="#">Rfx2</a>	<a href="#">Rfx2.cSep08</a>	<a href="#">301121</a>	22188	807	1	269	regulatory factor X, 2 (influences HLA class II expression) (Rfx2) alternative variant cSep08, mRNA.
<a href="#">Rfx3</a>	<a href="#">Rfx3.bSep08</a>	<a href="#">361746</a>	77865	1155	2	385	regulatory factor X, 3 (influences HLA class II expression) (Rfx3) alternative variant bSep08, mRNA.
<a href="#">Rfx4</a>	<a href="#">Rfx4.aSep08</a>	<a href="#">500818</a>	132709	1509	13	502	regulatory factor X, 4 (influences HLA class II expression) (Rfx4) alternative variant aSep08, mRNA.
<a href="#">Rfx4</a>	<a href="#">Rfx4.bSep08</a>	<a href="#">500818</a>	27809	537	1	140	regulatory factor X, 4 (influences HLA class II expression) (Rfx4) alternative variant bSep08, mRNA.
<a href="#">Rfx5</a>	<a href="#">Rfx5.bSep08</a>	<a href="#">310659</a>	3444	3372	2	351	regulatory factor X, 5 (influences HLA class II expression) (35.6 kD) (Rfx5) alternative variant bSep08, mRNA.
<a href="#">Rfx5</a>	<a href="#">Rfx5.cSep08</a>	<a href="#">310659</a>	2292	366	5	98	regulatory factor X, 5 (influences HLA class II expression) (Rfx5) alternative variant cSep08, mRNA.
<a href="#">Rfx5</a>	<a href="#">Rfx5.dSep08</a>	<a href="#">310659</a>	1104	417	4	75	regulatory factor X, 5 (influences HLA class II expression) (Rfx5) alternative variant dSep08, mRNA.
<a href="#">Rfxank</a>	<a href="#">Rfxank.bSep08</a>	<a href="#">306353</a>	7422	1348	8	211	regulatory factor X-associated ankyrin-containing protein (23.1 kD) (Rfxank) alternative variant bSep08, mRNA.
<a href="#">Rfxdc1</a>	<a href="#">Rfxdc1.bSep08</a>	<a href="#">294395</a>	8062	1775	1	381	regulatory factor X (Rfxdc1) alternative variant bSep08, mRNA.
<a href="#">Rfxdc1</a>	<a href="#">Rfxdc1.cSep08</a>	<a href="#">294395</a>	4327	700	2	211	regulatory factor X (Rfxdc1) alternative variant cSep08, mRNA.
<a href="#">Rfxdc2</a>	<a href="#">Rfxdc2.bSep08</a>	<a href="#">315804</a>	74603	1459	7	302	regulatory factor X (Rfxdc2) alternative variant bSep08, mRNA.

<a href="#">Rfxdc2</a>	<a href="#">Rfxdc2.cSep08</a>	<a href="#">315804</a>	631	469	2	41	putative protein (Rfxdc2) alternative variant cSep08, mRNA.
<a href="#">Rg9mtd2</a>	<a href="#">Rg9mtd2.bSep08</a>	<a href="#">295496</a>	14255	2711	7	285	tRNA (guanine-N1-)-methyltransferase (32.7 kD) (Rg9mtd2) alternative variant bSep08, mRNA.
<a href="#">Rg9mtd2</a>	<a href="#">Rg9mtd2.cSep08</a>	<a href="#">295496</a>	2583	854	2	149	putative protein of eukaryotic origin (16.5 kD) (Rg9mtd2) alternative variant cSep08, mRNA.
<a href="#">Rg9mtd2</a>	<a href="#">Rg9mtd2.dSep08</a>	<a href="#">295496</a>	2908	391	3	99	tRNA (guanine-N1-)-methyltransferase (Rg9mtd2) alternative variant dSep08, mRNA.
<a href="#">Rg9mtd3</a>	<a href="#">Rg9mtd3.bSep08</a>	<a href="#">298081</a>	1118	571	2	111	putative protein of eukaryotic origin (13.0 kD) (Rg9mtd3) alternative variant bSep08, mRNA.
<a href="#">RGD621352</a>	<a href="#">RGD621352.bSep08</a>	<a href="#">192229</a>	8049	872	4	158	similar to RIKEN cDNA 1500031L02 (RGD621352) alternative variant bSep08, mRNA.
<a href="#">RGD621352</a>	<a href="#">RGD621352.cSep08</a>	<a href="#">192229</a>	3144	590	2	156	similar to RIKEN cDNA 1500031L02 (RGD621352) alternative variant cSep08, mRNA.
<a href="#">RGD621352</a>	<a href="#">RGD621352.dSep08</a>	<a href="#">192229</a>	7776	584	4	137	similar to RIKEN cDNA 1500031L02 (RGD621352) alternative variant dSep08, mRNA.
<a href="#">RGD621352</a>	<a href="#">RGD621352.eSep08</a>	<a href="#">192229</a>	3265	2273	2	78	similar to RIKEN cDNA 1500031L02 (9.0 kD) (RGD621352) alternative variant eSep08, mRNA.
<a href="#">RGD708545</a>	<a href="#">RGD708545.cSep08</a>	<a href="#">56769</a>	979	818	3	101	nuclear protein E3-3 (10.9 kD) (RGD708545) alternative variant cSep08, mRNA.
<a href="#">RGD735029</a>	<a href="#">RGD735029.bSep08</a>	<a href="#">307480</a>	4252	609	5	202	putative protein of mammalian origin (RGD735029) alternative variant bSep08, mRNA.
<a href="#">RGD735029</a>	<a href="#">RGD735029.cSep08</a>	<a href="#">307480</a>	1672	763	2	90	CRA c like (RGD735029) alternative variant cSep08, mRNA.
<a href="#">RGD735029</a>	<a href="#">RGD735029.dSep08</a>	<a href="#">307480</a>	3617	1017	4	91	CRA c like (9.3 kD) (RGD735029) alternative variant dSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.aSep08</a>	<a href="#">294311</a>	26345	2375	6	595	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant aSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.bSep08</a>	<a href="#">294311</a>	30064	2743	8	348	similar to GI:13385412-like protein splice form I (39.6 kD) (RGD735065) alternative variant bSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.dSep08</a>	<a href="#">294311</a>	39221	1381	9	235	similar to GI:13385412-like protein splice form I (26.4 kD) (RGD735065) alternative variant dSep08, complete mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.eSep08</a>	<a href="#">294311</a>	20690	1193	4	185	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant eSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.fSep08</a>	<a href="#">294311</a>	31033	553	3	171	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant fSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.gSep08</a>	<a href="#">294311</a>	19887	780	2	101	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant gSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.hSep08</a>	<a href="#">294311</a>	19877	862	3	59	similar to GI:13385412-like protein splice form I (6.8 kD) (RGD735065) alternative variant hSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.iSep08</a>	<a href="#">294311</a>	19685	765	4	130	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant iSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.jSep08</a>	<a href="#">294311</a>	23894	725	5	82	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant jSep08, mRNA.
<a href="#">RGD735065</a>	<a href="#">RGD735065.kSep08</a>	<a href="#">294311</a>	9293	701	4	134	similar to GI:13385412-like protein splice form I (RGD735065) alternative variant kSep08, mRNA.

<a href="#">RGD735112</a>	<a href="#">RGD735112.bSep08</a>	<a href="#">304055</a>	14150	603	2	101	similar to RIKEN cDNA 5830404H04 (RGD735112) alternative variant bSep08, mRNA.
<a href="#">RGD735112</a>	<a href="#">RGD735112.cSep08</a>	<a href="#">304055</a>	4280	365	3	86	similar to RIKEN cDNA 5830404H04 (RGD735112) alternative variant cSep08, mRNA.
<a href="#">RGD735140</a>	<a href="#">RGD735140.bSep08</a>	<a href="#">362219</a>	12406	1256	10	297	hypothetical protein LK44 (RGD735140) alternative variant bSep08, mRNA.
<a href="#">RGD735140</a>	<a href="#">RGD735140.eSep08</a>	<a href="#">362219</a>	24747	545	5	37	hypothetical protein LK44 (RGD735140) alternative variant eSep08, mRNA.
<a href="#">RGD735140</a>	<a href="#">RGD735140.fSep08</a>	<a href="#">362219</a>	10000	386	3		
<a href="#">RGD735175</a>	<a href="#">RGD735175.aSep08</a>	<a href="#">316530</a>	6434	2771	3	408	hypothetical protein MGC:72616 (45.5 kD) (RGD735175) alternative variant aSep08, mRNA.
<a href="#">RGD735175</a>	<a href="#">RGD735175.cSep08</a>	<a href="#">316530</a>	4329	764	3	254	hypothetical protein MGC:72616 (RGD735175) alternative variant cSep08, mRNA.
<a href="#">RGD735175</a>	<a href="#">RGD735175.dSep08</a>	<a href="#">316530</a>	3644	648	1	216	hypothetical protein MGC:72616 (RGD735175) alternative variant dSep08, mRNA.
<a href="#">RGD1302996</a>	<a href="#">RGD1302996.bSep08</a>	<a href="#">294231</a>	1810	652	3	121	hypothetical protein MGC:15854 (14.2 kD) (RGD1302996) alternative variant bSep08, mRNA.
<a href="#">RGD1302996</a>	<a href="#">RGD1302996.cSep08</a>	<a href="#">294231</a>	1712	637	2	99	hypothetical protein MGC:15854 (RGD1302996) alternative variant cSep08, mRNA.
<a href="#">RGD1303003</a>	<a href="#">RGD1303003.bSep08</a>	<a href="#">294326</a>	7528	721	1	142	homolog of zebrafish ES1 (RGD1303003) alternative variant bSep08, mRNA.
<a href="#">RGD1303066</a>	<a href="#">RGD1303066.bSep08</a>	<a href="#">361789</a>	12948	1300	11	364	similar to RIKEN cDNA 2610110G12 (RGD1303066) alternative variant bSep08, mRNA.
<a href="#">RGD1303066</a>	<a href="#">RGD1303066.cSep08</a>	<a href="#">361789</a>	1450	702	5	234	similar to RIKEN cDNA 2610110G12 (RGD1303066) alternative variant cSep08, mRNA.
<a href="#">RGD1303066</a>	<a href="#">RGD1303066.dSep08</a>	<a href="#">361789</a>	8195	584	8	188	similar to RIKEN cDNA 2610110G12 (RGD1303066) alternative variant dSep08, mRNA.
<a href="#">RGD1303066</a>	<a href="#">RGD1303066.eSep08</a>	<a href="#">361789</a>	874	401	2	86	similar to RIKEN cDNA 2610110G12 (RGD1303066) alternative variant eSep08, mRNA.
<a href="#">RGD1303117</a>	<a href="#">RGD1303117.aSep08</a>	<a href="#">292764</a>	4216	1842	9	361	CRA a (RGD1303117) alternative variant aSep08, mRNA.
<a href="#">RGD1303117</a>	<a href="#">RGD1303117.cSep08</a>	<a href="#">292764</a>	798	625	3	147	CRA a (RGD1303117) alternative variant cSep08, mRNA.
<a href="#">RGD1303117</a>	<a href="#">RGD1303117.dSep08</a>	<a href="#">292764</a>	1054	718	2	121	putative cytoplasmic protein of mammalian origin (13.3 kD) (RGD1303117) alternative variant dSep08, mRNA.
<a href="#">RGD1303117</a>	<a href="#">RGD1303117.eSep08</a>	<a href="#">292764</a>	1844	630	4	102	putative protein of mammalian origin (11.3 kD) (RGD1303117) alternative variant eSep08, mRNA.
<a href="#">RGD1303117</a>	<a href="#">RGD1303117.gSep08</a>	<a href="#">292764</a>	753	655	2	64	CRA a like (6.4 kD) (RGD1303117) alternative variant gSep08, mRNA.
<a href="#">RGD1303127</a>	<a href="#">RGD1303127.bSep08</a>	<a href="#">300206</a>	16712	2870	8	259	similar to hypothetical protein FLJ20436 (29.2 kD) (RGD1303127) alternative variant bSep08, mRNA.
<a href="#">RGD1303127</a>	<a href="#">RGD1303127.cSep08</a>	<a href="#">300206</a>	2801	390	3	129	similar to hypothetical protein FLJ20436 (RGD1303127) alternative variant cSep08, mRNA.
<a href="#">RGD1303127</a>	<a href="#">RGD1303127.dSep08</a>	<a href="#">300206</a>	4772	957	2	59	similar to hypothetical protein FLJ20436 (RGD1303127) alternative variant dSep08, mRNA.
<a href="#">RGD1303130</a>	<a href="#">RGD1303130.bSep08</a>	<a href="#">295231</a>	2854	1020	4	205	kidney predominant protein NCU-G1 (RGD1303130) alternative variant bSep08, mRNA.
<a href="#">RGD1303144</a>	<a href="#">RGD1303144.bSep08</a>	<a href="#">311833</a>	39555	651	8	147	CRA a (RGD1303144) alternative variant bSep08, mRNA.

<a href="#">RGD1303232</a>	<a href="#">RGD1303232.bSep08</a>	<a href="#">309381</a>	7922	2792	6	254	probable oxidoreductase (27.0 kD) (RGD1303232) alternative variant bSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.cSep08</a>	<a href="#">309381</a>	4574	464	4	154	probable oxidoreductase (RGD1303232) alternative variant cSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.dSep08</a>	<a href="#">309381</a>	5973	830	5	135	probable oxidoreductase (RGD1303232) alternative variant dSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.fSep08</a>	<a href="#">309381</a>	2756	552	4	106	probable oxidoreductase (RGD1303232) alternative variant fSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.gSep08</a>	<a href="#">309381</a>	3423	464	2	104	putative protein (RGD1303232) alternative variant gSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.hSep08</a>	<a href="#">309381</a>	3126	551	4	102	probable oxidoreductase (11.0 kD) (RGD1303232) alternative variant hSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.iSep08</a>	<a href="#">309381</a>	2320	557	3	64	probable oxidoreductase (RGD1303232) alternative variant iSep08, mRNA.
<a href="#">RGD1303232</a>	<a href="#">RGD1303232.jSep08</a>	<a href="#">309381</a>	3074	1213	3	135	putative protein (14.3 kD) (RGD1303232) alternative variant jSep08, mRNA.
<a href="#">RGD1303271</a>	<a href="#">RGD1303271.bSep08</a>	<a href="#">313018</a>	8283	436	1	60	putative protein of mammalian origin (RGD1303271) alternative variant bSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.bSep08</a>	<a href="#">362134</a>	14363	913	6	235	similar to RIKEN cDNA 2010311D03 (26.5 kD) (RGD1303272) alternative variant bSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.cSep08</a>	<a href="#">362134</a>	8927	504	5	168	similar to RIKEN cDNA 2010311D03 (RGD1303272) alternative variant cSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.dSep08</a>	<a href="#">362134</a>	5514	767	3	124	similar to RIKEN cDNA 2010311D03 (RGD1303272) alternative variant dSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.eSep08</a>	<a href="#">362134</a>	1613	845	2	88	similar to RIKEN cDNA 2010311D03 (9.9 kD) (RGD1303272) alternative variant eSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.fSep08</a>	<a href="#">362134</a>	7606	750	2	50	similar to RIKEN cDNA 2010311D03 (RGD1303272) alternative variant fSep08, mRNA.
<a href="#">RGD1303272</a>	<a href="#">RGD1303272.gSep08</a>	<a href="#">362134</a>	10071	952	7	140	similar to RIKEN cDNA 2010311D03 (RGD1303272) alternative variant gSep08, mRNA.
<a href="#">RGD1304567</a>	<a href="#">RGD1304567.bSep08</a>	<a href="#">362671</a>	3588	724	3	216	similar to RIKEN cDNA A430005L14 (22.8 kD) (RGD1304567) alternative variant bSep08, complete mRNA.
<a href="#">RGD1304567</a>	<a href="#">RGD1304567.cSep08</a>	<a href="#">362671</a>	4239	777	3	215	similar to RIKEN cDNA A430005L14 (RGD1304567) alternative variant cSep08, mRNA.
<a href="#">RGD1304579</a>	<a href="#">RGD1304579.bSep08</a>	<a href="#">308906</a>	8589	1515	5	491	similar to 9230105E10Rik protein (RGD1304579) alternative variant bSep08, mRNA.
<a href="#">RGD1304579</a>	<a href="#">RGD1304579.cSep08</a>	<a href="#">308906</a>	9011	658	2	172	similar to 9230105E10Rik protein (RGD1304579) alternative variant cSep08, mRNA.
<a href="#">RGD1304580</a>	<a href="#">RGD1304580.bSep08</a>	<a href="#">292781</a>	2325	384		114	similar to Hypothetical protein MGC38513 (RGD1304580) alternative variant bSep08, mRNA.
<a href="#">RGD1304592</a>	<a href="#">RGD1304592.bSep08</a>	<a href="#">362461</a>	56267	3136	18	700	similar to KIAA0528 protein (RGD1304592) alternative variant bSep08, mRNA.
<a href="#">RGD1304592</a>	<a href="#">RGD1304592.cSep08</a>	<a href="#">362461</a>	7379	1644	5	203	similar to KIAA0528 protein (RGD1304592) alternative variant cSep08, mRNA.
<a href="#">RGD1304592</a>	<a href="#">RGD1304592.dSep08</a>	<a href="#">362461</a>	17619	709	5	136	similar to KIAA0528 protein (RGD1304592) alternative variant dSep08, mRNA.

<a href="#">RGD1304592</a>	<a href="#">RGD1304592.eSep08</a>	<a href="#">362461</a>	19555	414	4	106	similar to KIAA0528 protein (RGD1304592) alternative variant eSep08, mRNA.
<a href="#">RGD1304592</a>	<a href="#">RGD1304592.gSep08</a>	<a href="#">362461</a>	4504	824	3	52	similar to KIAA0528 protein (RGD1304592) alternative variant gSep08, mRNA.
<a href="#">RGD1304607</a>	<a href="#">RGD1304607.aSep08</a>	<a href="#">361727</a>	5148	1314	3	147	similar to thymus atrophy-related protein (RGD1304607) alternative variant aSep08, mRNA.
<a href="#">RGD1304607</a>	<a href="#">RGD1304607.bSep08</a>	<a href="#">361727</a>	3088	502	1	134	similar to thymus atrophy-related protein (RGD1304607) alternative variant bSep08, mRNA.
<a href="#">RGD1304607</a>	<a href="#">RGD1304607.cSep08</a>	<a href="#">361727</a>	4838	1018	3	87	similar to thymus atrophy-related protein (9.9 kD) (RGD1304607) alternative variant cSep08, mRNA.
<a href="#">RGD1304621</a>	<a href="#">RGD1304621.aSep08</a>	<a href="#">304104</a>	5236	1929		416	similar to Hypothetical protein KIAA0539 (46.3 kD) (RGD1304621) mRNA.
<a href="#">RGD1304622</a>	<a href="#">RGD1304622.aSep08</a>	<a href="#">305101</a>	10237	604		201	similar to 6820428L09 protein (RGD1304622) mRNA.
<a href="#">RGD1304624</a>	<a href="#">RGD1304624.bSep08</a>	<a href="#">314128</a>	27861	934	3	72	similar to RIKEN cDNA 2700097O09 (RGD1304624) alternative variant bSep08, mRNA.
<a href="#">RGD1304644</a>	<a href="#">RGD1304644.bSep08</a>	<a href="#">311536</a>	5979	782	4	203	similar to RIKEN cDNA 2310046K01 (RGD1304644) alternative variant bSep08, mRNA.
<a href="#">RGD1304644</a>	<a href="#">RGD1304644.cSep08</a>	<a href="#">311536</a>	3697	969	4	111	similar to RIKEN cDNA 2310046K01 (RGD1304644) alternative variant cSep08, mRNA.
<a href="#">RGD1304644</a>	<a href="#">RGD1304644.eSep08</a>	<a href="#">311536</a>	12220	1800	4	57	similar to RIKEN cDNA 2310046K01 (6.7 kD) (RGD1304644) alternative variant eSep08, mRNA.
<a href="#">RGD1304652</a>	<a href="#">RGD1304652.aSep08</a>	<a href="#">690349</a>	1973	632	2	135	similar to RIKEN cDNA 1810009A15 (RGD1304652) alternative variant aSep08, mRNA.
<a href="#">RGD1304652</a>	<a href="#">RGD1304652.cSep08</a>	<a href="#">690349</a>	1686	507	1	94	similar to RIKEN cDNA 1810009A15 (RGD1304652) alternative variant cSep08, mRNA.
<a href="#">RGD1304693</a>	<a href="#">RGD1304693.aSep08</a>	<a href="#">313108</a>	115007	1785	8	489	putative protein, with at least 2 transmembrane domains, of metazoan origin (RGD1304693) alternative variant aSep08, mRNA.
<a href="#">RGD1304693</a>	<a href="#">RGD1304693.bSep08</a>	<a href="#">313108</a>	50585	1203	5	390	putative protein of metazoan origin (RGD1304693) alternative variant bSep08, mRNA.
<a href="#">RGD1304693</a>	<a href="#">RGD1304693.cSep08</a>	<a href="#">313108</a>	17633	1299	7	347	putative protein (RGD1304693) alternative variant cSep08, mRNA.
<a href="#">RGD1304693</a>	<a href="#">RGD1304693.dSep08</a>	<a href="#">313108</a>	5995	1328	5	202	putative protein of metazoan origin (22.7 kD) (RGD1304693) alternative variant dSep08, mRNA.
<a href="#">RGD1304694</a>	<a href="#">RGD1304694.aSep08</a>	<a href="#">362974</a>	21186	2427	10	370	similar to CG9646-PA (41.9 kD) (RGD1304694) alternative variant aSep08, mRNA.
<a href="#">RGD1304694</a>	<a href="#">RGD1304694.bSep08</a>	<a href="#">362974</a>	28284	600	5	199	similar to CG9646-PA (RGD1304694) alternative variant bSep08, mRNA.
<a href="#">RGD1304694</a>	<a href="#">RGD1304694.dSep08</a>	<a href="#">362974</a>	4197	341	2	108	similar to CG9646-PA (RGD1304694) alternative variant dSep08, mRNA.
<a href="#">RGD1304704</a>	<a href="#">RGD1304704.bSep08</a>	<a href="#">302247</a>	10719	1006	8	278	homeobox prox 1 (RGD1304704) alternative variant bSep08, mRNA.
<a href="#">RGD1304704</a>	<a href="#">RGD1304704.cSep08</a>	<a href="#">302247</a>	7967	1921	5	161	putative protein of eukaryotic origin (19.0 kD) (RGD1304704) alternative variant cSep08, mRNA.
<a href="#">RGD1304704</a>	<a href="#">RGD1304704.dSep08</a>	<a href="#">302247</a>	5074	718	5	110	homeobox prox 1 (12.3 kD) (RGD1304704) alternative variant dSep08, mRNA.

<a href="#">RGD1304704</a>	<a href="#">RGD1304704.eSep08</a>	<a href="#">302247</a>	2282	274	3	46	homeobox prox 1 (RGD1304704) alternative variant eSep08, mRNA.
<a href="#">RGD1304704</a>	<a href="#">RGD1304704.fSep08</a>	<a href="#">302247</a>	777	660	2	33	putative protein (3.5 kD) (RGD1304704) alternative variant fSep08, mRNA.
<a href="#">RGD1304704</a>	<a href="#">RGD1304704.gSep08</a>	<a href="#">302247</a>	2794	328	3	24	putative protein (RGD1304704) alternative variant gSep08, mRNA.
<a href="#">RGD1304728</a>	<a href="#">RGD1304728.aSep08</a>	<a href="#">360560</a>	4223	427		141	similar to 4933427D14Rik protein (RGD1304728) mRNA.
<a href="#">RGD1304731</a>	<a href="#">RGD1304731.bSep08</a>	<a href="#">291549</a>	13847	956	5	141	CRA d (15.3 kD) (RGD1304731) alternative variant bSep08, mRNA.
<a href="#">RGD1304731</a>	<a href="#">RGD1304731.cSep08</a>	<a href="#">291549</a>	4109	456	3	123	CRA d (RGD1304731) alternative variant cSep08, mRNA.
<a href="#">RGD1304737</a>	<a href="#">RGD1304737.aSep08</a>	<a href="#">314639</a>	8970	548	5	182	similar to KIAA1086 protein (RGD1304737) alternative variant aSep08, mRNA.
<a href="#">RGD1304737</a>	<a href="#">RGD1304737.bSep08</a>	<a href="#">314639</a>	2193	380	2	110	similar to KIAA1086 protein (RGD1304737) alternative variant bSep08, mRNA.
<a href="#">RGD1304774</a>	<a href="#">RGD1304774.bSep08</a>	<a href="#">360969</a>	1814	1199	4	134	protein CRA a (RGD1304774) alternative variant bSep08, mRNA.
<a href="#">RGD1304774</a>	<a href="#">RGD1304774.cSep08</a>	<a href="#">360969</a>	2741	682	4	130	opposite strand transcription unit Stag3 (RGD1304774) alternative variant cSep08, mRNA.
<a href="#">RGD1304792</a>	<a href="#">RGD1304792.bSep08</a>	<a href="#">312474</a>	20136	1795	7	598	putative protein, with a coiled coil domain, of metazoan origin (RGD1304792) alternative variant bSep08, mRNA.
<a href="#">RGD1304792</a>	<a href="#">RGD1304792.cSep08</a>	<a href="#">312474</a>	7087	576	3	192	putative protein of mammalian origin (RGD1304792) alternative variant cSep08, mRNA.
<a href="#">RGD1304792</a>	<a href="#">RGD1304792.dSep08</a>	<a href="#">312474</a>	2192	1235	2	66	putative protein of mammalian origin (7.4 kD) (RGD1304792) alternative variant dSep08, mRNA.
<a href="#">RGD1304793</a>	<a href="#">RGD1304793.bSep08</a>	<a href="#">313021</a>	11692	1492	6	166	similar to hypothetical protein DKFZp564D0478 (18.9 kD) (RGD1304793) alternative variant bSep08, complete mRNA.
<a href="#">RGD1304793</a>	<a href="#">RGD1304793.cSep08</a>	<a href="#">313021</a>	2795	1436	3	157	similar to hypothetical protein DKFZp564D0478 (18.0 kD) (RGD1304793) alternative variant cSep08, mRNA.
<a href="#">RGD1304808</a>	<a href="#">RGD1304808.aSep08</a>	<a href="#">316440</a>	11008	400		115	similar to hypothetical protein FLJ20309 (RGD1304808) mRNA.
<a href="#">RGD1304810</a>	<a href="#">RGD1304810.bSep08</a>	<a href="#">306504</a>	34554	795	3	96	similar to 6430573F11Rik protein and similar to Ubiquitin-conjugating enzyme E2 E3 (Ubiquitin-protein ligase E3) (Ubiquitin carrier protein E3) (Ubiquitin-conjugating enzyme E2-23 kDa) (UbcM2) (RGD1304810) alternative variant bSep08, mRNA.
<a href="#">RGD1304810</a>	<a href="#">RGD1304810.bSep08</a>	<a href="#">498640</a>	34554	795	3	96	similar to 6430573F11Rik protein and similar to Ubiquitin-conjugating enzyme E2 E3 (Ubiquitin-protein ligase E3) (Ubiquitin carrier protein E3) (Ubiquitin-conjugating enzyme E2-23 kDa) (UbcM2) (RGD1304810) alternative variant bSep08, mRNA.
<a href="#">RGD1304810</a>	<a href="#">RGD1304810.cSep08</a>	<a href="#">306504</a>	2313	588	2	42	similar to 6430573F11Rik protein and similar to Ubiquitin-conjugating enzyme E2 E3 (Ubiquitin-protein ligase E3) (Ubiquitin carrier protein E3) (Ubiquitin-conjugating enzyme E2-23 kDa) (UbcM2) (4.9 kD) (RGD1304810) alternative variant cSep08, mRNA.

<a href="#">RGD1304810</a>	<a href="#">RGD1304810.cSep08</a>	<a href="#">498640</a>	2313	588	2	42	similar to 6430573F11Rik protein and similar to Ubiquitin-conjugating enzyme E2 E3 (Ubiquitin-protein ligase E3) (Ubiquitin carrier protein E3) (Ubiquitin-conjugating enzyme E2-23 kDa) (UbcM2) (4.9 kD) (RGD1304810) alternative variant cSep08, mRNA.
<a href="#">RGD1304822</a>	<a href="#">RGD1304822.aSep08</a>	<a href="#">295428</a>	16152	2609	11	493	similar to KIAA1627 protein (RGD1304822) alternative variant aSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.aSep08</a>	<a href="#">296550</a>	8678	1740	8	123	repeat domain 85 (13.4 kD) (RGD1304868) alternative variant aSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.bSep08</a>	<a href="#">296550</a>	3511	621	5	206	repeat domain 85 (RGD1304868) alternative variant bSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.cSep08</a>	<a href="#">296550</a>	2752	1385	2	162	putative protein of vertebrate origin (18.0 kD) (RGD1304868) alternative variant cSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.fSep08</a>	<a href="#">296550</a>	6220	1387	6	127	repeat domain 85 (RGD1304868) alternative variant fSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.iSep08</a>	<a href="#">296550</a>	844	370	2	33	putative protein (RGD1304868) alternative variant iSep08, mRNA.
<a href="#">RGD1304868</a>	<a href="#">RGD1304868.jSep08</a>	<a href="#">296550</a>	3414	305	3	91	putative protein (RGD1304868) alternative variant jSep08, mRNA.
<a href="#">RGD1304876</a>	<a href="#">RGD1304876.cSep08</a>	<a href="#">362368</a>	3587	648	2	61	similar to RIKEN cDNA A030007L17; EST AA673177 (RGD1304876) alternative variant cSep08, mRNA.
<a href="#">RGD1304878</a>	<a href="#">RGD1304878.aSep08</a>	<a href="#">314415</a>	21454	3902		819	similar to 2410024A21Rik protein (RGD1304878) mRNA.
<a href="#">RGD1304881</a>	<a href="#">RGD1304881.bSep08</a>	<a href="#">304653</a>	2674	1050	3	181	putative protein of eukaryotic origin (RGD1304881) alternative variant bSep08, mRNA.
<a href="#">RGD1304881</a>	<a href="#">RGD1304881.cSep08</a>	<a href="#">304653</a>	1862	1252	6	169	putative protein of vertebrate origin (19.4 kD) (RGD1304881) alternative variant cSep08, mRNA.
<a href="#">RGD1304881</a>	<a href="#">RGD1304881.dSep08</a>	<a href="#">304653</a>	2779	756	5	122	putative protein of eukaryotic origin (RGD1304881) alternative variant dSep08, mRNA.
<a href="#">RGD1304881</a>	<a href="#">RGD1304881.eSep08</a>	<a href="#">304653</a>	2331	790	2	77	putative protein of vertebrate origin (RGD1304881) alternative variant eSep08, mRNA.
<a href="#">RGD1304884</a>	<a href="#">RGD1304884.bSep08</a>	<a href="#">307907</a>	7447	625	6	139	CRA a (RGD1304884) alternative variant bSep08, mRNA.
<a href="#">RGD1304884</a>	<a href="#">RGD1304884.cSep08</a>	<a href="#">307907</a>	29802	608	2	104	putative protein (RGD1304884) alternative variant cSep08, mRNA.
<a href="#">RGD1304884</a>	<a href="#">RGD1304884.dSep08</a>	<a href="#">307907</a>	29930	352	2	94	putative protein (RGD1304884) alternative variant dSep08, mRNA.
<a href="#">RGD1304884</a>	<a href="#">RGD1304884.eSep08</a>	<a href="#">307907</a>	1157	415	2	61	putative protein (RGD1304884) alternative variant eSep08, mRNA.
<a href="#">RGD1304904</a>	<a href="#">RGD1304904.aSep08</a>	<a href="#">296130</a>	1663	678		176	similar to mitochondrial glycerol 3-phosphate acyltransferase (RGD1304904) mRNA.
<a href="#">RGD1304924</a>	<a href="#">RGD1304924.bSep08</a>	<a href="#">302982</a>	2768	1365	6	315	similar to hypothetical protein FLJ31364 (RGD1304924) alternative variant bSep08, mRNA.
<a href="#">RGD1304924</a>	<a href="#">RGD1304924.cSep08</a>	<a href="#">302982</a>	2633	2052	5	194	similar to hypothetical protein FLJ31364 (RGD1304924) alternative variant cSep08, mRNA.
<a href="#">RGD1304924</a>	<a href="#">RGD1304924.dSep08</a>	<a href="#">302982</a>	1593	687	3	179	similar to hypothetical protein FLJ31364 (RGD1304924) alternative variant dSep08, mRNA.
<a href="#">RGD1304924</a>	<a href="#">RGD1304924.eSep08</a>	<a href="#">302982</a>	1187	681	4	136	similar to hypothetical protein FLJ31364 (RGD1304924) alternative variant eSep08, mRNA.



<a href="#">RGD1304924</a>	<a href="#">RGD1304924.fSep08</a>	<a href="#">302982</a>	937	475	2	72	similar to hypothetical protein FLJ31364 (RGD1304924) alternative variant fSep08, mRNA.
<a href="#">RGD1304929</a>	<a href="#">RGD1304929.aSep08</a>	<a href="#">290403</a>	4613	762		217	putative protein of eukaryotic origin (RGD1304929) mRNA.
<a href="#">RGD1304952</a>	<a href="#">RGD1304952.aSep08</a>	<a href="#">312711</a>	4116	2041	3	304	similar to RIKEN cDNA C530028O21 gene (RGD1304952) alternative variant aSep08, mRNA.
<a href="#">RGD1304953</a>	<a href="#">RGD1304953.aSep08</a>	<a href="#">361979</a>	9572	426	1	141	similar to SSTK-interacting protein (RGD1304953) alternative variant aSep08, mRNA.
<a href="#">RGD1304963</a>	<a href="#">RGD1304963.aSep08</a>	<a href="#">313891</a>	4968	1178		119	similar to hypothetical protein MGC38716 (RGD1304963) mRNA.
<a href="#">RGD1304982</a>	<a href="#">RGD1304982.aSep08</a>	<a href="#">289138</a>	32080	1867	4	352	similar to RIKEN cDNA 2810025M15 (RGD1304982) alternative variant aSep08, mRNA.
<a href="#">RGD1304982</a>	<a href="#">RGD1304982.cSep08</a>	<a href="#">289138</a>	18707	735	2	165	similar to RIKEN cDNA 2810025M15 (RGD1304982) alternative variant cSep08, mRNA.
<a href="#">RGD1304999</a>	<a href="#">RGD1304999.bSep08</a>	<a href="#">289833</a>	2375	412	2	69	similar to 4930430E16Rik protein (RGD1304999) alternative variant bSep08, mRNA.
<a href="#">RGD1305007</a>	<a href="#">RGD1305007.cSep08</a>	<a href="#">303749</a>	1355	304	3	58	similar to RIKEN cDNA 1110031I02 (RGD1305007) alternative variant cSep08, mRNA.
<a href="#">RGD1305007</a>	<a href="#">RGD1305007.dSep08</a>	<a href="#">303749</a>	1661	693	2	78	similar to RIKEN cDNA 1110031I02 (8.8 kD) (RGD1305007) alternative variant dSep08, mRNA.
<a href="#">RGD1305020</a>	<a href="#">RGD1305020.bSep08</a>	<a href="#">311575</a>	62219	743	5	171	similar to Hepatocellular carcinoma-associated antigen 58 homolog (RGD1305020) alternative variant bSep08, mRNA.
<a href="#">RGD1305038</a>	<a href="#">RGD1305038.aSep08</a>	<a href="#">314627</a>	4370	1777		525	similar to Serine/threonine-protein kinase SNK (Serum inducible kinase) (RGD1305038) alternative variant aSep08, mRNA.
<a href="#">RGD1305038</a>	<a href="#">RGD1305038.bSep08</a>	<a href="#">314627</a>	2510	753		250	similar to Serine/threonine-protein kinase SNK (Serum inducible kinase) (RGD1305038) alternative variant bSep08, mRNA.
<a href="#">RGD1305038</a>	<a href="#">RGD1305038.cSep08</a>	<a href="#">314627</a>	1413	502		100	similar to Serine/threonine-protein kinase SNK (Serum inducible kinase) (RGD1305038) alternative variant cSep08, mRNA.
<a href="#">RGD1305045</a>	<a href="#">RGD1305045.aSep08</a>	<a href="#">296050</a>	10371	1102	2	241	similar to hypothetical protein (26.4 kD) (RGD1305045) alternative variant aSep08, mRNA.
<a href="#">RGD1305045</a>	<a href="#">RGD1305045.cSep08</a>	<a href="#">296050</a>	8882	352	1	116	similar to hypothetical protein (RGD1305045) alternative variant cSep08, mRNA.
<a href="#">RGD1305090</a>	<a href="#">RGD1305090.aSep08</a>	<a href="#">362598</a>	14600	1829	8	575	similar to CD2-associated protein (RGD1305090) alternative variant aSep08, mRNA.
<a href="#">RGD1305090</a>	<a href="#">RGD1305090.bSep08</a>	<a href="#">362598</a>	800	244	1	81	similar to CD2-associated protein (RGD1305090) alternative variant bSep08, mRNA.
<a href="#">RGD1305094</a>	<a href="#">RGD1305094.aSep08</a>	<a href="#">304649</a>	1990	1353	4	268	similar to CG2662-PA (RGD1305094) alternative variant aSep08, mRNA.
<a href="#">RGD1305094</a>	<a href="#">RGD1305094.bSep08</a>	<a href="#">304649</a>	906	822	1	91	similar to CG2662-PA (RGD1305094) alternative variant bSep08, mRNA.
<a href="#">RGD1305110</a>	<a href="#">RGD1305110.aSep08</a>	<a href="#">305579</a>	37796	1809	10	548	similar to KIAA1841 protein (RGD1305110) alternative variant aSep08, mRNA.
<a href="#">RGD1305138</a>	<a href="#">RGD1305138.bSep08</a>	<a href="#">315329</a>	1151	1067	2	286	similar to expressed sequence AW556797 (RGD1305138) alternative variant bSep08, mRNA.

<a href="#">RGD1305157</a>	<a href="#">RGD1305157.aSep08</a>	<a href="#">316090</a>	30560	2214		564	similar to hypothetical basic protein I-19 (62.5 kD) (RGD1305157) mRNA.
<a href="#">RGD1305158</a>	<a href="#">RGD1305158.aSep08</a>	<a href="#">297971</a>	8702	627	3	97	similar to RIKEN cDNA 1810030N24 (11.0 kD) (RGD1305158) alternative variant aSep08, complete mRNA.
<a href="#">RGD1305158</a>	<a href="#">RGD1305158.bSep08</a>	<a href="#">297971</a>	6529	520	2	95	similar to RIKEN cDNA 1810030N24 (10.8 kD) (RGD1305158) alternative variant bSep08, complete mRNA.
<a href="#">RGD1305158</a>	<a href="#">RGD1305158.cSep08</a>	<a href="#">297971</a>	8680	550	3	93	similar to RIKEN cDNA 1810030N24 (RGD1305158) alternative variant cSep08, mRNA.
<a href="#">RGD1305158</a>	<a href="#">RGD1305158.eSep08</a>	<a href="#">297971</a>	8821	589	2	6	similar to RIKEN cDNA 1810030N24 (0.7 kD) (RGD1305158) alternative variant eSep08, complete mRNA.
<a href="#">RGD1305178</a>	<a href="#">RGD1305178.aSep08</a>	<a href="#">311855</a>	8291	1321	4	289	similar to Hypothetical protein MGC11690 (33.5 kD) (RGD1305178) alternative variant aSep08, complete mRNA.
<a href="#">RGD1305211</a>	<a href="#">RGD1305211.bSep08</a>	<a href="#">365314</a>	1425	763	1	64	similar to RIKEN cDNA 2610209A20 (RGD1305211) alternative variant bSep08, mRNA.
<a href="#">RGD1305215</a>	<a href="#">RGD1305215.bSep08</a>	<a href="#">307643</a>	5371	385	1	128	similar to expressed sequence AA960436 (RGD1305215) alternative variant bSep08, mRNA.
<a href="#">RGD1305222</a>	<a href="#">RGD1305222.aSep08</a>	<a href="#">290686</a>	15944	974		221	similar to RIKEN cDNA 1810029B16 (25.8 kD) (RGD1305222) mRNA.
<a href="#">RGD1305225</a>	<a href="#">RGD1305225.bSep08</a>	<a href="#">297458</a>	15814	513	5	21	similar to RIKEN cDNA C130022K22 gene (2.6 kD) (RGD1305225) alternative variant bSep08, complete mRNA.
<a href="#">RGD1305235</a>	<a href="#">RGD1305235.bSep08</a>	<a href="#">292267</a>	20608	421	3	139	similar to RIKEN cDNA 1700052N19 (RGD1305235) alternative variant bSep08, mRNA.
<a href="#">RGD1305235</a>	<a href="#">RGD1305235.cSep08</a>	<a href="#">292267</a>	7262	880	3	136	similar to RIKEN cDNA 1700052N19 (16.0 kD) (RGD1305235) alternative variant cSep08, mRNA.
<a href="#">RGD1305235</a>	<a href="#">RGD1305235.dSep08</a>	<a href="#">292267</a>	5958	483	5	79	similar to RIKEN cDNA 1700052N19 (RGD1305235) alternative variant dSep08, mRNA.
<a href="#">RGD1305254</a>	<a href="#">RGD1305254.aSep08</a>	<a href="#">308797</a>	2592	350		116	similar to transmembrane protein 2 (RGD1305254) mRNA.
<a href="#">RGD1305256</a>	<a href="#">RGD1305256.aSep08</a>	<a href="#">311443</a>	7312	1185		394	neuronal leucine rich (RGD1305256) mRNA.
<a href="#">RGD1305269</a>	<a href="#">RGD1305269.aSep08</a>	<a href="#">305332</a>	87097	5787	21	914	similar to hypothetical protein (102.9 kD) (RGD1305269) alternative variant aSep08, mRNA.
<a href="#">RGD1305269</a>	<a href="#">RGD1305269.bSep08</a>	<a href="#">305332</a>	254110	1168	8	389	similar to hypothetical protein (RGD1305269) alternative variant bSep08, mRNA.
<a href="#">RGD1305269</a>	<a href="#">RGD1305269.cSep08</a>	<a href="#">305332</a>	13851	806	8	268	similar to hypothetical protein (RGD1305269) alternative variant cSep08, mRNA.
<a href="#">RGD1305269</a>	<a href="#">RGD1305269.dSep08</a>	<a href="#">305332</a>	23130	822	4	237	similar to hypothetical protein (RGD1305269) alternative variant dSep08, mRNA.
<a href="#">RGD1305269</a>	<a href="#">RGD1305269.eSep08</a>	<a href="#">305332</a>	10888	361	4	104	similar to hypothetical protein (RGD1305269) alternative variant eSep08, mRNA.
<a href="#">RGD1305283</a>	<a href="#">RGD1305283.aSep08</a>	<a href="#">363162</a>	58007	680	2	106	similar to RIKEN cDNA 2010110K16 (12.6 kD) (RGD1305283) alternative variant aSep08, mRNA.
<a href="#">RGD1305283</a>	<a href="#">RGD1305283.bSep08</a>	<a href="#">363162</a>	51304	425	1	83	similar to RIKEN cDNA 2010110K16 (RGD1305283) alternative variant bSep08, mRNA.

<a href="#">RGD1305288</a>	<a href="#">RGD1305288.aSep08</a>	<a href="#">305882</a>	7036	1248	7	326	uncharacterized protein (RGD1305288) alternative variant aSep08, mRNA.
<a href="#">RGD1305288</a>	<a href="#">RGD1305288.cSep08</a>	<a href="#">305882</a>	1331	900	3	177	putative nuclear protein, with a coiled coil domain, of metazoan origin (20.9 kD) (RGD1305288) alternative variant cSep08, mRNA.
<a href="#">RGD1305288</a>	<a href="#">RGD1305288.dSep08</a>	<a href="#">305882</a>	1332	798	4	176	uncharacterized protein (RGD1305288) alternative variant dSep08, mRNA.
<a href="#">RGD1305311</a>	<a href="#">RGD1305311.aSep08</a>	<a href="#">316616</a>	41582	1103		367	similar to hypothetical protein FLJ22527 (RGD1305311) mRNA.
<a href="#">RGD1305344</a>	<a href="#">RGD1305344.aSep08</a>	<a href="#">308557</a>	3851	633	5	144	similar to Opioid binding protein/cell adhesion molecule precursor (OBCAM) (Opioid-binding cell adhesion molecule) (OPCML) (RGD1305344) alternative variant aSep08, mRNA.
<a href="#">RGD1305347</a>	<a href="#">RGD1305347.aSep08</a>	<a href="#">362576</a>	2673	697	1	113	similar to RIKEN cDNA 2610528J11 (RGD1305347) alternative variant aSep08, mRNA.
<a href="#">RGD1305350</a>	<a href="#">RGD1305350.aSep08</a>	<a href="#">313699</a>	6108	2594	3	635	similar to RIKEN cDNA 2510039O18 (RGD1305350) alternative variant aSep08, mRNA.
<a href="#">RGD1305351</a>	<a href="#">RGD1305351.bSep08</a>	<a href="#">300663</a>	1682	838	1	67	hypothetical LOC300663 (7.3 kD) (RGD1305351) alternative variant bSep08, mRNA.
<a href="#">RGD1305362</a>	<a href="#">RGD1305362.aSep08</a>	<a href="#">303155</a>	13647	977	2	298	similar to RIKEN cDNA 9030624B09 (RGD1305362) alternative variant aSep08, mRNA.
<a href="#">RGD1305362</a>	<a href="#">RGD1305362.bSep08</a>	<a href="#">303155</a>	12165	842	2	253	similar to RIKEN cDNA 9030624B09 (RGD1305362) alternative variant bSep08, mRNA.
<a href="#">RGD1305362</a>	<a href="#">RGD1305362.cSep08</a>	<a href="#">303155</a>	12141	812	2	247	similar to RIKEN cDNA 9030624B09 (RGD1305362) alternative variant cSep08, mRNA.
<a href="#">RGD1305362</a>	<a href="#">RGD1305362.dSep08</a>	<a href="#">303155</a>	12118	792	2	237	similar to RIKEN cDNA 9030624B09 (RGD1305362) alternative variant dSep08, mRNA.
<a href="#">RGD1305415</a>	<a href="#">RGD1305415.aSep08</a>	<a href="#">362958</a>	15444	527		175	similar to CG11259-PA (RGD1305415) mRNA.
<a href="#">RGD1305420</a>	<a href="#">RGD1305420.bSep08</a>	<a href="#">298072</a>	2019	523		139	similar to Nef associated protein 1 (RGD1305420) alternative variant bSep08, mRNA.
<a href="#">RGD1305422</a>	<a href="#">RGD1305422.aSep08</a>	<a href="#">303885</a>	7589	3387	7	388	similar to mKIAA0226 protein (RGD1305422) alternative variant aSep08, mRNA.
<a href="#">RGD1305440</a>	<a href="#">RGD1305440.bSep08</a>	<a href="#">296161</a>	8588	462	2	101	similar to Rnf37-pending protein (RGD1305440) alternative variant bSep08, mRNA.
<a href="#">RGD1305440</a>	<a href="#">RGD1305440.cSep08</a>	<a href="#">296161</a>	41501	1128	5	96	similar to Rnf37-pending protein (10.7 kD) (RGD1305440) alternative variant cSep08, mRNA.
<a href="#">RGD1305440</a>	<a href="#">RGD1305440.dSep08</a>	<a href="#">296161</a>	31527	743	3	64	similar to Rnf37-pending protein (RGD1305440) alternative variant dSep08, mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.aSep08</a>	<a href="#">365407</a>	23951	3663	11	471	pre-mRNA cleavage factor I subunit (52.0 kD) (RGD1305441) alternative variant aSep08, complete mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.cSep08</a>	<a href="#">365407</a>	4763	1363	5	213	pre-mRNA cleavage factor I subunit (22.8 kD) (RGD1305441) alternative variant cSep08, mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.dSep08</a>	<a href="#">365407</a>	4165	738	5	202	pre-mRNA cleavage factor I subunit (RGD1305441) alternative variant dSep08, mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.eSep08</a>	<a href="#">365407</a>	3015	363	3	120	pre-mRNA cleavage factor I subunit (RGD1305441) alternative variant eSep08, mRNA.

<a href="#">RGD1305441</a>	<a href="#">RGD1305441.fSep08</a>	<a href="#">365407</a>	6987	578	3	99	pre-mRNA cleavage factor I subunit (11.0 kD) (RGD1305441) alternative variant fSep08, mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.gSep08</a>	<a href="#">365407</a>	7993	1888	2	69	putative protein (7.4 kD) (RGD1305441) alternative variant gSep08, mRNA.
<a href="#">RGD1305441</a>	<a href="#">RGD1305441.hSep08</a>	<a href="#">365407</a>	6735	768	3	64	putative protein (RGD1305441) alternative variant hSep08, mRNA.
<a href="#">RGD1305457</a>	<a href="#">RGD1305457.bSep08</a>	<a href="#">314730</a>	19443	1334	3	345	similar to RIKEN cDNA 1700023M03 (38.3 kD) (RGD1305457) alternative variant bSep08, mRNA.
<a href="#">RGD1305457</a>	<a href="#">RGD1305457.cSep08</a>	<a href="#">314730</a>	17562	896	4	105	similar to RIKEN cDNA 1700023M03 (11.6 kD) (RGD1305457) alternative variant cSep08, mRNA.
<a href="#">RGD1305481</a>	<a href="#">RGD1305481.aSep08</a>	<a href="#">294030</a>	4024	1363	4	280	hypothetical LOC294030 (RGD1305481) alternative variant aSep08, mRNA.
<a href="#">RGD1305481</a>	<a href="#">RGD1305481.cSep08</a>	<a href="#">294030</a>	3402	870	4	243	hypothetical LOC294030 (27.6 kD) (RGD1305481) alternative variant cSep08, mRNA.
<a href="#">RGD1305481</a>	<a href="#">RGD1305481.dSep08</a>	<a href="#">294030</a>	1748	604	2	200	hypothetical LOC294030 (RGD1305481) alternative variant dSep08, mRNA.
<a href="#">RGD1305481</a>	<a href="#">RGD1305481.eSep08</a>	<a href="#">294030</a>	1595	706	2	62	hypothetical LOC294030 (6.5 kD) (RGD1305481) alternative variant eSep08, complete mRNA.
<a href="#">RGD1305500</a>	<a href="#">RGD1305500.bSep08</a>	<a href="#">308004</a>	6008	476	3	108	similar to hypothetical protein FLJ13188 (RGD1305500) alternative variant bSep08, mRNA.
<a href="#">RGD1305500</a>	<a href="#">RGD1305500.cSep08</a>	<a href="#">308004</a>	9544	337	3	52	similar to hypothetical protein FLJ13188 (RGD1305500) alternative variant cSep08, mRNA.
<a href="#">RGD1305508</a>	<a href="#">RGD1305508.aSep08</a>	<a href="#">303275</a>	3668	1564		194	similar to hypothetical protein MGC23280 (RGD1305508) mRNA.
<a href="#">RGD1305526</a>	<a href="#">RGD1305526.bSep08</a>	<a href="#">294606</a>	168259	753	7	172	CRA a (RGD1305526) alternative variant bSep08, mRNA.
<a href="#">RGD1305526</a>	<a href="#">RGD1305526.cSep08</a>	<a href="#">294606</a>	33036	277	3	71	putative protein (RGD1305526) alternative variant cSep08, mRNA.
<a href="#">RGD1305534</a>	<a href="#">RGD1305534.aSep08</a>	<a href="#">315864</a>	14104	1872		356	similar to dJ202D23.2 (novel protein similar to C21ORF5 (KIAA0933)) (RGD1305534) mRNA.
<a href="#">RGD1305539</a>	<a href="#">RGD1305539.aSep08</a>	<a href="#">303920</a>	17315	936		311	similar to hypothetical protein (RGD1305539) mRNA.
<a href="#">RGD1305560</a>	<a href="#">RGD1305560.aSep08</a>	<a href="#">301561</a>	44747	372		123	similar to RIKEN cDNA 9430031J16 (RGD1305560) mRNA.
<a href="#">RGD1305572</a>	<a href="#">RGD1305572.bSep08</a>	<a href="#">289315</a>	7766	666	5	222	similar to hypothetical protein MGC30618 (RGD1305572) alternative variant bSep08, mRNA.
<a href="#">RGD1305572</a>	<a href="#">RGD1305572.cSep08</a>	<a href="#">289315</a>	3261	963	3	220	similar to hypothetical protein MGC30618 (RGD1305572) alternative variant cSep08, mRNA.
<a href="#">RGD1305587</a>	<a href="#">RGD1305587.aSep08</a>	<a href="#">294499</a>	2402	715	1	144	similar to RIKEN cDNA 2010107G23 (RGD1305587) alternative variant aSep08, mRNA.
<a href="#">RGD1305592</a>	<a href="#">RGD1305592.aSep08</a>	<a href="#">293500</a>	1832	945	2	227	similar to RIKEN cDNA 2900092E17 (RGD1305592) alternative variant aSep08, mRNA.
<a href="#">RGD1305604and dRGD1564921</a>	<a href="#">RGD1305604andRGD1 564921.aSep08</a>	<a href="#">362580</a>	16115	927	6	199	similar to CG2919-PA and similar to peptidyl prolyl isomerase H (RGD1305604andRGD1564921) alternative variant aSep08, mRNA.
<a href="#">RGD1305604and dRGD1564921</a>	<a href="#">RGD1305604andRGD1 564921.aSep08</a>	<a href="#">366461</a>	16115	927	6	199	similar to CG2919-PA and similar to peptidyl prolyl isomerase H (RGD1305604andRGD1564921) alternative variant aSep08, mRNA.

<a href="#">RGD1305604andRGD1564921</a>	<a href="#">RGD1305604andRGD1564921.bSep08</a>	<a href="#">362580</a>	919	563	2	38	similar to CG2919-PA and similar to peptidyl prolyl isomerase H (4.3 kD) (RGD1305604andRGD1564921) alternative variant bSep08, mRNA.
<a href="#">RGD1305604andRGD1564921</a>	<a href="#">RGD1305604andRGD1564921.bSep08</a>	<a href="#">366461</a>	919	563	2	38	similar to CG2919-PA and similar to peptidyl prolyl isomerase H (4.3 kD) (RGD1305604andRGD1564921) alternative variant bSep08, mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.aSep08</a>	<a href="#">361648</a>	4584	1991	9	342	spinster (36.7 kD) (RGD1305613) alternative variant aSep08, mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.cSep08</a>	<a href="#">361648</a>	7277	2926	10	270	spinster (29.1 kD) (RGD1305613) alternative variant cSep08, complete mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.dSep08</a>	<a href="#">361648</a>	4370	1476	9	229	spinster (24.2 kD) (RGD1305613) alternative variant dSep08, mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.eSep08</a>	<a href="#">361648</a>	2215	674	3	189	spinster (RGD1305613) alternative variant eSep08, mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.fSep08</a>	<a href="#">361648</a>	1933	653	4	133	spinster (RGD1305613) alternative variant fSep08, mRNA.
<a href="#">RGD1305613</a>	<a href="#">RGD1305613.gSep08</a>	<a href="#">361648</a>	1384	628	2	57	spinster (6.6 kD) (RGD1305613) alternative variant gSep08, mRNA.
<a href="#">RGD1305628</a>	<a href="#">RGD1305628.aSep08</a>	<a href="#">288204</a>	44496	3100	10	425	similar to Hypothetical protein 5031404N19 (RGD1305628) alternative variant aSep08, mRNA.
<a href="#">RGD1305628</a>	<a href="#">RGD1305628.bSep08</a>	<a href="#">288204</a>	1746	708	2	128	similar to Hypothetical protein 5031404N19 (RGD1305628) alternative variant bSep08, mRNA.
<a href="#">RGD1305645</a>	<a href="#">RGD1305645.aSep08</a>	<a href="#">363225</a>	19887	691	1	173	similar to RIKEN cDNA 1500015O10 (RGD1305645) alternative variant aSep08, mRNA.
<a href="#">RGD1305645</a>	<a href="#">RGD1305645.bSep08</a>	<a href="#">363225</a>	19560	814	1	148	similar to RIKEN cDNA 1500015O10 (16.9 kD) (RGD1305645) alternative variant bSep08, mRNA.
<a href="#">RGD1305645</a>	<a href="#">RGD1305645.cSep08</a>	<a href="#">363225</a>	55342	612	1	126	similar to RIKEN cDNA 1500015O10 (14.9 kD) (RGD1305645) alternative variant cSep08, mRNA.
<a href="#">RGD1305685</a>	<a href="#">RGD1305685.bSep08</a>	<a href="#">360811</a>	3542	395	3	107	similar to hypothetical protein FLJ13089 (RGD1305685) alternative variant bSep08, mRNA.
<a href="#">RGD1305685</a>	<a href="#">RGD1305685.cSep08</a>	<a href="#">360811</a>	3054	1002	2	83	similar to hypothetical protein FLJ13089 (RGD1305685) alternative variant cSep08, mRNA.
<a href="#">RGD1305685</a>	<a href="#">RGD1305685.eSep08</a>	<a href="#">360811</a>	1666	422	2	21	similar to hypothetical protein FLJ13089 (RGD1305685) alternative variant eSep08, mRNA.
<a href="#">RGD1305689</a>	<a href="#">RGD1305689.aSep08</a>	<a href="#">290529</a>	3053	545	1	142	similar to DNA segment, Chr 14, ERATO Doi 449, expressed (RGD1305689) alternative variant aSep08, mRNA.
<a href="#">RGD1305697</a>	<a href="#">RGD1305697.bSep08</a>	<a href="#">296846</a>	11952	741	3	238	similar to hypothetical protein FLJ25530 (RGD1305697) alternative variant bSep08, mRNA.
<a href="#">RGD1305713</a>	<a href="#">RGD1305713.bSep08</a>	<a href="#">293059</a>	6996	495	4	150	similar to RIKEN cDNA 3110040N11 (RGD1305713) alternative variant bSep08, mRNA.
<a href="#">RGD1305713</a>	<a href="#">RGD1305713.cSep08</a>	<a href="#">293059</a>	4613	607	4	150	similar to RIKEN cDNA 3110040N11 (RGD1305713) alternative variant cSep08, mRNA.
<a href="#">RGD1305725andRGD1561297</a>	<a href="#">RGD1305725andRGD1561297.aSep08</a>	<a href="#">362255</a>	27085	1103	1	165	similar to Rpl7a protein (RGD1305725andRGD1561297) alternative variant aSep08, mRNA.
<a href="#">RGD1305725andRGD1561297</a>	<a href="#">RGD1305725andRGD1561297.aSep08</a>	<a href="#">366239</a>	27085	1103	1	165	similar to Rpl7a protein (RGD1305725andRGD1561297) alternative variant aSep08, mRNA.
<a href="#">RGD1305725andRGD1561297</a>	<a href="#">RGD1305725andRGD1561297.bSep08</a>	<a href="#">362255</a>	7777	474	1	108	similar to Rpl7a protein (RGD1305725andRGD1561297) alternative variant bSep08, mRNA.

<a href="#">RGD1305725andRGD1561297</a>	<a href="#">RGD1305725andRGD1561297.bSep08</a>	<a href="#">366239</a>	7777	474	1	108	similar to Rpl7a protein (RGD1305725andRGD1561297) alternative variant bSep08, mRNA.
<a href="#">RGD1305727</a>	<a href="#">RGD1305727.bSep08</a>	<a href="#">306768</a>	1389	1304	1	96	similar to Protein CGI-117 (Protein HSPC111) (11.2 kD) (RGD1305727) alternative variant bSep08, complete mRNA.
<a href="#">RGD1305733</a>	<a href="#">RGD1305733.bSep08</a>	<a href="#">360459</a>	6914	555	1	167	similar to RIKEN cDNA 2900011O08 (RGD1305733) alternative variant bSep08, mRNA.
<a href="#">RGD1305793</a>	<a href="#">RGD1305793.bSep08</a>	<a href="#">309456</a>	44145	705	2	114	similar to hypothetical protein FLJ20154 (RGD1305793) alternative variant bSep08, mRNA.
<a href="#">RGD1305793</a>	<a href="#">RGD1305793.cSep08</a>	<a href="#">309456</a>	1176	757	2	100	similar to hypothetical protein FLJ20154 (RGD1305793) alternative variant cSep08, mRNA.
<a href="#">RGD1305793</a>	<a href="#">RGD1305793.eSep08</a>	<a href="#">309456</a>	30356	497	4	110	similar to hypothetical protein FLJ20154 (RGD1305793) alternative variant eSep08, mRNA.
<a href="#">RGD1305797</a>	<a href="#">RGD1305797.bSep08</a>	<a href="#">298186</a>	97696	1780	2	516	similar to hypothetical protein FLJ33868 (RGD1305797) alternative variant bSep08, mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.bSep08</a>	<a href="#">360461</a>	17483	871	3	170	similar to RIKEN cDNA 0610037P05 (RGD1305823) alternative variant bSep08, mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.cSep08</a>	<a href="#">360461</a>	22706	1405	4	140	similar to RIKEN cDNA 0610037P05 (16.1 kD) (RGD1305823) alternative variant cSep08, complete mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.dSep08</a>	<a href="#">360461</a>	22017	631	3	132	similar to RIKEN cDNA 0610037P05 (RGD1305823) alternative variant dSep08, mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.eSep08</a>	<a href="#">360461</a>	22163	838	4	132	similar to RIKEN cDNA 0610037P05 (15.2 kD) (RGD1305823) alternative variant eSep08, mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.fSep08</a>	<a href="#">360461</a>	22289	877	3	104	similar to RIKEN cDNA 0610037P05 (12.1 kD) (RGD1305823) alternative variant fSep08, mRNA.
<a href="#">RGD1305823</a>	<a href="#">RGD1305823.gSep08</a>	<a href="#">360461</a>	18334	400	1	54	similar to RIKEN cDNA 0610037P05 (RGD1305823) alternative variant gSep08, mRNA.
<a href="#">RGD1305844</a>	<a href="#">RGD1305844.aSep08</a>	<a href="#">294883</a>	284690	803	5	148	hypothetical LOC294883 (16.6 kD) (RGD1305844) alternative variant aSep08, mRNA.
<a href="#">RGD1305844</a>	<a href="#">RGD1305844.bSep08</a>	<a href="#">294883</a>	102393	1348	4	145	hypothetical LOC294883 (RGD1305844) alternative variant bSep08, mRNA.
<a href="#">RGD1305844</a>	<a href="#">RGD1305844.cSep08</a>	<a href="#">294883</a>	57153	319	1	96	hypothetical LOC294883 (RGD1305844) alternative variant cSep08, mRNA.
<a href="#">RGD1305898</a>	<a href="#">RGD1305898.aSep08</a>	<a href="#">361275</a>	78521	1289	8	414	similar to hypothetical protein FLJ40283 (RGD1305898) alternative variant aSep08, mRNA.
<a href="#">RGD1305898</a>	<a href="#">RGD1305898.bSep08</a>	<a href="#">361275</a>	10960	700	2	136	similar to hypothetical protein FLJ40283 (RGD1305898) alternative variant bSep08, mRNA.
<a href="#">RGD1305898</a>	<a href="#">RGD1305898.dSep08</a>	<a href="#">361275</a>	23265	436	3	53	similar to hypothetical protein FLJ40283 (RGD1305898) alternative variant dSep08, mRNA.
<a href="#">RGD1305899</a>	<a href="#">RGD1305899.aSep08</a>	<a href="#">311715</a>	18637	1058		352	similar to Protein C20orf158 (RGD1305899) mRNA.
<a href="#">RGD1305938</a>	<a href="#">RGD1305938.aSep08</a>	<a href="#">310362</a>	10582	1385		292	similar to expressed sequence AW549877 (RGD1305938) mRNA.
<a href="#">RGD1305939</a>	<a href="#">RGD1305939.aSep08</a>	<a href="#">300074</a>	12431	582	6	103	hypothetical LOC300074 (RGD1305939) alternative variant aSep08, mRNA.
<a href="#">RGD1306001</a>	<a href="#">RGD1306001.cSep08</a>	<a href="#">362975</a>	2251	615	2	61	similar to 2210021J22Rik protein (RGD1306001) alternative variant cSep08, mRNA.

<a href="#">RGD1306041</a>	<a href="#">RGD1306041.bSep08</a>	<a href="#">300125</a>	10948	719	7	156	similar to FLJ20699 protein (RGD1306041) alternative variant bSep08, mRNA.
<a href="#">RGD1306041</a>	<a href="#">RGD1306041.cSep08</a>	<a href="#">300125</a>	3173	727	4	93	similar to FLJ20699 protein (RGD1306041) alternative variant cSep08, mRNA.
<a href="#">RGD1306058</a>	<a href="#">RGD1306058.bSep08</a>	<a href="#">361224</a>	11724	768	6	198	similar to RIKEN cDNA 1110007C09 (RGD1306058) alternative variant bSep08, mRNA.
<a href="#">RGD1306058</a>	<a href="#">RGD1306058.cSep08</a>	<a href="#">361224</a>	4617	552	4	125	similar to RIKEN cDNA 1110007C09 (RGD1306058) alternative variant cSep08, mRNA.
<a href="#">RGD1306062</a>	<a href="#">RGD1306062.aSep08</a>	<a href="#">294595</a>	57964	1792	10	597	similar to KIAA0372 gene product (RGD1306062) alternative variant aSep08, mRNA.
<a href="#">RGD1306062</a>	<a href="#">RGD1306062.bSep08</a>	<a href="#">294595</a>	45840	1048	3	300	similar to KIAA0372 gene product (RGD1306062) alternative variant bSep08, mRNA.
<a href="#">RGD1306062</a>	<a href="#">RGD1306062.cSep08</a>	<a href="#">294595</a>	3780	501	2	66	similar to KIAA0372 gene product (RGD1306062) alternative variant cSep08, mRNA.
<a href="#">RGD1306062</a>	<a href="#">RGD1306062.dSep08</a>	<a href="#">294595</a>	3017	479	1	66	similar to KIAA0372 gene product (RGD1306062) alternative variant dSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.aSep08</a>	<a href="#">289928</a>	16880	943	7	135	similar to HT021 (15.9 kD) (RGD1306063) alternative variant aSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.dSep08</a>	<a href="#">289928</a>	15480	580	5	117	similar to HT021 (13.8 kD) (RGD1306063) alternative variant dSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.eSep08</a>	<a href="#">289928</a>	2326	543	2	89	similar to HT021 (10.2 kD) (RGD1306063) alternative variant eSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.fSep08</a>	<a href="#">289928</a>	16667	581	6	87	similar to HT021 (10.4 kD) (RGD1306063) alternative variant fSep08, complete mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.gSep08</a>	<a href="#">289928</a>	7539	570	3	71	similar to HT021 (8.2 kD) (RGD1306063) alternative variant gSep08, complete mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.hSep08</a>	<a href="#">289928</a>	8724	642	5	69	similar to HT021 (8.2 kD) (RGD1306063) alternative variant hSep08, complete mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.iSep08</a>	<a href="#">289928</a>	8396	311	5	69	similar to HT021 (8.2 kD) (RGD1306063) alternative variant iSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.jSep08</a>	<a href="#">289928</a>	3921	285	2	69	similar to HT021 (RGD1306063) alternative variant jSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.kSep08</a>	<a href="#">289928</a>	8806	800	5	69	similar to HT021 (8.2 kD) (RGD1306063) alternative variant kSep08, mRNA.
<a href="#">RGD1306063</a>	<a href="#">RGD1306063.lSep08</a>	<a href="#">289928</a>	8404	256	4	63	similar to HT021 (RGD1306063) alternative variant lSep08, mRNA.
<a href="#">RGD1306072</a>	<a href="#">RGD1306072.aSep08</a>	<a href="#">304654</a>	20924	2156	5	622	hypothetical LOC304654 (RGD1306072) alternative variant aSep08, mRNA.
<a href="#">RGD1306072</a>	<a href="#">RGD1306072.bSep08</a>	<a href="#">304654</a>	4520	843	2	170	hypothetical LOC304654 (RGD1306072) alternative variant bSep08, mRNA.
<a href="#">RGD1306072</a>	<a href="#">RGD1306072.cSep08</a>	<a href="#">304654</a>	14200	720	2	119	hypothetical LOC304654 (RGD1306072) alternative variant cSep08, mRNA.
<a href="#">RGD1306091</a>	<a href="#">RGD1306091.aSep08</a>	<a href="#">307950</a>	16007	421		139	similar to Mixed lineage kinase 4 (RGD1306091) mRNA.
<a href="#">RGD1306105</a>	<a href="#">RGD1306105.bSep08</a>	<a href="#">360916</a>	82241	424	6	140	similar to RIKEN cDNA 2610318G18 (RGD1306105) alternative variant bSep08, mRNA.
<a href="#">RGD1306105</a>	<a href="#">RGD1306105.cSep08</a>	<a href="#">360916</a>	82255	541	5	90	similar to RIKEN cDNA 2610318G18 (RGD1306105) alternative variant cSep08, mRNA.

<a href="#">RGD1306107</a>	<a href="#">RGD1306107.bSep08</a>	<a href="#">310640</a>	1923	499	5	165	putative protein of vertebrate origin (RGD1306107) alternative variant bSep08, mRNA.
<a href="#">RGD1306107</a>	<a href="#">RGD1306107.cSep08</a>	<a href="#">310640</a>	992	901	2	124	putative protein of mammalian origin (13.4 kD) (RGD1306107) alternative variant cSep08, mRNA.
<a href="#">RGD1306107</a>	<a href="#">RGD1306107.dSep08</a>	<a href="#">310640</a>	1312	759	3	110	putative protein of mammalian origin (RGD1306107) alternative variant dSep08, mRNA.
<a href="#">RGD1306107</a>	<a href="#">RGD1306107.eSep08</a>	<a href="#">310640</a>	1543	544	4	107	putative protein of vertebrate origin (RGD1306107) alternative variant eSep08, mRNA.
<a href="#">RGD1306116</a>	<a href="#">RGD1306116.aSep08</a>	<a href="#">361774</a>	16524	5042	14	679	LOC361774 (RGD1306116) alternative variant aSep08, mRNA.
<a href="#">RGD1306116</a>	<a href="#">RGD1306116.bSep08</a>	<a href="#">361774</a>	2733	835	4	154	LOC361774 (RGD1306116) alternative variant bSep08, mRNA.
<a href="#">RGD1306116</a>	<a href="#">RGD1306116.cSep08</a>	<a href="#">361774</a>	644	333	2	70	LOC361774 (RGD1306116) alternative variant cSep08, mRNA.
<a href="#">RGD1306119</a>	<a href="#">RGD1306119.aSep08</a>	<a href="#">314306</a>	10834	569	3	189	similar to transcriptional regulating protein 132 (RGD1306119) alternative variant aSep08, mRNA.
<a href="#">RGD1306148</a>	<a href="#">RGD1306148.aSep08</a>	<a href="#">313196</a>	31949	4505	25	883	similar to KIAA0368 (RGD1306148) alternative variant aSep08, mRNA.
<a href="#">RGD1306151</a>	<a href="#">RGD1306151.bSep08</a>	<a href="#">362455</a>	21441	1310	4	244	similar to hypothetical protein DKFZp761D0211 (RGD1306151) alternative variant bSep08, mRNA.
<a href="#">RGD1306153</a>	<a href="#">RGD1306153.aSep08</a>	<a href="#">361410</a>	18774	720		239	similar to predicted CDS, putative protein of bilateral origin (4J193) (RGD1306153) mRNA.
<a href="#">RGD1306157</a>	<a href="#">RGD1306157.aSep08</a>	<a href="#">291947</a>	34500	2742	10	841	similar to CG9882-PA (93.9 kD) (RGD1306157) alternative variant aSep08, mRNA.
<a href="#">RGD1306181</a>	<a href="#">RGD1306181.bSep08</a>	<a href="#">308346</a>	4347	708	2	91	similar to A630041N19 protein (RGD1306181) alternative variant bSep08, mRNA.
<a href="#">RGD1306192</a>	<a href="#">RGD1306192.aSep08</a>	<a href="#">361443</a>	13679	1775	2	539	similar to RIKEN cDNA 2310079N02 (RGD1306192) alternative variant aSep08, mRNA.
<a href="#">RGD1306215</a>	<a href="#">RGD1306215.aSep08</a>	<a href="#">296565</a>	1637	998	7	205	similar to hypothetical protein MGC36831 (22.1 kD) (RGD1306215) alternative variant aSep08, mRNA.
<a href="#">RGD1306215</a>	<a href="#">RGD1306215.bSep08</a>	<a href="#">296565</a>	1489	1155	4	127	similar to hypothetical protein MGC36831 (13.7 kD) (RGD1306215) alternative variant bSep08, mRNA.
<a href="#">RGD1306215</a>	<a href="#">RGD1306215.cSep08</a>	<a href="#">296565</a>	1482	1003	6	108	similar to hypothetical protein MGC36831 (11.4 kD) (RGD1306215) alternative variant cSep08, mRNA.
<a href="#">RGD1306215</a>	<a href="#">RGD1306215.eSep08</a>	<a href="#">296565</a>	780	455	4	85	similar to hypothetical protein MGC36831 (RGD1306215) alternative variant eSep08, mRNA.
<a href="#">RGD1306215</a>	<a href="#">RGD1306215.fSep08</a>	<a href="#">296565</a>	953	637	4	62	similar to hypothetical protein MGC36831 (6.5 kD) (RGD1306215) alternative variant fSep08, mRNA.
<a href="#">RGD1306228</a>	<a href="#">RGD1306228.aSep08</a>	<a href="#">362748</a>	3510	495	3	46	CRA b like (RGD1306228) alternative variant aSep08, mRNA.
<a href="#">RGD1306271</a>	<a href="#">RGD1306271.aSep08</a>	<a href="#">312246</a>	6841	577		192	similar to KIAA1549 protein (RGD1306271) mRNA.
<a href="#">RGD1306284</a>	<a href="#">RGD1306284.bSep08</a>	<a href="#">287918</a>	4679	991	3	147	similar to RIKEN cDNA 1110005A03 (16.1 kD) (RGD1306284) alternative variant bSep08, mRNA.
<a href="#">RGD1306284</a>	<a href="#">RGD1306284.cSep08</a>	<a href="#">287918</a>	889	806	1	114	similar to RIKEN cDNA 1110005A03 (13.2 kD) (RGD1306284) alternative variant cSep08, mRNA.



<a href="#">RGD1306284</a>	<a href="#">RGD1306284.dSep08</a>	<a href="#">287918</a>	4568	569	3	68	similar to RIKEN cDNA 1110005A03 (7.8 kD) (RGD1306284) alternative variant dSep08, complete mRNA.
<a href="#">RGD1306286</a>	<a href="#">RGD1306286.aSep08</a>	<a href="#">364902</a>	7221	692	4	230	similar to mKIAA1632 protein (RGD1306286) alternative variant aSep08, mRNA.
<a href="#">RGD1306286</a>	<a href="#">RGD1306286.bSep08</a>	<a href="#">364902</a>	6869	705	5	177	similar to mKIAA1632 protein (RGD1306286) alternative variant bSep08, mRNA.
<a href="#">RGD1306286</a>	<a href="#">RGD1306286.cSep08</a>	<a href="#">364902</a>	5706	2507	2	91	similar to mKIAA1632 protein (10.5 kD) (RGD1306286) alternative variant cSep08, mRNA.
<a href="#">RGD1306344</a>	<a href="#">RGD1306344.bSep08</a>	<a href="#">296384</a>	6139	372	3	62	similar to Ab1-133 (RGD1306344) alternative variant bSep08, mRNA.
<a href="#">RGD1306349</a>	<a href="#">RGD1306349.aSep08</a>	<a href="#">296512</a>	12298	403		134	similar to formin (RGD1306349) mRNA.
<a href="#">RGD1306353</a>	<a href="#">RGD1306353.aSep08</a>	<a href="#">305911</a>	19137	3733	7	628	putative nuclear protein, with a coiled coil domain, of eukaryotic origin (70.3 kD) (RGD1306353) alternative variant aSep08, mRNA.
<a href="#">RGD1306353</a>	<a href="#">RGD1306353.bSep08</a>	<a href="#">305911</a>	5718	1587	2	283	finger MYM-type 5 (32.4 kD) (RGD1306353) alternative variant bSep08, mRNA.
<a href="#">RGD1306353</a>	<a href="#">RGD1306353.cSep08</a>	<a href="#">305911</a>	2533	534	2	155	zinc finger (RGD1306353) alternative variant cSep08, mRNA.
<a href="#">RGD1306371</a>	<a href="#">RGD1306371.bSep08</a>	<a href="#">314061</a>	36022	1792	2	502	similar to putative nuclear protein, with 2 coiled coil-4 domains, of eukaryotic origin (5I282) (RGD1306371) alternative variant bSep08, mRNA.
<a href="#">RGD1306371</a>	<a href="#">RGD1306371.cSep08</a>	<a href="#">314061</a>	4163	1144	1	125	similar to putative nuclear protein, with 2 coiled coil-4 domains, of eukaryotic origin (5I282) (RGD1306371) alternative variant cSep08, mRNA.
<a href="#">RGD1306404</a>	<a href="#">RGD1306404.bSep08</a>	<a href="#">296733</a>	1829	1205	2	230	similar to mKIAA1402 protein (RGD1306404) alternative variant bSep08, mRNA.
<a href="#">RGD1306404</a>	<a href="#">RGD1306404.cSep08</a>	<a href="#">296733</a>	11428	572	2	160	similar to mKIAA1402 protein (RGD1306404) alternative variant cSep08, mRNA.
<a href="#">RGD1306410</a>	<a href="#">RGD1306410.aSep08</a>	<a href="#">360768</a>	22803	1451	13	391	uncharacterized protein c7orf28 homolog (RGD1306410) alternative variant aSep08, mRNA.
<a href="#">RGD1306410</a>	<a href="#">RGD1306410.cSep08</a>	<a href="#">360768</a>	20315	1824	9	202	uncharacterized protein c7orf28 homolog (22.8 kD) (RGD1306410) alternative variant cSep08, mRNA.
<a href="#">RGD1306410</a>	<a href="#">RGD1306410.dSep08</a>	<a href="#">360768</a>	12477	837	7	185	uncharacterized protein c7orf28 homolog (RGD1306410) alternative variant dSep08, mRNA.
<a href="#">RGD1306410</a>	<a href="#">RGD1306410.eSep08</a>	<a href="#">360768</a>	2118	527	2	52	uncharacterized protein c7orf28 homolog like (RGD1306410) alternative variant eSep08, mRNA.
<a href="#">RGD1306410</a>	<a href="#">RGD1306410.fSep08</a>	<a href="#">360768</a>	2046	622	2	49	uncharacterized protein c7orf28 homolog (5.8 kD) (RGD1306410) alternative variant fSep08, mRNA.
<a href="#">RGD1306441</a>	<a href="#">RGD1306441.bSep08</a>	<a href="#">290425</a>	3804	758	2	72	similar to RIKEN cDNA 4921530L21 (RGD1306441) alternative variant bSep08, mRNA.
<a href="#">RGD1306462</a>	<a href="#">RGD1306462.aSep08</a>	<a href="#">315214</a>	9969	856	4	285	similar to RIKEN cDNA 1700019P01 (RGD1306462) alternative variant aSep08, mRNA.
<a href="#">RGD1306474</a>	<a href="#">RGD1306474.bSep08</a>	<a href="#">362881</a>	1099	424	1	141	similar to RIKEN cDNA 9530003J23 (RGD1306474) alternative variant bSep08, mRNA.
<a href="#">RGD1306487</a>	<a href="#">RGD1306487.aSep08</a>	<a href="#">304759</a>	20116	3036	12	592	similar to RAB3 GTPase-activating protein (RGD1306487) alternative variant aSep08, mRNA.

<a href="#">RGD1306487</a>	<a href="#">RGD1306487.bSep08</a>	<a href="#">304759</a>	3483	629	5	117	similar to RAB3 GTPase-activating protein (13.2 kD) (RGD1306487) alternative variant bSep08, mRNA.
<a href="#">RGD1306507</a>	<a href="#">RGD1306507.aSep08</a>	<a href="#">313063</a>	39581	381		67	similar to RAD54B homolog isoform 1; RAD54, S. cerevisiae, homolog of, B (RGD1306507) mRNA.
<a href="#">RGD1306508</a>	<a href="#">RGD1306508.bSep08</a>	<a href="#">289874</a>	22797	1636	7	148	putative protein of eukaryotic origin (RGD1306508) alternative variant bSep08, mRNA.
<a href="#">RGD1306508</a>	<a href="#">RGD1306508.cSep08</a>	<a href="#">289874</a>	3270	528	2	35	putative protein of metazoan origin (3.8 kD) (RGD1306508) alternative variant cSep08, mRNA.
<a href="#">RGD1306520</a>	<a href="#">RGD1306520.bSep08</a>	<a href="#">310775</a>	96542	1606	3	279	similar to receptor-interacting factor 1 (RGD1306520) alternative variant bSep08, mRNA.
<a href="#">RGD1306520</a>	<a href="#">RGD1306520.cSep08</a>	<a href="#">310775</a>	13378	822	2	96	similar to receptor-interacting factor 1 (11.1 kD) (RGD1306520) alternative variant cSep08, mRNA.
<a href="#">RGD1306534</a>	<a href="#">RGD1306534.aSep08</a>	<a href="#">311647</a>	96076	722		240	similar to P-Rex1 (RGD1306534) mRNA.
<a href="#">RGD1306556</a>	<a href="#">RGD1306556.aSep08</a>	<a href="#">288744</a>	46953	698		174	similar to hypothetical protein A530094D01 (RGD1306556) mRNA.
<a href="#">RGD1306582</a>	<a href="#">RGD1306582.aSep08</a>	<a href="#">362103</a>	8156	754	4	251	similar to RIKEN cDNA 2610205E22 (RGD1306582) alternative variant aSep08, mRNA.
<a href="#">RGD1306582</a>	<a href="#">RGD1306582.cSep08</a>	<a href="#">362103</a>	1159	460	3	77	similar to RIKEN cDNA 2610205E22 (RGD1306582) alternative variant cSep08, mRNA.
<a href="#">RGD1306582</a>	<a href="#">RGD1306582.eSep08</a>	<a href="#">362103</a>	7539	391	2	29	similar to RIKEN cDNA 2610205E22 (RGD1306582) alternative variant eSep08, mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.bSep08</a>	<a href="#">294709</a>	21120	1127	9	327	protein C5orf44 homolog (RGD1306583) alternative variant bSep08, mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.cSep08</a>	<a href="#">294709</a>	31386	2343	11	299	protein CRA d (33.5 kD) (RGD1306583) alternative variant cSep08, complete mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.dSep08</a>	<a href="#">294709</a>	20848	787	6	174	protein CRA d (RGD1306583) alternative variant dSep08, mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.eSep08</a>	<a href="#">294709</a>	15827	745	5	143	protein CRA d (15.9 kD) (RGD1306583) alternative variant eSep08, mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.fSep08</a>	<a href="#">294709</a>	812	681	2	81	protein CRA e precursor (8.8 kD) (RGD1306583) alternative variant fSep08, mRNA.
<a href="#">RGD1306583</a>	<a href="#">RGD1306583.gSep08</a>	<a href="#">294709</a>	3711	956	2	70	putative protein (8.0 kD) (RGD1306583) alternative variant gSep08, mRNA.
<a href="#">RGD1306595</a>	<a href="#">RGD1306595.bSep08</a>	<a href="#">287554</a>	4329	2004	3	216	similar to hypothetical protein (24.5 kD) (RGD1306595) alternative variant bSep08, complete mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.cSep08</a>	<a href="#">50560</a>	1988	524	6	142	CRA a (RGD1306603andlhpk1) alternative variant cSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.cSep08</a>	<a href="#">363144</a>	1988	524	6	142	CRA a (RGD1306603andlhpk1) alternative variant cSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.dSep08</a>	<a href="#">50560</a>	661	464	2	128	CRA d like (RGD1306603andlhpk1) alternative variant dSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.dSep08</a>	<a href="#">363144</a>	661	464	2	128	CRA d like (RGD1306603andlhpk1) alternative variant dSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.eSep08</a>	<a href="#">50560</a>	29300	739	3	126	inositol hexaphosphate kinase 1 (RGD1306603andlhpk1) alternative variant eSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.eSep08</a>	<a href="#">363144</a>	29300	739	3	126	inositol hexaphosphate kinase 1 (RGD1306603andlhpk1) alternative variant eSep08, mRNA.

<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.fSep08</a>	<a href="#">50560</a>	4849	1751	7	120	CRA a like (13.4 kD) (RGD1306603andlhpk1) alternative variant fSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.fSep08</a>	<a href="#">363144</a>	4849	1751	7	120	CRA a like (13.4 kD) (RGD1306603andlhpk1) alternative variant fSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.gSep08</a>	<a href="#">50560</a>	1864	604	5	112	CRA a like (12.5 kD) (RGD1306603andlhpk1) alternative variant gSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.gSep08</a>	<a href="#">363144</a>	1864	604	5	112	CRA a like (12.5 kD) (RGD1306603andlhpk1) alternative variant gSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.hSep08</a>	<a href="#">50560</a>	849	709	2	71	putative protein (RGD1306603andlhpk1) alternative variant hSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.hSep08</a>	<a href="#">363144</a>	849	709	2	71	putative protein (RGD1306603andlhpk1) alternative variant hSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.iSep08</a>	<a href="#">50560</a>	34351	1227	5	59	CRA c like (6.6 kD) (RGD1306603andlhpk1) alternative variant iSep08, mRNA.
<a href="#">RGD1306603andlhpk1</a>	<a href="#">RGD1306603andlhpk1.iSep08</a>	<a href="#">363144</a>	34351	1227	5	59	CRA c like (6.6 kD) (RGD1306603andlhpk1) alternative variant iSep08, mRNA.
<a href="#">RGD1306613</a>	<a href="#">RGD1306613.bSep08</a>	<a href="#">306994</a>	3764	720	3	139	similar to RIKEN cDNA 1600012F09 (15.3 kD) (RGD1306613) alternative variant bSep08, mRNA.
<a href="#">RGD1306613</a>	<a href="#">RGD1306613.cSep08</a>	<a href="#">306994</a>	4265	1135	3	139	similar to RIKEN cDNA 1600012F09 (15.3 kD) (RGD1306613) alternative variant cSep08, complete mRNA.
<a href="#">RGD1306613</a>	<a href="#">RGD1306613.dSep08</a>	<a href="#">306994</a>	1455	718	2	30	similar to RIKEN cDNA 1600012F09 (3.5 kD) (RGD1306613) alternative variant dSep08, mRNA.
<a href="#">RGD1306622</a>	<a href="#">RGD1306622.aSep08</a>	<a href="#">293736</a>	4570	1271		314	similar to KIAA0954 protein (RGD1306622) mRNA.
<a href="#">RGD1306636</a>	<a href="#">RGD1306636.aSep08</a>	<a href="#">293498</a>	3195	1306	1	360	hypothetical LOC293498 (38.2 kD) (RGD1306636) alternative variant aSep08, mRNA.
<a href="#">RGD1306636</a>	<a href="#">RGD1306636.bSep08</a>	<a href="#">293498</a>	3600	762	1	193	hypothetical LOC293498 (20.9 kD) (RGD1306636) alternative variant bSep08, mRNA.
<a href="#">RGD1306682</a>	<a href="#">RGD1306682.bSep08</a>	<a href="#">360617</a>	742	313	3	92	similar to RIKEN cDNA 1810046J19 (RGD1306682) alternative variant bSep08, mRNA.
<a href="#">RGD1306682</a>	<a href="#">RGD1306682.cSep08</a>	<a href="#">360617</a>	794	681	2	72	similar to RIKEN cDNA 1810046J19 (RGD1306682) alternative variant cSep08, mRNA.
<a href="#">RGD1306704</a>	<a href="#">RGD1306704.aSep08</a>	<a href="#">295483</a>	8620	590		145	hypothetical LOC295483 (17.0 kD) (RGD1306704) mRNA.
<a href="#">RGD1306717</a>	<a href="#">RGD1306717.aSep08</a>	<a href="#">311257</a>	38764	3828	27	905	similar to hypothetical protein MGC25461 (101.4 kD) (RGD1306717) alternative variant aSep08, mRNA.
<a href="#">RGD1306717</a>	<a href="#">RGD1306717.cSep08</a>	<a href="#">311257</a>	1613	701	1	117	similar to hypothetical protein MGC25461 (RGD1306717) alternative variant cSep08, mRNA.
<a href="#">RGD1306730</a>	<a href="#">RGD1306730.bSep08</a>	<a href="#">309009</a>	15218	509	1	53	putative protein of vertebrate origin (6.1 kD) (RGD1306730) alternative variant bSep08, mRNA.
<a href="#">RGD1306746</a>	<a href="#">RGD1306746.bSep08</a>	<a href="#">312511</a>	5833	390	1	64	similar to Hypothetical protein MGC25529 (RGD1306746) alternative variant bSep08, mRNA.
<a href="#">RGD1306750</a>	<a href="#">RGD1306750.aSep08</a>	<a href="#">362451</a>	62183	694	3	150	LOC362451 (16.0 kD) (RGD1306750) alternative variant aSep08, mRNA.
<a href="#">RGD1306750</a>	<a href="#">RGD1306750.cSep08</a>	<a href="#">362451</a>	2805	387	2	92	LOC362451 (RGD1306750) alternative variant cSep08, mRNA.
<a href="#">RGD1306772</a>	<a href="#">RGD1306772.aSep08</a>	<a href="#">288683</a>	3640	1566	3	258	similar to RIKEN cDNA 1110008J03 (28.1 kD) (RGD1306772) alternative variant aSep08, mRNA.

<a href="#">RGD1306783</a>	<a href="#">RGD1306783.bSep08</a>	<a href="#">304928</a>	2791	1798	1	29	similar to 2810422O20Rik protein (3.4 kD) (RGD1306783) alternative variant bSep08, mRNA.
<a href="#">RGD1306809</a>	<a href="#">RGD1306809.bSep08</a>	<a href="#">365699</a>	14408	605	1	172	similar to hypothetical protein FLJ30596 (RGD1306809) alternative variant bSep08, mRNA.
<a href="#">RGD1306811</a>	<a href="#">RGD1306811.aSep08</a>	<a href="#">300517</a>	12400	443		147	similar to hypothetical protein FLJ25530 (RGD1306811) mRNA.
<a href="#">RGD1306820</a>	<a href="#">RGD1306820.bSep08</a>	<a href="#">309069</a>	5620	731	3	132	similar to erythroid differentiation-related factor 1 (RGD1306820) alternative variant bSep08, mRNA.
<a href="#">RGD1306839</a>	<a href="#">RGD1306839.aSep08</a>	<a href="#">293888</a>	8388	713	4	147	similar to RIKEN cDNA 5033414D02 (17.3 kD) (RGD1306839) alternative variant aSep08, mRNA.
<a href="#">RGD1306839</a>	<a href="#">RGD1306839.cSep08</a>	<a href="#">293888</a>	9914	476	5	126	similar to RIKEN cDNA 5033414D02 (RGD1306839) alternative variant cSep08, mRNA.
<a href="#">RGD1306861</a>	<a href="#">RGD1306861.aSep08</a>	<a href="#">362257</a>	36531	2432		621	similar to RIKEN cDNA B230339M05 gene (RGD1306861) mRNA.
<a href="#">RGD1306862</a>	<a href="#">RGD1306862.aSep08</a>	<a href="#">287596</a>	7692	1870	4	552	similar to RIKEN cDNA 1200011M11 (RGD1306862) alternative variant aSep08, mRNA.
<a href="#">RGD1306873</a>	<a href="#">RGD1306873.aSep08</a>	<a href="#">304285</a>	15339	3419	16	623	similar to RIKEN cDNA 2210010N04 gene (RGD1306873) alternative variant aSep08, mRNA.
<a href="#">RGD1306873</a>	<a href="#">RGD1306873.bSep08</a>	<a href="#">304285</a>	10191	1537	11	206	similar to RIKEN cDNA 2210010N04 gene (23.4 kD) (RGD1306873) alternative variant bSep08, mRNA.
<a href="#">RGD1306873</a>	<a href="#">RGD1306873.dSep08</a>	<a href="#">304285</a>	1644	406	2	42	similar to RIKEN cDNA 2210010N04 gene (4.6 kD) (RGD1306873) alternative variant dSep08, mRNA.
<a href="#">RGD1306917</a>	<a href="#">RGD1306917.bSep08</a>	<a href="#">361805</a>	11560	452	1	59	similar to RIKEN cDNA 2900010M23 (RGD1306917) alternative variant bSep08, mRNA.
<a href="#">RGD1306926</a>	<a href="#">RGD1306926.aSep08</a>	<a href="#">303742</a>	10296	1892	2	454	similar to hypothetical protein FLJ22175 (RGD1306926) alternative variant aSep08, mRNA.
<a href="#">RGD1306926</a>	<a href="#">RGD1306926.bSep08</a>	<a href="#">303742</a>	1112	405	2	134	similar to hypothetical protein FLJ22175 (RGD1306926) alternative variant bSep08, mRNA.
<a href="#">RGD1306926</a>	<a href="#">RGD1306926.cSep08</a>	<a href="#">303742</a>	1338	1122	2	125	similar to hypothetical protein FLJ22175 (13.4 kD) (RGD1306926) alternative variant cSep08, mRNA.
<a href="#">RGD1306936</a>	<a href="#">RGD1306936.bSep08</a>	<a href="#">297082</a>	8991	910	1	137	uncharacterized protein c7orf30 homolog (RGD1306936) alternative variant bSep08, mRNA.
<a href="#">RGD1306939</a>	<a href="#">RGD1306939.bSep08</a>	<a href="#">306934</a>	59087	1400	13	281	similar to mKIAA0386 protein (RGD1306939) alternative variant bSep08, mRNA.
<a href="#">RGD1306939</a>	<a href="#">RGD1306939.cSep08</a>	<a href="#">306934</a>	11442	2299	4	133	similar to mKIAA0386 protein (RGD1306939) alternative variant cSep08, mRNA.
<a href="#">RGD1306954</a>	<a href="#">RGD1306954.bSep08</a>	<a href="#">288269</a>	5559	757	3	175	similar to RIKEN cDNA 1110004E09 (RGD1306954) alternative variant bSep08, mRNA.
<a href="#">RGD1306954</a>	<a href="#">RGD1306954.cSep08</a>	<a href="#">288269</a>	1285	517	3	114	similar to RIKEN cDNA 1110004E09 (RGD1306954) alternative variant cSep08, mRNA.
<a href="#">RGD1306954</a>	<a href="#">RGD1306954.eSep08</a>	<a href="#">288269</a>	4563	749	2	91	similar to RIKEN cDNA 1110004E09 (RGD1306954) alternative variant eSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.bSep08</a>	<a href="#">361624</a>	7772	692	5	207	similar to C11orf17 protein (RGD1306959) alternative variant bSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.cSep08</a>	<a href="#">361624</a>	8123	1047	5	181	similar to C11orf17 protein (19.3 kD) (RGD1306959) alternative variant cSep08, mRNA.

<a href="#">RGD1306959</a>	<a href="#">RGD1306959.dSep08</a>	<a href="#">361624</a>	7637	523	5	174	similar to C11orf17 protein (RGD1306959) alternative variant dSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.eSep08</a>	<a href="#">361624</a>	7771	577	4	168	similar to C11orf17 protein (RGD1306959) alternative variant eSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.fSep08</a>	<a href="#">361624</a>	4898	494	5	164	similar to C11orf17 protein (RGD1306959) alternative variant fSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.gSep08</a>	<a href="#">361624</a>	2872	665	2	135	similar to C11orf17 protein (RGD1306959) alternative variant gSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.hSep08</a>	<a href="#">361624</a>	4912	360	3	120	similar to C11orf17 protein (RGD1306959) alternative variant hSep08, mRNA.
<a href="#">RGD1306959</a>	<a href="#">RGD1306959.jSep08</a>	<a href="#">361624</a>	2953	344	2	54	similar to C11orf17 protein (RGD1306959) alternative variant jSep08, mRNA.
<a href="#">RGD1306962</a>	<a href="#">RGD1306962.aSep08</a>	<a href="#">309570</a>	30511	1851	10	302	putative protein, with at least 7 transmembrane domains, of ancient origin (RGD1306962) alternative variant aSep08, mRNA.
<a href="#">RGD1306962</a>	<a href="#">RGD1306962.bSep08</a>	<a href="#">309570</a>	19511	581	3	111	putative protein of metazoan origin (RGD1306962) alternative variant bSep08, mRNA.
<a href="#">RGD1306962</a>	<a href="#">RGD1306962.cSep08</a>	<a href="#">309570</a>	1097	721	2	64	putative protein of mammalian origin (7.1 kD) (RGD1306962) alternative variant cSep08, mRNA.
<a href="#">RGD1306991</a>	<a href="#">RGD1306991.bSep08</a>	<a href="#">362220</a>	11266	1095	1	131	similar to Protein C20orf103 precursor (RGD1306991) alternative variant bSep08, mRNA.
<a href="#">RGD1307032</a>	<a href="#">RGD1307032.bSep08</a>	<a href="#">360486</a>	1264	479	7	109	similar to RIKEN cDNA 1520401A03 gene (RGD1307032) alternative variant bSep08, mRNA.
<a href="#">RGD1307034</a>	<a href="#">RGD1307034.aSep08</a>	<a href="#">304244</a>	22491	1406	7	337	furry homolog (RGD1307034) alternative variant aSep08, mRNA.
<a href="#">RGD1307034</a>	<a href="#">RGD1307034.bSep08</a>	<a href="#">304244</a>	1174	833	2	127	furry homolog (14.4 kD) (RGD1307034) alternative variant bSep08, mRNA.
<a href="#">RGD1307034</a>	<a href="#">RGD1307034.cSep08</a>	<a href="#">304244</a>	3830	432	2	96	CRA b like (RGD1307034) alternative variant cSep08, mRNA.
<a href="#">RGD1307041</a>	<a href="#">RGD1307041.aSep08</a>	<a href="#">361182</a>	24891	1789	14	408	similar to hypothetical protein FLJ11305 (RGD1307041) alternative variant aSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.aSep08</a>	<a href="#">362472</a>	17470	2048	12	475	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant aSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.bSep08</a>	<a href="#">362472</a>	39860	1490	12	461	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant bSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.cSep08</a>	<a href="#">362472</a>	71423	1778	6	450	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant cSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.dSep08</a>	<a href="#">362472</a>	26400	568	4	179	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant dSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.eSep08</a>	<a href="#">362472</a>	26344	359	3	109	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant eSep08, mRNA.
<a href="#">RGD1307055</a>	<a href="#">RGD1307055.fSep08</a>	<a href="#">362472</a>	5284	626	3	94	similar to hypothetical protein FLJ22490 (RGD1307055) alternative variant fSep08, mRNA.
<a href="#">RGD1307067</a>	<a href="#">RGD1307067.aSep08</a>	<a href="#">362840</a>	7005	1778	8	263	LOC362840 (29.8 kD) (RGD1307067) alternative variant aSep08, complete mRNA.
<a href="#">RGD1307071</a>	<a href="#">RGD1307071.aSep08</a>	<a href="#">311502</a>	111208	2554	12	713	similar to uncharacterized hypothalamus protein HT013 (RGD1307071) alternative variant aSep08, mRNA.

<a href="#">RGD1307071</a>	<a href="#">RGD1307071.bSep08</a>	<a href="#">311502</a>	13270	763	2	108	similar to uncharacterized hypothalamus protein HT013 (RGD1307071) alternative variant bSep08, mRNA.
<a href="#">RGD1307084</a>	<a href="#">RGD1307084.bSep08</a>	<a href="#">287598</a>	17016	458	2	107	similar to RIKEN cDNA 1110001A07 gene (RGD1307084) alternative variant bSep08, mRNA.
<a href="#">RGD1307100</a>	<a href="#">RGD1307100.aSep08</a>	<a href="#">294978</a>	17301	2949	13	822	similar to RIKEN cDNA D630029K19 (RGD1307100) alternative variant aSep08, mRNA.
<a href="#">RGD1307100</a>	<a href="#">RGD1307100.bSep08</a>	<a href="#">294978</a>	4404	754	3	251	similar to RIKEN cDNA D630029K19 (RGD1307100) alternative variant bSep08, mRNA.
<a href="#">RGD1307100</a>	<a href="#">RGD1307100.dSep08</a>	<a href="#">294978</a>	1216	775	2	73	similar to RIKEN cDNA D630029K19 (RGD1307100) alternative variant dSep08, mRNA.
<a href="#">RGD1307119</a>	<a href="#">RGD1307119.bSep08</a>	<a href="#">303743</a>	4950	1184	9	381	hypothetical LOC303743 (RGD1307119) alternative variant bSep08, mRNA.
<a href="#">RGD1307119</a>	<a href="#">RGD1307119.cSep08</a>	<a href="#">303743</a>	4242	899	8	299	hypothetical LOC303743 (RGD1307119) alternative variant cSep08, mRNA.
<a href="#">RGD1307119</a>	<a href="#">RGD1307119.dSep08</a>	<a href="#">303743</a>	1161	1050	2	161	hypothetical LOC303743 (18.2 kD) (RGD1307119) alternative variant dSep08, mRNA.
<a href="#">RGD1307122</a>	<a href="#">RGD1307122.bSep08</a>	<a href="#">314321</a>	2082	1331	1	403	similar to RIKEN cDNA 2810002I04 (RGD1307122) alternative variant bSep08, mRNA.
<a href="#">RGD1307155</a>	<a href="#">RGD1307155.bSep08</a>	<a href="#">302998</a>	1027	552	5	126	similar to CG18661-PA (RGD1307155) alternative variant bSep08, mRNA.
<a href="#">RGD1307158</a>	<a href="#">RGD1307158.aSep08</a>	<a href="#">361773</a>	5325	1797	4	590	similar to oocyte-testis gene 1 (RGD1307158) alternative variant aSep08, mRNA.
<a href="#">RGD1307158</a>	<a href="#">RGD1307158.bSep08</a>	<a href="#">361773</a>	13943	1731	7	577	similar to oocyte-testis gene 1 (RGD1307158) alternative variant bSep08, mRNA.
<a href="#">RGD1307158</a>	<a href="#">RGD1307158.cSep08</a>	<a href="#">361773</a>	7451	2683	5	220	similar to oocyte-testis gene 1 (RGD1307158) alternative variant cSep08, mRNA.
<a href="#">RGD1307158</a>	<a href="#">RGD1307158.dSep08</a>	<a href="#">361773</a>	2746	728	2	59	similar to oocyte-testis gene 1 (RGD1307158) alternative variant dSep08, mRNA.
<a href="#">RGD1307158</a>	<a href="#">RGD1307158.eSep08</a>	<a href="#">361773</a>	2507	1178	2	42	similar to oocyte-testis gene 1 (RGD1307158) alternative variant eSep08, mRNA.
<a href="#">RGD1307161</a>	<a href="#">RGD1307161.aSep08</a>	<a href="#">305031</a>	19798	3354	13	411	similar to 0610010K06Rik protein (46.2 kD) (RGD1307161) alternative variant aSep08, complete mRNA.
<a href="#">RGD1307161</a>	<a href="#">RGD1307161.bSep08</a>	<a href="#">305031</a>	13117	565	7	187	similar to 0610010K06Rik protein (RGD1307161) alternative variant bSep08, mRNA.
<a href="#">RGD1307177</a>	<a href="#">RGD1307177.aSep08</a>	<a href="#">365380</a>	3437	329		41	similar to Hypothetical protein KIAA0555 (RGD1307177) mRNA.
<a href="#">RGD1307179</a>	<a href="#">RGD1307179.bSep08</a>	<a href="#">299900</a>	4611	897	3	143	similar to RIKEN cDNA D530033C11 (16.1 kD) (RGD1307179) alternative variant bSep08, mRNA.
<a href="#">RGD1307182</a>	<a href="#">RGD1307182.aSep08</a>	<a href="#">316207</a>	644	403		59	similar to RIKEN cDNA B430306N03 gene (RGD1307182) mRNA.
<a href="#">RGD1307201</a>	<a href="#">RGD1307201.bSep08</a>	<a href="#">361047</a>	18979	2247	8	442	protein c13orf3 homolog (RGD1307201) alternative variant bSep08, mRNA.
<a href="#">RGD1307201</a>	<a href="#">RGD1307201.cSep08</a>	<a href="#">361047</a>	3819	726	1	103	protein c13orf3 homolog (RGD1307201) alternative variant cSep08, mRNA.
<a href="#">RGD1307218</a>	<a href="#">RGD1307218.bSep08</a>	<a href="#">362518</a>	13277	684	1	123	similar to RIKEN cDNA 2810432L12 (RGD1307218) alternative variant bSep08, mRNA.

<a href="#">RGD1307218</a>	<a href="#">RGD1307218.cSep08</a>	<a href="#">362518</a>	2113	327	1	29	similar to RIKEN cDNA 2810432L12 (RGD1307218) alternative variant cSep08, mRNA.
<a href="#">RGD1307222</a>	<a href="#">RGD1307222.aSep08</a>	<a href="#">303300</a>	9952	3114	14	851	similar to mKIAA0664 protein (RGD1307222) alternative variant aSep08, mRNA.
<a href="#">RGD1307222</a>	<a href="#">RGD1307222.bSep08</a>	<a href="#">303300</a>	5530	2715	11	421	similar to mKIAA0664 protein (47.6 kD) (RGD1307222) alternative variant bSep08, mRNA.
<a href="#">RGD1307222</a>	<a href="#">RGD1307222.cSep08</a>	<a href="#">303300</a>	2097	993	7	262	similar to mKIAA0664 protein (RGD1307222) alternative variant cSep08, mRNA.
<a href="#">RGD1307222</a>	<a href="#">RGD1307222.dSep08</a>	<a href="#">303300</a>	10248	546	5	156	similar to mKIAA0664 protein (RGD1307222) alternative variant dSep08, mRNA.
<a href="#">RGD1307222</a>	<a href="#">RGD1307222.eSep08</a>	<a href="#">303300</a>	1232	413	4	137	similar to mKIAA0664 protein (RGD1307222) alternative variant eSep08, mRNA.
<a href="#">RGD1307234</a>	<a href="#">RGD1307234.aSep08</a>	<a href="#">316732</a>	39074	2883	17	667	similar to RIKEN cDNA 4931400A14 (RGD1307234) alternative variant aSep08, mRNA.
<a href="#">RGD1307234</a>	<a href="#">RGD1307234.bSep08</a>	<a href="#">316732</a>	2721	1784	3	113	similar to RIKEN cDNA 4931400A14 (13.1 kD) (RGD1307234) alternative variant bSep08, mRNA.
<a href="#">RGD1307235</a>	<a href="#">RGD1307235.bSep08</a>	<a href="#">309053</a>	14543	616	7	204	similar to RIKEN cDNA 2310035C23 (RGD1307235) alternative variant bSep08, mRNA.
<a href="#">RGD1307235</a>	<a href="#">RGD1307235.cSep08</a>	<a href="#">309053</a>	10159	1436	5	115	similar to RIKEN cDNA 2310035C23 (RGD1307235) alternative variant cSep08, mRNA.
<a href="#">RGD1307235</a>	<a href="#">RGD1307235.dSep08</a>	<a href="#">309053</a>	7458	337	4	93	similar to RIKEN cDNA 2310035C23 (RGD1307235) alternative variant dSep08, mRNA.
<a href="#">RGD1307254</a>	<a href="#">RGD1307254.cSep08</a>	<a href="#">298712</a>	2885	633	2	88	similar to RIKEN cDNA 1200011I18 (RGD1307254) alternative variant cSep08, mRNA.
<a href="#">RGD1307264</a>	<a href="#">RGD1307264.aSep08</a>	<a href="#">309637</a>	25050	843	6	281	similar to hypothetical protein FLJ20302; similar to CG31653-PA (RGD1307264) alternative variant aSep08, mRNA.
<a href="#">RGD1307284</a>	<a href="#">RGD1307284.aSep08</a>	<a href="#">306811</a>	51771	1904	10	574	similar to protein kinase, lysine deficient 1; kinase deficient protein (RGD1307284) alternative variant aSep08, mRNA.
<a href="#">RGD1307284</a>	<a href="#">RGD1307284.bSep08</a>	<a href="#">306811</a>	43227	568	4	189	similar to protein kinase, lysine deficient 1; kinase deficient protein (RGD1307284) alternative variant bSep08, mRNA.
<a href="#">RGD1307288</a>	<a href="#">RGD1307288.aSep08</a>	<a href="#">287761</a>	2782	1298		187	similar to Protein C21orf58 (20.5 kD) (RGD1307288) mRNA.
<a href="#">RGD1307325</a>	<a href="#">RGD1307325.bSep08</a>	<a href="#">306469</a>	15152	363	3	103	similar to RIKEN cDNA 4933411K20 (RGD1307325) alternative variant bSep08, mRNA.
<a href="#">RGD1307336</a>	<a href="#">RGD1307336.aSep08</a>	<a href="#">293950</a>	13206	742	4	146	similar to hypothetical protein (16.2 kD) (RGD1307336) alternative variant aSep08, mRNA.
<a href="#">RGD1307355</a>	<a href="#">RGD1307355.aSep08</a>	<a href="#">311825</a>	19105	2541	2	674	similar to gene model 711 (76.4 kD) (RGD1307355) alternative variant aSep08, mRNA.
<a href="#">RGD1307357</a>	<a href="#">RGD1307357.bSep08</a>	<a href="#">291975</a>	3378	596	5	198	similar to hypothetical protein DKFZp434A1319 (RGD1307357) alternative variant bSep08, mRNA.
<a href="#">RGD1307357</a>	<a href="#">RGD1307357.cSep08</a>	<a href="#">291975</a>	831	646	3	147	similar to hypothetical protein DKFZp434A1319 (RGD1307357) alternative variant cSep08, mRNA.
<a href="#">RGD1307365</a>	<a href="#">RGD1307365.aSep08</a>	<a href="#">300880</a>	10949	1892	1	216	similar to KIAA1009 protein (25.1 kD) (RGD1307365) alternative variant aSep08, mRNA.
<a href="#">RGD1307365</a>	<a href="#">RGD1307365.bSep08</a>	<a href="#">300880</a>	11906	771	2	210	similar to KIAA1009 protein (RGD1307365) alternative variant bSep08, mRNA.

<a href="#">RGD1307381</a>	<a href="#">RGD1307381.bSep08</a>	<a href="#">360498</a>	2686	1979	9	270	similar to RIKEN cDNA 2610003J06 (30.6 kD) (RGD1307381) alternative variant bSep08, complete mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.cSep08</a>	<a href="#">360498</a>	1939	1197	9	257	similar to RIKEN cDNA 2610003J06 (RGD1307381) alternative variant cSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.dSep08</a>	<a href="#">360498</a>	1764	1216	7	256	similar to RIKEN cDNA 2610003J06 (29.1 kD) (RGD1307381) alternative variant dSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.eSep08</a>	<a href="#">360498</a>	1598	887	9	251	similar to RIKEN cDNA 2610003J06 (RGD1307381) alternative variant eSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.fSep08</a>	<a href="#">360498</a>	1382	911	6	187	similar to RIKEN cDNA 2610003J06 (RGD1307381) alternative variant fSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.gSep08</a>	<a href="#">360498</a>	2020	1640	5	153	similar to RIKEN cDNA 2610003J06 (16.7 kD) (RGD1307381) alternative variant gSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.hSep08</a>	<a href="#">360498</a>	1794	1140	8	130	similar to RIKEN cDNA 2610003J06 (14.7 kD) (RGD1307381) alternative variant hSep08, mRNA.
<a href="#">RGD1307381</a>	<a href="#">RGD1307381.iSep08</a>	<a href="#">360498</a>	920	770	3	98	similar to RIKEN cDNA 2610003J06 (11.3 kD) (RGD1307381) alternative variant iSep08, complete mRNA.
<a href="#">RGD1307390</a>	<a href="#">RGD1307390.bSep08</a>	<a href="#">292680</a>	1587	721	4	88	similar to BC282485 1 (10.0 kD) (RGD1307390) alternative variant bSep08, mRNA.
<a href="#">RGD1307392</a>	<a href="#">RGD1307392.bSep08</a>	<a href="#">299209</a>	1210	642	2	106	similar to 1700019E19Rik protein (12.5 kD) (RGD1307392) alternative variant bSep08, mRNA.
<a href="#">RGD1307392</a>	<a href="#">RGD1307392.cSep08</a>	<a href="#">299209</a>	802	678	1	92	similar to 1700019E19Rik protein (10.4 kD) (RGD1307392) alternative variant cSep08, mRNA.
<a href="#">RGD1307394</a>	<a href="#">RGD1307394.bSep08</a>	<a href="#">360667</a>	3166	1424	3	158	putative protein, with at least 3 transmembrane domains, of eukaryotic origin (RGD1307394) alternative variant bSep08, mRNA.
<a href="#">RGD1307394</a>	<a href="#">RGD1307394.cSep08</a>	<a href="#">360667</a>	5859	734	6	110	putative endoplasmic reticulum protein, with a transmembrane domain, of eukaryotic origin (12.1 kD) (RGD1307394) alternative variant cSep08, mRNA.
<a href="#">RGD1307394</a>	<a href="#">RGD1307394.dSep08</a>	<a href="#">360667</a>	9961	1047	7	100	putative endoplasmic reticulum protein, with a transmembrane domain, of eukaryotic origin (10.8 kD) (RGD1307394) alternative variant dSep08, mRNA.
<a href="#">RGD1307394</a>	<a href="#">RGD1307394.eSep08</a>	<a href="#">360667</a>	4399	721	5	84	putative protein of metazoan origin (RGD1307394) alternative variant eSep08, mRNA.
<a href="#">RGD1307394</a>	<a href="#">RGD1307394.hSep08</a>	<a href="#">360667</a>	649	411	2	59	putative protein (RGD1307394) alternative variant hSep08, mRNA.
<a href="#">RGD1307396</a>	<a href="#">RGD1307396.aSep08</a>	<a href="#">360757</a>	22234	2293	1	518	similar to RIKEN cDNA 6330406115 (RGD1307396) alternative variant aSep08, mRNA.
<a href="#">RGD1307396</a>	<a href="#">RGD1307396.bSep08</a>	<a href="#">360757</a>	20921	748		108	similar to RIKEN cDNA 6330406115 (RGD1307396) alternative variant bSep08, mRNA.
<a href="#">RGD1307396</a>	<a href="#">RGD1307396.cSep08</a>	<a href="#">360757</a>	3747	1343	1	60	similar to RIKEN cDNA 6330406115 (RGD1307396) alternative variant cSep08, mRNA.
<a href="#">RGD1307399</a>	<a href="#">RGD1307399.aSep08</a>	<a href="#">681315</a>	9349	1131	2	120	putative endoplasmic reticulum protein, with a transmembrane domain, of eukaryotic origin (13.2 kD) (RGD1307399) alternative variant aSep08, mRNA.



<a href="#">RGD1307410</a>	<a href="#">RGD1307410.aSep08</a>	<a href="#">360673</a>	6527	2668	7	553	similar to hypothetical protein FLJ31528 (RGD1307410) alternative variant aSep08, mRNA.
<a href="#">RGD1307443</a>	<a href="#">RGD1307443.aSep08</a>	<a href="#">361244</a>	27899	1612		333	similar to mKIAA0319 protein (RGD1307443) mRNA.
<a href="#">RGD1307465</a>	<a href="#">RGD1307465.bSep08</a>	<a href="#">296200</a>	7146	939	3	285	CRA a (RGD1307465) alternative variant bSep08, mRNA.
<a href="#">RGD1307465</a>	<a href="#">RGD1307465.cSep08</a>	<a href="#">296200</a>	1347	443	1	112	putative protein (RGD1307465) alternative variant cSep08, mRNA.
<a href="#">RGD1307500</a>	<a href="#">RGD1307500.aSep08</a>	<a href="#">363056</a>	10964	829	1	276	LOC363056 (RGD1307500) alternative variant aSep08, mRNA.
<a href="#">RGD1307500</a>	<a href="#">RGD1307500.bSep08</a>	<a href="#">363056</a>	15102	393	1	70	LOC363056 (RGD1307500) alternative variant bSep08, mRNA.
<a href="#">RGD1307503</a>	<a href="#">RGD1307503.aSep08</a>	<a href="#">314456</a>	3087	342		113	similar to Hypothetical protein KIAA0297/KIAA0329 (RGD1307503) mRNA.
<a href="#">RGD1307506</a>	<a href="#">RGD1307506.cSep08</a>	<a href="#">294744</a>	3526	1457	2	135	similar to RIKEN cDNA 2310016C16 (15.9 kD) (RGD1307506) alternative variant cSep08, mRNA.
<a href="#">RGD1307506</a>	<a href="#">RGD1307506.dSep08</a>	<a href="#">294744</a>	844	370	2	122	similar to RIKEN cDNA 2310016C16 (RGD1307506) alternative variant dSep08, mRNA.
<a href="#">RGD1307509</a>	<a href="#">RGD1307509.bSep08</a>	<a href="#">310348</a>	20757	3528	8	332	similar to RIKEN cDNA 1700108L22 and hypothetical protein LOC689154 (RGD1307509) alternative variant bSep08, mRNA.
<a href="#">RGD1307509</a>	<a href="#">RGD1307509.bSep08</a>	<a href="#">689154</a>	20757	3528	8	332	similar to RIKEN cDNA 1700108L22 and hypothetical protein LOC689154 (RGD1307509) alternative variant bSep08, mRNA.
<a href="#">RGD1307509</a>	<a href="#">RGD1307509.cSep08</a>	<a href="#">310348</a>	5091	434	4	74	similar to RIKEN cDNA 1700108L22 and hypothetical protein LOC689154 (RGD1307509) alternative variant cSep08, mRNA.
<a href="#">RGD1307509</a>	<a href="#">RGD1307509.cSep08</a>	<a href="#">689154</a>	5091	434	4	74	similar to RIKEN cDNA 1700108L22 and hypothetical protein LOC689154 (RGD1307509) alternative variant cSep08, mRNA.
<a href="#">RGD1307524</a>	<a href="#">RGD1307524.aSep08</a>	<a href="#">309415</a>	75013	2212	11	522	similar to Friedreich ataxia region gene X123 (RGD1307524) alternative variant aSep08, mRNA.
<a href="#">RGD1307525</a>	<a href="#">RGD1307525.aSep08</a>	<a href="#">308053</a>	18825	535	3	114	similar to intracellular protein transport like (XM453) (RGD1307525) alternative variant aSep08, mRNA.
<a href="#">RGD1307525</a>	<a href="#">RGD1307525.cSep08</a>	<a href="#">308053</a>	5923	406	2	46	similar to intracellular protein transport like (XM453) (RGD1307525) alternative variant cSep08, mRNA.
<a href="#">RGD1307526</a>	<a href="#">RGD1307526.aSep08</a>	<a href="#">315792</a>	27274	3187	17	861	transcription modulator (RGD1307526) alternative variant aSep08, mRNA.
<a href="#">RGD1307526</a>	<a href="#">RGD1307526.bSep08</a>	<a href="#">315792</a>	11344	1052	9	319	SAFB-like transcription modulator (RGD1307526) alternative variant bSep08, mRNA.
<a href="#">RGD1307526</a>	<a href="#">RGD1307526.cSep08</a>	<a href="#">315792</a>	3438	913	4	270	transcription modulator (RGD1307526) alternative variant cSep08, mRNA.
<a href="#">RGD1307526</a>	<a href="#">RGD1307526.dSep08</a>	<a href="#">315792</a>	27777	793	6	231	transcription modulator (RGD1307526) alternative variant dSep08, mRNA.
<a href="#">RGD1307526</a>	<a href="#">RGD1307526.eSep08</a>	<a href="#">315792</a>	642	551	2	65	SAFB-like transcription modulator (6.9 kD) (RGD1307526) alternative variant eSep08, mRNA.
<a href="#">RGD1307537</a>	<a href="#">RGD1307537.bSep08</a>	<a href="#">291077</a>	11856	446	1	108	similar to RIKEN cDNA 4933417A18 (RGD1307537) alternative variant bSep08, mRNA.

<a href="#">RGD1307554</a>	<a href="#">RGD1307554.bSep08</a>	<a href="#">292739</a>	24936	1996	11	397	similar to CG16812-PA (43.1 kD) (RGD1307554) alternative variant bSep08, mRNA.
<a href="#">RGD1307554</a>	<a href="#">RGD1307554.cSep08</a>	<a href="#">292739</a>	5225	388	4	121	similar to CG16812-PA (RGD1307554) alternative variant cSep08, mRNA.
<a href="#">RGD1307554</a>	<a href="#">RGD1307554.eSep08</a>	<a href="#">292739</a>	5460	368	3	59	similar to CG16812-PA (RGD1307554) alternative variant eSep08, mRNA.
<a href="#">RGD1307569</a>	<a href="#">RGD1307569.aSep08</a>	<a href="#">360695</a>	38520	1307	5	357	similar to Protein C21orf63 homolog precursor (RGD1307569) alternative variant aSep08, mRNA.
<a href="#">RGD1307569</a>	<a href="#">RGD1307569.bSep08</a>	<a href="#">360695</a>	60499	1188	2	209	similar to Protein C21orf63 homolog precursor (23.3 kD) (RGD1307569) alternative variant bSep08, mRNA.
<a href="#">RGD1307569</a>	<a href="#">RGD1307569.cSep08</a>	<a href="#">360695</a>	73042	1783	2	161	similar to Protein C21orf63 homolog precursor (RGD1307569) alternative variant cSep08, mRNA.
<a href="#">RGD1307597</a>	<a href="#">RGD1307597.bSep08</a>	<a href="#">299197</a>	27873	394	3	72	similar to mKIAA0317 protein (RGD1307597) alternative variant bSep08, mRNA.
<a href="#">RGD1307603</a>	<a href="#">RGD1307603.aSep08</a>	<a href="#">293656</a>	1984	725	2	222	similar to hypothetical protein MGC37914 (RGD1307603) alternative variant aSep08, mRNA.
<a href="#">RGD1307615</a>	<a href="#">RGD1307615.bSep08</a>	<a href="#">362084</a>	2276	1412	4	108	similar to hypothetical protein FLJ13045 (12.1 kD) (RGD1307615) alternative variant bSep08, mRNA.
<a href="#">RGD1307621</a>	<a href="#">RGD1307621.bSep08</a>	<a href="#">314168</a>	1338	679	2	142	hypothetical LOC314168 (RGD1307621) alternative variant bSep08, mRNA.
<a href="#">RGD1307621</a>	<a href="#">RGD1307621.cSep08</a>	<a href="#">314168</a>	1402	616	3	82	hypothetical LOC314168 (9.8 kD) (RGD1307621) alternative variant cSep08, mRNA.
<a href="#">RGD1307648</a>	<a href="#">RGD1307648.aSep08</a>	<a href="#">294004</a>	19030	765	6	225	similar to CG13901-PA (RGD1307648) alternative variant aSep08, mRNA.
<a href="#">RGD1307648</a>	<a href="#">RGD1307648.bSep08</a>	<a href="#">294004</a>	19310	931	6	203	similar to CG13901-PA (23.2 kD) (RGD1307648) alternative variant bSep08, mRNA.
<a href="#">RGD1307648</a>	<a href="#">RGD1307648.cSep08</a>	<a href="#">294004</a>	13860	767	3	173	similar to CG13901-PA (19.1 kD) (RGD1307648) alternative variant cSep08, mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.bSep08</a>	<a href="#">300675</a>	16425	1639	13	217	CRA a (24.5 kD) (RGD1307682) alternative variant bSep08, complete mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.cSep08</a>	<a href="#">300675</a>	14364	859	7	210	CRA a like (RGD1307682) alternative variant cSep08, mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.dSep08</a>	<a href="#">300675</a>	4411	745	5	181	CRA a (RGD1307682) alternative variant dSep08, mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.eSep08</a>	<a href="#">300675</a>	14145	728	6	143	CRA a like (RGD1307682) alternative variant eSep08, mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.fSep08</a>	<a href="#">300675</a>	5226	701	4	115	CRA a (RGD1307682) alternative variant fSep08, mRNA.
<a href="#">RGD1307682</a>	<a href="#">RGD1307682.gSep08</a>	<a href="#">300675</a>	1451	684	3	63	CRA a (7.0 kD) (RGD1307682) alternative variant gSep08, mRNA.
<a href="#">RGD1307722</a>	<a href="#">RGD1307722.bSep08</a>	<a href="#">362824</a>	983	459	3	110	similar to hypothetical protein MGC20700 (RGD1307722) alternative variant bSep08, mRNA.
<a href="#">RGD1307722</a>	<a href="#">RGD1307722.dSep08</a>	<a href="#">362824</a>	498	412	2	35	similar to hypothetical protein MGC20700 (3.7 kD) (RGD1307722) alternative variant dSep08, mRNA.
<a href="#">RGD1307739</a>	<a href="#">RGD1307739.bSep08</a>	<a href="#">362122</a>	24506	573		150	similar to CG3306-PA (RGD1307739) alternative variant bSep08, mRNA.
<a href="#">RGD1307749</a>	<a href="#">RGD1307749.bSep08</a>	<a href="#">299338</a>	2150	754	3	251	similar to RIKEN cDNA 1600013K19 (RGD1307749) alternative variant bSep08, mRNA.

<a href="#">RGD1307749</a>	<a href="#">RGD1307749.cSep08</a>	<a href="#">299338</a>	3174	1395	4	194	similar to RIKEN cDNA 1600013K19 (RGD1307749) alternative variant cSep08, mRNA.
<a href="#">RGD1307749</a>	<a href="#">RGD1307749.dSep08</a>	<a href="#">299338</a>	1946	460	2	122	similar to RIKEN cDNA 1600013K19 (RGD1307749) alternative variant dSep08, mRNA.
<a href="#">RGD1307752</a>	<a href="#">RGD1307752.bSep08</a>	<a href="#">296315</a>	1443	664	2	62	similar to RIKEN cDNA 1110008F13 (RGD1307752) alternative variant bSep08, mRNA.
<a href="#">RGD1307773</a>	<a href="#">RGD1307773.aSep08</a>	<a href="#">287115</a>	2614	2522	2	330	similar to RIKEN cDNA 1700012G19 gene (RGD1307773) alternative variant aSep08, mRNA.
<a href="#">RGD1307791</a>	<a href="#">RGD1307791.bSep08</a>	<a href="#">297968</a>	2878	1994	3	81	similar to hypothetical protein FLJ10342 (9.5 kD) (RGD1307791) alternative variant bSep08, mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.bSep08</a>	<a href="#">307833</a>	11746	2168	7	340	similar to RIKEN cDNA 2400003C14 (37.6 kD) (RGD1307799) alternative variant bSep08, mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.cSep08</a>	<a href="#">307833</a>	18653	680	7	226	similar to RIKEN cDNA 2400003C14 (RGD1307799) alternative variant cSep08, mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.dSep08</a>	<a href="#">307833</a>	11417	882	7	187	similar to RIKEN cDNA 2400003C14 (19.8 kD) (RGD1307799) alternative variant dSep08, mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.eSep08</a>	<a href="#">307833</a>	14912	212	3	70	similar to RIKEN cDNA 2400003C14 (RGD1307799) alternative variant eSep08, mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.fSep08</a>	<a href="#">307833</a>	22112	3374	8	141	similar to RIKEN cDNA 2400003C14 (14.8 kD) (RGD1307799) alternative variant fSep08, complete mRNA.
<a href="#">RGD1307799</a>	<a href="#">RGD1307799.iSep08</a>	<a href="#">307833</a>	18653	779	8	65	similar to RIKEN cDNA 2400003C14 (RGD1307799) alternative variant iSep08, mRNA.
<a href="#">RGD1307801</a>	<a href="#">RGD1307801.bSep08</a>	<a href="#">309656</a>	7983	1490	11	351	similar to RIKEN cDNA 1300018I05 (39.5 kD) (RGD1307801) alternative variant bSep08, mRNA.
<a href="#">RGD1307801</a>	<a href="#">RGD1307801.cSep08</a>	<a href="#">309656</a>	11885	429	5	142	similar to RIKEN cDNA 1300018I05 (RGD1307801) alternative variant cSep08, mRNA.
<a href="#">RGD1307801</a>	<a href="#">RGD1307801.dSep08</a>	<a href="#">309656</a>	18749	581	3	78	similar to RIKEN cDNA 1300018I05 (RGD1307801) alternative variant dSep08, mRNA.
<a href="#">RGD1307816</a>	<a href="#">RGD1307816.bSep08</a>	<a href="#">295446</a>	1683	751	2	47	similar to RIKEN cDNA A630047E20 (5.4 kD) (RGD1307816) alternative variant bSep08, mRNA.
<a href="#">RGD1307830</a>	<a href="#">RGD1307830.aSep08</a>	<a href="#">304863</a>	8333	1114		114	protein odr-4 homolog like (RGD1307830) mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.bSep08</a>	<a href="#">316085</a>	5232	2054	2	372	trafficking protein kinesin binding 1 CRA c like (40.5 kD) (RGD1307844) alternative variant bSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.cSep08</a>	<a href="#">316085</a>	3175	1276	3	259	trafficking protein kinesin binding 1 CRA e like (RGD1307844) alternative variant cSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.dSep08</a>	<a href="#">316085</a>	42037	560	3	168	trafficking protein kinesin binding 1 CRA a like (RGD1307844) alternative variant dSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.eSep08</a>	<a href="#">316085</a>	118219	693	4	149	trafficking protein kinesin binding 1 CRA d like (RGD1307844) alternative variant eSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.fSep08</a>	<a href="#">316085</a>	45214	434	5	144	trafficking protein kinesin binding 1 like (RGD1307844) alternative variant fSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.gSep08</a>	<a href="#">316085</a>	6632	990	2	128	trafficking protein kinesin binding 1 CRA b like (RGD1307844) alternative variant gSep08, mRNA.
<a href="#">RGD1307844</a>	<a href="#">RGD1307844.jSep08</a>	<a href="#">316085</a>	1671	315	2	86	trafficking protein kinesin binding 1 CRA d like (RGD1307844) alternative variant jSep08, mRNA.

<a href="#">RGD1307844</a>	<a href="#">RGD1307844.kSep08</a>	<a href="#">316085</a>	5628	460	5	74	putative protein (RGD1307844) alternative variant kSep08, mRNA.
<a href="#">RGD1307877</a>	<a href="#">RGD1307877.bSep08</a>	<a href="#">291846</a>	10820	765	1	145	similar to plasma kallikrein-like protein 4 precursor (RGD1307877) alternative variant bSep08, mRNA.
<a href="#">RGD1307882</a>	<a href="#">RGD1307882.aSep08</a>	<a href="#">315903</a>	29815	3694	15	596	similar to CG9346-PA (69.5 kD) (RGD1307882) alternative variant aSep08, mRNA.
<a href="#">RGD1307882</a>	<a href="#">RGD1307882.bSep08</a>	<a href="#">315903</a>	52622	1830	8	463	similar to CG9346-PA (RGD1307882) alternative variant bSep08, mRNA.
<a href="#">RGD1307882</a>	<a href="#">RGD1307882.cSep08</a>	<a href="#">315903</a>	8011	648	7	215	similar to CG9346-PA (RGD1307882) alternative variant cSep08, mRNA.
<a href="#">RGD1307882</a>	<a href="#">RGD1307882.dSep08</a>	<a href="#">315903</a>	16809	487	1	107	similar to CG9346-PA (RGD1307882) alternative variant dSep08, mRNA.
<a href="#">RGD1307887</a>	<a href="#">RGD1307887.aSep08</a>	<a href="#">366176</a>	1793	758		252	similar to RIKEN cDNA 4930424G05 (RGD1307887) mRNA.
<a href="#">RGD1307890</a>	<a href="#">RGD1307890.bSep08</a>	<a href="#">304851</a>	15677	606	7	201	similar to C1orf25 (RGD1307890) alternative variant bSep08, mRNA.
<a href="#">RGD1307890</a>	<a href="#">RGD1307890.cSep08</a>	<a href="#">304851</a>	6377	826	4	199	similar to C1orf25 (RGD1307890) alternative variant cSep08, mRNA.
<a href="#">RGD1307890</a>	<a href="#">RGD1307890.dSep08</a>	<a href="#">304851</a>	14561	606	5	96	similar to C1orf25 (RGD1307890) alternative variant dSep08, mRNA.
<a href="#">RGD1307890</a>	<a href="#">RGD1307890.eSep08</a>	<a href="#">304851</a>	4041	751	2	81	similar to C1orf25 (RGD1307890) alternative variant eSep08, mRNA.
<a href="#">RGD1307897</a>	<a href="#">RGD1307897.aSep08</a>	<a href="#">314172</a>	17480	2002		369	similar to RIKEN cDNA C730036B14 gene (RGD1307897) mRNA.
<a href="#">RGD1307929</a>	<a href="#">RGD1307929.aSep08</a>	<a href="#">303280</a>	14224	2939	16	760	similar to CG14967-PA (RGD1307929) alternative variant aSep08, mRNA.
<a href="#">RGD1307929</a>	<a href="#">RGD1307929.bSep08</a>	<a href="#">303280</a>	24422	359	3	72	similar to CG14967-PA (RGD1307929) alternative variant bSep08, mRNA.
<a href="#">RGD1307934</a>	<a href="#">RGD1307934.aSep08</a>	<a href="#">293953</a>	48146	1031		343	similar to DNA segment, Chr 19, ERATO Doi 386, expressed (RGD1307934) mRNA.
<a href="#">RGD1307935</a>	<a href="#">RGD1307935.bSep08</a>	<a href="#">360583</a>	5288	1840	3	141	similar to Hypothetical protein MGC18716 (RGD1307935) alternative variant bSep08, mRNA.
<a href="#">RGD1307935</a>	<a href="#">RGD1307935.cSep08</a>	<a href="#">360583</a>	8192	1040	3	82	similar to Hypothetical protein MGC18716 (RGD1307935) alternative variant cSep08, mRNA.
<a href="#">RGD1307937</a>	<a href="#">RGD1307937.bSep08</a>	<a href="#">289865</a>	15914	1038	5	233	similar to hypothetical protein FLJ31438 (RGD1307937) alternative variant bSep08, mRNA.
<a href="#">RGD1307947</a>	<a href="#">RGD1307947.bSep08</a>	<a href="#">314788</a>	5351	577	4	192	similar to RIKEN cDNA C430008C19 (RGD1307947) alternative variant bSep08, mRNA.
<a href="#">RGD1307983</a>	<a href="#">RGD1307983.aSep08</a>	<a href="#">298032</a>	12571	1513	4	118	similar to HSPC043 protein (RGD1307983) alternative variant aSep08, mRNA.
<a href="#">RGD1307983</a>	<a href="#">RGD1307983.bSep08</a>	<a href="#">298032</a>	11751	774	5	104	similar to HSPC043 protein (11.4 kD) (RGD1307983) alternative variant bSep08, mRNA.
<a href="#">RGD1308009</a>	<a href="#">RGD1308009.bSep08</a>	<a href="#">291841</a>	52592	3004	21	906	transcription complex (RGD1308009) alternative variant bSep08, mRNA.
<a href="#">RGD1308009</a>	<a href="#">RGD1308009.cSep08</a>	<a href="#">291841</a>	6562	2018	5	309	transcription complex (RGD1308009) alternative variant cSep08, mRNA.

<a href="#">RGD1308009</a>	<a href="#">RGD1308009.dSep08</a>	<a href="#">291841</a>	5176	708	5	159	transcription complex (RGD1308009) alternative variant dSep08, mRNA.
<a href="#">RGD1308009</a>	<a href="#">RGD1308009.eSep08</a>	<a href="#">291841</a>	2126	747	3	90	transcription complex (RGD1308009) alternative variant eSep08, mRNA.
<a href="#">RGD1308009</a>	<a href="#">RGD1308009.gSep08</a>	<a href="#">291841</a>	1462	600	2	61	transcription complex (RGD1308009) alternative variant gSep08, mRNA.
<a href="#">RGD1308023</a>	<a href="#">RGD1308023.aSep08</a>	<a href="#">296211</a>	4034	815	1	129	similar to CG5521-PA (RGD1308023) alternative variant aSep08, mRNA.
<a href="#">RGD1308023</a>	<a href="#">RGD1308023.bSep08</a>	<a href="#">296211</a>	66556	457	3	60	similar to CG5521-PA (RGD1308023) alternative variant bSep08, mRNA.
<a href="#">RGD1308026</a>	<a href="#">RGD1308026.aSep08</a>	<a href="#">363029</a>	1409	769	2	256	similar to 2310047B19Rik protein (RGD1308026) alternative variant aSep08, mRNA.
<a href="#">RGD1308031</a>	<a href="#">RGD1308031.aSep08</a>	<a href="#">300442</a>	8600	1747	10	397	similar to RIKEN cDNA 2510048L02 (45.0 kD) (RGD1308031) alternative variant aSep08, mRNA.
<a href="#">RGD1308031</a>	<a href="#">RGD1308031.cSep08</a>	<a href="#">300442</a>	7203	850	6	283	similar to RIKEN cDNA 2510048L02 (RGD1308031) alternative variant cSep08, mRNA.
<a href="#">RGD1308031</a>	<a href="#">RGD1308031.dSep08</a>	<a href="#">300442</a>	5369	791	4	142	similar to RIKEN cDNA 2510048L02 (RGD1308031) alternative variant dSep08, mRNA.
<a href="#">RGD1308048</a>	<a href="#">RGD1308048.aSep08</a>	<a href="#">298557</a>	10531	1526	6	242	similar to HT014 (RGD1308048) alternative variant aSep08, mRNA.
<a href="#">RGD1308048</a>	<a href="#">RGD1308048.bSep08</a>	<a href="#">298557</a>	2769	1225	3	98	similar to HT014 (11.4 kD) (RGD1308048) alternative variant bSep08, mRNA.
<a href="#">RGD1308065</a>	<a href="#">RGD1308065.bSep08</a>	<a href="#">287935</a>	8404	813	1	226	hypothetical LOC287935 (RGD1308065) alternative variant bSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.aSep08</a>	<a href="#">361034</a>	11907	3062	14	807	similar to FLJ00128 protein (RGD1308093) alternative variant aSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.bSep08</a>	<a href="#">361034</a>	800	685	2	227	similar to FLJ00128 protein (RGD1308093) alternative variant bSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.cSep08</a>	<a href="#">361034</a>	3392	817	6	179	similar to FLJ00128 protein (RGD1308093) alternative variant cSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.dSep08</a>	<a href="#">361034</a>	1103	638	3	159	similar to FLJ00128 protein (RGD1308093) alternative variant dSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.eSep08</a>	<a href="#">361034</a>	2357	1281	5	134	similar to FLJ00128 protein (13.7 kD) (RGD1308093) alternative variant eSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.fSep08</a>	<a href="#">361034</a>	1245	671	3	86	similar to FLJ00128 protein (9.0 kD) (RGD1308093) alternative variant fSep08, mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.gSep08</a>	<a href="#">361034</a>	1384	1076	2	94	similar to FLJ00128 protein (10.0 kD) (RGD1308093) alternative variant gSep08, complete mRNA.
<a href="#">RGD1308093</a>	<a href="#">RGD1308093.hSep08</a>	<a href="#">361034</a>	3283	925	6	116	similar to FLJ00128 protein (RGD1308093) alternative variant hSep08, mRNA.
<a href="#">RGD1308101</a>	<a href="#">RGD1308101.bSep08</a>	<a href="#">679640</a>	22556	398	1	132	similar to hypothetical protein FLJ20276 (RGD1308101) alternative variant bSep08, mRNA.
<a href="#">RGD1308101</a>	<a href="#">RGD1308101.cSep08</a>	<a href="#">679640</a>	995	313	1	75	similar to hypothetical protein FLJ20276 (RGD1308101) alternative variant cSep08, mRNA.
<a href="#">RGD1308106</a>	<a href="#">RGD1308106.bSep08</a>	<a href="#">361719</a>	698	395	3	131	LOC361719 (RGD1308106) alternative variant bSep08, mRNA.

<a href="#">RGD1308112</a>	<a href="#">RGD1308112.aSep08</a>	<a href="#">289038</a>	7825	788		189	similar to 40S ribosomal protein S7 (S8) (RGD1308112) mRNA.
<a href="#">RGD1308113</a>	<a href="#">RGD1308113.bSep08</a>	<a href="#">290224</a>	3810	571	4	168	similar to CGI-112 protein (19.0 kD) (RGD1308113) alternative variant bSep08, mRNA.
<a href="#">RGD1308113</a>	<a href="#">RGD1308113.cSep08</a>	<a href="#">290224</a>	4038	1233	4	98	similar to CGI-112 protein (RGD1308113) alternative variant cSep08, mRNA.
<a href="#">RGD1308114</a>	<a href="#">RGD1308114.aSep08</a>	<a href="#">361253</a>	57576	387	4	128	uncharacterized protein C7orf10 (RGD1308114) alternative variant aSep08, mRNA.
<a href="#">RGD1308114</a>	<a href="#">RGD1308114.bSep08</a>	<a href="#">361253</a>	290451	689	3	55	uncharacterized protein C7orf10 like (5.9 kD) (RGD1308114) alternative variant bSep08, mRNA.
<a href="#">RGD1308116</a>	<a href="#">RGD1308116.aSep08</a>	<a href="#">310376</a>	15880	1331		400	similar to hypothetical protein MGC42105 (RGD1308116) mRNA.
<a href="#">RGD1308117</a>	<a href="#">RGD1308117.bSep08</a>	<a href="#">361066</a>	1955	1390	3	127	similar to 9930012K11Rik protein (14.2 kD) (RGD1308117) alternative variant bSep08, mRNA.
<a href="#">RGD1308119</a>	<a href="#">RGD1308119.bSep08</a>	<a href="#">302032</a>	7272	834	1	121	similar to F-box protein FBL2 (14.0 kD) (RGD1308119) alternative variant bSep08, complete mRNA.
<a href="#">RGD1308124</a>	<a href="#">RGD1308124.aSep08</a>	<a href="#">308631</a>	37997	407		135	similar to KIAA1357 protein (RGD1308124) mRNA.
<a href="#">RGD1308127</a>	<a href="#">RGD1308127.bSep08</a>	<a href="#">365493</a>	18587	815	6	162	similar to 2700078E11Rik protein (RGD1308127) alternative variant bSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.bSep08</a>	<a href="#">287452</a>	2751	821	5	184	CRA b (RGD1308134) alternative variant bSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.cSep08</a>	<a href="#">287452</a>	2477	702	4	169	putative protein of metazoan origin (RGD1308134) alternative variant cSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.dSep08</a>	<a href="#">287452</a>	2386	610	4	119	CRA b (RGD1308134) alternative variant dSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.eSep08</a>	<a href="#">287452</a>	1185	568	4	88	CRA b like (8.8 kD) (RGD1308134) alternative variant eSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.fSep08</a>	<a href="#">287452</a>	1450	743	3	73	putative cytoplasmic protein of metazoan origin (7.8 kD) (RGD1308134) alternative variant fSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.gSep08</a>	<a href="#">287452</a>	1568	338	3	67	putative protein of vertebrate origin (RGD1308134) alternative variant gSep08, mRNA.
<a href="#">RGD1308134</a>	<a href="#">RGD1308134.jSep08</a>	<a href="#">287452</a>	960	432	2	57	CRA d like (RGD1308134) alternative variant jSep08, mRNA.
<a href="#">RGD1308147</a>	<a href="#">RGD1308147.bSep08</a>	<a href="#">307008</a>	5449	791	2	150	putative protein of eukaryotic origin (RGD1308147) alternative variant bSep08, mRNA.
<a href="#">RGD1308147</a>	<a href="#">RGD1308147.cSep08</a>	<a href="#">307008</a>	5439	701	2	140	putative protein of eukaryotic origin (RGD1308147) alternative variant cSep08, mRNA.
<a href="#">RGD1308147</a>	<a href="#">RGD1308147.dSep08</a>	<a href="#">307008</a>	5410	828	3	59	putative protein (RGD1308147) alternative variant dSep08, mRNA.
<a href="#">RGD1308154</a>	<a href="#">RGD1308154.aSep08</a>	<a href="#">363799</a>	125377	812		270	similar to CG11388-PA (RGD1308154) mRNA.
<a href="#">RGD1308160</a>	<a href="#">RGD1308160.aSep08</a>	<a href="#">307556</a>	20086	684		227	similar to Myosin heavy chain A (MHC A) (RGD1308160) mRNA.
<a href="#">RGD1308168</a>	<a href="#">RGD1308168.aSep08</a>	<a href="#">303735</a>	5566	646		214	CRA a (RGD1308168) mRNA.
<a href="#">RGD1308195</a>	<a href="#">RGD1308195.bSep08</a>	<a href="#">315074</a>	1383	525	1	93	similar to secreted Ly6/uPAR related protein 2 (RGD1308195) alternative variant bSep08, mRNA.
<a href="#">RGD1308215</a>	<a href="#">RGD1308215.bSep08</a>	<a href="#">293493</a>	32703	455	3	91	similar to hypothetical protein DKFZp434I2117 (RGD1308215) alternative variant bSep08, mRNA.

<a href="#">RGD1308221</a>	<a href="#">RGD1308221.bSep08</a>	<a href="#">304645</a>	2784	368	1	122	similar to TBC1 domain family, member 8 (with GRAM domain); vascular Rab-GAP/TBC-containing (RGD1308221) alternative variant bSep08, mRNA.
<a href="#">RGD1308226</a>	<a href="#">RGD1308226.bSep08</a>	<a href="#">296968</a>	21002	401	4	133	similar to hypothetical protein FLJ32786 (RGD1308226) alternative variant bSep08, mRNA.
<a href="#">RGD1308226</a>	<a href="#">RGD1308226.cSep08</a>	<a href="#">296968</a>	5350	310	3	103	similar to hypothetical protein FLJ32786 (RGD1308226) alternative variant cSep08, mRNA.
<a href="#">RGD1308226</a>	<a href="#">RGD1308226.dSep08</a>	<a href="#">296968</a>	35446	761	1	53	similar to hypothetical protein FLJ32786 (RGD1308226) alternative variant dSep08, mRNA.
<a href="#">RGD1308234</a>	<a href="#">RGD1308234.aSep08</a>	<a href="#">309150</a>	3757	641	3	213	hypothetical LOC309150 (RGD1308234) alternative variant aSep08, mRNA.
<a href="#">RGD1308234</a>	<a href="#">RGD1308234.bSep08</a>	<a href="#">309150</a>	2534	460	2	152	hypothetical LOC309150 (RGD1308234) alternative variant bSep08, mRNA.
<a href="#">RGD1308234</a>	<a href="#">RGD1308234.cSep08</a>	<a href="#">309150</a>	2515	384	1	127	hypothetical LOC309150 (RGD1308234) alternative variant cSep08, mRNA.
<a href="#">RGD1308251</a>	<a href="#">RGD1308251.bSep08</a>	<a href="#">362676</a>	19538	396	1	79	similar to RIKEN cDNA 2810405K02 (RGD1308251) alternative variant bSep08, mRNA.
<a href="#">RGD1308257</a>	<a href="#">RGD1308257.bSep08</a>	<a href="#">362437</a>	13762	3078	9	309	similar to intermediate filament-like protein MGC:2625 isoform 2; HOM-TES-103 tumor antigen-like (35.6 kD) (RGD1308257) alternative variant bSep08, mRNA.
<a href="#">RGD1308257</a>	<a href="#">RGD1308257.cSep08</a>	<a href="#">362437</a>	12209	1930	9	199	similar to intermediate filament-like protein MGC:2625 isoform 2; HOM-TES-103 tumor antigen-like (22.8 kD) (RGD1308257) alternative variant cSep08, mRNA.
<a href="#">RGD1308257</a>	<a href="#">RGD1308257.dSep08</a>	<a href="#">362437</a>	8698	693	3	140	similar to intermediate filament-like protein MGC:2625 isoform 2; HOM-TES-103 tumor antigen-like (RGD1308257) alternative variant dSep08, mRNA.
<a href="#">RGD1308257</a>	<a href="#">RGD1308257.eSep08</a>	<a href="#">362437</a>	3801	1091	5	118	similar to intermediate filament-like protein MGC:2625 isoform 2; HOM-TES-103 tumor antigen-like (RGD1308257) alternative variant eSep08, mRNA.
<a href="#">RGD1308268</a>	<a href="#">RGD1308268.aSep08</a>	<a href="#">365869</a>	11761	811		270	similar to KIAA0460 protein (RGD1308268) mRNA.
<a href="#">RGD1308290</a>	<a href="#">RGD1308290.bSep08</a>	<a href="#">360845</a>	12973	680	7	226	similar to RIKEN cDNA 5730454B08 (RGD1308290) alternative variant bSep08, mRNA.
<a href="#">RGD1308290</a>	<a href="#">RGD1308290.cSep08</a>	<a href="#">360845</a>	12056	990	4	107	similar to RIKEN cDNA 5730454B08 (RGD1308290) alternative variant cSep08, mRNA.
<a href="#">RGD1308290</a>	<a href="#">RGD1308290.dSep08</a>	<a href="#">360845</a>	6445	756	2	42	similar to RIKEN cDNA 5730454B08 (RGD1308290) alternative variant dSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.aSep08</a>	<a href="#">306137</a>	39998	2665	15	689	RNA binding motif protein 26 like (RGD1308297) alternative variant aSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.bSep08</a>	<a href="#">306137</a>	28389	2080	10	374	rna binding motif protein 26 CRA a like (RGD1308297) alternative variant bSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.cSep08</a>	<a href="#">306137</a>	12133	563	6	175	rna binding motif protein 26 CRA a like (RGD1308297) alternative variant cSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.dSep08</a>	<a href="#">306137</a>	11886	955	5	172	RNA binding motif protein 26 like (RGD1308297) alternative variant dSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.eSep08</a>	<a href="#">306137</a>	1250	413	3	137	RNA binding motif protein 26 like (RGD1308297) alternative variant eSep08, mRNA.

<a href="#">RGD1308297</a>	<a href="#">RGD1308297.fSep08</a>	<a href="#">306137</a>	19080	1373	4	77	putative protein (RGD1308297) alternative variant fSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.gSep08</a>	<a href="#">306137</a>	4759	568	3	42	putative protein (5.1 kD) (RGD1308297) alternative variant gSep08, mRNA.
<a href="#">RGD1308297</a>	<a href="#">RGD1308297.iSep08</a>	<a href="#">306137</a>	2319	655	3	24	putative protein (2.9 kD) (RGD1308297) alternative variant iSep08, mRNA.
<a href="#">RGD1308299</a>	<a href="#">RGD1308299.bSep08</a>	<a href="#">367214</a>	6247	791	6	152	solution Structure protein Domain (17.1 kD) (RGD1308299) alternative variant bSep08, mRNA.
<a href="#">RGD1308299</a>	<a href="#">RGD1308299.cSep08</a>	<a href="#">367214</a>	7429	748	7	152	solution Structure protein Domain (17.1 kD) (RGD1308299) alternative variant cSep08, mRNA.
<a href="#">RGD1308302</a>	<a href="#">RGD1308302.aSep08</a>	<a href="#">308911</a>	2741	1776	2	478	similar to RIKEN cDNA 1500003O22 (RGD1308302) alternative variant aSep08, mRNA.
<a href="#">RGD1308302</a>	<a href="#">RGD1308302.cSep08</a>	<a href="#">308911</a>	2642	739	2	222	similar to RIKEN cDNA 1500003O22 (RGD1308302) alternative variant cSep08, mRNA.
<a href="#">RGD1308319</a>	<a href="#">RGD1308319.aSep08</a>	<a href="#">316764</a>	5305	811		140	similar to KIAA0802 protein (RGD1308319) mRNA.
<a href="#">RGD1308333</a>	<a href="#">RGD1308333.bSep08</a>	<a href="#">292138</a>	12970	2402	7	289	similar to enolase (46.6 kD) (2J223) (31.9 kD) (RGD1308333) alternative variant bSep08, mRNA.
<a href="#">RGD1308333</a>	<a href="#">RGD1308333.cSep08</a>	<a href="#">292138</a>	13552	746	3	184	similar to enolase (46.6 kD) (2J223) (RGD1308333) alternative variant cSep08, mRNA.
<a href="#">RGD1308380</a>	<a href="#">RGD1308380.bSep08</a>	<a href="#">305443</a>	5335	1152	2	108	similar to RIKEN cDNA 2310079F23 (12.5 kD) (RGD1308380) alternative variant bSep08, mRNA.
<a href="#">RGD1308396</a>	<a href="#">RGD1308396.bSep08</a>	<a href="#">290235</a>	1523	1147	2	76	putative protein of metazoan origin (8.4 kD) (RGD1308396) alternative variant bSep08, mRNA.
<a href="#">RGD1308428</a>	<a href="#">RGD1308428.aSep08</a>	<a href="#">308509</a>	43223	1050		349	similar to RIKEN cDNA 4931406P16 (RGD1308428) mRNA.
<a href="#">RGD1308430</a>	<a href="#">RGD1308430.aSep08</a>	<a href="#">361038</a>	4630	983	1	142	similar to 1700123O20Rik protein (15.9 kD) (RGD1308430) alternative variant aSep08, mRNA.
<a href="#">RGD1308430</a>	<a href="#">RGD1308430.cSep08</a>	<a href="#">361038</a>	4065	604	1	65	similar to 1700123O20Rik protein (RGD1308430) alternative variant cSep08, mRNA.
<a href="#">RGD1308432</a>	<a href="#">RGD1308432.aSep08</a>	<a href="#">362324</a>	18135	2131	8	441	WD repeat-containing protein mio like (RGD1308432) alternative variant aSep08, mRNA.
<a href="#">RGD1308432</a>	<a href="#">RGD1308432.bSep08</a>	<a href="#">362324</a>	1598	702	1	83	WD repeat-containing protein mio like (9.3 kD) (RGD1308432) alternative variant bSep08, mRNA.
<a href="#">RGD1308461</a>	<a href="#">RGD1308461.aSep08</a>	<a href="#">307901</a>	8024	368		101	similar to RIKEN cDNA 4632415K11 (RGD1308461) mRNA.
<a href="#">RGD1308470</a>	<a href="#">RGD1308470.bSep08</a>	<a href="#">362778</a>	22853	2089	2	139	similar to RIKEN cDNA 4933433P14 gene (15.6 kD) (RGD1308470) alternative variant bSep08, mRNA.
<a href="#">RGD1308489</a>	<a href="#">RGD1308489.aSep08</a>	<a href="#">313340</a>	27216	2810	11	671	similar to hypothetical protein (RGD1308489) alternative variant aSep08, mRNA.
<a href="#">RGD1308489</a>	<a href="#">RGD1308489.bSep08</a>	<a href="#">313340</a>	7506	1513	4	145	similar to hypothetical protein (16.8 kD) (RGD1308489) alternative variant bSep08, mRNA.
<a href="#">RGD1308517</a>	<a href="#">RGD1308517.aSep08</a>	<a href="#">290722</a>	16269	1801	8	546	similar to KIAA1712 protein (RGD1308517) alternative variant aSep08, mRNA.
<a href="#">RGD1308517</a>	<a href="#">RGD1308517.cSep08</a>	<a href="#">290722</a>	20049	2356	9	271	similar to KIAA1712 protein (30.3 kD) (RGD1308517) alternative variant cSep08, mRNA.
<a href="#">RGD1308517</a>	<a href="#">RGD1308517.dSep08</a>	<a href="#">290722</a>	4261	480	4	159	similar to KIAA1712 protein (RGD1308517) alternative variant dSep08, mRNA.



<a href="#">RGD1308523</a>	<a href="#">RGD1308523.aSep08</a>	<a href="#">289502</a>	3554	455	1	109	similar to RIKEN cDNA 4932413O14 gene (RGD1308523) alternative variant aSep08, mRNA.
<a href="#">RGD1308523</a>	<a href="#">RGD1308523.bSep08</a>	<a href="#">289502</a>	3443	357	1	77	similar to RIKEN cDNA 4932413O14 gene (RGD1308523) alternative variant bSep08, mRNA.
<a href="#">RGD1308523</a>	<a href="#">RGD1308523.cSep08</a>	<a href="#">289502</a>	3427	349	1	74	similar to RIKEN cDNA 4932413O14 gene (RGD1308523) alternative variant cSep08, mRNA.
<a href="#">RGD1308541</a>	<a href="#">RGD1308541.aSep08</a>	<a href="#">308768</a>	1498	776		237	hypothetical LOC308768 (RGD1308541) mRNA.
<a href="#">RGD1308544</a>	<a href="#">RGD1308544.aSep08</a>	<a href="#">361192</a>	24751	612		114	LOC361192 (RGD1308544) mRNA.
<a href="#">RGD1308557</a>	<a href="#">RGD1308557.cSep08</a>	<a href="#">312965</a>	6331	389	3	16	similar to homolog of rat p47 (2.0 kD) (RGD1308557) alternative variant cSep08, mRNA.
<a href="#">RGD1308564</a>	<a href="#">RGD1308564.aSep08</a>	<a href="#">287305</a>	7021	731		153	similar to L-amino acid oxidase 1 (RGD1308564) mRNA.
<a href="#">RGD1308584</a>	<a href="#">RGD1308584.aSep08</a>	<a href="#">289083</a>	15352	1081	5	183	similar to RIKEN cDNA 5730449L18 (RGD1308584) alternative variant aSep08, mRNA.
<a href="#">RGD1308601</a>	<a href="#">RGD1308601.bSep08</a>	<a href="#">307249</a>	43376	854	1	230	similar to hypothetical protein (RGD1308601) alternative variant bSep08, mRNA.
<a href="#">RGD1308612</a>	<a href="#">RGD1308612.aSep08</a>	<a href="#">311713</a>	3412	1767	4	222	similar to Protein C20orf20 (RGD1308612) alternative variant aSep08, mRNA.
<a href="#">RGD1308612</a>	<a href="#">RGD1308612.bSep08</a>	<a href="#">311713</a>	3063	1109	5	212	similar to Protein C20orf20 (RGD1308612) alternative variant bSep08, mRNA.
<a href="#">RGD1308616</a>	<a href="#">RGD1308616.aSep08</a>	<a href="#">362573</a>	8219	1939		538	similar to KIAA0467 protein (RGD1308616) alternative variant aSep08, mRNA.
<a href="#">RGD1308616</a>	<a href="#">RGD1308616.bSep08</a>	<a href="#">362573</a>	10040	1518		403	similar to KIAA0467 protein (RGD1308616) alternative variant bSep08, mRNA.
<a href="#">RGD1308626</a>	<a href="#">RGD1308626.aSep08</a>	<a href="#">361087</a>	3672	929	1	309	similar to 9630044O09Rik protein (RGD1308626) alternative variant aSep08, mRNA.
<a href="#">RGD1308635</a>	<a href="#">RGD1308635.bSep08</a>	<a href="#">296587</a>	1766	786	3	162	similar to CG12379-PA (17.2 kD) (RGD1308635) alternative variant bSep08, complete mRNA.
<a href="#">RGD1308635</a>	<a href="#">RGD1308635.cSep08</a>	<a href="#">296587</a>	989	760	2	119	similar to CG12379-PA (RGD1308635) alternative variant cSep08, mRNA.
<a href="#">RGD1308635</a>	<a href="#">RGD1308635.dSep08</a>	<a href="#">296587</a>	1747	1562	2	112	similar to CG12379-PA (11.9 kD) (RGD1308635) alternative variant dSep08, complete mRNA.
<a href="#">RGD1308695</a>	<a href="#">RGD1308695.bSep08</a>	<a href="#">289088</a>	24977	440	6	98	CRA b like (11.3 kD) (RGD1308695) alternative variant bSep08, mRNA.
<a href="#">RGD1308695</a>	<a href="#">RGD1308695.cSep08</a>	<a href="#">289088</a>	15149	721	7	88	putative protein of vertebrate origin (RGD1308695) alternative variant cSep08, mRNA.
<a href="#">RGD1308695</a>	<a href="#">RGD1308695.dSep08</a>	<a href="#">289088</a>	6007	425	2	83	CRA b like (RGD1308695) alternative variant dSep08, mRNA.
<a href="#">RGD1308699</a>	<a href="#">RGD1308699.aSep08</a>	<a href="#">309790</a>	8576	2177	3	371	similar to 1700060H10Rik protein (41.4 kD) (RGD1308699) alternative variant aSep08, mRNA.
<a href="#">RGD1308706</a>	<a href="#">RGD1308706.bSep08</a>	<a href="#">291925</a>	25656	701	4	134	similar to RIKEN cDNA 4921524J17 (15.7 kD) (RGD1308706) alternative variant bSep08, mRNA.
<a href="#">RGD1308706</a>	<a href="#">RGD1308706.cSep08</a>	<a href="#">291925</a>	26306	768	3	107	similar to RIKEN cDNA 4921524J17 (RGD1308706) alternative variant cSep08, mRNA.
<a href="#">RGD1308706</a>	<a href="#">RGD1308706.dSep08</a>	<a href="#">291925</a>	20310	407	3	12	similar to RIKEN cDNA 4921524J17 (RGD1308706) alternative variant dSep08, mRNA.
<a href="#">RGD1308720</a>	<a href="#">RGD1308720.aSep08</a>	<a href="#">362945</a>	6473	2148	4	521	similar to Peroxidasin CG12002-PA (56.3 kD) (RGD1308720) alternative variant aSep08, mRNA.

<a href="#">RGD1308720</a>	<a href="#">RGD1308720.bSep08</a>	<a href="#">362945</a>	1669	546	1	117	similar to Peroxidasin CG12002-PA (12.3 kD) (RGD1308720) alternative variant bSep08, mRNA.
<a href="#">RGD1308722</a>	<a href="#">RGD1308722.bSep08</a>	<a href="#">361822</a>	5818	784	5	199	putative protein of ancient origin (RGD1308722) alternative variant bSep08, mRNA.
<a href="#">RGD1308723</a>	<a href="#">RGD1308723.bSep08</a>	<a href="#">292160</a>	19753	804	9	268	similar to hypothetical protein FLJ20729 (RGD1308723) alternative variant bSep08, mRNA.
<a href="#">RGD1308723</a>	<a href="#">RGD1308723.cSep08</a>	<a href="#">292160</a>	16557	794	7	264	similar to hypothetical protein FLJ20729 (RGD1308723) alternative variant cSep08, mRNA.
<a href="#">RGD1308723</a>	<a href="#">RGD1308723.dSep08</a>	<a href="#">292160</a>	6560	681	6	186	similar to hypothetical protein FLJ20729 (RGD1308723) alternative variant dSep08, mRNA.
<a href="#">RGD1308723</a>	<a href="#">RGD1308723.eSep08</a>	<a href="#">292160</a>	5338	1348	2	97	similar to hypothetical protein FLJ20729 (RGD1308723) alternative variant eSep08, mRNA.
<a href="#">RGD1308729</a>	<a href="#">RGD1308729.bSep08</a>	<a href="#">314584</a>	56924	623	3	72	similar to ZFP-like protein (8.3 kD) (RGD1308729) alternative variant bSep08, mRNA.
<a href="#">RGD1308742</a>	<a href="#">RGD1308742.aSep08</a>	<a href="#">362120</a>	11350	1777		528	similar to Complement C5 precursor (RGD1308742) mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.bSep08</a>	<a href="#">290668</a>	6450	3921	6	182	CRA a (20.5 kD) (RGD1308759) alternative variant bSep08, mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.cSep08</a>	<a href="#">290668</a>	4085	778	5	157	CRA c (RGD1308759) alternative variant cSep08, mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.dSep08</a>	<a href="#">290668</a>	14676	1466	5	105	CRA c (RGD1308759) alternative variant dSep08, mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.eSep08</a>	<a href="#">290668</a>	2773	682	3	91	CRA c like (RGD1308759) alternative variant eSep08, mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.fSep08</a>	<a href="#">290668</a>	889	613	2	80	CRA c like (8.9 kD) (RGD1308759) alternative variant fSep08, mRNA.
<a href="#">RGD1308759</a>	<a href="#">RGD1308759.gSep08</a>	<a href="#">290668</a>	4899	446	2	56	CRA c like (RGD1308759) alternative variant gSep08, mRNA.
<a href="#">RGD1308772</a>	<a href="#">RGD1308772.aSep08</a>	<a href="#">290381</a>	127007	2602		510	similar to KIAA0564 protein (RGD1308772) mRNA.
<a href="#">RGD1308782</a>	<a href="#">RGD1308782.aSep08</a>	<a href="#">308318</a>	2321	572	3	190	similar to Zinc finger protein OZF (POZF-1) (RGD1308782) alternative variant aSep08, mRNA.
<a href="#">RGD1308813</a>	<a href="#">RGD1308813.bSep08</a>	<a href="#">303606</a>	3466	796	4	108	similar to adipocyte-specific protein 4 (13.1 kD) (RGD1308813) alternative variant bSep08, mRNA.
<a href="#">RGD1308813</a>	<a href="#">RGD1308813.cSep08</a>	<a href="#">303606</a>	1825	610	3	66	similar to adipocyte-specific protein 4 (RGD1308813) alternative variant cSep08, mRNA.
<a href="#">RGD1308818</a>	<a href="#">RGD1308818.bSep08</a>	<a href="#">311113</a>	18796	1441	1	412	similar to hypothetical protein (49.6 kD) (RGD1308818) alternative variant bSep08, complete mRNA.
<a href="#">RGD1308872</a>	<a href="#">RGD1308872.bSep08</a>	<a href="#">291787</a>	13690	334	3	111	similar to Retinoblastoma-binding protein 8 (RBBP-8) (CtBP interacting protein) (CtIP) (Retinoblastoma-interacting protein and myosin-like) (RIM) (RGD1308872) alternative variant bSep08, mRNA.
<a href="#">RGD1308872</a>	<a href="#">RGD1308872.cSep08</a>	<a href="#">291787</a>	11974	314	5	104	similar to Retinoblastoma-binding protein 8 (RBBP-8) (CtBP interacting protein) (CtIP) (Retinoblastoma-interacting protein and myosin-like) (RIM) (RGD1308872) alternative variant cSep08, mRNA.
<a href="#">RGD1308872</a>	<a href="#">RGD1308872.dSep08</a>	<a href="#">291787</a>	21142	814	5	94	similar to Retinoblastoma-binding protein 8 (RBBP-8) (CtBP interacting protein) (CtIP) (Retinoblastoma-interacting protein and myosin-like) (RIM) (10.9 kD) (RGD1308872) alternative variant dSep08, mRNA.

<a href="#">RGD1308874</a>	<a href="#">RGD1308874.bSep08</a>	<a href="#">366227</a>	38021	526	3	148	similar to RIKEN cDNA 2310001A20 (RGD1308874) alternative variant bSep08, mRNA.
<a href="#">RGD1308874</a>	<a href="#">RGD1308874.cSep08</a>	<a href="#">366227</a>	15014	697	2	145	similar to RIKEN cDNA 2310001A20 (RGD1308874) alternative variant cSep08, mRNA.
<a href="#">RGD1308876</a>	<a href="#">RGD1308876.bSep08</a>	<a href="#">362597</a>	1367	869	2	219	similar to 2610027C15Rik protein (RGD1308876) alternative variant bSep08, mRNA.
<a href="#">RGD1308876</a>	<a href="#">RGD1308876.cSep08</a>	<a href="#">362597</a>	1357	941	2	103	similar to 2610027C15Rik protein (RGD1308876) alternative variant cSep08, mRNA.
<a href="#">RGD1308907</a>	<a href="#">RGD1308907.bSep08</a>	<a href="#">314325</a>	27275	3225	1	90	similar to FLJ20689 (RGD1308907) alternative variant bSep08, mRNA.
<a href="#">RGD1308908</a>	<a href="#">RGD1308908.aSep08</a>	<a href="#">288514</a>	11949	3314	23	511	similar to KIAA1440 protein (57.4 kD) (RGD1308908) alternative variant aSep08, mRNA.
<a href="#">RGD1308908</a>	<a href="#">RGD1308908.bSep08</a>	<a href="#">288514</a>	1339	944	2	196	similar to KIAA1440 protein (RGD1308908) alternative variant bSep08, mRNA.
<a href="#">RGD1308908</a>	<a href="#">RGD1308908.cSep08</a>	<a href="#">288514</a>	1287	624	4	156	similar to KIAA1440 protein (RGD1308908) alternative variant cSep08, mRNA.
<a href="#">RGD1308908</a>	<a href="#">RGD1308908.eSep08</a>	<a href="#">288514</a>	702	406	3	96	similar to KIAA1440 protein (RGD1308908) alternative variant eSep08, mRNA.
<a href="#">RGD1308915</a>	<a href="#">RGD1308915.bSep08</a>	<a href="#">363545</a>	3214	2039	2	147	similar to RIKEN cDNA 1200013P24 (RGD1308915) alternative variant bSep08, mRNA.
<a href="#">RGD1308918</a>	<a href="#">RGD1308918.bSep08</a>	<a href="#">293570</a>	21334	682	1	153	similar to KIAA0157 gene product is novel (RGD1308918) alternative variant bSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.aSep08</a>	<a href="#">361614</a>	21227	1783	5	586	CRA b like (RGD1308929) alternative variant aSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.cSep08</a>	<a href="#">361614</a>	139735	585	6	194	CRA b like (RGD1308929) alternative variant cSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.dSep08</a>	<a href="#">361614</a>	9663	551	4	177	protein fam168a (RGD1308929) alternative variant dSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.eSep08</a>	<a href="#">361614</a>	133649	559	4	153	putative protein (RGD1308929) alternative variant eSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.fSep08</a>	<a href="#">361614</a>	123228	433	4	96	CRA e like (RGD1308929) alternative variant fSep08, mRNA.
<a href="#">RGD1308929</a>	<a href="#">RGD1308929.gSep08</a>	<a href="#">361614</a>	39026	678	3	82	putative protein (RGD1308929) alternative variant gSep08, mRNA.
<a href="#">RGD1308952</a>	<a href="#">RGD1308952.aSep08</a>	<a href="#">303002</a>	9321	2608	7	278	similar to mKIAA0665 protein (RGD1308952) alternative variant aSep08, mRNA.
<a href="#">RGD1308952</a>	<a href="#">RGD1308952.bSep08</a>	<a href="#">303002</a>	17119	627	2	155	similar to mKIAA0665 protein (RGD1308952) alternative variant bSep08, mRNA.
<a href="#">RGD1308952</a>	<a href="#">RGD1308952.cSep08</a>	<a href="#">303002</a>	23782	1779	2	50	similar to mKIAA0665 protein (RGD1308952) alternative variant cSep08, mRNA.
<a href="#">RGD1308958</a>	<a href="#">RGD1308958.aSep08</a>	<a href="#">298020</a>	30470	1174		391	putative protein, with at least 8 transmembrane domains, of metazoan origin (RGD1308958) mRNA.
<a href="#">RGD1308977</a>	<a href="#">RGD1308977.aSep08</a>	<a href="#">291312</a>	9625	791	4	146	similar to RIKEN cDNA 1110017116 (RGD1308977) alternative variant aSep08, mRNA.
<a href="#">RGD1308977</a>	<a href="#">RGD1308977.cSep08</a>	<a href="#">291312</a>	4940	978	1	85	similar to RIKEN cDNA 1110017116 (10.7 kD) (RGD1308977) alternative variant cSep08, mRNA.

<a href="#">RGD1309019</a>	<a href="#">RGD1309019.bSep08</a>	<a href="#">314596</a>	1205	762	2	246	similar to Ras GTPase-activating protein nGAP (RAS protein activator like 1) (RGD1309019) alternative variant bSep08, mRNA.
<a href="#">RGD1309019</a>	<a href="#">RGD1309019.cSep08</a>	<a href="#">314596</a>	4109	662	3	220	similar to Ras GTPase-activating protein nGAP (RAS protein activator like 1) (RGD1309019) alternative variant cSep08, mRNA.
<a href="#">RGD1309020</a>	<a href="#">RGD1309020.bSep08</a>	<a href="#">291060</a>	14852	902	6	245	similar to RIKEN cDNA 6530403A03 (RGD1309020) alternative variant bSep08, mRNA.
<a href="#">RGD1309020</a>	<a href="#">RGD1309020.cSep08</a>	<a href="#">291060</a>	9883	752	5	144	similar to RIKEN cDNA 6530403A03 (16.6 kD) (RGD1309020) alternative variant cSep08, mRNA.
<a href="#">RGD1309020</a>	<a href="#">RGD1309020.fSep08</a>	<a href="#">291060</a>	1995	197	2	31	similar to RIKEN cDNA 6530403A03 (RGD1309020) alternative variant fSep08, mRNA.
<a href="#">RGD1309028</a>	<a href="#">RGD1309028.aSep08</a>	<a href="#">299265</a>	1040	383		127	similar to RIKEN cDNA A830059I20 (RGD1309028) mRNA.
<a href="#">RGD1309036</a>	<a href="#">RGD1309036.aSep08</a>	<a href="#">292874</a>	2703	746	3	199	hypothetical LOC292874 (RGD1309036) alternative variant aSep08, mRNA.
<a href="#">RGD1309051</a>	<a href="#">RGD1309051.bSep08</a>	<a href="#">299153</a>	8385	924	2	240	CRA a like (RGD1309051) alternative variant bSep08, mRNA.
<a href="#">RGD1309051</a>	<a href="#">RGD1309051.cSep08</a>	<a href="#">299153</a>	4794	898	5	237	putative protein of eukaryotic origin (RGD1309051) alternative variant cSep08, mRNA.
<a href="#">RGD1309051</a>	<a href="#">RGD1309051.dSep08</a>	<a href="#">299153</a>	15810	586	3	195	CRA a (RGD1309051) alternative variant dSep08, mRNA.
<a href="#">RGD1309058</a>	<a href="#">RGD1309058.aSep08</a>	<a href="#">314311</a>	3758	346	2	115	similar to RIKEN cDNA 9830169C18 (RGD1309058) alternative variant aSep08, mRNA.
<a href="#">RGD1309085</a>	<a href="#">RGD1309085.aSep08</a>	<a href="#">297821</a>	15643	587		195	similar to F23N19.9 (RGD1309085) mRNA.
<a href="#">RGD1309102</a>	<a href="#">RGD1309102.bSep08</a>	<a href="#">291750</a>	1527	658	1	133	similar to TRS85 homolog (RGD1309102) alternative variant bSep08, mRNA.
<a href="#">RGD1309104</a>	<a href="#">RGD1309104.bSep08</a>	<a href="#">289084</a>	118246	561	3	111	putative protein (RGD1309104) alternative variant bSep08, mRNA.
<a href="#">RGD1309108</a>	<a href="#">RGD1309108.aSep08</a>	<a href="#">315578</a>	3762	422		140	similar to hypothetical protein FLJ23554 (RGD1309108) mRNA.
<a href="#">RGD1309148</a>	<a href="#">RGD1309148.aSep08</a>	<a href="#">298147</a>	1101	876	2	119	similar to RIKEN cDNA 3110001D03 (RGD1309148) alternative variant aSep08, mRNA.
<a href="#">RGD1309148</a>	<a href="#">RGD1309148.bSep08</a>	<a href="#">298147</a>	787	472	2	104	similar to RIKEN cDNA 3110001D03 (RGD1309148) alternative variant bSep08, mRNA.
<a href="#">RGD1309170</a>	<a href="#">RGD1309170.aSep08</a>	<a href="#">362047</a>	24024	857		202	similar to hypothetical protein DKFz434G072 (RGD1309170) mRNA.
<a href="#">RGD1309172</a>	<a href="#">RGD1309172.aSep08</a>	<a href="#">315979</a>	9549	637		212	similar to RIKEN cDNA E330026B02 (RGD1309172) mRNA.
<a href="#">RGD1309198</a>	<a href="#">RGD1309198.bSep08</a>	<a href="#">313056</a>	12872	1491	1	77	similar to U5 snRNP-specific protein (Prp8-binding) (8.5 kD) (RGD1309198) alternative variant bSep08, mRNA.
<a href="#">RGD1309216</a>	<a href="#">RGD1309216.bSep08</a>	<a href="#">361726</a>	28018	1412	5	154	similar to hypothetical protein FLJ20487 (18.5 kD) (RGD1309216) alternative variant bSep08, mRNA.
<a href="#">RGD1309216</a>	<a href="#">RGD1309216.cSep08</a>	<a href="#">361726</a>	23679	784	5	132	similar to hypothetical protein FLJ20487 (15.5 kD) (RGD1309216) alternative variant cSep08, complete mRNA.
<a href="#">RGD1309216</a>	<a href="#">RGD1309216.dSep08</a>	<a href="#">361726</a>	27651	717	3	33	similar to hypothetical protein FLJ20487 (RGD1309216) alternative variant dSep08, mRNA.

<a href="#">RGD1309216</a>	<a href="#">RGD1309216.eSep08</a>	<a href="#">361726</a>	23844	562	2	37	similar to hypothetical protein FLJ20487 (RGD1309216) alternative variant eSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.bSep08</a>	<a href="#">316328</a>	3885	734	6	244	protein CRA g (RGD1309220) alternative variant bSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.dSep08</a>	<a href="#">316328</a>	17337	941	4	116	uncharacterized protein (11.6 kD) (RGD1309220) alternative variant dSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.eSep08</a>	<a href="#">316328</a>	1322	734	2	105	putative protein (RGD1309220) alternative variant eSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.gSep08</a>	<a href="#">316328</a>	3466	399	3	72	protein CRA e (8.4 kD) (RGD1309220) alternative variant gSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.hSep08</a>	<a href="#">316328</a>	578	393	2	72	protein CRA e (8.4 kD) (RGD1309220) alternative variant hSep08, mRNA.
<a href="#">RGD1309220</a>	<a href="#">RGD1309220.iSep08</a>	<a href="#">316328</a>	1140	374	2	68	protein CRA e (RGD1309220) alternative variant iSep08, mRNA.
<a href="#">RGD1309228</a>	<a href="#">RGD1309228.bSep08</a>	<a href="#">298851</a>	7967	770	1	127	similar to putative protein, with at least 9 transmembrane domains, of eukaryotic origin (43.9 kD) (2G415) (RGD1309228) alternative variant bSep08, mRNA.
<a href="#">RGD1309285</a>	<a href="#">RGD1309285.aSep08</a>	<a href="#">303599</a>	17618	379		126	similar to KIAA1636 protein (RGD1309285) mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.bSep08</a>	<a href="#">361044</a>	2837	1643	5	264	putative cytoplasmic protein of ancient origin (28.6 kD) (RGD1309307) alternative variant bSep08, complete mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.cSep08</a>	<a href="#">361044</a>	2846	638	6	164	putative protein of ancient origin (18.1 kD) (RGD1309307) alternative variant cSep08, complete mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.dSep08</a>	<a href="#">361044</a>	2096	617	6	163	putative protein of ancient origin (RGD1309307) alternative variant dSep08, mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.eSep08</a>	<a href="#">361044</a>	1858	761	4	143	putative protein of ancient origin (RGD1309307) alternative variant eSep08, mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.gSep08</a>	<a href="#">361044</a>	2801	1783	5	79	putative protein of ancient origin (8.4 kD) (RGD1309307) alternative variant gSep08, mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.hSep08</a>	<a href="#">361044</a>	1026	685	3	67	putative protein of metazoan origin (RGD1309307) alternative variant hSep08, mRNA.
<a href="#">RGD1309307</a>	<a href="#">RGD1309307.iSep08</a>	<a href="#">361044</a>	743	657	2	36	putative protein (3.8 kD) (RGD1309307) alternative variant iSep08, complete mRNA.
<a href="#">RGD1309308</a>	<a href="#">RGD1309308.bSep08</a>	<a href="#">313115</a>	1279	866	1	62	similar to RIKEN cDNA 1810074P20 (RGD1309308) alternative variant bSep08, mRNA.
<a href="#">RGD1309313</a>	<a href="#">RGD1309313.bSep08</a>	<a href="#">309454</a>	3316	1302	2	249	similar to RIKEN cDNA 4930538D17 (RGD1309313) alternative variant bSep08, mRNA.
<a href="#">RGD1309313</a>	<a href="#">RGD1309313.cSep08</a>	<a href="#">309454</a>	1320	858	2	216	similar to RIKEN cDNA 4930538D17 (23.8 kD) (RGD1309313) alternative variant cSep08, mRNA.
<a href="#">RGD1309313</a>	<a href="#">RGD1309313.dSep08</a>	<a href="#">309454</a>	4500	445	4	147	similar to RIKEN cDNA 4930538D17 (RGD1309313) alternative variant dSep08, mRNA.
<a href="#">RGD1309326</a>	<a href="#">RGD1309326.aSep08</a>	<a href="#">308568</a>	1705	750	3	214	similar to RIKEN cDNA 2410002F23 (RGD1309326) alternative variant aSep08, mRNA.
<a href="#">RGD1309326</a>	<a href="#">RGD1309326.cSep08</a>	<a href="#">308568</a>	1451	661	2	99	similar to RIKEN cDNA 2410002F23 (RGD1309326) alternative variant cSep08, mRNA.
<a href="#">RGD1309326</a>	<a href="#">RGD1309326.dSep08</a>	<a href="#">308568</a>	5153	2146	5	167	similar to RIKEN cDNA 2410002F23 (18.8 kD) (RGD1309326) alternative variant dSep08, mRNA.

<a href="#">RGD1309326</a>	<a href="#">RGD1309326.eSep08</a>	<a href="#">308568</a>	5946	1952	6	284	similar to RIKEN cDNA 2410002F23 (32.4 kD) (RGD1309326) alternative variant eSep08, mRNA.
<a href="#">RGD1309326</a>	<a href="#">RGD1309326.gSep08</a>	<a href="#">308568</a>	4588	760	7	76	similar to RIKEN cDNA 2410002F23 (RGD1309326) alternative variant gSep08, mRNA.
<a href="#">RGD1309350</a>	<a href="#">RGD1309350.bSep08</a>	<a href="#">293613</a>	3110	919	2	118	similar to transthyretin (4L369) (13.5 kD) (RGD1309350) alternative variant bSep08, mRNA.
<a href="#">RGD1309350</a>	<a href="#">RGD1309350.cSep08</a>	<a href="#">293613</a>	2714	502	2	106	similar to transthyretin (4L369) (RGD1309350) alternative variant cSep08, mRNA.
<a href="#">RGD1309360</a>	<a href="#">RGD1309360.bSep08</a>	<a href="#">294715</a>	92087	1619	6	170	hypothetical LOC294715 (19.3 kD) (RGD1309360) alternative variant bSep08, mRNA.
<a href="#">RGD1309362</a>	<a href="#">RGD1309362.bSep08</a>	<a href="#">307415</a>	7919	342	1	79	similar to interferon-inducible GTPase (RGD1309362) alternative variant bSep08, mRNA.
<a href="#">RGD1309368</a>	<a href="#">RGD1309368.aSep08</a>	<a href="#">366031</a>	6656	403		133	similar to 5930402A21 protein (RGD1309368) mRNA.
<a href="#">RGD1309387</a>	<a href="#">RGD1309387.aSep08</a>	<a href="#">313777</a>	11783	2600	19	748	nucleolar complex associated 2 homolog (85.8 kD) (RGD1309387) alternative variant aSep08, complete mRNA.
<a href="#">RGD1309387</a>	<a href="#">RGD1309387.cSep08</a>	<a href="#">313777</a>	2459	1481	3	170	nucleolar complex associated 2 homolog (19.7 kD) (RGD1309387) alternative variant cSep08, mRNA.
<a href="#">RGD1309387</a>	<a href="#">RGD1309387.dSep08</a>	<a href="#">313777</a>	1783	437	3	69	nucleolar complex associated 2 homolog (RGD1309387) alternative variant dSep08, mRNA.
<a href="#">RGD1309403</a>	<a href="#">RGD1309403.bSep08</a>	<a href="#">310417</a>	1326	750	2	90	similar to hypothetical protein FLJ12661 (9.2 kD) (RGD1309403) alternative variant bSep08, mRNA.
<a href="#">RGD1309410</a>	<a href="#">RGD1309410.bSep08</a>	<a href="#">363020</a>	70648	752	2	111	LOC363020 (6.3 kD) (RGD1309410) alternative variant bSep08, mRNA.
<a href="#">RGD1309414</a>	<a href="#">RGD1309414.aSep08</a>	<a href="#">361004</a>	5198	2711	11	790	similar to KIAA0913 protein (RGD1309414) alternative variant aSep08, mRNA.
<a href="#">RGD1309414</a>	<a href="#">RGD1309414.bSep08</a>	<a href="#">361004</a>	8755	1220	8	327	similar to KIAA0913 protein (RGD1309414) alternative variant bSep08, mRNA.
<a href="#">RGD1309414</a>	<a href="#">RGD1309414.cSep08</a>	<a href="#">361004</a>	2025	1648	4	145	similar to KIAA0913 protein (15.5 kD) (RGD1309414) alternative variant cSep08, mRNA.
<a href="#">RGD1309437</a>	<a href="#">RGD1309437.bSep08</a>	<a href="#">288176</a>	36740	395	1	66	similar to RIKEN cDNA 2610528E23 (RGD1309437) alternative variant bSep08, mRNA.
<a href="#">RGD1309459</a>	<a href="#">RGD1309459.bSep08</a>	<a href="#">360477</a>	20344	461	7	153	similar to RIKEN cDNA 3930401K13 (RGD1309459) alternative variant bSep08, mRNA.
<a href="#">RGD1309471</a>	<a href="#">RGD1309471.bSep08</a>	<a href="#">293719</a>	3887	940	9	281	similar to hypothetical protein MGC6696 (RGD1309471) alternative variant bSep08, mRNA.
<a href="#">RGD1309471</a>	<a href="#">RGD1309471.cSep08</a>	<a href="#">293719</a>	3571	515	4	140	similar to hypothetical protein MGC6696 (RGD1309471) alternative variant cSep08, mRNA.
<a href="#">RGD1309471</a>	<a href="#">RGD1309471.dSep08</a>	<a href="#">293719</a>	862	687	1	74	similar to hypothetical protein MGC6696 (8.3 kD) (RGD1309471) alternative variant dSep08, mRNA.
<a href="#">RGD1309482</a>	<a href="#">RGD1309482.bSep08</a>	<a href="#">365458</a>	31887	1464	14	292	putative protein of eukaryotic origin (RGD1309482) alternative variant bSep08, mRNA.
<a href="#">RGD1309482</a>	<a href="#">RGD1309482.cSep08</a>	<a href="#">365458</a>	5047	711	4	115	putative protein (RGD1309482) alternative variant cSep08, mRNA.
<a href="#">RGD1309482</a>	<a href="#">RGD1309482.dSep08</a>	<a href="#">365458</a>	33110	701	7	102	putative protein of eukaryotic origin (RGD1309482) alternative variant dSep08, mRNA.

<a href="#">RGD1309482</a>	<a href="#">RGD1309482.gSep08</a>	<a href="#">365458</a>	2923	268	2	20	putative protein (RGD1309482) alternative variant gSep08, mRNA.
<a href="#">RGD1309483</a>	<a href="#">RGD1309483.aSep08</a>	<a href="#">366915</a>	16323	454		150	similar to hypothetical protein MGC39715 (RGD1309483) mRNA.
<a href="#">RGD1309492</a>	<a href="#">RGD1309492.aSep08</a>	<a href="#">314330</a>	22822	3985	4	468	similar to mKIAA1737 protein (RGD1309492) alternative variant aSep08, mRNA.
<a href="#">RGD1309492</a>	<a href="#">RGD1309492.dSep08</a>	<a href="#">314330</a>	19150	407	4	135	similar to mKIAA1737 protein (RGD1309492) alternative variant dSep08, mRNA.
<a href="#">RGD1309492</a>	<a href="#">RGD1309492.eSep08</a>	<a href="#">314330</a>	8106	445	2	92	similar to mKIAA1737 protein (RGD1309492) alternative variant eSep08, mRNA.
<a href="#">RGD1309501</a>	<a href="#">RGD1309501.bSep08</a>	<a href="#">305552</a>	68508	949		316	hypothetical LOC305552 (RGD1309501) alternative variant bSep08, mRNA.
<a href="#">RGD1309501</a>	<a href="#">RGD1309501.cSep08</a>	<a href="#">305552</a>	39511	518		119	hypothetical LOC305552 (RGD1309501) alternative variant cSep08, mRNA.
<a href="#">RGD1309522</a>	<a href="#">RGD1309522.bSep08</a>	<a href="#">306102</a>	3191	737	2	200	similar to hypothetical protein FLJ22624 (RGD1309522) alternative variant bSep08, mRNA.
<a href="#">RGD1309522</a>	<a href="#">RGD1309522.cSep08</a>	<a href="#">306102</a>	6362	853	5	140	similar to hypothetical protein FLJ22624 (RGD1309522) alternative variant cSep08, mRNA.
<a href="#">RGD1309534</a>	<a href="#">RGD1309534.bSep08</a>	<a href="#">363016</a>	3464	335	2	72	similar to RIKEN cDNA 4931406C07 (RGD1309534) alternative variant bSep08, mRNA.
<a href="#">RGD1309534</a>	<a href="#">RGD1309534.dSep08</a>	<a href="#">363016</a>	6221	437	4	40	similar to RIKEN cDNA 4931406C07 (RGD1309534) alternative variant dSep08, mRNA.
<a href="#">RGD1309540</a>	<a href="#">RGD1309540.bSep08</a>	<a href="#">295930</a>	3681	1069	1	70	similar to hypothetical protein MGC40841; similar to hypothetical protein MGC4707 (7.7 kD) (RGD1309540) alternative variant bSep08, mRNA.
<a href="#">RGD1309543</a>	<a href="#">RGD1309543.aSep08</a>	<a href="#">361790</a>	9789	2461	4	658	similar to 2310014H01Rik protein (71.7 kD) (RGD1309543) alternative variant aSep08, mRNA.
<a href="#">RGD1309576</a>	<a href="#">RGD1309576.aSep08</a>	<a href="#">287405</a>	9005	1176	6	166	similar to RIKEN cDNA A730055C05 gene (RGD1309576) alternative variant aSep08, mRNA.
<a href="#">RGD1309576</a>	<a href="#">RGD1309576.bSep08</a>	<a href="#">287405</a>	8310	390	4	130	similar to RIKEN cDNA A730055C05 gene (RGD1309576) alternative variant bSep08, mRNA.
<a href="#">RGD1309578</a>	<a href="#">RGD1309578.bSep08</a>	<a href="#">361637</a>	6239	900	3	139	similar to Aa2-174 (16.0 kD) (RGD1309578) alternative variant bSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.aSep08</a>	<a href="#">290963</a>	23414	1141	5	198	similar to hypothetical protein FLJ14675 (RGD1309592) alternative variant aSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.bSep08</a>	<a href="#">290963</a>	96315	665	8	197	similar to hypothetical protein FLJ14675 (RGD1309592) alternative variant bSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.dSep08</a>	<a href="#">290963</a>	29310	1330	5	108	similar to hypothetical protein FLJ14675 (12.9 kD) (RGD1309592) alternative variant dSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.eSep08</a>	<a href="#">290963</a>	1108	632	2	108	similar to hypothetical protein FLJ14675 (RGD1309592) alternative variant eSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.fSep08</a>	<a href="#">290963</a>	9170	488	2	26	similar to hypothetical protein FLJ14675 (2.9 kD) (RGD1309592) alternative variant fSep08, mRNA.
<a href="#">RGD1309592</a>	<a href="#">RGD1309592.gSep08</a>	<a href="#">290963</a>	9057	390	2	43	similar to hypothetical protein FLJ14675 (RGD1309592) alternative variant gSep08, mRNA.

<a href="#">RGD1309594</a>	<a href="#">RGD1309594.bSep08</a>	<a href="#">309681</a>	5818	704	7	185	similar to RIKEN cDNA 1810043G02; DNA segment, Chr 10, Johns Hopkins University 13, expressed (RGD1309594) alternative variant bSep08, mRNA.
<a href="#">RGD1309594</a>	<a href="#">RGD1309594.cSep08</a>	<a href="#">309681</a>	1842	759	3	130	similar to RIKEN cDNA 1810043G02; DNA segment, Chr 10, Johns Hopkins University 13, expressed (RGD1309594) alternative variant cSep08, mRNA.
<a href="#">RGD1309594</a>	<a href="#">RGD1309594.dSep08</a>	<a href="#">309681</a>	4496	754	5	96	similar to RIKEN cDNA 1810043G02; DNA segment, Chr 10, Johns Hopkins University 13, expressed (RGD1309594) alternative variant dSep08, mRNA.
<a href="#">RGD1309594</a>	<a href="#">RGD1309594.eSep08</a>	<a href="#">309681</a>	2899	414	3	56	similar to RIKEN cDNA 1810043G02; DNA segment, Chr 10, Johns Hopkins University 13, expressed (RGD1309594) alternative variant eSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.bSep08</a>	<a href="#">291320</a>	59490	1436	10	272	putative secreted or extracellular protein precursor of eukaryotic origin (30.6 kD) (RGD1309605) alternative variant bSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.cSep08</a>	<a href="#">291320</a>	77658	1311	12	245	putative cytoplasmic protein of eukaryotic origin (27.4 kD) (RGD1309605) alternative variant cSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.dSep08</a>	<a href="#">291320</a>	56169	992	10	177	putative cytoplasmic protein of eukaryotic origin (19.5 kD) (RGD1309605) alternative variant dSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.eSep08</a>	<a href="#">291320</a>	68846	728	7	144	putative protein of eukaryotic origin (RGD1309605) alternative variant eSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.fSep08</a>	<a href="#">291320</a>	70447	750	8	122	putative protein of eukaryotic origin (RGD1309605) alternative variant fSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.gSep08</a>	<a href="#">291320</a>	70426	870	10	105	putative protein of eukaryotic origin (RGD1309605) alternative variant gSep08, mRNA.
<a href="#">RGD1309605</a>	<a href="#">RGD1309605.hSep08</a>	<a href="#">291320</a>	16687	826	4	52	putative protein of eukaryotic origin (RGD1309605) alternative variant hSep08, mRNA.
<a href="#">RGD1309621</a>	<a href="#">RGD1309621.aSep08</a>	<a href="#">316982</a>	7582	3366	3	885	similar to hypothetical protein FLJ10652 (RGD1309621) alternative variant aSep08, mRNA.
<a href="#">RGD1309621</a>	<a href="#">RGD1309621.cSep08</a>	<a href="#">316982</a>	5031	671	4	89	similar to hypothetical protein FLJ10652 (10.2 kD) (RGD1309621) alternative variant cSep08, mRNA.
<a href="#">RGD1309634</a>	<a href="#">RGD1309634.aSep08</a>	<a href="#">305452</a>	48977	3584	12	957	hypothetical LOC305452 (RGD1309634) alternative variant aSep08, mRNA.
<a href="#">RGD1309634</a>	<a href="#">RGD1309634.bSep08</a>	<a href="#">305452</a>	50545	1814	10	604	hypothetical LOC305452 (RGD1309634) alternative variant bSep08, mRNA.
<a href="#">RGD1309651</a>	<a href="#">RGD1309651.aSep08</a>	<a href="#">361424</a>	26408	937	4	111	similar to 1190005106Rik protein (12.0 kD) (RGD1309651) alternative variant aSep08, mRNA.
<a href="#">RGD1309676</a>	<a href="#">RGD1309676.aSep08</a>	<a href="#">361118</a>	13291	1008	1	229	similar to RIKEN cDNA 5730469M10 (25.8 kD) (RGD1309676) alternative variant aSep08, mRNA.
<a href="#">RGD1309696</a>	<a href="#">RGD1309696.aSep08</a>	<a href="#">314478</a>	1303	920		306	similar to KIAA2019 protein (RGD1309696) mRNA.
<a href="#">RGD1309701</a>	<a href="#">RGD1309701.aSep08</a>	<a href="#">316729</a>	638	591		196	similar to RIKEN cDNA 9130005N14 (RGD1309701) mRNA.
<a href="#">RGD1309707</a>	<a href="#">RGD1309707.bSep08</a>	<a href="#">690516</a>	2667	734	2	112	similar to RIKEN cDNA 4930431E10 (RGD1309707) alternative variant bSep08, mRNA.
<a href="#">RGD1309710</a>	<a href="#">RGD1309710.bSep08</a>	<a href="#">293700</a>	6219	441	4	112	similar to RIKEN cDNA 0610038D11 (RGD1309710) alternative variant bSep08, mRNA.



<a href="#">RGD1309730</a>	<a href="#">RGD1309730.bSep08</a>	<a href="#">295952</a>	15412	910	2	99	similar to RIKEN cDNA B230118H07 (10.7 kD) (RGD1309730) alternative variant bSep08, mRNA.
<a href="#">RGD1309735</a>	<a href="#">RGD1309735.bSep08</a>	<a href="#">360776</a>	3723	945	1	91	similar to CG14977-PA (9.9 kD) (RGD1309735) alternative variant bSep08, mRNA.
<a href="#">RGD1309744</a>	<a href="#">RGD1309744.bSep08</a>	<a href="#">311405</a>	2008	1119	3	173	similar to hypothetical protein FLJ20507 (19.1 kD) (RGD1309744) alternative variant bSep08, mRNA.
<a href="#">RGD1309744</a>	<a href="#">RGD1309744.cSep08</a>	<a href="#">311405</a>	2326	982	2	136	similar to hypothetical protein FLJ20507 (RGD1309744) alternative variant cSep08, mRNA.
<a href="#">RGD1309747</a>	<a href="#">RGD1309747.aSep08</a>	<a href="#">306665</a>	13766	2449	7	476	similar to KIAA0947 protein (RGD1309747) alternative variant aSep08, mRNA.
<a href="#">RGD1309748</a>	<a href="#">RGD1309748.bSep08</a>	<a href="#">302913</a>	11591	831	2	252	similar to CG4768-PA (RGD1309748) alternative variant bSep08, mRNA.
<a href="#">RGD1309748</a>	<a href="#">RGD1309748.cSep08</a>	<a href="#">302913</a>	25330	884	3	171	similar to CG4768-PA (19.0 kD) (RGD1309748) alternative variant cSep08, mRNA.
<a href="#">RGD1309759</a>	<a href="#">RGD1309759.aSep08</a>	<a href="#">361448</a>	16456	1566	8	339	putative protein of vertebrate origin (RGD1309759) alternative variant aSep08, mRNA.
<a href="#">RGD1309759</a>	<a href="#">RGD1309759.bSep08</a>	<a href="#">361448</a>	3823	707	3	128	putative protein of vertebrate origin (RGD1309759) alternative variant bSep08, mRNA.
<a href="#">RGD1309762</a>	<a href="#">RGD1309762.aSep08</a>	<a href="#">304503</a>	9025	3432	9	486	similar to KIAA0614 protein (RGD1309762) alternative variant aSep08, mRNA.
<a href="#">RGD1309765</a>	<a href="#">RGD1309765.aSep08</a>	<a href="#">290746</a>	15820	1451	1	217	similar to hypothetical protein (RGD1309765) alternative variant aSep08, mRNA.
<a href="#">RGD1309765</a>	<a href="#">RGD1309765.bSep08</a>	<a href="#">290746</a>	15130	750	1	142	similar to hypothetical protein (16.2 kD) (RGD1309765) alternative variant bSep08, mRNA.
<a href="#">RGD1309784</a>	<a href="#">RGD1309784.bSep08</a>	<a href="#">363099</a>	7416	973	5	141	similar to ribosomal protein L24-like; 60S ribosomal protein L30 isolog; my024 protein; homolog of yeast ribosomal like protein 24 (17.1 kD) (RGD1309784) alternative variant bSep08, mRNA.
<a href="#">RGD1309792</a>	<a href="#">RGD1309792.bSep08</a>	<a href="#">312199</a>	46775	759	5	219	similar to scruin like at the midline CG5186-PA (RGD1309792) alternative variant bSep08, mRNA.
<a href="#">RGD1309804</a>	<a href="#">RGD1309804.bSep08</a>	<a href="#">361751</a>	2514	796	4	126	similar to hypothetical protein FLJ11218 (13.8 kD) (RGD1309804) alternative variant bSep08, mRNA.
<a href="#">RGD1309804</a>	<a href="#">RGD1309804.cSep08</a>	<a href="#">361751</a>	71635	542	3	123	similar to hypothetical protein FLJ11218 (RGD1309804) alternative variant cSep08, mRNA.
<a href="#">RGD1309807</a>	<a href="#">RGD1309807.aSep08</a>	<a href="#">362378</a>	62565	750		250	similar to Fam13a1 protein (RGD1309807) alternative variant aSep08, mRNA.
<a href="#">RGD1309823</a>	<a href="#">RGD1309823.aSep08</a>	<a href="#">313525</a>	4579	1166	1	388	similar to hypothetical protein FLJ21156 (RGD1309823) alternative variant aSep08, mRNA.
<a href="#">RGD1309823</a>	<a href="#">RGD1309823.bSep08</a>	<a href="#">313525</a>	2168	697	3	232	similar to hypothetical protein FLJ21156 (RGD1309823) alternative variant bSep08, mRNA.
<a href="#">RGD1309823</a>	<a href="#">RGD1309823.cSep08</a>	<a href="#">313525</a>	2396	670	1	217	similar to hypothetical protein FLJ21156 (RGD1309823) alternative variant cSep08, mRNA.
<a href="#">RGD1309823</a>	<a href="#">RGD1309823.dSep08</a>	<a href="#">313525</a>	1400	1283	2	48	similar to hypothetical protein FLJ21156 (RGD1309823) alternative variant dSep08, mRNA.
<a href="#">RGD1309829</a>	<a href="#">RGD1309829.bSep08</a>	<a href="#">296190</a>	26067	842	9	194	CRA b (RGD1309829) alternative variant bSep08, mRNA.
<a href="#">RGD1309829</a>	<a href="#">RGD1309829.cSep08</a>	<a href="#">296190</a>	14414	749	7	132	CRA b (14.9 kD) (RGD1309829) alternative variant cSep08, mRNA.

<a href="#">RGD1309829</a>	<a href="#">RGD1309829.dSep08</a>	<a href="#">296190</a>	1122	660	3	83	CRA a like (9.4 kD) (RGD1309829) alternative variant dSep08, mRNA.
<a href="#">RGD1309870</a>	<a href="#">RGD1309870.aSep08</a>	<a href="#">289778</a>	31922	924	3	237	hypothetical LOC289778 (27.1 kD) (RGD1309870) alternative variant aSep08, mRNA.
<a href="#">RGD1309871</a>	<a href="#">RGD1309871.bSep08</a>	<a href="#">309187</a>	17724	445	3	148	similar to RIKEN cDNA 5730596K20 (RGD1309871) alternative variant bSep08, mRNA.
<a href="#">RGD1309888</a>	<a href="#">RGD1309888.bSep08</a>	<a href="#">365215</a>	2539	941	5	180	similar to RIKEN cDNA 1500002O20 (RGD1309888) alternative variant bSep08, mRNA.
<a href="#">RGD1309888</a>	<a href="#">RGD1309888.cSep08</a>	<a href="#">365215</a>	5520	408	1	79	similar to RIKEN cDNA 1500002O20 (RGD1309888) alternative variant cSep08, mRNA.
<a href="#">RGD1309906</a>	<a href="#">RGD1309906.aSep08</a>	<a href="#">287406</a>	13114	1796	4	337	similar to RIKEN cDNA 2310004I24 gene (38.7 kD) (RGD1309906) alternative variant aSep08, complete mRNA.
<a href="#">RGD1309922</a>	<a href="#">RGD1309922.aSep08</a>	<a href="#">306007</a>	9852	1590	11	529	similar to 2610301G19Rik protein (RGD1309922) alternative variant aSep08, mRNA.
<a href="#">RGD1309922</a>	<a href="#">RGD1309922.bSep08</a>	<a href="#">306007</a>	8432	1043	7	347	similar to 2610301G19Rik protein (RGD1309922) alternative variant bSep08, mRNA.
<a href="#">RGD1309922</a>	<a href="#">RGD1309922.cSep08</a>	<a href="#">306007</a>	2108	399	5	132	similar to 2610301G19Rik protein (RGD1309922) alternative variant cSep08, mRNA.
<a href="#">RGD1309926</a>	<a href="#">RGD1309926.aSep08</a>	<a href="#">303021</a>	92003	1494	12	315	similar to RIKEN cDNA G431001E03 gene (RGD1309926) alternative variant aSep08, mRNA.
<a href="#">RGD1309926</a>	<a href="#">RGD1309926.bSep08</a>	<a href="#">303021</a>	5975	357	1	85	similar to RIKEN cDNA G431001E03 gene (RGD1309926) alternative variant bSep08, mRNA.
<a href="#">RGD1309926</a>	<a href="#">RGD1309926.cSep08</a>	<a href="#">303021</a>	2273	456	1	41	similar to RIKEN cDNA G431001E03 gene (RGD1309926) alternative variant cSep08, mRNA.
<a href="#">RGD1309930</a>	<a href="#">RGD1309930.aSep08</a>	<a href="#">316426</a>	4554	1006		164	similar to 2810022L02Rik protein (RGD1309930) mRNA.
<a href="#">RGD1309931</a>	<a href="#">RGD1309931.aSep08</a>	<a href="#">362538</a>	27859	1086		361	similar to hypothetical protein (RGD1309931) mRNA.
<a href="#">RGD1309969</a>	<a href="#">RGD1309969.bSep08</a>	<a href="#">362171</a>	24338	410	3	75	similar to RIKEN cDNA 2600010E01 (RGD1309969) alternative variant bSep08, mRNA.
<a href="#">RGD1309985</a>	<a href="#">RGD1309985.aSep08</a>	<a href="#">361931</a>	3837	695	2	231	similar to RIKEN cDNA 4932438A13 (RGD1309985) alternative variant aSep08, mRNA.
<a href="#">RGD1309985</a>	<a href="#">RGD1309985.bSep08</a>	<a href="#">361931</a>	1939	798	1	167	similar to RIKEN cDNA 4932438A13 (RGD1309985) alternative variant bSep08, mRNA.
<a href="#">RGD1309995</a>	<a href="#">RGD1309995.aSep08</a>	<a href="#">314690</a>	14197	2666	9	309	similar to CG13957-PA (RGD1309995) alternative variant aSep08, mRNA.
<a href="#">RGD1309995</a>	<a href="#">RGD1309995.bSep08</a>	<a href="#">314690</a>	18261	1047	3	50	similar to CG13957-PA (RGD1309995) alternative variant bSep08, mRNA.
<a href="#">RGD1310012</a>	<a href="#">RGD1310012.aSep08</a>	<a href="#">313490</a>	5639	939	1	241	similar to NAG-5 protein (RGD1310012) alternative variant aSep08, mRNA.
<a href="#">RGD1310012</a>	<a href="#">RGD1310012.bSep08</a>	<a href="#">313490</a>	12861	753	3	175	similar to NAG-5 protein (RGD1310012) alternative variant bSep08, mRNA.
<a href="#">RGD1310016</a>	<a href="#">RGD1310016.aSep08</a>	<a href="#">309306</a>	2945	1964	3	127	similar to hypothetical protein (RGD1310016) alternative variant aSep08, mRNA.
<a href="#">RGD1310039</a>	<a href="#">RGD1310039.aSep08</a>	<a href="#">361747</a>	46679	1783	5	594	similar to hypothetical protein FLJ10058 (RGD1310039) alternative variant aSep08, mRNA.
<a href="#">RGD1310039</a>	<a href="#">RGD1310039.cSep08</a>	<a href="#">361747</a>	17676	712	7	237	similar to hypothetical protein FLJ10058 (RGD1310039) alternative variant cSep08, mRNA.

<a href="#">RGD1310039</a>	<a href="#">RGD1310039.dSep08</a>	<a href="#">361747</a>	21519	704	9	234	similar to hypothetical protein FLJ10058 (RGD1310039) alternative variant dSep08, mRNA.
<a href="#">RGD1310039</a>	<a href="#">RGD1310039.fSep08</a>	<a href="#">361747</a>	18391	697	7	131	similar to hypothetical protein FLJ10058 (RGD1310039) alternative variant fSep08, mRNA.
<a href="#">RGD1310061</a>	<a href="#">RGD1310061.bSep08</a>	<a href="#">290912</a>	37887	3669	1	227	similar to hypothetical protein FLJ10154 (27.2 kD) (RGD1310061) alternative variant bSep08, mRNA.
<a href="#">RGD1310081</a>	<a href="#">RGD1310081.aSep08</a>	<a href="#">310137</a>	24987	2207	13	645	similar to hypothetical protein FLJ13231 (RGD1310081) alternative variant aSep08, mRNA.
<a href="#">RGD1310110</a>	<a href="#">RGD1310110.bSep08</a>	<a href="#">361032</a>	85715	1780	1	558	similar to 3632451O06Rik protein (RGD1310110) alternative variant bSep08, mRNA.
<a href="#">RGD1310111</a>	<a href="#">RGD1310111.aSep08</a>	<a href="#">292875</a>	2250	1078	6	293	similar to RIKEN cDNA 0610012D14 (RGD1310111) alternative variant aSep08, complete mRNA.
<a href="#">RGD1310127</a>	<a href="#">RGD1310127.bSep08</a>	<a href="#">361654</a>	25876	1568	9	261	putative endoplasmic reticulum protein, with a transmembrane domain, of eukaryotic origin (28.5 kD) (RGD1310127) alternative variant bSep08, mRNA.
<a href="#">RGD1310127</a>	<a href="#">RGD1310127.cSep08</a>	<a href="#">361654</a>	2605	877	4	157	putative protein (17.4 kD) (RGD1310127) alternative variant cSep08, mRNA.
<a href="#">RGD1310127</a>	<a href="#">RGD1310127.dSep08</a>	<a href="#">361654</a>	4284	1281	7	151	putative protein (RGD1310127) alternative variant dSep08, mRNA.
<a href="#">RGD1310127</a>	<a href="#">RGD1310127.eSep08</a>	<a href="#">361654</a>	10307	685	2	126	putative protein of metazoan origin (RGD1310127) alternative variant eSep08, mRNA.
<a href="#">RGD1310127</a>	<a href="#">RGD1310127.fSep08</a>	<a href="#">361654</a>	1314	398	3	125	putative protein of mammalian origin (RGD1310127) alternative variant fSep08, mRNA.
<a href="#">RGD1310133</a>	<a href="#">RGD1310133.bSep08</a>	<a href="#">292760</a>	8086	729	8	243	similar to RIKEN cDNA 5830482F20 gene (RGD1310133) alternative variant bSep08, mRNA.
<a href="#">RGD1310159</a>	<a href="#">RGD1310159.aSep08</a>	<a href="#">304500</a>	11634	1250	8	319	similar to acetyl-coA dehydrogenase -related (111.6 kD) (5G231) (RGD1310159) alternative variant aSep08, mRNA.
<a href="#">RGD1310159</a>	<a href="#">RGD1310159.bSep08</a>	<a href="#">304500</a>	1380	745	3	126	similar to acetyl-coA dehydrogenase -related (111.6 kD) (5G231) (13.5 kD) (RGD1310159) alternative variant bSep08, mRNA.
<a href="#">RGD1310166</a>	<a href="#">RGD1310166.aSep08</a>	<a href="#">303395</a>	8851	568		135	similar to Chromodomain-helicase-DNA-binding protein 1 (CHD-1) (RGD1310166) mRNA.
<a href="#">RGD1310169and dCox6b2</a>	<a href="#">RGD1310169andCox6b2.aSep08</a>	<a href="#">308344</a>	17344	2604	13	356	putative protein of vertebrate origin (40.3 kD) (RGD1310169andCox6b2) alternative variant aSep08, mRNA.
<a href="#">RGD1310169and dCox6b2</a>	<a href="#">RGD1310169andCox6b2.aSep08</a>	<a href="#">654441</a>	17344	2604	13	356	putative protein of vertebrate origin (40.3 kD) (RGD1310169andCox6b2) alternative variant aSep08, mRNA.
<a href="#">RGD1310169and dCox6b2</a>	<a href="#">RGD1310169andCox6b2.cSep08</a>	<a href="#">308344</a>	4925	739	5	148	putative protein of mammalian origin (RGD1310169andCox6b2) alternative variant cSep08, mRNA.
<a href="#">RGD1310169and dCox6b2</a>	<a href="#">RGD1310169andCox6b2.cSep08</a>	<a href="#">654441</a>	4925	739	5	148	putative protein of mammalian origin (RGD1310169andCox6b2) alternative variant cSep08, mRNA.
<a href="#">RGD1310169and dCox6b2</a>	<a href="#">RGD1310169andCox6b2.dSep08</a>	<a href="#">308344</a>	1196	412	4	91	cytochrome c oxidase VIb (RGD1310169andCox6b2) alternative variant dSep08, mRNA.

<a href="#">RGD1310169andCox6b2</a>	<a href="#">RGD1310169andCox6b2.dSep08</a>	<a href="#">654441</a>	1196	412	4	91	cytochrome c oxidase VIb (RGD1310169andCox6b2) alternative variant dSep08, mRNA.
<a href="#">RGD1310185</a>	<a href="#">RGD1310185.aSep08</a>	<a href="#">313790</a>	1502	550		156	similar to RIKEN cDNA 9030625A04 (RGD1310185) alternative variant aSep08, mRNA.
<a href="#">RGD1310185</a>	<a href="#">RGD1310185.cSep08</a>	<a href="#">313790</a>	12320	1795			
<a href="#">RGD1310199</a>	<a href="#">RGD1310199.bSep08</a>	<a href="#">291737</a>	7683	744	3	183	similar to RIKEN cDNA D030070L09 (RGD1310199) alternative variant bSep08, mRNA.
<a href="#">RGD1310199</a>	<a href="#">RGD1310199.cSep08</a>	<a href="#">291737</a>	13628	1214	2	105	similar to RIKEN cDNA D030070L09 (RGD1310199) alternative variant cSep08, mRNA.
<a href="#">RGD1310199</a>	<a href="#">RGD1310199.dSep08</a>	<a href="#">291737</a>	4839	551	2	51	similar to RIKEN cDNA D030070L09 (5.7 kD) (RGD1310199) alternative variant dSep08, mRNA.
<a href="#">RGD1310205</a>	<a href="#">RGD1310205.aSep08</a>	<a href="#">366890</a>	9197	727		137	similar to RIKEN cDNA 4921508O11 (RGD1310205) mRNA.
<a href="#">RGD1310209</a>	<a href="#">RGD1310209.aSep08</a>	<a href="#">362019</a>	78532	3741	22	1013	similar to KIAA1324 protein (111.1 kD) (RGD1310209) alternative variant aSep08, mRNA.
<a href="#">RGD1310209</a>	<a href="#">RGD1310209.bSep08</a>	<a href="#">362019</a>	53433	673	6	224	similar to KIAA1324 protein (RGD1310209) alternative variant bSep08, mRNA.
<a href="#">RGD1310209</a>	<a href="#">RGD1310209.cSep08</a>	<a href="#">362019</a>	736	658	2	131	similar to KIAA1324 protein (RGD1310209) alternative variant cSep08, mRNA.
<a href="#">RGD1310209</a>	<a href="#">RGD1310209.dSep08</a>	<a href="#">362019</a>	39789	351	2	111	similar to KIAA1324 protein (RGD1310209) alternative variant dSep08, mRNA.
<a href="#">RGD1310209</a>	<a href="#">RGD1310209.eSep08</a>	<a href="#">362019</a>	776	243	2	80	similar to KIAA1324 protein (RGD1310209) alternative variant eSep08, mRNA.
<a href="#">RGD1310212</a>	<a href="#">RGD1310212.aSep08</a>	<a href="#">299557</a>	1588	326	1	68	similar to KIAA1111-like protein (RGD1310212) alternative variant aSep08, mRNA.
<a href="#">RGD1310212</a>	<a href="#">RGD1310212.bSep08</a>	<a href="#">299557</a>	5525	329	1	67	similar to KIAA1111-like protein (RGD1310212) alternative variant bSep08, mRNA.
<a href="#">RGD1310224</a>	<a href="#">RGD1310224.bSep08</a>	<a href="#">291076</a>	14866	715	7	181	similar to RIKEN cDNA 1810022C23 (RGD1310224) alternative variant bSep08, mRNA.
<a href="#">RGD1310224</a>	<a href="#">RGD1310224.cSep08</a>	<a href="#">291076</a>	1594	598	2	147	similar to RIKEN cDNA 1810022C23 (RGD1310224) alternative variant cSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.aSep08</a>	<a href="#">301563</a>	33413	3297	10	399	similar to RIKEN cDNA 5230400G24 (44.1 kD) (RGD1310230) alternative variant aSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.bSep08</a>	<a href="#">301563</a>	27482	1045	8	325	similar to RIKEN cDNA 5230400G24 (RGD1310230) alternative variant bSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.cSep08</a>	<a href="#">301563</a>	23986	1649	6	271	similar to RIKEN cDNA 5230400G24 (30.8 kD) (RGD1310230) alternative variant cSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.dSep08</a>	<a href="#">301563</a>	27549	1168	8	244	similar to RIKEN cDNA 5230400G24 (27.5 kD) (RGD1310230) alternative variant dSep08, complete mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.eSep08</a>	<a href="#">301563</a>	17603	926	7	222	similar to RIKEN cDNA 5230400G24 (RGD1310230) alternative variant eSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.fSep08</a>	<a href="#">301563</a>	28120	1303	6	221	similar to RIKEN cDNA 5230400G24 (RGD1310230) alternative variant fSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.hSep08</a>	<a href="#">301563</a>	25096	514	6	74	similar to RIKEN cDNA 5230400G24 (RGD1310230) alternative variant hSep08, mRNA.

<a href="#">RGD1310230</a>	<a href="#">RGD1310230.iSep08</a>	<a href="#">301563</a>	2457	1529	2	55	similar to RIKEN cDNA 5230400G24 (6.8 kD) (RGD1310230) alternative variant iSep08, mRNA.
<a href="#">RGD1310230</a>	<a href="#">RGD1310230.jSep08</a>	<a href="#">301563</a>	5307	841	3	55	similar to RIKEN cDNA 5230400G24 (6.8 kD) (RGD1310230) alternative variant jSep08, mRNA.
<a href="#">RGD1310262</a>	<a href="#">RGD1310262.bSep08</a>	<a href="#">304650</a>	1199	1068	2	200	hypothetical LOC304650 (RGD1310262) alternative variant bSep08, mRNA.
<a href="#">RGD1310262</a>	<a href="#">RGD1310262.cSep08</a>	<a href="#">304650</a>	1029	813	2	37	hypothetical LOC304650 (RGD1310262) alternative variant cSep08, mRNA.
<a href="#">RGD1310270</a>	<a href="#">RGD1310270.bSep08</a>	<a href="#">314798</a>	10554	766	4	211	similar to CG33154-PB (RGD1310270) alternative variant bSep08, mRNA.
<a href="#">RGD1310270</a>	<a href="#">RGD1310270.cSep08</a>	<a href="#">314798</a>	37778	619	6	164	similar to CG33154-PB (RGD1310270) alternative variant cSep08, mRNA.
<a href="#">RGD1310270</a>	<a href="#">RGD1310270.dSep08</a>	<a href="#">314798</a>	10171	611	4	87	similar to CG33154-PB (RGD1310270) alternative variant dSep08, mRNA.
<a href="#">RGD1310271</a>	<a href="#">RGD1310271.aSep08</a>	<a href="#">313778</a>	5776	910	7	277	similar to hypothetical protein MGC45873 (RGD1310271) alternative variant aSep08, mRNA.
<a href="#">RGD1310311</a>	<a href="#">RGD1310311.bSep08</a>	<a href="#">362746</a>	3472	720	1	210	uncharacterized protein homolog (RGD1310311) alternative variant bSep08, mRNA.
<a href="#">RGD1310324</a>	<a href="#">RGD1310324.aSep08</a>	<a href="#">287388</a>	6215	607		109	hypothetical LOC287388 (RGD1310324) mRNA.
<a href="#">RGD1310335</a>	<a href="#">RGD1310335.aSep08</a>	<a href="#">360711</a>	4959	716		237	similar to RIKEN cDNA C330027C09 (RGD1310335) mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.aSep08</a>	<a href="#">360738</a>	42843	1064	7	312	similar to Ser/Thr-rich protein T10 in DGCR region (RGD1310348) alternative variant aSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.bSep08</a>	<a href="#">360738</a>	43075	1080	7	280	similar to Ser/Thr-rich protein T10 in DGCR region (31.2 kD) (RGD1310348) alternative variant bSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.dSep08</a>	<a href="#">360738</a>	36800	756	4	171	similar to Ser/Thr-rich protein T10 in DGCR region (19.0 kD) (RGD1310348) alternative variant dSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.eSep08</a>	<a href="#">360738</a>	40702	1281	4	171	similar to Ser/Thr-rich protein T10 in DGCR region (19.0 kD) (RGD1310348) alternative variant eSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.fSep08</a>	<a href="#">360738</a>	43661	752	5	169	similar to Ser/Thr-rich protein T10 in DGCR region (RGD1310348) alternative variant fSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.gSep08</a>	<a href="#">360738</a>	40112	583	4	132	similar to Ser/Thr-rich protein T10 in DGCR region (RGD1310348) alternative variant gSep08, mRNA.
<a href="#">RGD1310348</a>	<a href="#">RGD1310348.hSep08</a>	<a href="#">360738</a>	10971	477	2	84	similar to Ser/Thr-rich protein T10 in DGCR region (RGD1310348) alternative variant hSep08, mRNA.
<a href="#">RGD1310351</a>	<a href="#">RGD1310351.bSep08</a>	<a href="#">298425</a>	13371	605	6	178	similar to RIKEN cDNA 4732418C07 (RGD1310351) alternative variant bSep08, mRNA.
<a href="#">RGD1310352</a>	<a href="#">RGD1310352.bSep08</a>	<a href="#">303122</a>	4643	1419	2	169	similar to HTGN29 protein; keratinocytes associated transmembrane protein 2 (RGD1310352) alternative variant bSep08, mRNA.
<a href="#">RGD1310352</a>	<a href="#">RGD1310352.cSep08</a>	<a href="#">303122</a>	967	607	2	65	similar to HTGN29 protein; keratinocytes associated transmembrane protein 2 (7.0 kD) (RGD1310352) alternative variant cSep08, mRNA.
<a href="#">RGD1310358</a>	<a href="#">RGD1310358.bSep08</a>	<a href="#">308537</a>	37742	481		102	similar to NNX3 (RGD1310358) alternative variant bSep08, mRNA.
<a href="#">RGD1310364</a>	<a href="#">RGD1310364.aSep08</a>	<a href="#">313111</a>	185993	945		130	similar to KIAA1900 protein (14.3 kD) (RGD1310364) mRNA.

<a href="#">RGD1310371</a>	<a href="#">RGD1310371.bSep08</a>	<a href="#">308794</a>	1216	692	2	140	similar to RIKEN cDNA 1700026D08 (RGD1310371) alternative variant bSep08, mRNA.
<a href="#">RGD1310371</a>	<a href="#">RGD1310371.dSep08</a>	<a href="#">308794</a>	4629	1718	3	76	similar to RIKEN cDNA 1700026D08 (RGD1310371) alternative variant dSep08, mRNA.
<a href="#">RGD1310376</a>	<a href="#">RGD1310376.aSep08</a>	<a href="#">298404</a>	3804	1854	9	344	vav-like protein (39.4 kD) (RGD1310376) alternative variant aSep08, mRNA.
<a href="#">RGD1310376</a>	<a href="#">RGD1310376.cSep08</a>	<a href="#">298404</a>	1682	700	4	125	vav-like protein (RGD1310376) alternative variant cSep08, mRNA.
<a href="#">RGD1310376</a>	<a href="#">RGD1310376.eSep08</a>	<a href="#">298404</a>	1019	808	2	84	novel protein containing RhoGEF PH domains (RGD1310376) alternative variant eSep08, mRNA.
<a href="#">RGD1310384</a>	<a href="#">RGD1310384.aSep08</a>	<a href="#">289530</a>	15670	1162		322	hypothetical LOC289530 (34.6 kD) (RGD1310384) mRNA.
<a href="#">RGD1310393</a>	<a href="#">RGD1310393.aSep08</a>	<a href="#">298197</a>	1781	408	2	100	similar to hypothetical protein 5930437A14 (RGD1310393) alternative variant aSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.aSep08</a>	<a href="#">308762</a>	15933	3427	15	854	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant aSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.aSep08</a>	<a href="#">499191</a>	15933	3427	15	854	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant aSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.aSep08</a>	<a href="#">685955</a>	15933	3427	15	854	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant aSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.cSep08</a>	<a href="#">308762</a>	5839	1702	2	230	neugrin (26.2 kD) (RGD1310399_predictedandNgrn) alternative variant cSep08, complete mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.cSep08</a>	<a href="#">499191</a>	5839	1702	2	230	neugrin (26.2 kD) (RGD1310399_predictedandNgrn) alternative variant cSep08, complete mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.cSep08</a>	<a href="#">685955</a>	5839	1702	2	230	neugrin (26.2 kD) (RGD1310399_predictedandNgrn) alternative variant cSep08, complete mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.dSep08</a>	<a href="#">308762</a>	12671	716	5	188	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant dSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.dSep08</a>	<a href="#">499191</a>	12671	716	5	188	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant dSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.dSep08</a>	<a href="#">685955</a>	12671	716	5	188	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant dSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.eSep08</a>	<a href="#">308762</a>	1299	666	2	73	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant eSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.eSep08</a>	<a href="#">499191</a>	1299	666	2	73	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant eSep08, mRNA.

<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.eSep08</a>	<a href="#">685955</a>	1299	666	2	73	tubulin tyrosine ligase-like family member 13 (RGD1310399_predictedandNgrn) alternative variant eSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.fSep08</a>	<a href="#">308762</a>	5088	1335	2	65	putative protein (7.0 kD) (RGD1310399_predictedandNgrn) alternative variant fSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.fSep08</a>	<a href="#">499191</a>	5088	1335	2	65	putative protein (7.0 kD) (RGD1310399_predictedandNgrn) alternative variant fSep08, mRNA.
<a href="#">RGD1310399_predictedandNgrn</a>	<a href="#">RGD1310399_predictedandNgrn.fSep08</a>	<a href="#">685955</a>	5088	1335	2	65	putative protein (7.0 kD) (RGD1310399_predictedandNgrn) alternative variant fSep08, mRNA.
<a href="#">RGD1310414</a>	<a href="#">RGD1310414.bSep08</a>	<a href="#">290811</a>	906	445	3	148	similar to hypothetical protein FLJ23263 (RGD1310414) alternative variant bSep08, mRNA.
<a href="#">RGD1310423</a>	<a href="#">RGD1310423.aSep08</a>	<a href="#">361112</a>	24203	1455		98	similar to hypothetical protein FLJ31737 (RGD1310423) mRNA.
<a href="#">RGD1310425</a>	<a href="#">RGD1310425.bSep08</a>	<a href="#">298398</a>	1117	756	2	20	hypothetical LOC298398 (2.1 kD) (RGD1310425) alternative variant bSep08, mRNA.
<a href="#">RGD1310429</a>	<a href="#">RGD1310429.aSep08</a>	<a href="#">303764</a>	78857	2806		479	similar to Protein Njmu-R1 (53.5 kD) (RGD1310429) mRNA.
<a href="#">RGD1310440</a>	<a href="#">RGD1310440.aSep08</a>	<a href="#">362594</a>	24088	2109	8	205	similar to hypothetical protein (RGD1310440) alternative variant aSep08, mRNA.
<a href="#">RGD1310440</a>	<a href="#">RGD1310440.bSep08</a>	<a href="#">362594</a>	23164	1155	7	193	similar to hypothetical protein (RGD1310440) alternative variant bSep08, mRNA.
<a href="#">RGD1310440</a>	<a href="#">RGD1310440.cSep08</a>	<a href="#">362594</a>	22999	1077	9	190	similar to hypothetical protein (RGD1310440) alternative variant cSep08, mRNA.
<a href="#">RGD1310440</a>	<a href="#">RGD1310440.dSep08</a>	<a href="#">362594</a>	11479	372	3	124	similar to hypothetical protein (RGD1310440) alternative variant dSep08, mRNA.
<a href="#">RGD1310440</a>	<a href="#">RGD1310440.eSep08</a>	<a href="#">362594</a>	6925	467	2	64	similar to hypothetical protein (RGD1310440) alternative variant eSep08, mRNA.
<a href="#">RGD1310444</a>	<a href="#">RGD1310444.aSep08</a>	<a href="#">363015</a>	25900	808		269	LOC363015 (RGD1310444) mRNA.
<a href="#">RGD1310474</a>	<a href="#">RGD1310474.aSep08</a>	<a href="#">314169</a>	32678	3268	3	593	similar to KIAA0423 (66.4 kD) (RGD1310474) alternative variant aSep08, mRNA.
<a href="#">RGD1310474</a>	<a href="#">RGD1310474.bSep08</a>	<a href="#">314169</a>	32730	1794	8	549	similar to KIAA0423 (RGD1310474) alternative variant bSep08, mRNA.
<a href="#">RGD1310474</a>	<a href="#">RGD1310474.cSep08</a>	<a href="#">314169</a>	20895	1160	5	386	similar to KIAA0423 (RGD1310474) alternative variant cSep08, mRNA.
<a href="#">RGD1310474</a>	<a href="#">RGD1310474.dSep08</a>	<a href="#">314169</a>	12165	770	1	208	similar to KIAA0423 (RGD1310474) alternative variant dSep08, mRNA.
<a href="#">RGD1310474</a>	<a href="#">RGD1310474.eSep08</a>	<a href="#">314169</a>	8237	740	1	91	similar to KIAA0423 (RGD1310474) alternative variant eSep08, mRNA.
<a href="#">RGD1310475</a>	<a href="#">RGD1310475.bSep08</a>	<a href="#">293949</a>	11751	1509	5	188	similar to RIKEN cDNA 0610010D20 (RGD1310475) alternative variant bSep08, mRNA.
<a href="#">RGD1310475</a>	<a href="#">RGD1310475.cSep08</a>	<a href="#">293949</a>	6945	275	3	91	similar to RIKEN cDNA 0610010D20 (RGD1310475) alternative variant cSep08, mRNA.
<a href="#">RGD1310484</a>	<a href="#">RGD1310484.bSep08</a>	<a href="#">291694</a>	11606	550	6	183	similar to hypothetical protein MGC37079 (RGD1310484) alternative variant bSep08, mRNA.

<a href="#">RGD1310484</a>	<a href="#">RGD1310484.cSep08</a>	<a href="#">291694</a>	6641	684	5	148	similar to hypothetical protein MGC37079 (RGD1310484) alternative variant cSep08, mRNA.
<a href="#">RGD1310484</a>	<a href="#">RGD1310484.dSep08</a>	<a href="#">291694</a>	1614	388	3	73	similar to hypothetical protein MGC37079 (RGD1310484) alternative variant dSep08, mRNA.
<a href="#">RGD1310495</a>	<a href="#">RGD1310495.aSep08</a>	<a href="#">309809</a>	12995	2508		473	similar to KIAA1919 protein (RGD1310495) mRNA.
<a href="#">RGD1310507</a>	<a href="#">RGD1310507.bSep08</a>	<a href="#">315963</a>	38721	2350	17	683	similar to RIKEN cDNA 1300017J02 (74.8 kD) (RGD1310507) alternative variant bSep08, mRNA.
<a href="#">RGD1310507</a>	<a href="#">RGD1310507.cSep08</a>	<a href="#">315963</a>	22676	514	4	69	similar to RIKEN cDNA 1300017J02 (RGD1310507) alternative variant cSep08, mRNA.
<a href="#">RGD1310552</a>	<a href="#">RGD1310552.bSep08</a>	<a href="#">300836</a>	6714	761	3	105	putative protein of eukaryotic origin (12.4 kD) (RGD1310552) alternative variant bSep08, mRNA.
<a href="#">RGD1310552</a>	<a href="#">RGD1310552.cSep08</a>	<a href="#">300836</a>	915	417	3	93	putative protein (10.6 kD) (RGD1310552) alternative variant cSep08, mRNA.
<a href="#">RGD1310572</a>	<a href="#">RGD1310572.bSep08</a>	<a href="#">304646</a>	19980	629	4	106	similar to calmegin (RGD1310572) alternative variant bSep08, mRNA.
<a href="#">RGD1310587</a>	<a href="#">RGD1310587.aSep08</a>	<a href="#">360894</a>	9732	2695		142	similar to hypothetical protein FLJ14146 (RGD1310587) mRNA.
<a href="#">RGD1310592</a>	<a href="#">RGD1310592.aSep08</a>	<a href="#">297823</a>	40672	2436	21	780	integrator complex (RGD1310592) alternative variant aSep08, mRNA.
<a href="#">RGD1310592</a>	<a href="#">RGD1310592.bSep08</a>	<a href="#">297823</a>	7898	693	7	148	integrator complex (RGD1310592) alternative variant bSep08, mRNA.
<a href="#">RGD1310592</a>	<a href="#">RGD1310592.cSep08</a>	<a href="#">297823</a>	47403	479	4	131	integrator complex (RGD1310592) alternative variant cSep08, mRNA.
<a href="#">RGD1310592</a>	<a href="#">RGD1310592.dSep08</a>	<a href="#">297823</a>	8695	372	3	66	integrator complex (RGD1310592) alternative variant dSep08, mRNA.
<a href="#">RGD1310592</a>	<a href="#">RGD1310592.eSep08</a>	<a href="#">297823</a>	9522	544	5	65	integrator complex (RGD1310592) alternative variant eSep08, mRNA.
<a href="#">RGD1310597</a>	<a href="#">RGD1310597.bSep08</a>	<a href="#">294667</a>	2373	723	2	48	similar to RIKEN cDNA 1200014M14 (5.4 kD) (RGD1310597) alternative variant bSep08, complete mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.aSep08</a>	<a href="#">297728</a>	32008	2918	14	377	similar to RIKEN cDNA 1200009B18; EST AA408438 (42.5 kD) (RGD1310606) alternative variant aSep08, mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.bSep08</a>	<a href="#">297728</a>	30460	1142	12	301	similar to RIKEN cDNA 1200009B18; EST AA408438 (33.8 kD) (RGD1310606) alternative variant bSep08, complete mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.cSep08</a>	<a href="#">297728</a>	14351	765	8	228	similar to RIKEN cDNA 1200009B18; EST AA408438 (RGD1310606) alternative variant cSep08, mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.eSep08</a>	<a href="#">297728</a>	28909	830	11	152	similar to RIKEN cDNA 1200009B18; EST AA408438 (RGD1310606) alternative variant eSep08, mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.fSep08</a>	<a href="#">297728</a>	9503	398	4	103	similar to RIKEN cDNA 1200009B18; EST AA408438 (RGD1310606) alternative variant fSep08, mRNA.
<a href="#">RGD1310606</a>	<a href="#">RGD1310606.gSep08</a>	<a href="#">297728</a>	467	337	2	69	similar to RIKEN cDNA 1200009B18; EST AA408438 (RGD1310606) alternative variant gSep08, mRNA.
<a href="#">RGD1310641</a>	<a href="#">RGD1310641.bSep08</a>	<a href="#">366347</a>	14432	813	1	244	similar to hypothetical protein (RGD1310641) alternative variant bSep08, mRNA.
<a href="#">RGD1310651</a>	<a href="#">RGD1310651.aSep08</a>	<a href="#">362297</a>	21073	688		228	similar to hypothetical protein MGC20460 (RGD1310651) mRNA.



<a href="#">RGD1310656</a>	<a href="#">RGD1310656.aSep08</a>	<a href="#">288664</a>	3754	750		170	similar to hypothetical protein FLJ32356 (RGD1310656) mRNA.
<a href="#">RGD1310660</a>	<a href="#">RGD1310660.bSep08</a>	<a href="#">296470</a>	1697	652	3	115	similar to RIKEN cDNA 2700038C09 (12.1 kD) (RGD1310660) alternative variant bSep08, mRNA.
<a href="#">RGD1310660</a>	<a href="#">RGD1310660.cSep08</a>	<a href="#">296470</a>	1239	738	3	115	similar to RIKEN cDNA 2700038C09 (12.1 kD) (RGD1310660) alternative variant cSep08, mRNA.
<a href="#">RGD1310660</a>	<a href="#">RGD1310660.dSep08</a>	<a href="#">296470</a>	833	602	2	67	similar to RIKEN cDNA 2700038C09 (RGD1310660) alternative variant dSep08, mRNA.
<a href="#">RGD1310660</a>	<a href="#">RGD1310660.eSep08</a>	<a href="#">296470</a>	1206	777	2	48	similar to RIKEN cDNA 2700038C09 (5.1 kD) (RGD1310660) alternative variant eSep08, complete mRNA.
<a href="#">RGD1310686</a>	<a href="#">RGD1310686.aSep08</a>	<a href="#">360480</a>	21077	1191	2	240	putative protein of vertebrate origin (RGD1310686) alternative variant aSep08, complete mRNA.
<a href="#">RGD1310686</a>	<a href="#">RGD1310686.cSep08</a>	<a href="#">360480</a>	20659	450	1	98	putative protein of mammalian origin (RGD1310686) alternative variant cSep08, mRNA.
<a href="#">RGD1310693</a>	<a href="#">RGD1310693.aSep08</a>	<a href="#">316238</a>	13497	904	5	279	similar to RIKEN cDNA 1700027N10 (RGD1310693) alternative variant aSep08, mRNA.
<a href="#">RGD1310693</a>	<a href="#">RGD1310693.bSep08</a>	<a href="#">316238</a>	4860	605	2	94	similar to RIKEN cDNA 1700027N10 (RGD1310693) alternative variant bSep08, mRNA.
<a href="#">RGD1310693</a>	<a href="#">RGD1310693.dSep08</a>	<a href="#">316238</a>	11206	329	4	86	similar to RIKEN cDNA 1700027N10 (RGD1310693) alternative variant dSep08, mRNA.
<a href="#">RGD1310712</a>	<a href="#">RGD1310712.aSep08</a>	<a href="#">361607</a>	6226	1568	4	449	similar to EMSY protein (RGD1310712) alternative variant aSep08, mRNA.
<a href="#">RGD1310712</a>	<a href="#">RGD1310712.bSep08</a>	<a href="#">361607</a>	6932	1794	3	261	similar to EMSY protein (RGD1310712) alternative variant bSep08, mRNA.
<a href="#">RGD1310722</a>	<a href="#">RGD1310722.bSep08</a>	<a href="#">312248</a>	2804	383	4	127	similar to RIKEN cDNA D130059P03 gene (RGD1310722) alternative variant bSep08, mRNA.
<a href="#">RGD1310727</a>	<a href="#">RGD1310727.aSep08</a>	<a href="#">363070</a>	558	369	1	122	LOC363070 (RGD1310727) alternative variant aSep08, mRNA.
<a href="#">RGD1310769</a>	<a href="#">RGD1310769.aSep08</a>	<a href="#">299207</a>	10329	739	2	140	similar to HSPC288 (15.8 kD) (RGD1310769) alternative variant aSep08, mRNA.
<a href="#">RGD1310769</a>	<a href="#">RGD1310769.cSep08</a>	<a href="#">299207</a>	9878	390	1	129	similar to HSPC288 (RGD1310769) alternative variant cSep08, mRNA.
<a href="#">RGD1310784</a>	<a href="#">RGD1310784.aSep08</a>	<a href="#">291441</a>	9890	1089	7	278	similar to RIKEN cDNA 2810433K01 (32.2 kD) (RGD1310784) alternative variant aSep08, mRNA.
<a href="#">RGD1310784</a>	<a href="#">RGD1310784.bSep08</a>	<a href="#">291441</a>	10222	1812	6	257	similar to RIKEN cDNA 2810433K01 (29.2 kD) (RGD1310784) alternative variant bSep08, mRNA.
<a href="#">RGD1310784</a>	<a href="#">RGD1310784.dSep08</a>	<a href="#">291441</a>	7470	774	5	149	similar to RIKEN cDNA 2810433K01 (17.0 kD) (RGD1310784) alternative variant dSep08, mRNA.
<a href="#">RGD1310794</a>	<a href="#">RGD1310794.bSep08</a>	<a href="#">296840</a>	4254	805		267	similar to RIKEN cDNA C030048B08 (RGD1310794) alternative variant bSep08, mRNA.
<a href="#">RGD1310799</a>	<a href="#">RGD1310799.bSep08</a>	<a href="#">309060</a>	82567	831	5	69	similar to RIKEN cDNA A930008G19 (RGD1310799) alternative variant bSep08, mRNA.
<a href="#">RGD1310810</a>	<a href="#">RGD1310810.bSep08</a>	<a href="#">306549</a>	2080	1015	4	90	similar to RIKEN cDNA 4930444A02 (9.8 kD) (RGD1310810) alternative variant bSep08, mRNA.
<a href="#">RGD1310819</a>	<a href="#">RGD1310819.aSep08</a>	<a href="#">301351</a>	2746	997		126	similar to putative protein (5S487) (RGD1310819) mRNA.
<a href="#">RGD1310845</a>	<a href="#">RGD1310845.aSep08</a>	<a href="#">313928</a>	2711	519		172	similar to KIAA0953 protein (RGD1310845) mRNA.

<a href="#">RGD1310852</a>	<a href="#">RGD1310852.bSep08</a>	<a href="#">314992</a>	5141	860	4	153	similar to RIKEN cDNA 9130401M01 (RGD1310852) alternative variant bSep08, mRNA.
<a href="#">RGD1310861</a>	<a href="#">RGD1310861.aSep08</a>	<a href="#">288667</a>	9250	1796	1	362	similar to RIKEN cDNA 1500011H22 (RGD1310861) alternative variant aSep08, complete mRNA.
<a href="#">RGD1310861</a>	<a href="#">RGD1310861.cSep08</a>	<a href="#">288667</a>	6772	1122	1	221	similar to RIKEN cDNA 1500011H22 (25.2 kD) (RGD1310861) alternative variant cSep08, mRNA.
<a href="#">RGD1310862</a>	<a href="#">RGD1310862.aSep08</a>	<a href="#">303016</a>	5071	403		99	similar to adult retina protein (RGD1310862) mRNA.
<a href="#">RGD1310868</a>	<a href="#">RGD1310868.bSep08</a>	<a href="#">303702</a>	3752	1488	2	227	similar to RIKEN cDNA D230014K01 (RGD1310868) alternative variant bSep08, mRNA.
<a href="#">RGD1310877</a>	<a href="#">RGD1310877.aSep08</a>	<a href="#">305667</a>	5554	670		156	similar to RIKEN cDNA 1810063B07 gene (RGD1310877) mRNA.
<a href="#">RGD1310893</a>	<a href="#">RGD1310893.bSep08</a>	<a href="#">362514</a>	14319	1165	6	300	similar to hypothetical protein 3010020C06 (33.9 kD) (RGD1310893) alternative variant bSep08, mRNA.
<a href="#">RGD1310893</a>	<a href="#">RGD1310893.cSep08</a>	<a href="#">362514</a>	14645	1657	5	195	similar to hypothetical protein 3010020C06 (22.2 kD) (RGD1310893) alternative variant cSep08, complete mRNA.
<a href="#">RGD1310899</a>	<a href="#">RGD1310899.bSep08</a>	<a href="#">299198</a>	1448	620	2	64	similar to CGI-35 protein (7.6 kD) (RGD1310899) alternative variant bSep08, mRNA.
<a href="#">RGD1310899</a>	<a href="#">RGD1310899.cSep08</a>	<a href="#">299198</a>	9391	1799	5	8	similar to CGI-35 protein (RGD1310899) alternative variant cSep08, mRNA.
<a href="#">RGD1310922</a>	<a href="#">RGD1310922.aSep08</a>	<a href="#">287170</a>	1760	819	3	175	putative protein of metazoan origin (RGD1310922) alternative variant aSep08, mRNA.
<a href="#">RGD1310922</a>	<a href="#">RGD1310922.bSep08</a>	<a href="#">287170</a>	3300	749	5	115	CRA a like (12.7 kD) (RGD1310922) alternative variant bSep08, mRNA.
<a href="#">RGD1310922</a>	<a href="#">RGD1310922.cSep08</a>	<a href="#">287170</a>	3152	495	4	91	u11 U12 protein snRNP (RGD1310922) alternative variant cSep08, mRNA.
<a href="#">RGD1310935</a>	<a href="#">RGD1310935.aSep08</a>	<a href="#">360707</a>	8806	687	2	156	similar to Dermal papilla derived protein 7 (RGD1310935) alternative variant aSep08, mRNA.
<a href="#">RGD1310935</a>	<a href="#">RGD1310935.bSep08</a>	<a href="#">360707</a>	7374	562	1	91	similar to Dermal papilla derived protein 7 (RGD1310935) alternative variant bSep08, mRNA.
<a href="#">RGD1310942</a>	<a href="#">RGD1310942.aSep08</a>	<a href="#">292767</a>	12012	1758	13	585	similar to R27328 1 (RGD1310942) alternative variant aSep08, mRNA.
<a href="#">RGD1310942</a>	<a href="#">RGD1310942.bSep08</a>	<a href="#">292767</a>	5383	676	6	224	similar to R27328 1 (RGD1310942) alternative variant bSep08, mRNA.
<a href="#">RGD1310942</a>	<a href="#">RGD1310942.cSep08</a>	<a href="#">292767</a>	1482	353	1	75	similar to R27328 1 (RGD1310942) alternative variant cSep08, mRNA.
<a href="#">RGD1310945</a>	<a href="#">RGD1310945.aSep08</a>	<a href="#">292266</a>	34782	492	1	164	similar to hypothetical protein FLJ23305 (RGD1310945) alternative variant aSep08, mRNA.
<a href="#">RGD1310945</a>	<a href="#">RGD1310945.bSep08</a>	<a href="#">292266</a>	3479	564	1	129	similar to hypothetical protein FLJ23305 (RGD1310945) alternative variant bSep08, mRNA.
<a href="#">RGD1310950</a>	<a href="#">RGD1310950.bSep08</a>	<a href="#">289400</a>	5742	595	5	37	similar to KIAA1078 protein (4.3 kD) (RGD1310950) alternative variant bSep08, mRNA.
<a href="#">RGD1310958</a>	<a href="#">RGD1310958.aSep08</a>	<a href="#">305307</a>	24301	1196		278	similar to RIKEN cDNA C130090K23 (RGD1310958) mRNA.
<a href="#">RGD1311019</a>	<a href="#">RGD1311019.bSep08</a>	<a href="#">298694</a>	1493	1014	4	158	similar to hypothetical protein DKFzP434H2010 (RGD1311019) alternative variant bSep08, mRNA.

<a href="#">RGD1311019</a>	<a href="#">RGD1311019.cSep08</a>	<a href="#">298694</a>	847	669	3	90	similar to hypothetical protein DKFZp434H2010 (RGD1311019) alternative variant cSep08, mRNA.
<a href="#">RGD1311019</a>	<a href="#">RGD1311019.dSep08</a>	<a href="#">298694</a>	490	392	2	31	similar to hypothetical protein DKFZp434H2010 (3.5 kD) (RGD1311019) alternative variant dSep08, mRNA.
<a href="#">RGD1311021</a>	<a href="#">RGD1311021.bSep08</a>	<a href="#">308765</a>	28375	1017	3	102	hypothetical LOC308765 (RGD1311021) alternative variant bSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.aSep08</a>	<a href="#">360947</a>	31600	1783	5	463	similar to RIKEN cDNA 4933428G09 (RGD1311045) alternative variant aSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.bSep08</a>	<a href="#">360947</a>	10812	1055	7	284	similar to RIKEN cDNA 4933428G09 (RGD1311045) alternative variant bSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.cSep08</a>	<a href="#">360947</a>	31130	847	6	241	similar to RIKEN cDNA 4933428G09 (RGD1311045) alternative variant cSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.dSep08</a>	<a href="#">360947</a>	25663	1102	5	144	similar to RIKEN cDNA 4933428G09 (RGD1311045) alternative variant dSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.eSep08</a>	<a href="#">360947</a>	31129	914	7	115	similar to RIKEN cDNA 4933428G09 (15.9 kD) (RGD1311045) alternative variant eSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.fSep08</a>	<a href="#">360947</a>	8983	696	5	92	similar to RIKEN cDNA 4933428G09 (9.9 kD) (RGD1311045) alternative variant fSep08, mRNA.
<a href="#">RGD1311045</a>	<a href="#">RGD1311045.gSep08</a>	<a href="#">360947</a>	4246	376	3	72	similar to RIKEN cDNA 4933428G09 (RGD1311045) alternative variant gSep08, mRNA.
<a href="#">RGD1311066</a>	<a href="#">RGD1311066.bSep08</a>	<a href="#">296312</a>	22587	3620	4	384	similar to RIKEN cDNA 0610011L14 gene (43.5 kD) (RGD1311066) alternative variant bSep08, mRNA.
<a href="#">RGD1311066</a>	<a href="#">RGD1311066.cSep08</a>	<a href="#">296312</a>	20435	666	3	97	similar to RIKEN cDNA 0610011L14 gene (11.0 kD) (RGD1311066) alternative variant cSep08, complete mRNA.
<a href="#">RGD1311077</a>	<a href="#">RGD1311077.bSep08</a>	<a href="#">367021</a>	11477	1078		166	similar to RIKEN cDNA 2810485I05 (RGD1311077) alternative variant bSep08, mRNA.
<a href="#">RGD1311078</a>	<a href="#">RGD1311078.aSep08</a>	<a href="#">360664</a>	11175	564		154	LOC360664 (RGD1311078) mRNA.
<a href="#">RGD1311080</a>	<a href="#">RGD1311080.aSep08</a>	<a href="#">312401</a>	37242	3321		486	similar to RIKEN cDNA A930038C07 (RGD1311080) mRNA.
<a href="#">RGD1311084</a>	<a href="#">RGD1311084.bSep08</a>	<a href="#">311852</a>	594	249	1	43	similar to 1700113K14Rik protein (RGD1311084) alternative variant bSep08, mRNA.
<a href="#">RGD1311095</a>	<a href="#">RGD1311095.bSep08</a>	<a href="#">301004</a>	8176	1390	2	463	similar to hypothetical protein FLJ20259 (RGD1311095) alternative variant bSep08, mRNA.
<a href="#">RGD1311095</a>	<a href="#">RGD1311095.cSep08</a>	<a href="#">301004</a>	11333	590	5	190	similar to hypothetical protein FLJ20259 (RGD1311095) alternative variant cSep08, mRNA.
<a href="#">RGD1311098</a>	<a href="#">RGD1311098.bSep08</a>	<a href="#">362998</a>	2053	723	2	138	similar to RIKEN cDNA 2810451A06 (RGD1311098) alternative variant bSep08, mRNA.
<a href="#">RGD1311103</a>	<a href="#">RGD1311103.aSep08</a>	<a href="#">301100</a>	1135	654	3	161	similar to RIKEN cDNA 2410146L05 (RGD1311103) alternative variant aSep08, mRNA.
<a href="#">RGD1311107</a>	<a href="#">RGD1311107.bSep08</a>	<a href="#">308428</a>	3695	1653	2	446	similar to hypothetical protein R30953 1 (48.7 kD) (RGD1311107) alternative variant bSep08, complete mRNA.
<a href="#">RGD1311107</a>	<a href="#">RGD1311107.cSep08</a>	<a href="#">308428</a>	2946	828	2	238	similar to hypothetical protein R30953 1 (RGD1311107) alternative variant cSep08, mRNA.
<a href="#">RGD1311107</a>	<a href="#">RGD1311107.dSep08</a>	<a href="#">308428</a>	2683	664	2	128	similar to hypothetical protein R30953 1 (14.2 kD) (RGD1311107) alternative variant dSep08, mRNA.

<a href="#">RGD1311107</a>	<a href="#">RGD1311107.eSep08</a>	<a href="#">308428</a>	1289	510	2	36	similar to hypothetical protein R30953 1 (RGD1311107) alternative variant eSep08, mRNA.
<a href="#">RGD1311117</a>	<a href="#">RGD1311117.aSep08</a>	<a href="#">314401</a>	44645	2100		570	similar to KIAA1409 protein (RGD1311117) alternative variant aSep08, mRNA.
<a href="#">RGD1311117</a>	<a href="#">RGD1311117.bSep08</a>	<a href="#">314401</a>	76546	1788		467	similar to KIAA1409 protein (RGD1311117) alternative variant bSep08, mRNA.
<a href="#">RGD1311117</a>	<a href="#">RGD1311117.cSep08</a>	<a href="#">314401</a>	21950	1020		339	similar to KIAA1409 protein (RGD1311117) alternative variant cSep08, mRNA.
<a href="#">RGD1311122</a>	<a href="#">RGD1311122.bSep08</a>	<a href="#">364154</a>	3070	676	1	54	similar to RIKEN cDNA 1110003E01 (RGD1311122) alternative variant bSep08, mRNA.
<a href="#">RGD1311126</a>	<a href="#">RGD1311126.bSep08</a>	<a href="#">298366</a>	30756	2327	2	578	similar to RIKEN cDNA 4922503N01 (65.9 kD) (RGD1311126) alternative variant bSep08, complete mRNA.
<a href="#">RGD1311154</a>	<a href="#">RGD1311154.aSep08</a>	<a href="#">300317</a>	8160	1796	5	573	similar to hypothetical protein FLJ12242 (RGD1311154) alternative variant aSep08, mRNA.
<a href="#">RGD1311154</a>	<a href="#">RGD1311154.cSep08</a>	<a href="#">300317</a>	10649	1381	8	203	similar to hypothetical protein FLJ12242 (RGD1311154) alternative variant cSep08, mRNA.
<a href="#">RGD1311154</a>	<a href="#">RGD1311154.dSep08</a>	<a href="#">300317</a>	10042	751	6	183	similar to hypothetical protein FLJ12242 (RGD1311154) alternative variant dSep08, mRNA.
<a href="#">RGD1311154</a>	<a href="#">RGD1311154.eSep08</a>	<a href="#">300317</a>	9403	1207	6	148	similar to hypothetical protein FLJ12242 (RGD1311154) alternative variant eSep08, mRNA.
<a href="#">RGD1311186</a>	<a href="#">RGD1311186.aSep08</a>	<a href="#">293587</a>	2867	1798	4	480	similar to RIKEN cDNA 1810014F10 gene (RGD1311186) alternative variant aSep08, mRNA.
<a href="#">RGD1311186</a>	<a href="#">RGD1311186.cSep08</a>	<a href="#">293587</a>	2280	796	4	88	similar to RIKEN cDNA 1810014F10 gene (RGD1311186) alternative variant cSep08, mRNA.
<a href="#">RGD1311186</a>	<a href="#">RGD1311186.eSep08</a>	<a href="#">293587</a>	2213	462	4	76	similar to RIKEN cDNA 1810014F10 gene (RGD1311186) alternative variant eSep08, mRNA.
<a href="#">RGD1311186</a>	<a href="#">RGD1311186.fSep08</a>	<a href="#">293587</a>	3015	993	7	75	similar to RIKEN cDNA 1810014F10 gene (8.6 kD) (RGD1311186) alternative variant fSep08, mRNA.
<a href="#">RGD1311188</a>	<a href="#">RGD1311188.aSep08</a>	<a href="#">315088</a>	6955	1851	1	390	similar to 1500031N24Rik protein (44.5 kD) (RGD1311188) alternative variant aSep08, mRNA.
<a href="#">RGD1311188</a>	<a href="#">RGD1311188.bSep08</a>	<a href="#">315088</a>	5951	844	1	281	similar to 1500031N24Rik protein (RGD1311188) alternative variant bSep08, mRNA.
<a href="#">RGD1311224</a>	<a href="#">RGD1311224.aSep08</a>	<a href="#">295714</a>	5016	892		110	similar to fatty acid desaturase 2; linoleoyl-CoA desaturase (delta-6-desaturase)-like 2; delta-6 fatty acid desaturase (RGD1311224) mRNA.
<a href="#">RGD1311249</a>	<a href="#">RGD1311249.bSep08</a>	<a href="#">298201</a>	695	448	1	49	putative protein (5.5 kD) (RGD1311249) alternative variant bSep08, mRNA.
<a href="#">RGD1311249</a>	<a href="#">RGD1311249.cSep08</a>	<a href="#">298201</a>	9957	630	2	49	putative protein (5.5 kD) (RGD1311249) alternative variant cSep08, mRNA.
<a href="#">RGD1311251</a>	<a href="#">RGD1311251.bSep08</a>	<a href="#">315665</a>	7824	1088	4	311	similar to RIKEN cDNA 4930550C14 (RGD1311251) alternative variant bSep08, mRNA.
<a href="#">RGD1311251</a>	<a href="#">RGD1311251.cSep08</a>	<a href="#">315665</a>	15814	772	6	156	similar to RIKEN cDNA 4930550C14 (RGD1311251) alternative variant cSep08, mRNA.
<a href="#">RGD1311257</a>	<a href="#">RGD1311257.bSep08</a>	<a href="#">294333</a>	14916	758	2	197	similar to C21orf70 protein (RGD1311257) alternative variant bSep08, mRNA.
<a href="#">RGD1311260</a>	<a href="#">RGD1311260.aSep08</a>	<a href="#">303211</a>	3759	2279		124	hypothetical LOC303211 (13.8 kD) (RGD1311260) mRNA.

<a href="#">RGD1311265</a>	<a href="#">RGD1311265.bSep08</a>	<a href="#">361976</a>	4944	708	5	211	similar to CGI-41 protein (RGD1311265) alternative variant bSep08, mRNA.
<a href="#">RGD1311265</a>	<a href="#">RGD1311265.cSep08</a>	<a href="#">361976</a>	4552	397	3	132	similar to CGI-41 protein (RGD1311265) alternative variant cSep08, mRNA.
<a href="#">RGD1311265</a>	<a href="#">RGD1311265.dSep08</a>	<a href="#">361976</a>	5009	678	5	93	similar to CGI-41 protein (RGD1311265) alternative variant dSep08, mRNA.
<a href="#">RGD1311265</a>	<a href="#">RGD1311265.fSep08</a>	<a href="#">361976</a>	4195	2320	3	47	similar to CGI-41 protein (5.3 kD) (RGD1311265) alternative variant fSep08, mRNA.
<a href="#">RGD1311267</a>	<a href="#">RGD1311267.bSep08</a>	<a href="#">311429</a>	1102	803	1	69	similar to RIKEN cDNA 4931426K16 gene (7.7 kD) (RGD1311267) alternative variant bSep08, mRNA.
<a href="#">RGD1311269</a>	<a href="#">RGD1311269.aSep08</a>	<a href="#">301419</a>	18466	1800	4	542	similar to hypothetical protein FLJ37953 (RGD1311269) alternative variant aSep08, mRNA.
<a href="#">RGD1311269</a>	<a href="#">RGD1311269.bSep08</a>	<a href="#">301419</a>	12396	400	4	133	similar to hypothetical protein FLJ37953 (RGD1311269) alternative variant bSep08, mRNA.
<a href="#">RGD1311269</a>	<a href="#">RGD1311269.cSep08</a>	<a href="#">301419</a>	18743	1456	7	121	similar to hypothetical protein FLJ37953 (14.0 kD) (RGD1311269) alternative variant cSep08, mRNA.
<a href="#">RGD1311273</a>	<a href="#">RGD1311273.aSep08</a>	<a href="#">685545</a>	5815	1379	5	433	similar to RIKEN cDNA 9530058B02 (RGD1311273) alternative variant aSep08, mRNA.
<a href="#">RGD1311273</a>	<a href="#">RGD1311273.cSep08</a>	<a href="#">685545</a>	4922	883	4	134	similar to RIKEN cDNA 9530058B02 (15.0 kD) (RGD1311273) alternative variant cSep08, mRNA.
<a href="#">RGD1311273</a>	<a href="#">RGD1311273.dSep08</a>	<a href="#">685545</a>	1893	656	4	127	similar to RIKEN cDNA 9530058B02 (RGD1311273) alternative variant dSep08, mRNA.
<a href="#">RGD1311273</a>	<a href="#">RGD1311273.eSep08</a>	<a href="#">685545</a>	4742	416	3	109	similar to RIKEN cDNA 9530058B02 (RGD1311273) alternative variant eSep08, mRNA.
<a href="#">RGD1311294</a>	<a href="#">RGD1311294.aSep08</a>	<a href="#">361853</a>	22748	1798	8	514	similar to Hypothetical protein C6orf60 (RGD1311294) alternative variant aSep08, mRNA.
<a href="#">RGD1311294</a>	<a href="#">RGD1311294.bSep08</a>	<a href="#">361853</a>	20112	889	6	296	similar to Hypothetical protein C6orf60 (RGD1311294) alternative variant bSep08, mRNA.
<a href="#">RGD1311294</a>	<a href="#">RGD1311294.cSep08</a>	<a href="#">361853</a>	20043	709	4	126	similar to Hypothetical protein C6orf60 (RGD1311294) alternative variant cSep08, mRNA.
<a href="#">RGD1311294</a>	<a href="#">RGD1311294.dSep08</a>	<a href="#">361853</a>	6174	520	2	104	similar to Hypothetical protein C6orf60 (11.8 kD) (RGD1311294) alternative variant dSep08, mRNA.
<a href="#">RGD1311294</a>	<a href="#">RGD1311294.eSep08</a>	<a href="#">361853</a>	2660	1258	3	54	similar to Hypothetical protein C6orf60 (6.5 kD) (RGD1311294) alternative variant eSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.aSep08</a>	<a href="#">290071</a>	344095	2437	9	294	T cell receptor V delta 6 (32.3 kD) (RGD1311300andRGD1359684) alternative variant aSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.aSep08</a>	<a href="#">364378</a>	344095	2437	9	294	T cell receptor V delta 6 (32.3 kD) (RGD1311300andRGD1359684) alternative variant aSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.bSep08</a>	<a href="#">290071</a>	571920	1311	7	277	T-cell receptor like (RGD1311300andRGD1359684) alternative variant bSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.bSep08</a>	<a href="#">364378</a>	571920	1311	7	277	T-cell receptor like (RGD1311300andRGD1359684) alternative variant bSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.cSep08</a>	<a href="#">290071</a>	571503	888	7	275	T-cell receptor like (RGD1311300andRGD1359684) alternative variant cSep08, mRNA.

<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.cSep08</a>	<a href="#">364378</a>	571503	888	7	275	T-cell receptor like (RGD1311300andRGD1359684) alternative variant cSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.dSep08</a>	<a href="#">290071</a>	478702	1360	7	269	T cell receptor precursor (29.7 kD) (RGD1311300andRGD1359684) alternative variant dSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.dSep08</a>	<a href="#">364378</a>	478702	1360	7	269	T cell receptor precursor (29.7 kD) (RGD1311300andRGD1359684) alternative variant dSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.eSep08</a>	<a href="#">290071</a>	343464	1055	5	224	T-cell receptor like (RGD1311300andRGD1359684) alternative variant eSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.eSep08</a>	<a href="#">364378</a>	343464	1055	5	224	T-cell receptor like (RGD1311300andRGD1359684) alternative variant eSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.fSep08</a>	<a href="#">290071</a>	13516	1363	6	186	T cell receptor V delta (20.3 kD) (RGD1311300andRGD1359684) alternative variant fSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.fSep08</a>	<a href="#">364378</a>	13516	1363	6	186	T cell receptor V delta (20.3 kD) (RGD1311300andRGD1359684) alternative variant fSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.gSep08</a>	<a href="#">290071</a>	7762	964	5	162	T-cell receptor like (RGD1311300andRGD1359684) alternative variant gSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.gSep08</a>	<a href="#">364378</a>	7762	964	5	162	T-cell receptor like (RGD1311300andRGD1359684) alternative variant gSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.hSep08</a>	<a href="#">290071</a>	60922	953	5	158	T-cell receptor like (RGD1311300andRGD1359684) alternative variant hSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.hSep08</a>	<a href="#">364378</a>	60922	953	5	158	T-cell receptor like (RGD1311300andRGD1359684) alternative variant hSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.iSep08</a>	<a href="#">290071</a>	39261	944	5	157	T-cell receptor like (RGD1311300andRGD1359684) alternative variant iSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.iSep08</a>	<a href="#">364378</a>	39261	944	5	157	T-cell receptor like (RGD1311300andRGD1359684) alternative variant iSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.jSep08</a>	<a href="#">290071</a>	45481	947	5	156	T-cell receptor like (RGD1311300andRGD1359684) alternative variant jSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.jSep08</a>	<a href="#">364378</a>	45481	947	5	156	T-cell receptor like (RGD1311300andRGD1359684) alternative variant jSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.kSep08</a>	<a href="#">290071</a>	29379	944	5	155	T-cell receptor like (RGD1311300andRGD1359684) alternative variant kSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.kSep08</a>	<a href="#">364378</a>	29379	944	5	155	T-cell receptor like (RGD1311300andRGD1359684) alternative variant kSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.lSep08</a>	<a href="#">290071</a>	50588	930	5	153	T-cell receptor like (RGD1311300andRGD1359684) alternative variant lSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.lSep08</a>	<a href="#">364378</a>	50588	930	5	153	T-cell receptor like (RGD1311300andRGD1359684) alternative variant lSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.mSep08</a>	<a href="#">290071</a>	988	667	2	146	T cell receptor V delta 2 (RGD1311300andRGD1359684) alternative variant mSep08, mRNA.
<a href="#">RGD1311300andRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.mSep08</a>	<a href="#">364378</a>	988	667	2	146	T cell receptor V delta 2 (RGD1311300andRGD1359684) alternative variant mSep08, mRNA.

<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.nSep08</a>	<a href="#">290071</a>	50523	434	4	144	T-cell receptor like (RGD1311300andRGD1359684) alternative variant nSep08, mRNA.
<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.nSep08</a>	<a href="#">364378</a>	50523	434	4	144	T-cell receptor like (RGD1311300andRGD1359684) alternative variant nSep08, mRNA.
<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.oSep08</a>	<a href="#">290071</a>	33728	359	3	119	T-cell receptor like (RGD1311300andRGD1359684) alternative variant oSep08, mRNA.
<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.oSep08</a>	<a href="#">364378</a>	33728	359	3	119	T-cell receptor like (RGD1311300andRGD1359684) alternative variant oSep08, mRNA.
<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.gSep08</a>	<a href="#">290071</a>	14275	336	2	111	T-cell receptor like (RGD1311300andRGD1359684) alternative variant qSep08, mRNA.
<a href="#">RGD1311300andRGD1359684.dRGD1359684</a>	<a href="#">RGD1311300andRGD1359684.gSep08</a>	<a href="#">364378</a>	14275	336	2	111	T-cell receptor like (RGD1311300andRGD1359684) alternative variant qSep08, mRNA.
<a href="#">RGD1311307</a>	<a href="#">RGD1311307.bSep08</a>	<a href="#">361238</a>	3099	860	2	114	similar to 1300014I06Rik protein (13.0 kD) (RGD1311307) alternative variant bSep08, mRNA.
<a href="#">RGD1311307</a>	<a href="#">RGD1311307.cSep08</a>	<a href="#">361238</a>	11165	788	4	99	similar to 1300014I06Rik protein (RGD1311307) alternative variant cSep08, mRNA.
<a href="#">RGD1311309</a>	<a href="#">RGD1311309.aSep08</a>	<a href="#">364144</a>	22614	1467		433	similar to 2510002A14Rik protein (RGD1311309) mRNA.
<a href="#">RGD1311310</a>	<a href="#">RGD1311310.aSep08</a>	<a href="#">315166</a>	3281	786	3	184	similar to hypothetical protein supported by AL449243 (RGD1311310) alternative variant aSep08, mRNA.
<a href="#">RGD1311310</a>	<a href="#">RGD1311310.bSep08</a>	<a href="#">315166</a>	3105	561	3	110	similar to hypothetical protein supported by AL449243 (RGD1311310) alternative variant bSep08, mRNA.
<a href="#">RGD1311343</a>	<a href="#">RGD1311343.bSep08</a>	<a href="#">360506</a>	10906	521	1	112	similar to RIKEN cDNA 4930524B15 (RGD1311343) alternative variant bSep08, mRNA.
<a href="#">RGD1311344</a>	<a href="#">RGD1311344.aSep08</a>	<a href="#">311486</a>	4209	390		68	similar to RIKEN cDNA 2810039F03 (RGD1311344) mRNA.
<a href="#">RGD1311345</a>	<a href="#">RGD1311345.aSep08</a>	<a href="#">361201</a>	7671	2653	3	355	similar to CG9752-PA (RGD1311345) alternative variant aSep08, mRNA.
<a href="#">RGD1311345</a>	<a href="#">RGD1311345.bSep08</a>	<a href="#">361201</a>	15953	481	2	64	similar to CG9752-PA (RGD1311345) alternative variant bSep08, mRNA.
<a href="#">RGD1311345</a>	<a href="#">RGD1311345.cSep08</a>	<a href="#">361201</a>	5119	772	1	75	similar to CG9752-PA (8.9 kD) (RGD1311345) alternative variant cSep08, mRNA.
<a href="#">RGD1311350</a>	<a href="#">RGD1311350.aSep08</a>	<a href="#">293823</a>	22999	610	6	145	similar to KIAA0367 (RGD1311350) alternative variant aSep08, mRNA.
<a href="#">RGD1311350</a>	<a href="#">RGD1311350.bSep08</a>	<a href="#">293823</a>	23659	1267	6	120	similar to KIAA0367 (RGD1311350) alternative variant bSep08, mRNA.
<a href="#">RGD1311361</a>	<a href="#">RGD1311361.bSep08</a>	<a href="#">305171</a>	17355	1176	2	245	similar to hypothetical protein (RGD1311361) alternative variant bSep08, mRNA.
<a href="#">RGD1311361</a>	<a href="#">RGD1311361.cSep08</a>	<a href="#">305171</a>	8392	804	3	209	similar to hypothetical protein (RGD1311361) alternative variant cSep08, mRNA.
<a href="#">RGD1311361</a>	<a href="#">RGD1311361.dSep08</a>	<a href="#">305171</a>	19786	859	3	163	similar to hypothetical protein (RGD1311361) alternative variant dSep08, mRNA.
<a href="#">RGD1311362</a>	<a href="#">RGD1311362.aSep08</a>	<a href="#">362914</a>	5841	990	3	114	similar to hypothetical protein FLJ10204 (RGD1311362) alternative variant aSep08, mRNA.
<a href="#">RGD1311362</a>	<a href="#">RGD1311362.bSep08</a>	<a href="#">362914</a>	5382	1461	2	57	similar to hypothetical protein FLJ10204 (6.5 kD) (RGD1311362) alternative variant bSep08, mRNA.

<a href="#">RGD1311364</a>	<a href="#">RGD1311364.aSep08</a>	<a href="#">300516</a>	15967	1582	2	115	similar to RIKEN cDNA 1810021J13 (12.5 kD) (RGD1311364) alternative variant aSep08, complete mRNA.
<a href="#">RGD1311378</a>	<a href="#">RGD1311378.bSep08</a>	<a href="#">296408</a>	9423	3454	2	131	similar to RIKEN cDNA 2010011I20 (14.0 kD) (RGD1311378) alternative variant bSep08, mRNA.
<a href="#">RGD1311422</a>	<a href="#">RGD1311422.aSep08</a>	<a href="#">287822</a>	13985	1955	12	529	similar to CG8841-PA (RGD1311422) alternative variant aSep08, mRNA.
<a href="#">RGD1311422</a>	<a href="#">RGD1311422.bSep08</a>	<a href="#">287822</a>	7188	1863	9	374	similar to CG8841-PA (RGD1311422) alternative variant bSep08, mRNA.
<a href="#">RGD1311424</a>	<a href="#">RGD1311424.bSep08</a>	<a href="#">362685</a>	2457	718	4	133	similar to hypothetical protein FLJ38348 (RGD1311424) alternative variant bSep08, mRNA.
<a href="#">RGD1311424</a>	<a href="#">RGD1311424.cSep08</a>	<a href="#">362685</a>	3663	705	4	91	similar to hypothetical protein FLJ38348 (RGD1311424) alternative variant cSep08, mRNA.
<a href="#">RGD1311429</a>	<a href="#">RGD1311429.aSep08</a>	<a href="#">360642</a>	16609	2616	5	331	similar to KIAA1267 protein (RGD1311429) alternative variant aSep08, mRNA.
<a href="#">RGD1311429</a>	<a href="#">RGD1311429.bSep08</a>	<a href="#">360642</a>	14815	895	2	298	similar to KIAA1267 protein (RGD1311429) alternative variant bSep08, mRNA.
<a href="#">RGD1311429</a>	<a href="#">RGD1311429.cSep08</a>	<a href="#">360642</a>	13820	835	4	229	similar to KIAA1267 protein (RGD1311429) alternative variant cSep08, mRNA.
<a href="#">RGD1311429</a>	<a href="#">RGD1311429.dSep08</a>	<a href="#">360642</a>	3171	2136	4	217	similar to KIAA1267 protein (RGD1311429) alternative variant dSep08, mRNA.
<a href="#">RGD1311433</a>	<a href="#">RGD1311433.aSep08</a>	<a href="#">287054</a>	17542	1337	11	393	CRA b (43.7 kD) (RGD1311433) alternative variant aSep08, complete mRNA.
<a href="#">RGD1311433</a>	<a href="#">RGD1311433.bSep08</a>	<a href="#">287054</a>	14052	819	9	213	CRA b (RGD1311433) alternative variant bSep08, mRNA.
<a href="#">RGD1311433</a>	<a href="#">RGD1311433.dSep08</a>	<a href="#">287054</a>	2181	673	2	76	LP8272 (RGD1311433) alternative variant dSep08, mRNA.
<a href="#">RGD1311435</a>	<a href="#">RGD1311435.aSep08</a>	<a href="#">362037</a>	5152	1098	2	257	similar to hypothetical protein PRO0971 (RGD1311435) alternative variant aSep08, mRNA.
<a href="#">RGD1311435</a>	<a href="#">RGD1311435.bSep08</a>	<a href="#">362037</a>	9462	2171	4	232	similar to hypothetical protein PRO0971 (RGD1311435) alternative variant bSep08, mRNA.
<a href="#">RGD1311435</a>	<a href="#">RGD1311435.cSep08</a>	<a href="#">362037</a>	7760	370	3	123	similar to hypothetical protein PRO0971 (RGD1311435) alternative variant cSep08, mRNA.
<a href="#">RGD1311447</a>	<a href="#">RGD1311447.bSep08</a>	<a href="#">363276</a>	1020	895	2	129	LOC363276 (14.5 kD) (RGD1311447) alternative variant bSep08, mRNA.
<a href="#">RGD1311447</a>	<a href="#">RGD1311447.cSep08</a>	<a href="#">363276</a>	2636	551	4	119	LOC363276 (12.0 kD) (RGD1311447) alternative variant cSep08, mRNA.
<a href="#">RGD1311456</a>	<a href="#">RGD1311456.aSep08</a>	<a href="#">363089</a>	43276	748		249	similar to RIKEN cDNA B230380D07 (RGD1311456) mRNA.
<a href="#">RGD1311463</a>	<a href="#">RGD1311463.bSep08</a>	<a href="#">311279</a>	9076	890	5	259	similar to RIKEN cDNA 2700007P21 (29.2 kD) (RGD1311463) alternative variant bSep08, mRNA.
<a href="#">RGD1311463</a>	<a href="#">RGD1311463.dSep08</a>	<a href="#">311279</a>	2630	419	2	135	similar to RIKEN cDNA 2700007P21 (RGD1311463) alternative variant dSep08, mRNA.
<a href="#">RGD1311463</a>	<a href="#">RGD1311463.eSep08</a>	<a href="#">311279</a>	9121	304	2	95	similar to RIKEN cDNA 2700007P21 (RGD1311463) alternative variant eSep08, mRNA.
<a href="#">RGD1311490</a>	<a href="#">RGD1311490.bSep08</a>	<a href="#">361697</a>	6751	1576	9	320	similar to DKFZP434P1750 protein (35.3 kD) (RGD1311490) alternative variant bSep08, complete mRNA.



<a href="#">RGD1311490</a>	<a href="#">RGD1311490.cSep08</a>	<a href="#">361697</a>	4399	656	6	218	similar to DKFZP434P1750 protein (RGD1311490) alternative variant cSep08, mRNA.
<a href="#">RGD1311490</a>	<a href="#">RGD1311490.fSep08</a>	<a href="#">361697</a>	1222	776	3	99	similar to DKFZP434P1750 protein (RGD1311490) alternative variant fSep08, mRNA.
<a href="#">RGD1311493</a>	<a href="#">RGD1311493.aSep08</a>	<a href="#">287734</a>	5690	3407		342	similar to CG13379-PA (RGD1311493) mRNA.
<a href="#">RGD1311501</a>	<a href="#">RGD1311501.bSep08</a>	<a href="#">296599</a>	9309	2099	3	108	putative protein of metazoan origin (RGD1311501) alternative variant bSep08, complete mRNA.
<a href="#">RGD1311517</a>	<a href="#">RGD1311517.aSep08</a>	<a href="#">313775</a>	14092	692	7	146	similar to RIKEN cDNA 9430015G10 (RGD1311517) alternative variant aSep08, mRNA.
<a href="#">RGD1311517</a>	<a href="#">RGD1311517.bSep08</a>	<a href="#">313775</a>	4141	2304	2	47	similar to RIKEN cDNA 9430015G10 (RGD1311517) alternative variant bSep08, mRNA.
<a href="#">RGD1311517</a>	<a href="#">RGD1311517.cSep08</a>	<a href="#">313775</a>	18264	808	5	43	similar to RIKEN cDNA 9430015G10 (4.9 kD) (RGD1311517) alternative variant cSep08, complete mRNA.
<a href="#">RGD1311517</a>	<a href="#">RGD1311517.dSep08</a>	<a href="#">313775</a>	8176	636	5	58	similar to RIKEN cDNA 9430015G10 (6.2 kD) (RGD1311517) alternative variant dSep08, mRNA.
<a href="#">RGD1311519</a>	<a href="#">RGD1311519.aSep08</a>	<a href="#">311496</a>	54379	276		92	hypothetical LOC311496 (RGD1311519) mRNA.
<a href="#">RGD1311558</a>	<a href="#">RGD1311558.bSep08</a>	<a href="#">292139</a>	55512	741	8	236	similar to 4930506M07Rik protein (RGD1311558) alternative variant bSep08, mRNA.
<a href="#">RGD1311558</a>	<a href="#">RGD1311558.cSep08</a>	<a href="#">292139</a>	30956	832	8	226	similar to 4930506M07Rik protein (RGD1311558) alternative variant cSep08, mRNA.
<a href="#">RGD1311558</a>	<a href="#">RGD1311558.dSep08</a>	<a href="#">292139</a>	26114	2205	4	194	similar to 4930506M07Rik protein (RGD1311558) alternative variant dSep08, mRNA.
<a href="#">RGD1311563</a>	<a href="#">RGD1311563.aSep08</a>	<a href="#">363160</a>	37450	3428	14	505	similar to Oligosaccharyl transferase 3 CG7748-PA (RGD1311563) alternative variant aSep08, mRNA.
<a href="#">RGD1311563</a>	<a href="#">RGD1311563.bSep08</a>	<a href="#">363160</a>	66960	3044	13	505	similar to Oligosaccharyl transferase 3 CG7748-PA (RGD1311563) alternative variant bSep08, mRNA.
<a href="#">RGD1311564</a>	<a href="#">RGD1311564.bSep08</a>	<a href="#">360590</a>	21077	356	3	52	LOC360590 (RGD1311564) alternative variant bSep08, mRNA.
<a href="#">RGD1311575</a>	<a href="#">RGD1311575.aSep08</a>	<a href="#">289568</a>	10418	2096	1	323	hypothetical LOC289568 (RGD1311575) alternative variant aSep08, mRNA.
<a href="#">RGD1311575</a>	<a href="#">RGD1311575.bSep08</a>	<a href="#">289568</a>	3919	675	1	168	hypothetical LOC289568 (18.5 kD) (RGD1311575) alternative variant bSep08, mRNA.
<a href="#">RGD1311578</a>	<a href="#">RGD1311578.bSep08</a>	<a href="#">298748</a>	6597	843	7	260	upf0511 protein c2orf56 homolog mitochondrial (29.2 kD) (RGD1311578) alternative variant bSep08, mRNA.
<a href="#">RGD1311578</a>	<a href="#">RGD1311578.cSep08</a>	<a href="#">298748</a>	4873	677	5	165	upf0511 protein c2orf56 homolog mitochondrial (RGD1311578) alternative variant cSep08, mRNA.
<a href="#">RGD1311578</a>	<a href="#">RGD1311578.dSep08</a>	<a href="#">298748</a>	1033	635	2	99	upf0511 protein c2orf56 homolog mitochondrial (11.0 kD) (RGD1311578) alternative variant dSep08, mRNA.
<a href="#">RGD1311605</a>	<a href="#">RGD1311605.bSep08</a>	<a href="#">298841</a>	1978	821	1	115	similar to apoptosis related protein APR-3; p18 protein (12.4 kD) (RGD1311605) alternative variant bSep08, mRNA.
<a href="#">RGD1311612</a>	<a href="#">RGD1311612.aSep08</a>	<a href="#">361184</a>	37778	1823	14	373	cysteinyI-tRNA synthetase 2 (RGD1311612) alternative variant aSep08, mRNA.
<a href="#">RGD1311612</a>	<a href="#">RGD1311612.bSep08</a>	<a href="#">361184</a>	4932	1269	6	205	cysteinyI-tRNA synthetase 2 (22.5 kD) (RGD1311612) alternative variant bSep08, mRNA.
<a href="#">RGD1311612</a>	<a href="#">RGD1311612.cSep08</a>	<a href="#">361184</a>	7485	1080	3	163	CRA d (RGD1311612) alternative variant cSep08, mRNA.

<a href="#">RGD1311612</a>	<a href="#">RGD1311612.dSep08</a>	<a href="#">361184</a>	20600	576	6	114	cysteinyI-tRNA synthetase 2 (RGD1311612) alternative variant dSep08, mRNA.
<a href="#">RGD1311612</a>	<a href="#">RGD1311612.eSep08</a>	<a href="#">361184</a>	709	431	2	95	cysteinyI-tRNA synthetase 2 (10.5 kD) (RGD1311612) alternative variant eSep08, mRNA.
<a href="#">RGD1311615</a>	<a href="#">RGD1311615.aSep08</a>	<a href="#">366531</a>	7221	766		159	similar to hypothetical protein FLJ33868 (RGD1311615) mRNA.
<a href="#">RGD1311624</a>	<a href="#">RGD1311624.aSep08</a>	<a href="#">309001</a>	1729	1288	3	429	similar to KIAA0339 protein (RGD1311624) alternative variant aSep08, mRNA.
<a href="#">RGD1311624</a>	<a href="#">RGD1311624.bSep08</a>	<a href="#">309001</a>	3647	1622	6	324	similar to KIAA0339 protein (RGD1311624) alternative variant bSep08, mRNA.
<a href="#">RGD1311625</a>	<a href="#">RGD1311625.aSep08</a>	<a href="#">310859</a>	7960	299		99	similar to KIAA1546 protein (RGD1311625) mRNA.
<a href="#">RGD1311634</a>	<a href="#">RGD1311634.aSep08</a>	<a href="#">293155</a>	3174	1001	4	132	similar to RIKEN cDNA 3200002M19 (15.4 kD) (RGD1311634) alternative variant aSep08, mRNA.
<a href="#">RGD1311634</a>	<a href="#">RGD1311634.bSep08</a>	<a href="#">293155</a>	3655	890	4	121	similar to RIKEN cDNA 3200002M19 (14.3 kD) (RGD1311634) alternative variant bSep08, mRNA.
<a href="#">RGD1311634</a>	<a href="#">RGD1311634.cSep08</a>	<a href="#">293155</a>	27174	1175	4	121	similar to RIKEN cDNA 3200002M19 (13.9 kD) (RGD1311634) alternative variant cSep08, mRNA.
<a href="#">RGD1311640</a>	<a href="#">RGD1311640.aSep08</a>	<a href="#">314787</a>	5716	1090		363	similar to Hypothetical protein KIAA0373 (RGD1311640) mRNA.
<a href="#">RGD1311642</a>	<a href="#">RGD1311642.aSep08</a>	<a href="#">499941</a>	3729	745	3	180	similar to uncharacterized hypothalamus protein HSMNP1 (RGD1311642) alternative variant aSep08, mRNA.
<a href="#">RGD1311642</a>	<a href="#">RGD1311642.cSep08</a>	<a href="#">499941</a>	2255	364	2	120	similar to uncharacterized hypothalamus protein HSMNP1 (RGD1311642) alternative variant cSep08, mRNA.
<a href="#">RGD1311648</a>	<a href="#">RGD1311648.bSep08</a>	<a href="#">313949</a>	56815	742	5	225	similar to hypothetical protein FLJ21820 (RGD1311648) alternative variant bSep08, mRNA.
<a href="#">RGD1311660</a>	<a href="#">RGD1311660.aSep08</a>	<a href="#">288518</a>	17220	2266	8	330	similar to RIKEN cDNA 1110007L15 (36.8 kD) (RGD1311660) alternative variant aSep08, mRNA.
<a href="#">RGD1311660</a>	<a href="#">RGD1311660.bSep08</a>	<a href="#">288518</a>	13941	750	6	247	similar to RIKEN cDNA 1110007L15 (RGD1311660) alternative variant bSep08, mRNA.
<a href="#">RGD1311660</a>	<a href="#">RGD1311660.cSep08</a>	<a href="#">288518</a>	800	681	1	73	similar to RIKEN cDNA 1110007L15 (RGD1311660) alternative variant cSep08, mRNA.
<a href="#">RGD1311678</a>	<a href="#">RGD1311678.aSep08</a>	<a href="#">296311</a>	45751	2038	2	452	similar to 4921517L17Rik protein (RGD1311678) alternative variant aSep08, mRNA.
<a href="#">RGD1311678</a>	<a href="#">RGD1311678.bSep08</a>	<a href="#">296311</a>	30214	740	2	246	similar to 4921517L17Rik protein (RGD1311678) alternative variant bSep08, mRNA.
<a href="#">RGD1311678</a>	<a href="#">RGD1311678.cSep08</a>	<a href="#">296311</a>	29900	728	1	242	similar to 4921517L17Rik protein (RGD1311678) alternative variant cSep08, mRNA.
<a href="#">RGD1311703</a>	<a href="#">RGD1311703.aSep08</a>	<a href="#">293160</a>	10886	1204	5	232	similar to sid2057p (RGD1311703) alternative variant aSep08, mRNA.
<a href="#">RGD1311723</a>	<a href="#">RGD1311723.aSep08</a>	<a href="#">363018</a>	5848	1581	8	480	similar to KIAA1731 protein (RGD1311723) alternative variant aSep08, mRNA.
<a href="#">RGD1311723</a>	<a href="#">RGD1311723.bSep08</a>	<a href="#">363018</a>	866	332	3	81	similar to KIAA1731 protein (9.9 kD) (RGD1311723) alternative variant bSep08, mRNA.
<a href="#">RGD1311723</a>	<a href="#">RGD1311723.dSep08</a>	<a href="#">363018</a>	317	203	2	38	similar to KIAA1731 protein (4.7 kD) (RGD1311723) alternative variant dSep08, mRNA.
<a href="#">RGD1311730</a>	<a href="#">RGD1311730.bSep08</a>	<a href="#">292811</a>	17434	1399	9	465	similar to RIKEN cDNA 2610507L03 (RGD1311730) alternative variant bSep08, mRNA.

<a href="#">RGD1311730</a>	<a href="#">RGD1311730.cSep08</a>	<a href="#">292811</a>	23057	1072	6	311	similar to RIKEN cDNA 2610507L03 (RGD1311730) alternative variant cSep08, mRNA.
<a href="#">RGD1311730</a>	<a href="#">RGD1311730.dSep08</a>	<a href="#">292811</a>	3249	465	2	109	similar to RIKEN cDNA 2610507L03 (RGD1311730) alternative variant dSep08, mRNA.
<a href="#">RGD1311730</a>	<a href="#">RGD1311730.eSep08</a>	<a href="#">292811</a>	3158	398	1	44	similar to RIKEN cDNA 2610507L03 (RGD1311730) alternative variant eSep08, mRNA.
<a href="#">RGD1311730</a>	<a href="#">RGD1311730.fSep08</a>	<a href="#">292811</a>	10148	962	3		
<a href="#">RGD1311739</a>	<a href="#">RGD1311739.bSep08</a>	<a href="#">311428</a>	13140	735	7	116	similar to RIKEN cDNA 1700037H04 (RGD1311739) alternative variant bSep08, mRNA.
<a href="#">RGD1311739</a>	<a href="#">RGD1311739.cSep08</a>	<a href="#">311428</a>	6783	877	3	84	similar to RIKEN cDNA 1700037H04 (9.2 kD) (RGD1311739) alternative variant cSep08, mRNA.
<a href="#">RGD1311739</a>	<a href="#">RGD1311739.dSep08</a>	<a href="#">311428</a>	975	793	2	79	similar to RIKEN cDNA 1700037H04 (RGD1311739) alternative variant dSep08, mRNA.
<a href="#">RGD1311742</a>	<a href="#">RGD1311742.bSep08</a>	<a href="#">291676</a>	3943	441	5	135	similar to RIKEN cDNA 5133400G04 (RGD1311742) alternative variant bSep08, mRNA.
<a href="#">RGD1311742</a>	<a href="#">RGD1311742.cSep08</a>	<a href="#">291676</a>	6210	292	3	88	similar to RIKEN cDNA 5133400G04 (RGD1311742) alternative variant cSep08, mRNA.
<a href="#">RGD1311744</a>	<a href="#">RGD1311744.aSep08</a>	<a href="#">300608</a>	7152	736		245	similar to RIKEN cDNA 5830475I06 (RGD1311744) mRNA.
<a href="#">RGD1311747</a>	<a href="#">RGD1311747.aSep08</a>	<a href="#">290706</a>	16492	1329	5	368	similar to 2700029M09Rik protein (RGD1311747) alternative variant aSep08, mRNA.
<a href="#">RGD1311747</a>	<a href="#">RGD1311747.bSep08</a>	<a href="#">290706</a>	13232	1078	3	335	similar to 2700029M09Rik protein (RGD1311747) alternative variant bSep08, mRNA.
<a href="#">RGD1311747</a>	<a href="#">RGD1311747.cSep08</a>	<a href="#">290706</a>	19671	979	3	179	similar to 2700029M09Rik protein (20.3 kD) (RGD1311747) alternative variant cSep08, mRNA.
<a href="#">RGD1311747</a>	<a href="#">RGD1311747.dSep08</a>	<a href="#">290706</a>	2252	529	2	78	similar to 2700029M09Rik protein (8.8 kD) (RGD1311747) alternative variant dSep08, mRNA.
<a href="#">RGD1311756</a>	<a href="#">RGD1311756.aSep08</a>	<a href="#">362769</a>	47514	982		222	similar to hypothetical protein FLJ20950 (RGD1311756) mRNA.
<a href="#">RGD1311783</a>	<a href="#">RGD1311783.bSep08</a>	<a href="#">294012</a>	13278	1033	4	83	similar to RIKEN cDNA 2010012O05 (RGD1311783) alternative variant bSep08, mRNA.
<a href="#">RGD1311783</a>	<a href="#">RGD1311783.cSep08</a>	<a href="#">294012</a>	11798	277	3	83	similar to RIKEN cDNA 2010012O05 (9.0 kD) (RGD1311783) alternative variant cSep08, complete mRNA.
<a href="#">RGD1311784</a>	<a href="#">RGD1311784.bSep08</a>	<a href="#">287871</a>	14563	1781	8	150	similar to p150 target of rapamycin (TOR)-scaffold protein containing WD-repeats (RGD1311784) alternative variant bSep08, mRNA.
<a href="#">RGD1311784</a>	<a href="#">RGD1311784.cSep08</a>	<a href="#">287871</a>	11491	318	4	105	similar to p150 target of rapamycin (TOR)-scaffold protein containing WD-repeats (RGD1311784) alternative variant cSep08, mRNA.
<a href="#">RGD1311805</a>	<a href="#">RGD1311805.bSep08</a>	<a href="#">291784</a>	848	434	2	67	similar to RIKEN cDNA 2400010D15 (7.7 kD) (RGD1311805) alternative variant bSep08, mRNA.
<a href="#">RGD1311847</a>	<a href="#">RGD1311847.bSep08</a>	<a href="#">290615</a>	10462	488	2	72	similar to 1700030K09Rik protein (RGD1311847) alternative variant bSep08, mRNA.
<a href="#">RGD1311849</a>	<a href="#">RGD1311849.aSep08</a>	<a href="#">313346</a>	97341	1477		330	similar to mKIAA1797 protein (RGD1311849) mRNA.
<a href="#">RGD1311861</a>	<a href="#">RGD1311861.aSep08</a>	<a href="#">288042</a>	12301	938		230	similar to hypothetical protein MGC19444 (RGD1311861) mRNA.

<a href="#">RGD1311870</a>	<a href="#">RGD1311870.aSep08</a>	<a href="#">298563</a>	4658	400		122	similar to RIKEN cDNA 4930549C01 (RGD1311870) mRNA.
<a href="#">RGD1311874</a>	<a href="#">RGD1311874.bSep08</a>	<a href="#">300751</a>	158152	686	5	189	hypothetical LOC300751 (RGD1311874) alternative variant bSep08, mRNA.
<a href="#">RGD1311874</a>	<a href="#">RGD1311874.cSep08</a>	<a href="#">300751</a>	8476	1424	2	118	hypothetical LOC300751 (RGD1311874) alternative variant cSep08, mRNA.
<a href="#">RGD1311893</a>	<a href="#">RGD1311893.bSep08</a>	<a href="#">288478</a>	2044	422	2	113	hypothetical LOC288478 (RGD1311893) alternative variant bSep08, mRNA.
<a href="#">RGD1311893</a>	<a href="#">RGD1311893.cSep08</a>	<a href="#">288478</a>	4613	614	2	85	hypothetical LOC288478 (9.5 kD) (RGD1311893) alternative variant cSep08, mRNA.
<a href="#">RGD1311899</a>	<a href="#">RGD1311899.bSep08</a>	<a href="#">288704</a>	5575	1491	6	218	similar to RIKEN cDNA 2210016L21 gene (24.0 kD) (RGD1311899) alternative variant bSep08, complete mRNA.
<a href="#">RGD1311899</a>	<a href="#">RGD1311899.cSep08</a>	<a href="#">288704</a>	3551	487	4	157	similar to RIKEN cDNA 2210016L21 gene (RGD1311899) alternative variant cSep08, mRNA.
<a href="#">RGD1311899</a>	<a href="#">RGD1311899.eSep08</a>	<a href="#">288704</a>	1143	962	2	56	similar to RIKEN cDNA 2210016L21 gene (RGD1311899) alternative variant eSep08, mRNA.
<a href="#">RGD1311906</a>	<a href="#">RGD1311906.aSep08</a>	<a href="#">292746</a>	25963	6820		2223	similar to Fc fragment of IgG binding protein; IgG Fc binding protein (RGD1311906) mRNA.
<a href="#">RGD1311910</a>	<a href="#">RGD1311910.aSep08</a>	<a href="#">307235</a>	9193	1361	5	387	similar to hypothetical p38 protein (RGD1311910) alternative variant aSep08, mRNA.
<a href="#">RGD1311933</a>	<a href="#">RGD1311933.aSep08</a>	<a href="#">308056</a>	5458	574		117	similar to RIKEN cDNA 2310057J18 (13.5 kD) (RGD1311933) mRNA.
<a href="#">RGD1311940</a>	<a href="#">RGD1311940.bSep08</a>	<a href="#">362287</a>	2579	1127	4	136	putative protein, with a transmembrane domain, of ancient origin (RGD1311940) alternative variant bSep08, mRNA.
<a href="#">RGD1311940</a>	<a href="#">RGD1311940.cSep08</a>	<a href="#">362287</a>	8493	412	4	108	uncharacterized mfs-type transporter homolog like (11.9 kD) (RGD1311940) alternative variant cSep08, mRNA.
<a href="#">RGD1311940</a>	<a href="#">RGD1311940.dSep08</a>	<a href="#">362287</a>	6771	381	2	94	uncharacterized mfs-type transporter homolog like (RGD1311940) alternative variant dSep08, mRNA.
<a href="#">RGD1311952</a>	<a href="#">RGD1311952.aSep08</a>	<a href="#">311692</a>	5859	501	1	166	similar to Protein C20orf177 (RGD1311952) alternative variant aSep08, mRNA.
<a href="#">RGD1311952</a>	<a href="#">RGD1311952.bSep08</a>	<a href="#">311692</a>	5703	386	1	128	similar to Protein C20orf177 (RGD1311952) alternative variant bSep08, mRNA.
<a href="#">RGD1311993</a>	<a href="#">RGD1311993.aSep08</a>	<a href="#">315072</a>	4869	776	6	228	hypothetical LOC315072 (RGD1311993) alternative variant aSep08, mRNA.
<a href="#">RGD1311993</a>	<a href="#">RGD1311993.bSep08</a>	<a href="#">315072</a>	5448	687	6	206	hypothetical LOC315072 (RGD1311993) alternative variant bSep08, mRNA.
<a href="#">RGD1311993</a>	<a href="#">RGD1311993.cSep08</a>	<a href="#">315072</a>	495	243	2	74	hypothetical LOC315072 (RGD1311993) alternative variant cSep08, mRNA.
<a href="#">RGD1312005</a>	<a href="#">RGD1312005.aSep08</a>	<a href="#">291580</a>	14533	1970	3	193	similar to DD1 (21.7 kD) (RGD1312005) alternative variant aSep08, mRNA.
<a href="#">RGD1312005</a>	<a href="#">RGD1312005.cSep08</a>	<a href="#">291580</a>	13145	662	4	151	similar to DD1 (17.0 kD) (RGD1312005) alternative variant cSep08, complete mRNA.
<a href="#">RGD1312005</a>	<a href="#">RGD1312005.dSep08</a>	<a href="#">291580</a>	3617	383	2	127	similar to DD1 (RGD1312005) alternative variant dSep08, mRNA.
<a href="#">RGD1312005</a>	<a href="#">RGD1312005.eSep08</a>	<a href="#">291580</a>	13432	981	5	117	similar to DD1 (13.5 kD) (RGD1312005) alternative variant eSep08, mRNA.

<a href="#">RGD1312005</a>	<a href="#">RGD1312005.fSep08</a>	<a href="#">291580</a>	9556	358	4	80	similar to DD1 (RGD1312005) alternative variant fSep08, mRNA.
<a href="#">RGD1312026</a>	<a href="#">RGD1312026.bSep08</a>	<a href="#">315686</a>	98913	496	4	35	similar to RIKEN cDNA C230081A13 (3.8 kD) (RGD1312026) alternative variant bSep08, mRNA.
<a href="#">RGD1312038</a>	<a href="#">RGD1312038.aSep08</a>	<a href="#">362732</a>	337396	2328	7	497	similar to putative protein, with at least 6 transmembrane domains, of ancient origin (58.5 kD) (3N884) (RGD1312038) alternative variant aSep08, mRNA.
<a href="#">RGD1312038</a>	<a href="#">RGD1312038.bSep08</a>	<a href="#">362732</a>	181645	1039	2	235	similar to putative protein, with at least 6 transmembrane domains, of ancient origin (58.5 kD) (3N884) (26.9 kD) (RGD1312038) alternative variant bSep08, mRNA.
<a href="#">RGD1359108</a>	<a href="#">RGD1359108.bSep08</a>	<a href="#">313155</a>	2340	767	2	50	similar to RIKEN cDNA 3110043O21 (RGD1359108) alternative variant bSep08, mRNA.
<a href="#">RGD1359158</a>	<a href="#">RGD1359158.bSep08</a>	<a href="#">361740</a>	3567	633	2	60	similar to RIKEN cDNA 1110059E24 (RGD1359158) alternative variant bSep08, mRNA.
<a href="#">RGD1359191</a>	<a href="#">RGD1359191.aSep08</a>	<a href="#">314462</a>	5394	1699	1	290	trna-methyltransferase TRM61 (31.6 kD) (RGD1359191) alternative variant aSep08, mRNA.
<a href="#">RGD1359201</a>	<a href="#">RGD1359201.aSep08</a>	<a href="#">289595</a>	21303	1790		395	similar to nuclear transcription factor, X-box binding-like 1 (RGD1359201) alternative variant aSep08, mRNA.
<a href="#">RGD1359310</a>	<a href="#">RGD1359310.bSep08</a>	<a href="#">300240</a>	9135	1237	2	218	similar to RIKEN cDNA 9430023L20 (25.0 kD) (RGD1359310) alternative variant bSep08, complete mRNA.
<a href="#">RGD1359310</a>	<a href="#">RGD1359310.cSep08</a>	<a href="#">300240</a>	3377	710	1	28	similar to RIKEN cDNA 9430023L20 (RGD1359310) alternative variant cSep08, mRNA.
<a href="#">RGD1359310</a>	<a href="#">RGD1359310.dSep08</a>	<a href="#">300240</a>	3244	390	2	58	similar to RIKEN cDNA 9430023L20 (RGD1359310) alternative variant dSep08, mRNA.
<a href="#">RGD1359380</a>	<a href="#">RGD1359380.bSep08</a>	<a href="#">303922</a>	20828	853	6	236	similar to hypothetical protein MGC7537 (26.4 kD) (RGD1359380) alternative variant bSep08, mRNA.
<a href="#">RGD1359380</a>	<a href="#">RGD1359380.cSep08</a>	<a href="#">303922</a>	14502	734	5	125	similar to hypothetical protein MGC7537 (RGD1359380) alternative variant cSep08, mRNA.
<a href="#">RGD1359449</a>	<a href="#">RGD1359449.bSep08</a>	<a href="#">314959</a>	5760	773	5	212	putative protein of vertebrate origin (RGD1359449) alternative variant bSep08, mRNA.
<a href="#">RGD1359449</a>	<a href="#">RGD1359449.cSep08</a>	<a href="#">314959</a>	3584	761	6	163	putative protein of vertebrate origin (RGD1359449) alternative variant cSep08, mRNA.
<a href="#">RGD1359449</a>	<a href="#">RGD1359449.dSep08</a>	<a href="#">314959</a>	1348	786	2	141	putative protein, with a coiled coil domain, of mammalian origin (15.6 kD) (RGD1359449) alternative variant dSep08, mRNA.
<a href="#">RGD1359452</a>	<a href="#">RGD1359452.bSep08</a>	<a href="#">296118</a>	37197	793	9	264	similar to hypothetical protein FLJ32800 (RGD1359452) alternative variant bSep08, mRNA.
<a href="#">RGD1359452</a>	<a href="#">RGD1359452.cSep08</a>	<a href="#">296118</a>	6537	700	5	208	similar to hypothetical protein FLJ32800 (RGD1359452) alternative variant cSep08, mRNA.
<a href="#">RGD1359452</a>	<a href="#">RGD1359452.dSep08</a>	<a href="#">296118</a>	37167	740	10	177	similar to hypothetical protein FLJ32800 (RGD1359452) alternative variant dSep08, mRNA.
<a href="#">RGD1359460</a>	<a href="#">RGD1359460.bSep08</a>	<a href="#">289562</a>	12431	957	1	255	putative protein of ancient origin (RGD1359460) alternative variant bSep08, mRNA.
<a href="#">RGD1359508</a>	<a href="#">RGD1359508.bSep08</a>	<a href="#">361941</a>	21896	752		103	similar to protein C33A12.3 (12.2 kD) (RGD1359508) alternative variant bSep08, mRNA.

<a href="#">RGD1359529</a>	<a href="#">RGD1359529.bSep08</a>	<a href="#">362626</a>	3713	2969	3	251	putative nuclear protein of mammalian origin (29.3 kD) (RGD1359529) alternative variant bSep08, complete mRNA.
<a href="#">RGD1359529</a>	<a href="#">RGD1359529.cSep08</a>	<a href="#">362626</a>	3224	1151	5	227	DNA segment Chr 4 Wayne State University 53 expressed like (RGD1359529) alternative variant cSep08, mRNA.
<a href="#">RGD1359529</a>	<a href="#">RGD1359529.dSep08</a>	<a href="#">362626</a>	2587	1555	3	169	DNA segment Chr 4 Wayne State University 53 expressed like (RGD1359529) alternative variant dSep08, mRNA.
<a href="#">RGD1359616</a>	<a href="#">RGD1359616.bSep08</a>	<a href="#">300782</a>	5367	1014	4	163	putative mitochondrial protein (18.8 kD) (RGD1359616) alternative variant bSep08, mRNA.
<a href="#">RGD1359616</a>	<a href="#">RGD1359616.cSep08</a>	<a href="#">300782</a>	1033	587	2	132	putative protein (RGD1359616) alternative variant cSep08, mRNA.
<a href="#">RGD1359616</a>	<a href="#">RGD1359616.eSep08</a>	<a href="#">300782</a>	3202	743	2	92	putative protein of metazoan origin (RGD1359616) alternative variant eSep08, mRNA.
<a href="#">RGD1359634</a>	<a href="#">RGD1359634.aSep08</a>	<a href="#">315126</a>	7551	1192	2	235	similar to RIKEN cDNA 1700088E04 (RGD1359634) alternative variant aSep08, mRNA.
<a href="#">RGD1359634</a>	<a href="#">RGD1359634.cSep08</a>	<a href="#">315126</a>	7110	790	3	181	similar to RIKEN cDNA 1700088E04 (20.8 kD) (RGD1359634) alternative variant cSep08, mRNA.
<a href="#">RGD1359713</a>	<a href="#">RGD1359713.bSep08</a>	<a href="#">305340</a>	38068	1783	1	440	hypothetical RNA binding protein RGD1359713 (RGD1359713) alternative variant bSep08, mRNA.
<a href="#">RGD1359713</a>	<a href="#">RGD1359713.cSep08</a>	<a href="#">305340</a>	30678	815	1	201	hypothetical RNA binding protein RGD1359713 (RGD1359713) alternative variant cSep08, mRNA.
<a href="#">RGD1559150</a>	<a href="#">RGD1559150.bSep08</a>	<a href="#">311353</a>	1171	719	2	92	sim to KOX31-like Zfp (9.8 kD) (RGD1559150) alternative variant bSep08, mRNA.
<a href="#">RGD1559441</a>	<a href="#">RGD1559441.aSep08</a>	<a href="#">500410</a>	47723	1427		164	similar to MIC2L1 (17.1 kD) (RGD1559441) mRNA.
<a href="#">RGD1559482</a>	<a href="#">RGD1559482.aSep08</a>	<a href="#">498022</a>	4799	844	3	211	similar to immunoglobulin superfamily, member 7 (23.2 kD) (RGD1559482) alternative variant aSep08, mRNA.
<a href="#">RGD1559482</a>	<a href="#">RGD1559482.bSep08</a>	<a href="#">498022</a>	2177	382	1	88	similar to immunoglobulin superfamily, member 7 (RGD1559482) alternative variant bSep08, mRNA.
<a href="#">RGD1559493</a>	<a href="#">RGD1559493.bSep08</a>	<a href="#">500516</a>	25902	1950	5	402	similar to Hypothetical protein MGC58608 (45.8 kD) (RGD1559493) alternative variant bSep08, complete mRNA.
<a href="#">RGD1559493</a>	<a href="#">RGD1559493.cSep08</a>	<a href="#">500516</a>	3664	724	1	172	similar to Hypothetical protein MGC58608 (RGD1559493) alternative variant cSep08, mRNA.
<a href="#">RGD1559494</a>	<a href="#">RGD1559494.aSep08</a>	<a href="#">499535</a>	6918	452	3	93	RGD1559494 (RGD1559494) alternative variant aSep08, mRNA.
<a href="#">RGD1559494</a>	<a href="#">RGD1559494.bSep08</a>	<a href="#">499535</a>	6871	1673	3	31	RGD1559494 (3.6 kD) (RGD1559494) alternative variant bSep08, mRNA.
<a href="#">RGD1559494</a>	<a href="#">RGD1559494.cSep08</a>	<a href="#">499535</a>	5684	955	2	31	RGD1559494 (3.6 kD) (RGD1559494) alternative variant cSep08, mRNA.
<a href="#">RGD1559496</a>	<a href="#">RGD1559496.bSep08</a>	<a href="#">292101</a>	4176	1171	3	390	similar to hypothetical protein (RGD1559496) alternative variant bSep08, mRNA.
<a href="#">RGD1559496</a>	<a href="#">RGD1559496.cSep08</a>	<a href="#">292101</a>	3473	716	3	161	similar to hypothetical protein (RGD1559496) alternative variant cSep08, mRNA.
<a href="#">RGD1559497</a>	<a href="#">RGD1559497.aSep08</a>	<a href="#">303684</a>	11668	1194	1	256	similar to tripartite motif-containing 65 (28.5 kD) (RGD1559497) alternative variant aSep08, mRNA.
<a href="#">RGD1559497</a>	<a href="#">RGD1559497.bSep08</a>	<a href="#">303684</a>	3178	397	2	131	similar to tripartite motif-containing 65 (RGD1559497) alternative variant bSep08, mRNA.

<a href="#">RGD1559530</a>	<a href="#">RGD1559530.aSep08</a>	<a href="#">308653</a>	6740	364		108	similar to nuclear RNA export factor 2 (RGD1559530) mRNA.
<a href="#">RGD1559536</a>	<a href="#">RGD1559536.aSep08</a>	<a href="#">498855</a>	3373	353		113	similar to vitellogenin-like 1 precursor (RGD1559536) mRNA.
<a href="#">RGD1559548</a>	<a href="#">RGD1559548.aSep08</a>	<a href="#">501199</a>	296856	2322		617	similar to Tes13-L (71.9 kD) (RGD1559548) mRNA.
<a href="#">RGD1559575</a>	<a href="#">RGD1559575.aSep08</a>	<a href="#">287231</a>	15350	1540		401	similar to novel protein (RGD1559575) mRNA.
<a href="#">RGD1559586</a>	<a href="#">RGD1559586.aSep08</a>	<a href="#">502112</a>	3631	838		66	similar to cathepsin Q2 (RGD1559586) mRNA.
<a href="#">RGD1559599</a>	<a href="#">RGD1559599.aSep08</a>	<a href="#">317624</a>	997	359		96	putative protein of vertebrate origin (RGD1559599) mRNA.
<a href="#">RGD1559600</a>	<a href="#">RGD1559600.aSep08</a>	<a href="#">499256</a>	12307	1120	1	169	RGD1559600 (RGD1559600) alternative variant aSep08, mRNA.
<a href="#">RGD1559604</a>	<a href="#">RGD1559604.aSep08</a>	<a href="#">498790</a>	10502	1207		323	similar to protein RAKd (37.0 kD) (RGD1559604) mRNA.
<a href="#">RGD1559610</a>	<a href="#">RGD1559610.bSep08</a>	<a href="#">313581</a>	1330	637	1	29	similar to CGI-94 protein (RGD1559610) alternative variant bSep08, mRNA.
<a href="#">RGD1559613</a>	<a href="#">RGD1559613.bSep08</a>	<a href="#">499067</a>	17993	671	1	109	RGD1559613 (12.3 kD) (RGD1559613) alternative variant bSep08, mRNA.
<a href="#">RGD1559640andRGD1562674</a>	<a href="#">RGD1559640andRGD1562674.aSep08</a>	<a href="#">288691</a>	303310	1798	6	553	kinase suppressor of ras 2 CRA a (RGD1559640andRGD1562674) alternative variant aSep08, mRNA.
<a href="#">RGD1559640andRGD1562674</a>	<a href="#">RGD1559640andRGD1562674.aSep08</a>	<a href="#">501840</a>	303310	1798	6	553	kinase suppressor of ras 2 CRA a (RGD1559640andRGD1562674) alternative variant aSep08, mRNA.
<a href="#">RGD1559643</a>	<a href="#">RGD1559643.bSep08</a>	<a href="#">498100</a>	42086	326	1	52	similar to hypothetical protein A430031N04 (RGD1559643) alternative variant bSep08, mRNA.
<a href="#">RGD1559651</a>	<a href="#">RGD1559651.aSep08</a>	<a href="#">500637</a>	18520	609		175	similar to Ab2-162 (RGD1559651) mRNA.
<a href="#">RGD1559683</a>	<a href="#">RGD1559683.bSep08</a>	<a href="#">500620</a>	10941	751	1	56	similar to RIKEN cDNA 1700001C02 (RGD1559683) alternative variant bSep08, mRNA.
<a href="#">RGD1559695</a>	<a href="#">RGD1559695.aSep08</a>	<a href="#">500889</a>	4123	643		213	similar to FLJ43860 protein (RGD1559695) mRNA.
<a href="#">RGD1559696</a>	<a href="#">RGD1559696.aSep08</a>	<a href="#">289309</a>	9923	730		243	similar to kinesin-like protein (103.5 kD) (klp-6) (RGD1559696) mRNA.
<a href="#">RGD1559709</a>	<a href="#">RGD1559709.aSep08</a>	<a href="#">360615</a>	1573	729		92	similar to F-box protein 47 (RGD1559709) mRNA.
<a href="#">RGD1559710</a>	<a href="#">RGD1559710.aSep08</a>	<a href="#">288386</a>	4325	387		129	similar to DC-SIGN (RGD1559710) mRNA.
<a href="#">RGD1559732</a>	<a href="#">RGD1559732.aSep08</a>	<a href="#">497912</a>	1008	489		93	similar to butyrophilin related 1 (10.7 kD) (RGD1559732) mRNA.
<a href="#">RGD1559747</a>	<a href="#">RGD1559747.bSep08</a>	<a href="#">312310</a>	1708	494	2	103	zinc finger, C2H2-type (RGD1559747) alternative variant bSep08, mRNA.
<a href="#">RGD1559748</a>	<a href="#">RGD1559748.bSep08</a>	<a href="#">311559</a>	1486	614	1	35	similar to Palate lung and nasal carcinoma-like protein precursor (Tongue plunc-like protein) (RGD1559748) alternative variant bSep08, mRNA.
<a href="#">RGD1559786</a>	<a href="#">RGD1559786.bSep08</a>	<a href="#">298384</a>	2534	387	5	92	similar to RIKEN cDNA 0610037L13 (10.4 kD) (RGD1559786) alternative variant bSep08, mRNA.
<a href="#">RGD1559811</a>	<a href="#">RGD1559811.aSep08</a>	<a href="#">498657</a>	4156	1869		223	similar to RIKEN cDNA 1700112P19 (27.6 kD) (RGD1559811) mRNA.
<a href="#">RGD1559812</a>	<a href="#">RGD1559812.bSep08</a>	<a href="#">297832</a>	45278	384	1	120	similar to contactin associated protein-like 5 isoform 1 (RGD1559812) alternative variant bSep08, mRNA.
<a href="#">RGD1559841</a>	<a href="#">RGD1559841.aSep08</a>	<a href="#">307816</a>	139946	1515	4	318	similar to expressed sequence AW413431 (RGD1559841) alternative variant aSep08, mRNA.

<a href="#">RGD1559841</a>	<a href="#">RGD1559841.bSep08</a>	<a href="#">307816</a>	43589	618	3	205	similar to expressed sequence AW413431 (RGD1559841) alternative variant bSep08, mRNA.
<a href="#">RGD1559856</a>	<a href="#">RGD1559856.aSep08</a>	<a href="#">291618</a>	89086	1103		178	CRA a (19.9 kD) (RGD1559856) complete mRNA.
<a href="#">RGD1559859</a>	<a href="#">RGD1559859.aSep08</a>	<a href="#">363244</a>	9891	688		130	RGD1559859 (RGD1559859) mRNA.
<a href="#">RGD1559862</a>	<a href="#">RGD1559862.aSep08</a>	<a href="#">499189</a>	20162	720		239	similar to hypothetical protein from EUROIMAGE 384293 (RGD1559862) mRNA.
<a href="#">RGD1559864</a>	<a href="#">RGD1559864.aSep08</a>	<a href="#">500446</a>	7766	3037	4	337	similar to mKIAA1045 protein (RGD1559864) alternative variant aSep08, mRNA.
<a href="#">RGD1559864</a>	<a href="#">RGD1559864.bSep08</a>	<a href="#">500446</a>	1205	563	1	67	similar to mKIAA1045 protein (RGD1559864) alternative variant bSep08, mRNA.
<a href="#">RGD1559871</a>	<a href="#">RGD1559871.aSep08</a>	<a href="#">500135</a>	24542	1233		410	similar to Glycyl-tRNA synthetase (RGD1559871) mRNA.
<a href="#">RGD1559875</a>	<a href="#">RGD1559875.aSep08</a>	<a href="#">497970</a>	4757	766		232	similar to novel protein (RGD1559875) mRNA.
<a href="#">RGD1559879</a>	<a href="#">RGD1559879.aSep08</a>	<a href="#">309854</a>	12018	527		175	adenylate kinase (RGD1559879) mRNA.
<a href="#">RGD1559888and dPlxdc2</a>	<a href="#">RGD1559888andPlxdc2 .bSep08</a>	<a href="#">361282</a>	65640	575	6	191	similar to apical early endosomal glycoprotein (RGD1559888andPlxdc2) alternative variant bSep08, mRNA.
<a href="#">RGD1559888and dPlxdc2</a>	<a href="#">RGD1559888andPlxdc2 .bSep08</a>	<a href="#">502147</a>	65640	575	6	191	similar to apical early endosomal glycoprotein (RGD1559888andPlxdc2) alternative variant bSep08, mRNA.
<a href="#">RGD1559904</a>	<a href="#">RGD1559904.bSep08</a>	<a href="#">313061</a>	32859	1942	6	610	CRA b (RGD1559904) alternative variant bSep08, mRNA.
<a href="#">RGD1559904</a>	<a href="#">RGD1559904.cSep08</a>	<a href="#">313061</a>	5285	1384	4	256	CRA b (RGD1559904) alternative variant cSep08, mRNA.
<a href="#">RGD1559904</a>	<a href="#">RGD1559904.dSep08</a>	<a href="#">313061</a>	63665	651	5	181	CRA b (RGD1559904) alternative variant dSep08, mRNA.
<a href="#">RGD1559904</a>	<a href="#">RGD1559904.eSep08</a>	<a href="#">313061</a>	2017	1279	2	87	putative protein of mammalian origin (RGD1559904) alternative variant eSep08, mRNA.
<a href="#">RGD1559904</a>	<a href="#">RGD1559904.fSep08</a>	<a href="#">313061</a>	1879	264	2	65	CRA b (RGD1559904) alternative variant fSep08, mRNA.
<a href="#">RGD1559909</a>	<a href="#">RGD1559909.aSep08</a>	<a href="#">362592</a>	1343	872	2	265	RGD1559909 (RGD1559909) alternative variant aSep08, mRNA.
<a href="#">RGD1559909</a>	<a href="#">RGD1559909.bSep08</a>	<a href="#">362592</a>	817	565	1	166	RGD1559909 (RGD1559909) alternative variant bSep08, mRNA.
<a href="#">RGD1559923</a>	<a href="#">RGD1559923.bSep08</a>	<a href="#">498489</a>	16059	589	2	104	N-acetyltransferase 12 (12.0 kD) (RGD1559923) alternative variant bSep08, mRNA.
<a href="#">RGD1559930</a>	<a href="#">RGD1559930.aSep08</a>	<a href="#">296115</a>	19764	5183	9	628	similar to mKIAA0256 protein (RGD1559930) alternative variant aSep08, mRNA.
<a href="#">RGD1559930</a>	<a href="#">RGD1559930.bSep08</a>	<a href="#">296115</a>	7300	616	5	205	similar to mKIAA0256 protein (RGD1559930) alternative variant bSep08, mRNA.
<a href="#">RGD1559933</a>	<a href="#">RGD1559933.aSep08</a>	<a href="#">498584</a>	6965	471		122	RGD1559933 (RGD1559933) mRNA.
<a href="#">RGD1559942</a>	<a href="#">RGD1559942.bSep08</a>	<a href="#">363782</a>	18054	781	1	226	similar to hypothetical protein (RGD1559942) alternative variant bSep08, mRNA.
<a href="#">RGD1559954</a>	<a href="#">RGD1559954.aSep08</a>	<a href="#">304353</a>	7110	698		232	similar to family with sequence similarity 55, member C (RGD1559954) mRNA.
<a href="#">RGD1559958</a>	<a href="#">RGD1559958.aSep08</a>	<a href="#">362058</a>	10947	1344		189	similar to RIKEN cDNA C030011O14 gene (RGD1559958) mRNA.
<a href="#">RGD1559961</a>	<a href="#">RGD1559961.aSep08</a>	<a href="#">497974</a>	13935	1207	11	294	similar to novel protein (RGD1559961) alternative variant aSep08, mRNA.
<a href="#">RGD1559961</a>	<a href="#">RGD1559961.bSep08</a>	<a href="#">497974</a>	4146	421	2	116	similar to novel protein (RGD1559961) alternative variant bSep08, mRNA.



<a href="#">RGD1559961</a>	<a href="#">RGD1559961.cSep08</a>	<a href="#">497974</a>	1540	421	2	45	similar to novel protein (RGD1559961) alternative variant cSep08, mRNA.
<a href="#">RGD1559980</a>	<a href="#">RGD1559980.aSep08</a>	<a href="#">499158</a>	3900	363		72	RGD1559980 (RGD1559980) mRNA.
<a href="#">RGD1559984</a>	<a href="#">RGD1559984.aSep08</a>	<a href="#">499399</a>	5454	423		140	RGD1559984 (RGD1559984) mRNA.
<a href="#">RGD1559998</a>	<a href="#">RGD1559998.aSep08</a>	<a href="#">287131</a>	1066	710		112	similar to centrosomal protein 2 (RGD1559998) mRNA.
<a href="#">RGD1560011</a>	<a href="#">RGD1560011.aSep08</a>	<a href="#">315732</a>	309706	2951	4	895	similar to Nuclear membrane binding protein NUCLING (RGD1560011) alternative variant aSep08, mRNA.
<a href="#">RGD1560020_predicted</a>	<a href="#">RGD1560020_predicted.aSep08</a>	<a href="#">498982</a>	22076	2144	9	379	similar to Myb proto-oncogene protein (C-myb) (predicted) (RGD1560020_predicted) alternative variant aSep08, mRNA.
<a href="#">RGD1560020_predicted</a>	<a href="#">RGD1560020_predicted.bSep08</a>	<a href="#">498982</a>	8452	763	7	134	similar to Myb proto-oncogene protein (C-myb) (predicted) (RGD1560020_predicted) alternative variant bSep08, mRNA.
<a href="#">RGD1560020_predicted</a>	<a href="#">RGD1560020_predicted.dSep08</a>	<a href="#">498982</a>	1409	563	2	27	similar to Myb proto-oncogene protein (C-myb) (predicted) (RGD1560020_predicted) alternative variant dSep08, mRNA.
<a href="#">RGD1560065</a>	<a href="#">RGD1560065.aSep08</a>	<a href="#">499724</a>	6603	1518	3	184	similar to RIKEN cDNA 2410004B18 (RGD1560065) alternative variant aSep08, mRNA.
<a href="#">RGD1560065</a>	<a href="#">RGD1560065.bSep08</a>	<a href="#">499724</a>	5506	738	3	83	similar to RIKEN cDNA 2410004B18 (10.2 kD) (RGD1560065) alternative variant bSep08, mRNA.
<a href="#">RGD1560070</a>	<a href="#">RGD1560070.bSep08</a>	<a href="#">287847</a>	14412	668	9	181	similar to ataxin 2-binding protein 1 isoform 2 (RGD1560070) alternative variant bSep08, mRNA.
<a href="#">RGD1560070</a>	<a href="#">RGD1560070.eSep08</a>	<a href="#">287847</a>	9969	484	2	51	similar to ataxin 2-binding protein 1 isoform 2 (5.3 kD) (RGD1560070) alternative variant eSep08, mRNA.
<a href="#">RGD1560071</a>	<a href="#">RGD1560071.aSep08</a>	<a href="#">498691</a>	4582	1307		330	similar to Cathepsin L-like (36.8 kD) (RGD1560071) mRNA.
<a href="#">RGD1560112</a>	<a href="#">RGD1560112.aSep08</a>	<a href="#">363044</a>	3623	531		106	similar to Urinary protein 2 precursor (RUP-2) (RGD1560112) mRNA.
<a href="#">RGD1560125</a>	<a href="#">RGD1560125.aSep08</a>	<a href="#">360825</a>	6399	523		173	similar to OTTHUMP00000028561 (RGD1560125) mRNA.
<a href="#">RGD1560137</a>	<a href="#">RGD1560137.aSep08</a>	<a href="#">290372</a>	11575	554	3	123	similar to expressed sequence AU021034 (14.2 kD) (RGD1560137) alternative variant aSep08, mRNA.
<a href="#">RGD1560151</a>	<a href="#">RGD1560151.aSep08</a>	<a href="#">501105</a>	10075	983	2	153	similar to predicted CDS, mechanosensory transduction channel NOMPC (1O503) (RGD1560151) mRNA.
<a href="#">RGD1560155</a>	<a href="#">RGD1560155.bSep08</a>	<a href="#">307067</a>	254727	774	5	208	similar to mKIAA0934 protein (23.2 kD) (RGD1560155) alternative variant bSep08, mRNA.
<a href="#">RGD1560155</a>	<a href="#">RGD1560155.cSep08</a>	<a href="#">307067</a>	358131	419	3	103	similar to mKIAA0934 protein (RGD1560155) alternative variant cSep08, mRNA.
<a href="#">RGD1560174</a>	<a href="#">RGD1560174.aSep08</a>	<a href="#">501051</a>	1861	508		164	similar to IQ motif containing F4 (RGD1560174) mRNA.
<a href="#">RGD1560175</a>	<a href="#">RGD1560175.aSep08</a>	<a href="#">303946</a>	3091	405	2	135	similar to hypothetical protein KIAA2018 (RGD1560175) alternative variant aSep08, mRNA.
<a href="#">RGD1560182</a>	<a href="#">RGD1560182.aSep08</a>	<a href="#">362480</a>	22122	640		126	similar to lipoxygenase homology domains 1 (RGD1560182) mRNA.
<a href="#">RGD1560187</a>	<a href="#">RGD1560187.aSep08</a>	<a href="#">362641</a>	26333	600	5	116	similar to Hypothetical UPF0327 protein (12.7 kD) (RGD1560187) alternative variant aSep08, mRNA.
<a href="#">RGD1560187</a>	<a href="#">RGD1560187.bSep08</a>	<a href="#">362641</a>	1620	860	2	99	similar to Hypothetical UPF0327 protein (RGD1560187) alternative variant bSep08, mRNA.

<a href="#">RGD1560187</a>	<a href="#">RGD1560187.cSep08</a>	<a href="#">362641</a>	26274	504	4	76	similar to Hypothetical UPF0327 protein (8.5 kD) (RGD1560187) alternative variant cSep08, complete mRNA.
<a href="#">RGD1560187</a>	<a href="#">RGD1560187.dSep08</a>	<a href="#">362641</a>	2486	714	3	48	similar to Hypothetical UPF0327 protein (5.6 kD) (RGD1560187) alternative variant dSep08, mRNA.
<a href="#">RGD1560205</a>	<a href="#">RGD1560205.aSep08</a>	<a href="#">310548</a>	30063	948		131	similar to hypothetical protein MGC27016 (RGD1560205) mRNA.
<a href="#">RGD1560210</a>	<a href="#">RGD1560210.aSep08</a>	<a href="#">295622</a>	8428	351		95	RGD1560210 (10.9 kD) (RGD1560210) mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.aSep08</a>	<a href="#">498890</a>	6392	1537	3	309	N-acetyltransferase ARD1 (RGD1560212) alternative variant aSep08, mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.bSep08</a>	<a href="#">498890</a>	15784	1271	6	220	N-acetyltransferase ARD1 (25.8 kD) (RGD1560212) alternative variant bSep08, complete mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.cSep08</a>	<a href="#">498890</a>	15774	719	4	142	thioredoxin-like 4A (16.8 kD) (RGD1560212) alternative variant cSep08, complete mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.dSep08</a>	<a href="#">498890</a>	15764	716	4	142	thioredoxin-like 4A (16.8 kD) (RGD1560212) alternative variant dSep08, complete mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.eSep08</a>	<a href="#">498890</a>	714	629	2	129	N-acetyltransferase ARD1 (RGD1560212) alternative variant eSep08, mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.fSep08</a>	<a href="#">498890</a>	6004	1225	4	112	putative protein (RGD1560212) alternative variant fSep08, mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.hSep08</a>	<a href="#">498890</a>	5467	739	4	101	putative protein (RGD1560212) alternative variant hSep08, mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.iSep08</a>	<a href="#">498890</a>	5491	705	4	100	putative protein (RGD1560212) alternative variant iSep08, mRNA.
<a href="#">RGD1560212</a>	<a href="#">RGD1560212.jSep08</a>	<a href="#">498890</a>	15738	752	5	98	putative protein (RGD1560212) alternative variant jSep08, mRNA.
<a href="#">RGD1560213</a>	<a href="#">RGD1560213.aSep08</a>	<a href="#">498189</a>	25719	760		253	similar to RIKEN cDNA 1500001A10 (RGD1560213) mRNA.
<a href="#">RGD1560214</a>	<a href="#">RGD1560214.aSep08</a>	<a href="#">498026</a>	889	679	2	150	similar to mKIAA1783 protein (RGD1560214) alternative variant aSep08, mRNA.
<a href="#">RGD1560214</a>	<a href="#">RGD1560214.bSep08</a>	<a href="#">498026</a>	714	399	3	133	similar to mKIAA1783 protein (RGD1560214) alternative variant bSep08, mRNA.
<a href="#">RGD1560220</a>	<a href="#">RGD1560220.aSep08</a>	<a href="#">502525</a>	4385	579	4	154	similar to homolog of yeast TIM14 isoform c (RGD1560220) alternative variant aSep08, mRNA.
<a href="#">RGD1560221</a>	<a href="#">RGD1560221.aSep08</a>	<a href="#">498780</a>	4672	815		224	similar to serine/threonine kinase (RGD1560221) mRNA.
<a href="#">RGD1560244</a>	<a href="#">RGD1560244.aSep08</a>	<a href="#">302996</a>	3898	1561	12	224	similar to hypothetical protein FLJ34512 (25.6 kD) (RGD1560244) alternative variant aSep08, mRNA.
<a href="#">RGD1560244</a>	<a href="#">RGD1560244.bSep08</a>	<a href="#">302996</a>	1268	508	4	132	similar to hypothetical protein FLJ34512 (RGD1560244) alternative variant bSep08, mRNA.
<a href="#">RGD1560248</a>	<a href="#">RGD1560248.aSep08</a>	<a href="#">499797</a>	18579	1664	10	446	similar to formin-like 2 isoform B (RGD1560248) alternative variant aSep08, mRNA.
<a href="#">RGD1560248</a>	<a href="#">RGD1560248.bSep08</a>	<a href="#">499797</a>	7967	2263	3	101	similar to formin-like 2 isoform B (RGD1560248) alternative variant bSep08, mRNA.
<a href="#">RGD1560248</a>	<a href="#">RGD1560248.cSep08</a>	<a href="#">499797</a>	5886	707	3	78	similar to formin-like 2 isoform B (RGD1560248) alternative variant cSep08, mRNA.
<a href="#">RGD1560248</a>	<a href="#">RGD1560248.dSep08</a>	<a href="#">499797</a>	5306	581	3	46	similar to formin-like 2 isoform B (RGD1560248) alternative variant dSep08, mRNA.

<a href="#">RGD1560257</a>	<a href="#">RGD1560257.aSep08</a>	<a href="#">499923</a>	10084	757		252	similar to hypothetical protein A630008I04 (RGD1560257) mRNA.
<a href="#">RGD1560269</a>	<a href="#">RGD1560269.aSep08</a>	<a href="#">289369</a>	35324	420		139	similar to usherin isoform B (RGD1560269) mRNA.
<a href="#">RGD1560271</a>	<a href="#">RGD1560271.bSep08</a>	<a href="#">501097</a>	16878	1539	3	247	similar to inhibitor of MyoD family-a (25.2 kD) (RGD1560271) alternative variant bSep08, complete mRNA.
<a href="#">RGD1560271</a>	<a href="#">RGD1560271.cSep08</a>	<a href="#">501097</a>	5143	763	1	110	similar to inhibitor of MyoD family-a (RGD1560271) alternative variant cSep08, mRNA.
<a href="#">RGD1560273</a>	<a href="#">RGD1560273.aSep08</a>	<a href="#">498405</a>	8107	1101	4	121	putative protein (12.9 kD) (RGD1560273) alternative variant aSep08, mRNA.
<a href="#">RGD1560273</a>	<a href="#">RGD1560273.bSep08</a>	<a href="#">498405</a>	1967	531	2	47	putative protein (RGD1560273) alternative variant bSep08, mRNA.
<a href="#">RGD1560286</a>	<a href="#">RGD1560286.cSep08</a>	<a href="#">500575</a>	4528	501	3	133	similar to DNA segment, Chr 4, ERATO Doi 22, expressed (RGD1560286) alternative variant cSep08, mRNA.
<a href="#">RGD1560286</a>	<a href="#">RGD1560286.dSep08</a>	<a href="#">500575</a>	4436	515	4	117	similar to DNA segment, Chr 4, ERATO Doi 22, expressed (RGD1560286) alternative variant dSep08, mRNA.
<a href="#">RGD1560286</a>	<a href="#">RGD1560286.eSep08</a>	<a href="#">500575</a>	24478	3337	5	107	similar to DNA segment, Chr 4, ERATO Doi 22, expressed (RGD1560286) alternative variant eSep08, mRNA.
<a href="#">RGD1560286</a>	<a href="#">RGD1560286.fSep08</a>	<a href="#">500575</a>	21937	391	2	48	similar to DNA segment, Chr 4, ERATO Doi 22, expressed (5.5 kD) (RGD1560286) alternative variant fSep08, mRNA.
<a href="#">RGD1560289</a>	<a href="#">RGD1560289.bSep08</a>	<a href="#">500258</a>	6706	978	6	325	putative protein of mammalian origin (RGD1560289) alternative variant bSep08, mRNA.
<a href="#">RGD1560289</a>	<a href="#">RGD1560289.cSep08</a>	<a href="#">500258</a>	11237	754	5	251	putative protein of metazoan origin (RGD1560289) alternative variant cSep08, mRNA.
<a href="#">RGD1560300</a>	<a href="#">RGD1560300.bSep08</a>	<a href="#">499360</a>	11918	3888	6	150	CRA b (RGD1560300) alternative variant bSep08, mRNA.
<a href="#">RGD1560314</a>	<a href="#">RGD1560314.aSep08</a>	<a href="#">362846</a>	5806	1690		266	RGD1560314 (RGD1560314) mRNA.
<a href="#">RGD1560328</a>	<a href="#">RGD1560328.aSep08</a>	<a href="#">499317</a>	3808	388	4	91	similar to UPF0197 protein C11orf10 homolog (RGD1560328) alternative variant aSep08, mRNA.
<a href="#">RGD1560364</a>	<a href="#">RGD1560364.aSep08</a>	<a href="#">363087</a>	13812	1252		417	similar to vacuolar protein sorting 13C protein (RGD1560364) mRNA.
<a href="#">RGD1560368</a>	<a href="#">RGD1560368.aSep08</a>	<a href="#">310544</a>	9773	1204		320	similar to hypothetical protein FLJ21159 (RGD1560368) mRNA.
<a href="#">RGD1560386</a>	<a href="#">RGD1560386.aSep08</a>	<a href="#">294521</a>	3254	702		140	similar to novel protein (RGD1560386) mRNA.
<a href="#">RGD1560391_predicted</a>	<a href="#">RGD1560391_predicted.aSep08</a>	<a href="#">499883</a>	42492	1712	3	383	similar to GA binding protein transcription factor, beta subunit 2 (GABPB2) (predicted) (41.4 kD) (RGD1560391_predicted) alternative variant aSep08, mRNA.
<a href="#">RGD1560391_predicted</a>	<a href="#">RGD1560391_predicted.cSep08</a>	<a href="#">499883</a>	11483	653	1	162	similar to GA binding protein transcription factor, beta subunit 2 (GABPB2) (predicted) (RGD1560391_predicted) alternative variant cSep08, mRNA.
<a href="#">RGD1560394</a>	<a href="#">RGD1560394.aSep08</a>	<a href="#">289728</a>	1606	478	2	143	RGD1560394 (RGD1560394) alternative variant aSep08, mRNA.
<a href="#">RGD1560398</a>	<a href="#">RGD1560398.bSep08</a>	<a href="#">498192</a>	4687	1251	2	81	RGD1560398 (RGD1560398) alternative variant bSep08, mRNA.
<a href="#">RGD1560398</a>	<a href="#">RGD1560398.cSep08</a>	<a href="#">498192</a>	4177	870	3	73	RGD1560398 (7.6 kD) (RGD1560398) alternative variant cSep08, complete mRNA.

<a href="#">RGD1560408</a>	<a href="#">RGD1560408.aSep08</a>	<a href="#">499166</a>	910	691		43	similar to Mannoside acetylglucosaminyltransferase 4, isoenzyme A (5.0 kD) (RGD1560408) mRNA.
<a href="#">RGD1560433</a>	<a href="#">RGD1560433.aSep08</a>	<a href="#">296750</a>	15958	1572		460	similar to 1500019C06Rik protein (RGD1560433) mRNA.
<a href="#">RGD1560436</a>	<a href="#">RGD1560436.bSep08</a>	<a href="#">500546</a>	3356	780	1	136	similar to hypothetical protein FLJ20508 (RGD1560436) alternative variant bSep08, mRNA.
<a href="#">RGD1560449andRGD1563545</a>	<a href="#">RGD1560449andRGD1563545.aSep08</a>	<a href="#">499004</a>	47302	1692	1	138	similar to nidogen 2 and similar to GTPase activating protein testicular GAP1 (RGD1560449andRGD1563545) alternative variant aSep08, mRNA.
<a href="#">RGD1560449andRGD1563545</a>	<a href="#">RGD1560449andRGD1563545.aSep08</a>	<a href="#">499005</a>	47302	1692	1	138	similar to nidogen 2 and similar to GTPase activating protein testicular GAP1 (RGD1560449andRGD1563545) alternative variant aSep08, mRNA.
<a href="#">RGD1560449andRGD1563545</a>	<a href="#">RGD1560449andRGD1563545.bSep08</a>	<a href="#">499004</a>	56946	565	3	45	similar to nidogen 2 and similar to GTPase activating protein testicular GAP1 (4.9 kD) (RGD1560449andRGD1563545) alternative variant bSep08, mRNA.
<a href="#">RGD1560449andRGD1563545</a>	<a href="#">RGD1560449andRGD1563545.bSep08</a>	<a href="#">499005</a>	56946	565	3	45	similar to nidogen 2 and similar to GTPase activating protein testicular GAP1 (4.9 kD) (RGD1560449andRGD1563545) alternative variant bSep08, mRNA.
<a href="#">RGD1560470</a>	<a href="#">RGD1560470.aSep08</a>	<a href="#">362083</a>	2686	1181		285	similar to Gene model 996 (RGD1560470) mRNA.
<a href="#">RGD1560481</a>	<a href="#">RGD1560481.bSep08</a>	<a href="#">500409</a>	21167	832	5	265	similar to hypothetical protein FLJ20171 (RGD1560481) alternative variant bSep08, mRNA.
<a href="#">RGD1560481</a>	<a href="#">RGD1560481.cSep08</a>	<a href="#">500409</a>	27758	1997	4	154	similar to hypothetical protein FLJ20171 (RGD1560481) alternative variant cSep08, mRNA.
<a href="#">RGD1560481</a>	<a href="#">RGD1560481.dSep08</a>	<a href="#">500409</a>	24613	895	4	137	similar to hypothetical protein FLJ20171 (RGD1560481) alternative variant dSep08, mRNA.
<a href="#">RGD1560492</a>	<a href="#">RGD1560492.aSep08</a>	<a href="#">361039</a>	1883	882		259	similar to leishmanolysin-like (metallopeptidase M8 family) (RGD1560492) mRNA.
<a href="#">RGD1560493</a>	<a href="#">RGD1560493.aSep08</a>	<a href="#">315365</a>	8460	489	1	162	putative protein, with a coiled coil domain, of vertebrate origin (RGD1560493) alternative variant aSep08, mRNA.
<a href="#">RGD1560493</a>	<a href="#">RGD1560493.bSep08</a>	<a href="#">315365</a>	9194	420	2	140	putative protein of vertebrate origin (RGD1560493) alternative variant bSep08, mRNA.
<a href="#">RGD1560511</a>	<a href="#">RGD1560511.aSep08</a>	<a href="#">306991</a>	169537	3026	28	853	similar to Vps41 protein (RGD1560511) alternative variant aSep08, mRNA.
<a href="#">RGD1560511</a>	<a href="#">RGD1560511.bSep08</a>	<a href="#">306991</a>	20159	743	8	247	similar to Vps41 protein (RGD1560511) alternative variant bSep08, mRNA.
<a href="#">RGD1560511</a>	<a href="#">RGD1560511.cSep08</a>	<a href="#">306991</a>	21081	656	6	218	similar to Vps41 protein (RGD1560511) alternative variant cSep08, mRNA.
<a href="#">RGD1560511</a>	<a href="#">RGD1560511.dSep08</a>	<a href="#">306991</a>	81082	799	7	164	similar to Vps41 protein (RGD1560511) alternative variant dSep08, mRNA.
<a href="#">RGD1560511</a>	<a href="#">RGD1560511.eSep08</a>	<a href="#">306991</a>	925	822	2	54	similar to Vps41 protein (6.2 kD) (RGD1560511) alternative variant eSep08, mRNA.
<a href="#">RGD1560516</a>	<a href="#">RGD1560516.aSep08</a>	<a href="#">499038</a>	11795	755		147	RGD1560516 (16.2 kD) (RGD1560516) mRNA.
<a href="#">RGD1560544</a>	<a href="#">RGD1560544.aSep08</a>	<a href="#">293689</a>	3710	2121	8	663	putative protein, with a coiled coil domain, of eukaryotic origin (RGD1560544) mRNA.
<a href="#">RGD1560557</a>	<a href="#">RGD1560557.aSep08</a>	<a href="#">499437</a>	7357	437		145	putative protein of eukaryotic origin (RGD1560557) mRNA.

<a href="#">RGD1560565</a>	<a href="#">RGD1560565.aSep08</a>	<a href="#">499287</a>	1798	725		159	similar to Tumor protein p53 inducible protein 5 (RGD1560565) mRNA.
<a href="#">RGD1560566</a>	<a href="#">RGD1560566.aSep08</a>	<a href="#">308986</a>	38333	1472	9	290	RGD1560566 (31.0 kD) (RGD1560566) alternative variant aSep08, mRNA.
<a href="#">RGD1560566</a>	<a href="#">RGD1560566.bSep08</a>	<a href="#">308986</a>	6284	929	3	221	RGD1560566 (RGD1560566) alternative variant bSep08, mRNA.
<a href="#">RGD1560566</a>	<a href="#">RGD1560566.cSep08</a>	<a href="#">308986</a>	32523	2512	9	184	RGD1560566 (19.9 kD) (RGD1560566) alternative variant cSep08, mRNA.
<a href="#">RGD1560566</a>	<a href="#">RGD1560566.fSep08</a>	<a href="#">308986</a>	27373	768	4	41	RGD1560566 (RGD1560566) alternative variant fSep08, mRNA.
<a href="#">RGD1560600</a>	<a href="#">RGD1560600.bSep08</a>	<a href="#">289810</a>	4601	994	4	141	similar to small unique nuclear receptor co-repressor (16.0 kD) (RGD1560600) alternative variant bSep08, mRNA.
<a href="#">RGD1560601</a>	<a href="#">RGD1560601.aSep08</a>	<a href="#">317432</a>	25905	859	7	281	similar to Jumonji/ARID domain-containing protein 1C (SmcX protein) (RGD1560601) alternative variant aSep08, mRNA.
<a href="#">RGD1560601</a>	<a href="#">RGD1560601.bSep08</a>	<a href="#">317432</a>	4007	537	2	76	similar to Jumonji/ARID domain-containing protein 1C (SmcX protein) (RGD1560601) alternative variant bSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.aSep08</a>	<a href="#">362166</a>	118883	1207	9	402	similar to PHF21A protein (RGD1560612) alternative variant aSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.bSep08</a>	<a href="#">362166</a>	146402	1894	11	388	similar to PHF21A protein (RGD1560612) alternative variant bSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.cSep08</a>	<a href="#">362166</a>	10255	2224	5	278	similar to PHF21A protein (RGD1560612) alternative variant cSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.dSep08</a>	<a href="#">362166</a>	3075	462	3	154	similar to PHF21A protein (RGD1560612) alternative variant dSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.eSep08</a>	<a href="#">362166</a>	2316	666	2	48	similar to PHF21A protein (RGD1560612) alternative variant eSep08, mRNA.
<a href="#">RGD1560612</a>	<a href="#">RGD1560612.gSep08</a>	<a href="#">362166</a>	1862	712	3	47	similar to PHF21A protein (RGD1560612) alternative variant gSep08, mRNA.
<a href="#">RGD1560620</a>	<a href="#">RGD1560620.aSep08</a>	<a href="#">291940</a>	42422	472		156	similar to hypothetical protein (RGD1560620) mRNA.
<a href="#">RGD1560622</a>	<a href="#">RGD1560622.aSep08</a>	<a href="#">501182</a>	5717	1138	2	114	RGD1560622 (12.8 kD) (RGD1560622) alternative variant aSep08, mRNA.
<a href="#">RGD1560622</a>	<a href="#">RGD1560622.bSep08</a>	<a href="#">501182</a>	471	381	1	61	RGD1560622 (RGD1560622) alternative variant bSep08, mRNA.
<a href="#">RGD1560626</a>	<a href="#">RGD1560626.aSep08</a>	<a href="#">500588</a>	5027	1064		130	RGD1560626 (13.9 kD) (RGD1560626) mRNA.
<a href="#">RGD1560629</a>	<a href="#">RGD1560629.aSep08</a>	<a href="#">499561</a>	11721	1124	2	245	similar to RIKEN cDNA A930016P21 (27.4 kD) (RGD1560629) alternative variant aSep08, mRNA.
<a href="#">RGD1560629</a>	<a href="#">RGD1560629.cSep08</a>	<a href="#">499561</a>	9261	667	2	70	similar to RIKEN cDNA A930016P21 (RGD1560629) alternative variant cSep08, mRNA.
<a href="#">RGD1560629</a>	<a href="#">RGD1560629.dSep08</a>	<a href="#">499561</a>	3171	347	2	65	similar to RIKEN cDNA A930016P21 (RGD1560629) alternative variant dSep08, mRNA.
<a href="#">RGD1560636</a>	<a href="#">RGD1560636.aSep08</a>	<a href="#">305467</a>	5698	1890	11	442	similar to novel protein (RGD1560636) alternative variant aSep08, mRNA.
<a href="#">RGD1560636</a>	<a href="#">RGD1560636.bSep08</a>	<a href="#">305467</a>	2230	1248	4	187	similar to novel protein (RGD1560636) alternative variant bSep08, mRNA.

<a href="#">RGD1560638</a>	<a href="#">RGD1560638.bSep08</a>	<a href="#">500233</a>	204292	2455	1	410	similar to Exocyst complex component Sec15B (48.2 kD) (RGD1560638) alternative variant bSep08, mRNA.
<a href="#">RGD1560638</a>	<a href="#">RGD1560638.cSep08</a>	<a href="#">500233</a>	42241	266	2	88	similar to Exocyst complex component Sec15B (RGD1560638) alternative variant cSep08, mRNA.
<a href="#">RGD1560652</a>	<a href="#">RGD1560652.aSep08</a>	<a href="#">500352</a>	3174	405		108	RGD1560652 (RGD1560652) mRNA.
<a href="#">RGD1560658</a>	<a href="#">RGD1560658.aSep08</a>	<a href="#">306891</a>	5233	942		313	similar to serine (or cysteine) proteinase inhibitor, clade B, member 1b (RGD1560658) mRNA.
<a href="#">RGD1560666</a>	<a href="#">RGD1560666.aSep08</a>	<a href="#">317439</a>	50190	753		250	similar to KIAA1280 protein (RGD1560666) mRNA.
<a href="#">RGD1560686</a>	<a href="#">RGD1560686.aSep08</a>	<a href="#">304297</a>	94710	2067	14	689	similar to sidekick 1 (RGD1560686) alternative variant aSep08, mRNA.
<a href="#">RGD1560686</a>	<a href="#">RGD1560686.bSep08</a>	<a href="#">304297</a>	3151	392	1	29	similar to sidekick 1 (RGD1560686) alternative variant bSep08, mRNA.
<a href="#">RGD1560700</a>	<a href="#">RGD1560700.aSep08</a>	<a href="#">362585</a>	4305	517		107	similar to palmitoyl-protein thioesterase (RGD1560700) mRNA.
<a href="#">RGD1560705.1</a>	<a href="#">RGD1560705.1.aSep08</a>	<a href="#">498550</a>	25696	551		117	similar to LRRGT00152 (RGD1560705.1) mRNA.
<a href="#">RGD1560717</a>	<a href="#">RGD1560717.bSep08</a>	<a href="#">363169</a>	8435	2082	4	226	similar to hypothetical protein DKFZp313N0621 (RGD1560717) alternative variant bSep08, mRNA.
<a href="#">RGD1560717</a>	<a href="#">RGD1560717.cSep08</a>	<a href="#">363169</a>	19890	661	6	126	similar to hypothetical protein DKFZp313N0621 (14.7 kD) (RGD1560717) alternative variant cSep08, mRNA.
<a href="#">RGD1560717</a>	<a href="#">RGD1560717.dSep08</a>	<a href="#">363169</a>	19639	696	6	126	similar to hypothetical protein DKFZp313N0621 (14.7 kD) (RGD1560717) alternative variant dSep08, mRNA.
<a href="#">RGD1560717</a>	<a href="#">RGD1560717.eSep08</a>	<a href="#">363169</a>	7567	516	4	65	similar to hypothetical protein DKFZp313N0621 (7.9 kD) (RGD1560717) alternative variant eSep08, complete mRNA.
<a href="#">RGD1560720</a>	<a href="#">RGD1560720.aSep08</a>	<a href="#">313311</a>	684	513		122	RGD1560720 (RGD1560720) mRNA.
<a href="#">RGD1560724</a>	<a href="#">RGD1560724.aSep08</a>	<a href="#">501127</a>	19289	1242	2	360	similar to FLJ42986 protein (RGD1560724) alternative variant aSep08, mRNA.
<a href="#">RGD1560736</a>	<a href="#">RGD1560736.aSep08</a>	<a href="#">363115</a>	68745	910		200	similar to solute carrier family 9 (sodium/hydrogen exchanger), isoform 9 (RGD1560736) mRNA.
<a href="#">RGD1560755</a>	<a href="#">RGD1560755.aSep08</a>	<a href="#">361165</a>	14624	1294		430	similar to D8Ertd354e protein (RGD1560755) alternative variant aSep08, mRNA.
<a href="#">RGD1560755</a>	<a href="#">RGD1560755.bSep08</a>	<a href="#">361165</a>	20908	1191		327	similar to D8Ertd354e protein (RGD1560755) alternative variant bSep08, mRNA.
<a href="#">RGD1560755</a>	<a href="#">RGD1560755.cSep08</a>	<a href="#">361165</a>	11785	2519		204	similar to D8Ertd354e protein (RGD1560755) alternative variant cSep08, mRNA.
<a href="#">RGD1560775</a>	<a href="#">RGD1560775.aSep08</a>	<a href="#">501031</a>	30197	747	2	229	similar to RIKEN cDNA 4930579C12 gene (RGD1560775) alternative variant aSep08, mRNA.
<a href="#">RGD1560775</a>	<a href="#">RGD1560775.bSep08</a>	<a href="#">501031</a>	31928	743	2	229	similar to RIKEN cDNA 4930579C12 gene (RGD1560775) alternative variant bSep08, mRNA.
<a href="#">RGD1560796</a>	<a href="#">RGD1560796.bSep08</a>	<a href="#">315798</a>	24359	2057	4	224	similar to suppressor of hairy wing homolog 4 isoform 1 (23.5 kD) (RGD1560796) alternative variant bSep08, mRNA.
<a href="#">RGD1560796</a>	<a href="#">RGD1560796.cSep08</a>	<a href="#">315798</a>	9714	544	4	181	similar to suppressor of hairy wing homolog 4 isoform 1 (RGD1560796) alternative variant cSep08, mRNA.
<a href="#">RGD1560803</a>	<a href="#">RGD1560803.aSep08</a>	<a href="#">501583</a>	675	498		118	RGD1560803 (RGD1560803) mRNA.

<a href="#">RGD1560846</a>	<a href="#">RGD1560846.bSep08</a>	<a href="#">498133</a>	684	271	1	62	similar to hypothetical protein MGC40178 (RGD1560846) alternative variant bSep08, mRNA.
<a href="#">RGD1560854</a>	<a href="#">RGD1560854.aSep08</a>	<a href="#">499106</a>	1577	560	3	171	similar to FLJ41131 protein (RGD1560854) alternative variant aSep08, mRNA.
<a href="#">RGD1560860</a>	<a href="#">RGD1560860.aSep08</a>	<a href="#">498767</a>	53538	1388	4	396	putative protein, with 4 coiled coil domains, of mammalian origin (RGD1560860) alternative variant aSep08, complete mRNA.
<a href="#">RGD1560860</a>	<a href="#">RGD1560860.bSep08</a>	<a href="#">498767</a>	45779	700	1	186	putative protein, with 2 coiled coil domains, of mammalian origin (RGD1560860) alternative variant bSep08, mRNA.
<a href="#">RGD1560873</a>	<a href="#">RGD1560873.aSep08</a>	<a href="#">306238</a>	7074	1528	5	467	similar to RIKEN cDNA E230015L20 gene (RGD1560873) alternative variant aSep08, mRNA.
<a href="#">RGD1560873</a>	<a href="#">RGD1560873.bSep08</a>	<a href="#">306238</a>	15024	1464	8	462	similar to RIKEN cDNA E230015L20 gene (RGD1560873) alternative variant bSep08, mRNA.
<a href="#">RGD1560873</a>	<a href="#">RGD1560873.cSep08</a>	<a href="#">306238</a>	8442	367	4	122	similar to RIKEN cDNA E230015L20 gene (RGD1560873) alternative variant cSep08, mRNA.
<a href="#">RGD1560902and dCldn7</a>	<a href="#">RGD1560902andCldn7.aSep08</a>	<a href="#">65132</a>	2204	1217		211	claudin 7 and similar to Cofilin, non-muscle isoform (Cofilin-1) (22.4 kD) (RGD1560902andCldn7) complete mRNA.
<a href="#">RGD1560902and dCldn7</a>	<a href="#">RGD1560902andCldn7.aSep08</a>	<a href="#">363635</a>	2204	1217		211	claudin 7 and similar to Cofilin, non-muscle isoform (Cofilin-1) (22.4 kD) (RGD1560902andCldn7) complete mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.aSep08</a>	<a href="#">361288</a>	4005	1082	5	198	FUN14 (RGD1560916) alternative variant aSep08, mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.bSep08</a>	<a href="#">361288</a>	5467	1564	6	196	FUN14 (21.6 kD) (RGD1560916) alternative variant bSep08, mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.cSep08</a>	<a href="#">361288</a>	5246	983	5	151	FUN14 (16.3 kD) (RGD1560916) alternative variant cSep08, mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.dSep08</a>	<a href="#">361288</a>	2933	755	3	138	FUN14 (14.9 kD) (RGD1560916) alternative variant dSep08, mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.eSep08</a>	<a href="#">361288</a>	2787	1420	3	115	FUN14 (RGD1560916) alternative variant eSep08, mRNA.
<a href="#">RGD1560916</a>	<a href="#">RGD1560916.fSep08</a>	<a href="#">361288</a>	1907	346	2	88	putative protein of vertebrate origin (RGD1560916) alternative variant fSep08, mRNA.
<a href="#">RGD1560925</a>	<a href="#">RGD1560925.aSep08</a>	<a href="#">501196</a>	109320	994		303	similar to 2610034M16Rik protein (RGD1560925) mRNA.
<a href="#">RGD1560927</a>	<a href="#">RGD1560927.bSep08</a>	<a href="#">501507</a>	1271	429	1	109	RGD1560927 (RGD1560927) alternative variant bSep08, mRNA.
<a href="#">RGD1560940</a>	<a href="#">RGD1560940.bSep08</a>	<a href="#">501060</a>	4514	1812	5	377	similar to testis specific serine proteinase 3 (RGD1560940) alternative variant bSep08, mRNA.
<a href="#">RGD1560958</a>	<a href="#">RGD1560958.aSep08</a>	<a href="#">499277</a>	1341	528		122	similar to RIKEN cDNA 1700063117 (RGD1560958) mRNA.
<a href="#">RGD1560978</a>	<a href="#">RGD1560978.aSep08</a>	<a href="#">500693</a>	16478	1061	5	353	similar to hypothetical protein and hypothetical protein LOC685784 (RGD1560978) alternative variant aSep08, mRNA.
<a href="#">RGD1560978</a>	<a href="#">RGD1560978.aSep08</a>	<a href="#">685784</a>	16478	1061	5	353	similar to hypothetical protein and hypothetical protein LOC685784 (RGD1560978) alternative variant aSep08, mRNA.
<a href="#">RGD1560978</a>	<a href="#">RGD1560978.cSep08</a>	<a href="#">500693</a>	3665	671	3	205	similar to hypothetical protein and hypothetical protein LOC685784 (RGD1560978) alternative variant cSep08, mRNA.
<a href="#">RGD1560978</a>	<a href="#">RGD1560978.cSep08</a>	<a href="#">685784</a>	3665	671	3	205	similar to hypothetical protein and hypothetical protein LOC685784 (RGD1560978) alternative variant cSep08, mRNA.

<a href="#">RGD1560986</a>	<a href="#">RGD1560986.aSep08</a>	<a href="#">292786</a>	3591	736		245	similar to Gene model 1082 (RGD1560986) mRNA.
<a href="#">RGD1560989</a>	<a href="#">RGD1560989.aSep08</a>	<a href="#">313060</a>	1206	377		125	putative protein of vertebrate origin (RGD1560989) mRNA.
<a href="#">RGD1561004</a>	<a href="#">RGD1561004.aSep08</a>	<a href="#">501539</a>	84929	374		124	similar to AMME syndrome candidate gene 1 protein homolog (RGD1561004) mRNA.
<a href="#">RGD1561014</a>	<a href="#">RGD1561014.aSep08</a>	<a href="#">497966</a>	4283	417		112	similar to T-complex protein 1, zeta-2 subunit (TCP-1-zeta-2) (RGD1561014) mRNA.
<a href="#">RGD1561023</a>	<a href="#">RGD1561023.aSep08</a>	<a href="#">498429</a>	6563	866		247	similar to RIKEN cDNA 4931440F15 gene (RGD1561023) mRNA.
<a href="#">RGD1561034</a>	<a href="#">RGD1561034.aSep08</a>	<a href="#">308908</a>	771	407		29	similar to hypothetical protein MGC34805 (RGD1561034) mRNA.
<a href="#">RGD1561039</a>	<a href="#">RGD1561039.bSep08</a>	<a href="#">498870</a>	3303	259	1	44	similar to RIKEN cDNA 1700065I17 (RGD1561039) alternative variant bSep08, mRNA.
<a href="#">RGD1561042</a>	<a href="#">RGD1561042.aSep08</a>	<a href="#">498386</a>	6352	558	4	185	similar to RIKEN cDNA 5730509K17 gene (RGD1561042) alternative variant aSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.aSep08</a>	<a href="#">498642</a>	134567	1176	8	241	similar to RNA binding protein gene with multiple splicing (26.4 kD) (RGD1561067) alternative variant aSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.bSep08</a>	<a href="#">498642</a>	130974	1999	7	230	similar to RNA binding protein gene with multiple splicing (26.3 kD) (RGD1561067) alternative variant bSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.cSep08</a>	<a href="#">498642</a>	65855	695	6	198	similar to RNA binding protein gene with multiple splicing (RGD1561067) alternative variant cSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.dSep08</a>	<a href="#">498642</a>	84869	760	8	175	similar to RNA binding protein gene with multiple splicing (RGD1561067) alternative variant dSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.eSep08</a>	<a href="#">498642</a>	84821	578	7	169	similar to RNA binding protein gene with multiple splicing (RGD1561067) alternative variant eSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.fSep08</a>	<a href="#">498642</a>	51931	426	4	98	similar to RNA binding protein gene with multiple splicing (RGD1561067) alternative variant fSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.gSep08</a>	<a href="#">498642</a>	64164	436	4	91	similar to RNA binding protein gene with multiple splicing (RGD1561067) alternative variant gSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.hSep08</a>	<a href="#">498642</a>	10900	652	3	52	similar to RNA binding protein gene with multiple splicing (5.4 kD) (RGD1561067) alternative variant hSep08, mRNA.
<a href="#">RGD1561067</a>	<a href="#">RGD1561067.jSep08</a>	<a href="#">498642</a>	12391	602	4	31	similar to RNA binding protein gene with multiple splicing (3.2 kD) (RGD1561067) alternative variant jSep08, mRNA.
<a href="#">RGD1561069</a>	<a href="#">RGD1561069.aSep08</a>	<a href="#">498959</a>	13671	430		143	similar to F-box only protein 31 (RGD1561069) mRNA.
<a href="#">RGD1561074</a>	<a href="#">RGD1561074.aSep08</a>	<a href="#">300884</a>	3850	633		210	similar to tripartite motif-containing 43 (RGD1561074) mRNA.
<a href="#">RGD1561090</a>	<a href="#">RGD1561090.aSep08</a>	<a href="#">313278</a>	80673	780		260	similar to protein tyrosine phosphatase, receptor type, D (RGD1561090) mRNA.
<a href="#">RGD1561111and Crct1</a>	<a href="#">RGD1561111andCrct1.aSep08</a>	<a href="#">310585</a>	43143	1159		23	similar to Eno1 protein and cysteine-rich C-terminal 1 (7.8 kD) (RGD1561111andCrct1) mRNA.
<a href="#">RGD1561111and Crct1</a>	<a href="#">RGD1561111andCrct1.aSep08</a>	<a href="#">688401</a>	43143	1159		23	similar to Eno1 protein and cysteine-rich C-terminal 1 (7.8 kD) (RGD1561111andCrct1) mRNA.
<a href="#">RGD1561113</a>	<a href="#">RGD1561113.aSep08</a>	<a href="#">499780</a>	2755	2220	1	133	similar to Hypothetical UPF0184 protein C9orf16 homolog (15.3 kD) (RGD1561113) alternative variant aSep08, mRNA.



<a href="#">RGD1561113</a>	<a href="#">RGD1561113.bSep08</a>	<a href="#">499780</a>	2784	651	1	125	similar to Hypothetical UPF0184 protein C9orf16 homolog (RGD1561113) alternative variant bSep08, mRNA.
<a href="#">RGD1561143</a>	<a href="#">RGD1561143.aSep08</a>	<a href="#">304382</a>	5146	684		121	similar to cell surface receptor FDFACT (RGD1561143) mRNA.
<a href="#">RGD1561145</a>	<a href="#">RGD1561145.bSep08</a>	<a href="#">498580</a>	32195	1042	4	109	similar to novel protein (RGD1561145) alternative variant bSep08, mRNA.
<a href="#">RGD1561145</a>	<a href="#">RGD1561145.cSep08</a>	<a href="#">498580</a>	21577	315	2	105	similar to novel protein (RGD1561145) alternative variant cSep08, mRNA.
<a href="#">RGD1561147</a>	<a href="#">RGD1561147.aSep08</a>	<a href="#">499468</a>	3026	722		71	similar to hypothetical protein FLJ37396 (RGD1561147) mRNA.
<a href="#">RGD1561157</a>	<a href="#">RGD1561157.bSep08</a>	<a href="#">360487</a>	762	601	2	103	RGD1561157 (11.5 kD) (RGD1561157) alternative variant bSep08, mRNA.
<a href="#">RGD1561161</a>	<a href="#">RGD1561161.aSep08</a>	<a href="#">294747</a>	2764	1640		546	similar to BC067074 protein (RGD1561161) mRNA.
<a href="#">RGD1561162</a>	<a href="#">RGD1561162.aSep08</a>	<a href="#">297962</a>	22743	585	3	195	similar to 35 kDa SR repressor protein (SRrp35) (RGD1561162) alternative variant aSep08, mRNA.
<a href="#">RGD1561162</a>	<a href="#">RGD1561162.bSep08</a>	<a href="#">297962</a>	5594	813	1	161	similar to 35 kDa SR repressor protein (SRrp35) (RGD1561162) alternative variant bSep08, mRNA.
<a href="#">RGD1561176</a>	<a href="#">RGD1561176.aSep08</a>	<a href="#">363155</a>	35229	1780	4	593	similar to Programmed cell death 6 interacting protein (ALG-2 interacting protein X) (RGD1561176) mRNA.
<a href="#">RGD1561196_predicted</a>	<a href="#">RGD1561196_predicted_aSep08</a>	<a href="#">498560</a>	5405	520		82	similar to ubiquitin carboxyl-terminal hydrolase l3 (predicted) (9.3 kD) (RGD1561196_predicted) mRNA.
<a href="#">RGD1561200</a>	<a href="#">RGD1561200.aSep08</a>	<a href="#">500230</a>	6064	1714		510	similar to hypothetical protein FLJ12056 (57.5 kD) (RGD1561200) mRNA.
<a href="#">RGD1561205</a>	<a href="#">RGD1561205.aSep08</a>	<a href="#">500557</a>	2109	753	2	122	similar to RIKEN cDNA 2610200G18 (13.4 kD) (RGD1561205) alternative variant aSep08, mRNA.
<a href="#">RGD1561205</a>	<a href="#">RGD1561205.bSep08</a>	<a href="#">500557</a>	7627	2659	4	117	similar to RIKEN cDNA 2610200G18 (13.3 kD) (RGD1561205) alternative variant bSep08, mRNA.
<a href="#">RGD1561205</a>	<a href="#">RGD1561205.cSep08</a>	<a href="#">500557</a>	2828	859	4	74	similar to RIKEN cDNA 2610200G18 (RGD1561205) alternative variant cSep08, mRNA.
<a href="#">RGD1561238</a>	<a href="#">RGD1561238.aSep08</a>	<a href="#">502091</a>	18247	748	1	248	similar to ring finger protein 122 homolog (RGD1561238) alternative variant aSep08, mRNA.
<a href="#">RGD1561238</a>	<a href="#">RGD1561238.bSep08</a>	<a href="#">502091</a>	18575	1079	1	218	similar to ring finger protein 122 homolog (RGD1561238) alternative variant bSep08, mRNA.
<a href="#">RGD1561238</a>	<a href="#">RGD1561238.cSep08</a>	<a href="#">502091</a>	15353	734	2	178	similar to ring finger protein 122 homolog (RGD1561238) alternative variant cSep08, mRNA.
<a href="#">RGD1561238</a>	<a href="#">RGD1561238.dSep08</a>	<a href="#">502091</a>	4926	1229	1	95	similar to ring finger protein 122 homolog (RGD1561238) alternative variant dSep08, mRNA.
<a href="#">RGD1561251</a>	<a href="#">RGD1561251.aSep08</a>	<a href="#">499352</a>	14603	413		118	RGD1561251 (RGD1561251) mRNA.
<a href="#">RGD1561277</a>	<a href="#">RGD1561277.aSep08</a>	<a href="#">497911</a>	8029	427		104	RGD1561277 (RGD1561277) mRNA.
<a href="#">RGD1561282</a>	<a href="#">RGD1561282.bSep08</a>	<a href="#">499958</a>	827	647	2	167	RGD1561282 (RGD1561282) alternative variant bSep08, mRNA.
<a href="#">RGD1561303</a>	<a href="#">RGD1561303.aSep08</a>	<a href="#">304509</a>	4218	556		185	RGD1561303 (RGD1561303) mRNA.
<a href="#">RGD1561306</a>	<a href="#">RGD1561306.aSep08</a>	<a href="#">502904</a>	7180	419	2	61	similar to immunoreceptor Ly49si3 (RGD1561306) alternative variant aSep08, mRNA.
<a href="#">RGD1561306</a>	<a href="#">RGD1561306.bSep08</a>	<a href="#">502904</a>	1074	995	1	65	similar to immunoreceptor Ly49si3 (7.9 kD) (RGD1561306) alternative variant bSep08, mRNA.

<a href="#">RGD1561327</a>	<a href="#">RGD1561327.aSep08</a>	<a href="#">317593</a>	1783	1706		419	similar to melanoma antigen family A, 10 (49.2 kD) (RGD1561327) mRNA.
<a href="#">RGD1561339</a>	<a href="#">RGD1561339.aSep08</a>	<a href="#">502271</a>	1495	785		79	similar to putative protein kinase (RGD1561339) mRNA.
<a href="#">RGD1561347</a>	<a href="#">RGD1561347.bSep08</a>	<a href="#">362612</a>	16686	3358	6	468	similar to Sfrs4 protein (RGD1561347) alternative variant bSep08, mRNA.
<a href="#">RGD1561357</a>	<a href="#">RGD1561357.aSep08</a>	<a href="#">497798</a>	57954	1420		145	similar to LIM domain only 3 (16.6 kD) (RGD1561357) mRNA.
<a href="#">RGD1561367</a>	<a href="#">RGD1561367.aSep08</a>	<a href="#">363031</a>	2928	2077		105	similar to Anillin (12.5 kD) (RGD1561367) mRNA.
<a href="#">RGD1561388</a>	<a href="#">RGD1561388.aSep08</a>	<a href="#">499042</a>	4331	686		104	similar to Afadin (Af-6 protein) (11.6 kD) (RGD1561388) mRNA.
<a href="#">RGD1561393</a>	<a href="#">RGD1561393.aSep08</a>	<a href="#">361932</a>	6365	1262		420	similar to CG15133-PA (RGD1561393) mRNA.
<a href="#">RGD1561394</a>	<a href="#">RGD1561394.aSep08</a>	<a href="#">311743</a>	83912	864		170	similar to MLTK-beta (RGD1561394) mRNA.
<a href="#">RGD1561413</a>	<a href="#">RGD1561413.aSep08</a>	<a href="#">500585</a>	1293	377		125	similar to BC021442 protein (RGD1561413) mRNA.
<a href="#">RGD1561415</a>	<a href="#">RGD1561415.bSep08</a>	<a href="#">498943</a>	670	570	1	149	RGD1561415 (RGD1561415) alternative variant bSep08, mRNA.
<a href="#">RGD1561416</a>	<a href="#">RGD1561416.aSep08</a>	<a href="#">500536</a>	650	318		105	similar to novel protein (HT036) (RGD1561416) mRNA.
<a href="#">RGD1561425</a>	<a href="#">RGD1561425.bSep08</a>	<a href="#">301378</a>	3844	425	1	119	similar to RIKEN cDNA 4832428D23 gene (RGD1561425) alternative variant bSep08, mRNA.
<a href="#">RGD1561426</a>	<a href="#">RGD1561426.aSep08</a>	<a href="#">362633</a>	8379	395		131	RGD1561426 (RGD1561426) mRNA.
<a href="#">RGD1561442</a>	<a href="#">RGD1561442.aSep08</a>	<a href="#">311416</a>	18209	716	1	212	similar to Vinculin (Metavinculin) (RGD1561442) alternative variant aSep08, mRNA.
<a href="#">RGD1561442</a>	<a href="#">RGD1561442.bSep08</a>	<a href="#">311416</a>	18183	474		128	similar to Vinculin (Metavinculin) (RGD1561442) alternative variant bSep08, mRNA.
<a href="#">RGD1561444</a>	<a href="#">RGD1561444.aSep08</a>	<a href="#">315473</a>	13470	1233		259	similar to RIKEN cDNA 9530077C05 (RGD1561444) mRNA.
<a href="#">RGD1561445</a>	<a href="#">RGD1561445.aSep08</a>	<a href="#">298320</a>	4945	712		143	similar to novel protein (RGD1561445) mRNA.
<a href="#">RGD1561459</a>	<a href="#">RGD1561459.aSep08</a>	<a href="#">361606</a>	30643	703	1	124	similar to RIKEN cDNA 1810020D17 (13.5 kD) (RGD1561459) alternative variant aSep08, mRNA.
<a href="#">RGD1561459</a>	<a href="#">RGD1561459.cSep08</a>	<a href="#">361606</a>	30626	536		92	similar to RIKEN cDNA 1810020D17 (RGD1561459) alternative variant cSep08, mRNA.
<a href="#">RGD1561474</a>	<a href="#">RGD1561474.aSep08</a>	<a href="#">314824</a>	21059	943	8	314	similar to oxysterol-binding protein-like protein 8 isoform a (RGD1561474) alternative variant aSep08, mRNA.
<a href="#">RGD1561491</a>	<a href="#">RGD1561491.bSep08</a>	<a href="#">365362</a>	30586	1154	4	384	similar to ATP-binding cassette transporter sub-family A member 14 (RGD1561491) alternative variant bSep08, mRNA.
<a href="#">RGD1561494</a>	<a href="#">RGD1561494.bSep08</a>	<a href="#">500226</a>	3228	993	9	291	similar to D3Mm3e (RGD1561494) alternative variant bSep08, mRNA.
<a href="#">RGD1561494</a>	<a href="#">RGD1561494.cSep08</a>	<a href="#">500226</a>	1345	426	5	128	similar to D3Mm3e (RGD1561494) alternative variant cSep08, mRNA.
<a href="#">RGD1561494</a>	<a href="#">RGD1561494.dSep08</a>	<a href="#">500226</a>	1183	747	3	109	similar to D3Mm3e (RGD1561494) alternative variant dSep08, mRNA.
<a href="#">RGD1561494</a>	<a href="#">RGD1561494.eSep08</a>	<a href="#">500226</a>	599	240	2	79	similar to D3Mm3e (RGD1561494) alternative variant eSep08, mRNA.
<a href="#">RGD1561494</a>	<a href="#">RGD1561494.fSep08</a>	<a href="#">500226</a>	17371	494	2	42	similar to D3Mm3e (5.2 kD) (RGD1561494) alternative variant fSep08, mRNA.

<a href="#">RGD1561503</a>	<a href="#">RGD1561503.aSep08</a>	<a href="#">499536</a>	1982	1554		55	similar to hypothetical protein AN1443.2 (RGD1561503) mRNA.
<a href="#">RGD1561507</a>	<a href="#">RGD1561507.bSep08</a>	<a href="#">292078</a>	12307	562	1	105	similar to hypothetical protein FLJ31606 (12.0 kD) (RGD1561507) alternative variant bSep08, complete mRNA.
<a href="#">RGD1561507</a>	<a href="#">RGD1561507.cSep08</a>	<a href="#">292078</a>	5769	311	1	71	similar to hypothetical protein FLJ31606 (7.6 kD) (RGD1561507) alternative variant cSep08, mRNA.
<a href="#">RGD1561513</a>	<a href="#">RGD1561513.aSep08</a>	<a href="#">362232</a>	11561	283	3	94	similar to ventral prostate-specific protein (RGD1561513) alternative variant aSep08, mRNA.
<a href="#">RGD1561513</a>	<a href="#">RGD1561513.bSep08</a>	<a href="#">362232</a>	9004	735	3	91	similar to ventral prostate-specific protein (11.1 kD) (RGD1561513) alternative variant bSep08, mRNA.
<a href="#">RGD1561513</a>	<a href="#">RGD1561513.dSep08</a>	<a href="#">362232</a>	1408	632	2	32	similar to ventral prostate-specific protein (3.7 kD) (RGD1561513) alternative variant dSep08, mRNA.
<a href="#">RGD1561530</a>	<a href="#">RGD1561530.aSep08</a>	<a href="#">299637</a>	3536	1095	6	321	similar to Tle6 protein (RGD1561530) alternative variant aSep08, mRNA.
<a href="#">RGD1561537</a>	<a href="#">RGD1561537.aSep08</a>	<a href="#">361197</a>	30795	2168		445	similar to putative repair and recombination helicase RAD26L (RGD1561537) alternative variant aSep08, mRNA.
<a href="#">RGD1561537</a>	<a href="#">RGD1561537.bSep08</a>	<a href="#">361197</a>	62810	1971		434	similar to putative repair and recombination helicase RAD26L (RGD1561537) alternative variant bSep08, mRNA.
<a href="#">RGD1561551</a>	<a href="#">RGD1561551.aSep08</a>	<a href="#">500359</a>	8413	448	1	148	similar to Hypothetical protein MGC75664 (RGD1561551) alternative variant aSep08, mRNA.
<a href="#">RGD1561551</a>	<a href="#">RGD1561551.bSep08</a>	<a href="#">500359</a>	11825	269	1	89	similar to Hypothetical protein MGC75664 (RGD1561551) alternative variant bSep08, mRNA.
<a href="#">RGD1561560</a>	<a href="#">RGD1561560.aSep08</a>	<a href="#">502711</a>	2715	800	1	266	similar to Gpd1l protein (RGD1561560) alternative variant aSep08, mRNA.
<a href="#">RGD1561560</a>	<a href="#">RGD1561560.bSep08</a>	<a href="#">502711</a>	2524	818	1	224	similar to Gpd1l protein (RGD1561560) alternative variant bSep08, mRNA.
<a href="#">RGD1561574</a>	<a href="#">RGD1561574.aSep08</a>	<a href="#">501715</a>	1837	172		56	similar to Hypothetical protein MGC76322 (RGD1561574) mRNA.
<a href="#">RGD1561605</a>	<a href="#">RGD1561605.aSep08</a>	<a href="#">289717</a>	11288	3581		420	similar to hypothetical protein (48.6 kD) (RGD1561605) mRNA.
<a href="#">RGD1561609</a>	<a href="#">RGD1561609.aSep08</a>	<a href="#">306117</a>	6475	942		229	similar to TBC1 domain family member 4 (RGD1561609) mRNA.
<a href="#">RGD1561648</a>	<a href="#">RGD1561648.bSep08</a>	<a href="#">500841</a>	18617	707	4	160	RGD1561648 (RGD1561648) alternative variant bSep08, mRNA.
<a href="#">RGD1561648</a>	<a href="#">RGD1561648.cSep08</a>	<a href="#">500841</a>	8408	790	5	46	RGD1561648 (5.1 kD) (RGD1561648) alternative variant cSep08, mRNA.
<a href="#">RGD1561662</a>	<a href="#">RGD1561662.aSep08</a>	<a href="#">301232</a>	3741	3337		821	similar to Al661453 protein (RGD1561662) mRNA.
<a href="#">RGD1561665</a>	<a href="#">RGD1561665.aSep08</a>	<a href="#">499429</a>	13577	1182	8	307	similar to spatial-delta (RGD1561665) alternative variant aSep08, mRNA.
<a href="#">RGD1561665</a>	<a href="#">RGD1561665.bSep08</a>	<a href="#">499429</a>	13501	980	8	264	similar to spatial-delta (RGD1561665) alternative variant bSep08, mRNA.
<a href="#">RGD1561672</a>	<a href="#">RGD1561672.aSep08</a>	<a href="#">305610</a>	14323	456		151	similar to novel protein (RGD1561672) mRNA.
<a href="#">RGD1561676</a>	<a href="#">RGD1561676.aSep08</a>	<a href="#">307917</a>	33892	1800	7	526	putative protein of eukaryotic origin (RGD1561676) alternative variant aSep08, mRNA.

<a href="#">RGD1561676</a>	<a href="#">RGD1561676.bSep08</a>	<a href="#">307917</a>	28819	1694	9	307	kelch repeat containing protein and kelch (RGD1561676) alternative variant bSep08, mRNA.
<a href="#">RGD1561676</a>	<a href="#">RGD1561676.cSep08</a>	<a href="#">307917</a>	24646	742	6	246	kelch repeat containing protein and kelch (RGD1561676) alternative variant cSep08, mRNA.
<a href="#">RGD1561676</a>	<a href="#">RGD1561676.dSep08</a>	<a href="#">307917</a>	1530	536	3	105	CRA a (RGD1561676) alternative variant dSep08, mRNA.
<a href="#">RGD1561676</a>	<a href="#">RGD1561676.fSep08</a>	<a href="#">307917</a>	13551	419	4	47	putative protein of eukaryotic origin (RGD1561676) alternative variant fSep08, mRNA.
<a href="#">RGD1561678</a>	<a href="#">RGD1561678.aSep08</a>	<a href="#">298168</a>	14559	424		141	similar to Ten-m1 (RGD1561678) mRNA.
<a href="#">RGD1561693</a>	<a href="#">RGD1561693.aSep08</a>	<a href="#">501597</a>	6865	1155	9	385	similar to Gene model 784 (RGD1561693) alternative variant aSep08, mRNA.
<a href="#">RGD1561708</a>	<a href="#">RGD1561708.bSep08</a>	<a href="#">367838</a>	15315	736	2	157	hepatocellular carcinoma-associated antigen like (RGD1561708) alternative variant bSep08, mRNA.
<a href="#">RGD1561724</a>	<a href="#">RGD1561724.aSep08</a>	<a href="#">366608</a>	132786	1791		453	similar to mKIAA0716 protein (RGD1561724) alternative variant aSep08, mRNA.
<a href="#">RGD1561732</a>	<a href="#">RGD1561732.aSep08</a>	<a href="#">363554</a>	2914	1063		300	similar to hypothetical protein 4732467B22 (RGD1561732) mRNA.
<a href="#">RGD1561734</a>	<a href="#">RGD1561734.aSep08</a>	<a href="#">501201</a>	3840	1065		232	similar to KIAA1913 (RGD1561734) mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.bSep08</a>	<a href="#">287156</a>	2576	867	5	289	CRA a (RGD1561785andRhot2) alternative variant bSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.bSep08</a>	<a href="#">287157</a>	2576	867	5	289	CRA a (RGD1561785andRhot2) alternative variant bSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.cSep08</a>	<a href="#">287156</a>	2710	1402	9	232	ras homolog gene family member T2 CRA a (26.3 kD) (RGD1561785andRhot2) alternative variant cSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.cSep08</a>	<a href="#">287157</a>	2710	1402	9	232	ras homolog gene family member T2 CRA a (26.3 kD) (RGD1561785andRhot2) alternative variant cSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.dSep08</a>	<a href="#">287156</a>	513	386	2	128	CRA a like (RGD1561785andRhot2) alternative variant dSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.dSep08</a>	<a href="#">287157</a>	513	386	2	128	CRA a like (RGD1561785andRhot2) alternative variant dSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.eSep08</a>	<a href="#">287156</a>	2410	796	7	110	ras homolog gene family member T2 CRA c (12.3 kD) (RGD1561785andRhot2) alternative variant eSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.eSep08</a>	<a href="#">287157</a>	2410	796	7	110	ras homolog gene family member T2 CRA c (12.3 kD) (RGD1561785andRhot2) alternative variant eSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.gSep08</a>	<a href="#">287156</a>	1700	501	3	48	putative protein (RGD1561785andRhot2) alternative variant gSep08, mRNA.
<a href="#">RGD1561785andRhot2.dRhot2</a>	<a href="#">RGD1561785andRhot2.gSep08</a>	<a href="#">287157</a>	1700	501	3	48	putative protein (RGD1561785andRhot2) alternative variant gSep08, mRNA.
<a href="#">RGD1561796</a>	<a href="#">RGD1561796.aSep08</a>	<a href="#">360483</a>	5733	1465	3	139	RGD1561796 (14.8 kD) (RGD1561796) alternative variant aSep08, mRNA.
<a href="#">RGD1561796</a>	<a href="#">RGD1561796.bSep08</a>	<a href="#">360483</a>	1246	703	2	117	RGD1561796 (RGD1561796) alternative variant bSep08, mRNA.
<a href="#">RGD1561797</a>	<a href="#">RGD1561797.aSep08</a>	<a href="#">499655</a>	4760	1790		79	RGD1561797 (9.6 kD) (RGD1561797) mRNA.

<a href="#">RGD1561817</a>	<a href="#">RGD1561817.bSep08</a>	<a href="#">294917</a>	58731	1800	6	591	similar to Traf2 and NCK interacting kinase, splice variant 4 (RGD1561817) alternative variant bSep08, mRNA.
<a href="#">RGD1561817</a>	<a href="#">RGD1561817.cSep08</a>	<a href="#">294917</a>	12870	725	5	232	similar to Traf2 and NCK interacting kinase, splice variant 4 (RGD1561817) alternative variant cSep08, mRNA.
<a href="#">RGD1561817</a>	<a href="#">RGD1561817.dSep08</a>	<a href="#">294917</a>	8838	3187	5	192	similar to Traf2 and NCK interacting kinase, splice variant 4 (RGD1561817) alternative variant dSep08, mRNA.
<a href="#">RGD1561828</a>	<a href="#">RGD1561828.aSep08</a>	<a href="#">679595</a>	1202	541		139	RGD1561828 (RGD1561828) mRNA.
<a href="#">RGD1561852</a>	<a href="#">RGD1561852.bSep08</a>	<a href="#">499893</a>	3004	1218	3	79	similar to Protein C20orf29 (RGD1561852) alternative variant bSep08, mRNA.
<a href="#">RGD1561916</a>	<a href="#">RGD1561916.aSep08</a>	<a href="#">500441</a>	2742	683		127	similar to testes development-related NYD-SP22 isoform 1 (14.4 kD) (RGD1561916) mRNA.
<a href="#">RGD1561931</a>	<a href="#">RGD1561931.aSep08</a>	<a href="#">302396</a>	3790	1130	3	208	similar to KIAA2022 protein (RGD1561931) alternative variant aSep08, mRNA.
<a href="#">RGD1561932</a>	<a href="#">RGD1561932.aSep08</a>	<a href="#">287346</a>	9875	563		137	similar to novel protein (16.5 kD) (RGD1561932) mRNA.
<a href="#">RGD1561934</a>	<a href="#">RGD1561934.aSep08</a>	<a href="#">499940</a>	1065	465		105	similar to TP53-target gene 5 protein (TP53-inducible gene 5 protein) (RGD1561934) mRNA.
<a href="#">RGD1561940</a>	<a href="#">RGD1561940.aSep08</a>	<a href="#">364686</a>	19849	2347	6	288	RAN binding protein 9 like (RGD1561940) alternative variant aSep08, mRNA.
<a href="#">RGD1561940</a>	<a href="#">RGD1561940.bSep08</a>	<a href="#">364686</a>	72600	1647	10	273	RAN binding protein like (RGD1561940) alternative variant bSep08, mRNA.
<a href="#">RGD1561940</a>	<a href="#">RGD1561940.cSep08</a>	<a href="#">364686</a>	48677	448	4	149	RAN binding protein like (RGD1561940) alternative variant cSep08, mRNA.
<a href="#">RGD1561940</a>	<a href="#">RGD1561940.dSep08</a>	<a href="#">364686</a>	11271	753	3	97	RAN binding protein 9 like (RGD1561940) alternative variant dSep08, mRNA.
<a href="#">RGD1561955</a>	<a href="#">RGD1561955.aSep08</a>	<a href="#">361076</a>	24908	782	3	236	similar to diacylglycerol kinase eta (RGD1561955) alternative variant aSep08, mRNA.
<a href="#">RGD1561955</a>	<a href="#">RGD1561955.bSep08</a>	<a href="#">361076</a>	5703	401	1	110	similar to diacylglycerol kinase eta (RGD1561955) alternative variant bSep08, mRNA.
<a href="#">RGD1561958</a>	<a href="#">RGD1561958.bSep08</a>	<a href="#">363490</a>	43601	734	1	135	similar to RIKEN cDNA 2010106E10 (RGD1561958) alternative variant bSep08, mRNA.
<a href="#">RGD1561962</a>	<a href="#">RGD1561962.aSep08</a>	<a href="#">304201</a>	15543	2290	10	688	similar to ecotropic viral integration site 5-like (RGD1561962) alternative variant aSep08, mRNA.
<a href="#">RGD1561962</a>	<a href="#">RGD1561962.bSep08</a>	<a href="#">304201</a>	1615	1187	5	230	similar to ecotropic viral integration site 5-like (RGD1561962) alternative variant bSep08, mRNA.
<a href="#">RGD1561962</a>	<a href="#">RGD1561962.dSep08</a>	<a href="#">304201</a>	5899	219	2	52	similar to ecotropic viral integration site 5-like (6.1 kD) (RGD1561962) alternative variant dSep08, mRNA.
<a href="#">RGD1561962</a>	<a href="#">RGD1561962.eSep08</a>	<a href="#">304201</a>	456	365	2	48	similar to ecotropic viral integration site 5-like (RGD1561962) alternative variant eSep08, mRNA.
<a href="#">RGD1561963</a>	<a href="#">RGD1561963.aSep08</a>	<a href="#">301556</a>	26041	2032	13	527	similar to Dedicator of cytokinesis protein 10 (Protein zizimin 3) (RGD1561963) alternative variant aSep08, mRNA.
<a href="#">RGD1561963</a>	<a href="#">RGD1561963.bSep08</a>	<a href="#">301556</a>	7476	1186	4	183	similar to Dedicator of cytokinesis protein 10 (Protein zizimin 3) (RGD1561963) alternative variant bSep08, mRNA.
<a href="#">RGD1561963</a>	<a href="#">RGD1561963.cSep08</a>	<a href="#">301556</a>	6321	506	5	168	similar to Dedicator of cytokinesis protein 10 (Protein zizimin 3) (RGD1561963) alternative variant cSep08, mRNA.

<a href="#">RGD1561963</a>	<a href="#">RGD1561963.dSep08</a>	<a href="#">301556</a>	4446	809	3	115	similar to Deducator of cytokinesis protein 10 (Protein zizimin 3) (12.9 kD) (RGD1561963) alternative variant dSep08, mRNA.
<a href="#">RGD1561972</a>	<a href="#">RGD1561972.aSep08</a>	<a href="#">308581</a>	632	444		94	similar to hypothetical protein FLJ32658 (RGD1561972) mRNA.
<a href="#">RGD1561988</a>	<a href="#">RGD1561988.aSep08</a>	<a href="#">307449</a>	47104	551		108	similar to Colorectal mutant cancer protein (MCC protein) (RGD1561988) mRNA.
<a href="#">RGD1561996</a>	<a href="#">RGD1561996.aSep08</a>	<a href="#">300821</a>	15482	787	3	123	similar to 60S ribosomal protein L35 (RGD1561996) alternative variant aSep08, mRNA.
<a href="#">RGD1561996</a>	<a href="#">RGD1561996.bSep08</a>	<a href="#">300821</a>	4458	727	2	53	similar to 60S ribosomal protein L35 (RGD1561996) alternative variant bSep08, mRNA.
<a href="#">RGD1561997</a>	<a href="#">RGD1561997.aSep08</a>	<a href="#">317271</a>	69084	1579		316	similar to Smage-1 protein (34.3 kD) (RGD1561997) mRNA.
<a href="#">RGD1562010</a>	<a href="#">RGD1562010.aSep08</a>	<a href="#">500286</a>	3706	469		61	similar to Fls485 protein (RGD1562010) mRNA.
<a href="#">RGD1562011</a>	<a href="#">RGD1562011.aSep08</a>	<a href="#">499297</a>	3463	2014	2	301	similar to MAS-related G-protein coupled receptor, member G (33.9 kD) (RGD1562011) alternative variant aSep08, mRNA.
<a href="#">RGD1562011</a>	<a href="#">RGD1562011.bSep08</a>	<a href="#">499297</a>	1865	383	2	107	similar to MAS-related G-protein coupled receptor, member G (RGD1562011) alternative variant bSep08, mRNA.
<a href="#">RGD1562012</a>	<a href="#">RGD1562012.aSep08</a>	<a href="#">497962</a>	12953	776	1	93	RGD1562012 (RGD1562012) alternative variant aSep08, mRNA.
<a href="#">RGD1562012</a>	<a href="#">RGD1562012.bSep08</a>	<a href="#">497962</a>	12957	759		78	RGD1562012 (9.4 kD) (RGD1562012) alternative variant bSep08, complete mRNA.
<a href="#">RGD1562018</a>	<a href="#">RGD1562018.aSep08</a>	<a href="#">361030</a>	15213	1479	7	298	similar to Protein C14orf101 homolog (RGD1562018) alternative variant aSep08, mRNA.
<a href="#">RGD1562018</a>	<a href="#">RGD1562018.bSep08</a>	<a href="#">361030</a>	12864	614	5	204	similar to Protein C14orf101 homolog (RGD1562018) alternative variant bSep08, mRNA.
<a href="#">RGD1562018</a>	<a href="#">RGD1562018.cSep08</a>	<a href="#">361030</a>	20483	822	6	204	similar to Protein C14orf101 homolog (RGD1562018) alternative variant cSep08, mRNA.
<a href="#">RGD1562018</a>	<a href="#">RGD1562018.dSep08</a>	<a href="#">361030</a>	11333	663	5	179	similar to Protein C14orf101 homolog (RGD1562018) alternative variant dSep08, mRNA.
<a href="#">RGD1562018</a>	<a href="#">RGD1562018.eSep08</a>	<a href="#">361030</a>	3816	264	2	87	similar to Protein C14orf101 homolog (RGD1562018) alternative variant eSep08, mRNA.
<a href="#">RGD1562026</a>	<a href="#">RGD1562026.aSep08</a>	<a href="#">500074</a>	4354	738		127	RGD1562026 (RGD1562026) mRNA.
<a href="#">RGD1562029</a>	<a href="#">RGD1562029.aSep08</a>	<a href="#">501150</a>	46659	688		222	similar to KIAA2012 protein (RGD1562029) mRNA.
<a href="#">RGD1562037</a>	<a href="#">RGD1562037.aSep08</a>	<a href="#">498764</a>	4444	711		106	similar to OTTHUMP00000046255 (RGD1562037) mRNA.
<a href="#">RGD1562044</a>	<a href="#">RGD1562044.bSep08</a>	<a href="#">499068</a>	18915	1250	4	368	similar to Zfp583 protein (RGD1562044) alternative variant bSep08, mRNA.
<a href="#">RGD1562044</a>	<a href="#">RGD1562044.dSep08</a>	<a href="#">499068</a>	9592	405	4	68	similar to Zfp583 protein (RGD1562044) alternative variant dSep08, mRNA.
<a href="#">RGD1562071</a>	<a href="#">RGD1562071.aSep08</a>	<a href="#">499931</a>	4273	770		144	similar to Src-like adaptor protein-2 (RGD1562071) mRNA.
<a href="#">RGD1562080</a>	<a href="#">RGD1562080.aSep08</a>	<a href="#">498827</a>	879	728		175	similar to Hypothetical protein CBG10141 (RGD1562080) mRNA.
<a href="#">RGD1562084</a>	<a href="#">RGD1562084.bSep08</a>	<a href="#">501002</a>	1349	315	1	86	similar to expressed sequence AI118078 (RGD1562084) alternative variant bSep08, mRNA.
<a href="#">RGD1562091</a>	<a href="#">RGD1562091.bSep08</a>	<a href="#">292690</a>	1680	678	2	69	similar to expressed sequence C79127 (7.3 kD) (RGD1562091) alternative variant bSep08, mRNA.

<a href="#">RGD1562095_predicted</a>	<a href="#">RGD1562095_predicted_bSep08</a>	<a href="#">499818</a>	61043	541	3	39	similar to protein phosphatase 1, regulatory (inhibitor) subunit 1C (predicted) (RGD1562095_predicted) alternative variant bSep08, mRNA.
<a href="#">RGD1562097</a>	<a href="#">RGD1562097.aSep08</a>	<a href="#">498718</a>	15380	395	1	88	RGD1562097 (RGD1562097) alternative variant aSep08, mRNA.
<a href="#">RGD1562097</a>	<a href="#">RGD1562097.bSep08</a>	<a href="#">498718</a>	18262	328	1	77	RGD1562097 (RGD1562097) alternative variant bSep08, mRNA.
<a href="#">RGD1562099</a>	<a href="#">RGD1562099.aSep08</a>	<a href="#">290325</a>	5964	713		173	similar to putative protein product of HMFN0672 (RGD1562099) mRNA.
<a href="#">RGD1562101</a>	<a href="#">RGD1562101.aSep08</a>	<a href="#">294614</a>	13648	901	4	175	similar to very large G-protein coupled receptor 1 (18.8 kD) (RGD1562101) alternative variant aSep08, mRNA.
<a href="#">RGD1562101</a>	<a href="#">RGD1562101.bSep08</a>	<a href="#">294614</a>	55891	419	3	139	similar to very large G-protein coupled receptor 1 (RGD1562101) alternative variant bSep08, mRNA.
<a href="#">RGD1562107</a>	<a href="#">RGD1562107.aSep08</a>	<a href="#">363205</a>	7072	1775		510	similar to class-alpha glutathione S-transferase (RGD1562107) alternative variant aSep08, mRNA.
<a href="#">RGD1562114</a>	<a href="#">RGD1562114.bSep08</a>	<a href="#">500795</a>	1401	1012	1	89	RGD1562114 (10.4 kD) (RGD1562114) alternative variant bSep08, mRNA.
<a href="#">RGD1562135</a>	<a href="#">RGD1562135.aSep08</a>	<a href="#">499979</a>	28987	1842	1	95	RGD1562135 (RGD1562135) alternative variant aSep08, mRNA.
<a href="#">RGD1562135</a>	<a href="#">RGD1562135.bSep08</a>	<a href="#">499979</a>	18668	743	1	50	RGD1562135 (5.7 kD) (RGD1562135) alternative variant bSep08, mRNA.
<a href="#">RGD1562146</a>	<a href="#">RGD1562146.aSep08</a>	<a href="#">500612</a>	5310	667		87	RGD1562146 (9.5 kD) (RGD1562146) mRNA.
<a href="#">RGD1562161</a>	<a href="#">RGD1562161.cSep08</a>	<a href="#">501559</a>	14629	557	3	52	putative protein of mammalian origin (RGD1562161) alternative variant cSep08, mRNA.
<a href="#">RGD1562174</a>	<a href="#">RGD1562174.aSep08</a>	<a href="#">310538</a>	55721	446		148	similar to mKIAA1450 protein (RGD1562174) mRNA.
<a href="#">RGD1562181</a>	<a href="#">RGD1562181.aSep08</a>	<a href="#">498853</a>	77558	673		92	similar to ribosomal protein S2 (RGD1562181) mRNA.
<a href="#">RGD1562200</a>	<a href="#">RGD1562200.aSep08</a>	<a href="#">363471</a>	3523	1025	8	252	similar to GS2 gene (27.4 kD) (RGD1562200) alternative variant aSep08, mRNA.
<a href="#">RGD1562200</a>	<a href="#">RGD1562200.bSep08</a>	<a href="#">363471</a>	1034	401	2	110	similar to GS2 gene (12.2 kD) (RGD1562200) alternative variant bSep08, mRNA.
<a href="#">RGD1562200</a>	<a href="#">RGD1562200.cSep08</a>	<a href="#">363471</a>	1823	780	3	87	similar to GS2 gene (RGD1562200) alternative variant cSep08, mRNA.
<a href="#">RGD1562218</a>	<a href="#">RGD1562218.bSep08</a>	<a href="#">292100</a>	8454	735	4	245	similar to RIKEN cDNA 0610039J04 (RGD1562218) alternative variant bSep08, mRNA.
<a href="#">RGD1562230</a>	<a href="#">RGD1562230.aSep08</a>	<a href="#">361839</a>	717596	817	6	243	similar to catenin alpha 3 (RGD1562230) alternative variant aSep08, mRNA.
<a href="#">RGD1562230</a>	<a href="#">RGD1562230.bSep08</a>	<a href="#">361839</a>	95617	731	3	130	similar to catenin alpha 3 (RGD1562230) alternative variant bSep08, mRNA.
<a href="#">RGD1562236</a>	<a href="#">RGD1562236.aSep08</a>	<a href="#">315019</a>	2399	1886		310	similar to breast cancer membrane protein 101 (34.6 kD) (RGD1562236) mRNA.
<a href="#">RGD1562252</a>	<a href="#">RGD1562252.bSep08</a>	<a href="#">315646</a>	4684	2106	2	547	similar to hypothetical gene supported by AK085276 (RGD1562252) alternative variant bSep08, mRNA.
<a href="#">RGD1562263</a>	<a href="#">RGD1562263.aSep08</a>	<a href="#">295092</a>	71222	773		222	similar to GTPase activating protein testicular GAP1 and similar to GTPase activating protein testicular GAP1 (RGD1562263) mRNA.

<a href="#">RGD1562263</a>	<a href="#">RGD1562263.aSep08</a>	<a href="#">691033</a>	71222	773		222	similar to GTPase activating protein testicular GAP1 and similar to GTPase activating protein testicular GAP1 (RGD1562263) mRNA.
<a href="#">RGD1562276</a>	<a href="#">RGD1562276.aSep08</a>	<a href="#">360567</a>	17527	441		97	similar to novel protein (RGD1562276) mRNA.
<a href="#">RGD1562284</a>	<a href="#">RGD1562284.aSep08</a>	<a href="#">313837</a>	32237	1795	2	447	similar to Glutaminy-peptide cyclotransferase precursor (QC) (RGD1562284) alternative variant aSep08, complete mRNA.
<a href="#">RGD1562284</a>	<a href="#">RGD1562284.cSep08</a>	<a href="#">313837</a>	24576	740	1	246	similar to Glutaminy-peptide cyclotransferase precursor (QC) (RGD1562284) alternative variant cSep08, mRNA.
<a href="#">RGD1562301</a>	<a href="#">RGD1562301.aSep08</a>	<a href="#">501055</a>	2357	1043	5	315	similar to glutaminyl-tRNA synthetase (RGD1562301) alternative variant aSep08, mRNA.
<a href="#">RGD1562301</a>	<a href="#">RGD1562301.bSep08</a>	<a href="#">501055</a>	1752	691	2	175	similar to glutaminyl-tRNA synthetase (19.7 kD) (RGD1562301) alternative variant bSep08, mRNA.
<a href="#">RGD1562301</a>	<a href="#">RGD1562301.cSep08</a>	<a href="#">501055</a>	1843	672	3	117	similar to glutaminyl-tRNA synthetase (RGD1562301) alternative variant cSep08, mRNA.
<a href="#">RGD1562310</a>	<a href="#">RGD1562310.bSep08</a>	<a href="#">498188</a>	6423	743	3	89	similar to hypothetical protein FLJ21415 (RGD1562310) alternative variant bSep08, mRNA.
<a href="#">RGD1562311</a>	<a href="#">RGD1562311.aSep08</a>	<a href="#">365169</a>	2646	231		54	similar to PIRA5 (RGD1562311) mRNA.
<a href="#">RGD1562317</a>	<a href="#">RGD1562317.bSep08</a>	<a href="#">301388</a>	2564	369	1	66	similar to expressed sequence AW212394 (RGD1562317) alternative variant bSep08, mRNA.
<a href="#">RGD1562323</a>	<a href="#">RGD1562323.aSep08</a>	<a href="#">499984</a>	18655	2068		373	similar to fatty acid translocase/CD36 (RGD1562323) mRNA.
<a href="#">RGD1562326</a>	<a href="#">RGD1562326.aSep08</a>	<a href="#">499877</a>	12464	2908	6	190	similar to ubiquitin-protein ligase E3-alpha (RGD1562326) mRNA.
<a href="#">RGD1562335</a>	<a href="#">RGD1562335.aSep08</a>	<a href="#">498703</a>	32622	1797	5	475	similar to mKIAA1931 protein (RGD1562335) alternative variant aSep08, mRNA.
<a href="#">RGD1562335</a>	<a href="#">RGD1562335.bSep08</a>	<a href="#">498703</a>	20884	1444	7	357	similar to mKIAA1931 protein (RGD1562335) alternative variant bSep08, mRNA.
<a href="#">RGD1562335</a>	<a href="#">RGD1562335.cSep08</a>	<a href="#">498703</a>	27517	719	2	179	similar to mKIAA1931 protein (RGD1562335) alternative variant cSep08, mRNA.
<a href="#">RGD1562335</a>	<a href="#">RGD1562335.dSep08</a>	<a href="#">498703</a>	4090	530	3	176	similar to mKIAA1931 protein (RGD1562335) alternative variant dSep08, mRNA.
<a href="#">RGD1562335</a>	<a href="#">RGD1562335.gSep08</a>	<a href="#">498703</a>	16155	530	4	76	similar to mKIAA1931 protein (RGD1562335) alternative variant gSep08, mRNA.
<a href="#">RGD1562344</a>	<a href="#">RGD1562344.aSep08</a>	<a href="#">295337</a>	5133	1236	4	188	similar to Gm566 protein (20.8 kD) (RGD1562344) alternative variant aSep08, mRNA.
<a href="#">RGD1562344</a>	<a href="#">RGD1562344.cSep08</a>	<a href="#">295337</a>	10740	1783	4	21	similar to Gm566 protein (RGD1562344) alternative variant cSep08, mRNA.
<a href="#">RGD1562351</a>	<a href="#">RGD1562351.bSep08</a>	<a href="#">499990</a>	18199	2806	2	82	putative protein, with a transmembrane domain, of vertebrate origin (8.8 kD) (RGD1562351) alternative variant bSep08, mRNA.
<a href="#">RGD1562352</a>	<a href="#">RGD1562352.aSep08</a>	<a href="#">498787</a>	5809	1037	1	294	similar to isopentenyl diphosphate delta-isomerase type 2 (RGD1562352) alternative variant aSep08, mRNA.
<a href="#">RGD1562352</a>	<a href="#">RGD1562352.bSep08</a>	<a href="#">498787</a>	3985	716	1	89	similar to isopentenyl diphosphate delta-isomerase type 2 (10.5 kD) (RGD1562352) alternative variant bSep08, mRNA.



<a href="#">RGD1562356</a>	<a href="#">RGD1562356.bSep08</a>	<a href="#">498456</a>	12696	502	3	92	similar to hypothetical protein A430083B19 (RGD1562356) alternative variant bSep08, mRNA.
<a href="#">RGD1562356</a>	<a href="#">RGD1562356.cSep08</a>	<a href="#">498456</a>	643	524	1	74	similar to hypothetical protein A430083B19 (RGD1562356) alternative variant cSep08, mRNA.
<a href="#">RGD1562359</a>	<a href="#">RGD1562359.aSep08</a>	<a href="#">500049</a>	10548	1067		355	similar to hypothetical protein 4932408B21 (RGD1562359) mRNA.
<a href="#">RGD1562371</a>	<a href="#">RGD1562371.aSep08</a>	<a href="#">498819</a>	11688	424		61	similar to GREB1 protein isoform a (RGD1562371) mRNA.
<a href="#">RGD1562376</a>	<a href="#">RGD1562376.aSep08</a>	<a href="#">499055</a>	2310	496	3	165	CRA d (RGD1562376) alternative variant aSep08, mRNA.
<a href="#">RGD1562376</a>	<a href="#">RGD1562376.bSep08</a>	<a href="#">499055</a>	5429	1211	4	157	CRA d (18.3 kD) (RGD1562376) alternative variant bSep08, mRNA.
<a href="#">RGD1562376</a>	<a href="#">RGD1562376.cSep08</a>	<a href="#">499055</a>	3706	1019	3	75	CRA d like (8.0 kD) (RGD1562376) alternative variant cSep08, mRNA.
<a href="#">RGD1562376</a>	<a href="#">RGD1562376.dSep08</a>	<a href="#">499055</a>	1243	392	2	59	CRA c like (6.3 kD) (RGD1562376) alternative variant dSep08, mRNA.
<a href="#">RGD1562376</a>	<a href="#">RGD1562376.eSep08</a>	<a href="#">499055</a>	4810	444	3	19	putative protein (RGD1562376) alternative variant eSep08, mRNA.
<a href="#">RGD1562390</a>	<a href="#">RGD1562390.aSep08</a>	<a href="#">307797</a>	1819	890		228	similar to RGD, leucine-rich repeat, tropomodulin and proline-rich containing protein (RGD1562390) mRNA.
<a href="#">RGD1562406</a>	<a href="#">RGD1562406.aSep08</a>	<a href="#">288559</a>	1019	834	3	252	similar to FLJ00248 protein (RGD1562406) alternative variant aSep08, mRNA.
<a href="#">RGD1562449</a>	<a href="#">RGD1562449.aSep08</a>	<a href="#">366333</a>	8824	662		131	similar to hypothetical protein MGC2817 (RGD1562449) mRNA.
<a href="#">RGD1562451</a>	<a href="#">RGD1562451.aSep08</a>	<a href="#">288398</a>	3578	3023	1	331	similar to Pabpc4 predicted protein (35.5 kD) (RGD1562451) alternative variant aSep08, mRNA.
<a href="#">RGD1562451</a>	<a href="#">RGD1562451.bSep08</a>	<a href="#">288398</a>	775	640	2	213	similar to Pabpc4 predicted protein (RGD1562451) alternative variant bSep08, mRNA.
<a href="#">RGD1562465</a>	<a href="#">RGD1562465.aSep08</a>	<a href="#">499006</a>	14227	656		58	similar to GTPase activating protein testicular GAP1 (RGD1562465) mRNA.
<a href="#">RGD1562481andGpr123</a>	<a href="#">RGD1562481andGpr123.bSep08</a>	<a href="#">309097</a>	15540	828	1	120	putative protein (RGD1562481andGpr123) alternative variant bSep08, mRNA.
<a href="#">RGD1562481andGpr123</a>	<a href="#">RGD1562481andGpr123.bSep08</a>	<a href="#">499283</a>	15540	828	1	120	putative protein (RGD1562481andGpr123) alternative variant bSep08, mRNA.
<a href="#">RGD1562488_predicted</a>	<a href="#">RGD1562488_predicted.aSep08</a>	<a href="#">294268</a>	1069	671		127	similar to butyrophilin-like 8 (predicted) (RGD1562488_predicted) mRNA.
<a href="#">RGD1562492</a>	<a href="#">RGD1562492.bSep08</a>	<a href="#">499151</a>	11405	2560	4	676	similar to Orphan sodium- and chloride-dependent neurotransmitter transporter NTT5 (Solute carrier family 6 member 16) (76.7 kD) (RGD1562492) alternative variant bSep08, mRNA.
<a href="#">RGD1562494</a>	<a href="#">RGD1562494.aSep08</a>	<a href="#">500935</a>	925	366		122	similar to keratin 6 alpha (RGD1562494) mRNA.
<a href="#">RGD1562502</a>	<a href="#">RGD1562502.bSep08</a>	<a href="#">363485</a>	3582	1057	4	102	similar to RIKEN cDNA 2610029G23 (RGD1562502) alternative variant bSep08, mRNA.
<a href="#">RGD1562511</a>	<a href="#">RGD1562511.aSep08</a>	<a href="#">500571</a>	20800	1756		301	similar to MmKIF17 (RGD1562511) mRNA.
<a href="#">RGD1562526</a>	<a href="#">RGD1562526.aSep08</a>	<a href="#">499821</a>	50954	1242		414	similar to RIKEN cDNA D430039N05 gene (RGD1562526) mRNA.
<a href="#">RGD1562529</a>	<a href="#">RGD1562529.aSep08</a>	<a href="#">311372</a>	12487	2378		764	similar to hypothetical protein FLJ21439 (RGD1562529) mRNA.

<a href="#">RGD1562532</a>	<a href="#">RGD1562532.aSep08</a>	<a href="#">500513</a>	66403	2433		441	hypothetical gene supported by BC079057 (50.6 kD) (RGD1562532) mRNA.
<a href="#">RGD1562533</a>	<a href="#">RGD1562533.bSep08</a>	<a href="#">498136</a>	17670	971	3	172	similar to mKIAA0774 protein (RGD1562533) alternative variant bSep08, mRNA.
<a href="#">RGD1562540</a>	<a href="#">RGD1562540.bSep08</a>	<a href="#">500678</a>	13681	479	4	88	RGD1562540 (RGD1562540) alternative variant bSep08, mRNA.
<a href="#">RGD1562543</a>	<a href="#">RGD1562543.aSep08</a>	<a href="#">308719</a>	828	744	2	101	similar to ribosomal protein L27a (RGD1562543) mRNA.
<a href="#">RGD1562557</a>	<a href="#">RGD1562557.aSep08</a>	<a href="#">500723</a>	1672	722		240	putative protein of mammalian origin (RGD1562557) mRNA.
<a href="#">RGD1562582</a>	<a href="#">RGD1562582.bSep08</a>	<a href="#">499935</a>	30738	537	4	179	similar to KIAA0406-like protein (RGD1562582) alternative variant bSep08, mRNA.
<a href="#">RGD1562590</a>	<a href="#">RGD1562590.aSep08</a>	<a href="#">312365</a>	42362	1027		342	similar to hypothetical protein (RGD1562590) mRNA.
<a href="#">RGD1562608</a>	<a href="#">RGD1562608.aSep08</a>	<a href="#">498831</a>	30836	715		77	similar to KIAA1328 protein (RGD1562608) mRNA.
<a href="#">RGD1562618</a>	<a href="#">RGD1562618.bSep08</a>	<a href="#">501007</a>	3932	994	2	103	similar to RIKEN cDNA 6030419C18 gene (11.7 kD) (RGD1562618) alternative variant bSep08, mRNA.
<a href="#">RGD1562619</a>	<a href="#">RGD1562619.aSep08</a>	<a href="#">499147</a>	1559	345	1	109	similar to SCRL protein variant 1 (RGD1562619) alternative variant aSep08, mRNA.
<a href="#">RGD1562619</a>	<a href="#">RGD1562619.bSep08</a>	<a href="#">499147</a>	1557	326	1	75	similar to SCRL protein variant 1 (RGD1562619) alternative variant bSep08, mRNA.
<a href="#">RGD1562622</a>	<a href="#">RGD1562622.bSep08</a>	<a href="#">503035</a>	1204	568	2	47	similar to RIKEN cDNA 6330442E10 gene (5.1 kD) (RGD1562622) alternative variant bSep08, mRNA.
<a href="#">RGD1562626</a>	<a href="#">RGD1562626.aSep08</a>	<a href="#">287622</a>	44284	2073	16	474	similar to adaptor molecule SRCASM (RGD1562626) alternative variant aSep08, mRNA.
<a href="#">RGD1562626</a>	<a href="#">RGD1562626.bSep08</a>	<a href="#">287622</a>	4439	404	3	50	similar to adaptor molecule SRCASM (RGD1562626) alternative variant bSep08, mRNA.
<a href="#">RGD1562629</a>	<a href="#">RGD1562629.aSep08</a>	<a href="#">361948</a>	138028	780	6	260	similar to neurobeachin (RGD1562629) alternative variant aSep08, mRNA.
<a href="#">RGD1562629</a>	<a href="#">RGD1562629.bSep08</a>	<a href="#">361948</a>	3237	515	2	31	similar to neurobeachin (3.8 kD) (RGD1562629) alternative variant bSep08, mRNA.
<a href="#">RGD1562639</a>	<a href="#">RGD1562639.aSep08</a>	<a href="#">315756</a>	28017	532		60	similar to c-myc promoter binding protein (RGD1562639) mRNA.
<a href="#">RGD1562646</a>	<a href="#">RGD1562646.aSep08</a>	<a href="#">305392</a>	14790	1457		485	condensin I complex (RGD1562646) mRNA.
<a href="#">RGD1562657</a>	<a href="#">RGD1562657.bSep08</a>	<a href="#">361709</a>	507	425	2	121	similar to hypothetical protein DKFZp761E198 (RGD1562657) alternative variant bSep08, mRNA.
<a href="#">RGD1562665</a>	<a href="#">RGD1562665.aSep08</a>	<a href="#">362936</a>	6612	1657	4	385	similar to 1500031N24Rik protein (RGD1562665) alternative variant aSep08, mRNA.
<a href="#">RGD1562665</a>	<a href="#">RGD1562665.bSep08</a>	<a href="#">362936</a>	10817	1868	5	369	similar to 1500031N24Rik protein (41.7 kD) (RGD1562665) alternative variant bSep08, mRNA.
<a href="#">RGD1562673</a>	<a href="#">RGD1562673.bSep08</a>	<a href="#">363551</a>	14632	1634	3	97	similar to Prostatic spermine-binding protein precursor (SBP) (10.8 kD) (RGD1562673) alternative variant bSep08, mRNA.
<a href="#">RGD1562691</a>	<a href="#">RGD1562691.aSep08</a>	<a href="#">361185</a>	9492	1221	5	273	similar to RIKEN cDNA 0710008K08 (RGD1562691) alternative variant aSep08, mRNA.
<a href="#">RGD1562691</a>	<a href="#">RGD1562691.bSep08</a>	<a href="#">361185</a>	3150	630	3	130	similar to RIKEN cDNA 0710008K08 (RGD1562691) alternative variant bSep08, mRNA.
<a href="#">RGD1562691</a>	<a href="#">RGD1562691.cSep08</a>	<a href="#">361185</a>	13198	334	3	70	similar to RIKEN cDNA 0710008K08 (RGD1562691) alternative variant cSep08, mRNA.

<a href="#">RGD1562691</a>	<a href="#">RGD1562691.dSep08</a>	<a href="#">361185</a>	9407	1553	3	126	similar to RIKEN cDNA 0710008K08 (13.3 kD) (RGD1562691) alternative variant dSep08, mRNA.
<a href="#">RGD1562717</a>	<a href="#">RGD1562717.aSep08</a>	<a href="#">363767</a>	99610	448		149	similar to ABI gene family, member 3 (NESH) binding protein (RGD1562717) mRNA.
<a href="#">RGD1562767</a>	<a href="#">RGD1562767.bSep08</a>	<a href="#">362133</a>	7874	578	1	192	similar to RIKEN cDNA 2310010M24 (RGD1562767) alternative variant bSep08, mRNA.
<a href="#">RGD1562767</a>	<a href="#">RGD1562767.cSep08</a>	<a href="#">362133</a>	11137	1902	2	172	similar to RIKEN cDNA 2310010M24 (19.6 kD) (RGD1562767) alternative variant cSep08, mRNA.
<a href="#">RGD1562768</a>	<a href="#">RGD1562768.aSep08</a>	<a href="#">302327</a>	6092	620		136	similar to ZNF6 protein (RGD1562768) mRNA.
<a href="#">RGD1562811</a>	<a href="#">RGD1562811.aSep08</a>	<a href="#">315651</a>	11254	1250		220	RGD1562811 (RGD1562811) mRNA.
<a href="#">RGD1562844</a>	<a href="#">RGD1562844.aSep08</a>	<a href="#">306892</a>	31461	997	5	332	similar to serine (or cysteine) proteinase inhibitor, clade B, member 9 (RGD1562844) alternative variant aSep08, mRNA.
<a href="#">RGD1562844</a>	<a href="#">RGD1562844.bSep08</a>	<a href="#">306892</a>	8023	1612	4	329	similar to serine (or cysteine) proteinase inhibitor, clade B, member 9 (RGD1562844) alternative variant bSep08, mRNA.
<a href="#">RGD1562844</a>	<a href="#">RGD1562844.cSep08</a>	<a href="#">306892</a>	4741	761		150	similar to serine (or cysteine) proteinase inhibitor, clade B, member 9 (RGD1562844) alternative variant cSep08, mRNA.
<a href="#">RGD1562846</a>	<a href="#">RGD1562846.aSep08</a>	<a href="#">502694</a>	31168	541		178	similar to Docking protein 5 (Downstream of tyrosine kinase 5) (Protein dok-5) (RGD1562846) alternative variant aSep08, mRNA.
<a href="#">RGD1562848</a>	<a href="#">RGD1562848.aSep08</a>	<a href="#">362565</a>	157860	594		104	RGD1562848 (RGD1562848) mRNA.
<a href="#">RGD1562860</a>	<a href="#">RGD1562860.aSep08</a>	<a href="#">360945</a>	15343	1353		181	similar to RIKEN cDNA 2310045A20 (RGD1562860) mRNA.
<a href="#">RGD1562865</a>	<a href="#">RGD1562865.aSep08</a>	<a href="#">313125</a>	12954	1032		344	similar to BTB and CNC homology 1, basic leucine zipper transcription factor 2 (RGD1562865) mRNA.
<a href="#">RGD1562877</a>	<a href="#">RGD1562877.aSep08</a>	<a href="#">296780</a>	3771	713		144	RGD1562877 (17.3 kD) (RGD1562877) mRNA.
<a href="#">RGD1562911</a>	<a href="#">RGD1562911.bSep08</a>	<a href="#">500534</a>	1727	724	4	159	similar to hypothetical protein MGC45441 (RGD1562911) alternative variant bSep08, mRNA.
<a href="#">RGD1562911</a>	<a href="#">RGD1562911.cSep08</a>	<a href="#">500534</a>	961	434	4	132	similar to hypothetical protein MGC45441 (RGD1562911) alternative variant cSep08, mRNA.
<a href="#">RGD1562943</a>	<a href="#">RGD1562943.bSep08</a>	<a href="#">499113</a>	7224	1056	6	229	similar to selenoprotein V (RGD1562943) alternative variant bSep08, mRNA.
<a href="#">RGD1562943</a>	<a href="#">RGD1562943.cSep08</a>	<a href="#">499113</a>	6713	470	5	59	similar to selenoprotein V (RGD1562943) alternative variant cSep08, mRNA.
<a href="#">RGD1562943</a>	<a href="#">RGD1562943.dSep08</a>	<a href="#">499113</a>	4228	272	2	32	similar to selenoprotein V (RGD1562943) alternative variant dSep08, mRNA.
<a href="#">RGD1562952</a>	<a href="#">RGD1562952.aSep08</a>	<a href="#">365661</a>	12639	1102	9	300	similar to ErbB2 interacting protein isoform 2 (34.6 kD) (RGD1562952) alternative variant aSep08, mRNA.
<a href="#">RGD1562963</a>	<a href="#">RGD1562963.aSep08</a>	<a href="#">498729</a>	12656	738	4	157	putative protein of vertebrate origin (18.0 kD) (RGD1562963) alternative variant aSep08, mRNA.
<a href="#">RGD1562965</a>	<a href="#">RGD1562965.aSep08</a>	<a href="#">498650</a>	30398	5113	4	479	similar to Rab coupling protein isoform 3 (RGD1562965) alternative variant aSep08, mRNA.
<a href="#">RGD1562965</a>	<a href="#">RGD1562965.bSep08</a>	<a href="#">498650</a>	5685	904	1	273	similar to Rab coupling protein isoform 3 (RGD1562965) alternative variant bSep08, mRNA.

<a href="#">RGD1562979</a>	<a href="#">RGD1562979.aSep08</a>	<a href="#">305501</a>	78547	579		192	similar to DNA-binding protein Ikaros form 1 - mouse (RGD1562979) mRNA.
<a href="#">RGD1562987</a>	<a href="#">RGD1562987.aSep08</a>	<a href="#">498886</a>	8721	1028	3	118	CRA a (RGD1562987) alternative variant aSep08, mRNA.
<a href="#">RGD1562988</a>	<a href="#">RGD1562988.aSep08</a>	<a href="#">500464</a>	87225	1800	1	533	similar to EHM2 (RGD1562988) alternative variant aSep08, mRNA.
<a href="#">RGD1563034</a>	<a href="#">RGD1563034.aSep08</a>	<a href="#">292720</a>	1343	597		199	similar to ETS domain transcription factor ERF (Ets2 repressor factor) (RGD1563034) mRNA.
<a href="#">RGD1563037</a>	<a href="#">RGD1563037.aSep08</a>	<a href="#">309357</a>	24711	1997		407	similar to family with sequence similarity 11, member A (RGD1563037) alternative variant aSep08, mRNA.
<a href="#">RGD1563037</a>	<a href="#">RGD1563037.bSep08</a>	<a href="#">309357</a>	3641	1930		172	similar to family with sequence similarity 11, member A (20.0 kD) (RGD1563037) alternative variant bSep08, mRNA.
<a href="#">RGD1563060</a>	<a href="#">RGD1563060.bSep08</a>	<a href="#">291608</a>	11625	319	2	80	similar to AVLV472 (RGD1563060) alternative variant bSep08, mRNA.
<a href="#">RGD1563092</a>	<a href="#">RGD1563092.aSep08</a>	<a href="#">500642</a>	85852	610		93	similar to histone deacetylase-related protein (RGD1563092) mRNA.
<a href="#">RGD1563095</a>	<a href="#">RGD1563095.aSep08</a>	<a href="#">288485</a>	6651	1139	3	187	similar to hypothetical protein DKFZp434J1015.1 - human (fragment) (RGD1563095) alternative variant aSep08, mRNA.
<a href="#">RGD1563095</a>	<a href="#">RGD1563095.bSep08</a>	<a href="#">288485</a>	1607	751	2	148	similar to hypothetical protein DKFZp434J1015.1 - human (fragment) (RGD1563095) alternative variant bSep08, mRNA.
<a href="#">RGD1563102</a>	<a href="#">RGD1563102.aSep08</a>	<a href="#">498044</a>	76308	523		100	RGD1563102 (RGD1563102) mRNA.
<a href="#">RGD1563106</a>	<a href="#">RGD1563106.bSep08</a>	<a href="#">303238</a>	7639	2448	13	570	similar to novel protein (63.2 kD) (RGD1563106) alternative variant bSep08, mRNA.
<a href="#">RGD1563107</a>	<a href="#">RGD1563107.aSep08</a>	<a href="#">499034</a>	14473	805		170	RGD1563107 (18.7 kD) (RGD1563107) mRNA.
<a href="#">RGD1563109</a>	<a href="#">RGD1563109.aSep08</a>	<a href="#">499586</a>	18063	763	4	216	RGD1563109 (RGD1563109) alternative variant aSep08, mRNA.
<a href="#">RGD1563109</a>	<a href="#">RGD1563109.bSep08</a>	<a href="#">499586</a>	33478	449	4	141	RGD1563109 (RGD1563109) alternative variant bSep08, mRNA.
<a href="#">RGD1563109</a>	<a href="#">RGD1563109.cSep08</a>	<a href="#">499586</a>	6547	672	3	128	RGD1563109 (RGD1563109) alternative variant cSep08, mRNA.
<a href="#">RGD1563109</a>	<a href="#">RGD1563109.dSep08</a>	<a href="#">499586</a>	21883	275	2	83	RGD1563109 (RGD1563109) alternative variant dSep08, mRNA.
<a href="#">RGD1563110</a>	<a href="#">RGD1563110.aSep08</a>	<a href="#">502905</a>	10038	764	3	159	similar to immunoreceptor Ly49si3 (RGD1563110) alternative variant aSep08, mRNA.
<a href="#">RGD1563110</a>	<a href="#">RGD1563110.bSep08</a>	<a href="#">502905</a>	4705	346	1	76	similar to immunoreceptor Ly49si3 (RGD1563110) alternative variant bSep08, mRNA.
<a href="#">RGD1563120</a>	<a href="#">RGD1563120.aSep08</a>	<a href="#">499895</a>	172348	700		194	similar to RIKEN cDNA 2210009G21 (RGD1563120) mRNA.
<a href="#">RGD1563136</a>	<a href="#">RGD1563136.aSep08</a>	<a href="#">296230</a>	6343	735		163	similar to Cystatin-related protein 2 precursor (Prostatic 22 kDa glycoprotein P22K15) (RGD1563136) mRNA.
<a href="#">RGD1563148</a>	<a href="#">RGD1563148.bSep08</a>	<a href="#">362447</a>	2407	848	3	116	similar to osteoclast inhibitory lectin (RGD1563148) alternative variant bSep08, mRNA.
<a href="#">RGD1563148</a>	<a href="#">RGD1563148.cSep08</a>	<a href="#">362447</a>	2883	1927	2	51	similar to osteoclast inhibitory lectin (RGD1563148) alternative variant cSep08, mRNA.

<a href="#">RGD1563148</a>	<a href="#">RGD1563148.dSep08</a>	<a href="#">362447</a>	756	368	1	49	similar to osteoclast inhibitory lectin (RGD1563148) alternative variant dSep08, mRNA.
<a href="#">RGD1563154</a>	<a href="#">RGD1563154.aSep08</a>	<a href="#">296788</a>	8576	368		122	similar to cadherin EGF LAG seven-pass G-type receptor 2 (RGD1563154) mRNA.
<a href="#">RGD1563180</a>	<a href="#">RGD1563180.aSep08</a>	<a href="#">291815</a>	6971	422		140	similar to amyloid beta precursor protein binding protein 1 (RGD1563180) mRNA.
<a href="#">RGD1563200</a>	<a href="#">RGD1563200.bSep08</a>	<a href="#">500938</a>	6152	2475	1	109	putative nuclear protein of eukaryotic origin (12.4 kD) (RGD1563200) alternative variant bSep08, complete mRNA.
<a href="#">RGD1563216</a>	<a href="#">RGD1563216.bSep08</a>	<a href="#">500694</a>	1576	1172	2	123	iron-sulfur cluster assembly 2 (13.6 kD) (RGD1563216) alternative variant bSep08, mRNA.
<a href="#">RGD1563217</a>	<a href="#">RGD1563217.bSep08</a>	<a href="#">499265</a>	4623	746	2	98	similar to RIKEN cDNA 4930451I11 (RGD1563217) alternative variant bSep08, mRNA.
<a href="#">RGD1563224</a>	<a href="#">RGD1563224.aSep08</a>	<a href="#">500874</a>	105862	1956	2	227	similar to 4930438D12Rik protein (RGD1563224) alternative variant aSep08, mRNA.
<a href="#">RGD1563224</a>	<a href="#">RGD1563224.cSep08</a>	<a href="#">500874</a>	42869	3940	7	178	similar to 4930438D12Rik protein (19.1 kD) (RGD1563224) alternative variant cSep08, mRNA.
<a href="#">RGD1563224</a>	<a href="#">RGD1563224.dSep08</a>	<a href="#">500874</a>	669	374	2	124	similar to 4930438D12Rik protein (RGD1563224) alternative variant dSep08, mRNA.
<a href="#">RGD1563236</a>	<a href="#">RGD1563236.aSep08</a>	<a href="#">498608</a>	1704	449		149	similar to BC028663 protein (RGD1563236) mRNA.
<a href="#">RGD1563239</a>	<a href="#">RGD1563239.bSep08</a>	<a href="#">502316</a>	9084	741	1	178	similar to Zinc finger protein 566 (RGD1563239) alternative variant bSep08, mRNA.
<a href="#">RGD1563266</a>	<a href="#">RGD1563266.aSep08</a>	<a href="#">361143</a>	27899	821	5	118	RGD1563266 (RGD1563266) alternative variant aSep08, mRNA.
<a href="#">RGD1563266</a>	<a href="#">RGD1563266.bSep08</a>	<a href="#">361143</a>	11506	495	1	103	RGD1563266 (RGD1563266) alternative variant bSep08, mRNA.
<a href="#">RGD1563270</a>	<a href="#">RGD1563270.aSep08</a>	<a href="#">302172</a>	6273	522		94	similar to synaptonemal complex protein 3 (RGD1563270) mRNA.
<a href="#">RGD1563278</a>	<a href="#">RGD1563278.aSep08</a>	<a href="#">303963</a>	31008	2367	19	713	zinc finger DAZ interacting protein 3 CRA a (RGD1563278) alternative variant aSep08, mRNA.
<a href="#">RGD1563278</a>	<a href="#">RGD1563278.bSep08</a>	<a href="#">303963</a>	36512	1707	15	430	DAZ interacting protein 3 zinc finger (RGD1563278) alternative variant bSep08, mRNA.
<a href="#">RGD1563278</a>	<a href="#">RGD1563278.cSep08</a>	<a href="#">303963</a>	9962	749	5	163	DAZ interacting protein 3 zinc finger (18.8 kD) (RGD1563278) alternative variant cSep08, mRNA.
<a href="#">RGD1563278</a>	<a href="#">RGD1563278.dSep08</a>	<a href="#">303963</a>	1902	508	2	96	DAZ interacting protein 3 zinc finger (RGD1563278) alternative variant dSep08, mRNA.
<a href="#">RGD1563284</a>	<a href="#">RGD1563284.aSep08</a>	<a href="#">290691</a>	19416	688		68	similar to actin (8.3 kD) (RGD1563284) mRNA.
<a href="#">RGD1563285</a>	<a href="#">RGD1563285.aSep08</a>	<a href="#">499273</a>	6536	328		97	similar to RIKEN cDNA 1700022C21 (RGD1563285) mRNA.
<a href="#">RGD1563302</a>	<a href="#">RGD1563302.aSep08</a>	<a href="#">361049</a>	8698	709	2	172	RGD1563302 (RGD1563302) alternative variant aSep08, mRNA.
<a href="#">RGD1563302</a>	<a href="#">RGD1563302.bSep08</a>	<a href="#">361049</a>	8364	373	2	63	RGD1563302 (RGD1563302) alternative variant bSep08, mRNA.
<a href="#">RGD1563309</a>	<a href="#">RGD1563309.aSep08</a>	<a href="#">368088</a>	11222	1319	11	439	similar to diacylglycerol kinase, delta 130kDa isoform 1 (RGD1563309) alternative variant aSep08, mRNA.
<a href="#">RGD1563309</a>	<a href="#">RGD1563309.bSep08</a>	<a href="#">368088</a>	482	402	1	54	similar to diacylglycerol kinase, delta 130kDa isoform 1 (RGD1563309) alternative variant bSep08, mRNA.

<a href="#">RGD1563319</a>	<a href="#">RGD1563319.aSep08</a>	<a href="#">293632</a>	15468	1153		230	similar to RIKEN cDNA 6330512M04 gene (RGD1563319) mRNA.
<a href="#">RGD1563323</a>	<a href="#">RGD1563323.aSep08</a>	<a href="#">361702</a>	439	359		91	similar to hypothetical protein FLJ22531 (RGD1563323) mRNA.
<a href="#">RGD1563325</a>	<a href="#">RGD1563325.aSep08</a>	<a href="#">362857</a>	18331	3482	3	187	similar to hypothetical protein MGC17943 (RGD1563325) alternative variant aSep08, mRNA.
<a href="#">RGD1563342</a>	<a href="#">RGD1563342.aSep08</a>	<a href="#">304572</a>	15920	1124	8	374	similar to RIKEN cDNA 2410025L10 (RGD1563342) alternative variant aSep08, mRNA.
<a href="#">RGD1563347_predicted</a>	<a href="#">RGD1563347_predicted_aSep08</a>	<a href="#">501095</a>	54205	1739		244	similar to RIKEN cDNA 2310015N21 (predicted) (RGD1563347_predicted) mRNA.
<a href="#">RGD1563348</a>	<a href="#">RGD1563348.aSep08</a>	<a href="#">502642</a>	1002	792	3	71	similar to Selenoprotein H (RGD1563348) alternative variant aSep08, mRNA.
<a href="#">RGD1563348</a>	<a href="#">RGD1563348.cSep08</a>	<a href="#">502642</a>	1474	622	4	70	similar to Selenoprotein H (RGD1563348) alternative variant cSep08, mRNA.
<a href="#">RGD1563348</a>	<a href="#">RGD1563348.dSep08</a>	<a href="#">502642</a>	1433	609	5	45	similar to Selenoprotein H (RGD1563348) alternative variant dSep08, mRNA.
<a href="#">RGD1563348</a>	<a href="#">RGD1563348.eSep08</a>	<a href="#">502642</a>	1481	607	4	61	similar to Selenoprotein H (RGD1563348) alternative variant eSep08, mRNA.
<a href="#">RGD1563351</a>	<a href="#">RGD1563351.aSep08</a>	<a href="#">294350</a>	2356	1005		178	RGD1563351 (4.5 kD) (RGD1563351) mRNA.
<a href="#">RGD1563354</a>	<a href="#">RGD1563354.aSep08</a>	<a href="#">311592</a>	7204	1301		365	similar to hypothetical protein D630003M21 (RGD1563354) mRNA.
<a href="#">RGD1563398</a>	<a href="#">RGD1563398.aSep08</a>	<a href="#">301075</a>	4223	535		154	RGD1563398 (RGD1563398) mRNA.
<a href="#">RGD1563400</a>	<a href="#">RGD1563400.aSep08</a>	<a href="#">360777</a>	3003	1029	2	50	similar to paired immunoglobulin-like type 2 receptor beta (RGD1563400) alternative variant aSep08, mRNA.
<a href="#">RGD1563400</a>	<a href="#">RGD1563400.bSep08</a>	<a href="#">360777</a>	885	328	2	46	similar to paired immunoglobulin-like type 2 receptor beta (5.1 kD) (RGD1563400) alternative variant bSep08, mRNA.
<a href="#">RGD1563422</a>	<a href="#">RGD1563422.bSep08</a>	<a href="#">289182</a>	19364	854	4	108	similar to Brain protein 44 (12.0 kD) (RGD1563422) alternative variant bSep08, mRNA.
<a href="#">RGD1563437</a>	<a href="#">RGD1563437.aSep08</a>	<a href="#">291356</a>	9601	664	6	221	similar to KIAA1217 (RGD1563437) alternative variant aSep08, mRNA.
<a href="#">RGD1563438</a>	<a href="#">RGD1563438.aSep08</a>	<a href="#">287442</a>	1123	485	3	113	similar to novel protein of unknown function (DUF423) family member (11.8 kD) (RGD1563438) alternative variant aSep08, mRNA.
<a href="#">RGD1563438</a>	<a href="#">RGD1563438.cSep08</a>	<a href="#">287442</a>	41146	444	3	22	similar to novel protein of unknown function (DUF423) family member (2.4 kD) (RGD1563438) alternative variant cSep08, mRNA.
<a href="#">RGD1563440</a>	<a href="#">RGD1563440.bSep08</a>	<a href="#">298301</a>	17692	1469	11	483	similar to hypothetical protein (RGD1563440) alternative variant bSep08, mRNA.
<a href="#">RGD1563440</a>	<a href="#">RGD1563440.cSep08</a>	<a href="#">298301</a>	1108	390	2	129	similar to hypothetical protein (RGD1563440) alternative variant cSep08, mRNA.
<a href="#">RGD1563440</a>	<a href="#">RGD1563440.dSep08</a>	<a href="#">298301</a>	8725	708	5	106	similar to hypothetical protein (RGD1563440) alternative variant dSep08, mRNA.
<a href="#">RGD1563444</a>	<a href="#">RGD1563444.bSep08</a>	<a href="#">500577</a>	7703	1032	5	229	similar to RIKEN cDNA 2900090M10 (26.3 kD) (RGD1563444) alternative variant bSep08, mRNA.
<a href="#">RGD1563444</a>	<a href="#">RGD1563444.cSep08</a>	<a href="#">500577</a>	4777	549	3	113	similar to RIKEN cDNA 2900090M10 (RGD1563444) alternative variant cSep08, mRNA.

<a href="#">RGD1563444</a>	<a href="#">RGD1563444.dSep08</a>	<a href="#">500577</a>	14026	555	2	50	similar to RIKEN cDNA 2900090M10 (RGD1563444) alternative variant dSep08, mRNA.
<a href="#">RGD1563482</a>	<a href="#">RGD1563482.bSep08</a>	<a href="#">498179</a>	17497	3728	4	214	similar to hypothetical protein FLJ38663 (RGD1563482) alternative variant bSep08, mRNA.
<a href="#">RGD1563482</a>	<a href="#">RGD1563482.cSep08</a>	<a href="#">498179</a>	13620	1014	3	106	similar to hypothetical protein FLJ38663 (RGD1563482) alternative variant cSep08, mRNA.
<a href="#">RGD1563507</a>	<a href="#">RGD1563507.aSep08</a>	<a href="#">500238</a>	6336	3092	3	99	similar to RIKEN cDNA 1700019G17 and similar to Camello-like 2 (11.4 kD) (RGD1563507) alternative variant aSep08, mRNA.
<a href="#">RGD1563507</a>	<a href="#">RGD1563507.aSep08</a>	<a href="#">681227</a>	6336	3092	3	99	similar to RIKEN cDNA 1700019G17 and similar to Camello-like 2 (11.4 kD) (RGD1563507) alternative variant aSep08, mRNA.
<a href="#">RGD1563507</a>	<a href="#">RGD1563507.bSep08</a>	<a href="#">500238</a>	4121	750	3	79	similar to RIKEN cDNA 1700019G17 and similar to Camello-like 2 (RGD1563507) alternative variant bSep08, mRNA.
<a href="#">RGD1563507</a>	<a href="#">RGD1563507.bSep08</a>	<a href="#">681227</a>	4121	750	3	79	similar to RIKEN cDNA 1700019G17 and similar to Camello-like 2 (RGD1563507) alternative variant bSep08, mRNA.
<a href="#">RGD1563510</a>	<a href="#">RGD1563510.aSep08</a>	<a href="#">362240</a>	64408	582	5	193	similar to RIKEN cDNA 8430427H17 gene (RGD1563510) alternative variant aSep08, mRNA.
<a href="#">RGD1563510</a>	<a href="#">RGD1563510.bSep08</a>	<a href="#">362240</a>	24502	483	3	160	similar to RIKEN cDNA 8430427H17 gene (RGD1563510) alternative variant bSep08, mRNA.
<a href="#">RGD1563510</a>	<a href="#">RGD1563510.cSep08</a>	<a href="#">362240</a>	24062	458	3	152	similar to RIKEN cDNA 8430427H17 gene (RGD1563510) alternative variant cSep08, mRNA.
<a href="#">RGD1563510</a>	<a href="#">RGD1563510.dSep08</a>	<a href="#">362240</a>	6061	347	2	107	similar to RIKEN cDNA 8430427H17 gene (RGD1563510) alternative variant dSep08, mRNA.
<a href="#">RGD1563520</a>	<a href="#">RGD1563520.aSep08</a>	<a href="#">361091</a>	27914	908	1	302	similar to UDP-glucose ceramide glucosyltransferase-like 2 (RGD1563520) alternative variant aSep08, mRNA.
<a href="#">RGD1563520</a>	<a href="#">RGD1563520.bSep08</a>	<a href="#">361091</a>	17273	648	1	215	similar to UDP-glucose ceramide glucosyltransferase-like 2 (RGD1563520) alternative variant bSep08, mRNA.
<a href="#">RGD1563533</a>	<a href="#">RGD1563533.aSep08</a>	<a href="#">313713</a>	31139	998	2	311	similar to novel protein (RGD1563533) alternative variant aSep08, mRNA.
<a href="#">RGD1563533</a>	<a href="#">RGD1563533.bSep08</a>	<a href="#">313713</a>	48094	997	3	306	similar to novel protein (RGD1563533) alternative variant bSep08, mRNA.
<a href="#">RGD1563547</a>	<a href="#">RGD1563547.bSep08</a>	<a href="#">360478</a>	1474	625	1	85	RGD1563547 (9.5 kD) (RGD1563547) alternative variant bSep08, mRNA.
<a href="#">RGD1563550</a>	<a href="#">RGD1563550.aSep08</a>	<a href="#">300834</a>	5940	262		62	similar to hypothetical protein FLJ38736 (RGD1563550) mRNA.
<a href="#">RGD1563556</a>	<a href="#">RGD1563556.aSep08</a>	<a href="#">315409</a>	2572	1634		544	similar to mKIAA1377 protein (RGD1563556) mRNA.
<a href="#">RGD1563560</a>	<a href="#">RGD1563560.aSep08</a>	<a href="#">499321</a>	13052	762		140	similar to ribosomal protein L27a (RGD1563560) mRNA.
<a href="#">RGD1563578</a>	<a href="#">RGD1563578.aSep08</a>	<a href="#">501003</a>	8674	407		135	similar to PGC7 (RGD1563578) mRNA.
<a href="#">RGD1563583</a>	<a href="#">RGD1563583.aSep08</a>	<a href="#">299208</a>	87064	1570	7	523	similar to mKIAA0998 protein (RGD1563583) alternative variant aSep08, mRNA.
<a href="#">RGD1563583</a>	<a href="#">RGD1563583.bSep08</a>	<a href="#">299208</a>	6394	697	1	221	similar to mKIAA0998 protein (RGD1563583) alternative variant bSep08, mRNA.
<a href="#">RGD1563600</a>	<a href="#">RGD1563600.aSep08</a>	<a href="#">364093</a>	10775	450		70	similar to ribosomal protein S11 (7.9 kD) (RGD1563600) mRNA.

<a href="#">RGD1563612</a>	<a href="#">RGD1563612.aSep08</a>	<a href="#">500026</a>	227860	4005	9	430	glucocorticoid induced transcript 1 like (47.6 kD) (RGD1563612) alternative variant aSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.aSep08</a>	<a href="#">680817</a>	227860	4005	9	430	glucocorticoid induced transcript 1 like (47.6 kD) (RGD1563612) alternative variant aSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.bSep08</a>	<a href="#">500026</a>	169488	955	4	133	CRA b (14.3 kD) (RGD1563612) alternative variant bSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.bSep08</a>	<a href="#">680817</a>	169488	955	4	133	CRA b (14.3 kD) (RGD1563612) alternative variant bSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.cSep08</a>	<a href="#">500026</a>	169356	882	5	107	CRA b (11.4 kD) (RGD1563612) alternative variant cSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.cSep08</a>	<a href="#">680817</a>	169356	882	5	107	CRA b (11.4 kD) (RGD1563612) alternative variant cSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.dSep08</a>	<a href="#">500026</a>	665	357	2	73	putative protein (RGD1563612) alternative variant dSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.dSep08</a>	<a href="#">680817</a>	665	357	2	73	putative protein (RGD1563612) alternative variant dSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.eSep08</a>	<a href="#">500026</a>	21590	290	3	50	CRA b like (RGD1563612) alternative variant eSep08, mRNA.
<a href="#">RGD1563612</a>	<a href="#">RGD1563612.eSep08</a>	<a href="#">680817</a>	21590	290	3	50	CRA b like (RGD1563612) alternative variant eSep08, mRNA.
<a href="#">RGD1563628</a>	<a href="#">RGD1563628.bSep08</a>	<a href="#">304554</a>	20909	475	3	157	similar to Seizure 6-like protein precursor (RGD1563628) alternative variant bSep08, mRNA.
<a href="#">RGD1563628</a>	<a href="#">RGD1563628.cSep08</a>	<a href="#">304554</a>	5275	420	5	89	similar to Seizure 6-like protein precursor (RGD1563628) alternative variant cSep08, mRNA.
<a href="#">RGD1563634</a>	<a href="#">RGD1563634.aSep08</a>	<a href="#">500790</a>	5435	3226	7	764	similar to R31449 3 (89.4 kD) (RGD1563634) alternative variant aSep08, mRNA.
<a href="#">RGD1563634</a>	<a href="#">RGD1563634.cSep08</a>	<a href="#">500790</a>	1040	795	1	183	similar to R31449 3 (RGD1563634) alternative variant cSep08, mRNA.
<a href="#">RGD1563652</a>	<a href="#">RGD1563652.aSep08</a>	<a href="#">291332</a>	40305	548		182	putative protein of eukaryotic origin (RGD1563652) mRNA.
<a href="#">RGD1563703</a>	<a href="#">RGD1563703.aSep08</a>	<a href="#">361206</a>	13294	401		92	similar to hypothetical protein E230025K15 (RGD1563703) mRNA.
<a href="#">RGD1563706</a>	<a href="#">RGD1563706.bSep08</a>	<a href="#">362042</a>	5153	386	1	107	similar to Zgc:92184 (RGD1563706) alternative variant bSep08, mRNA.
<a href="#">RGD1563706</a>	<a href="#">RGD1563706.cSep08</a>	<a href="#">362042</a>	5010	389	2	67	similar to Zgc:92184 (RGD1563706) alternative variant cSep08, mRNA.
<a href="#">RGD1563714</a>	<a href="#">RGD1563714.bSep08</a>	<a href="#">500531</a>	23357	748	1	125	RGD1563714 (RGD1563714) alternative variant bSep08, mRNA.
<a href="#">RGD1563719</a>	<a href="#">RGD1563719.aSep08</a>	<a href="#">500324</a>	17304	281		93	similar to hypothetical protein MGC4266 (RGD1563719) mRNA.
<a href="#">RGD1563746</a>	<a href="#">RGD1563746.aSep08</a>	<a href="#">303572</a>	4852	1460		321	similar to nuclear protein with a coiled coil-4 domain of bilateral origin like (3L720) (RGD1563746) mRNA.
<a href="#">RGD1563782</a>	<a href="#">RGD1563782.aSep08</a>	<a href="#">498115</a>	2439	1703	7	368	RGD1563782 (RGD1563782) alternative variant aSep08, complete mRNA.
<a href="#">RGD1563782</a>	<a href="#">RGD1563782.bSep08</a>	<a href="#">498115</a>	1039	952	3	302	RGD1563782 (RGD1563782) alternative variant bSep08, mRNA.
<a href="#">RGD1563815</a>	<a href="#">RGD1563815.aSep08</a>	<a href="#">365549</a>	43991	2689	3	628	similar to sodium-glucose cotransporter-like 1 (69.9 kD) (RGD1563815) alternative variant aSep08, mRNA.



<a href="#">RGD1563818</a>	<a href="#">RGD1563818.aSep08</a>	<a href="#">296356</a>	2211	606		128	similar to secretory leukocyte protease inhibitor (RGD1563818) mRNA.
<a href="#">RGD1563821</a>	<a href="#">RGD1563821.aSep08</a>	<a href="#">500984</a>	7187	497		105	RGD1563821 (RGD1563821) mRNA.
<a href="#">RGD1563838</a>	<a href="#">RGD1563838.aSep08</a>	<a href="#">361583</a>	118893	1783		547	similar to leucine zipper protein 2 (RGD1563838) mRNA.
<a href="#">RGD1563846</a>	<a href="#">RGD1563846.aSep08</a>	<a href="#">317224</a>	9655	1778		487	similar to Gene model 784 (RGD1563846) alternative variant aSep08, mRNA.
<a href="#">RGD1563853</a>	<a href="#">RGD1563853.bSep08</a>	<a href="#">361181</a>	9970	896	6	222	cullin 4A (RGD1563853) alternative variant bSep08, mRNA.
<a href="#">RGD1563853</a>	<a href="#">RGD1563853.cSep08</a>	<a href="#">361181</a>	15743	546	5	181	cullin 4 (RGD1563853) alternative variant cSep08, mRNA.
<a href="#">RGD1563853</a>	<a href="#">RGD1563853.dSep08</a>	<a href="#">361181</a>	5478	2371	4	159	cullin 4A (18.6 kD) (RGD1563853) alternative variant dSep08, mRNA.
<a href="#">RGD1563853</a>	<a href="#">RGD1563853.eSep08</a>	<a href="#">361181</a>	7065	515	6	58	cullin 4A (RGD1563853) alternative variant eSep08, mRNA.
<a href="#">RGD1563866</a>	<a href="#">RGD1563866.aSep08</a>	<a href="#">315652</a>	4556	972		204	RGD1563866 (22.5 kD) (RGD1563866) mRNA.
<a href="#">RGD1563870</a>	<a href="#">RGD1563870.aSep08</a>	<a href="#">500895</a>	460	392		119	similar to CG3104-PA, isoform A (RGD1563870) mRNA.
<a href="#">RGD1563874</a>	<a href="#">RGD1563874.aSep08</a>	<a href="#">302900</a>	1018	678		109	RGD1563874 (RGD1563874) mRNA.
<a href="#">RGD1563888</a>	<a href="#">RGD1563888.bSep08</a>	<a href="#">360692</a>	956	549	2	129	similar to DNA segment, Chr 16, ERATO Doi 472, expressed (RGD1563888) alternative variant bSep08, mRNA.
<a href="#">RGD1563888</a>	<a href="#">RGD1563888.cSep08</a>	<a href="#">360692</a>	25757	381	3	127	similar to DNA segment, Chr 16, ERATO Doi 472, expressed (RGD1563888) alternative variant cSep08, mRNA.
<a href="#">RGD1563912</a>	<a href="#">RGD1563912.aSep08</a>	<a href="#">362256</a>	7557	1129	6	97	RGD1563912 (10.6 kD) (RGD1563912) alternative variant aSep08, mRNA.
<a href="#">RGD1563912</a>	<a href="#">RGD1563912.bSep08</a>	<a href="#">362256</a>	8333	3528	5	97	RGD1563912 (10.6 kD) (RGD1563912) alternative variant bSep08, mRNA.
<a href="#">RGD1563912</a>	<a href="#">RGD1563912.cSep08</a>	<a href="#">362256</a>	1028	332	2	59	RGD1563912 (RGD1563912) alternative variant cSep08, mRNA.
<a href="#">RGD1563912</a>	<a href="#">RGD1563912.dSep08</a>	<a href="#">362256</a>	833	746	2	24	RGD1563912 (2.4 kD) (RGD1563912) alternative variant dSep08, mRNA.
<a href="#">RGD1563912</a>	<a href="#">RGD1563912.eSep08</a>	<a href="#">362256</a>	970	597	3	43	RGD1563912 (4.8 kD) (RGD1563912) alternative variant eSep08, mRNA.
<a href="#">RGD1563913</a>	<a href="#">RGD1563913.aSep08</a>	<a href="#">500691</a>	66883	770		77	RGD1563913 (8.7 kD) (RGD1563913) mRNA.
<a href="#">RGD1563917</a>	<a href="#">RGD1563917.aSep08</a>	<a href="#">501176</a>	13743	591		155	similar to Nuclear autoantigen Sp-100 (Speckled 100 kDa) (RGD1563917) mRNA.
<a href="#">RGD1563945</a>	<a href="#">RGD1563945.aSep08</a>	<a href="#">299305</a>	13910	632		210	similar to mKIAA0215 protein (RGD1563945) mRNA.
<a href="#">RGD1563946</a>	<a href="#">RGD1563946.aSep08</a>	<a href="#">306340</a>	2792	1390	6	462	similar to mKIAA1623 protein (RGD1563946) alternative variant aSep08, mRNA.
<a href="#">RGD1563946</a>	<a href="#">RGD1563946.bSep08</a>	<a href="#">306340</a>	914	497	1	165	similar to mKIAA1623 protein (RGD1563946) alternative variant bSep08, mRNA.
<a href="#">RGD1563952</a>	<a href="#">RGD1563952.bSep08</a>	<a href="#">363463</a>	21084	580	6	162	similar to Mospd2 protein (RGD1563952) alternative variant bSep08, mRNA.
<a href="#">RGD1563955</a>	<a href="#">RGD1563955.bSep08</a>	<a href="#">363274</a>	1119	790	2	179	similar to transmembrane serine protease 9 (19.5 kD) (RGD1563955) alternative variant bSep08, mRNA.
<a href="#">RGD1563955</a>	<a href="#">RGD1563955.cSep08</a>	<a href="#">363274</a>	1563	474	5	158	similar to transmembrane serine protease 9 (RGD1563955) alternative variant cSep08, mRNA.

<a href="#">RGD1563973</a>	<a href="#">RGD1563973.aSep08</a>	<a href="#">366466</a>	4609	1598		452	similar to Gene model 50 (RGD1563973) alternative variant aSep08, mRNA.
<a href="#">RGD1563979</a>	<a href="#">RGD1563979.aSep08</a>	<a href="#">316212</a>	7486	1147	6	328	similar to over-expressed breast tumor protein (RGD1563979) alternative variant aSep08, mRNA.
<a href="#">RGD1563979</a>	<a href="#">RGD1563979.bSep08</a>	<a href="#">316212</a>	4551	678	5	225	similar to over-expressed breast tumor protein (RGD1563979) alternative variant bSep08, mRNA.
<a href="#">RGD1563996</a>	<a href="#">RGD1563996.aSep08</a>	<a href="#">315163</a>	3949	607		202	similar to Protein UNQ9166/PRO28631 precursor (RGD1563996) mRNA.
<a href="#">RGD1564005</a>	<a href="#">RGD1564005.bSep08</a>	<a href="#">360575</a>	725	639	2	92	similar to novel protein (RGD1564005) alternative variant bSep08, mRNA.
<a href="#">RGD1564016</a>	<a href="#">RGD1564016.aSep08</a>	<a href="#">498969</a>	838	745	1	90	RGD1564016 (9.9 kD) (RGD1564016) alternative variant aSep08, mRNA.
<a href="#">RGD1564016</a>	<a href="#">RGD1564016.bSep08</a>	<a href="#">498969</a>	50664	770	3	85	RGD1564016 (9.5 kD) (RGD1564016) alternative variant bSep08, mRNA.
<a href="#">RGD1564036</a>	<a href="#">RGD1564036.bSep08</a>	<a href="#">497895</a>	24562	740	2	115	similar to RIKEN cDNA 3010026O09 (RGD1564036) alternative variant bSep08, mRNA.
<a href="#">RGD1564036</a>	<a href="#">RGD1564036.cSep08</a>	<a href="#">497895</a>	26880	1204	2	224	similar to RIKEN cDNA 3010026O09 (24.5 kD) (RGD1564036) alternative variant cSep08, mRNA.
<a href="#">RGD1564046</a>	<a href="#">RGD1564046.aSep08</a>	<a href="#">500823</a>	34520	1468	11	489	similar to RIKEN cDNA 4933417K05 gene (RGD1564046) alternative variant aSep08, mRNA.
<a href="#">RGD1564046</a>	<a href="#">RGD1564046.bSep08</a>	<a href="#">500823</a>	3709	611	2	166	similar to RIKEN cDNA 4933417K05 gene (19.6 kD) (RGD1564046) alternative variant bSep08, mRNA.
<a href="#">RGD1564053</a>	<a href="#">RGD1564053.bSep08</a>	<a href="#">500390</a>	129437	496	4	73	similar to hypothetical protein (RGD1564053) alternative variant bSep08, mRNA.
<a href="#">RGD1564058</a>	<a href="#">RGD1564058.bSep08</a>	<a href="#">288925</a>	1562	880	2	106	putative nuclear protein of eukaryotic origin (12.1 kD) (RGD1564058) alternative variant bSep08, mRNA.
<a href="#">RGD1564058</a>	<a href="#">RGD1564058.cSep08</a>	<a href="#">288925</a>	1326	716	1	90	putative protein of eukaryotic origin (RGD1564058) alternative variant cSep08, mRNA.
<a href="#">RGD1564060</a>	<a href="#">RGD1564060.aSep08</a>	<a href="#">315981</a>	16841	742		247	similar to procollagen, type VI, alpha 3 isoform 4 (RGD1564060) mRNA.
<a href="#">RGD1564074</a>	<a href="#">RGD1564074.aSep08</a>	<a href="#">500514</a>	80906	1156	15	384	similar to novel protein (RGD1564074) alternative variant aSep08, mRNA.
<a href="#">RGD1564081</a>	<a href="#">RGD1564081.aSep08</a>	<a href="#">500943</a>	130201	409		136	similar to novel protein similar to human oligophrenin 1 (OPHN1) (RGD1564081) mRNA.
<a href="#">RGD1564086</a>	<a href="#">RGD1564086.aSep08</a>	<a href="#">497862</a>	1016	303		95	RGD1564086 (RGD1564086) mRNA.
<a href="#">RGD1564091</a>	<a href="#">RGD1564091.aSep08</a>	<a href="#">500633</a>	10157	1801		600	similar to Kiaa0575 (RGD1564091) alternative variant aSep08, mRNA.
<a href="#">RGD1564091</a>	<a href="#">RGD1564091.bSep08</a>	<a href="#">500633</a>	5901	534		177	similar to Kiaa0575 (RGD1564091) alternative variant bSep08, mRNA.
<a href="#">RGD1564114</a>	<a href="#">RGD1564114.bSep08</a>	<a href="#">499765</a>	109069	774	1	225	similar to FLJ46082 protein (RGD1564114) alternative variant bSep08, mRNA.
<a href="#">RGD1564117</a>	<a href="#">RGD1564117.aSep08</a>	<a href="#">317253</a>	3840	1098		169	similar to novel protein similar to multidomain presynaptic cytomatrix protein piccolo (presynaptic cytomatrix protein) (RGD1564117) alternative variant aSep08, mRNA.
<a href="#">RGD1564120</a>	<a href="#">RGD1564120.aSep08</a>	<a href="#">499461</a>	2197	527		112	similar to WNT1 inducible signaling pathway protein 3 precursor (WISP-3) (RGD1564120) mRNA.
<a href="#">RGD1564122</a>	<a href="#">RGD1564122.aSep08</a>	<a href="#">362888</a>	16292	765		240	RGD1564122 (RGD1564122) mRNA.

<a href="#">RGD1564129</a>	<a href="#">RGD1564129.aSep08</a>	<a href="#">361257</a>	2475	1063		243	similar to hypothetical protein 4930474N05 (RGD1564129) mRNA.
<a href="#">RGD1564140</a>	<a href="#">RGD1564140.bSep08</a>	<a href="#">498832</a>	57657	1483	2	155	similar to AW554918 protein (RGD1564140) alternative variant bSep08, mRNA.
<a href="#">RGD1564161</a>	<a href="#">RGD1564161.aSep08</a>	<a href="#">362562</a>	422311	466	2	154	atp gtp binding protein-like 4 (RGD1564161) alternative variant aSep08, mRNA.
<a href="#">RGD1564161</a>	<a href="#">RGD1564161.aSep08</a>	<a href="#">689771</a>	422311	466	2	154	atp gtp binding protein-like 4 (RGD1564161) alternative variant aSep08, mRNA.
<a href="#">RGD1564161</a>	<a href="#">RGD1564161.bSep08</a>	<a href="#">362562</a>	211012	480	1	106	atp gtp binding protein-like 4 (RGD1564161) alternative variant bSep08, mRNA.
<a href="#">RGD1564161</a>	<a href="#">RGD1564161.bSep08</a>	<a href="#">689771</a>	211012	480	1	106	atp gtp binding protein-like 4 (RGD1564161) alternative variant bSep08, mRNA.
<a href="#">RGD1564162</a>	<a href="#">RGD1564162.aSep08</a>	<a href="#">499421</a>	2706	1608		158	similar to Homo sapiens fetal lung specific expression~unknown (17.6 kD) (RGD1564162) mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.aSep08</a>	<a href="#">361985</a>	16527	1870	10	253	similar to NICE-3 and RGD1564171 (28.8 kD) (RGD1564171) alternative variant aSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.aSep08</a>	<a href="#">499656</a>	16527	1870	10	253	similar to NICE-3 and RGD1564171 (28.8 kD) (RGD1564171) alternative variant aSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.cSep08</a>	<a href="#">361985</a>	8140	1801	5	186	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant cSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.cSep08</a>	<a href="#">499656</a>	8140	1801	5	186	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant cSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.dSep08</a>	<a href="#">361985</a>	8319	761	5	183	similar to NICE-3 and RGD1564171 (20.9 kD) (RGD1564171) alternative variant dSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.dSep08</a>	<a href="#">499656</a>	8319	761	5	183	similar to NICE-3 and RGD1564171 (20.9 kD) (RGD1564171) alternative variant dSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.eSep08</a>	<a href="#">361985</a>	8346	625	6	164	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant eSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.eSep08</a>	<a href="#">499656</a>	8346	625	6	164	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant eSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.fSep08</a>	<a href="#">361985</a>	8429	658	6	133	similar to NICE-3 and RGD1564171 (15.2 kD) (RGD1564171) alternative variant fSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.fSep08</a>	<a href="#">499656</a>	8429	658	6	133	similar to NICE-3 and RGD1564171 (15.2 kD) (RGD1564171) alternative variant fSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.gSep08</a>	<a href="#">361985</a>	12289	857	5	126	similar to NICE-3 and RGD1564171 (14.3 kD) (RGD1564171) alternative variant gSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.gSep08</a>	<a href="#">499656</a>	12289	857	5	126	similar to NICE-3 and RGD1564171 (14.3 kD) (RGD1564171) alternative variant gSep08, mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.iSep08</a>	<a href="#">361985</a>	12908	1368	4	114	similar to NICE-3 and RGD1564171 (13.0 kD) (RGD1564171) alternative variant iSep08, complete mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.iSep08</a>	<a href="#">499656</a>	12908	1368	4	114	similar to NICE-3 and RGD1564171 (13.0 kD) (RGD1564171) alternative variant iSep08, complete mRNA.
<a href="#">RGD1564171</a>	<a href="#">RGD1564171.kSep08</a>	<a href="#">361985</a>	12010	897	4	100	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant kSep08, mRNA.

<a href="#">RGD1564171</a>	<a href="#">RGD1564171.kSep08</a>	<a href="#">499656</a>	12010	897	4	100	similar to NICE-3 and RGD1564171 (RGD1564171) alternative variant kSep08, mRNA.
<a href="#">RGD1564174</a>	<a href="#">RGD1564174.aSep08</a>	<a href="#">360980</a>	147910	1947	13	430	similar to novel protein similar to Tensin Tns (48.1 kD) (RGD1564174) alternative variant aSep08, mRNA.
<a href="#">RGD1564174</a>	<a href="#">RGD1564174.bSep08</a>	<a href="#">360980</a>	7473	525	1	108	similar to novel protein similar to Tensin Tns (RGD1564174) alternative variant bSep08, mRNA.
<a href="#">RGD1564177</a>	<a href="#">RGD1564177.aSep08</a>	<a href="#">501592</a>	633	369	1	105	RGD1564177 (RGD1564177) alternative variant aSep08, mRNA.
<a href="#">RGD1564177</a>	<a href="#">RGD1564177.bSep08</a>	<a href="#">501592</a>	722	396	1	75	RGD1564177 (8.7 kD) (RGD1564177) alternative variant bSep08, mRNA.
<a href="#">RGD1564195</a>	<a href="#">RGD1564195.aSep08</a>	<a href="#">361651</a>	1736	1227	3	195	similar to hypothetical protein (RGD1564195) alternative variant aSep08, mRNA.
<a href="#">RGD1564195</a>	<a href="#">RGD1564195.dSep08</a>	<a href="#">361651</a>	2430	433	4	110	similar to hypothetical protein (RGD1564195) alternative variant dSep08, mRNA.
<a href="#">RGD1564195</a>	<a href="#">RGD1564195.eSep08</a>	<a href="#">361651</a>	2446	459	3	79	similar to hypothetical protein (RGD1564195) alternative variant eSep08, mRNA.
<a href="#">RGD1564200</a>	<a href="#">RGD1564200.bSep08</a>	<a href="#">366311</a>	12404	442		90	putative mitochondrial protein (9.7 kD) (RGD1564200) alternative variant bSep08, mRNA.
<a href="#">RGD1564209</a>	<a href="#">RGD1564209.aSep08</a>	<a href="#">367046</a>	3084	1111	3	103	similar to Acyl-CoA dehydrogenase family member 8, mitochondrial precursor (ACAD-8) (RGD1564209) alternative variant aSep08, mRNA.
<a href="#">RGD1564214</a>	<a href="#">RGD1564214.aSep08</a>	<a href="#">499093</a>	5077	598	4	102	similar to Zfp93 protein (RGD1564214) alternative variant aSep08, mRNA.
<a href="#">RGD1564214</a>	<a href="#">RGD1564214.bSep08</a>	<a href="#">499093</a>	4974	413	3	65	similar to Zfp93 protein (RGD1564214) alternative variant bSep08, mRNA.
<a href="#">RGD1564214</a>	<a href="#">RGD1564214.dSep08</a>	<a href="#">499093</a>	1180	351	2	42	similar to Zfp93 protein (RGD1564214) alternative variant dSep08, mRNA.
<a href="#">RGD1564241</a>	<a href="#">RGD1564241.bSep08</a>	<a href="#">363022</a>	15582	1083	5	97	similar to zinc finger protein 426 (11.5 kD) (RGD1564241) alternative variant bSep08, mRNA.
<a href="#">RGD1564257</a>	<a href="#">RGD1564257.aSep08</a>	<a href="#">500595</a>	2691	453	3	133	similar to hypothetical protein FLJ32825 (RGD1564257) alternative variant aSep08, mRNA.
<a href="#">RGD1564257</a>	<a href="#">RGD1564257.bSep08</a>	<a href="#">500595</a>	2602	519	4	115	similar to hypothetical protein FLJ32825 (RGD1564257) alternative variant bSep08, mRNA.
<a href="#">RGD1564257</a>	<a href="#">RGD1564257.cSep08</a>	<a href="#">500595</a>	2349	606	2	59	similar to hypothetical protein FLJ32825 (RGD1564257) alternative variant cSep08, mRNA.
<a href="#">RGD1564259</a>	<a href="#">RGD1564259.aSep08</a>	<a href="#">292477</a>	35247	1229		386	similar to OTTHUMP00000040155 (RGD1564259) mRNA.
<a href="#">RGD1564263</a>	<a href="#">RGD1564263.aSep08</a>	<a href="#">499873</a>	9159	1829		241	similar to Opa-interacting protein 5 (RGD1564263) mRNA.
<a href="#">RGD1564272</a>	<a href="#">RGD1564272.aSep08</a>	<a href="#">499065</a>	1052	283		94	similar to RIKEN cDNA 9930022N03 gene (RGD1564272) mRNA.
<a href="#">RGD1564289</a>	<a href="#">RGD1564289.aSep08</a>	<a href="#">367793</a>	14598	670		169	similar to transcriptional repressor Scml2 (RGD1564289) mRNA.
<a href="#">RGD1564300</a>	<a href="#">RGD1564300.aSep08</a>	<a href="#">361661</a>	10393	1785	4	496	CRA c (RGD1564300) alternative variant aSep08, mRNA.
<a href="#">RGD1564300</a>	<a href="#">RGD1564300.bSep08</a>	<a href="#">361661</a>	10362	1082	6	218	CRA d (RGD1564300) alternative variant bSep08, mRNA.
<a href="#">RGD1564300</a>	<a href="#">RGD1564300.cSep08</a>	<a href="#">361661</a>	15282	612	3	75	CRA d (RGD1564300) alternative variant cSep08, mRNA.
<a href="#">RGD1564300</a>	<a href="#">RGD1564300.dSep08</a>	<a href="#">361661</a>	3589	606	2	71	CRA c (8.4 kD) (RGD1564300) alternative variant dSep08, mRNA.

<a href="#">RGD1564300</a>	<a href="#">RGD1564300.fSep08</a>	<a href="#">361661</a>	8173	1281	4	69	CRA c (RGD1564300) alternative variant fSep08, mRNA.
<a href="#">RGD1564300</a>	<a href="#">RGD1564300.gSep08</a>	<a href="#">361661</a>	8170	262	2	32	putative protein (4.0 kD) (RGD1564300) alternative variant gSep08, mRNA.
<a href="#">RGD1564308</a>	<a href="#">RGD1564308.aSep08</a>	<a href="#">502086</a>	3253	383		78	similar to LOC495042 protein (RGD1564308) mRNA.
<a href="#">RGD1564311</a>	<a href="#">RGD1564311.aSep08</a>	<a href="#">300461</a>	21970	748		52	similar to hypothetical protein FLJ32949 (RGD1564311) alternative variant aSep08, mRNA.
<a href="#">RGD1564322</a>	<a href="#">RGD1564322.aSep08</a>	<a href="#">500917</a>	979	652	2	133	similar to shippo 1 (RGD1564322) alternative variant aSep08, mRNA.
<a href="#">RGD1564322</a>	<a href="#">RGD1564322.bSep08</a>	<a href="#">500917</a>	729	374	2	105	similar to shippo 1 (RGD1564322) alternative variant bSep08, mRNA.
<a href="#">RGD1564324</a>	<a href="#">RGD1564324.bSep08</a>	<a href="#">290217</a>	1923	404	1	62	similar to dehydrogenase/reductase member 2 (RGD1564324) alternative variant bSep08, mRNA.
<a href="#">RGD1564327</a>	<a href="#">RGD1564327.aSep08</a>	<a href="#">364786</a>	47731	3249	3	190	similar to integrin alpha 8 (RGD1564327) alternative variant aSep08, mRNA.
<a href="#">RGD1564327</a>	<a href="#">RGD1564327.bSep08</a>	<a href="#">364786</a>	22303	491	1	133	similar to integrin alpha 8 (RGD1564327) alternative variant bSep08, mRNA.
<a href="#">RGD1564342</a>	<a href="#">RGD1564342.aSep08</a>	<a href="#">305710</a>	5335	547		182	similar to hypothetical protein FLJ32685 (RGD1564342) mRNA.
<a href="#">RGD1564357</a>	<a href="#">RGD1564357.aSep08</a>	<a href="#">502306</a>	1860	818		35	RGD1564357 (RGD1564357) mRNA.
<a href="#">RGD1564369</a>	<a href="#">RGD1564369.aSep08</a>	<a href="#">302832</a>	70602	714		85	RGD1564369 (9.6 kD) (RGD1564369) mRNA.
<a href="#">RGD1564379</a>	<a href="#">RGD1564379.aSep08</a>	<a href="#">499758</a>	1610	715	2	126	putative protein, with at least 2 transmembrane domains, of vertebrate origin (RGD1564379) alternative variant aSep08, mRNA.
<a href="#">RGD1564380</a>	<a href="#">RGD1564380.aSep08</a>	<a href="#">292711</a>	1133	765		183	similar to BC049730 protein (RGD1564380) mRNA.
<a href="#">RGD1564391</a>	<a href="#">RGD1564391.aSep08</a>	<a href="#">361199</a>	3222	755	1	140	RGD1564391 (RGD1564391) alternative variant aSep08, mRNA.
<a href="#">RGD1564391</a>	<a href="#">RGD1564391.bSep08</a>	<a href="#">361199</a>	4880	1048	2	89	RGD1564391 (10.1 kD) (RGD1564391) alternative variant bSep08, mRNA.
<a href="#">RGD1564406</a>	<a href="#">RGD1564406.aSep08</a>	<a href="#">501586</a>	593	416		91	similar to RIKEN cDNA 1700019M22 (RGD1564406) mRNA.
<a href="#">RGD1564419</a>	<a href="#">RGD1564419.bSep08</a>	<a href="#">500128</a>	4625	1215	2	138	similar to hypothetical gene supported by BC025338 (14.7 kD) (RGD1564419) alternative variant bSep08, mRNA.
<a href="#">RGD1564419</a>	<a href="#">RGD1564419.dSep08</a>	<a href="#">500128</a>	1645	1443	2	76	similar to hypothetical gene supported by BC025338 (7.9 kD) (RGD1564419) alternative variant dSep08, mRNA.
<a href="#">RGD1564428</a>	<a href="#">RGD1564428.aSep08</a>	<a href="#">498866</a>	21372	424		108	similar to FLJ32921 protein (12.4 kD) (RGD1564428) alternative variant aSep08, mRNA.
<a href="#">RGD1564450</a>	<a href="#">RGD1564450.bSep08</a>	<a href="#">294291</a>	1752	379	5	125	RGD1564450 (RGD1564450) alternative variant bSep08, mRNA.
<a href="#">RGD1564454</a>	<a href="#">RGD1564454.aSep08</a>	<a href="#">499690</a>	5855	2813		135	similar to RIKEN cDNA 2010200O16 (RGD1564454) mRNA.
<a href="#">RGD1564456</a>	<a href="#">RGD1564456.aSep08</a>	<a href="#">361269</a>	40437	2764		828	peripheral benzodiazepine receptor associated protein (RGD1564456) mRNA.
<a href="#">RGD1564482</a>	<a href="#">RGD1564482.aSep08</a>	<a href="#">500568</a>	1565	550	2	114	RGD1564482 (RGD1564482) alternative variant aSep08, mRNA.
<a href="#">RGD1564482</a>	<a href="#">RGD1564482.bSep08</a>	<a href="#">500568</a>	1429	675	1	110	RGD1564482 (RGD1564482) alternative variant bSep08, mRNA.

<a href="#">RGD1564491</a>	<a href="#">RGD1564491.bSep08</a>	<a href="#">303812</a>	3578	1272	5	327	RGD1564491 (RGD1564491) alternative variant bSep08, mRNA.
<a href="#">RGD1564491</a>	<a href="#">RGD1564491.cSep08</a>	<a href="#">303812</a>	3502	537	3	179	RGD1564491 (RGD1564491) alternative variant cSep08, mRNA.
<a href="#">RGD1564496</a>	<a href="#">RGD1564496.aSep08</a>	<a href="#">499650</a>	1795	542		151	RGD1564496 (RGD1564496) mRNA.
<a href="#">RGD1564515</a>	<a href="#">RGD1564515.aSep08</a>	<a href="#">299963</a>	1618	750		244	similar to alpha 1B-glycoprotein (RGD1564515) mRNA.
<a href="#">RGD1564516</a>	<a href="#">RGD1564516.aSep08</a>	<a href="#">497891</a>	12537	431		52	similar to smuckler (6.0 kD) (RGD1564516) mRNA.
<a href="#">RGD1564522</a>	<a href="#">RGD1564522.aSep08</a>	<a href="#">305567</a>	1427	618		99	similar to Hypothetical protein CBG23547 (RGD1564522) mRNA.
<a href="#">RGD1564560</a>	<a href="#">RGD1564560.bSep08</a>	<a href="#">500988</a>	8404	717	4	103	dead box polypeptide 6 CRA a (10.8 kD) (RGD1564560) alternative variant bSep08, mRNA.
<a href="#">RGD1564560</a>	<a href="#">RGD1564560.cSep08</a>	<a href="#">500988</a>	3599	557	2	83	putative protein (RGD1564560) alternative variant cSep08, mRNA.
<a href="#">RGD1564560</a>	<a href="#">RGD1564560.fSep08</a>	<a href="#">500988</a>	2979	433	2	52	dead box polypeptide 6 CRA a (RGD1564560) alternative variant fSep08, mRNA.
<a href="#">RGD1564560</a>	<a href="#">RGD1564560.gSep08</a>	<a href="#">500988</a>	1280	289	2	36	putative protein (RGD1564560) alternative variant gSep08, mRNA.
<a href="#">RGD1564567</a>	<a href="#">RGD1564567.aSep08</a>	<a href="#">305453</a>	5621	653	1	217	similar to CG14998-PC, isoform C (RGD1564567) alternative variant aSep08, mRNA.
<a href="#">RGD1564567</a>	<a href="#">RGD1564567.bSep08</a>	<a href="#">305453</a>	3738	706	1	155	similar to CG14998-PC, isoform C (RGD1564567) alternative variant bSep08, mRNA.
<a href="#">RGD1564579</a>	<a href="#">RGD1564579.aSep08</a>	<a href="#">293491</a>	3303	954	5	185	similar to yippee-like 3 (RGD1564579) alternative variant aSep08, mRNA.
<a href="#">RGD1564579</a>	<a href="#">RGD1564579.bSep08</a>	<a href="#">293491</a>	1322	447	5	149	similar to yippee-like 3 (RGD1564579) alternative variant bSep08, mRNA.
<a href="#">RGD1564579</a>	<a href="#">RGD1564579.cSep08</a>	<a href="#">293491</a>	2884	426	4	138	similar to yippee-like 3 (RGD1564579) alternative variant cSep08, mRNA.
<a href="#">RGD1564579</a>	<a href="#">RGD1564579.dSep08</a>	<a href="#">293491</a>	2146	559	3	82	similar to yippee-like 3 (RGD1564579) alternative variant dSep08, mRNA.
<a href="#">RGD1564579</a>	<a href="#">RGD1564579.eSep08</a>	<a href="#">293491</a>	3293	1819	3	54	similar to yippee-like 3 (6.2 kD) (RGD1564579) alternative variant eSep08, complete mRNA.
<a href="#">RGD1564591</a>	<a href="#">RGD1564591.bSep08</a>	<a href="#">501125</a>	23215	732	1	91	putative protein of mammalian origin (RGD1564591) alternative variant bSep08, mRNA.
<a href="#">RGD1564614</a>	<a href="#">RGD1564614.bSep08</a>	<a href="#">498241</a>	24017	746	2	76	similar to complement factor H-related protein (RGD1564614) alternative variant bSep08, mRNA.
<a href="#">RGD1564615</a>	<a href="#">RGD1564615.aSep08</a>	<a href="#">312371</a>	14344	905	8	203	similar to mKIAA0241 protein (23.3 kD) (RGD1564615) alternative variant aSep08, mRNA.
<a href="#">RGD1564615</a>	<a href="#">RGD1564615.bSep08</a>	<a href="#">312371</a>	7911	349		64	similar to mKIAA0241 protein (RGD1564615) alternative variant bSep08, mRNA.
<a href="#">RGD1564620</a>	<a href="#">RGD1564620.aSep08</a>	<a href="#">365962</a>	15391	623		207	similar to hypothetical protein MGC34032 (RGD1564620) mRNA.
<a href="#">RGD1564635</a>	<a href="#">RGD1564635.aSep08</a>	<a href="#">498355</a>	58551	510	3	124	similar to CG12206-PA, isoform A (RGD1564635) alternative variant aSep08, mRNA.
<a href="#">RGD1564635</a>	<a href="#">RGD1564635.cSep08</a>	<a href="#">498355</a>	58520	413	2	96	similar to CG12206-PA, isoform A (RGD1564635) alternative variant cSep08, mRNA.
<a href="#">RGD1564638</a>	<a href="#">RGD1564638.aSep08</a>	<a href="#">287060</a>	22080	579		192	similar to KIAA0420 (RGD1564638) mRNA.

<a href="#">RGD1564641</a>	<a href="#">RGD1564641.aSep08</a>	<a href="#">303891</a>	8866	422		140	similar to Neurogenic locus notch homolog protein 2 precursor (Notch 2) (hN2) (RGD1564641) mRNA.
<a href="#">RGD1564651</a>	<a href="#">RGD1564651.aSep08</a>	<a href="#">292572</a>	18874	729		197	similar to oocyte specific homeobox 3 (RGD1564651) mRNA.
<a href="#">RGD1564655</a>	<a href="#">RGD1564655.aSep08</a>	<a href="#">292014</a>	10538	1067		294	similar to Tle2 protein (RGD1564655) mRNA.
<a href="#">RGD1564657</a>	<a href="#">RGD1564657.aSep08</a>	<a href="#">306718</a>	2730	675		204	similar to cathepsin 1 precursor (RGD1564657) mRNA.
<a href="#">RGD1564666</a>	<a href="#">RGD1564666.aSep08</a>	<a href="#">499627</a>	1221	572		119	RGD1564666 (RGD1564666) mRNA.
<a href="#">RGD1564672</a>	<a href="#">RGD1564672.aSep08</a>	<a href="#">499520</a>	12692	239		50	similar to hypothetical protein MGC52498 (RGD1564672) mRNA.
<a href="#">RGD1564680</a>	<a href="#">RGD1564680.aSep08</a>	<a href="#">312115</a>	2072	811		116	similar to matrilin 2 precursor (RGD1564680) mRNA.
<a href="#">RGD1564709</a>	<a href="#">RGD1564709.bSep08</a>	<a href="#">305142</a>	19043	357	1	115	similar to ATP-binding cassette, sub-family G (WHITE), member 3 (RGD1564709) alternative variant bSep08, mRNA.
<a href="#">RGD1564722</a>	<a href="#">RGD1564722.aSep08</a>	<a href="#">362832</a>	2615	797	9	126	similar to C19orf36 protein (14.8 kD) (RGD1564722) alternative variant aSep08, mRNA.
<a href="#">RGD1564722</a>	<a href="#">RGD1564722.cSep08</a>	<a href="#">362832</a>	832	651	3	66	similar to C19orf36 protein (6.7 kD) (RGD1564722) alternative variant cSep08, mRNA.
<a href="#">RGD1564731</a>	<a href="#">RGD1564731.aSep08</a>	<a href="#">363824</a>	2679	756		62	similar to 60S ribosomal protein L7a (RGD1564731) mRNA.
<a href="#">RGD1564739</a>	<a href="#">RGD1564739.aSep08</a>	<a href="#">296782</a>	52949	685		173	similar to spermatogenesis associated glutamate (E)-rich protein 4d (RGD1564739) mRNA.
<a href="#">RGD1564743</a>	<a href="#">RGD1564743.aSep08</a>	<a href="#">499534</a>	6220	1500		59	RGD1564743 (RGD1564743) mRNA.
<a href="#">RGD1564752</a>	<a href="#">RGD1564752.aSep08</a>	<a href="#">363050</a>	96933	598	3	80	RGD1564752 (9.2 kD) (RGD1564752) mRNA.
<a href="#">RGD1564763</a>	<a href="#">RGD1564763.aSep08</a>	<a href="#">367747</a>	5057	943		201	similar to nudix (nucleoside diphosphate linked moiety X)-type motif 11 (RGD1564763) mRNA.
<a href="#">RGD1564770</a>	<a href="#">RGD1564770.aSep08</a>	<a href="#">500335</a>	10917	1567		185	similar to CD69 antigen (p60, early T-cell activation antigen) (21.6 kD) (RGD1564770) complete mRNA.
<a href="#">RGD1564773</a>	<a href="#">RGD1564773.aSep08</a>	<a href="#">361142</a>	59819	567		148	similar to GalNAc transferase 10 isoform a (RGD1564773) mRNA.
<a href="#">RGD1564776</a>	<a href="#">RGD1564776.bSep08</a>	<a href="#">500550</a>	1759	886	1	107	similar to ornithine decarboxylase-like protein (RGD1564776) alternative variant bSep08, mRNA.
<a href="#">RGD1564778</a>	<a href="#">RGD1564778.bSep08</a>	<a href="#">303514</a>	5135	2034	3	453	similar to RIKEN cDNA 4121402D02 (RGD1564778) alternative variant bSep08, mRNA.
<a href="#">RGD1564778</a>	<a href="#">RGD1564778.cSep08</a>	<a href="#">303514</a>	5907	1225	8	339	similar to RIKEN cDNA 4121402D02 (RGD1564778) alternative variant cSep08, mRNA.
<a href="#">RGD1564788</a>	<a href="#">RGD1564788.aSep08</a>	<a href="#">290805</a>	8268	880	4	200	similar to Werner syndrome helicase homolog (RGD1564788) alternative variant aSep08, mRNA.
<a href="#">RGD1564788</a>	<a href="#">RGD1564788.bSep08</a>	<a href="#">290805</a>	13222	917	4	187	similar to Werner syndrome helicase homolog (RGD1564788) alternative variant bSep08, mRNA.
<a href="#">RGD1564788</a>	<a href="#">RGD1564788.cSep08</a>	<a href="#">290805</a>	11410	1256	4	170	similar to Werner syndrome helicase homolog (18.9 kD) (RGD1564788) alternative variant cSep08, mRNA.
<a href="#">RGD1564791</a>	<a href="#">RGD1564791.aSep08</a>	<a href="#">498766</a>	6755	778	5	122	similar to hypothetical protein 4930474N05 (RGD1564791) mRNA.
<a href="#">RGD1564792</a>	<a href="#">RGD1564792.bSep08</a>	<a href="#">499976</a>	8644	611	2	53	RGD1564792 (RGD1564792) alternative variant bSep08, mRNA.

<a href="#">RGD1564793</a>	<a href="#">RGD1564793.aSep08</a>	<a href="#">691337</a>	50434	903		184	similar to KIAA0965 protein (21.4 kD) (RGD1564793) mRNA.
<a href="#">RGD1564803</a>	<a href="#">RGD1564803.aSep08</a>	<a href="#">498181</a>	1144	484		142	RGD1564803 (RGD1564803) mRNA.
<a href="#">RGD1564804</a>	<a href="#">RGD1564804.bSep08</a>	<a href="#">313551</a>	3157	1461	2	88	putative nuclear protein of ancient origin (10.1 kD) (RGD1564804) alternative variant bSep08, mRNA.
<a href="#">RGD1564809</a>	<a href="#">RGD1564809.aSep08</a>	<a href="#">313208</a>	32589	529	6	146	similar to fibronectin type 3 and SPRY domain-containing protein (RGD1564809) alternative variant aSep08, mRNA.
<a href="#">RGD1564811</a>	<a href="#">RGD1564811.bSep08</a>	<a href="#">316157</a>	11827	818	1	164	putative protein of metazoan origin (RGD1564811) alternative variant bSep08, mRNA.
<a href="#">RGD1564827</a>	<a href="#">RGD1564827.aSep08</a>	<a href="#">498689</a>	4760	426		31	similar to cathepsin M (RGD1564827) mRNA.
<a href="#">RGD1564833</a>	<a href="#">RGD1564833.aSep08</a>	<a href="#">498297</a>	17018	3411	2	387	similar to 9630058J23Rik protein (RGD1564833) alternative variant aSep08, mRNA.
<a href="#">RGD1564833</a>	<a href="#">RGD1564833.bSep08</a>	<a href="#">498297</a>	12118	414	1	131	similar to 9630058J23Rik protein (RGD1564833) alternative variant bSep08, mRNA.
<a href="#">RGD1564841</a>	<a href="#">RGD1564841.aSep08</a>	<a href="#">361498</a>	5917	671		117	similar to NALP-alpha (RGD1564841) mRNA.
<a href="#">RGD1564845</a>	<a href="#">RGD1564845.aSep08</a>	<a href="#">363507</a>	4152	739		143	similar to Xlr-like (RGD1564845) mRNA.
<a href="#">RGD1564854</a>	<a href="#">RGD1564854.bSep08</a>	<a href="#">502617</a>	17586	782	6	135	similar to divalent cation tolerant protein CUTA (15.4 kD) (RGD1564854) alternative variant bSep08, mRNA.
<a href="#">RGD1564859</a>	<a href="#">RGD1564859.aSep08</a>	<a href="#">362536</a>	42562	1349	2	117	RGD1564859 (13.2 kD) (RGD1564859) alternative variant aSep08, complete mRNA.
<a href="#">RGD1564859</a>	<a href="#">RGD1564859.bSep08</a>	<a href="#">362536</a>	33360	619	2	72	RGD1564859 (8.1 kD) (RGD1564859) alternative variant bSep08, mRNA.
<a href="#">RGD1564859</a>	<a href="#">RGD1564859.dSep08</a>	<a href="#">362536</a>	41846	967	4	81	RGD1564859 (RGD1564859) alternative variant dSep08, mRNA.
<a href="#">RGD1564865</a>	<a href="#">RGD1564865.aSep08</a>	<a href="#">498789</a>	6123	413	2	64	similar to 20-alpha-hydroxysteroid dehydrogenase (RGD1564865) alternative variant aSep08, mRNA.
<a href="#">RGD1564865</a>	<a href="#">RGD1564865.bSep08</a>	<a href="#">498789</a>	1808	294	1	24	similar to 20-alpha-hydroxysteroid dehydrogenase (RGD1564865) alternative variant bSep08, mRNA.
<a href="#">RGD1564871</a>	<a href="#">RGD1564871.aSep08</a>	<a href="#">315947</a>	20697	707	5	171	thioredoxin domain containing protein (RGD1564871) alternative variant aSep08, mRNA.
<a href="#">RGD1564871</a>	<a href="#">RGD1564871.bSep08</a>	<a href="#">315947</a>	7199	1520	2	58	putative protein (RGD1564871) alternative variant bSep08, mRNA.
<a href="#">RGD1564877</a>	<a href="#">RGD1564877.aSep08</a>	<a href="#">501581</a>	675	498		118	RGD1564877 (RGD1564877) mRNA.
<a href="#">RGD1564878</a>	<a href="#">RGD1564878.aSep08</a>	<a href="#">502002</a>	769	381		88	similar to natural killer cell protease 7 (RGD1564878) mRNA.
<a href="#">RGD1564887</a>	<a href="#">RGD1564887.aSep08</a>	<a href="#">499363</a>	81845	2079		187	similar to 9130011E15Rik protein (RGD1564887) mRNA.
<a href="#">RGD1564907</a>	<a href="#">RGD1564907.aSep08</a>	<a href="#">498723</a>	15400	705		95	RGD1564907 (RGD1564907) mRNA.
<a href="#">RGD1564927and dRGD1560115</a>	<a href="#">RGD1564927andRGD1 560115.aSep08</a>	<a href="#">499929</a>	18275	2888	1	237	similar to TGFB-induced factor 2 and similar to TGFB-induced factor 2 (RGD1564927andRGD1560115) alternative variant aSep08, mRNA.
<a href="#">RGD1564927and dRGD1560115</a>	<a href="#">RGD1564927andRGD1 560115.aSep08</a>	<a href="#">499930</a>	18275	2888	1	237	similar to TGFB-induced factor 2 and similar to TGFB-induced factor 2 (RGD1564927andRGD1560115) alternative variant aSep08, mRNA.
<a href="#">RGD1564927and dRGD1560115</a>	<a href="#">RGD1564927andRGD1 560115.bSep08</a>	<a href="#">499929</a>	13052	756	2	179	similar to TGFB-induced factor 2 and similar to TGFB-induced factor 2 (RGD1564927andRGD1560115) alternative variant bSep08, mRNA.



<a href="#">RGD1564927andRGD1560115</a>	<a href="#">RGD1564927andRGD1560115.bSep08</a>	<a href="#">499930</a>	13052	756	2	179	similar to TGFB-induced factor 2 and similar to TGFB-induced factor 2 (RGD1564927andRGD1560115) alternative variant bSep08, mRNA.
<a href="#">RGD1564936</a>	<a href="#">RGD1564936.aSep08</a>	<a href="#">361875</a>	20704	599	1	188	similar to Vacuolar ATP synthase subunit S1 precursor (V-ATPase S1 subunit) (RGD1564936) alternative variant aSep08, mRNA.
<a href="#">RGD1564936</a>	<a href="#">RGD1564936.bSep08</a>	<a href="#">361875</a>	24173	645	4	163	similar to Vacuolar ATP synthase subunit S1 precursor (V-ATPase S1 subunit) (RGD1564936) alternative variant bSep08, mRNA.
<a href="#">RGD1564941</a>	<a href="#">RGD1564941.aSep08</a>	<a href="#">364538</a>	90362	809		269	ankyrin (RGD1564941) mRNA.
<a href="#">RGD1564942</a>	<a href="#">RGD1564942.aSep08</a>	<a href="#">298409</a>	28188	1005	1	247	RGD1564942 (RGD1564942) alternative variant aSep08, mRNA.
<a href="#">RGD1564942</a>	<a href="#">RGD1564942.bSep08</a>	<a href="#">298409</a>	1633	482	1	17	RGD1564942 (RGD1564942) alternative variant bSep08, mRNA.
<a href="#">RGD1564952</a>	<a href="#">RGD1564952.aSep08</a>	<a href="#">304761</a>	58670	461		153	putative protein of ancient origin (RGD1564952) mRNA.
<a href="#">RGD1564955</a>	<a href="#">RGD1564955.aSep08</a>	<a href="#">501642</a>	31584	1926		566	similar to fibrous sheath interacting protein 2 (RGD1564955) mRNA.
<a href="#">RGD1564956</a>	<a href="#">RGD1564956.bSep08</a>	<a href="#">500462</a>	11593	729	3	138	similar to calcium binding protein P22 (RGD1564956) alternative variant bSep08, complete mRNA.
<a href="#">RGD1564964</a>	<a href="#">RGD1564964.bSep08</a>	<a href="#">315843</a>	11847	897	6	298	similar to WD repeat domain 11 protein (RGD1564964) alternative variant bSep08, mRNA.
<a href="#">RGD1564964</a>	<a href="#">RGD1564964.dSep08</a>	<a href="#">315843</a>	921	319	2	106	similar to WD repeat domain 11 protein (RGD1564964) alternative variant dSep08, mRNA.
<a href="#">RGD1564982</a>	<a href="#">RGD1564982.aSep08</a>	<a href="#">308556</a>	2132	767	1	228	RGD1564982 (RGD1564982) alternative variant aSep08, mRNA.
<a href="#">RGD1564982</a>	<a href="#">RGD1564982.bSep08</a>	<a href="#">308556</a>	2111	453	1	151	RGD1564982 (RGD1564982) alternative variant bSep08, mRNA.
<a href="#">RGD1564998</a>	<a href="#">RGD1564998.aSep08</a>	<a href="#">498323</a>	3702	420		121	hypothetical gene supported by BC082068 (RGD1564998) mRNA.
<a href="#">RGD1564999</a>	<a href="#">RGD1564999.aSep08</a>	<a href="#">291266</a>	4472	749		156	similar to isopentenyl-diphosphate delta isomerase 2 (RGD1564999) mRNA.
<a href="#">RGD1565002</a>	<a href="#">RGD1565002.aSep08</a>	<a href="#">299135</a>	15722	1481	7	418	similar to Dehydrogenase/reductase SDR family member 7 precursor (Retinal short-chain dehydrogenase/reductase 4) (RGD1565002) alternative variant aSep08, mRNA.
<a href="#">RGD1565007</a>	<a href="#">RGD1565007.aSep08</a>	<a href="#">365834</a>	100349	556		66	similar to RIKEN cDNA 4632419K20 (RGD1565007) mRNA.
<a href="#">RGD1565014</a>	<a href="#">RGD1565014.aSep08</a>	<a href="#">290790</a>	2141	407		114	similar to Low-density lipoprotein receptor-related protein 4 precursor (LDLR dan) (RGD1565014) mRNA.
<a href="#">RGD1565032</a>	<a href="#">RGD1565032.aSep08</a>	<a href="#">367701</a>	19279	235		78	similar to transcription elongation factor B (SIII), polypeptide 2 (RGD1565032) mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.aSep08</a>	<a href="#">498014</a>	3911	2018	3	265	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant aSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.aSep08</a>	<a href="#">688846</a>	3911	2018	3	265	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant aSep08, mRNA.

<a href="#">RGD1565033</a>	<a href="#">RGD1565033.bSep08</a>	<a href="#">498014</a>	2416	621	2	192	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant bSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.bSep08</a>	<a href="#">688846</a>	2416	621	2	192	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant bSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.dSep08</a>	<a href="#">498014</a>	1621	722	3	169	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant dSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.dSep08</a>	<a href="#">688846</a>	1621	722	3	169	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant dSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.eSep08</a>	<a href="#">498014</a>	1877	818	2	123	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant eSep08, mRNA.
<a href="#">RGD1565033</a>	<a href="#">RGD1565033.eSep08</a>	<a href="#">688846</a>	1877	818	2	123	similar to hypothetical protein LOC284018 isoform b and hypothetical protein LOC688846 (RGD1565033) alternative variant eSep08, mRNA.
<a href="#">RGD1565037</a>	<a href="#">RGD1565037.bSep08</a>	<a href="#">498398</a>	1714	381	1	65	similar to selenoprotein SeIM (RGD1565037) alternative variant bSep08, mRNA.
<a href="#">RGD1565043</a>	<a href="#">RGD1565043.aSep08</a>	<a href="#">306618</a>	3873	386		128	similar to Rho guanine nucleotide exchange factor (GEF) 10 (RGD1565043) mRNA.
<a href="#">RGD1565052</a>	<a href="#">RGD1565052.bSep08</a>	<a href="#">499589</a>	5165	688	4	136	similar to hypothetical protein MGC27085 (RGD1565052) alternative variant bSep08, mRNA.
<a href="#">RGD1565057</a>	<a href="#">RGD1565057.aSep08</a>	<a href="#">309300</a>	1939	601		74	similar to hypothetical protein FLJ32871 (RGD1565057) mRNA.
<a href="#">RGD1565058</a>	<a href="#">RGD1565058.aSep08</a>	<a href="#">501587</a>	752	572		145	similar to RIKEN cDNA 1700019M22 (RGD1565058) mRNA.
<a href="#">RGD1565082</a>	<a href="#">RGD1565082.aSep08</a>	<a href="#">314381</a>	21550	630		210	similar to Protein C14orf102 homolog (RGD1565082) mRNA.
<a href="#">RGD1565095</a>	<a href="#">RGD1565095.aSep08</a>	<a href="#">503252</a>	13705	3869		74	similar to hypothetical protein MGC52110 (8.4 kD) (RGD1565095) mRNA.
<a href="#">RGD1565096</a>	<a href="#">RGD1565096.aSep08</a>	<a href="#">499255</a>	3412	1250		416	similar to TSG118.1 (RGD1565096) mRNA.
<a href="#">RGD1565132</a>	<a href="#">RGD1565132.aSep08</a>	<a href="#">503462</a>	82659	902	2	79	similar to RIKEN cDNA 4732479N06 (RGD1565132) alternative variant aSep08, mRNA.
<a href="#">RGD1565143</a>	<a href="#">RGD1565143.aSep08</a>	<a href="#">296256</a>	4984	3805	1	620	similar to serine/threonine kinase (RGD1565143) alternative variant aSep08, mRNA.
<a href="#">RGD1565143</a>	<a href="#">RGD1565143.bSep08</a>	<a href="#">296256</a>	20319	5131	8	586	similar to serine/threonine kinase (65.8 kD) (RGD1565143) alternative variant bSep08, mRNA.
<a href="#">RGD1565146</a>	<a href="#">RGD1565146.aSep08</a>	<a href="#">498150</a>	3513	2781		300	similar to hypothetical gene supported by AF226663 (33.2 kD) (RGD1565146) mRNA.
<a href="#">RGD1565149</a>	<a href="#">RGD1565149.aSep08</a>	<a href="#">307923</a>	13623	2710	15	647	vacuolar sorting protein 9 (71.0 kD) (RGD1565149) alternative variant aSep08, complete mRNA.
<a href="#">RGD1565149</a>	<a href="#">RGD1565149.cSep08</a>	<a href="#">307923</a>	3045	1417	6	123	putative protein, with a coiled coil domain, of vertebrate origin (RGD1565149) alternative variant cSep08, mRNA.
<a href="#">RGD1565149</a>	<a href="#">RGD1565149.dSep08</a>	<a href="#">307923</a>	4275	411	6	91	CRA b like (9.4 kD) (RGD1565149) alternative variant dSep08, mRNA.

<a href="#">RGD1565149</a>	<a href="#">RGD1565149.eSep08</a>	<a href="#">307923</a>	1392	401	4	67	putative protein, with a coiled coil domain, of vertebrate origin (7.5 kD) (RGD1565149) alternative variant eSep08, mRNA.
<a href="#">RGD1565164</a>	<a href="#">RGD1565164.aSep08</a>	<a href="#">361582</a>	2366	378		125	similar to associated molecule with the SH3 domain of STAM (RGD1565164) mRNA.
<a href="#">RGD1565192</a>	<a href="#">RGD1565192.bSep08</a>	<a href="#">501923</a>	9830	269		75	similar to 1810013D10Rik protein (RGD1565192) alternative variant bSep08, mRNA.
<a href="#">RGD1565196_predicted</a>	<a href="#">RGD1565196_predicted_bSep08</a>	<a href="#">500629</a>	1162	763	2	137	similar to alcohol dehydrogenase PAN2 (predicted) (14.0 kD) (RGD1565196_predicted) alternative variant bSep08, mRNA.
<a href="#">RGD1565203</a>	<a href="#">RGD1565203.aSep08</a>	<a href="#">363523</a>	1999	706	2	187	similar to ESO3 protein (RGD1565203) alternative variant aSep08, mRNA.
<a href="#">RGD1565210</a>	<a href="#">RGD1565210.aSep08</a>	<a href="#">499072</a>	4721	462		113	RGD1565210 (RGD1565210) mRNA.
<a href="#">RGD1565215</a>	<a href="#">RGD1565215.bSep08</a>	<a href="#">498998</a>	414	341	2	75	similar to 60S ribosomal protein L26 (RGD1565215) alternative variant bSep08, mRNA.
<a href="#">RGD1565218</a>	<a href="#">RGD1565218.aSep08</a>	<a href="#">498697</a>	14915	1546		198	similar to Kelch-like protein 3 (RGD1565218) mRNA.
<a href="#">RGD1565232</a>	<a href="#">RGD1565232.aSep08</a>	<a href="#">500443</a>	1975	1129		141	RGD1565232 (RGD1565232) mRNA.
<a href="#">RGD1565261</a>	<a href="#">RGD1565261.aSep08</a>	<a href="#">361672</a>	5583	1079		359	similar to kinase non-catalytic C-lobe domain (KIND) containing 1 isoform b (RGD1565261) mRNA.
<a href="#">RGD1565283</a>	<a href="#">RGD1565283.bSep08</a>	<a href="#">498274</a>	19143	705	1	169	similar to novel protein (19.2 kD) (RGD1565283) alternative variant bSep08, mRNA.
<a href="#">RGD1565289</a>	<a href="#">RGD1565289.aSep08</a>	<a href="#">363474</a>	104826	894		204	similar to RIKEN cDNA 0610008C08 (RGD1565289) mRNA.
<a href="#">RGD1565309</a>	<a href="#">RGD1565309.aSep08</a>	<a href="#">498292</a>	25273	367		114	similar to hypothetical protein MGC33370 (RGD1565309) mRNA.
<a href="#">RGD1565310</a>	<a href="#">RGD1565310.aSep08</a>	<a href="#">362697</a>	43348	2224		523	similar to RIKEN cDNA 1110018J12 (RGD1565310) mRNA.
<a href="#">RGD1565316</a>	<a href="#">RGD1565316.aSep08</a>	<a href="#">315091</a>	2539	1992	6	545	similar to sphingomyelin phosphodiesterase 3, neutral membrane (RGD1565316) alternative variant aSep08, mRNA.
<a href="#">RGD1565316</a>	<a href="#">RGD1565316.bSep08</a>	<a href="#">315091</a>	1610	1117	3	371	similar to sphingomyelin phosphodiesterase 3, neutral membrane (RGD1565316) alternative variant bSep08, mRNA.
<a href="#">RGD1565316</a>	<a href="#">RGD1565316.cSep08</a>	<a href="#">315091</a>	1388	609	5	76	similar to sphingomyelin phosphodiesterase 3, neutral membrane (RGD1565316) alternative variant cSep08, mRNA.
<a href="#">RGD1565344</a>	<a href="#">RGD1565344.aSep08</a>	<a href="#">308654</a>	39840	616		98	similar to zinc finger, matrin type 1 (RGD1565344) mRNA.
<a href="#">RGD1565350</a>	<a href="#">RGD1565350.aSep08</a>	<a href="#">362513</a>	106442	1569	2	300	similar to Shb protein (RGD1565350) alternative variant aSep08, mRNA.
<a href="#">RGD1565350</a>	<a href="#">RGD1565350.bSep08</a>	<a href="#">362513</a>	45389	694	1	230	similar to Shb protein (RGD1565350) alternative variant bSep08, mRNA.
<a href="#">RGD1565355_predicted</a>	<a href="#">RGD1565355_predicted_bSep08</a>	<a href="#">499985</a>	5574	568	3	128	similar to fatty acid translocase/CD36 (predicted) (RGD1565355_predicted) alternative variant bSep08, mRNA.
<a href="#">RGD1565355_predicted</a>	<a href="#">RGD1565355_predicted_cSep08</a>	<a href="#">499985</a>	1680	749	2	70	similar to fatty acid translocase/CD36 (predicted) (RGD1565355_predicted) alternative variant cSep08, mRNA.

<a href="#">RGD1565361</a>	<a href="#">RGD1565361.aSep08</a>	<a href="#">500010</a>	3335	1880	3	601	similar to male-enhanced antigen-2 (RGD1565361) alternative variant aSep08, mRNA.
<a href="#">RGD1565362_predicted</a>	<a href="#">RGD1565362_predicted.aSep08</a>	<a href="#">313382</a>	88423	3472	13	484	similar to channel-interacting PDZ domain protein isoform 1 (predicted) (RGD1565362_predicted) alternative variant aSep08, mRNA.
<a href="#">RGD1565362_predicted</a>	<a href="#">RGD1565362_predicted.bSep08</a>	<a href="#">313382</a>	13993	601	2	56	similar to channel-interacting PDZ domain protein isoform 1 (predicted) (RGD1565362_predicted) alternative variant bSep08, mRNA.
<a href="#">RGD1565362_predicted</a>	<a href="#">RGD1565362_predicted.cSep08</a>	<a href="#">313382</a>	8726	671	2	66	similar to channel-interacting PDZ domain protein isoform 1 (predicted) (7.3 kD) (RGD1565362_predicted) alternative variant cSep08, mRNA.
<a href="#">RGD1565367</a>	<a href="#">RGD1565367.aSep08</a>	<a href="#">312226</a>	22483	722	4	236	similar to Solute carrier family 23, member 2 (Sodium-dependent vitamin C transporter 2) (RGD1565367) alternative variant aSep08, mRNA.
<a href="#">RGD1565367</a>	<a href="#">RGD1565367.cSep08</a>	<a href="#">312226</a>	27234	390	3	130	similar to Solute carrier family 23, member 2 (Sodium-dependent vitamin C transporter 2) (RGD1565367) alternative variant cSep08, mRNA.
<a href="#">RGD1565373</a>	<a href="#">RGD1565373.aSep08</a>	<a href="#">296985</a>	22845	1561	5	254	C-type lectin (RGD1565373) alternative variant aSep08, mRNA.
<a href="#">RGD1565373</a>	<a href="#">RGD1565373.cSep08</a>	<a href="#">296985</a>	785	665	2	104	putative secreted or extracellular protein precursor (11.0 kD) (RGD1565373) alternative variant cSep08, mRNA.
<a href="#">RGD1565374</a>	<a href="#">RGD1565374.aSep08</a>	<a href="#">498128</a>	2241	703	4	192	similar to hypothetical protein LOC199675 (RGD1565374) alternative variant aSep08, mRNA.
<a href="#">RGD1565374</a>	<a href="#">RGD1565374.cSep08</a>	<a href="#">498128</a>	3593	1782	4	72	similar to hypothetical protein LOC199675 (RGD1565374) alternative variant cSep08, mRNA.
<a href="#">RGD1565378</a>	<a href="#">RGD1565378.aSep08</a>	<a href="#">360565</a>	810	427		95	similar to novel protein (RGD1565378) mRNA.
<a href="#">RGD1565385</a>	<a href="#">RGD1565385.aSep08</a>	<a href="#">362153</a>	23821	1897	1	539	similar to mKIAA1604 protein (62.6 kD) (RGD1565385) alternative variant aSep08, mRNA.
<a href="#">RGD1565385</a>	<a href="#">RGD1565385.bSep08</a>	<a href="#">362153</a>	16099	829	1	275	similar to mKIAA1604 protein (RGD1565385) alternative variant bSep08, mRNA.
<a href="#">RGD1565407</a>	<a href="#">RGD1565407.aSep08</a>	<a href="#">290628</a>	1744	1245	1	265	similar to RIKEN cDNA 9430098E02 (29.8 kD) (RGD1565407) alternative variant aSep08, mRNA.
<a href="#">RGD1565407</a>	<a href="#">RGD1565407.cSep08</a>	<a href="#">290628</a>	1310	474		92	similar to RIKEN cDNA 9430098E02 (RGD1565407) alternative variant cSep08, mRNA.
<a href="#">RGD1565411</a>	<a href="#">RGD1565411.bSep08</a>	<a href="#">363000</a>	810	728	2	88	similar to 2610317D23Rik protein (RGD1565411) alternative variant bSep08, mRNA.
<a href="#">RGD1565411</a>	<a href="#">RGD1565411.dSep08</a>	<a href="#">363000</a>	524	344	3	75	similar to 2610317D23Rik protein (RGD1565411) alternative variant dSep08, mRNA.
<a href="#">RGD1565411</a>	<a href="#">RGD1565411.eSep08</a>	<a href="#">363000</a>	928	748	2	31	similar to 2610317D23Rik protein (3.4 kD) (RGD1565411) alternative variant eSep08, mRNA.
<a href="#">RGD1565416</a>	<a href="#">RGD1565416.aSep08</a>	<a href="#">315776</a>	98292	968	1	264	similar to talin 2 (RGD1565416) alternative variant aSep08, mRNA.
<a href="#">RGD1565416</a>	<a href="#">RGD1565416.bSep08</a>	<a href="#">315776</a>	250898	575	2	108	similar to talin 2 (RGD1565416) alternative variant bSep08, mRNA.
<a href="#">RGD1565416</a>	<a href="#">RGD1565416.cSep08</a>	<a href="#">315776</a>	145903	398	1	77	similar to talin 2 (RGD1565416) alternative variant cSep08, mRNA.

RGD1565419	RGD1565419.bSep08	500950	19038	2141	6	596	similar to zinc finger protein 75 (RGD1565419) alternative variant bSep08, mRNA.
RGD1565419	RGD1565419.cSep08	500950	31564	1760	5		
RGD1565419	RGD1565419.dSep08	500950	31609	1484	4		
RGD1565421	RGD1565421.aSep08	501086	4220	1898	4	172	similar to Sulfotransferase K2 (rSULT1C2A) (RGD1565421) alternative variant aSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.bSep08	361953	52664	584	5	194	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant bSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.bSep08	499631	52664	584	5	194	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant bSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.bSep08	691104	52664	584	5	194	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant bSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.cSep08	361953	8116	463	3	86	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant cSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.cSep08	499631	8116	463	3	86	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant cSep08, mRNA.
RGD1565425andLekr1	RGD1565425andLekr1.cSep08	691104	8116	463	3	86	similar to Restin and leucine, glutamate and lysine rich 1 and hypothetical protein LOC691104 (RGD1565425andLekr1) alternative variant cSep08, mRNA.
RGD1565432	RGD1565432.aSep08	499157	5716	1205	3	125	similar to hypothetical protein (RGD1565432) alternative variant aSep08, mRNA.
RGD1565432	RGD1565432.bSep08	499157	2304	610	1	73	similar to hypothetical protein (RGD1565432) alternative variant bSep08, mRNA.
RGD1565457	RGD1565457.aSep08	288589	10506	1757	5	492	similar to Rasa4 protein (RGD1565457) alternative variant aSep08, mRNA.
RGD1565457	RGD1565457.bSep08	288589	10021	1762	2	414	similar to Rasa4 protein (RGD1565457) alternative variant bSep08, mRNA.
RGD1565474	RGD1565474.aSep08	362353	8861	2297		625	similar to mKIAA0738 protein (RGD1565474) mRNA.
RGD1565477	RGD1565477.aSep08	498430	2332	831	2	98	similar to echinoderm microtubule associated protein like 5 (11.3 kD) (RGD1565477) alternative variant aSep08, mRNA.
RGD1565480	RGD1565480.aSep08	500511	10166	773		152	similar to hypothetical protein MGC35130 (RGD1565480) mRNA.
RGD1565482	RGD1565482.aSep08	291535	6460	462		109	RGD1565482 (RGD1565482) mRNA.
RGD1565487	RGD1565487.aSep08	292625	2166	480		132	RGD1565487 (RGD1565487) mRNA.

RGD1565493	RGD1565493.aSep08	500853	43345	919	2	305	similar to DKFZP434I092 protein (RGD1565493) alternative variant aSep08, mRNA.
RGD1565493	RGD1565493.bSep08	500853	32264	772	1	257	similar to DKFZP434I092 protein (RGD1565493) alternative variant bSep08, mRNA.
RGD1565496	RGD1565496.bSep08	300783	9701	2203		188	similar to Butyrate-induced transcript 1 (RGD1565496) alternative variant bSep08, mRNA.
RGD1565498	RGD1565498.bSep08	500843	55390	556	3	128	similar to Hypothetical protein LOC270802 (RGD1565498) alternative variant bSep08, mRNA.
RGD1565502	RGD1565502.aSep08	498531	3851	1308	3	102	RGD1565502 (RGD1565502) alternative variant aSep08, mRNA.
RGD1565512	RGD1565512.bSep08	499759	6813	681	1	226	similar to hypothetical protein 4932418E24 (RGD1565512) alternative variant bSep08, mRNA.
RGD1565514	RGD1565514.aSep08	317614	11544	617		205	similar to RIKEN cDNA 4933424A10 gene (RGD1565514) mRNA.
RGD1565533	RGD1565533.bSep08	498004	1417	264	1	41	RGD1565533 (4.8 kD) (RGD1565533) alternative variant bSep08, mRNA.
RGD1565536	RGD1565536.aSep08	311323	816	734		99	similar to hypothetical protein (RGD1565536) mRNA.
RGD1565551	RGD1565551.aSep08	499807	24744	1774	11	365	similar to RIKEN cDNA 4833418A01 (40.6 kD) (RGD1565551) alternative variant aSep08, mRNA.
RGD1565551	RGD1565551.bSep08	499807	2014	421	1	90	similar to RIKEN cDNA 4833418A01 (RGD1565551) alternative variant bSep08, mRNA.
RGD1565556	RGD1565556.aSep08	314721	143826	3251	11	448	similar to cajalin 2 isoform a (RGD1565556) alternative variant aSep08, mRNA.
RGD1565556	RGD1565556.bSep08	314721	73991	1139	9	334	similar to cajalin 2 isoform a (RGD1565556) alternative variant bSep08, mRNA.
RGD1565556	RGD1565556.cSep08	314721	18062	551	3	97	similar to cajalin 2 isoform a (RGD1565556) alternative variant cSep08, mRNA.
RGD1565556	RGD1565556.dSep08	314721	970	352	2	60	similar to cajalin 2 isoform a (RGD1565556) alternative variant dSep08, mRNA.
RGD1565556	RGD1565556.fSep08	314721	1526	403	2	35	similar to cajalin 2 isoform a (3.7 kD) (RGD1565556) alternative variant fSep08, mRNA.
RGD1565557	RGD1565557.aSep08	500247	20041	1442	4	200	similar to RIKEN cDNA 2010301N04 (RGD1565557) alternative variant aSep08, mRNA.
RGD1565557	RGD1565557.bSep08	500247	10305	1224	2	70	similar to RIKEN cDNA 2010301N04 (8.2 kD) (RGD1565557) alternative variant bSep08, mRNA.
RGD1565557	RGD1565557.dSep08	500247	5806	711	2	83	similar to RIKEN cDNA 2010301N04 (9.2 kD) (RGD1565557) alternative variant dSep08, mRNA.
RGD1565561	RGD1565561.bSep08	498741	9457	2062	1	108	putative protein, with at least 2 transmembrane domains, of bilateral origin (RGD1565561) alternative variant bSep08, mRNA.
RGD1565584	RGD1565584.aSep08	293112	3057	1816	3	129	similar to tyrosine kinase-associated leucine zipper protein LAZipII (15.8 kD) (RGD1565584) alternative variant aSep08, complete mRNA.
RGD1565584	RGD1565584.eSep08	293112	1950	920	2	112	similar to tyrosine kinase-associated leucine zipper protein LAZipII (12.9 kD) (RGD1565584) alternative variant eSep08, mRNA.

RGD1565591	RGD1565591.aSep08	313757	5570	1981		480	similar to Ski protein (RGD1565591) alternative variant aSep08, mRNA.
RGD1565598	RGD1565598.aSep08	502118	1312	544		121	similar to serine/threonine kinase (RGD1565598) mRNA.
RGD1565616	RGD1565616.aSep08	499891	20938	1011	11	318	RGD1565616 (RGD1565616) alternative variant aSep08, mRNA.
RGD1565616	RGD1565616.bSep08	499891	4000	851	5	170	RGD1565616 (RGD1565616) alternative variant bSep08, mRNA.
RGD1565629	RGD1565629.aSep08	307358	6142	449		78	similar to collagen and calcium binding EGF domains 1 (RGD1565629) mRNA.
RGD1565635	RGD1565635.aSep08	291802	4454	625		176	similar to zinc finger protein 124 (RGD1565635) mRNA.
RGD1565641	RGD1565641.bSep08	499567	14001	314	3	45	RGD1565641 (RGD1565641) alternative variant bSep08, mRNA.
RGD1565652	RGD1565652.aSep08	498583	16052	644		214	putative protein (RGD1565652) mRNA.
RGD1565655	RGD1565655.aSep08	499097	4449	1496		201	similar to BC049730 protein (21.2 kD) (RGD1565655) complete mRNA.
RGD1565672	RGD1565672.bSep08	500870	65947	2107		536	similar to MDM2 Binding protein (RGD1565672) alternative variant bSep08, mRNA.
RGD1565675	RGD1565675.aSep08	290925	5343	784		173	similar to RIKEN cDNA 2410022L05 (18.5 kD) (RGD1565675) complete mRNA.
RGD1565676	RGD1565676.aSep08	501128	9698	413		83	uncharacterized protein like (RGD1565676) mRNA.
RGD1565682	RGD1565682.aSep08	294097	7826	713		226	lipase Gastric (RGD1565682) mRNA.
RGD1565685	RGD1565685.bSep08	317344	8446	827	4	119	similar to RIKEN cDNA 1810030O07 (RGD1565685) alternative variant bSep08, mRNA.
RGD1565687	RGD1565687.aSep08	301596	3578	720		191	similar to hypothetical protein FLJ40243 (RGD1565687) mRNA.
RGD1565688	RGD1565688.aSep08	498517	8567	1558		470	similar to mKIAA0323 protein (RGD1565688) mRNA.
RGD1565690	RGD1565690.aSep08	502784	84483	465		86	similar to mKIAA2027 protein (9.4 kD) (RGD1565690) mRNA.
RGD1565705	RGD1565705.aSep08	299488	47256	1211	12	403	similar to chr2 synaptotagmin (RGD1565705) alternative variant aSep08, mRNA.
RGD1565705	RGD1565705.bSep08	299488	16517	739	3	245	similar to chr2 synaptotagmin (RGD1565705) alternative variant bSep08, mRNA.
RGD1565705	RGD1565705.cSep08	299488	5808	1206	1	85	similar to chr2 synaptotagmin (RGD1565705) alternative variant cSep08, mRNA.
RGD1565709	RGD1565709.aSep08	362442	4701	700		170	similar to ovostatin-2 (RGD1565709) mRNA.
RGD1565726	RGD1565726.aSep08	363167	44415	1056		276	similar to hypothetical protein A730098P15 (RGD1565726) mRNA.
RGD1565744	RGD1565744.bSep08	360494	2132	1402	3	228	similar to RIKEN cDNA 0610007P22 (25.4 kD) (RGD1565744) alternative variant bSep08, mRNA.
RGD1565744	RGD1565744.cSep08	360494	1541	658	3	208	similar to RIKEN cDNA 0610007P22 (RGD1565744) alternative variant cSep08, mRNA.
RGD1565744	RGD1565744.dSep08	360494	2335	897	4	188	similar to RIKEN cDNA 0610007P22 (20.8 kD) (RGD1565744) alternative variant dSep08, mRNA.
RGD1565775	RGD1565775.aSep08	361980	35216	2608	13	544	similar to RIKEN cDNA 2810403A07 (58.4 kD) (RGD1565775) alternative variant aSep08, mRNA.

RGD1565775	RGD1565775.bSep08	361980	21789	735	6	244	similar to RIKEN cDNA 2810403A07 (RGD1565775) alternative variant bSep08, mRNA.
RGD1565775	RGD1565775.cSep08	361980	22697	3543	3	115	similar to RIKEN cDNA 2810403A07 (RGD1565775) alternative variant cSep08, mRNA.
RGD1565775	RGD1565775.eSep08	361980	21500	655	5	71	similar to RIKEN cDNA 2810403A07 (RGD1565775) alternative variant eSep08, mRNA.
RGD1565779	RGD1565779.aSep08	502155	1799	696		166	similar to hypothetical protein E230025N22 (19.1 kD) (RGD1565779) mRNA.
RGD1565796	RGD1565796.bSep08	498394	3049	340	2	77	similar to RIKEN cDNA A930005I04 gene (RGD1565796) alternative variant bSep08, mRNA.
RGD1565800	RGD1565800.aSep08	304529	9240	2658		148	similar to hypothetical protein FLJ20674 (RGD1565800) mRNA.
RGD1565804	RGD1565804.aSep08	501047	2255	543		181	similar to procollagen, type VI, alpha 3 (RGD1565804) mRNA.
RGD1565820	RGD1565820.aSep08	364029	12931	773	2	204	similar to novel protein (RGD1565820) alternative variant aSep08, mRNA.
RGD1565820	RGD1565820.bSep08	364029	14933	689	2	182	similar to novel protein (RGD1565820) alternative variant bSep08, mRNA.
RGD1565840	RGD1565840.aSep08	295975	17610	1264		374	similar to Dendritic cell protein GA17 (42.5 kD) (RGD1565840) complete mRNA.
RGD1565847	RGD1565847.aSep08	367033	17125	3735	10	956	similar to zinc finger protein 560 (108.7 kD) (RGD1565847) alternative variant aSep08, mRNA.
RGD1565858	RGD1565858.aSep08	303965	46689	395		131	similar to KIAA1000 protein (RGD1565858) mRNA.
RGD1565866	RGD1565866.aSep08	314740	22796	392		130	similar to FLJ44112 protein (RGD1565866) mRNA.
RGD1565886	RGD1565886.aSep08	290286	7178	484		32	RGD1565886 (3.4 kD) (RGD1565886) mRNA.
RGD1565895	RGD1565895.aSep08	308850	2878	743		206	similar to DRE1 protein (RGD1565895) mRNA.
RGD1565923	RGD1565923.aSep08	500539	26166	558	4	131	similar to hypothetical protein FLJ20972 (RGD1565923) alternative variant aSep08, mRNA.
RGD1565926	RGD1565926.aSep08	361629	3210	509	2	106	RGD1565926 (RGD1565926) alternative variant aSep08, mRNA.
RGD1565926	RGD1565926.bSep08	361629	4497	1210	2	85	RGD1565926 (RGD1565926) alternative variant bSep08, mRNA.
RGD1565926	RGD1565926.cSep08	361629	3189	439	2	82	RGD1565926 (RGD1565926) alternative variant cSep08, mRNA.
RGD1565926	RGD1565926.dSep08	361629	3584	235	1	70	RGD1565926 (RGD1565926) alternative variant dSep08, mRNA.
RGD1565947	RGD1565947.bSep08	299737	6137	388	3	129	similar to netrin 4 (RGD1565947) alternative variant bSep08, mRNA.
RGD1565953	RGD1565953.aSep08	306166	15373	561		186	similar to multidrug resistance-associated protein 4 (RGD1565953) mRNA.
RGD1565967	RGD1565967.aSep08	302819	2322	589		178	similar to hypothetical protein FLJ30058 (RGD1565967) mRNA.
RGD1565970	RGD1565970.aSep08	498519	1733	777		229	similar to mast cell protease 8 (RGD1565970) mRNA.
RGD1565975	RGD1565975.aSep08	291903	9766	557		137	RGD1565975 (RGD1565975) mRNA.
RGD1565983	RGD1565983.bSep08	317628	19778	694	1	184	similar to apurinic/apyrimidinic endonuclease 2 (RGD1565983) alternative variant bSep08, mRNA.



RGD1565997	RGD1565997.aSep08	500294	4187	580	1	63	putative protein (RGD1565997) alternative variant aSep08, mRNA.
RGD1566006	RGD1566006.aSep08	501828	3103	451		44	similar to paired immunoglobulin-like type 2 receptor beta (RGD1566006) mRNA.
RGD1566010	RGD1566010.aSep08	499195	733	610		157	RGD1566010 (RGD1566010) mRNA.
RGD1566016	RGD1566016.aSep08	360840	212515	3542	22	858	similar to KIAA0456 protein (98.4 kD) (RGD1566016) alternative variant aSep08, mRNA.
RGD1566016	RGD1566016.bSep08	360840	48941	275	4	91	similar to KIAA0456 protein (RGD1566016) alternative variant bSep08, mRNA.
RGD1566017	RGD1566017.aSep08	365037	10619	792		263	sterile alpha motif homology 2 and sterile alpha motif SAM (RGD1566017) mRNA.
RGD1566029	RGD1566029.aSep08	500913	89419	756		188	similar to mKIAA1644 protein (RGD1566029) mRNA.
RGD1566036	RGD1566036.aSep08	498119	88077	703		234	similar to RIKEN cDNA 2310008H04 (RGD1566036) mRNA.
RGD1566052	RGD1566052.aSep08	499851	108991	591	3	117	similar to elongation protein 4 homolog (RGD1566052) alternative variant aSep08, mRNA.
RGD1566083	RGD1566083.aSep08	298602	4420	862		160	similar to hypothetical protein MGC24047 (16.6 kD) (RGD1566083) mRNA.
RGD1566084	RGD1566084.aSep08	501795	2518	1304		372	similar to Hypothetical protein LOC73072 (RGD1566084) mRNA.
RGD1566085	RGD1566085.aSep08	361819	4366	612	3	203	similar to pyridoxal (pyridoxine, vitamin B6) kinase (RGD1566085) alternative variant aSep08, mRNA.
RGD1566085	RGD1566085.bSep08	361819	3603	644	1	71	similar to pyridoxal (pyridoxine, vitamin B6) kinase (RGD1566085) alternative variant bSep08, mRNA.
RGD1566086	RGD1566086.aSep08	360795	3835	471		130	similar to gtf2ird2 (RGD1566086) mRNA.
RGD1566102	RGD1566102.aSep08	499613	2630	913	4	118	RGD1566102 (RGD1566102) alternative variant aSep08, mRNA.
RGD1566114	RGD1566114.aSep08	500671	12645	1779	3	593	c virus f protein-binding protein 2 like (RGD1566114) alternative variant aSep08, mRNA.
RGD1566118	RGD1566118.aSep08	361797	2713	761	6	122	RGD1566118 (RGD1566118) alternative variant aSep08, mRNA.
RGD1566118	RGD1566118.bSep08	361797	1320	739	4	84	RGD1566118 (RGD1566118) alternative variant bSep08, mRNA.
RGD1566118	RGD1566118.cSep08	361797	999	752	2	102	RGD1566118 (RGD1566118) alternative variant cSep08, mRNA.
RGD1566120	RGD1566120.aSep08	497904	16484	684		228	similar to novel protein (RGD1566120) mRNA.
RGD1566124	RGD1566124.aSep08	501861	1714	364		57	similar to hypothetical protein FLJ32940 isoform 1 (RGD1566124) mRNA.
RGD1566127	RGD1566127.bSep08	499270	5325	2211	9	324	similar to BC039632 protein (34.3 kD) (RGD1566127) alternative variant bSep08, mRNA.
RGD1566127	RGD1566127.cSep08	499270	816	409	1	90	similar to BC039632 protein (RGD1566127) alternative variant cSep08, mRNA.
RGD1566132	RGD1566132.aSep08	502360	799	446		87	similar to Tripartite motif protein 30-like (10.1 kD) (RGD1566132) mRNA.
RGD1566133	RGD1566133.bSep08	361219	17794	1717	10	406	similar to Fbxw17 protein (45.0 kD) (RGD1566133) alternative variant bSep08, complete mRNA.

RGD1566133	RGD1566133.cSep08	361219	2107	862	4	183	similar to Fbxw17 protein (RGD1566133) alternative variant cSep08, mRNA.
RGD1566133	RGD1566133.dSep08	361219	905	338	3	89	similar to Fbxw17 protein (RGD1566133) alternative variant dSep08, mRNA.
RGD1566138	RGD1566138.aSep08	498167	2629	565		138	similar to Zinc finger, CW type with PWWP domain 1 (RGD1566138) mRNA.
RGD1566141	RGD1566141.aSep08	288417	4056	1624		515	similar to CG016 (RGD1566141) mRNA.
RGD1566144	RGD1566144.aSep08	499562	11219	393	2	130	similar to hypothetical protein A230042K10 (RGD1566144) alternative variant aSep08, mRNA.
RGD1566144	RGD1566144.bSep08	499562	34237	396	2	111	similar to hypothetical protein A230042K10 (RGD1566144) alternative variant bSep08, mRNA.
RGD1566144	RGD1566144.cSep08	499562	15573	506	2	94	similar to hypothetical protein A230042K10 (RGD1566144) alternative variant cSep08, mRNA.
RGD1566155	RGD1566155.aSep08	501658	9096	638	4	200	similar to 2610030H06Rik protein (RGD1566155) alternative variant aSep08, mRNA.
RGD1566155	RGD1566155.bSep08	501658	5402	744	4	120	similar to 2610030H06Rik protein (13.4 kD) (RGD1566155) alternative variant bSep08, mRNA.
RGD1566155	RGD1566155.cSep08	501658	7551	3270	3	101	similar to 2610030H06Rik protein (11.4 kD) (RGD1566155) alternative variant cSep08, mRNA.
RGD1566180	RGD1566180.aSep08	311956	11401	1266		322	RGD1566180 (RGD1566180) mRNA.
RGD1566215	RGD1566215.bSep08	301742	55830	692	8	230	similar to Coatomer gamma-2 subunit (Gamma-2 coat protein) (Gamma-2 COP) (RGD1566215) alternative variant bSep08, mRNA.
RGD1566220	RGD1566220.aSep08	500692	6099	436		109	similar to hypothetical MGC48595 (RGD1566220) mRNA.
RGD1566239	RGD1566239.aSep08	306348	2454	650	5	168	similar to RIKEN cDNA 2810428115 (RGD1566239) alternative variant aSep08, mRNA.
RGD1566239	RGD1566239.bSep08	306348	1462	923	2	145	similar to RIKEN cDNA 2810428115 (RGD1566239) alternative variant bSep08, mRNA.
RGD1566239	RGD1566239.cSep08	306348	2410	683	4	138	similar to RIKEN cDNA 2810428115 (15.7 kD) (RGD1566239) alternative variant cSep08, mRNA.
RGD1566239	RGD1566239.eSep08	306348	2359	1190	3	44	similar to RIKEN cDNA 2810428115 (4.9 kD) (RGD1566239) alternative variant eSep08, mRNA.
RGD1566248	RGD1566248.aSep08	499060	34361	419		139	similar to hypothetical protein FLJ14345 (RGD1566248) mRNA.
RGD1566254	RGD1566254.aSep08	499318	3012	882		120	RGD1566254 (RGD1566254) mRNA.
RGD1566266	RGD1566266.aSep08	304486	4616	1140		312	similar to hypothetical protein FLJ21127 (RGD1566266) alternative variant aSep08, mRNA.
RGD1566296	RGD1566296.aSep08	315728	385039	596		198	similar to RIKEN cDNA B230114P05 (RGD1566296) mRNA.
RGD1566307	RGD1566307.aSep08	308350	1830	652	1	127	similar to PIRB1 (RGD1566307) alternative variant aSep08, mRNA.
RGD1566307	RGD1566307.bSep08	308350	2551	694	4	59	similar to PIRB1 (RGD1566307) alternative variant bSep08, mRNA.
RGD1566311	RGD1566311.aSep08	500963	968	697		94	RGD1566311 (RGD1566311) mRNA.
RGD1566313	RGD1566313.aSep08	297572	15839	1079		359	similar to Murinoglobulin 1 homolog (RGD1566313) mRNA.

RGD1566314	RGD1566314.aSep08	288106	35618	1573	6	524	similar to KIAA1407 (RGD1566314) alternative variant aSep08, mRNA.
RGD1566314	RGD1566314.bSep08	288106	3416	276		84	similar to KIAA1407 (RGD1566314) alternative variant bSep08, mRNA.
RGD1566319	RGD1566319.bSep08	502988	11455	917	1	60	similar to Sestrin 2 (Hi95) (6.6 kD) (RGD1566319) alternative variant bSep08, mRNA.
RGD1566320	RGD1566320.bSep08	296207	1280	493	2	68	RGD1566320 (7.9 kD) (RGD1566320) alternative variant bSep08, complete mRNA.
RGD1566325	RGD1566325.aSep08	498682	9144	579	4	192	similar to regulator of sex-limitation candidate 16 (RGD1566325) alternative variant aSep08, mRNA.
RGD1566325	RGD1566325.bSep08	498682	9029	340	3	35	similar to regulator of sex-limitation candidate 16 (RGD1566325) alternative variant bSep08, mRNA.
RGD1566368	RGD1566368.aSep08	301083	3140	623		207	similar to Solute carrier family 6 (neurotransmitter transporter), member 20 (RGD1566368) mRNA.
RGD1566383	RGD1566383.aSep08	362233	3455	344		99	similar to Cystatin S precursor (LM protein) (RGD1566383) mRNA.
RGD1566386	RGD1566386.aSep08	304336	35002	2922	1	677	similar to Hypothetical protein A430033K04 (79.7 kD) (RGD1566386) alternative variant aSep08, complete mRNA.
RGD1566386	RGD1566386.bSep08	304336	32805	698	1	216	similar to Hypothetical protein A430033K04 (RGD1566386) alternative variant bSep08, mRNA.
RGD1566400	RGD1566400.aSep08	289001	527	403		37	similar to hypothetical protein FLJ23074 (RGD1566400) mRNA.
RGD1566401	RGD1566401.aSep08	500717	23557	1216	5	129	putative protein (13.9 kD) (RGD1566401) alternative variant aSep08, mRNA.
RGD1566401	RGD1566401.bSep08	500717	15215	6449	5	129	putative protein (13.9 kD) (RGD1566401) alternative variant bSep08, mRNA.
RGD1566401	RGD1566401.cSep08	500717	13518	924	5	108	CRA a like (11.7 kD) (RGD1566401) alternative variant cSep08, mRNA.
RGD1566401	RGD1566401.dSep08	500717	13751	1184	5	108	CRA a like (11.7 kD) (RGD1566401) alternative variant dSep08, mRNA.
RGD1566401	RGD1566401.eSep08	500717	24531	1784	5	104	CRA a like (11.3 kD) (RGD1566401) alternative variant eSep08, mRNA.
RGD1566401	RGD1566401.fSep08	500717	26868	699	5	104	CRA a like (11.3 kD) (RGD1566401) alternative variant fSep08, mRNA.
RGD1566401	RGD1566401.gSep08	500717	4356	1516	3	90	CRA a like (9.5 kD) (RGD1566401) alternative variant gSep08, complete mRNA.
RGD1566401	RGD1566401.jSep08	500717	4241	1374	3	81	CRA b like (8.7 kD) (RGD1566401) alternative variant jSep08, mRNA.
RGD1566401	RGD1566401.kSep08	500717	6270	359	4	44	CRA a like (4.8 kD) (RGD1566401) alternative variant kSep08, mRNA.
RGD1566401	RGD1566401.nSep08	500717	3450	504	2	14	putative protein (1.6 kD) (RGD1566401) alternative variant nSep08, mRNA.
RGD1566401	RGD1566401.oSep08	500717	6395	487	3	39	putative protein (4.1 kD) (RGD1566401) alternative variant oSep08, mRNA.
RGD1566401	RGD1566401.pSep08	500717	6251	417	4	39	putative protein (4.1 kD) (RGD1566401) alternative variant pSep08, mRNA.

RGD1566401	RGD1566401.qSep08	500717	2031	354	2	20	putative protein (RGD1566401) alternative variant qSep08, mRNA.
RGD1566401	RGD1566401.rSep08	500717	5897	347	3	39	putative protein (4.1 kD) (RGD1566401) alternative variant rSep08, mRNA.
RGD1566401	RGD1566401.sSep08	500717	6112	343	3	46	CRA a like (RGD1566401) alternative variant sSep08, mRNA.
RGD1566403	RGD1566403.bSep08	502228	4462	1156	5	132	similar to OTTHUMP00000040081 (RGD1566403) alternative variant bSep08, mRNA.
Rgl1	Rgl1.aSep08	289080	8395	3017	3	158	ral guanine nucleotide dissociation stimulator,-like 1 (Rgl1) alternative variant aSep08, mRNA.
Rgl3	Rgl3.bSep08	300444	5104	1104	2	78	ral guanine nucleotide dissociation stimulator-like 3 (Rgl3) alternative variant bSep08, mRNA.
Rgma	Rgma.bSep08	308739	27725	3253	1	433	RGM domain family, member A (47.5 kD) (Rgma) alternative variant bSep08, mRNA.
Rgn	Rgn.bSep08	25106	16201	795	1	264	regucalcin (Rgn) alternative variant bSep08, mRNA.
Rgnef	Rgnef.bSep08	361882	28193	757	4	252	rho-guanine nucleotide exchange factor (Rgnef) alternative variant bSep08, mRNA.
Rgnef	Rgnef.cSep08	361882	27536	465	5	88	rho-guanine nucleotide exchange factor (Rgnef) alternative variant cSep08, mRNA.
Rgp1	Rgp1.aSep08	313493	1368	379	1	126	RGP1 retrograde golgi transport homolog (S. cerevisiae) (Rgp1) alternative variant aSep08, mRNA.
Rgp1	Rgp1.bSep08	313493	1590	419	2	98	RGP1 retrograde golgi transport homolog (S. cerevisiae) (Rgp1) alternative variant bSep08, mRNA.
Rgr	Rgr.bSep08	306307	4982	285	1	88	retinal G protein coupled receptor (Rgr) alternative variant bSep08, mRNA.
Rgs2	Rgs2.bSep08	84583	2517	707	2	142	regulator of G-protein signaling 2 (Rgs2) alternative variant bSep08, mRNA.
Rgs3	Rgs3.bSep08	54293	39933	4124	11	433	regulator of G-protein 3 (48.4 kD) (Rgs3) alternative variant bSep08, complete mRNA.
Rgs3	Rgs3.cSep08	54293	2892	963	5	192	regulator of G-protein 3 (22.4 kD) (Rgs3) alternative variant cSep08, mRNA.
Rgs3	Rgs3.dSep08	54293	16202	1198	2	184	putative nuclear protein (20.0 kD) (Rgs3) alternative variant dSep08, mRNA.
Rgs3	Rgs3.eSep08	54293	15341	524	2	174	regulator of G-protein 3 (Rgs3) alternative variant eSep08, mRNA.
Rgs3	Rgs3.fSep08	54293	2142	420	3	109	regulator of G-protein 3 (Rgs3) alternative variant fSep08, mRNA.
Rgs3	Rgs3.gSep08	54293	6338	313	4	103	regulator of G-protein 3 (Rgs3) alternative variant gSep08, mRNA.
Rgs3	Rgs3.hSep08	54293	1470	418	4	100	regulator of G-protein 3 (Rgs3) alternative variant hSep08, mRNA.
Rgs3	Rgs3.iSep08	54293	1439	563	3	83	regulator of G-protein 3 (Rgs3) alternative variant iSep08, mRNA.
Rgs3	Rgs3.jSep08	54293	6199	375	2	54	putative protein (Rgs3) alternative variant jSep08, mRNA.
Rgs7	Rgs7.bSep08	54296	17375	737	1	55	regulator of G-protein signaling 7 (6.2 kD) (Rgs7) alternative variant bSep08, mRNA.

Rgs7	Rgs7.cSep08	54296	7680	623	1	34	regulator of G-protein signaling 7 (Rgs7) alternative variant cSep08, mRNA.
Rgs8	Rgs8.aSep08	54297	28268	900	6	180	regulator of G-protein signaling 8 (20.9 kD) (Rgs8) alternative variant aSep08, mRNA.
Rgs8	Rgs8.cSep08	54297	4813	451	1	47	regulator of G-protein signaling 8 (Rgs8) alternative variant cSep08, mRNA.
Rgs9	Rgs9.bSep08	29481	16665	1263	6	331	regulator of G-protein signaling 9 (Rgs9) alternative variant bSep08, mRNA.
Rgs9	Rgs9.cSep08	29481	11522	285	3	94	regulator of G-protein signaling 9 (Rgs9) alternative variant cSep08, mRNA.
Rgs9	Rgs9.dSep08	29481	3144	347	2	73	regulator of G-protein signaling 9 (Rgs9) alternative variant dSep08, mRNA.
Rgs9	Rgs9.eSep08	29481	20114	293	3	70	regulator of G-protein signaling 9 (Rgs9) alternative variant eSep08, mRNA.
Rgs9	Rgs9.fSep08	29481	716	388	2	58	regulator of G-protein signaling 9 (Rgs9) alternative variant fSep08, mRNA.
Rgs11	Rgs11.aSep08	54291	6717	1783	12	184	regulator of G-protein signaling 11 (21.0 kD) (Rgs11) alternative variant aSep08, mRNA.
Rgs11	Rgs11.bSep08	54291	3879	833	10	141	regulator of G-protein signaling 11 (16.4 kD) (Rgs11) alternative variant bSep08, mRNA.
Rgs12	Rgs12.bSep08	54292	47022	754	4	234	regulator of G-protein signaling 12 (Rgs12) alternative variant bSep08, mRNA.
Rgs12	Rgs12.cSep08	54292	30805	731	2	146	regulator of G-protein signaling 12 (Rgs12) alternative variant cSep08, mRNA.
Rgs13	Rgs13.aSep08	54289	3622	541	1	179	regulator of G-protein signaling 1 and regulator of G-protein signaling 13 (Rgs13) alternative variant aSep08, mRNA.
Rgs13	Rgs13.aSep08	498246	3622	541	1	179	regulator of G-protein signaling 1 and regulator of G-protein signaling 13 (Rgs13) alternative variant aSep08, mRNA.
Rgs14	Rgs14.bSep08	114705	3670	704	5	234	regulator of G-protein signaling 14 (Rgs14) alternative variant bSep08, mRNA.
Rgs14	Rgs14.cSep08	114705	7773	662	5	220	regulator of G-protein signaling 14 (Rgs14) alternative variant cSep08, mRNA.
Rgs14	Rgs14.dSep08	114705	1501	743	2	107	regulator of G-protein signaling 14 (Rgs14) alternative variant dSep08, mRNA.
Rgs17	Rgs17.bSep08	308118	80338	523		127	regulator of G-protein signaling 17 (Rgs17) alternative variant bSep08, mRNA.
Rgs19	Rgs19.aSep08	59293	4800	1225	3	295	regulator of G-protein signaling 19 (Rgs19) alternative variant aSep08, mRNA.
Rgs19	Rgs19.cSep08	59293	2341	648	2	152	regulator of G-protein signaling 19 (Rgs19) alternative variant cSep08, mRNA.
Rgs19	Rgs19.dSep08	59293	4447	368	2	82	regulator of G-protein signaling 19 (Rgs19) alternative variant dSep08, mRNA.
Rhbdd1	Rhbdd1.aSep08	316557	116361	1799	1	507	putative protein of metazoan origin (Rhbdd1) alternative variant aSep08, mRNA.
Rhbdd1	Rhbdd1.cSep08	316557	24650	460	1	104	putative protein (Rhbdd1) alternative variant cSep08, mRNA.

Rhbdd2	Rhbdd2.aSep08	360793	10663	1786	4	398	peptidase S54, rhomboid (Rhbdd2) alternative variant aSep08, complete mRNA.
Rhbdd2	Rhbdd2.bSep08	360793	9982	1223	4	379	peptidase S54, rhomboid (Rhbdd2) alternative variant bSep08, mRNA.
Rhbdd2	Rhbdd2.cSep08	360793	10692	1095	5	317	peptidase S54, rhomboid (Rhbdd2) alternative variant cSep08, mRNA.
Rhbdd3	Rhbdd3.bSep08	289753	6024	1782	5	30	putative protein (Rhbdd3) alternative variant bSep08, mRNA.
Rhbdd3	Rhbdd3.cSep08	289753	4586	1195	4	103	putative protein (Rhbdd3) alternative variant cSep08, mRNA.
Rhbdd3	Rhbdd3.dSep08	289753	5304	1000	5	72	putative protein (Rhbdd3) alternative variant dSep08, mRNA.
Rhbdd3	Rhbdd3.eSep08	289753	3867	738	4	69	putative protein of mammalian origin (Rhbdd3) alternative variant eSep08, mRNA.
Rhbdf1	Rhbdf1.bSep08	303008	1574	719	5	195	rhomboid family 1 (Drosophila) (Rhbdf1) alternative variant bSep08, mRNA.
Rhbdf2	Rhbdf2.cSep08	303690	1158	748	4	67	rhomboid 5 homolog 2 (Drosophila) (Rhbdf2) alternative variant cSep08, mRNA.
Rhbdl1	Rhbdl1.aSep08	117025	1649	614	4	204	rhomboid, veinlet-like 1 (Drosophila) (Rhbdl1) alternative variant aSep08, mRNA.
Rhbdl1	Rhbdl1.bSep08	117025	1088	456	3	151	rhomboid, veinlet-like 1 (Drosophila) (Rhbdl1) alternative variant bSep08, mRNA.
Rhbdl2	Rhbdl2.bSep08	298512	5908	838	2	71	rhomboid, veinlet-like 2 (Drosophila) (Rhbdl2) alternative variant bSep08, mRNA.
Rhbdl3	Rhbdl3.bSep08	287556	12125	674	4	149	rhomboid, veinlet-like 3 (Drosophila) and hypothetical protein LOC686138 (Rhbdl3) alternative variant bSep08, mRNA.
Rhbdl3	Rhbdl3.bSep08	686138	12125	674	4	149	rhomboid, veinlet-like 3 (Drosophila) and hypothetical protein LOC686138 (Rhbdl3) alternative variant bSep08, mRNA.
Rhbdl3	Rhbdl3.cSep08	287556	332	262	2	87	rhomboid, veinlet-like 3 (Drosophila) and hypothetical protein LOC686138 (Rhbdl3) alternative variant cSep08, mRNA.
Rhbdl3	Rhbdl3.cSep08	686138	332	262	2	87	rhomboid, veinlet-like 3 (Drosophila) and hypothetical protein LOC686138 (Rhbdl3) alternative variant cSep08, mRNA.
Rhbg	Rhbg.aSep08	310625	12153	1785	2	499	rhesus blood group-associated B glycoprotein (Rhbg) alternative variant aSep08, mRNA.
Rhbg	Rhbg.cSep08	310625	3091	720	1	214	rhesus blood group-associated B glycoprotein (Rhbg) alternative variant cSep08, mRNA.
Rhbg	Rhbg.dSep08	310625	1657	1508	2	68	rhesus blood group-associated B glycoprotein (7.4 kD) (Rhbg) alternative variant dSep08, mRNA.
Rhd	Rhd.bSep08	60414	5278	446	4	116	rh blood group, D antigen (Rhd) alternative variant bSep08, mRNA.
Rhebl1	Rhebl1.bSep08	359959	3304	1098	2	87	ras homolog enriched in brain like 1 (9.7 kD) (Rhebl1) alternative variant bSep08, mRNA.

Rhoa	Rhoa.aSep08	117273	34097	1648	2	281	ras homolog gene family, member A (Rhoa) alternative variant aSep08, mRNA.
Rhobtb1	Rhobtb1.aSep08	309722	13265	1245	4	415	putative protein of eukaryotic origin (Rhobtb1) alternative variant aSep08, mRNA.
Rhobtb3	Rhobtb3.bSep08	309922	4392	406	2	78	putative protein of vertebrate origin (Rhobtb3) alternative variant bSep08, mRNA.
Rhoc	Rhoc.aSep08	295342	6324	1246	1	193	ras homolog gene family, member C (22.0 kD) (Rhoc) alternative variant aSep08, mRNA.
Rhod	Rhod.bSep08	293660	10015	719	2	51	ras homolog gene family, member D (Rhod) alternative variant bSep08, mRNA.
Rhog	Rhog.bSep08	308875	10773	775	1	185	ras homolog gene family, member G (Rhog) alternative variant bSep08, mRNA.
Rhog	Rhog.cSep08	308875	10932	734	2	178	ras homolog gene family, member G (Rhog) alternative variant cSep08, mRNA.
RhoGAP.0	RhoGAP.0.aSep08		8215	1415	8	274	dual-specificity Rho- Arf-GTPase activating protein 1 (RhoGAP.0) alternative variant aSep08, mRNA.
RhoGAP.1	RhoGAP.1.aSep08		35966	834		278	rho gtpase-activating protein rich2 (RhoGAP.1) mRNA.
RhoGAP.2	RhoGAP.2.aSep08		9335	792		263	gtpase-activating protein (RhoGAP.2) mRNA.
RhoGAP.3	RhoGAP.3.aSep08		4048	1057		287	binding protein 1 like (RhoGAP.3) mRNA.
RhoGAP.4	RhoGAP.4.aSep08		1378	415		138	regulatory (RhoGAP.4) mRNA.
RhoGEF.0	RhoGEF.0.aSep08		35804	1580		335	active -related (RhoGEF.0) mRNA.
RhoGEF.1	RhoGEF.1.aSep08		40264	1537		297	1 -specific guanine nucleotide-releasing factor (RhoGEF.1) mRNA.
Rhoj	Rhoj.bSep08	299145	11796	394	3	105	ras homolog gene family, member J (Rhoj) alternative variant bSep08, mRNA.
Rhot1	Rhot1.bSep08	303351	21093	1994	9	350	ras homolog gene family, member T1 (Rhot1) alternative variant bSep08, mRNA.
Rhot1	Rhot1.cSep08	303351	13660	886	4	173	ras homolog gene family, member T1 (Rhot1) alternative variant cSep08, mRNA.
Rhot1	Rhot1.dSep08	303351	2456	895	2	65	ras homolog gene family, member T1 (Rhot1) alternative variant dSep08, mRNA.
Rhpn1	Rhpn1.aSep08	300030	1957	784	4	192	rhophilin, Rho GTPase binding protein 1 (Rhpn1) alternative variant aSep08, mRNA.
Rhpn1	Rhpn1.bSep08	300030	1046	532	2	129	rhophilin, Rho GTPase binding protein 1 (Rhpn1) alternative variant bSep08, mRNA.
Rhpn1	Rhpn1.cSep08	300030	1340	1105	2	113	rhophilin, Rho GTPase binding protein 1 (Rhpn1) alternative variant cSep08, mRNA.
Ribc1	Ribc1.bSep08	317431	10865	1229	7	351	RIB43A domain with coiled-coils 1 (Ribc1) alternative variant bSep08, mRNA.
Ribc1	Ribc1.cSep08	317431	9969	805	6	212	RIB43A domain with coiled-coils 1 (Ribc1) alternative variant cSep08, mRNA.
Ribc1	Ribc1.dSep08	317431	8727	756	6	183	RIB43A domain with coiled-coils 1 (Ribc1) alternative variant dSep08, mRNA.
Ribc1	Ribc1.eSep08	317431	8521	632	5	103	RIB43A domain with coiled-coils 1 (Ribc1) alternative variant eSep08, mRNA.

Ribc2	Ribc2.bSep08	300122	8657	678	5	9	RIB43A domain with coiled-coils 2 (1.1 kD) (Ribc2) alternative variant bSep08, mRNA.
Ribosomal_L7Ae.0	Ribosomal_L7Ae.0.aSep08		647	511	2	132	ribosomal protein L7Ae/L30e/S12e/Gadd45 (14.5 kD) (Ribosomal_L7Ae.0) alternative variant aSep08, complete mRNA.
Ribosomal_L32e.0	Ribosomal_L32e.0.aSep08		3433	374	3	103	ribosomal protein L32e (Ribosomal_L32e.0) alternative variant aSep08, mRNA.
Ribosomal_L32e.0	Ribosomal_L32e.0.bSep08		3414	1158	2	47	putative protein of eukaryotic origin (Ribosomal_L32e.0) alternative variant bSep08, mRNA.
Ribosomal_L37e.2	Ribosomal_L37e.2.aSep08		30777	1004	3	107	putative protein (12.5 kD) (Ribosomal_L37e.2) alternative variant aSep08, mRNA.
Ribosomal_L37e.2	Ribosomal_L37e.2.cSep08		23511	614	3	79	ribosomal protein L37e (8.8 kD) (Ribosomal_L37e.2) alternative variant cSep08, mRNA.
Ribosomal_L39.4	Ribosomal_L39.4.aSep08		4116	508		67	putative protein (Ribosomal_L39.4) mRNA.
Ribosomal_S5_C.1	Ribosomal_S5_C.1.bSep08		466	367	2	102	ribosomal protein S5, C-terminal (Ribosomal_S5_C.1) alternative variant bSep08, mRNA.
Ribosomal_S5_C.1	Ribosomal_S5_C.1.cSep08		682	371	2	56	putative protein of eukaryotic origin (6.4 kD) (Ribosomal_S5_C.1) alternative variant cSep08, mRNA.
Ric8	Ric8.bSep08	293614	842	564	2	112	resistance inhibitors of cholinesterase 8 homolog A (Ric8) alternative variant bSep08, mRNA.
Ric8	Ric8.cSep08	293614	869	780	1	108	resistance inhibitors of cholinesterase 8 homolog A (Ric8) alternative variant cSep08, mRNA.
Ric8b	Ric8b.bSep08	314681	20550	470	3	93	resistance inhibitors of cholinesterase 8B (Ric8b) alternative variant bSep08, mRNA.
Ric8b	Ric8b.cSep08	314681	8915	3515	2	74	putative mitochondrial protein (8.9 kD) (Ric8b) alternative variant cSep08, mRNA.
Ric8b	Ric8b.dSep08	314681	1920	1572	2	65	putative protein (7.3 kD) (Ric8b) alternative variant dSep08, mRNA.
Ricin_B_lectin.0	Ricin_B_lectin.0.aSep08		4044	679	5	225	N-acetylgalactosaminyltransferase (Ricin_B_lectin.0) alternative variant aSep08, mRNA.
Rictor	Rictor.aSep08	310131	41629	405		134	rapamycin-insensitive companion of mTOR (Rictor) mRNA.
Rif1	Rif1.aSep08	295602	4233	1667		159	rap1 interacting factor 1 homolog (yeast) (Rif1) mRNA.
RIH_assoc.0	RIH_assoc.0.aSep08		20167	1470	2	427	ryanodine receptor (RIH_assoc.0) alternative variant aSep08, mRNA.
RIH_assoc.0	RIH_assoc.0.bSep08		16550	628	1	186	ryanodine receptor (RIH_assoc.0) alternative variant bSep08, mRNA.
RIH_assoc.0	RIH_assoc.0.cSep08		7834	516	5	171	ryanodine receptor (RIH_assoc.0) alternative variant cSep08, mRNA.
Rilpl1	Rilpl1.aSep08	304469	37313	1870	8	476	rab interacting lysosomal protein-like 1 (Rilpl1) alternative variant aSep08, mRNA.
Rilpl1	Rilpl1.bSep08	304469	8251	842	2	78	rab interacting lysosomal protein-like 1 (Rilpl1) alternative variant bSep08, mRNA.
Rilpl2	Rilpl2.bSep08	288652	4791	689	1	86	rab interacting lysosomal protein-like 2 (Rilpl2) alternative variant bSep08, mRNA.
RimK.0	RimK.0.aSep08		3796	1828		226	family with sequence similarity 80 member A (RimK.0) mRNA.



Rims2	Rims2.cSep08	116839	50069	326	3	108	regulating synaptic membrane exocytosis 2 (Rims2) alternative variant cSep08, mRNA.
Rin2	Rin2.bSep08	311494	21432	1237	4	209	ras and Rab interactor 2 (24.2 kD) (Rin2) alternative variant bSep08, mRNA.
Rin2	Rin2.cSep08	311494	47612	1253	3	53	ras and Rab interactor 2 (6.0 kD) (Rin2) alternative variant cSep08, mRNA.
Rin3	Rin3.bSep08	314397	14434	1921	3	629	ras and Rab interactor 3 (Rin3) alternative variant bSep08, mRNA.
Rin3	Rin3.cSep08	314397	863	627	2	208	ras and Rab interactor 3 (Rin3) alternative variant cSep08, mRNA.
Rin3	Rin3.dSep08	314397	44446	407	3	135	ras and Rab interactor 3 (Rin3) alternative variant dSep08, mRNA.
Ring1	Ring1.bSep08	309626	912	821	2	128	ring finger protein 1 (Ring1) alternative variant bSep08, mRNA.
Ring1	Ring1.cSep08	309626	1752	482	3	111	ring finger protein 1 (Ring1) alternative variant cSep08, mRNA.
Riok1	Riok1.aSep08	291061	22901	2602	17	566	RIO kinase 1 (yeast) (65.2 kD) (Riok1) alternative variant aSep08, mRNA.
Riok1	Riok1.bSep08	291061	9752	756	6	79	RIO kinase 1 (yeast) (Riok1) alternative variant bSep08, mRNA.
Riok2	Riok2.cSep08	308201	1265	362	2	24	RIO kinase 2 (yeast) (Riok2) alternative variant cSep08, mRNA.
Riok3	Riok3.cSep08	361293	7894	351	3	104	RIO kinase 3 (Riok3) alternative variant cSep08, mRNA.
Riok3	Riok3.dSep08	361293	4320	1052	4	98	RIO kinase 3 (10.9 kD) (Riok3) alternative variant dSep08, mRNA.
Riok3	Riok3.gSep08	361293	2179	1084	2	87	putative protein (9.3 kD) (Riok3) alternative variant gSep08, mRNA.
Riok3	Riok3.hSep08	361293	2072	360	3	31	RIO kinase 3 (Riok3) alternative variant hSep08, mRNA.
Ripk1	Ripk1.bSep08	306886	18363	1776	2	493	receptor (TNFRSF)-interacting serine-threonine kinase 1 (Ripk1) alternative variant bSep08, mRNA.
Ripk2	Ripk2.aSep08	362491	17168	1326	1	381	receptor (TNFRSF)-interacting serine-threonine kinase 2 (Ripk2) alternative variant aSep08, mRNA.
Ripk2	Ripk2.bSep08	362491	17088	1782		110	receptor (TNFRSF)-interacting serine-threonine kinase 2 (Ripk2) alternative variant bSep08, mRNA.
Ripk3	Ripk3.bSep08	246240	2055	895	5	210	receptor-interacting serine-threonine kinase 3 (Ripk3) alternative variant bSep08, mRNA.
Ripk3	Ripk3.cSep08	246240	1410	693	4	180	receptor-interacting serine-threonine kinase 3 (Ripk3) alternative variant cSep08, mRNA.
Ripk3	Ripk3.dSep08	246240	1504	749	3	108	receptor-interacting serine-threonine kinase 3 (Ripk3) alternative variant dSep08, mRNA.
Ripply2	Ripply2.aSep08	363111	4005	525		134	rippy2 homolog (zebrafish) (Ripply2) mRNA.
Rit1	Rit1.aSep08	499652	13010	1619	3	219	ras-like without CAAX 1 (25.2 kD) (Rit1) alternative variant aSep08, mRNA.
Rit1	Rit1.bSep08	499652	891	454	1	79	ras-like without CAAX 1 (8.6 kD) (Rit1) alternative variant bSep08, mRNA.
Rit2	Rit2.bSep08	291713	365856	845		198	ras-like without CAAX 2 (22.5 kD) (Rit2) alternative variant bSep08, mRNA.

Rkhd1	Rkhd1.aSep08	299613	6826	2449	2	577	KH, type 1 and zinc finger, RING-type (Rkhd1) alternative variant aSep08, mRNA.
Rkhd2	Rkhd2.bSep08	307271	42307	693	2	230	hypothetical protein LOC680362 (Rkhd2) alternative variant bSep08, mRNA.
Rkhd2	Rkhd2.bSep08	680362	42307	693	2	230	hypothetical protein LOC680362 (Rkhd2) alternative variant bSep08, mRNA.
Rlbp112	Rlbp112.bSep08	361459	95505	621	4	206	retinaldehyde binding protein 1-like 2 (Rlbp112) alternative variant bSep08, mRNA.
Rlf	Rlf.aSep08	313566	34837	614		204	rearranged L-myc fusion sequence (Rlf) mRNA.
Rmi1	Rmi1.aSep08	306734	3508	2559	1	617	RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae) (68.2 kD) (Rmi1) alternative variant aSep08, mRNA.
Rmi1	Rmi1.bSep08	306734	7202	2416	2	617	RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae) (68.2 kD) (Rmi1) alternative variant bSep08, mRNA.
Rmnd1	Rmnd1.bSep08	292268	10724	521	6	158	required for meiotic nuclear division 1 homolog (S. cerevisiae) (Rmnd1) alternative variant bSep08, mRNA.
Rmnd1	Rmnd1.cSep08	292268	8563	1782	5	123	required for meiotic nuclear division 1 homolog (S. cerevisiae) (Rmnd1) alternative variant cSep08, mRNA.
Rmnd5a	Rmnd5a.aSep08	312439	56583	4659	2	401	required for meiotic nuclear division 5 homolog A (S. cerevisiae) (Rmnd5a) alternative variant aSep08, mRNA.
Rmnd5a	Rmnd5a.bSep08	312439	6645	3081	1	86	required for meiotic nuclear division 5 homolog A (S. cerevisiae) (Rmnd5a) alternative variant bSep08, mRNA.
Rmnd5a	Rmnd5a.cSep08	312439	2935	402	2	41	required for meiotic nuclear division 5 homolog A (S. cerevisiae) (Rmnd5a) alternative variant cSep08, mRNA.
Rnase4andAng1	Rnase4andAng1.cSep08	56759	11743	1375	2	147	putative protein (Rnase4andAng1) alternative variant cSep08, mRNA.
Rnase4andAng1	Rnase4andAng1.cSep08	305843	11743	1375	2	147	putative protein (Rnase4andAng1) alternative variant cSep08, mRNA.
Rnase4andAng1	Rnase4andAng1.dSep08	56759	15908	685	3	147	ribonuclease 4 precursor (16.9 kD) (Rnase4andAng1) alternative variant dSep08, mRNA.
Rnase4andAng1	Rnase4andAng1.dSep08	305843	15908	685	3	147	ribonuclease 4 precursor (16.9 kD) (Rnase4andAng1) alternative variant dSep08, mRNA.
RnaseA.0	RnaseA.0.aSep08		809	649		171	eosinophil-associated ribonuclease A family member (RnaseA.0) mRNA.
Rnaseh1	Rnaseh1.bSep08	298933	2565	506	2	54	ribonuclease H1 (Rnaseh1) alternative variant bSep08, mRNA.
Rnaseh1	Rnaseh1.eSep08	298933	1712	284	2	28	ribonuclease H1 (3.2 kD) (Rnaseh1) alternative variant eSep08, mRNA.
Rnaseh2a	Rnaseh2a.aSep08	364974	9936	1903	5	339	ribonuclease H2, large subunit (Rnaseh2a) alternative variant aSep08, mRNA.
Rnaseh2a	Rnaseh2a.cSep08	364974	8072	1389	7	183	ribonuclease H2, large subunit (Rnaseh2a) alternative variant cSep08, mRNA.
Rnaseh2a	Rnaseh2a.dSep08	364974	3662	763	4	180	ribonuclease H2, large subunit (Rnaseh2a) alternative variant dSep08, mRNA.
Rnaseh2a	Rnaseh2a.eSep08	364974	2215	551	4	164	ribonuclease H2, large subunit (Rnaseh2a) alternative variant eSep08, mRNA.

Rnaseh2a	Rnaseh2a.fSep08	364974	3781	671	4	132	ribonuclease H2, large subunit (Rnaseh2a) alternative variant fSep08, mRNA.
Rnaseh2a	Rnaseh2a.hSep08	364974	3984	740	5	115	ribonuclease H2, large subunit (13.2 kD) (Rnaseh2a) alternative variant hSep08, mRNA.
Rnaseh2b	Rnaseh2b.bSep08	361056	32962	1276	8	226	ribonuclease H2, subunit B (Rnaseh2b) alternative variant bSep08, mRNA.
Rnaseh2b	Rnaseh2b.cSep08	361056	24837	960	8	214	ribonuclease H2, subunit B (Rnaseh2b) alternative variant cSep08, mRNA.
Rnaseh2b	Rnaseh2b.dSep08	361056	30565	736	7	172	ribonuclease H2, subunit B (Rnaseh2b) alternative variant dSep08, mRNA.
Rnasek	Rnasek.aSep08	287453	1816	687	1	98	ribonuclease, RNase K (11.0 kD) (Rnasek) alternative variant aSep08, mRNA.
Rnasek	Rnasek.bSep08	287453	1627	658	1	55	ribonuclease, RNase K (Rnasek) alternative variant bSep08, mRNA.
Rnasel	Rnasel.bSep08	359726	1838	332	2	78	putative cytoplasmic protein (8.6 kD) (Rnasel) alternative variant bSep08, mRNA.
Rnasen	Rnasen.bSep08	310159	63304	1940	20	646	ribonuclease III, nuclear (Rnasen) alternative variant bSep08, mRNA.
Rnasen	Rnasen.cSep08	310159	9853	1721	4	284	ribonuclease III, nuclear (32.3 kD) (Rnasen) alternative variant cSep08, mRNA.
Rnasen	Rnasen.dSep08	310159	1625	895	2	35	ribonuclease III, nuclear (Rnasen) alternative variant dSep08, mRNA.
RNase_H1_sml.0	RNase_H1_sml.0.aSep08		1619	1177	2	179	ribonuclease H2 (RNase_H1_sml.0) alternative variant aSep08, mRNA.
RNase_H1_sml.0	RNase_H1_sml.0.bSep08		1034	846	1	178	ribonuclease H2 (RNase_H1_sml.0) alternative variant bSep08, mRNA.
RNase_PH.0	RNase_PH.0.aSep08		5820	545	7	181	polyribonucleotide nucleotidyltransferase 1 (RNase_PH.0) mRNA.
RNA_pol_Rpb1_1.0	RNA_pol_Rpb1_1.0.aSep08		3160	561		187	polymerase III (RNA_pol_Rpb1_1.0) mRNA.
RNA_pol_Rpb1_2.0	RNA_pol_Rpb1_2.0.aSep08		4037	1722	5	574	RNA polymerase II (RNA_pol_Rpb1_2.0) alternative variant aSep08, mRNA.
RNA_pol_Rpb1_2.0	RNA_pol_Rpb1_2.0.bSep08		580	468	1	116	polymerase II (RNA_pol_Rpb1_2.0) alternative variant bSep08, mRNA.
RNA_pol_Rpb1_5.0	RNA_pol_Rpb1_5.0.aSep08		13331	1346		448	polymerase III polypeptide a 155kDa (RNA_pol_Rpb1_5.0) mRNA.
RNA_pol_Rpb1_R.0	RNA_pol_Rpb1_R.0.aSep08		1118	844		281	zinc finger protein 768 (RNA_pol_Rpb1_R.0) mRNA.
Rnf2	Rnf2.bSep08	304850	7275	2752	2	297	ring finger protein 2 (Rnf2) alternative variant bSep08, mRNA.
Rnf4	Rnf4.aSep08	29274	18680	835	6	113	CRA b (Rnf4) alternative variant aSep08, mRNA.
Rnf4	Rnf4.cSep08	29274	18635	641	6	74	ring finger protein 4 (8.2 kD) (Rnf4) alternative variant cSep08, mRNA.
Rnf4	Rnf4.dSep08	29274	21949	2905	7	116	ring finger protein 4 (13.1 kD) (Rnf4) alternative variant dSep08, mRNA.
Rnf4	Rnf4.fSep08	29274	19557	770	7	91	ring finger protein 4 (Rnf4) alternative variant fSep08, mRNA.

Rnf6	Rnf6.cSep08	304271	4181	810	3	123	ring finger protein (C3H2C3 type) 6 (Rnf6) alternative variant cSep08, mRNA.
Rnf7	Rnf7.cSep08	300948	8404	709	3	42	ring finger protein 7 (4.8 kD) (Rnf7) alternative variant cSep08, mRNA.
Rnf8	Rnf8.aSep08	361815	18002	1348	6	413	ring finger protein 8 (Rnf8) alternative variant aSep08, mRNA.
Rnf10	Rnf10.bSep08	288710	9747	1779	9	522	ring finger protein 10 (Rnf10) alternative variant bSep08, mRNA.
Rnf10	Rnf10.cSep08	288710	6868	1111	8	297	ring finger protein 10 (Rnf10) alternative variant cSep08, mRNA.
Rnf10	Rnf10.dSep08	288710	4483	894	6	216	ring finger protein 10 (Rnf10) alternative variant dSep08, mRNA.
Rnf10	Rnf10.eSep08	288710	802	387	2	115	ring finger protein 10 (Rnf10) alternative variant eSep08, mRNA.
Rnf10	Rnf10.fSep08	288710	1714	703	3	74	ring finger protein 10 (Rnf10) alternative variant fSep08, mRNA.
Rnf11	Rnf11.aSep08	689577	20707	632	3	175	ring finger protein 11 (Rnf11) alternative variant aSep08, mRNA.
Rnf11	Rnf11.bSep08	689577	32822	3044	5	154	ring finger protein 11 (Rnf11) alternative variant bSep08, mRNA.
Rnf12	Rnf12.aSep08	317241	16685	3031	6	603	ring finger protein 12 (67.0 kD) (Rnf12) alternative variant aSep08, mRNA.
Rnf17	Rnf17.aSep08	305908	15460	690		230	ring finger protein 17 (Rnf17) mRNA.
Rnf19a	Rnf19a.bSep08	362900	3265	355	3	98	ring finger protein 19A (Rnf19a) alternative variant bSep08, mRNA.
Rnf19b	Rnf19b.aSep08	313806	23620	2451	2	686	ring finger protein 19B (Rnf19b) alternative variant aSep08, mRNA.
Rnf20	Rnf20.bSep08	313216	2484	1469	4	192	ring finger protein 20 (Rnf20) alternative variant bSep08, mRNA.
Rnf24	Rnf24.aSep08	362218	51219	585		194	ring finger protein 24 (Rnf24) mRNA.
Rnf25	Rnf25.bSep08	301515	6382	1015	9	335	ring finger protein 25 (Rnf25) alternative variant bSep08, mRNA.
Rnf25	Rnf25.cSep08	301515	4922	758	6	221	ring finger protein 25 (25.2 kD) (Rnf25) alternative variant cSep08, mRNA.
Rnf25	Rnf25.dSep08	301515	5921	774	8	221	ring finger protein 25 (Rnf25) alternative variant dSep08, mRNA.
Rnf25	Rnf25.eSep08	301515	5126	1572	7	195	ring finger protein 25 (Rnf25) alternative variant eSep08, mRNA.
Rnf25	Rnf25.fSep08	301515	2782	301	3	42	ring finger protein 25 (Rnf25) alternative variant fSep08, mRNA.
Rnf31	Rnf31.bSep08	364386	4764	1159	8	256	ring finger protein 31 (Rnf31) alternative variant bSep08, mRNA.
Rnf31	Rnf31.cSep08	364386	2079	424	5	140	ring finger protein 31 (Rnf31) alternative variant cSep08, mRNA.
Rnf31	Rnf31.dSep08	364386	896	386	3	128	ring finger protein 31 (Rnf31) alternative variant dSep08, mRNA.

Rnf31	Rnf31.eSep08	364386	4746	879	8	121	ring finger protein 31 (13.4 kD) (Rnf31) alternative variant eSep08, mRNA.
Rnf32	Rnf32.bSep08	311936	10099	771	7	220	ring finger protein 32 (Rnf32) alternative variant bSep08, mRNA.
Rnf32	Rnf32.cSep08	311936	10076	745	7	208	ring finger protein 32 (Rnf32) alternative variant cSep08, mRNA.
Rnf32	Rnf32.dSep08	311936	9622	543	5	103	ring finger protein 32 (11.2 kD) (Rnf32) alternative variant dSep08, complete mRNA.
Rnf32	Rnf32.eSep08	311936	1849	360	3	59	ring finger protein 32 (6.7 kD) (Rnf32) alternative variant eSep08, mRNA.
Rnf34	Rnf34.cSep08	282845	953	751	2	45	ring finger protein 34 (4.8 kD) (Rnf34) alternative variant cSep08, mRNA.
Rnf34	Rnf34.dSep08	282845	12105	302	3	52	ring finger protein 34 (6.0 kD) (Rnf34) alternative variant dSep08, mRNA.
Rnf38	Rnf38.bSep08	171501	25945	783	3	217	ring finger protein 38 (Rnf38) alternative variant bSep08, mRNA.
Rnf38	Rnf38.cSep08	171501	18406	710	1	172	ring finger protein 38 (Rnf38) alternative variant cSep08, mRNA.
Rnf39	Rnf39.aSep08	171387	1956	439	2	145	ring finger protein 39 (Rnf39) alternative variant aSep08, mRNA.
Rnf39	Rnf39.bSep08	171387	1228	423	2	140	ring finger protein 39 (Rnf39) alternative variant bSep08, mRNA.
Rnf39	Rnf39.cSep08	171387	655	477	2	57	ring finger protein 39 (6.1 kD) (Rnf39) alternative variant cSep08, mRNA.
Rnf41	Rnf41.bSep08	362814	24368	1241	7	289	ring finger protein 41 (Rnf41) alternative variant bSep08, mRNA.
Rnf41	Rnf41.cSep08	362814	1399	731	2	101	ring finger protein 41 CRA b (11.3 kD) (Rnf41) alternative variant cSep08, mRNA.
Rnf41	Rnf41.dSep08	362814	21889	717	5	100	putative mitochondrial protein (10.7 kD) (Rnf41) alternative variant dSep08, mRNA.
Rnf43	Rnf43.aSep08	303412	1682	401		133	ring finger protein 43 (Rnf43) mRNA.
Rnf44	Rnf44.aSep08	361212	8433	1783	6	357	ring finger protein 44 (Rnf44) alternative variant aSep08, mRNA.
Rnf44	Rnf44.dSep08	361212	1165	405	3	134	ring finger protein 44 (Rnf44) alternative variant dSep08, mRNA.
Rnf44	Rnf44.eSep08	361212	1084	785	3	107	ring finger protein (12.2 kD) (Rnf44) alternative variant eSep08, mRNA.
Rnf44	Rnf44.fSep08	361212	10099	720	4	100	ring finger protein (11.0 kD) (Rnf44) alternative variant fSep08, mRNA.
Rnf44	Rnf44.gSep08	361212	4573	400	4	45	ring finger protein 44 like (4.9 kD) (Rnf44) alternative variant gSep08, mRNA.
Rnf111	Rnf111.bSep08	300813	11550	2209	6	219	ring finger 111 (Rnf111) alternative variant bSep08, mRNA.
Rnf111	Rnf111.cSep08	300813	33888	403	2	112	ring finger 111 (Rnf111) alternative variant cSep08, mRNA.
Rnf111	Rnf111.dSep08	300813	33628	659	2	73	ring finger 111 (Rnf111) alternative variant dSep08, mRNA.
Rnf121	Rnf121.bSep08	308871	51283	764	6	122	ring finger protein 121 (Rnf121) alternative variant bSep08, mRNA.

Rnf126	Rnf126.bSep08	314613	7233	799	4	266	ring finger protein 126 (Rnf126) alternative variant bSep08, mRNA.
Rnf126	Rnf126.cSep08	314613	5911	751	7	249	ring finger protein 126 (Rnf126) alternative variant cSep08, mRNA.
Rnf128	Rnf128.aSep08	315911	58090	2813	6	519	ring finger protein 128 (Rnf128) alternative variant aSep08, mRNA.
Rnf128	Rnf128.bSep08	315911	7204	757	2	36	ring finger protein 128 (4.1 kD) (Rnf128) alternative variant bSep08, mRNA.
Rnf138	Rnf138.aSep08	94196	22344	772	2	214	ring finger protein 138 (Rnf138) alternative variant aSep08, mRNA.
Rnf138	Rnf138.cSep08	94196	16609	2412	1	117	ring finger protein 138 (Rnf138) alternative variant cSep08, mRNA.
Rnf138	Rnf138.dSep08	94196	7962	464	2	114	ring finger protein 138 (Rnf138) alternative variant dSep08, mRNA.
rnf141	rnf141.bSep08	308900	22345	3949	7	230	ring finger protein 141 (25.5 kD) (rnf141) alternative variant bSep08, mRNA.
rnf141	rnf141.cSep08	308900	2346	604	2	72	ring finger protein 141 (8.1 kD) (rnf141) alternative variant cSep08, mRNA.
Rnf144a	Rnf144a.bSep08	500636	91690	579	4	171	ring finger protein 144A (Rnf144a) alternative variant bSep08, mRNA.
Rnf144a	Rnf144a.cSep08	500636	8742	571	2	61	ring finger protein 144A (Rnf144a) alternative variant cSep08, mRNA.
Rnf145	Rnf145.bSep08	287212	29056	1798	5	599	ring finger protein 145 (Rnf145) alternative variant bSep08, mRNA.
Rnf145	Rnf145.cSep08	287212	2064	990	2	228	ring finger protein 145 (Rnf145) alternative variant cSep08, mRNA.
Rnf145	Rnf145.dSep08	287212	6587	672	4	215	ring finger protein 145 (Rnf145) alternative variant dSep08, mRNA.
Rnf145	Rnf145.eSep08	287212	20664	507	4	124	ring finger protein 145 (Rnf145) alternative variant eSep08, mRNA.
Rnf145	Rnf145.fSep08	287212	4379	812	3	119	ring finger protein 145 (Rnf145) alternative variant fSep08, mRNA.
Rnf146	Rnf146.aSep08	308051	16870	1923	3	355	ring finger protein 146 (38.6 kD) (Rnf146) alternative variant aSep08, mRNA.
Rnf146	Rnf146.cSep08	308051	15829	940	4	246	ring finger protein 146 (Rnf146) alternative variant cSep08, mRNA.
Rnf149	Rnf149.aSep08	363222	25336	3726	3	475	ring finger protein 149 (Rnf149) alternative variant aSep08, mRNA.
Rnf149	Rnf149.bSep08	363222	12742	774	2	157	ring finger protein 149 (Rnf149) alternative variant bSep08, mRNA.
Rnf149	Rnf149.cSep08	363222	12838	749	1	81	ring finger protein 149 (Rnf149) alternative variant cSep08, mRNA.
Rnf150	Rnf150.aSep08	364983	49867	3046		66	ring finger protein 150 (Rnf150) alternative variant aSep08, mRNA.
Rnf165	Rnf165.aSep08	307251	114391	3449	8	359	ring finger protein 165 (Rnf165) alternative variant aSep08, mRNA.

Rnf165	Rnf165.cSep08	307251	2302	645	1	42	ring finger protein 165 (Rnf165) alternative variant cSep08, mRNA.
Rnf166	Rnf166.bSep08	365022	7429	787	1	216	ring finger protein 166 (Rnf166) alternative variant bSep08, mRNA.
Rnf167	Rnf167.bSep08	360554	2687	415	6	124	ring finger protein 167 (Rnf167) alternative variant bSep08, mRNA.
Rnf167	Rnf167.cSep08	360554	1306	388	4	119	ring finger protein 167 (Rnf167) alternative variant cSep08, mRNA.
Rnf170	Rnf170.aSep08	364654	2696	1003		126	ring finger protein 170 (Rnf170) mRNA.
Rnf180	Rnf180.aSep08	685384	67726	1410		183	ring finger protein 180 (Rnf180) mRNA.
Rnf181	Rnf181.bSep08	297337	2695	1493	5	129	ring finger protein 181 (Rnf181) alternative variant bSep08, mRNA.
Rnf181	Rnf181.cSep08	297337	1838	569	5	118	ring finger protein 181 (13.8 kD) (Rnf181) alternative variant cSep08, mRNA.
Rnf181	Rnf181.dSep08	297337	2046	742	5	62	ring finger protein 181 (7.4 kD) (Rnf181) alternative variant dSep08, complete mRNA.
Rnf181	Rnf181.eSep08	297337	2115	753	4	62	ring finger protein 181 (7.4 kD) (Rnf181) alternative variant eSep08, complete mRNA.
Rnf181	Rnf181.fSep08	297337	1286	993	2	30	ring finger protein 181 (3.7 kD) (Rnf181) alternative variant fSep08, mRNA.
Rnf185	Rnf185.bSep08	360967	12392	2385	3	143	ring finger protein 185 (Rnf185) alternative variant bSep08, mRNA.
Rnf187	Rnf187.aSep08	360533	5946	1937	2	277	ring finger protein 187 (Rnf187) alternative variant aSep08, mRNA.
Rnf187	Rnf187.bSep08	360533	4493	1050	1	153	ring finger protein 187 (Rnf187) alternative variant bSep08, mRNA.
Rnf190	Rnf190.bSep08	303596	58685	1786	5	595	ring finger protein 190 (Rnf190) alternative variant bSep08, mRNA.
Rnf190	Rnf190.dSep08	303596	33085	310	3	103	ring finger protein 190 (Rnf190) alternative variant dSep08, mRNA.
Rnf190	Rnf190.eSep08	303596	10832	537	3	32	ring finger protein 190 (3.6 kD) (Rnf190) alternative variant eSep08, mRNA.
Rnf207	Rnf207.aSep08	691246	12972	2441	17	635	ring finger protein 207 (70.6 kD) (Rnf207) alternative variant aSep08, mRNA.
Rnf207	Rnf207.bSep08	691246	7080	1596	10	168	ring finger protein 207 (18.8 kD) (Rnf207) alternative variant bSep08, mRNA.
Rnf207	Rnf207.dSep08	691246	4324	698	3	104	ring finger protein 207 (Rnf207) alternative variant dSep08, mRNA.
Rnf207	Rnf207.eSep08	691246	2490	879	5	59	ring finger protein 207 (Rnf207) alternative variant eSep08, mRNA.
Rnf207	Rnf207.fSep08	691246	891	493	2	74	ring finger protein 207 (Rnf207) alternative variant fSep08, mRNA.
Rnf215	Rnf215.bSep08	305478	1302	493	2	20	ring finger protein 215 (2.7 kD) (Rnf215) alternative variant bSep08, mRNA.
Rnf216	Rnf216.bSep08	304294	6828	330	2	87	ring finger protein 216 (Rnf216) alternative variant bSep08, mRNA.
Rnft1	Rnft1.aSep08	360595	9358	1736		288	ring finger protein, transmembrane 1 (Rnft1) mRNA.

Rnft2	Rnft2.cSep08	304521	12745	2958	4	129	ring finger protein, transmembrane 2 (Rnft2) alternative variant cSep08, mRNA.
Rnft2	Rnft2.dSep08	304521	8380	397	2	48	ring finger protein, transmembrane 2 (Rnft2) alternative variant dSep08, mRNA.
Rngtt	Rngtt.bSep08	313131	206346	3960	14	512	RNA guanylyltransferase and 5'-phosphatase (58.9 kD) (Rngtt) alternative variant bSep08, complete mRNA.
Rngtt	Rngtt.cSep08	313131	22820	525	5	114	RNA guanylyltransferase and 5'-phosphatase (Rngtt) alternative variant cSep08, mRNA.
Rnh1	Rnh1.bSep08	245964	9731	917	7	305	ribonuclease/angiogenin inhibitor 1 (Rnh1) alternative variant bSep08, mRNA.
Rnh1	Rnh1.cSep08	245964	5024	830	6	235	ribonuclease/angiogenin inhibitor 1 (Rnh1) alternative variant cSep08, mRNA.
Rnh1	Rnh1.dSep08	245964	4100	750	3	86	ribonuclease/angiogenin inhibitor 1 (9.6 kD) (Rnh1) alternative variant dSep08, mRNA.
Rnmt	Rnmt.bSep08	291534	10888	874	1	257	RNA (guanine-7-) methyltransferase (29.1 kD) (Rnmt) alternative variant bSep08, mRNA.
Rnpep	Rnpep.bSep08	81761	4352	597	3	162	arginyl aminopeptidase (aminopeptidase B) (Rnpep) alternative variant bSep08, mRNA.
Rnpep	Rnpep.cSep08	81761	6000	812	4	122	arginyl aminopeptidase (aminopeptidase B) (Rnpep) alternative variant cSep08, mRNA.
Rnps1	Rnps1.bSep08	287113	11479	2284	7	282	ribonucleic acid binding protein S1 (31.7 kD) (Rnps1) alternative variant bSep08, mRNA.
Rnps1	Rnps1.cSep08	287113	7563	735	6	202	ribonucleic acid binding protein S1 (Rnps1) alternative variant cSep08, mRNA.
Rnps1	Rnps1.dSep08	287113	7516	869	7	200	ribonucleic acid binding protein S1 (Rnps1) alternative variant dSep08, mRNA.
Rnps1	Rnps1.eSep08	287113	5868	999	4	139	ribonucleic acid binding protein S1 (Rnps1) alternative variant eSep08, mRNA.
Rnps1	Rnps1.fSep08	287113	661	345	2	84	ribonucleic acid binding protein S1 (Rnps1) alternative variant fSep08, mRNA.
Rnps1	Rnps1.hSep08	287113	7031	578	5	26	ribonucleic acid binding protein S1 (2.9 kD) (Rnps1) alternative variant hSep08, complete mRNA.
Robo1	Robo1.aSep08	58946	215361	576		191	roundabout homolog 1 (Drosophila) (Robo1) mRNA.
Robo2	Robo2.aSep08	84409	19492	1767	2	509	roundabout, axon guidance receptor, homolog 2 (Drosophila) (Robo2) alternative variant aSep08, mRNA.
Robo2	Robo2.bSep08	84409	10545	2030	1	185	roundabout, axon guidance receptor, homolog 2 (Drosophila) (Robo2) alternative variant bSep08, mRNA.
Robo2	Robo2.cSep08	84409	3441	615	1	125	roundabout, axon guidance receptor, homolog 2 (Drosophila) (Robo2) alternative variant cSep08, mRNA.
Robo3	Robo3.bSep08	315564	1891	892	5	210	roundabout homolog 3 (Drosophila) (Robo3) alternative variant bSep08, mRNA.
Robo3	Robo3.cSep08	315564	1929	711	4	130	roundabout homolog 3 (Drosophila) (Robo3) alternative variant cSep08, mRNA.
Robo4	Robo4.bSep08	300518	3407	1802	1	433	roundabout homolog 4 (Drosophila) (Robo4) alternative variant bSep08, mRNA.
roby	roby.aSep08		1044	399		107	putative protein (12.5 kD) (roby) mRNA.
rochy	rochy.aSep08		6120	326		108	CRA a (rochy) mRNA.



Rock1	Rock1.bSep08	81762	16290	2115	9	326	rho-associated coiled-coil containing protein kinase 1 (38.1 kD) (Rock1) alternative variant bSep08, mRNA.
Rock2	Rock2.bSep08	25537	3913	414	5	82	rho-associated coiled-coil containing protein kinase 2 (Rock2) alternative variant bSep08, mRNA.
Rock2	Rock2.cSep08	25537	2458	537	2	6	rho-associated coiled-coil containing protein kinase 2 (0.7 kD) (Rock2) alternative variant cSep08, mRNA.
Rod1	Rod1.bSep08	83515	3780	332	3	110	ROD1 regulator of differentiation 1 (S. pombe) (Rod1) alternative variant bSep08, mRNA.
Rod1	Rod1.cSep08	83515	44595	381	4	68	ROD1 regulator of differentiation 1 (S. pombe) (Rod1) alternative variant cSep08, mRNA.
rodar	rodar.aSep08		2482	619		41	putative protein (4.6 kD) (rodar) mRNA.
rofer	rofer.aSep08		2006	292		40	putative protein (rofer) mRNA.
roflo	roflo.aSep08		8661	371		53	polycomb group ring finger 6 (roflo) mRNA.
roflu	roflu.aSep08		11944	714		62	uncharacterized protein like (6.8 kD) (roflu) mRNA.
rogar	rogar.aSep08		2803	1098		67	putative protein (7.4 kD) (rogar) mRNA.
Rogdi	Rogdi.bSep08	287061	3229	730	7	123	rogdi homolog (Rogdi) alternative variant bSep08, mRNA.
Rogdi	Rogdi.cSep08	287061	1947	943	4	107	rogdi homolog (11.3 kD) (Rogdi) alternative variant cSep08, mRNA.
Rogdi	Rogdi.dSep08	287061	1703	831	3	86	rogdi homolog CRA a (Rogdi) alternative variant dSep08, mRNA.
roja	roja.aSep08		21369	558		185	protein CRA g (roja) mRNA.
rojey	rojey.aSep08		1976	462		65	CRA a like (rojey) mRNA.
rokee	rokee.aSep08		12667	494		41	putative protein (4.8 kD) (rokee) mRNA.
rokler	rokler.aSep08		13181	487		81	CRA b (9.3 kD) (rokler) mRNA.
rolo	rolo.aSep08		2724	849		62	putative protein (7.3 kD) (rolo) mRNA.
romeo	romeo.aSep08		6247	303			
romer	romer.aSep08		1713	780		259	ring finger protein 213 (romer) mRNA.
ronoy	ronoy.aSep08		1623	402		133	putative protein of metazoan origin (ronoy) mRNA.
Ropn1l	Ropn1l.aSep08	685646	3323	353		79	ropporin 1-like (Ropn1l) mRNA.
ropor	ropor.aSep08		3927	216		56	putative protein (ropor) mRNA.
Ror1	Ror1.bSep08	362550	2229	503	2	162	receptor tyrosine kinase-like orphan receptor 1 (Ror1) alternative variant bSep08, mRNA.
Ror1	Ror1.cSep08	362550	1734	479	3	128	receptor tyrosine kinase-like orphan receptor 1 (Ror1) alternative variant cSep08, mRNA.
Rora	Rora.bSep08	300807	6874	409	1	136	RAR-related orphan receptor alpha (Rora) alternative variant bSep08, mRNA.
Rorb	Rorb.aSep08	309288	17141	541		100	RAR-related orphan receptor beta (Rorb) mRNA.
rorby	rorby.aSep08		35717	556		30	putative protein (rorby) mRNA.
RorcandLingo4	RorcandLingo4.aSep08	368158	17887	2951	10	413	RAR-related orphan receptor C (46.3 kD) (RorcandLingo4) alternative variant aSep08, mRNA.
RorcandLingo4	RorcandLingo4.aSep08	499668	17887	2951	10	413	RAR-related orphan receptor C (46.3 kD) (RorcandLingo4) alternative variant aSep08, mRNA.
RorcandLingo4	RorcandLingo4.cSep08	368158	726	476	2	158	RAR-related orphan receptor C (RorcandLingo4) alternative variant cSep08, mRNA.

RorcandLingo4	RorcandLingo4.cSep08	499668	726	476	2	158	RAR-related orphan receptor C (RorcandLingo4) alternative variant cSep08, mRNA.
rorchy	rorchy.aSep08		792	397		94	putative protein (rorchy) mRNA.
rordar	rordar.aSep08		8224	728		242	transmembrane coiled-coil domains 7 (rordar) mRNA.
rorfer	rorfer.aSep08		1037	538		110	enhancer trap locus 4 (rorfer) mRNA.
rorflo	rorflo.aSep08		47692	706		29	putative protein (3.5 kD) (rorflo) mRNA.
rorflu	rorflu.aSep08		4092	648		108	putative nuclear protein (11.6 kD) (rorflu) mRNA.
rorgar	rorgar.aSep08		3345	568		98	putative protein (rorgar) mRNA.
rorja	rorja.aSep08		5261	504		82	putative protein (9.0 kD) (rorja) mRNA.
rorjey	rorjey.aSep08		44308	313		34	putative protein (rorjey) mRNA.
rorkee	rorkee.aSep08		6503	1067	4	113	putative nuclear protein of mammalian origin (13.2 kD) (rorkee) alternative variant aSep08, mRNA.
rorkler	rorkler.aSep08		742	571		84	putative secreted or extracellular protein precursor (9.3 kD) (rorkler) mRNA.
rorlo	rorlo.aSep08		7224	1757	2	175	furry homolog (rorlo) alternative variant aSep08, mRNA.
rorlo	rorlo.bSep08		1198	943	1	161	furry homolog (18.2 kD) (rorlo) alternative variant bSep08, mRNA.
ormee	ormee.aSep08		1801	420		84	putative protein (ormee) mRNA.
ormer	ormer.bSep08		7450	504	4	23	putative protein (ormer) alternative variant bSep08, mRNA.
ormer	ormer.dSep08		2354	645	3		
ormer	ormer.eSep08		7585	591	3		
ormer	ormer.fSep08		3269	486	3		
rornoy	rornoy.bSep08		4812	250	2	50	putative protein (rornoy) alternative variant bSep08, mRNA.
rorpor	rorpor.aSep08		43582	410		38	putative protein (4.1 kD) (rorpor) mRNA.
rorsa	rorsa.aSep08		5690	511		170	E1A binding protein p300 like (rorsa) mRNA.
rorshee	rorshee.aSep08		5067	932		99	putative secreted or extracellular protein precursor (11.3 kD) (rorshee) mRNA.
rortu	rortu.aSep08		1689	1163		81	putative mitochondrial protein (9.3 kD) (rortu) mRNA.
rorvar	rorvar.aSep08		1033	831		75	putative protein (rorvar) mRNA.
rorwey	rorwey.aSep08		1418	446		33	putative protein (3.8 kD) (rorwey) mRNA.
rosa	rosa.aSep08		1995	583		71	putative protein of vertebrate origin (7.3 kD) (rosa) mRNA.
roshee	roshee.aSep08		26810	802	1	36	putative protein (4.1 kD) (roshee) alternative variant aSep08, mRNA.
roshee	roshee.bSep08		27282	410	1	47	putative protein (roshee) alternative variant bSep08, mRNA.
Rotamase.0	Rotamase.0.aSep08		6061	515		80	protein NIMA-interacting 4 (8.9 kD) (Rotamase.0) alternative variant aSep08, mRNA.
Rotamase.0	Rotamase.0.bSep08		794	482		80	protein NIMA-interacting 4 (8.9 kD) (Rotamase.0) alternative variant bSep08, mRNA.
rotu	rotu.aSep08		9109	1034		276	putative protein (rotu) alternative variant aSep08, mRNA.
rovar	rovar.aSep08	689675	13774	3310		368	selenoprotein N 1 (rovar) mRNA.
rowey	rowey.aSep08		394	290		62	putative protein (7.0 kD) (rowey) mRNA.

royby	royby.aSep08		5681	708		236	transcriptional regulator atrx (royby) mRNA.
roychy	roychy.aSep08		7016	399		41	putative protein (roychy) mRNA.
roydar	roydar.aSep08		1090	536		64	putative protein (6.8 kD) (roydar) mRNA.
royfer	royfer.aSep08		2519	1800	3	173	enhancer trap locus 4 (royfer) alternative variant aSep08, mRNA.
royflo	royflo.aSep08		4395	755		68	putative protein (royflo) mRNA.
royflu	royflu.aSep08		871	297		98	putative protein (royflu) mRNA.
roygar	roygar.aSep08		48582	632		49	putative protein (5.7 kD) (roygar) mRNA.
royja	royja.aSep08		29144	378		125	putative protein of metazoan origin (royja) mRNA.
royjey	royjey.aSep08		47552	533		72	putative secreted or extracellular protein precursor (8.2 kD) (royjey) mRNA.
roykee	roykee.aSep08		58610	807		88	putative protein (10.1 kD) (roykee) mRNA.
roykler	roykler.aSep08		2233	1816		106	putative protein (11.1 kD) (roykler) mRNA.
roylo	roylo.aSep08		10169	703		234	furry homolog (roylo) mRNA.
roymee	roymee.aSep08		1467	298		28	putative protein (roymee) mRNA.
roymer	roymer.aSep08		2019	1175		110	putative protein (roymer) mRNA.
roynoy	roynoy.aSep08		792	410	2	76	obscurin-like 1 (roynoy) alternative variant aSep08, mRNA.
roypor	roypor.aSep08		3896	2382		73	adam metallopeptidase domain 10 (roypor) alternative variant aSep08, mRNA.
roysa	roysa.aSep08		2236	747	3	118	CRA a (roysa) alternative variant aSep08, mRNA.
roysa	roysa.bSep08		7655	578	3	117	CRA a (roysa) alternative variant bSep08, mRNA.
roysa	roysa.dSep08		7535	407	3	73	CRA a (roysa) alternative variant dSep08, mRNA.
royshee	royshee.aSep08		2164	710		17	putative protein (royshee) mRNA.
roytu	roytu.aSep08		548	447	2	128	uncharacterized protein c2orf53 homolog like (roytu) alternative variant aSep08, mRNA.
royvar	royvar.aSep08		408	294		91	putative protein (royvar) mRNA.
roywey	roywey.aSep08		744	656		99	putative protein (roywey) mRNA.
Rp2h	Rp2h.aSep08	367714	584	240		65	retinitis pigmentosa 2 homolog (human) (Rp2h) mRNA.
rp9	rp9.aSep08	363032	11989	861	3	170	retinitis pigmentosa 9 (human) (rp9) alternative variant aSep08, mRNA.
rp9	rp9.cSep08	363032	5201	433	1	103	retinitis pigmentosa 9 (human) (rp9) alternative variant cSep08, mRNA.
Rpa1	Rpa1.bSep08	287524	23117	2514	13	523	replication protein A1 (Rpa1) alternative variant bSep08, mRNA.
Rpa1	Rpa1.cSep08	287524	495	351	2	117	replication protein A1 (Rpa1) alternative variant cSep08, mRNA.
Rpa1	Rpa1.dSep08	287524	8086	401	4	81	replication protein A1 (Rpa1) alternative variant dSep08, mRNA.
Rpa2	Rpa2.bSep08	59102	7577	744	6	178	replication protein A2 (19.3 kD) (Rpa2) alternative variant bSep08, mRNA.
Rpa2	Rpa2.cSep08	59102	8476	742	6	111	replication protein A2 (Rpa2) alternative variant cSep08, mRNA.
Rpa2	Rpa2.dSep08	59102	3740	858	3	91	replication protein A2 (9.9 kD) (Rpa2) alternative variant dSep08, complete mRNA.

Rpa2	Rpa2.eSep08	59102	1191	868	2	82	replication protein A2 (Rpa2) alternative variant eSep08, mRNA.
Rpa3	Rpa3.cSep08	296883	1807	733	2	62	replication protein A3 (6.9 kD) (Rpa3) alternative variant cSep08, mRNA.
Rpa3	Rpa3.dSep08	296883	811	304	2	26	replication protein A3 (Rpa3) alternative variant dSep08, mRNA.
Rpain	Rpain.aSep08	287463	7467	1029	6	314	RPA interacting protein (Rpain) alternative variant aSep08, mRNA.
Rpain	Rpain.bSep08	287463	1542	450	1	91	RPA interacting protein (10.3 kD) (Rpain) alternative variant bSep08, mRNA.
Rpap1	Rpap1.bSep08	311338	1049	818	3	159	RNA polymerase II associated protein 1 (Rpap1) alternative variant bSep08, mRNA.
Rpap2	Rpap2.bSep08	305120	120444	2056	11	530	RNA polymerase II associated protein 2 (59.2 kD) (Rpap2) alternative variant bSep08, mRNA.
Rpap2	Rpap2.cSep08	305120	29116	527	5	175	RNA polymerase II associated protein 2 (Rpap2) alternative variant cSep08, mRNA.
Rpap2	Rpap2.dSep08	305120	15688	908	3	122	RNA polymerase II associated protein 2 (Rpap2) alternative variant dSep08, mRNA.
Rpap3	Rpap3.bSep08	300189	24346	1502	12	500	RNA polymerase II associated protein 3 (Rpap3) alternative variant bSep08, mRNA.
Rpap3	Rpap3.cSep08	300189	12211	1034	6	264	RNA polymerase II associated protein 3 (Rpap3) alternative variant cSep08, mRNA.
Rpap3	Rpap3.dSep08	300189	10388	1232	6	242	RNA polymerase II associated protein 3 (Rpap3) alternative variant dSep08, mRNA.
Rpap3	Rpap3.eSep08	300189	5029	523	4	174	RNA polymerase II associated protein 3 (Rpap3) alternative variant eSep08, mRNA.
Rpap3	Rpap3.iSep08	300189	2496	818	2	96	RNA polymerase II associated protein 3 (Rpap3) alternative variant iSep08, mRNA.
Rpe	Rpe.aSep08	501157	18805	828	6	228	ribulose-5-phosphate-3-epimerase (Rpe) alternative variant aSep08, mRNA.
Rpe	Rpe.bSep08	501157	19153	1187	6	228	ribulose-5-phosphate-3-epimerase (24.9 kD) (Rpe) alternative variant bSep08, mRNA.
Rpe	Rpe.dSep08	501157	16015	399	4	131	ribulose-5-phosphate-3-epimerase (Rpe) alternative variant dSep08, mRNA.
Rpe65	Rpe65.bSep08	89826	2243	408	3	136	retinal pigment epithelium 65 (Rpe65) alternative variant bSep08, mRNA.
Rpe65	Rpe65.cSep08	89826	2649	903	4	118	retinal pigment epithelium 65 (Rpe65) alternative variant cSep08, mRNA.
Rpe65	Rpe65.dSep08	89826	443	310	1	54	retinal pigment epithelium 65 (Rpe65) alternative variant dSep08, mRNA.
Rpresp	Rpresp.aSep08	297757	31191	1692	5	285	RPE-spondin (Rpresp) alternative variant aSep08, mRNA.
Rpresp	Rpresp.bSep08	297757	30420	857	6	239	RPE-spondin (Rpresp) alternative variant bSep08, mRNA.
Rpgrip1	Rpgrip1.bSep08	305850	11735	619	4	205	retinitis pigmentosa GTPase regulator interacting protein 1 (Rpgrip1) alternative variant bSep08, mRNA.
Rpgrip1	Rpgrip1.cSep08	305850	14017	322	4	107	retinitis pigmentosa GTPase regulator interacting protein 1 (Rpgrip1) alternative variant cSep08, mRNA.
Rpgrip1l	Rpgrip1l.bSep08	307724	14982	755	1	181	rprip1-like (Rpgrip1l) alternative variant bSep08, mRNA.

Rph3a	Rph3a.bSep08	171039	9290	561	6	187	rabphilin 3A (Rph3a) alternative variant bSep08, mRNA.
Rpia	Rpia.bSep08	362383	24768	766	6	244	ribose 5-phosphate isomerase A (25.9 kD) (Rpia) alternative variant bSep08, mRNA.
Rpia	Rpia.cSep08	362383	24632	747	8	216	ribose 5-phosphate isomerase A (23.7 kD) (Rpia) alternative variant cSep08, mRNA.
Rpia	Rpia.dSep08	362383	22971	741	9	192	ribose 5-phosphate isomerase A (Rpia) alternative variant dSep08, mRNA.
Rpia	Rpia.eSep08	362383	3152	285	4	43	ribose 5-phosphate isomerase A (Rpia) alternative variant eSep08, mRNA.
Rpl3	Rpl3.bSep08	300079	3707	957	6	180	ribosomal protein L3 (19.8 kD) (Rpl3) alternative variant bSep08, mRNA.
Rpl3	Rpl3.cSep08	300079	2513	1094	3	122	ribosomal protein L3 (13.9 kD) (Rpl3) alternative variant cSep08, mRNA.
Rpl3	Rpl3.dSep08	300079	10483	294	3	97	putative protein of ancient origin (Rpl3) alternative variant dSep08, mRNA.
Rpl3l	Rpl3l.aSep08	287122	10551	1343		407	ribosomal protein L3 (46.4 kD) (Rpl3l) complete mRNA.
Rpl4	Rpl4.bSep08	64302	2097	969	4	137	ribosomal protein L4/L1e (Rpl4) alternative variant bSep08, mRNA.
Rpl4	Rpl4.cSep08	64302	3615	1019	5	103	ribosomal protein L4/L1e (10.9 kD) (Rpl4) alternative variant cSep08, complete mRNA.
Rpl7	Rpl7.aSep08	297755	3142	1621	5	290	ribosomal L30, N-terminal and ribosomal protein L30 (33.7 kD) (Rpl7) alternative variant aSep08, mRNA.
Rpl7	Rpl7.cSep08	297755	2450	1278	4	212	ribosomal L30, N-terminal and ribosomal protein L30 (24.6 kD) (Rpl7) alternative variant cSep08, complete mRNA.
Rpl7a.1	Rpl7a.1.bSep08	296596	2628	1355	6	256	ribosomal protein L7Ae/L30e/S12e/Gadd45 (28.7 kD) (Rpl7a.1) alternative variant bSep08, mRNA.
Rpl7a.1	Rpl7a.1.cSep08	296596	2141	896	5	186	putative protein of eukaryotic origin (Rpl7a.1) alternative variant cSep08, mRNA.
Rpl7a.1	Rpl7a.1.dSep08	296596	2468	1327	6	160	ribosomal protein L7Ae/L30e/S12e/Gadd45 (17.7 kD) (Rpl7a.1) alternative variant dSep08, complete mRNA.
Rpl7a.1	Rpl7a.1.eSep08	296596	1661	463	4	154	ribosomal protein L7Ae/L30e/S12e/Gadd45 (Rpl7a.1) alternative variant eSep08, mRNA.
Rpl7a.1	Rpl7a.1.fSep08	296596	959	520	3	105	putative protein of eukaryotic origin (Rpl7a.1) alternative variant fSep08, mRNA.
Rpl711	Rpl711.aSep08	317275	7738	2219	4	257	ribosomal protein L30 (Rpl711) alternative variant aSep08, mRNA.
Rpl711	Rpl711.bSep08	317275	5587	771	3	221	ribosomal protein L30 (26.0 kD) (Rpl711) alternative variant bSep08, mRNA.
Rpl711	Rpl711.cSep08	317275	1065	929	1	78	putative protein of eukaryotic origin (Rpl711) alternative variant cSep08, mRNA.
Rpl711	Rpl711.dSep08	317275	978	273		37	putative protein of vertebrate origin (Rpl711) alternative variant dSep08, mRNA.
Rpl8	Rpl8.cSep08	26962	898	771	2	86	putative protein (Rpl8) alternative variant cSep08, mRNA.
Rpl9	Rpl9.aSep08	29257	2797	701	7	207	ribosomal protein L6 (Rpl9) alternative variant aSep08, mRNA.
Rpl9	Rpl9.cSep08	29257	1169	480	3	90	ribosomal protein L6 (Rpl9) alternative variant cSep08, mRNA.

Rpl9	Rpl9.dSep08	29257	808	494	2	80	ribosomal protein L6 (9.1 kD) (Rpl9) alternative variant dSep08, mRNA.
Rpl10	Rpl10.bSep08	81764	2073	1189	4	110	ribosomal protein L16 (12.7 kD) (Rpl10) alternative variant bSep08, complete mRNA.
Rpl10	Rpl10.cSep08	81764	1567	1287	3	73	putative protein (8.6 kD) (Rpl10) alternative variant cSep08, mRNA.
Rpl10a	Rpl10a.bSep08	81729	2520	1112	5	133	ribosomal protein L1 (15.1 kD) (Rpl10a) alternative variant bSep08, mRNA.
Rpl10a	Rpl10a.cSep08	81729	583	401	2	74	ribosomal protein L1 (8.5 kD) (Rpl10a) alternative variant cSep08, mRNA.
Rpl10a	Rpl10a.dSep08	81729	1637	1342	3	58	putative protein of eukaryotic origin (6.8 kD) (Rpl10a) alternative variant dSep08, mRNA.
Rpl11	Rpl11.aSep08	362631	3703	759	6	185	ribosomal protein L5 (Rpl11) alternative variant aSep08, mRNA.
Rpl13	Rpl13.aSep08	81765	1770	806	5	233	ribosomal protein L13e (26.2 kD) (Rpl13) alternative variant aSep08, mRNA.
Rpl13	Rpl13.cSep08	81765	1479	1388	2	125	putative protein (Rpl13) alternative variant cSep08, mRNA.
Rpl13	Rpl13.dSep08	81765	1680	910	4	72	putative protein (Rpl13) alternative variant dSep08, mRNA.
Rpl13a	Rpl13a.bSep08	317646	3061	1107	7	183	ribosomal protein L13 (Rpl13a) alternative variant bSep08, mRNA.
Rpl13a	Rpl13a.cSep08	317646	2687	960	6	165	ribosomal protein L13 (18.9 kD) (Rpl13a) alternative variant cSep08, complete mRNA.
Rpl13a	Rpl13a.dSep08	317646	2975	952	8	148	ribosomal protein L13 (16.5 kD) (Rpl13a) alternative variant dSep08, complete mRNA.
Rpl13a	Rpl13a.eSep08	317646	1576	793	6	142	ribosomal protein L13 (16.6 kD) (Rpl13a) alternative variant eSep08, mRNA.
Rpl13a	Rpl13a.fSep08	317646	1628	699	6	142	ribosomal protein L13 (16.6 kD) (Rpl13a) alternative variant fSep08, mRNA.
Rpl13a	Rpl13a.gSep08	317646	2697	821	8	142	ribosomal protein L13 (16.6 kD) (Rpl13a) alternative variant gSep08, complete mRNA.
Rpl13a	Rpl13a.hSep08	317646	798	595	2	90	putative protein (Rpl13a) alternative variant hSep08, mRNA.
Rpl14	Rpl14.aSep08	65043	3771	1614	5	219	ribosomal protein L14 (23.8 kD) (Rpl14) alternative variant aSep08, complete mRNA.
Rpl14	Rpl14.bSep08	65043	1315	750	1	65	putative protein of eukaryotic origin (Rpl14) alternative variant bSep08, mRNA.
Rpl14	Rpl14.cSep08	65043	756	677	2	58	ribosomal protein L14 (Rpl14) alternative variant cSep08, mRNA.
Rpl14	Rpl14.dSep08	65043	2387	869	3	56	ribosomal protein L14 (6.8 kD) (Rpl14) alternative variant dSep08, mRNA.
Rpl15	Rpl15.aSep08	245981	1985	748	4	204	ribosomal protein L15e (24.1 kD) (Rpl15) alternative variant aSep08, mRNA.
Rpl15	Rpl15.dSep08	245981	760	519	2	89	ribosomal protein L15e (Rpl15) alternative variant dSep08, mRNA.
Rpl17	Rpl17.bSep08	291434	2273	576	6	171	ribosomal protein L22/L17 (20.0 kD) (Rpl17) alternative variant bSep08, complete mRNA.

Rpl17	Rpl17.cSep08	291434	3101	556	6	146	protein ribosomal L17 (17.1 kD) (Rpl17) alternative variant cSep08, complete mRNA.
Rpl17	Rpl17.dSep08	291434	1432	820	4	125	ribosomal protein L22/L17 (Rpl17) alternative variant dSep08, mRNA.
Rpl17	Rpl17.eSep08	291434	2160	877	4	116	protein ribosomal L17 (13.6 kD) (Rpl17) alternative variant eSep08, mRNA.
Rpl17	Rpl17.fSep08	291434	1459	1354	2	32	putative protein of eukaryotic origin (3.8 kD) (Rpl17) alternative variant fSep08, mRNA.
Rpl18	Rpl18.aSep08	81766	2156	716	6	193	ribosomal protein L18e (22.2 kD) (Rpl18) alternative variant aSep08, mRNA.
Rpl18	Rpl18.dSep08	81766	1158	719	3	93	ribosomal protein L18e (Rpl18) alternative variant dSep08, mRNA.
Rpl18	Rpl18.eSep08	81766	2149	813	5	75	ribosomal protein L18e (Rpl18) alternative variant eSep08, mRNA.
Rpl18	Rpl18.fSep08	81766	984	372	3	71	putative nuclear protein of eukaryotic origin (8.1 kD) (Rpl18) alternative variant fSep08, mRNA.
Rpl18	Rpl18.gSep08	81766	2620	1380	5	113	ribosomal protein L18e (12.6 kD) (Rpl18) alternative variant gSep08, complete mRNA.
Rpl18a	Rpl18a.bSep08	290641	865	682	1	102	ribosomal L18ae protein (Rpl18a) alternative variant bSep08, mRNA.
Rpl19	Rpl19.bSep08	81767	1515	854	3	108	ribosomal protein L19e (13.4 kD) (Rpl19) alternative variant bSep08, mRNA.
Rpl19	Rpl19.dSep08	81767	1590	251	3	83	ribosomal protein L19e (Rpl19) alternative variant dSep08, mRNA.
Rpl21	Rpl21.aSep08	79449	3051	778	6	160	ribosomal protein L21e (18.6 kD) (Rpl21) alternative variant aSep08, mRNA.
Rpl21	Rpl21.bSep08	79449	3707	1327	6	160	ribosomal protein L21e (18.6 kD) (Rpl21) alternative variant bSep08, complete mRNA.
Rpl21	Rpl21.dSep08	79449	1438	683	3	60	ribosomal protein L21e (Rpl21) alternative variant dSep08, mRNA.
Rpl21	Rpl21.eSep08	79449	1211	601	2	42	putative protein (Rpl21) alternative variant eSep08, mRNA.
Rpl22	Rpl22.aSep08	81768	8160	1889	4	137	ribosomal L22e protein (Rpl22) alternative variant aSep08, mRNA.
Rpl22	Rpl22.bSep08	81768	6841	729	5	122	ribosomal L22e protein (14.4 kD) (Rpl22) alternative variant bSep08, mRNA.
Rpl22	Rpl22.cSep08	81768	6691	1952	3	95	ribosomal L22e protein (11.2 kD) (Rpl22) alternative variant cSep08, complete mRNA.
Rpl22l1	Rpl22l1.bSep08	361923	869	517	3	82	ribosomal L22e protein (9.7 kD) (Rpl22l1) alternative variant bSep08, complete mRNA.
Rpl22l1	Rpl22l1.cSep08	361923	1814	858	2	43	putative protein of vertebrate origin (Rpl22l1) alternative variant cSep08, mRNA.
Rpl24	Rpl24.bSep08	64307	5298	597	5	131	ribosomal protein L24E (14.8 kD) (Rpl24) alternative variant bSep08, mRNA.
Rpl24	Rpl24.cSep08	64307	5191	1635	4	79	ribosomal protein L24E (Rpl24) alternative variant cSep08, mRNA.
Rpl24	Rpl24.dSep08	64307	3271	615	2	57	CRA c like (6.2 kD) (Rpl24) alternative variant dSep08, mRNA.

Rpl26	Rpl26.bSep08	287417	3111	527	4	145	KOW (17.3 kD) (Rpl26) alternative variant bSep08, complete mRNA.
Rpl26	Rpl26.cSep08	287417	5903	668	4	110	KOW (Rpl26) alternative variant cSep08, mRNA.
Rpl27	Rpl27.bSep08	64306	3752	515	5	136	KOW and ribosomal protein L27e (15.8 kD) (Rpl27) alternative variant bSep08, mRNA.
Rpl27	Rpl27.cSep08	64306	3728	551	5	109	KOW (Rpl27) alternative variant cSep08, mRNA.
Rpl27a	Rpl27a.bSep08	293418	1257	640	1	47	putative protein of fungal and metazoan origin (5.4 kD) (Rpl27a) alternative variant bSep08, mRNA.
Rpl28	Rpl28.aSep08	64638	1495	788	4	245	ribosomal L28e protein (Rpl28) alternative variant aSep08, mRNA.
Rpl28	Rpl28.cSep08	64638	1288	663	3	74	putative mitochondrial protein of vertebrate origin (8.6 kD) (Rpl28) alternative variant cSep08, mRNA.
Rpl29	Rpl29.bSep08	29283	1944	774	4	156	ribosomal L29e protein (17.3 kD) (Rpl29) alternative variant bSep08, mRNA.
Rpl29	Rpl29.cSep08	29283	2009	598	4	156	ribosomal L29e protein (17.3 kD) (Rpl29) alternative variant cSep08, complete mRNA.
Rpl29	Rpl29.dSep08	29283	866	574	2	37	ribosomal L29e protein (Rpl29) alternative variant dSep08, mRNA.
Rpl30	Rpl30.aSep08	64640	3063	740	3	115	ribosomal protein L7Ae/L30e/S12e/Gadd45 (12.8 kD) (Rpl30) alternative variant aSep08, mRNA.
Rpl30	Rpl30.cSep08	64640	2821	1302	2	114	ribosomal protein L7Ae/L30e/S12e/Gadd45 (12.8 kD) (Rpl30) alternative variant cSep08, mRNA.
Rpl31	Rpl31.aSep08	64298	14522	825	4	233	ribosomal protein L31e (26.5 kD) (Rpl31) alternative variant aSep08, complete mRNA.
Rpl31	Rpl31.bSep08	64298	3461	591	4	125	ribosomal protein L31e (14.5 kD) (Rpl31) alternative variant bSep08, complete mRNA.
Rpl34	Rpl34.aSep08	362041	3710	603	1	117	ribosomal protein L34e (13.3 kD) (Rpl34) alternative variant aSep08, mRNA.
Rpl34	Rpl34.bSep08	362041	3749	446	1	117	ribosomal protein L34e (13.3 kD) (Rpl34) alternative variant bSep08, complete mRNA.
Rpl35	Rpl35.bSep08	296709	2941	2224	3	126	ribosomal protein L29 (14.0 kD) (Rpl35) alternative variant bSep08, complete mRNA.
Rpl35	Rpl35.cSep08	296709	2914	643	3	124	ribosomal protein L29 (14.7 kD) (Rpl35) alternative variant cSep08, mRNA.
Rpl35a.1	Rpl35a.1.aSep08	57809	3871	458	5	140	ribosomal protein L35Ae (Rpl35a.1) alternative variant aSep08, mRNA.
Rpl35a.1	Rpl35a.1.bSep08	57809	3239	1146	4	135	ribosomal protein L35Ae (15.2 kD) (Rpl35a.1) alternative variant bSep08, mRNA.
Rpl35a.1	Rpl35a.1.dSep08	57809	3895	436	5	110	ribosomal protein L35Ae (12.6 kD) (Rpl35a.1) alternative variant dSep08, complete mRNA.
Rpl36.1	Rpl36.1.bSep08	58927	5392	467	4	105	ribosomal protein L36E (12.3 kD) (Rpl36.1) alternative variant bSep08, mRNA.
Rpl36.1	Rpl36.1.cSep08	58927	995	737	3	105	ribosomal protein L36E (12.3 kD) (Rpl36.1) alternative variant cSep08, mRNA.
Rpl36a	Rpl36a.cSep08	292964	944	630	2	50	CRA a like (5.5 kD) (Rpl36a) alternative variant cSep08, mRNA.



Rpl36al	Rpl36al.aSep08	81769	1158	499	1	106	ribosomal protein L44E (12.4 kD) (Rpl36al) alternative variant aSep08, mRNA.
Rpl37	Rpl37.bSep08	81770	1939	1181	3	98	ribosomal protein L37e (11.3 kD) (Rpl37) alternative variant bSep08, complete mRNA.
Rpl37a_predicted	Rpl37a_predicted.bSep08	363248	1336	942	2	82	ribosomal L37ae protein (9.2 kD) (Rpl37a_predicted) alternative variant bSep08, mRNA.
Rpl37a_predicted	Rpl37a_predicted.cSep08	363248	683	340	2	75	ribosomal L37ae protein (Rpl37a_predicted) alternative variant cSep08, mRNA.
Rpl39	Rpl39.bSep08	25347	2319	402	3	26	putative protein (Rpl39) alternative variant bSep08, mRNA.
Rpl41.1	Rpl41.1.aSep08	124440	1061	426	3	25	putative protein (3.5 kD) (Rpl41.1) alternative variant aSep08, complete mRNA.
Rpl41.1	Rpl41.1.bSep08	124440	981	460	2	25	putative protein (3.5 kD) (Rpl41.1) alternative variant bSep08, complete mRNA.
Rpl41.1	Rpl41.1.cSep08	124440	1048	716	2	40	putative protein, with a coiled coil domain (4.6 kD) (Rpl41.1) alternative variant cSep08, complete mRNA.
Rplp1.1	Rplp1.1.aSep08	140661	1344	506	3	114	ribosomal protein 60S (11.5 kD) (Rplp1.1) alternative variant aSep08, complete mRNA.
Rplp2	Rplp2.aSep08	140662	2501	657	3	115	ribosomal protein 60S (11.7 kD) (Rplp2) alternative variant aSep08, mRNA.
Rplp2	Rplp2.bSep08	140662	2271	424	3	114	ribosomal protein 60S (11.6 kD) (Rplp2) alternative variant bSep08, complete mRNA.
Rplp2	Rplp2.cSep08	140662	1625	746	1	70	putative protein of eukaryotic origin (Rplp2) alternative variant cSep08, mRNA.
Rpn1	Rpn1.aSep08	25596	11103	1589	7	396	ribophorin I (Rpn1) alternative variant aSep08, mRNA.
Rpn1	Rpn1.eSep08	25596	14048	298	3	32	ribophorin I (Rpn1) alternative variant eSep08, mRNA.
Rpn2	Rpn2.bSep08	64701	32155	1439	10	479	ribophorin II (Rpn2) alternative variant bSep08, mRNA.
Rpn2	Rpn2.cSep08	64701	20614	945	9	314	ribophorin II (Rpn2) alternative variant cSep08, mRNA.
Rpn2	Rpn2.dSep08	64701	16986	734	7	240	ribophorin II (Rpn2) alternative variant dSep08, mRNA.
Rpn2	Rpn2.eSep08	64701	11315	1264	8	201	ribophorin II (22.8 kD) (Rpn2) alternative variant eSep08, mRNA.
Rpo1-4	Rpo1-4.bSep08	83581	6285	703	1	225	RNA polymerase 1-4 (Rpo1-4) alternative variant bSep08, mRNA.
Rpp14	Rpp14.aSep08	361020	9443	1302	4	122	ribonuclease P 14 subunit (human) (13.4 kD) (Rpp14) alternative variant aSep08, mRNA.
Rpp14	Rpp14.cSep08	361020	9616	1783	2	44	ribonuclease P 14 subunit (human) (5.1 kD) (Rpp14) alternative variant cSep08, mRNA.
Rpp21	Rpp21.aSep08	406230	1867	605	3	187	ribonuclease P 21 subunit (human) (Rpp21) alternative variant aSep08, mRNA.
Rpp21	Rpp21.dSep08	406230	1929	591	4	117	ribonuclease P 21 subunit (human) (13.4 kD) (Rpp21) alternative variant dSep08, complete mRNA.
Rpp21	Rpp21.eSep08	406230	1932	576	4	117	ribonuclease P 21 subunit (human) (13.6 kD) (Rpp21) alternative variant eSep08, complete mRNA.
Rpp21	Rpp21.fSep08	406230	1918	567	4	98	ribonuclease P 21 subunit (human) (11.3 kD) (Rpp21) alternative variant fSep08, complete mRNA.
Rpp21	Rpp21.gSep08	406230	1915	488	5	98	ribonuclease P 21 subunit (human) (11.3 kD) (Rpp21) alternative variant gSep08, complete mRNA.

Rpp30	Rpp30.aSep08	685332	19322	1151	8	192	ribonuclease P/MRP 30 subunit (human) (21.2 kD) (Rpp30) alternative variant aSep08, mRNA.
Rpp30	Rpp30.bSep08	685332	11822	1220	6	156	ribonuclease P/MRP 30 subunit (human) (17.3 kD) (Rpp30) alternative variant bSep08, mRNA.
Rpp30	Rpp30.dSep08	685332	22530	1012	10	92	ribonuclease P/MRP 30 subunit (human) (10.4 kD) (Rpp30) alternative variant dSep08, complete mRNA.
Rpp38	Rpp38.aSep08	291317	3516	1481	1	272	ribonuclease P/MRP 38 subunit (human) (30.1 kD) (Rpp38) alternative variant aSep08, mRNA.
Rpp38	Rpp38.cSep08	291317	3293	778	1	194	ribonuclease P/MRP 38 subunit (human) (Rpp38) alternative variant cSep08, mRNA.
Rpp40	Rpp40.aSep08	291071	9443	1785	7	512	ribonuclease P 40 subunit (human) (Rpp40) alternative variant aSep08, mRNA.
Rpp40	Rpp40.cSep08	291071	7396	763	6	254	ribonuclease P 40 subunit (human) (Rpp40) alternative variant cSep08, mRNA.
Rpp40	Rpp40.dSep08	291071	2351	630	2	129	ribonuclease P 40 subunit (human) (Rpp40) alternative variant dSep08, mRNA.
Rps2	Rps2.aSep08	83789	1872	959	7	302	ribosomal protein S5, N-terminal and ribosomal protein S5, C-terminal (Rps2) alternative variant aSep08, mRNA.
Rps2	Rps2.bSep08	83789	1697	840	7	266	ribosomal protein S5, N-terminal and ribosomal protein S5, C-terminal (Rps2) alternative variant bSep08, mRNA.
Rps2	Rps2.cSep08	83789	1726	976	6	260	ribosomal protein S5, N-terminal and ribosomal protein S5, C-terminal (27.4 kD) (Rps2) alternative variant cSep08, mRNA.
Rps2	Rps2.dSep08	83789	1376	745	5	238	ribosomal protein S5, N-terminal and ribosomal protein S5, C-terminal (25.0 kD) (Rps2) alternative variant dSep08, mRNA.
Rps2	Rps2.eSep08	83789	1151	905	3	127	CRA b (13.4 kD) (Rps2) alternative variant eSep08, mRNA.
Rps2	Rps2.fSep08	83789	1279	1112	2	55	putative protein of ancient origin (6.4 kD) (Rps2) alternative variant fSep08, mRNA.
Rps2.1	Rps2.1.bSep08	83789	464	365	2	102	ribosomal protein S5, C-terminal (Rps2.1) alternative variant bSep08, mRNA.
Rps2.1	Rps2.1.cSep08	83789	680	369	2	56	putative protein of eukaryotic origin (6.4 kD) (Rps2.1) alternative variant cSep08, mRNA.
Rps3	Rps3.bSep08	140654	4031	828	5	183	KH, type 2 and ribosomal protein S3, C-terminal (Rps3) alternative variant bSep08, mRNA.
Rps3	Rps3.cSep08	140654	4557	837	5	180	KH, type 2 and ribosomal protein S3, C-terminal (Rps3) alternative variant cSep08, mRNA.
Rps3	Rps3.dSep08	140654	5300	487	6	151	KH, type 2 (16.6 kD) (Rps3) alternative variant dSep08, complete mRNA.
Rps4x	Rps4x.bSep08	29426	4054	949	4	237	ribosomal protein S4E, N-terminal and RNA-binding S4 and ribosomal protein S4E, central and KOW (26.7 kD) (Rps4x) alternative variant bSep08, mRNA.
Rps4x	Rps4x.cSep08	29426	1929	731	1	89	ribosomal protein S4E, N-terminal (10.2 kD) (Rps4x) alternative variant cSep08, mRNA.
Rps5	Rps5.aSep08	25538	4298	765	6	236	ribosomal protein S7 (Rps5) alternative variant aSep08, mRNA.

Rps6	Rps6.bSep08	29304	3084	1536	4	218	ribosomal protein S6e (25.0 kD) (Rps6) alternative variant bSep08, mRNA.
Rps6ka1	Rps6ka1.bSep08	81771	2724	737	2	81	putative protein of metazoan origin (9.1 kD) (Rps6ka1) alternative variant bSep08, mRNA.
Rps6ka3	Rps6ka3.aSep08	501560	26276	1661	9	348	protein kinase and tyrosine protein kinase (Rps6ka3) alternative variant aSep08, mRNA.
Rps6ka3	Rps6ka3.bSep08	501560	39868	562	7	187	tyrosine protein kinase (Rps6ka3) alternative variant bSep08, mRNA.
Rps6ka3	Rps6ka3.cSep08	501560	100983	671	8	162	putative protein of eukaryotic origin (Rps6ka3) alternative variant cSep08, mRNA.
Rps6ka6	Rps6ka6.aSep08	317203	26005	1682	8	391	protein kinase, C-terminal and tyrosine protein kinase (Rps6ka6) alternative variant aSep08, mRNA.
Rps6ka6	Rps6ka6.bSep08	317203	36891	843	6	153	putative protein of eukaryotic origin (17.2 kD) (Rps6ka6) alternative variant bSep08, mRNA.
Rps6kb1	Rps6kb1.bSep08	83840	27268	718	8	230	protein S6 kinase (Rps6kb1) alternative variant bSep08, mRNA.
Rps6kb1	Rps6kb1.cSep08	83840	22530	875	9	177	putative cytoplasmic protein of eukaryotic origin (19.9 kD) (Rps6kb1) alternative variant cSep08, mRNA.
Rps6kb1	Rps6kb1.dSep08	83840	4555	516	5	149	kinase 70 ribosomal S6 (Rps6kb1) alternative variant dSep08, mRNA.
Rps6kb2	Rps6kb2.cSep08	361696	730	620	2	117	kinase 70 ribosomal S6 (Rps6kb2) alternative variant cSep08, mRNA.
Rps6kb2	Rps6kb2.dSep08	361696	1673	700	7	111	kinase ribosomal S6 70 (Rps6kb2) alternative variant dSep08, mRNA.
Rps6kc1	Rps6kc1.aSep08	289342	65773	2774	5	855	putative protein of metazoan origin (Rps6kc1) alternative variant aSep08, mRNA.
Rps6kc1	Rps6kc1.bSep08	289342	24579	1595	5	294	putative protein of eukaryotic origin (Rps6kc1) alternative variant bSep08, mRNA.
Rps6kc1	Rps6kc1.cSep08	289342	4616	463	3	54	putative protein of vertebrate origin (Rps6kc1) alternative variant cSep08, mRNA.
Rps6kc1	Rps6kc1.dSep08	289342	3373	139	2	46	putative protein (Rps6kc1) alternative variant dSep08, mRNA.
Rps6kl1	Rps6kl1.aSep08	299202	4022	956	6	318	putative protein of eukaryotic origin (Rps6kl1) alternative variant aSep08, mRNA.
Rps7.1	Rps7.1.aSep08	29258	45939	1377	7	288	collectin sub-family member 11 (Rps7.1) alternative variant aSep08, mRNA.
Rps7.1	Rps7.1.aSep08	497813	45939	1377	7	288	collectin sub-family member 11 (Rps7.1) alternative variant aSep08, mRNA.
Rps7.1	Rps7.1.bSep08	29258	35317	950	7	271	collectin sub-family member 11 precursor (28.9 kD) (Rps7.1) alternative variant bSep08, mRNA.
Rps7.1	Rps7.1.bSep08	497813	35317	950	7	271	collectin sub-family member 11 precursor (28.9 kD) (Rps7.1) alternative variant bSep08, mRNA.
Rps7.1	Rps7.1.dSep08	29258	1987	845	3	105	ribosomal protein S7E (11.7 kD) (Rps7.1) alternative variant dSep08, mRNA.
Rps7.1	Rps7.1.dSep08	497813	1987	845	3	105	ribosomal protein S7E (11.7 kD) (Rps7.1) alternative variant dSep08, mRNA.

Rps7.1	Rps7.1.eSep08	29258	14861	934	3	79	collectin sub-family member 11 (Rps7.1) alternative variant eSep08, mRNA.
Rps7.1	Rps7.1.eSep08	497813	14861	934	3	79	collectin sub-family member 11 (Rps7.1) alternative variant eSep08, mRNA.
Rps8	Rps8.bSep08	65136	1706	742	3	101	ribosomal protein S8E (Rps8) alternative variant bSep08, mRNA.
Rps8	Rps8.cSep08	65136	975	595	2	88	putative nuclear protein of eukaryotic origin (9.9 kD) (Rps8) alternative variant cSep08, mRNA.
Rps8	Rps8.dSep08	65136	889	306	3	63	CRA b like (6.9 kD) (Rps8) alternative variant dSep08, mRNA.
Rps9	Rps9.bSep08	81772	3402	929	6	139	ribosomal protein S4 (16.6 kD) (Rps9) alternative variant bSep08, mRNA.
Rps9	Rps9.cSep08	81772	3362	1460	4	136	ribosomal protein S4 (16.3 kD) (Rps9) alternative variant cSep08, mRNA.
Rps9	Rps9.dSep08	81772	853	635	2	91	ribosomal protein S4 (Rps9) alternative variant dSep08, mRNA.
Rps10	Rps10.aSep08	81773	4585	577	6	165	plectin/S10, N-terminal (18.9 kD) (Rps10) alternative variant aSep08, complete mRNA.
Rps10	Rps10.cSep08	81773	2571	964	2	83	putative protein (Rps10) alternative variant cSep08, mRNA.
Rps11	Rps11.bSep08	81774	1719	1231	3	140	ribosomal protein S17 (16.4 kD) (Rps11) alternative variant bSep08, complete mRNA.
Rps11	Rps11.cSep08	81774	2157	926	4	134	ribosomal protein S17 (Rps11) alternative variant cSep08, mRNA.
Rps11	Rps11.dSep08	81774	1452	1108	2	79	ribosomal protein S17 (9.5 kD) (Rps11) alternative variant dSep08, mRNA.
Rps11	Rps11.eSep08	81774	1128	354	3	38	putative protein (4.3 kD) (Rps11) alternative variant eSep08, mRNA.
Rps12	Rps12.aSep08	65139	1098	862	3	133	putative protein human specific (Rps12) alternative variant aSep08, mRNA.
Rps12	Rps12.bSep08	65139	2074	572	4	132	ribosomal protein L7Ae/L30e/S12e/Gadd45 (14.5 kD) (Rps12) alternative variant bSep08, mRNA.
Rps12	Rps12.dSep08	65139	1951	1545	1	98	CRA c like (11.3 kD) (Rps12) alternative variant dSep08, mRNA.
Rps13	Rps13.bSep08	161477	1639	760	4	136	ribosomal S13S15 N-terminal and ribosomal protein S15 (15.4 kD) (Rps13) alternative variant bSep08, mRNA.
Rps13	Rps13.cSep08	161477	2168	526	4	124	ribosomal S13S15 N-terminal and ribosomal protein S15 (14.2 kD) (Rps13) alternative variant cSep08, mRNA.
Rps14	Rps14.bSep08	29284	4400	745	5	151	ribosomal protein S11 (16.3 kD) (Rps14) alternative variant bSep08, mRNA.
Rps15a.1	Rps15a.1.aSep08	117053	6819	541	4	130	ribosomal protein S8 (14.8 kD) (Rps15a.1) alternative variant aSep08, complete mRNA.
Rps15a.1	Rps15a.1.bSep08	117053	7235	850	4	130	ribosomal protein S8 (14.8 kD) (Rps15a.1) alternative variant bSep08, complete mRNA.
Rps15a.1	Rps15a.1.cSep08	117053	2009	820	2	81	ribosomal protein S8 (Rps15a.1) alternative variant cSep08, complete mRNA.
Rps16	Rps16.aSep08	140655	2944	817	5	170	ribosomal protein S9 (Rps16) alternative variant aSep08, mRNA.

Rps16	Rps16.cSep08	140655	2639	644	6	87	ribosomal protein S9 (10.0 kD) (Rps16) alternative variant cSep08, complete mRNA.
Rps17	Rps17.bSep08	29286	2599	708	4	78	ribosomal protein S17e (8.9 kD) (Rps17) alternative variant bSep08, complete mRNA.
Rps18	Rps18.aSep08	294282	2020	1826	3	82	ribosomal protein S13 (9.8 kD) (Rps18) alternative variant aSep08, mRNA.
Rps18	Rps18.bSep08	294282	2700	230	2	76	ribosomal protein S13 (Rps18) alternative variant bSep08, mRNA.
Rps19	Rps19.bSep08	29287	1312	551	1	98	putative mitochondrial protein (10.7 kD) (Rps19) alternative variant bSep08, mRNA.
Rps19	Rps19.cSep08	29287	1779	585	2	71	ribosomal protein S19e (7.9 kD) (Rps19) alternative variant cSep08, mRNA.
Rps19bp1	Rps19bp1.aSep08	500907	3279	703	4	138	protein S19 binding like (Rps19bp1) alternative variant aSep08, mRNA.
Rps20	Rps20.aSep08	122772	1391	731	3	119	ribosomal protein S10 (13.4 kD) (Rps20) alternative variant aSep08, complete mRNA.
Rps20	Rps20.bSep08	122772	711	436	1	91	protein ribosomal 20 (10.1 kD) (Rps20) alternative variant bSep08, mRNA.
Rps20	Rps20.cSep08	122772	907	436	2	64	ribosomal protein S10 (7.2 kD) (Rps20) alternative variant cSep08, mRNA.
Rps21.1	Rps21.1.aSep08	81775	1058	380	2	105	ribosomal protein S21e (Rps21.1) alternative variant aSep08, mRNA.
Rps21.1	Rps21.1.bSep08	81775	1022	499	1	83	ribosomal protein S21e (9.1 kD) (Rps21.1) alternative variant bSep08, complete mRNA.
Rps24	Rps24.aSep08	81776	4652	2739	4	193	ribosomal protein S24e (Rps24) alternative variant aSep08, mRNA.
Rps24	Rps24.cSep08	81776	4635	551	7	131	ribosomal protein S24e (15.2 kD) (Rps24) alternative variant cSep08, complete mRNA.
Rps24	Rps24.dSep08	81776	4648	546	6	130	ribosomal protein S24e (15.1 kD) (Rps24) alternative variant dSep08, complete mRNA.
Rps24	Rps24.eSep08	81776	4590	797	6	118	ribosomal protein S24e (13.6 kD) (Rps24) alternative variant eSep08, mRNA.
Rps24	Rps24.fSep08	81776	990	667	2	37	putative protein (4.7 kD) (Rps24) alternative variant fSep08, mRNA.
Rps26	Rps26.bSep08	27139	1480	587	3	111	ribosomal protein S26E (12.5 kD) (Rps26) alternative variant bSep08, mRNA.
Rps26	Rps26.cSep08	27139	10663	444	5	81	putative protein (8.6 kD) (Rps26) alternative variant cSep08, mRNA.
Rps27	Rps27.bSep08	94266	1067	612	2	52	ribosomal protein S27E (5.6 kD) (Rps27) alternative variant bSep08, complete mRNA.
Rps27	Rps27.cSep08	94266	689	470	2	50	CRA b like (5.9 kD) (Rps27) alternative variant cSep08, mRNA.
Rps27a	Rps27a.bSep08	81777	2702	1515	5	156	protein ribosomal S27 (17.9 kD) (Rps27a) alternative variant bSep08, complete mRNA.
Rpsa	Rpsa.aSep08	29236	3712	1475	6	295	ribosomal protein S2 (32.8 kD) (Rpsa) alternative variant aSep08, mRNA.

Rpsa	Rpsa.cSep08	29236	3859	1015	7	294	ribosomal protein S2 (32.7 kD) (Rpsa) alternative variant cSep08, complete mRNA.
Rpsa	Rpsa.dSep08	29236	2662	1106	3	194	laminin Receptor (Rpsa) alternative variant dSep08, mRNA.
Rpsa	Rpsa.eSep08	29236	1629	751	2	114	putative protein of eukaryotic origin (12.4 kD) (Rpsa) alternative variant eSep08, mRNA.
Rpsa	Rpsa.fSep08	29236	3780	664	6	112	putative protein of eukaryotic origin (Rpsa) alternative variant fSep08, mRNA.
Rpusd1	Rpusd1.bSep08	287148	3843	1757	6	306	pseudouridine synthase (34.3 kD) (Rpusd1) alternative variant bSep08, complete mRNA.
Rpusd1	Rpusd1.cSep08	287148	2471	902	4	260	pseudouridine synthase (Rpusd1) alternative variant cSep08, mRNA.
Rpusd1	Rpusd1.dSep08	287148	1464	417	4	112	pseudouridine synthase (Rpusd1) alternative variant dSep08, mRNA.
Rpusd1	Rpusd1.fSep08	287148	962	374	3	3	putative protein (0.3 kD) (Rpusd1) alternative variant fSep08, mRNA.
Rpusd3	Rpusd3.aSep08	362416	3734	991	9	330	pseudouridine synthase (Rpusd3) alternative variant aSep08, mRNA.
Rpusd3	Rpusd3.cSep08	362416	1954	889	4	157	putative protein of ancient origin (17.5 kD) (Rpusd3) alternative variant cSep08, mRNA.
Rpusd3	Rpusd3.dSep08	362416	5328	745	2	117	CRA b like (Rpusd3) alternative variant dSep08, mRNA.
Rpusd3	Rpusd3.fSep08	362416	1479	599	3	81	CRA d (Rpusd3) alternative variant fSep08, mRNA.
Rqcd1	Rqcd1.bSep08	301513	4925	907	3	100	rcd1 (required for cell differentiation) homolog 1 (S. pombe) (Rqcd1) alternative variant bSep08, mRNA.
Rrad	Rrad.bSep08	83521	1770	872	1	215	ras-related associated with diabetes (Rrad) alternative variant bSep08, mRNA.
RragB	RragB.bSep08	117043	30244	794	4	87	ras-related GTP binding B (RragB) alternative variant bSep08, mRNA.
Rragc	Rragc.bSep08	298514	7152	1502	5	116	ras-related GTP binding C (Rragc) alternative variant bSep08, mRNA.
Rragc	Rragc.cSep08	298514	15809	736	6	107	ras-related GTP binding C (Rragc) alternative variant cSep08, mRNA.
Rrbp1	Rrbp1.aSep08	311483	39696	4127	23	1207	ribosome-binding protein 1 like (Rrbp1) alternative variant aSep08, mRNA.
Rrbp1	Rrbp1.bSep08	311483	13055	1300	14	347	ribosome binding protein 1 like (39.2 kD) (Rrbp1) alternative variant bSep08, mRNA.
Rrbp1	Rrbp1.cSep08	311483	2649	734	2	141	ribosome receptor (Rrbp1) alternative variant cSep08, mRNA.
Rrbp1	Rrbp1.dSep08	311483	1724	626	3	128	ribosome binding protein 1 like (Rrbp1) alternative variant dSep08, mRNA.
Rrm2andRGD1559671	Rrm2andRGD1559671.aSep08	304657	8830	2594	1	390	similar to M2 ribonucleotide reductase and ribonucleotide reductase M2 (Rrm2andRGD1559671) alternative variant aSep08, mRNA.
Rrm2andRGD1559671	Rrm2andRGD1559671.aSep08	362720	8830	2594	1	390	similar to M2 ribonucleotide reductase and ribonucleotide reductase M2 (Rrm2andRGD1559671) alternative variant aSep08, mRNA.

Rrm2andRGD1559671	Rrm2andRGD1559671.bSep08	304657	4117	940	1	285	similar to M2 ribonucleotide reductase and ribonucleotide reductase M2 (Rrm2andRGD1559671) alternative variant bSep08, mRNA.
Rrm2andRGD1559671	Rrm2andRGD1559671.bSep08	362720	4117	940	1	285	similar to M2 ribonucleotide reductase and ribonucleotide reductase M2 (Rrm2andRGD1559671) alternative variant bSep08, mRNA.
Rrm2b	Rrm2b.bSep08	299976	1155	287		83	ribonucleotide reductase M2 B (TP53 inducible) (Rrm2b) alternative variant bSep08, mRNA.
RRM_1.6	RRM_1.6.aSep08		3157	1010	2	114	bruno-like RNA binding protein (RRM_1.6) alternative variant aSep08, mRNA.
RRM_1.6	RRM_1.6.cSep08		2756	1785	2	98	bruno-like 5 RNA binding protein (10.9 kD) (RRM_1.6) alternative variant cSep08, mRNA.
RRM_1.9	RRM_1.9.aSep08		35282	655		218	ribonucleoprotein PTB-binding 2 CRA a like (RRM_1.9) mRNA.
RRM_1.12	RRM_1.12.aSep08		5894	526		175	RNA-binding region RNP-1 (RNA recognition motif) (RRM_1.12) mRNA.
Rrn3	Rrn3.aSep08	304714	35038	3423	13	661	RRN3 RNA polymerase I transcription factor homolog (yeast) (75.1 kD) (Rrn3) alternative variant aSep08, complete mRNA.
Rrn3	Rrn3.bSep08	304714	12797	784	1	260	RRN3 RNA polymerase I transcription factor homolog (yeast) (Rrn3) alternative variant bSep08, mRNA.
Rrp1	Rrp1.bSep08	309674	11145	1003	9	290	ribosomal RNA processing 1 homolog (S. cerevisiae) (32.9 kD) (Rrp1) alternative variant bSep08, complete mRNA.
Rrp1b	Rrp1b.aSep08	309673	25573	4420	16	749	ribosomal RNA processing 1 homolog B (S. cerevisiae) (Rrp1b) alternative variant aSep08, mRNA.
Rrp1b	Rrp1b.cSep08	309673	9211	404	2	82	ribosomal RNA processing 1 homolog B (S. cerevisiae) (9.4 kD) (Rrp1b) alternative variant cSep08, mRNA.
Rrp9	Rrp9.aSep08	363134	8990	1544	15	478	RRP9, small subunit (SSU) processome component, homolog (yeast) (52.5 kD) (Rrp9) alternative variant aSep08, complete mRNA.
Rrp9	Rrp9.bSep08	363134	4198	399	6	121	RRP9, small subunit (SSU) processome component, homolog (yeast) (Rrp9) alternative variant bSep08, mRNA.
Rrp9	Rrp9.cSep08	363134	905	734	3	109	RRP9, small subunit (SSU) processome component, homolog (yeast) (Rrp9) alternative variant cSep08, mRNA.
RR_TM4-6.0	RR_TM4-6.0.aSep08		2617	486		161	ryanodine receptor (RR_TM4-6.0) mRNA.
Rsb66	Rsb66.aSep08	296610	9518	826	5	254	rsb-66 protein (Rsb66) alternative variant aSep08, mRNA.
Rsb66	Rsb66.cSep08	296610	1017	463	2	130	rsb-66 protein (Rsb66) alternative variant cSep08, mRNA.
Rsb1	Rsb1.aSep08	310749	13788	642	2	214	rosbin, round spermatid basic protein 1 (Rsb1) alternative variant aSep08, mRNA.
Rsb1l	Rsb1l.aSep08	311987	61645	1735	7	578	round spermatid basic protein 1-like (Rsb1l) alternative variant aSep08, mRNA.
Rsb1l	Rsb1l.bSep08	311987	49966	1093	7	364	round spermatid basic protein 1-like (Rsb1l) alternative variant bSep08, mRNA.
Rsb1l	Rsb1l.cSep08	311987	829	750	1	77	round spermatid basic protein 1-like (Rsb1l) alternative variant cSep08, mRNA.
Rsf1	Rsf1.aSep08	308839	94028	2131	4	710	remodeling and spacing factor 1 (Rsf1) alternative variant aSep08, mRNA.

Rsf1	Rsf1.bSep08	308839	60244	391	1	99	remodeling and spacing factor 1 (Rsf1) alternative variant bSep08, mRNA.
Rshl2	Rshl2.aSep08	361476	39392	2027	7	571	radial spokehead-like 2 (64.6 kD) (Rshl2) alternative variant aSep08, mRNA.
Rshl2	Rshl2.cSep08	361476	4665	358	2	85	radial spokehead-like 2 (Rshl2) alternative variant cSep08, mRNA.
Rshl3	Rshl3.bSep08	309767	7936	1135	1	369	radial spokehead-like 3 (Rshl3) alternative variant bSep08, mRNA.
Rsl1d1	Rsl1d1.bSep08	302898	11887	1951	9	453	ribosomal protein L1 (50.9 kD) (Rsl1d1) alternative variant bSep08, complete mRNA.
Rsl1d1	Rsl1d1.cSep08	302898	10158	976	7	311	putative protein, with a coiled coil domain, of eukaryotic origin (Rsl1d1) alternative variant cSep08, mRNA.
Rspo1	Rspo1.bSep08	313589	21337	1834	1	262	R-spondin homolog (Xenopus laevis) (28.9 kD) (Rspo1) alternative variant bSep08, mRNA.
Rspo2	Rspo2.aSep08	500863	137831	3036	3	250	R-spondin 2 homolog (Xenopus laevis) and hypothetical protein LOC688674 (28.9 kD) (Rspo2) alternative variant aSep08, mRNA.
Rspo2	Rspo2.aSep08	688674	137831	3036	3	250	R-spondin 2 homolog (Xenopus laevis) and hypothetical protein LOC688674 (28.9 kD) (Rspo2) alternative variant aSep08, mRNA.
Rsrc1	Rsrc1.bSep08	361956	194801	2498	5	157	arginine/serine-rich coiled-coil 1 (Rsrc1) alternative variant bSep08, mRNA.
Rsrc1	Rsrc1.cSep08	361956	126505	686	5	133	arginine/serine-rich coiled-coil 1 (Rsrc1) alternative variant cSep08, mRNA.
Rsu1	Rsu1.bSep08	680419	103793	1081	2	243	ras suppressor protein 1 (27.5 kD) (Rsu1) alternative variant bSep08, complete mRNA.
RT1-149	RT1-149.bSep08	414784	7340	1494	6	334	class I precursor (38.9 kD) (RT1-149) alternative variant bSep08, complete mRNA.
RT1-149	RT1-149.cSep08	414784	6905	803	4	255	histocompatibility 2 T region locus 24 (RT1-149) alternative variant cSep08, mRNA.
RT1-149	RT1-149.dSep08	414784	6075	868	5	240	class I (RT1-149) alternative variant dSep08, mRNA.
RT1-149	RT1-149.eSep08	414784	41516	1323	8	238	MHC class I (26.5 kD) (RT1-149) alternative variant eSep08, complete mRNA.
RT1-149	RT1-149.fSep08	414784	8055	731	3	176	class I (20.1 kD) (RT1-149) alternative variant fSep08, mRNA.
RT1-149	RT1-149.gSep08	414784	7381	807	4	67	CRA a like (7.8 kD) (RT1-149) alternative variant gSep08, mRNA.
RT1-Ba	RT1-Ba.bSep08	309621	1208	782	2	223	RT1 class II, locus Ba (RT1-Ba) alternative variant bSep08, mRNA.
RT1-Ba	RT1-Ba.cSep08	309621	3736	619	3	130	RT1 class II, locus Ba (14.7 kD) (RT1-Ba) alternative variant cSep08, mRNA.
RT1-Bb	RT1-Bb.aSep08	309622	5089	951	6	295	RT1 class II, locus Bb (RT1-Bb) alternative variant aSep08, mRNA.
RT1-Bb	RT1-Bb.cSep08	309622	9155	1247	4	240	RT1 class II, locus Bb (27.5 kD) (RT1-Bb) alternative variant cSep08, mRNA.
RT1-Bb	RT1-Bb.dSep08	309622	2130	786	2	223	RT1 class II, locus Bb (24.3 kD) (RT1-Bb) alternative variant dSep08, mRNA.



RT1-Bb	RT1-Bb.eSep08	309622	3237	1387	3	181	RT1 class II, locus Bb (RT1-Bb) alternative variant eSep08, mRNA.
RT1-CE1	RT1-CE1.aSep08	309603	3360	1440	2	330	RT1 class I, CE1 (RT1-CE1) alternative variant aSep08, mRNA.
RT1-CE1	RT1-CE1.bSep08	309603	3351	1708	2	249	RT1 class I, CE1 (27.6 kD) (RT1-CE1) alternative variant bSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.cSep08	309607	69834	1437	7	327	class I (RT1-CE2andRT1-CE5) alternative variant cSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.cSep08	414779	69834	1437	7	327	class I (RT1-CE2andRT1-CE5) alternative variant cSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.dSep08	309607	787	584	2	116	class I precursor (13.4 kD) (RT1-CE2andRT1-CE5) alternative variant dSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.dSep08	414779	787	584	2	116	class I precursor (13.4 kD) (RT1-CE2andRT1-CE5) alternative variant dSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.eSep08	309607	1251	573	4	66	histocompatibility class I (RT1-CE2andRT1-CE5) alternative variant eSep08, mRNA.
RT1-CE2andRT1-CE5	RT1-CE2andRT1-CE5.eSep08	414779	1251	573	4	66	histocompatibility class I (RT1-CE2andRT1-CE5) alternative variant eSep08, mRNA.
RT1-CE3	RT1-CE3.bSep08	414793	2655	914	4	256	RT1 class I, CE3 (RT1-CE3) alternative variant bSep08, mRNA.
RT1-CE7	RT1-CE7.bSep08	368153	35606	1936	6	245	RT1 class I, CE7 (27.1 kD) (RT1-CE7) alternative variant bSep08, mRNA.
RT1-CE7	RT1-CE7.dSep08	368153	1254	709	3	49	RT1 class I, CE7 (RT1-CE7) alternative variant dSep08, mRNA.
RT1-CE7	RT1-CE7.eSep08	368153	1039	546	3	37	RT1 class I, CE7 (RT1-CE7) alternative variant eSep08, mRNA.
RT1-CE10	RT1-CE10.bSep08	414792	4088	1919	5	328	RT1 class I, CE10 (RT1-CE10) alternative variant bSep08, mRNA.
RT1-CE10	RT1-CE10.cSep08	414792	916	713	2	116	RT1 class I, CE10 (13.5 kD) (RT1-CE10) alternative variant cSep08, mRNA.
RT1-CE11	RT1-CE11.bSep08	414791	963	761	1	119	RT1 class I, CE11 (RT1-CE11) alternative variant bSep08, mRNA.
RT1-CE12	RT1-CE12.bSep08	309600	2587	906	4	226	RT1 class I, CE12 (RT1-CE12) alternative variant bSep08, mRNA.
RT1-CE12	RT1-CE12.cSep08	309600	563	436	1	75	RT1 class I, CE12 (RT1-CE12) alternative variant cSep08, mRNA.
RT1-CE13andRT1-CE14	RT1-CE13andRT1-CE14.cSep08	414270	1732	755	2	171	class I (19.2 kD) (RT1-CE13andRT1-CE14) alternative variant cSep08, mRNA.

RT1-CE13andRT1-CE14	RT1-CE13andRT1-CE14.cSep08	414790	1732	755	2	171	class I (19.2 kD) (RT1-CE13andRT1-CE14) alternative variant cSep08, mRNA.
RT1-CE15	RT1-CE15.bSep08	414789	1708	1159	1	41	RT1 class I, CE15 (RT1-CE15) alternative variant bSep08, mRNA.
RT1-CE16andRT1-CI	RT1-CE16andRT1-CI.bSep08	24977	3073	1277	3	255	class I (RT1-CE16andRT1-CI) alternative variant bSep08, mRNA.
RT1-CE16andRT1-CI	RT1-CE16andRT1-CI.bSep08	414819	3073	1277	3	255	class I (RT1-CE16andRT1-CI) alternative variant bSep08, mRNA.
RT1-CE16andRT1-CI	RT1-CE16andRT1-CI.cSep08	24977	1772	805	1	170	class I (RT1-CE16andRT1-CI) alternative variant cSep08, mRNA.
RT1-CE16andRT1-CI	RT1-CE16andRT1-CI.cSep08	414819	1772	805	1	170	class I (RT1-CE16andRT1-CI) alternative variant cSep08, mRNA.
RT1-Db1	RT1-Db1.bSep08	294270	9559	1865	5	223	RT1 class II, locus Db1 (26.0 kD) (RT1-Db1) alternative variant bSep08, complete mRNA.
RT1-Db1	RT1-Db1.cSep08	294270	4694	917	2	166	RT1 class II, locus Db1 (RT1-Db1) alternative variant cSep08, mRNA.
RT1-Db1	RT1-Db1.eSep08	294270	4426	596	2	61	RT1 class II, locus Db1 (RT1-Db1) alternative variant eSep08, mRNA.
RT1-Db1	RT1-Db1.hSep08	294270	1827	278	2	31	RT1 class II, locus Db1 (RT1-Db1) alternative variant hSep08, mRNA.
RT1-DOa	RT1-DOa.bSep08	24984	3347	756	2	120	RT1 class II, locus DOa (RT1-DOa) alternative variant bSep08, mRNA.
RT1-DOb	RT1-DOb.aSep08	365542	5206	696	4	231	major Histocompatibility complex class II DO beta (RT1-DOb) alternative variant aSep08, mRNA.
RT1-DOb	RT1-DOb.bSep08	365542	1559	632	2	196	major Histocompatibility complex class II DO beta (RT1-DOb) alternative variant bSep08, mRNA.
RT1-DOb	RT1-DOb.cSep08	365542	2895	873	2	52	histocompatibility 2 O region beta locus like (RT1-DOb) alternative variant cSep08, mRNA.
RT1-Ha	RT1-Ha.bSep08	24986	1077	776	1	198	RT1 class II, locus Ha (RT1-Ha) alternative variant bSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.bSep08	294281	1663	940	6	312	solute carrier family 39 member 7 (RT1-Ke4andHsd17b8) alternative variant bSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.bSep08	361802	1663	940	6	312	solute carrier family 39 member 7 (RT1-Ke4andHsd17b8) alternative variant bSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.dSep08	294281	1971	895	8	239	hydroxysteroid dehydrogenase 8 CRA a (RT1-Ke4andHsd17b8) alternative variant dSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.dSep08	361802	1971	895	8	239	hydroxysteroid dehydrogenase 8 CRA a (RT1-Ke4andHsd17b8) alternative variant dSep08, mRNA.

RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.eSep08	294281	2250	2140	2	162	solute carrier family 39 member 7 (17.0 kD) (RT1-Ke4andHsd17b8) alternative variant eSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.eSep08	361802	2250	2140	2	162	solute carrier family 39 member 7 (17.0 kD) (RT1-Ke4andHsd17b8) alternative variant eSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.fSep08	294281	596	386	3	128	solute carrier family 39 member 7 (RT1-Ke4andHsd17b8) alternative variant fSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.fSep08	361802	596	386	3	128	solute carrier family 39 member 7 (RT1-Ke4andHsd17b8) alternative variant fSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.gSep08	294281	834	518	3	68	hydroxysteroid dehydrogenase 8 CRA b (RT1-Ke4andHsd17b8) alternative variant gSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.gSep08	361802	834	518	3	68	hydroxysteroid dehydrogenase 8 CRA b (RT1-Ke4andHsd17b8) alternative variant gSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.hSep08	294281	846	316	3	46	hydroxysteroid dehydrogenase 8 CRA a (RT1-Ke4andHsd17b8) alternative variant hSep08, mRNA.
RT1-Ke4andHsd17b8	RT1-Ke4andHsd17b8.hSep08	361802	846	316	3	46	hydroxysteroid dehydrogenase 8 CRA a (RT1-Ke4andHsd17b8) alternative variant hSep08, mRNA.
RT1-M3	RT1-M3.bSep08	24747	600	439	2	72	RT1 class Ib, locus M3 (8.4 kD) (RT1-M3) alternative variant bSep08, mRNA.
RT1-M3	RT1-M3.cSep08	24747	129293	733	3	15	RT1 class Ib, locus M3 (1.5 kD) (RT1-M3) alternative variant cSep08, mRNA.
RT1-M5	RT1-M5.bSep08	499400	1923	982	1	326	RT1 class Ib, locus M5 (RT1-M5) alternative variant bSep08, mRNA.
RT1-M6-1	RT1-M6-1.bSep08	414785	1820	657	2	218	RT1 class I, M6, gene 1 (RT1-M6-1) alternative variant bSep08, mRNA.
RT1-M6-1	RT1-M6-1.cSep08	414785	723	541	1	179	RT1 class I, M6, gene 1 (RT1-M6-1) alternative variant cSep08, mRNA.
RT1-M6-2	RT1-M6-2.bSep08	365527	1834	922	1	241	RT1 class I, M6, gene 2 (RT1-M6-2) alternative variant bSep08, mRNA.
RT1-M10-2andPpid-ps	RT1-M10-2andPpid-ps.aSep08	414807	17446	570		34	RT1 class 1, M10, pseudogene 2 and RT1 class 1, Ppid pseudogene (3.7 kD) (RT1-M10-2andPpid-ps) mRNA.
RT1-M10-2andPpid-ps	RT1-M10-2andPpid-ps.aSep08	414810	17446	570		34	RT1 class 1, M10, pseudogene 2 and RT1 class 1, Ppid pseudogene (3.7 kD) (RT1-M10-2andPpid-ps) mRNA.
RT1-N1	RT1-N1.bSep08	24748	2586	1661	3	128	RT1 class Ib gene, H2-TL-like, grc region (N1) (14.2 kD) (RT1-N1) alternative variant bSep08, mRNA.
RT1-N1	RT1-N1.cSep08	24748	1806	784	3	89	RT1 class Ib gene, H2-TL-like, grc region (N1) (RT1-N1) alternative variant cSep08, mRNA.
RT1-N1	RT1-N1.dSep08	24748	1154	269	3	61	RT1 class Ib gene, H2-TL-like, grc region (N1) (RT1-N1) alternative variant dSep08, mRNA.

RT1-N3	RT1-N3.bSep08	24750	2274	1862	3	218	RT1 class Ib gene, H2-TL-like, grc region (N3) (25.5 kD) (RT1-N3) alternative variant bSep08, mRNA.
RT1-N3	RT1-N3.cSep08	24750	802	619	2	206	RT1 class Ib gene, H2-TL-like, grc region (N3) (RT1-N3) alternative variant cSep08, mRNA.
RT1-O	RT1-O.bSep08	24751	558	273	2	47	RT1 class Ib, locus H2-Q-like, grc region (RT1-O) alternative variant bSep08, mRNA.
RT1-O	RT1-O.cSep08	24751	3754	1050	4	42	RT1 class Ib, locus H2-Q-like, grc region (RT1-O) alternative variant cSep08, mRNA.
RT1-S3	RT1-S3.bSep08	294228	3119	1532	6	324	RT1 class Ib, locus S3 (35.6 kD) (RT1-S3) alternative variant bSep08, mRNA.
RT1-S3	RT1-S3.cSep08	294228	2338	826	6	221	RT1 class Ib, locus S3 (RT1-S3) alternative variant cSep08, mRNA.
Rtcd1	Rtcd1.bSep08	295395	15947	895	9	192	RNA terminal phosphate cyclase domain 1 (Rtcd1) alternative variant bSep08, mRNA.
Rtcd1	Rtcd1.cSep08	295395	9395	689	5	138	RNA terminal phosphate cyclase domain 1 (Rtcd1) alternative variant cSep08, mRNA.
Rtel1	Rtel1.aSep08	362288	5521	2166	13	589	DEAH helicase (Rtel1) alternative variant aSep08, mRNA.
Rtel1	Rtel1.bSep08	362288	8081	4059	15	466	DEAH helicase (51.3 kD) (Rtel1) alternative variant bSep08, mRNA.
Rtel1	Rtel1.cSep08	362288	22600	892	11	256	DEAH helicase (Rtel1) alternative variant cSep08, mRNA.
Rtel1	Rtel1.dSep08	362288	2794	753	6	250	DEAH helicase (Rtel1) alternative variant dSep08, mRNA.
Rtel1	Rtel1.eSep08	362288	2382	764	4	233	DEAH helicase (Rtel1) alternative variant eSep08, mRNA.
Rtel1	Rtel1.fSep08	362288	12957	631	6	162	DEAH helicase (Rtel1) alternative variant fSep08, mRNA.
Rtel1	Rtel1.gSep08	362288	35351	863	7	142	regulator of telomere elongation helicase 1 like (16.3 kD) (Rtel1) alternative variant gSep08, mRNA.
Rtel1	Rtel1.hSep08	362288	6959	605	5	123	DEAH helicase (13.6 kD) (Rtel1) alternative variant hSep08, mRNA.
Rtel1	Rtel1.iSep08	362288	8789	473	3	116	DEAH helicase (Rtel1) alternative variant iSep08, mRNA.
Rtel1	Rtel1.jSep08	362288	921	698	2	102	putative protein (Rtel1) alternative variant jSep08, mRNA.
Rtel1	Rtel1.kSep08	362288	447	350	2	63	putative protein (Rtel1) alternative variant kSep08, mRNA.
Rtel1	Rtel1.lSep08	362288	4028	402	2	47	DEAH helicase (Rtel1) alternative variant lSep08, mRNA.
Rtf1	Rtf1.aSep08	366169	51031	1457	4	485	rtf1, Paf1/RNA polymerase II complex component, homolog ( <i>S. cerevisiae</i> ) (Rtf1) alternative variant aSep08, mRNA.
Rtf1	Rtf1.cSep08	366169	24694	2618	3	233	rtf1, Paf1/RNA polymerase II complex component, homolog ( <i>S. cerevisiae</i> ) (Rtf1) alternative variant cSep08, mRNA.
Rtkn	Rtkn.bSep08	297383	7395	1059	7	353	rhotekin (Rtkn) alternative variant bSep08, mRNA.
Rtkn	Rtkn.cSep08	297383	7187	618	6	153	rhotekin (Rtkn) alternative variant cSep08, mRNA.
Rtkn	Rtkn.dSep08	297383	10357	527	5	131	rhotekin (Rtkn) alternative variant dSep08, mRNA.
Rtkn	Rtkn.eSep08	297383	11612	393	5	130	rhotekin (Rtkn) alternative variant eSep08, mRNA.
Rtkn	Rtkn.fSep08	297383	7960	586	2	81	rhotekin (8.5 kD) (Rtkn) alternative variant fSep08, mRNA.
Rtn1	Rtn1.bSep08	116644	29254	1530	7	255	reticulin 1 (Rtn1) alternative variant bSep08, mRNA.
Rtn2	Rtn2.bSep08	308410	5391	1029	7	216	reticulin 2 (Z-band associated protein) (Rtn2) alternative variant bSep08, mRNA.
Rtn3	Rtn3.bSep08	140945	25977	1616	3	500	reticulin 3 (Rtn3) alternative variant bSep08, mRNA.
Rtn3	Rtn3.cSep08	140945	52699	720	7	239	reticulin 3 (Rtn3) alternative variant cSep08, mRNA.

Rtn3	Rtn3.eSep08	140945	15095	1446	7	236	reticulon 3 (26.3 kD) (Rtn3) alternative variant eSep08, mRNA.
Rtn4	Rtn4.bSep08	83765	47475	2238	7	437	reticulon 4 (Rtn4) alternative variant bSep08, mRNA.
Rtn4	Rtn4.cSep08	83765	34350	851	7	204	reticulon 4 (Rtn4) alternative variant cSep08, mRNA.
Rtn4	Rtn4.dSep08	83765	21825	1415	7	199	reticulon 4 (22.4 kD) (Rtn4) alternative variant dSep08, mRNA.
Rtn4	Rtn4.eSep08	83765	6142	565	5	155	reticulon 4 (Rtn4) alternative variant eSep08, mRNA.
Rtn4ip1	Rtn4ip1.bSep08	309912	8084	444	1	147	reticulon 4 interacting protein 1 (Rtn4ip1) alternative variant bSep08, mRNA.
Rtn4rl2	Rtn4rl2.bSep08	311169	5946	618	2	206	reticulon 4 receptor-like 2 (Rtn4rl2) alternative variant bSep08, mRNA.
Rttn	Rttn.aSep08	291377	48362	1394		464	rotatin (Rttn) mRNA.
ruby	ruby.aSep08		9803	2545	2	86	putative nuclear protein (10.2 kD) (ruby) alternative variant aSep08, mRNA.
ruchy	ruchy.aSep08		18368	603		200	CRA b (ruchy) mRNA.
rudar	rudar.aSep08		1080	522		108	putative protein (12.0 kD) (rudar) mRNA.
rufer	rufer.aSep08		561	267		41	putative protein (rufer) mRNA.
ruflo	ruflo.aSep08		500	135		45	putative protein (ruflo) mRNA.
ruflo	ruflo.bSep08		26859	311	2	37	putative protein (ruflo) alternative variant bSep08, mRNA.
Rufy1	Rufy1.aSep08	360521	21680	1691	11	392	RUN FYVE domain-containing 1 (Rufy1) alternative variant aSep08, mRNA.
Rufy3	Rufy3.aSep08	360921	75917	3284	18	634	RUN and zinc finger, FYVE-type (Rufy3) alternative variant aSep08, mRNA.
Rufy3	Rufy3.bSep08	360921	57644	4353	19	487	RUN (Rufy3) alternative variant bSep08, mRNA.
Rufy3	Rufy3.dSep08	360921	13856	878	8	179	rap2 interacting protein x CRA e (20.3 kD) (Rufy3) alternative variant dSep08, mRNA.
Rufy3	Rufy3.eSep08	360921	28864	735	4	156	putative protein of metazoan origin (Rufy3) alternative variant eSep08, mRNA.
Rufy3	Rufy3.fSep08	360921	24841	684	2	100	putative protein of bilateral origin (Rufy3) alternative variant fSep08, mRNA.
Rufy3	Rufy3.gSep08	360921	3252	597	3	92	rap2 interacting protein x CRA c (Rufy3) alternative variant gSep08, mRNA.
rujar	rujar.aSep08		45623	893	7	297	CRA d (rujar) alternative variant aSep08, mRNA.
ruja	ruja.aSep08		14179	595		31	putative protein (ruja) mRNA.
rujey	rujey.aSep08		49870	456		45	putative protein (5.0 kD) (rujey) mRNA.
rukee	rukee.aSep08		25934	548		60	putative protein (7.1 kD) (rukee) mRNA.
rukler	rukler.aSep08		45226	520	1	113	putative protein (rukler) alternative variant aSep08, mRNA.
rukler	rukler.bSep08		19275	629	1	97	putative protein (rukler) alternative variant bSep08, mRNA.
rulo	rulo.aSep08		812	475	1	59	CRA a like (6.8 kD) (rulo) alternative variant aSep08, mRNA.
rulo	rulo.bSep08		7616	486	2	59	CRA a like (6.8 kD) (rulo) alternative variant bSep08, mRNA.
rumee	rumee.aSep08		7959	247		70	putative protein (8.1 kD) (rumee) mRNA.
rumer	rumer.aSep08		2521	373		124	CRA a like (rumer) mRNA.

RUN.0	RUN.0.aSep08		19188	694	7	210	RUN (RUN.0) alternative variant aSep08, mRNA.
RUN.1	RUN.1.aSep08		4718	587		187	RUN (RUN.1) mRNA.
RUN.2	RUN.2.aSep08		5852	1862	9	450	RUN and src homology-3 and variant SH3 (RUN.2) alternative variant aSep08, mRNA.
RUN.2	RUN.2.bSep08		6276	2495	8	380	RUN (40.6 kD) (RUN.2) alternative variant bSep08, mRNA.
RUN.2	RUN.2.cSep08		1150	927	3	168	putative protein of bilateral origin (RUN.2) alternative variant cSep08, mRNA.
RUN.2	RUN.2.eSep08		880	420	5	139	putative protein of bilateral origin (RUN.2) alternative variant eSep08, mRNA.
Rundc1	Rundc1.aSep08	303552	3495	2382	2	328	RUN (Rundc1) alternative variant aSep08, mRNA.
runoy	runoy.aSep08		933	427		142	putative protein, with a coiled coil domain (runoy) mRNA.
Runx1t1	Runx1t1.bSep08	362489	53513	296	2	43	runt-related transcription factor 1; translocated to, 1 (cyclin D-related) (Runx1t1) alternative variant bSep08, mRNA.
Rup2	Rup2.aSep08	619560	3565	413	3	74	urinary protein 2 (Rup2) alternative variant aSep08, mRNA.
Rup2	Rup2.bSep08	619560	1458	804	2	41	urinary protein 2 (Rup2) alternative variant bSep08, mRNA.
rupor	rupor.aSep08		1962	741	4	132	putative protein (rupor) alternative variant aSep08, mRNA.
rusa	rusa.aSep08		55500	559	3	80	putative protein of mammalian origin (rusa) alternative variant aSep08, mRNA.
rusa	rusa.bSep08		5786	489	2	32	putative protein (rusa) alternative variant bSep08, mRNA.
rushee	rushee.aSep08		37935	994		258	mediator of rna polymerase ii transcription (rushee) mRNA.
rutu	rutu.aSep08		28951	588		59	putative protein (rutu) mRNA.
ruvar	ruvar.aSep08		507	233		77	putative protein (ruvar) mRNA.
Ruvbl2	Ruvbl2.bSep08	292907	9664	770	8	224	RuvB-like protein 2 (Ruvbl2) alternative variant bSep08, mRNA.
Ruvbl2	Ruvbl2.cSep08	292907	2951	985	5	189	RuvB-like protein 2 (Ruvbl2) alternative variant cSep08, mRNA.
Ruvbl2	Ruvbl2.dSep08	292907	2753	386	4	128	RuvB-like protein 2 (Ruvbl2) alternative variant dSep08, mRNA.
Ruvbl2	Ruvbl2.eSep08	292907	3648	382	3	117	RuvB-like protein 2 (Ruvbl2) alternative variant eSep08, mRNA.
Ruvbl2	Ruvbl2.fSep08	292907	4404	233	3	70	RuvB-like protein 2 (Ruvbl2) alternative variant fSep08, mRNA.
ruwey	ruwey.aSep08		4923	892	2	297	retinoblastoma binding protein 2 like (ruwey) alternative variant aSep08, mRNA.
ruwey	ruwey.bSep08		2686	714	1	205	histone demethylase jarid1a (ruwey) alternative variant bSep08, mRNA.
rve.0	rve.0.aSep08		942	799		115	integrase like (12.8 kD) (rve.0) mRNA.
rve.1	rve.1.aSep08		3808	1425		105	ac1576 like (11.6 kD) (rve.1) mRNA.
rve.5	rve.5.aSep08		942	799		115	integrase like (12.8 kD) (rve.5) mRNA.
RWD.0	RWD.0.aSep08		15076	343		101	eukaryotic translation initiation factor 2 alpha kinase 4 CRA a (RWD.0) mRNA.
Rwdd2a	Rwdd2a.aSep08	363110	2789	1016		292	RWD (33.8 kD) (Rwdd2a) mRNA.
Rwdd3	Rwdd3.bSep08	65026	12720	685	2	212	RWD (Rwdd3) alternative variant bSep08, mRNA.
Rwdd4a	Rwdd4a.bSep08	502084	8669	401	3	44	putative protein of vertebrate origin (5.1 kD) (Rwdd4a) alternative variant bSep08, complete mRNA.

Rwdd4a	Rwdd4a.cSep08	502084	3828	1390	2	60	putative protein (6.3 kD) (Rwdd4a) alternative variant cSep08, mRNA.
Rxfp2	Rxfp2.bSep08	363866	3640	808	2	92	relaxin/insulin-like family peptide receptor 2 (10.3 kD) (Rxfp2) alternative variant bSep08, mRNA.
Rxra	Rxra.bSep08	25271	7653	729	2	130	retinoid X receptor alpha (Rxra) alternative variant bSep08, mRNA.
Rxrb	Rxrb.bSep08	361801	4040	1102	8	309	retinoid X receptor beta (Rxrb) alternative variant bSep08, mRNA.
Rxrb	Rxrb.cSep08	361801	5554	3044	8	239	retinoid X receptor beta (26.5 kD) (Rxrb) alternative variant cSep08, mRNA.
Rxrb	Rxrb.dSep08	361801	2159	1103	4	229	retinoid X receptor beta (25.6 kD) (Rxrb) alternative variant dSep08, mRNA.
Rxrb	Rxrb.eSep08	361801	688	515	2	144	retinoid X receptor beta (Rxrb) alternative variant eSep08, mRNA.
Rxrb	Rxrb.fSep08	361801	541	426	2	109	retinoid X receptor beta (Rxrb) alternative variant fSep08, mRNA.
Rxrg	Rxrg.bSep08	83574	16223	1436	2	292	retinoid X receptor gamma (33.0 kD) (Rxrg) alternative variant bSep08, mRNA.
RY2G5	RY2G5.aSep08	499925	5172	533		49	potential ligand-binding protein (5.3 kD) (RY2G5) mRNA.
ryby	ryby.aSep08		8408	4389		175	putative protein (19.5 kD) (ryby) mRNA.
rychy	rychy.aSep08		14949	294		61	putative protein (rychy) mRNA.
rydar	rydar.aSep08		26092	257		33	putative protein (rydar) mRNA.
RYDR_ITPR.0	RYDR_ITPR.0.aSep08		4505	429		142	ryanodine receptor (RYDR_ITPR.0) mRNA.
ryfer	ryfer.bSep08		5560	413		74	putative protein (ryfer) alternative variant bSep08, mRNA.
ryflo	ryflo.aSep08		17267	688		120	putative protein (ryflo) mRNA.
ryflu	ryflu.aSep08		11961	559	4	48	CRA c like (6.0 kD) (ryflu) alternative variant aSep08, mRNA.
ryflu	ryflu.bSep08		14588	1179	5	167	CRA b like (ryflu) alternative variant bSep08, mRNA.
rygar	rygar.aSep08		3314	694		230	CRA b (rygar) mRNA.
ryja	ryja.aSep08		156450	309		103	putative protein (ryja) mRNA.
ryjey	ryjey.aSep08		20783	1575		524	shroom (ryjey) mRNA.
Ryk	Ryk.bSep08	140585	32434	2145	2	390	receptor-like tyrosine kinase (Ryk) alternative variant bSep08, mRNA.
rykee	rykee.aSep08		37571	3854		92	putative protein (9.4 kD) (rykee) alternative variant aSep08, mRNA.
rykee	rykee.bSep08		34265	620	1	83	CRA b like (rykee) alternative variant bSep08, mRNA.
rykler	rykler.aSep08		7205	1843		123	putative protein of mammalian origin (rykler) mRNA.
rylo	rylo.aSep08		3547	402		41	LYST-interacting protein 8 like (rylo) mRNA.
rymee	rymee.aSep08		7796	267		34	putative protein (4.2 kD) (rymee) mRNA.
rymer	rymer.aSep08		1490	330		109	putative protein of mammalian origin (rymer) mRNA.
rynoy	rynoy.aSep08		360	261		66	putative protein of vertebrate origin (rynoy) mRNA.
rypor	rypor.aSep08		8864	341		113	talin 2 (rypor) mRNA.
RyR.0	RyR.0.aSep08		1051	352		117	ryanodine receptor (RyR.0) mRNA.
Ryr1	Ryr1.aSep08	114207	14057	1453		445	ryanodine receptor 1, skeletal muscle (Ryr1) mRNA.

Ryr2	Ryr2.bSep08	689560	108463	2232	15	666	ryanodine receptor 2, cardiac (Ryr2) alternative variant bSep08, mRNA.
Ryr2	Ryr2.cSep08	689560	11958	1794	6	597	ryanodine receptor 2, cardiac (Ryr2) alternative variant cSep08, mRNA.
Ryr2	Ryr2.dSep08	689560	13139	701	7	208	ryanodine receptor 2, cardiac (Ryr2) alternative variant dSep08, mRNA.
Ryr2	Ryr2.eSep08	689560	1848	246	2	81	ryanodine receptor 2, cardiac (Ryr2) alternative variant eSep08, mRNA.
Ryr3	Ryr3.aSep08	170546	5546	700		233	ryanodine receptor 3 (Ryr3) mRNA.
rysa	rysa.aSep08		1805	414		138	mical-like 1 (rysa) mRNA.
ryshee	ryshee.aSep08		4133	591		196	mediator of rna polymerase ii transcription (ryshee) mRNA.
rytu	rytu.aSep08		6196	244		39	putative protein (rytu) mRNA.
ryvar	ryvar.aSep08		6949	557		62	putative protein (ryvar) mRNA.
rywey	rywey.aSep08		5272	1448		157	putative protein, with 3 coiled coil domains, of eukaryotic origin (rywey) mRNA.
S100a3	S100a3.aSep08	114216	22027	578	1	141	s100 calcium binding protein A3 (S100a3) alternative variant aSep08, mRNA.
S100a3	S100a3.bSep08	114216	18535	690	1	124	s100 calcium binding protein A3 (S100a3) alternative variant bSep08, mRNA.
S100a3	S100a3.dSep08	114216	40520	1290	2	77	s100 calcium binding protein A3 (8.5 kD) (S100a3) alternative variant dSep08, mRNA.
S100a6	S100a6.aSep08	85247	1456	479	3	89	s100 calcium binding protein A6 (calcyclin) (10.0 kD) (S100a6) alternative variant aSep08, mRNA.
S100a6	S100a6.bSep08	85247	1236	945	2	89	s100 calcium binding protein A6 (calcyclin) (10.0 kD) (S100a6) alternative variant bSep08, complete mRNA.
S100a10	S100a10.bSep08	81778	8499	345	3	77	s100 calcium binding protein A10 (calpactin) (S100a10) alternative variant bSep08, mRNA.
S100a13	S100a13.aSep08	295213	6496	484	3	98	s100 calcium binding protein A13 (11.2 kD) (S100a13) alternative variant aSep08, mRNA.
S100a16	S100a16.aSep08	361991	1504	643	1	123	s100 calcium binding protein A16 (14.2 kD) (S100a16) alternative variant aSep08, mRNA.
S100a16	S100a16.bSep08	361991	3356	958	1	123	s100 calcium binding protein A16 (14.2 kD) (S100a16) alternative variant bSep08, mRNA.
S100b	S100b.aSep08	25742	7299	1988	2	135	s100 protein, beta polypeptide, neural (S100b) alternative variant aSep08, mRNA.
S100vp	S100vp.aSep08	295176	2946	555	3	125	s100 calcium-binding protein, ventral prostate (S100vp) alternative variant aSep08, mRNA.
S100vp	S100vp.cSep08	295176	639	332	2	34	s100 calcium-binding protein, ventral prostate (S100vp) alternative variant cSep08, mRNA.
S100vp	S100vp.dSep08	295176	522	272	2	48	s100 calcium-binding protein, ventral prostate (S100vp) alternative variant dSep08, mRNA.
S100vp	S100vp.eSep08	295176	531	272	3	48	s100 calcium-binding protein, ventral prostate (S100vp) alternative variant eSep08, mRNA.
SAA.0	SAA.0.aSep08		4651	1761	3	339	serum amyloid (SAA.0) alternative variant aSep08, mRNA.
SAA.0	SAA.0.bSep08		3440	551	2	139	serum amyloid (SAA.0) alternative variant bSep08, mRNA.



SAA.0	SAA.0.cSep08		3085	692	1	130	serum amyloid precursor (14.9 kD) (SAA.0) alternative variant cSep08, mRNA.
SAA.1	SAA.1.aSep08		2500	395		87	serum amyloid A (SAA.1) mRNA.
Saa4	Saa4.aSep08	365245	3684	1121	1	130	serum amyloid A 4 (15.0 kD) (Saa4) alternative variant aSep08, mRNA.
saby	saby.aSep08		9187	2916		323	alpha thalassemia mental retardation syndrome x-linked homolog like (saby) mRNA.
SAC3_GANP.0	SAC3_GANP.0.aSep08		1597	561		187	SAC3/GANP/Nin1/mts3/eIF-3 p25 (SAC3_GANP.0) mRNA.
sachy	sachy.aSep08		4892	177		34	putative protein (sachy) mRNA.
Sacm1l	Sacm1l.aSep08	116482	57093	3457	5	622	SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae) (Sacm1l) alternative variant aSep08, mRNA.
Sacm1l	Sacm1l.bSep08	116482	37721	762		156	SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae) (Sacm1l) alternative variant bSep08, mRNA.
Sacm1l	Sacm1l.cSep08	116482	4779	722	2	146	SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae) (Sacm1l) alternative variant cSep08, mRNA.
Sacm1l	Sacm1l.dSep08	116482	7239	397	1	131	SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae) (Sacm1l) alternative variant dSep08, mRNA.
sadar	sadar.aSep08		1550	738		37	putative protein (4.1 kD) (sadar) mRNA.
Sae1	Sae1.bSep08	308384	30838	799	1	136	SUMO1 activating enzyme subunit 1 (15.1 kD) (Sae1) alternative variant bSep08, mRNA.
Safb	Safb.bSep08	64196	2144	824	5	218	scaffold attachment factor B (Safb) alternative variant bSep08, mRNA.
Safb	Safb.cSep08	64196	2069	673	6	197	scaffold attachment factor B (Safb) alternative variant cSep08, mRNA.
safer	safer.aSep08		6063	388		128	gtpase-activating protein 21 (safer) mRNA.
saflo	saflo.aSep08		17350	762		93	putative nuclear protein (10.2 kD) (saflo) mRNA.
saflu	saflu.aSep08		2496	382		113	putative protein (saflu) mRNA.
Sag	Sag.bSep08	25539	817	252	1	38	retinal S-antigen (Sag) alternative variant bSep08, mRNA.
Sag	Sag.cSep08	25539	10038	711	3	34	retinal S-antigen (Sag) alternative variant cSep08, mRNA.
sagar	sagar.aSep08		3019	699		61	putative protein (sagar) mRNA.
saja	saja.aSep08		11950	625		138	CRA a (saja) mRNA.
sajey	sajey.aSep08		25142	497	2	52	putative protein (sajey) alternative variant aSep08, mRNA.
sajey	sajey.bSep08		79136	474	2	56	putative protein (6.3 kD) (sajey) alternative variant bSep08, mRNA.
sakee	sakee.aSep08		705	309		44	putative protein (sakee) mRNA.
sakler	sakler.aSep08		955	590		196	plasma membrane associated protein s3-12 (sakler) mRNA.
Sall1	Sall1.aSep08	307740	5042	3928	2	1242	sal-like 1 (Drosophila) (Sall1) alternative variant aSep08, mRNA.
Sall1	Sall1.cSep08	307740	2889	1775	3	462	sal-like 1 (Drosophila) (Sall1) alternative variant cSep08, mRNA.
Sall1	Sall1.dSep08	307740	14574	700	2	214	sal-like 1 (Drosophila) (24.5 kD) (Sall1) alternative variant dSep08, mRNA.
salo	salo.aSep08		3102	374		124	furry homolog CRA b (salo) mRNA.

Samd4	Samd4.aSep08	305826	104256	2410	12	610	sterile alpha motif homology 2 and sterile alpha motif SAM (67.1 kD) (Samd4) alternative variant aSep08, complete mRNA.
Samd4	Samd4.cSep08	305826	90576	1782	5	406	sterile alpha motif homology 2 and sterile alpha motif SAM (Samd4) alternative variant cSep08, mRNA.
Samd4	Samd4.dSep08	305826	57220	752	5	250	sterile alpha motif homology 2 and sterile alpha motif SAM (Samd4) alternative variant dSep08, mRNA.
Samd7	Samd7.aSep08	310257	8088	1000		43	putative protein (Samd7) mRNA.
Samd10	Samd10.aSep08	499957	4256	2142	5	238	sterile alpha motif homology 2 and sterile alpha motif SAM (Samd10) alternative variant aSep08, mRNA.
Samd10	Samd10.bSep08	499957	2840	759	5	174	sterile alpha motif homology 2 and sterile alpha motif SAM (19.6 kD) (Samd10) alternative variant bSep08, mRNA.
Samd10	Samd10.cSep08	499957	2996	931	5	157	sterile alpha motif homology 2 and sterile alpha motif SAM (17.8 kD) (Samd10) alternative variant cSep08, complete mRNA.
Samd10	Samd10.dSep08	499957	1756	419	2	139	putative protein of vertebrate origin (Samd10) alternative variant dSep08, mRNA.
Samd12	Samd12.aSep08	362910	517273	874	4	161	sterile alpha motif/pointed and sterile alpha motif homology 2 and sterile alpha motif SAM (18.2 kD) (Samd12) alternative variant aSep08, mRNA.
Samd12	Samd12.cSep08	362910	15314	449	1	36	putative protein (4.0 kD) (Samd12) alternative variant cSep08, mRNA.
Samd14	Samd14.bSep08	287637	9868	788	4	257	putative protein of vertebrate origin (Samd14) alternative variant bSep08, mRNA.
Samd14	Samd14.cSep08	287637	8888	945	1	74	putative protein of mammalian origin (8.2 kD) (Samd14) alternative variant cSep08, mRNA.
samee	samee.aSep08		5377	269		41	putative protein (4.7 kD) (samee) mRNA.
samer	samer.aSep08		1957	1030		154	apoptosis-associated tyrosine kinase (17.3 kD) (samer) mRNA.
Samhd1	Samhd1.aSep08	311580	13539	1659		346	SAM domain and HD domain, 1 (Samhd1) alternative variant aSep08, mRNA.
Samhd1	Samhd1.bSep08	311580	10618	936		311	SAM domain and HD domain, 1 (Samhd1) alternative variant bSep08, mRNA.
Samm50	Samm50.bSep08	300111	4744	1289	3	189	sorting and assembly machinery component 50 homolog ( <i>S. cerevisiae</i> ) (20.8 kD) (Samm50) alternative variant bSep08, mRNA.
Samm50	Samm50.cSep08	300111	2542	728	2	109	sorting and assembly machinery component 50 homolog ( <i>S. cerevisiae</i> ) (12.8 kD) (Samm50) alternative variant cSep08, mRNA.
SAM_1.0	SAM_1.0.aSep08		4751	635		172	sterile alpha motif homology 2 and sterile alpha motif SAM (SAM_1.0) mRNA.
SAM_1.1	SAM_1.1.aSep08		79187	393		58	cajalín 2 (SAM_1.1) mRNA.
SAM_1.2	SAM_1.2.aSep08		6571	376		125	kazrin (SAM_1.2) mRNA.
SAM_1.3	SAM_1.3.aSep08		22011	3403	13	583	ptprf interacting protein binding 1 like (SAM_1.3) alternative variant aSep08, mRNA.
SAM_1.3	SAM_1.3.bSep08		969	508	1	116	ptprf interacting protein binding 1 like (SAM_1.3) alternative variant bSep08, mRNA.

SAM_1.4	SAM_1.4.aSep08		2265	316		105	protein tyrosine phosphatase receptor type f polypeptide interacting alpha (SAM_1.4) mRNA.
SAM_2.0	SAM_2.0.bSep08		3054	2158		61	CRA b like (SAM_2.0) alternative variant bSep08, mRNA.
SAM_2.1	SAM_2.1.aSep08		1463	1190		178	sterile alpha motif homology 2 and sterile alpha motif SAM (18.9 kD) (SAM_2.1) mRNA.
SAM_2.2	SAM_2.2.aSep08		1764	827	5	205	eph receptor B6 (SAM_2.2) alternative variant aSep08, mRNA.
SAM_2.2	SAM_2.2.bSep08		740	385	3	63	eph receptor B6 (SAM_2.2) alternative variant bSep08, mRNA.
sanoy	sanoy.aSep08		1038	571		74	putative protein (sanoy) mRNA.
SAP.0	SAP.0.aSep08		20821	487	5	161	apoptotic chromatin condensation inducer 1 CRA a (SAP.0) alternative variant aSep08, mRNA.
SAP.1	SAP.1.aSep08		3830	1245		415	megakaryoblastic leukemia 1 like (SAP.1) mRNA.
Sap18	Sap18.bSep08	290284	810	476	2	136	sin3-associated polypeptide 18 (Sap18) alternative variant bSep08, mRNA.
Sap18	Sap18.cSep08	290284	900	772	2	80	sin3-associated polypeptide 18 (9.2 kD) (Sap18) alternative variant cSep08, mRNA.
Sap30bp	Sap30bp.aSep08	360662	32676	3196	11	308	SAP30 binding protein (33.8 kD) (Sap30bp) alternative variant aSep08, complete mRNA.
Sap30bp	Sap30bp.bSep08	360662	23551	395	5	131	SAP30 binding protein (Sap30bp) alternative variant bSep08, mRNA.
Sap30bp	Sap30bp.cSep08	360662	873	613	2	104	SAP30 binding protein (10.6 kD) (Sap30bp) alternative variant cSep08, mRNA.
Sap30l	Sap30l.aSep08	360531	7656	862		150	SAP30-like (Sap30l) mRNA.
Sap130	Sap130.aSep08	307527	44720	1138	6	339	Sin3A associated protein (Sap130) alternative variant aSep08, mRNA.
Sap130	Sap130.bSep08	307527	18233	673	4	224	Sin3A associated protein (Sap130) alternative variant bSep08, mRNA.
sapor	sapor.aSep08		3644	453		86	putative protein (sapor) mRNA.
Saps1	Saps1.aSep08	361502	12326	3129	22	772	SAPS domain family, member 1 (Saps1) alternative variant aSep08, mRNA.
Saps1	Saps1.bSep08	361502	5631	731	8	237	SAPS domain family, member 1 (Saps1) alternative variant bSep08, mRNA.
Saps1	Saps1.cSep08	361502	2169	557	5	143	SAPS domain family, member 1 (Saps1) alternative variant cSep08, mRNA.
Saps1	Saps1.dSep08	361502	1529	384	5	128	SAPS domain family, member 1 (Saps1) alternative variant dSep08, mRNA.
Saps1	Saps1.eSep08	361502	1078	381	2	62	SAPS domain family, member 1 (7.0 kD) (Saps1) alternative variant eSep08, mRNA.
Saps2	Saps2.aSep08	300146	33813	3665	21	940	saps domain family member 2 (Saps2) alternative variant aSep08, mRNA.
Saps2	Saps2.cSep08	300146	1745	740	3	114	saps domain family member 2 (Saps2) alternative variant cSep08, mRNA.
Saps2	Saps2.eSep08	300146	1196	394	3	78	putative mitochondrial protein (8.6 kD) (Saps2) alternative variant eSep08, mRNA.

Saps2	Saps2.fSep08	300146	676	431	2	72	saps domain family member 2 (Saps2) alternative variant fSep08, mRNA.
Saps3	Saps3.aSep08	309144	82856	1798	4	556	SAPS domain family, member 3 (Saps3) alternative variant aSep08, mRNA.
Saps3	Saps3.bSep08	309144	19364	955	7	317	SAPS domain family, member 3 (Saps3) alternative variant bSep08, mRNA.
Saps3	Saps3.cSep08	309144	10864	2519	2	150	SAPS domain family, member 3 (Saps3) alternative variant cSep08, mRNA.
Sar1a	Sar1a.aSep08	361842	10649	934	4	198	SAR1 gene homolog A (S. cerevisiae) (22.4 kD) (Sar1a) alternative variant aSep08, mRNA.
Sar1a	Sar1a.cSep08	361842	10459	708	5	196	SAR1 gene homolog A (S. cerevisiae) (Sar1a) alternative variant cSep08, mRNA.
Sar1a	Sar1a.dSep08	361842	10727	910	4	186	SAR1 gene homolog A (S. cerevisiae) (21.1 kD) (Sar1a) alternative variant dSep08, mRNA.
Sar1a	Sar1a.eSep08	361842	5870	671	2	116	SAR1 gene homolog A (S. cerevisiae) (Sar1a) alternative variant eSep08, mRNA.
Sar1b	Sar1b.bSep08	287276	29615	1171	4	155	SAR1 gene homolog B (S. cerevisiae) (17.6 kD) (Sar1b) alternative variant bSep08, complete mRNA.
Sar1b	Sar1b.cSep08	287276	25076	569	2	92	SAR1 gene homolog B (S. cerevisiae) (10.2 kD) (Sar1b) alternative variant cSep08, mRNA.
sarby	sarby.aSep08		1194	418		27	putative protein (3.3 kD) (sarby) mRNA.
sarchy	sarchy.aSep08		3293	405		134	kinesin family member 18A (sarchy) mRNA.
sardar	sardar.aSep08		2355	849			
Sardh	Sardh.bSep08	114123	24758	573	4	190	sarcosine dehydrogenase (Sardh) alternative variant bSep08, mRNA.
Sardh	Sardh.cSep08	114123	7791	1517	1	101	sarcosine dehydrogenase (Sardh) alternative variant cSep08, mRNA.
sarflo	sarflo.aSep08		837	349		62	putative protein (sarflo) mRNA.
sarflu	sarflu.aSep08		7293	983	11	327	anemia group I (sarflu) alternative variant aSep08, mRNA.
sarflu	sarflu.bSep08		4720	941	5	141	anemia group I (sarflu) alternative variant bSep08, mRNA.
sargar	sargar.aSep08		1551	598		11	putative protein (1.2 kD) (sargar) mRNA.
sarja	sarja.aSep08		11312	1525		32	putative protein (3.7 kD) (sarja) mRNA.
sarjey	sarjey.aSep08		25038	699	2	43	putative protein (sarjey) alternative variant aSep08, mRNA.
sarjey	sarjey.bSep08		19162	318	1	44	putative protein (sarjey) alternative variant bSep08, mRNA.
sarkee	sarkee.aSep08		3515	634		211	putative protein of vertebrate origin (sarkee) mRNA.
sarlo	sarlo.aSep08		5846	740		246	furry homolog CRA a (sarlo) mRNA.
sarmee	sarmee.aSep08		2506	415		32	putative protein (3.2 kD) (sarmee) mRNA.
sarmer	sarmer.aSep08		1731	567		189	fanconi anemia core complex subunit (sarmer) mRNA.
sarnoy	sarnoy.aSep08		732	269		87	putative protein (sarnoy) mRNA.
sarpor	sarpor.bSep08		3585	461	2	54	putative protein of mammalian origin (6.4 kD) (sarpor) alternative variant bSep08, mRNA.
Sars	Sars.bSep08	266975	3940	981	5	207	seryl-aminoacyl-tRNA synthetase (Sars) alternative variant bSep08, mRNA.
Sars	Sars.cSep08	266975	875	636	2	122	seryl-aminoacyl-tRNA synthetase (Sars) alternative variant cSep08, mRNA.

sarsa	sarsa.aSep08		1742	394		69	putative protein (sarsa) mRNA.
sarshee	sarshee.aSep08		1839	787		101	sucrase-isomaltase (sarshee) mRNA.
Sart1	Sart1.bSep08	29678	3837	1351	2	358	squamous cell carcinoma antigen recognized by T-cells 1 (Sart1) alternative variant bSep08, mRNA.
sartu	sartu.aSep08		16464	546		109	putative protein (sartu) mRNA.
sarvar	sarvar.aSep08		10947	468		62	putative protein (sarvar) mRNA.
sarwey	sarwey.aSep08		3954	1214	2	49	putative protein (5.2 kD) (sarwey) alternative variant aSep08, mRNA.
sarwey	sarwey.bSep08		7310	2041	3	49	putative protein (5.2 kD) (sarwey) alternative variant bSep08, mRNA.
sasa	sasa.aSep08		1594	737	3	156	polyprotein -pol (sasa) alternative variant aSep08, mRNA.
sasa	sasa.bSep08		3284	282	2	63	gag protein like (sasa) alternative variant bSep08, mRNA.
Sash3	Sash3.aSep08	317578	14720	2567	8	380	src homology-3 and variant SH3 and sterile alpha motif homology 2 and sterile alpha motif SAM (41.7 kD) (Sash3) alternative variant aSep08, mRNA.
Sash3	Sash3.bSep08	317578	1520	872	3	198	domain-containing protein 3 (Sash3) alternative variant bSep08, mRNA.
sashee	sashee.aSep08		1184	755		92	putative protein (10.1 kD) (sashee) mRNA.
Sass6	Sass6.bSep08	310807	9190	918	7	240	spindle assembly 6 (Sass6) alternative variant bSep08, mRNA.
Sass6	Sass6.cSep08	310807	6312	611	7	186	spindle assembly 6 (Sass6) alternative variant cSep08, mRNA.
Sass6	Sass6.dSep08	310807	6482	508	3	102	spindle assembly 6 (Sass6) alternative variant dSep08, mRNA.
Sat1	Sat1.bSep08	302642	3148	2457	1	114	putative protein of ancient origin (13.2 kD) (Sat1) alternative variant bSep08, complete mRNA.
Sat1	Sat1.cSep08	302642	3157	1145	3	71	spermidine spermine transferase CRA a (8.2 kD) (Sat1) alternative variant cSep08, complete mRNA.
Sat1	Sat1.dSep08	302642	3157	1058	2	96	spermidine spermine N1-acetyltransferase (11.4 kD) (Sat1) alternative variant dSep08, complete mRNA.
Sat2	Sat2.aSep08	360547	1601	775	5	192	spermidine/spermine N1-acetyl transferase 2 (Sat2) alternative variant aSep08, mRNA.
Sat2	Sat2.bSep08	360547	1586	566	6	185	spermidine/spermine N1-acetyl transferase 2 (Sat2) alternative variant bSep08, mRNA.
Sat2	Sat2.dSep08	360547	1486	562	4	105	spermidine/spermine N1-acetyl transferase 2 (12.0 kD) (Sat2) alternative variant dSep08, mRNA.
Sat2	Sat2.eSep08	360547	849	431	3	98	spermidine/spermine N1-acetyl transferase 2 (Sat2) alternative variant eSep08, mRNA.
Satb1	Satb1.bSep08	316164	39280	1797	5	500	special AT-rich sequence binding protein 1 like (Satb1) alternative variant bSep08, mRNA.
Satb1	Satb1.cSep08	316164	4021	2071	3	271	CRA a (Satb1) alternative variant cSep08, mRNA.
Satb1	Satb1.dSep08	316164	2781	735	2	194	special AT-rich sequence binding protein 1 CRA b like (Satb1) alternative variant dSep08, mRNA.
Satb1	Satb1.fSep08	316164	21375	789	3	41	putative protein (4.6 kD) (Satb1) alternative variant fSep08, mRNA.

satu	satu.aSep08		3800	2004	8	417	prolactin regulatory element binding like (45.4 kD) (satu) alternative variant aSep08, mRNA.
satu	satu.bSep08		2599	1757	5	270	prolactin regulatory element binding like (satu) alternative variant bSep08, mRNA.
satu	satu.cSep08		1621	1072	3	167	prolactin regulatory element binding CRA c like (satu) alternative variant cSep08, mRNA.
Sav1	Sav1.bSep08	299116	2225	406	2	50	salvador homolog 1 (Drosophila) (Sav1) alternative variant bSep08, mRNA.
savar	savar.aSep08		3192	734	2	96	putative protein (savar) alternative variant aSep08, mRNA.
savar	savar.bSep08		12166	821	4	60	putative protein (7.2 kD) (savar) alternative variant bSep08, mRNA.
sawby	sawby.aSep08		9515	295		73	putative protein of eukaryotic origin (sawby) mRNA.
sawchy	sawchy.aSep08		20042	337		28	putative protein (3.2 kD) (sawchy) mRNA.
sawdar	sawdar.bSep08		556	261		76	AT motif binding factor 1 like (sawdar) alternative variant bSep08, mRNA.
sawey	sawey.aSep08		45229	723		37	putative protein (sawey) mRNA.
sawflo	sawflo.aSep08		11613	750		77	putative mitochondrial protein (8.4 kD) (sawflo) mRNA.
sawflu	sawflu.aSep08		16044	815		74	putative protein (sawflu) mRNA.
sawgar	sawgar.bSep08		6002	557	2	41	CRA b like (4.6 kD) (sawgar) alternative variant bSep08, mRNA.
sawja	sawja.aSep08		737	567		80	putative protein (9.2 kD) (sawja) mRNA.
sawjey	sawjey.aSep08		15769	256		39	putative protein (sawjey) mRNA.
sawkee	sawkee.aSep08		3969	571		98	putative protein (sawkee) mRNA.
sawkler	sawkler.aSep08		31586	644		60	putative protein (sawkler) mRNA.
sawlo	sawlo.aSep08		4825	353		117	furry homolog (sawlo) mRNA.
sawmee	sawmee.aSep08		2952	738		54	putative protein (5.8 kD) (sawmee) mRNA.
sawmer	sawmer.aSep08		1674	663		78	putative protein of mammalian origin (sawmer) mRNA.
sawnoy	sawnoy.aSep08		6282	214		49	putative protein (sawnoy) mRNA.
sawpor	sawpor.aSep08		24006	451		36	putative protein (4.1 kD) (sawpor) mRNA.
sawsa	sawsa.aSep08		2224	284		34	putative protein (sawsa) mRNA.
sawshee	sawshee.aSep08		2072	447		41	putative protein (sawshee) mRNA.
sawtu	sawtu.cSep08		3371	288	3	52	putative protein (sawtu) alternative variant cSep08, mRNA.
sawvar	sawvar.aSep08		2088	805		151	CRA a like (sawvar) mRNA.
sawwey	sawwey.aSep08		2199	511		45	putative protein (sawwey) mRNA.
Sbds	Sbds.bSep08	288615	8635	735	5	210	shwachman-Bodian-Diamond syndrome homolog (human) (Sbds) alternative variant bSep08, mRNA.
Sbds	Sbds.cSep08	288615	4639	1198	3	197	shwachman-Bodian-Diamond syndrome homolog (human) (21.6 kD) (Sbds) alternative variant cSep08, complete mRNA.
Sbf1	Sbf1.aSep08	300147	7673	1994	12	548	SET binding factor 1 like (Sbf1) alternative variant aSep08, mRNA.
Sbf1	Sbf1.bSep08	300147	4851	807	3	257	CRA b (Sbf1) alternative variant bSep08, mRNA.
Sbf1	Sbf1.dSep08	300147	1328	712	5	93	SET binding factor 1 like (10.7 kD) (Sbf1) alternative variant dSep08, mRNA.

Sbk1	Sbk1.bSep08	113907	1621	1032	1	200	SH3-binding kinase 1 (Sbk1) alternative variant bSep08, mRNA.
Sbno1	Sbno1.bSep08	304470	7647	2191	7	273	sno, strawberry notch homolog 1 (Drosophila) (Sbno1) alternative variant bSep08, mRNA.
Sbno1	Sbno1.cSep08	304470	18249	733	5	195	sno, strawberry notch homolog 1 (Drosophila) (Sbno1) alternative variant cSep08, mRNA.
Sbno2	Sbno2.cSep08	314619	1719	543	6	132	strawberry notch homolog (Sbno2) alternative variant cSep08, mRNA.
Sbno2	Sbno2.dSep08	314619	2389	1004	2	64	putative protein (Sbno2) alternative variant dSep08, mRNA.
Sbno2	Sbno2.eSep08	314619	17542	360	3	61	putative protein (Sbno2) alternative variant eSep08, mRNA.
Sbp	Sbp.bSep08	25540	3862	1173	3	246	spermine binding protein (Sbp) alternative variant bSep08, mRNA.
Sbsn	Sbsn.aSep08	292793	3599	1324	4	392	suprabasin (Sbsn) alternative variant aSep08, mRNA.
Sbsn	Sbsn.dSep08	292793	4453	429	4	92	suprabasin (9.5 kD) (Sbsn) alternative variant dSep08, complete mRNA.
Sbsn	Sbsn.fSep08	292793	22302	556	6	54	suprabasin (Sbsn) alternative variant fSep08, mRNA.
Sc65	Sc65.bSep08	59101	3680	735	2	162	synaptonemal complex protein SC65 (Sc65) alternative variant bSep08, mRNA.
SCA7.0	SCA7.0.aSep08		6103	400		133	ataxin 7 (SCA7.0) mRNA.
SCA7.1	SCA7.1.aSep08		9385	699		179	ataxin 7-like (SCA7.1) mRNA.
Scaf1	Scaf1.bSep08	56081	1197	692	3	99	SR-related CTD-associated factor 1 (Scaf1) alternative variant bSep08, mRNA.
Scamp2	Scamp2.bSep08	65168	1839	1623	2	124	secretory carrier membrane protein 2 (Scamp2) alternative variant bSep08, mRNA.
Scamp2	Scamp2.dSep08	65168	5093	317	4	83	secretory carrier membrane protein 2 (Scamp2) alternative variant dSep08, mRNA.
Scamp3	Scamp3.aSep08	65169	5858	1432	1	413	secretory carrier membrane protein 3 (Scamp3) alternative variant aSep08, mRNA.
Scamp3	Scamp3.bSep08	65169	5790	1485	2	386	secretory carrier membrane protein 3 (Scamp3) alternative variant bSep08, mRNA.
Scamp4	Scamp4.aSep08	65170	12158	1825	7	288	secretory carrier membrane protein 4 (Scamp4) alternative variant aSep08, mRNA.
Scamp4	Scamp4.cSep08	65170	2776	922	4	131	secretory carrier membrane protein 4 (Scamp4) alternative variant cSep08, mRNA.
Scamp4	Scamp4.dSep08	65170	1824	654	2	131	secretory carrier membrane protein 4 (Scamp4) alternative variant dSep08, mRNA.
Scamp4	Scamp4.eSep08	65170	11385	686	7	102	secretory carrier membrane protein 4 (11.4 kD) (Scamp4) alternative variant eSep08, mRNA.
Scamp4	Scamp4.gSep08	65170	7122	760	3	59	secretory carrier membrane protein 4 (6.8 kD) (Scamp4) alternative variant gSep08, mRNA.
Scamp4	Scamp4.hSep08	65170	9512	590	6	97	secretory carrier membrane protein 4 (Scamp4) alternative variant hSep08, mRNA.
SCAN.0	SCAN.0.aSep08		11987	951		246	zinc finger protein 445 (SCAN.0) mRNA.
SCAN.1	SCAN.1.aSep08		1656	762		147	zinc finger (SCAN.1) mRNA.

SCAN.3	SCAN.3.aSep08		1302	441		61	putative protein (SCAN.3) mRNA.
Scap	Scap.bSep08	301024	44129	1074	6	246	SREBF chaperone (27.8 kD) (Scap) alternative variant bSep08, mRNA.
Scap	Scap.cSep08	301024	475	341	2	113	SREBF chaperone (Scap) alternative variant cSep08, mRNA.
Scara5	Scara5.aSep08	305974	77944	1052		350	scavenger receptor class A, member 5 (putative) (Scara5) mRNA.
Scarb1	Scarb1.bSep08	25073	43770	646	4	147	scavenger receptor class B, member 1 (Scarb1) alternative variant bSep08, mRNA.
Scarb1	Scarb1.cSep08	25073	42390	758	3	145	scavenger receptor class B, member 1 (16.5 kD) (Scarb1) alternative variant cSep08, mRNA.
Scarb1	Scarb1.dSep08	25073	9992	775	4	112	scavenger receptor class B, member 1 (Scarb1) alternative variant dSep08, mRNA.
Scarb1	Scarb1.fSep08	25073	1220	577	2	99	scavenger receptor class B, member 1 (Scarb1) alternative variant fSep08, mRNA.
Scarf1	Scarf1.bSep08	303313	5153	1168	5	281	scavenger receptor class F, member 1 (Scarf1) alternative variant bSep08, mRNA.
Scarf1	Scarf1.cSep08	303313	6526	3490	6	183	scavenger receptor class F, member 1 (Scarf1) alternative variant cSep08, mRNA.
Scarf1	Scarf1.dSep08	303313	774	690	2	95	scavenger receptor class F, member 1 (Scarf1) alternative variant dSep08, mRNA.
Scarf1	Scarf1.eSep08	303313	3161	2228	3	59	scavenger receptor class F, member 1 (6.3 kD) (Scarf1) alternative variant eSep08, mRNA.
Scarf2	Scarf2.cSep08	287949	3512	1377	2	81	scavenger receptor class F, member 2 (8.0 kD) (Scarf2) alternative variant cSep08, mRNA.
Sccpdh	Sccpdh.aSep08	305021	19456	1791	4	544	saccharopine dehydrogenase (putative) (Sccpdh) alternative variant aSep08, mRNA.
Sccpdh	Sccpdh.cSep08	305021	11959	825	6	181	saccharopine dehydrogenase (putative) (Sccpdh) alternative variant cSep08, mRNA.
Scd1	Scd1.bSep08	246074	1242	518	2	47	stearoyl-Coenzyme A desaturase 1 (Scd1) alternative variant bSep08, mRNA.
Scel	Scel.bSep08	361086	54616	1090	16	299	sciellin (Scel) alternative variant bSep08, mRNA.
Scel	Scel.cSep08	361086	35881	1376	8	163	sciellin (Scel) alternative variant cSep08, mRNA.
Scfd1	Scfd1.aSep08	54350	78973	2086	25	637	vesicle transport-related protein (72.3 kD) (Scfd1) alternative variant aSep08, mRNA.
Scg2	Scg2.bSep08	24765	5574	2435	3	579	secretogranin II (66.7 kD) (Scg2) alternative variant bSep08, mRNA.
Scg2	Scg2.cSep08	24765	3345	393	3	85	secretogranin II (Scg2) alternative variant cSep08, mRNA.
Scg2	Scg2.dSep08	24765	3269	383	2	76	secretogranin II (Scg2) alternative variant dSep08, mRNA.
Scg3	Scg3.aSep08	116635	16296	1779	1	578	secretogranin III (Scg3) alternative variant aSep08, mRNA.
Scg3	Scg3.cSep08	116635	8535	673	2	66	secretogranin III (Scg3) alternative variant cSep08, mRNA.
Scg5	Scg5.bSep08	25719	43609	755	2	197	secretogranin V (Scg5) alternative variant bSep08, mRNA.
Scgb2a1andScgb2a2	Scgb2a1andScgb2a2.aSep08	25010	135609	744	4	130	secretoglobin family 2A member 1 (Scgb2a1andScgb2a2) alternative variant aSep08, mRNA.
Scgb2a1andScgb2a2	Scgb2a1andScgb2a2.aSep08	361725	135609	744	4	130	secretoglobin family 2A member 1 (Scgb2a1andScgb2a2) alternative variant aSep08, mRNA.



Schip1	Schip1.bSep08	295105	1031	826	2	274	schwannomin interacting protein 1 (Schip1) alternative variant bSep08, mRNA.
Schip1	Schip1.cSep08	295105	52367	1594	7	261	schwannomin interacting protein 1 (29.4 kD) (Schip1) alternative variant cSep08, mRNA.
Schip1	Schip1.dSep08	295105	63116	1030	7	244	schwannomin interacting protein 1 (27.5 kD) (Schip1) alternative variant dSep08, mRNA.
Schip1	Schip1.eSep08	295105	26952	728	2	242	schwannomin interacting protein 1 (Schip1) alternative variant eSep08, mRNA.
Scly	Scly.bSep08	363285	11289	638	5	212	selenocysteine lyase (Scly) alternative variant bSep08, mRNA.
Scly	Scly.cSep08	363285	1602	796	2	137	selenocysteine lyase (14.8 kD) (Scly) alternative variant cSep08, mRNA.
Scly	Scly.dSep08	363285	9744	407	4	118	selenocysteine lyase (Scly) alternative variant dSep08, mRNA.
Scly	Scly.eSep08	363285	1142	377	2	58	selenocysteine lyase (Scly) alternative variant eSep08, mRNA.
Scmh1	Scmh1.bSep08	362581	12859	707	1	181	sex comb on midleg homolog 1 (Scmh1) alternative variant bSep08, mRNA.
Scml4	Scml4.aSep08	309859	15914	955		241	sex comb on midleg-like 4 (Drosophila) (Scml4) mRNA.
Scn1a	Scn1a.bSep08	81574	31004	1269	8	423	sodium channel, voltage-gated, type I, alpha (Scn1a) alternative variant bSep08, mRNA.
Scn1a	Scn1a.cSep08	81574	6205	3314	2	398	sodium channel, voltage-gated, type I, alpha (Scn1a) alternative variant cSep08, mRNA.
Scn1a	Scn1a.dSep08	81574	70005	341	1	52	sodium channel, voltage-gated, type I, alpha (Scn1a) alternative variant dSep08, mRNA.
Scn1b	Scn1b.bSep08	29686	9722	1487	5	260	sodium channel, voltage-gated, type I, beta (Scn1b) alternative variant bSep08, mRNA.
Scn2a1	Scn2a1.bSep08	24766	21574	1792	7	551	sodium channel, voltage-gated, type II, alpha 1 (Scn2a1) alternative variant bSep08, mRNA.
Scn2a1	Scn2a1.cSep08	24766	10891	513	4	98	sodium channel, voltage-gated, type II, alpha 1 (Scn2a1) alternative variant cSep08, mRNA.
Scn2a1	Scn2a1.dSep08	24766	10891	504	4	98	sodium channel, voltage-gated, type II, alpha 1 (Scn2a1) alternative variant dSep08, mRNA.
Scn2b	Scn2b.bSep08	25349	9281	890	3	177	sodium channel, voltage-gated, type II, beta (20.2 kD) (Scn2b) alternative variant bSep08, mRNA.
Scn3b	Scn3b.cSep08	245956	3169	2859	2	83	sodium channel, voltage-gated, type III, beta (Scn3b) alternative variant cSep08, mRNA.
Scn3b	Scn3b.dSep08	245956	8686	660	4	44	sodium channel, voltage-gated, type III, beta (Scn3b) alternative variant dSep08, mRNA.
Scn5a	Scn5a.bSep08	25665	80192	6196	22	1381	sodium channel, voltage-gated, type 5, alpha subunit (Scn5a) alternative variant bSep08, mRNA.
Scn5a	Scn5a.cSep08	25665	80192	6037	21	1328	sodium channel, voltage-gated, type 5, alpha subunit (Scn5a) alternative variant cSep08, mRNA.
Scn5a	Scn5a.dSep08	25665	6787	718	5	238	sodium channel, voltage-gated, type 5, alpha subunit (Scn5a) alternative variant dSep08, mRNA.
Scn7a	Scn7a.aSep08	64155	6105	2904		446	sodium channel, voltage-gated, type VII, alpha (Scn7a) mRNA.

Scn8a	Scn8a.cSep08	29710	4638	505	2	42	sodium channel, voltage-gated, type 8, alpha subunit (Scn8a) alternative variant cSep08, mRNA.
Scn8a	Scn8a.dSep08	29710	1228	525	2	68	sodium channel, voltage-gated, type 8, alpha subunit (7.5 kD) (Scn8a) alternative variant dSep08, mRNA.
Scn9a	Scn9a.bSep08	78956	19199	565	1	112	sodium channel, voltage-gated, type IX, alpha (Scn9a) alternative variant bSep08, mRNA.
Scn10a	Scn10a.aSep08	29571	8187	1024		341	sodium channel, voltage-gated, type X, alpha (Scn10a) mRNA.
Scnm1	Scnm1.bSep08	310662	3045	645	1	214	sodium channel modifier 1 (Scnm1) alternative variant bSep08, mRNA.
Scnn1a	Scnn1a.bSep08	25122	4920	767	2	235	sodium channel, nonvoltage-gated, type I, alpha (Scnn1a) alternative variant bSep08, mRNA.
Scnn1b	Scnn1b.aSep08	24767	3521	742	6	247	sodium channel, nonvoltage-gated 1 beta (Scnn1b) alternative variant aSep08, mRNA.
Scnn1b	Scnn1b.bSep08	24767	845	761	2	40	sodium channel, nonvoltage-gated 1 beta (4.6 kD) (Scnn1b) alternative variant bSep08, mRNA.
Sco1	Sco1.aSep08	497930	13153	2149		284	SCO cytochrome oxidase deficient homolog 1 (yeast) (31.8 kD) (Sco1) mRNA.
Scoc	Scoc.bSep08	364981	6275	1719	4	82	short coiled-coil protein (9.4 kD) (Scoc) alternative variant bSep08, complete mRNA.
Scoc	Scoc.eSep08	364981	2956	603	3	32	short coiled-coil protein (4.0 kD) (Scoc) alternative variant eSep08, mRNA.
SCP.0	SCP.0.aSep08		22065	833		206	cysteine-rich secretory protein 2 (SCP.0) mRNA.
Scp2	Scp2.bSep08	25541	33576	1104	10	184	sterol carrier protein 2 (20.4 kD) (Scp2) alternative variant bSep08, mRNA.
Scp2	Scp2.cSep08	25541	27083	755	4	154	sterol carrier protein 2 (Scp2) alternative variant cSep08, mRNA.
Scp2	Scp2.dSep08	25541	8574	390	3	60	sterol carrier protein 2 (Scp2) alternative variant dSep08, mRNA.
Scpep1	Scpep1.bSep08	114861	18187	728	9	240	serine carboxypeptidase 1 CRA b (Scpep1) alternative variant bSep08, mRNA.
Scpep1	Scpep1.cSep08	114861	13904	733	6	202	serine carboxypeptidase 1 CRA b (Scpep1) alternative variant cSep08, mRNA.
Scpep1	Scpep1.dSep08	114861	7692	617	3	102	serine carboxypeptidase 1 CRA b (Scpep1) alternative variant dSep08, mRNA.
Scpep1	Scpep1.eSep08	114861	1263	315	2	66	serine carboxypeptidase 1 (Scpep1) alternative variant eSep08, mRNA.
Scpep1	Scpep1.fSep08	114861	5894	993	2	40	serine carboxypeptidase 1 (4.4 kD) (Scpep1) alternative variant fSep08, mRNA.
Scrib	Scrib.aSep08	362938	16729	3525	21	715	scribbled homolog (Drosophila) (Scrib) alternative variant aSep08, mRNA.
Scrib	Scrib.bSep08	362938	7098	1787	5	595	scribbled homolog (Drosophila) (Scrib) alternative variant bSep08, mRNA.
Scrib	Scrib.cSep08	362938	10946	1879	11	548	scribbled homolog (Drosophila) (Scrib) alternative variant cSep08, mRNA.
Scrib	Scrib.dSep08	362938	10657	1980	13	493	scribbled homolog (Drosophila) (52.7 kD) (Scrib) alternative variant dSep08, mRNA.

Scrib	Scrib.eSep08	362938	10218	2000	10	351	scribbled homolog (Drosophila) (Scrib) alternative variant eSep08, mRNA.
Scrib	Scrib.fSep08	362938	3780	1379	8	238	scribbled homolog (Drosophila) (Scrib) alternative variant fSep08, mRNA.
Scrib	Scrib.gSep08	362938	891	626	4	208	scribbled homolog (Drosophila) (Scrib) alternative variant gSep08, mRNA.
Scrib	Scrib.hSep08	362938	3247	617	4	205	scribbled homolog (Drosophila) (Scrib) alternative variant hSep08, mRNA.
Scrib	Scrib.iSep08	362938	2333	764	4	159	scribbled homolog (Drosophila) (Scrib) alternative variant iSep08, mRNA.
Scrib	Scrib.jSep08	362938	1007	542	4	102	scribbled homolog (Drosophila) (Scrib) alternative variant jSep08, mRNA.
Scrib	Scrib.kSep08	362938	2100	596	2	90	scribbled homolog (Drosophila) (Scrib) alternative variant kSep08, mRNA.
Scrib	Scrib.lSep08	362938	836	417	3	60	scribbled homolog (Drosophila) (Scrib) alternative variant lSep08, mRNA.
Scrn1	Scrn1.bSep08	502776	4743	1791	2	131	secernin 1 (Scrn1) alternative variant bSep08, mRNA.
Scrn2	Scrn2.bSep08	360612	2722	753	1	229	secernin 2 (Scrn2) alternative variant bSep08, mRNA.
Scrn3	Scrn3.bSep08	311731	18963	2536	3	256	secernin 3 (Scrn3) alternative variant bSep08, mRNA.
Scube1	Scube1.bSep08	315174	5608	853	4	254	signal peptide, CUB domain, EGF-like 1 (Scube1) alternative variant bSep08, mRNA.
Scube1	Scube1.cSep08	315174	46922	702	4	197	signal peptide, CUB domain, EGF-like 1 (21.5 kD) (Scube1) alternative variant cSep08, complete mRNA.
Scx	Scx.cSep08	680712	1908	616	3	91	scleraxis (Scx) alternative variant cSep08, mRNA.
Scye1	Scye1.aSep08	114632	23824	1225	6	317	small inducible cytokine subfamily E, member 1 (Scye1) alternative variant aSep08, mRNA.
Scye1	Scye1.cSep08	114632	10961	438	2	36	small inducible cytokine subfamily E, member 1 (Scye1) alternative variant cSep08, mRNA.
Scyl1	Scyl1.bSep08	293684	2596	2026	5	175	CRA b (19.6 kD) (Scyl1) alternative variant bSep08, mRNA.
Scyl1	Scyl1.cSep08	293684	805	445	3	121	1 CRA a (Scyl1) alternative variant cSep08, mRNA.
Scyl1	Scyl1.dSep08	293684	487	387	2	40	putative protein (Scyl1) alternative variant dSep08, mRNA.
Scyl1bp1	Scyl1bp1.aSep08	304923	16620	2467		368	SCY1-like 1 binding protein 1 (41.6 kD) (Scyl1bp1) complete mRNA.
Scyl2	Scyl2.aSep08	314717	24980	833		185	SCY1-like 2 (S. cerevisiae) (Scyl2) mRNA.
Scyl3	Scyl3.aSep08	360866	7084	2887		392	SCY1-like 3 (S. cerevisiae) (Scyl3) mRNA.
Sdc2	Sdc2.bSep08	25615	59651	829	5	172	syndecan 2 (18.9 kD) (Sdc2) alternative variant bSep08, mRNA.
Sdc2	Sdc2.cSep08	25615	107669	730	4	134	syndecan 2 (14.6 kD) (Sdc2) alternative variant cSep08, mRNA.
Sdcbp	Sdcbp.bSep08	83841	25590	1656	8	282	syndecan binding protein (30.4 kD) (Sdcbp) alternative variant bSep08, mRNA.
Sdcbp	Sdcbp.cSep08	83841	25805	1164	9	278	syndecan binding protein (29.9 kD) (Sdcbp) alternative variant cSep08, mRNA.
Sdcbp	Sdcbp.dSep08	83841	24076	838	7	250	syndecan binding protein (Sdcbp) alternative variant dSep08, mRNA.

Sdcbp	Sdcbp.eSep08	83841	20597	641	6	185	syndecan binding protein (Sdcbp) alternative variant eSep08, mRNA.
Sdcbp	Sdcbp.fSep08	83841	6252	1554	4	111	syndecan binding protein (12.2 kD) (Sdcbp) alternative variant fSep08, mRNA.
Sdcbp	Sdcbp.gSep08	83841	1813	1019	2	68	syndecan binding protein (8.1 kD) (Sdcbp) alternative variant gSep08, mRNA.
Sdcbp2	Sdcbp2.bSep08	311532	1390	754	1	152	syndecan binding protein (syntenin) 2 (Sdcbp2) alternative variant bSep08, mRNA.
Sdccag3	Sdccag3.aSep08	306322	6096	1648	10	432	serologically defined colon cancer antigen 3 like (48.1 kD) (Sdccag3) alternative variant aSep08, mRNA.
Sdccag3	Sdccag3.bSep08	306322	6453	2409	6	306	serologically defined colon cancer antigen 3 like (34.1 kD) (Sdccag3) alternative variant bSep08, mRNA.
Sdccag3	Sdccag3.cSep08	306322	5765	849	5	257	serologically defined colon cancer antigen 3 like (Sdccag3) alternative variant cSep08, mRNA.
Sdccag3	Sdccag3.dSep08	306322	4355	1427	6	251	serologically defined colon cancer antigen 3 like (Sdccag3) alternative variant dSep08, mRNA.
Sdccag3	Sdccag3.eSep08	306322	6713	3561	5	240	serologically defined colon cancer antigen 3 like (26.5 kD) (Sdccag3) alternative variant eSep08, complete mRNA.
Sdccag3	Sdccag3.fSep08	306322	2414	739	4	217	serologically defined colon cancer antigen 3 like (Sdccag3) alternative variant fSep08, mRNA.
Sdccag3	Sdccag3.gSep08	306322	4143	698	4	216	serologically defined colon cancer antigen 3 like (Sdccag3) alternative variant gSep08, mRNA.
Sdccag3	Sdccag3.hSep08	306322	2573	536	2	127	serologically defined colon cancer antigen 3 like (Sdccag3) alternative variant hSep08, mRNA.
Sdccag3	Sdccag3.jSep08	306322	1889	381	2	72	serologically defined colon cancer antigen 3 like (8.1 kD) (Sdccag3) alternative variant jSep08, mRNA.
Sdccag8	Sdccag8.bSep08	305002	128975	508	4	169	serologically defined colon cancer antigen 8 (Sdccag8) alternative variant bSep08, mRNA.
Sdccag8	Sdccag8.cSep08	305002	11188	1076	3	63	serologically defined colon cancer antigen 8 (Sdccag8) alternative variant cSep08, mRNA.
Sdccag10	Sdccag10.bSep08	361887	14917	846	5	156	serologically defined colon cancer antigen 10 (Sdccag10) alternative variant bSep08, mRNA.
Sdccag10	Sdccag10.cSep08	361887	136886	701	1	102	serologically defined colon cancer antigen 10 (Sdccag10) alternative variant cSep08, mRNA.
Sdf4	Sdf4.bSep08	155173	15358	744	5	240	stromal cell derived factor 4 (Sdf4) alternative variant bSep08, mRNA.
Sdf4	Sdf4.cSep08	155173	15665	791	5	235	stromal cell derived factor 4 (26.9 kD) (Sdf4) alternative variant cSep08, mRNA.
Sdf4	Sdf4.dSep08	155173	3846	489	2	129	stromal cell derived factor 4 (14.6 kD) (Sdf4) alternative variant dSep08, mRNA.
Sdf4	Sdf4.eSep08	155173	2703	787	3	105	stromal cell derived factor 4 (Sdf4) alternative variant eSep08, mRNA.
Sdha	Sdha.bSep08	157074	14951	874	7	287	succinate dehydrogenase complex, subunit A, flavoprotein (Fp) (Sdha) alternative variant bSep08, mRNA.
Sdha	Sdha.cSep08	157074	9974	549	5	153	succinate dehydrogenase complex, subunit A, flavoprotein (Fp) (Sdha) alternative variant cSep08, mRNA.

Sdha	Sdha.dSep08	157074	4028	424	2	107	succinate dehydrogenase complex, subunit A, flavoprotein (Fp) (Sdha) alternative variant dSep08, mRNA.
Sdha	Sdha.fSep08	157074	5277	1034	3	87	succinate dehydrogenase complex, subunit A, flavoprotein (Fp) (Sdha) alternative variant fSep08, mRNA.
Sdhb	Sdhb.aSep08	298596	21422	1289	8	282	succinate dehydrogenase complex, subunit B, iron sulfur (lp) (31.8 kD) (Sdhb) alternative variant aSep08, mRNA.
Sdhb	Sdhb.bSep08	298596	2184	901	2	198	succinate dehydrogenase complex, subunit B, iron sulfur (lp) (Sdhb) alternative variant bSep08, mRNA.
Sdhb	Sdhb.cSep08	298596	3156	475	3	124	succinate dehydrogenase complex, subunit B, iron sulfur (lp) (Sdhb) alternative variant cSep08, mRNA.
Sdhb	Sdhb.dSep08	298596	12751	730	3	76	succinate dehydrogenase complex, subunit B, iron sulfur (lp) (Sdhb) alternative variant dSep08, mRNA.
Sdhc	Sdhc.bSep08	289217	20680	1283	8	132	succinate dehydrogenase complex, subunit C, integral membrane protein (14.2 kD) (Sdhc) alternative variant bSep08, complete mRNA.
Sdhd	Sdhd.bSep08	363061	8942	749	1	114	succinate dehydrogenase complex, subunit D, integral membrane protein (Sdhd) alternative variant bSep08, mRNA.
Sdk2	Sdk2.aSep08	360652	13105	679	4	216	sidekick homolog 2 (chicken) (Sdk2) alternative variant aSep08, mRNA.
Sds	Sds.bSep08	25044	814	705	1	56	serine dehydratase (Sds) alternative variant bSep08, mRNA.
Sds3.0	Sds3.0.aSep08		14170	1129	7	268	suppressor of defective silencing 3 (Sds3.0) alternative variant aSep08, mRNA.
Sec3l1	Sec3l1.bSep08	305287	16310	943	5	282	SEC3-like 1 ( <i>S. cerevisiae</i> ) (Sec3l1) alternative variant bSep08, mRNA.
Sec3l1	Sec3l1.cSep08	305287	19074	785	7	261	SEC3-like 1 ( <i>S. cerevisiae</i> ) (Sec3l1) alternative variant cSep08, mRNA.
Sec3l1	Sec3l1.dSep08	305287	1924	1415	3	133	SEC3-like 1 ( <i>S. cerevisiae</i> ) (14.9 kD) (Sec3l1) alternative variant dSep08, mRNA.
Sec7.0	Sec7.0.aSep08		8363	1382		341	IQ motif Sec7 domain 1 (Sec7.0) mRNA.
Sec8_exocyst.0	Sec8_exocyst.0.aSep08		58402	753		250	sec8 (Sec8_exocyst.0) mRNA.
Sec11a	Sec11a.bSep08	65166	11346	710	2	139	SEC11 homolog A ( <i>S. cerevisiae</i> ) (Sec11a) alternative variant bSep08, mRNA.
Sec11c	Sec11c.bSep08	266758	12122	859	3	133	SEC11 homolog C ( <i>S. cerevisiae</i> ) (Sec11c) alternative variant bSep08, mRNA.
Sec11c	Sec11c.cSep08	266758	3399	1020	3	49	SEC11 homolog C ( <i>S. cerevisiae</i> ) (5.6 kD) (Sec11c) alternative variant cSep08, mRNA.
Sec13	Sec13.bSep08	297522	5758	689	1	157	SEC13 homolog ( <i>S. cerevisiae</i> ) (Sec13) alternative variant bSep08, mRNA.
Sec14l1	Sec14l1.bSep08	360668	8367	1605	9	313	SEC14 -like 1 (35.0 kD) (Sec14l1) alternative variant bSep08, mRNA.
Sec14l1	Sec14l1.bSep08	690667	8367	1605	9	313	SEC14 -like 1 (35.0 kD) (Sec14l1) alternative variant bSep08, mRNA.
Sec14l1	Sec14l1.cSep08	360668	5906	852	4	220	SEC14 -like 1 (Sec14l1) alternative variant cSep08, mRNA.
Sec14l1	Sec14l1.cSep08	690667	5906	852	4	220	SEC14 -like 1 (Sec14l1) alternative variant cSep08, mRNA.

Sec1411	Sec1411.dSep08	360668	3159	618	3	166	SEC14 -like 1 (Sec1411) alternative variant dSep08, mRNA.
Sec1411	Sec1411.dSep08	690667	3159	618	3	166	SEC14 -like 1 (Sec1411) alternative variant dSep08, mRNA.
Sec1412	Sec1412.bSep08	116486	17607	742	1	247	SEC14-like 2 ( <i>S. cerevisiae</i> ) (Sec1412) alternative variant bSep08, mRNA.
Sec15.0	Sec15.0.aSep08		45195	1537		153	exocyst complex component 6 (Sec15.0) mRNA.
Sec16a	Sec16a.aSep08	114089	19398	3291	24	1005	SEC16 homolog A ( <i>S. cerevisiae</i> ) (Sec16a) alternative variant aSep08, mRNA.
Sec16a	Sec16a.bSep08	114089	10644	2528	7	263	SEC16 homolog A ( <i>S. cerevisiae</i> ) (27.6 kD) (Sec16a) alternative variant bSep08, mRNA.
Sec16a	Sec16a.cSep08	114089	5049	740	3	98	SEC16 homolog A ( <i>S. cerevisiae</i> ) (10.0 kD) (Sec16a) alternative variant cSep08, mRNA.
Sec16b	Sec16b.bSep08	89868	2973	618	3	139	SEC16 homolog B ( <i>S. cerevisiae</i> ) (15.6 kD) (Sec16b) alternative variant bSep08, mRNA.
Sec23a	Sec23a.bSep08	58817	12560	700	7	135	SEC23A ( <i>S. cerevisiae</i> ) (Sec23a) alternative variant bSep08, mRNA.
Sec23a	Sec23a.cSep08	58817	7032	419	4	119	SEC23A ( <i>S. cerevisiae</i> ) (Sec23a) alternative variant cSep08, mRNA.
Sec23a	Sec23a.dSep08	58817	6999	394	4	108	SEC23A ( <i>S. cerevisiae</i> ) (Sec23a) alternative variant dSep08, mRNA.
Sec23a	Sec23a.eSep08	58817	10169	549	3	80	SEC23A ( <i>S. cerevisiae</i> ) (9.2 kD) (Sec23a) alternative variant eSep08, mRNA.
Sec23b	Sec23b.aSep08	362226	28034	1552	14	517	protein transport Sec23 (Sec23b) alternative variant aSep08, mRNA.
Sec23b	Sec23b.cSep08	362226	3390	413	2	113	SEC23B (12.8 kD) (Sec23b) alternative variant cSep08, mRNA.
Sec23b	Sec23b.dSep08	362226	34280	382	3	31	SEC23B (3.3 kD) (Sec23b) alternative variant dSep08, mRNA.
Sec23ip	Sec23ip.bSep08	309010	23420	1783	6	396	SEC23 interacting protein (Sec23ip) alternative variant bSep08, mRNA.
Sec23ip	Sec23ip.cSep08	309010	7911	1665	5	154	SEC23 interacting protein (Sec23ip) alternative variant cSep08, mRNA.
Sec23ip	Sec23ip.dSep08	309010	2755	797	4	147	SEC23 interacting protein (Sec23ip) alternative variant dSep08, mRNA.
Sec23ip	Sec23ip.eSep08	309010	10645	434	3	144	SEC23 interacting protein (Sec23ip) alternative variant eSep08, mRNA.
Sec24a	Sec24a.bSep08	287275	12036	653	6	217	SEC24 related gene family, member A ( <i>S. cerevisiae</i> ) (Sec24a) alternative variant bSep08, mRNA.
Sec24a	Sec24a.cSep08	287275	4516	3112	2	59	SEC24 related gene family, member A ( <i>S. cerevisiae</i> ) (7.2 kD) (Sec24a) alternative variant cSep08, mRNA.
Sec24d	Sec24d.aSep08	310843	76122	3098	15	708	SEC24 related gene family, member D ( <i>S. cerevisiae</i> ) (Sec24d) alternative variant aSep08, mRNA.
Sec24d	Sec24d.cSep08	310843	4173	366	1	98	SEC24 related gene family, member D ( <i>S. cerevisiae</i> ) (Sec24d) alternative variant cSep08, mRNA.
Sec31a	Sec31a.bSep08	93646	56066	3669	24	1082	SEC31 homolog A ( <i>S. cerevisiae</i> ) (Sec31a) alternative variant bSep08, mRNA.

Sec31a	Sec31a.cSep08	93646	18618	1170	8	390	SEC31 homolog A ( <i>S. cerevisiae</i> ) (Sec31a) alternative variant cSep08, mRNA.
Sec31a	Sec31a.dSep08	93646	14068	748	6	249	SEC31 homolog A ( <i>S. cerevisiae</i> ) (Sec31a) alternative variant dSep08, mRNA.
Sec31a	Sec31a.eSep08	93646	5485	730	4	163	SEC31 homolog A ( <i>S. cerevisiae</i> ) (Sec31a) alternative variant eSep08, mRNA.
Sec31a	Sec31a.fSep08	93646	5378	384	4	127	SEC31 homolog A ( <i>S. cerevisiae</i> ) (Sec31a) alternative variant fSep08, mRNA.
Sec61a1	Sec61a1.bSep08	80843	6550	1179	6	350	sec61 (Sec61a1) alternative variant bSep08, mRNA.
Sec61a1	Sec61a1.dSep08	80843	3241	710	4	143	sec61 (Sec61a1) alternative variant dSep08, mRNA.
Sec61a2	Sec61a2.aSep08	361273	28526	2439	12	476	sec61, alpha subunit 2 ( <i>S. cerevisiae</i> ) (52.2 kD) (Sec61a2) alternative variant aSep08, mRNA.
Sec61a2	Sec61a2.bSep08	361273	22179	965	6	270	sec61, alpha subunit 2 ( <i>S. cerevisiae</i> ) (Sec61a2) alternative variant bSep08, mRNA.
Sec61a2	Sec61a2.cSep08	361273	22164	772	6	162	sec61, alpha subunit 2 ( <i>S. cerevisiae</i> ) (Sec61a2) alternative variant cSep08, mRNA.
Sec62	Sec62.bSep08	294912	26825	2584	8	481	SEC62 homolog ( <i>S. cerevisiae</i> ) (Sec62) alternative variant bSep08, mRNA.
Sec62	Sec62.cSep08	294912	17525	631	5	124	SEC62 homolog ( <i>S. cerevisiae</i> ) (Sec62) alternative variant cSep08, mRNA.
Sec63	Sec63.bSep08	309858	18331	1418	8	284	SEC63-like ( <i>S. cerevisiae</i> ) (Sec63) alternative variant bSep08, mRNA.
Sec63.1	Sec63.1.aSep08		16591	5510	28	1427	activating signal cointegrator 1 complex CRA c (Sec63.1) alternative variant aSep08, mRNA.
Sec63.1	Sec63.1.bSep08		1893	757	1	90	u5 snRNP-specific protein (Sec63.1) alternative variant bSep08, mRNA.
Secisbp2	Secisbp2.bSep08	79049	5412	807	5	252	SECIS binding protein 2 (Secisbp2) alternative variant bSep08, mRNA.
Secisbp2	Secisbp2.cSep08	79049	3607	756	4	219	SECIS binding protein 2 (Secisbp2) alternative variant cSep08, mRNA.
Secisbp2	Secisbp2.eSep08	79049	3545	504	3	96	SECIS binding protein 2 (Secisbp2) alternative variant eSep08, mRNA.
Sectm1b	Sectm1b.bSep08	287884	8159	827	3	186	secreted and transmembrane 1B (Sectm1b) alternative variant bSep08, mRNA.
Sectm1b	Sectm1b.cSep08	287884	1541	751	1	183	secreted and transmembrane 1B (Sectm1b) alternative variant cSep08, mRNA.
seeby	seeby.aSep08		31289	517		171	putative protein of vertebrate origin (seeby) mRNA.
seechy	seechy.aSep08		6236	3234		120	transmembrane protein 16C (seechy) mRNA.
seedar	seedar.aSep08		5820	477		86	putative protein (seedar) mRNA.
seeflo	seeflo.aSep08		2289	379		126	putative protein (seeflo) mRNA.
seeflu	seeflu.aSep08		2233	472		120	putative protein (seeflu) mRNA.
seegar	seegar.aSep08		3109	421		69	putative protein (seegar) mRNA.
seeja	seeja.aSep08		8073	446		64	putative protein (seeja) mRNA.
seejey	seejey.aSep08		10718	545		181	centrosomal protein 135 CRA a (seejey) mRNA.
seekee	seekee.aSep08		36611	245		20	putative protein (seekee) mRNA.

seekler	seekler.aSep08		36520	752		128	putative protein, with a coiled coil domain, of mammalian origin (15.2 kD) (seekler) mRNA.
seelo	seelo.aSep08		5597	627		142	putative protein (seelo) mRNA.
seemee	seemee.aSep08		22173	488		55	putative protein (seemee) mRNA.
seemer	seemer.aSep08		18950	2173	13	486	CRA a (54.5 kD) (seemer) alternative variant aSep08, mRNA.
seemer	seemer.bSep08		17336	1800	8	245	hexosaminidase containing CRA a (seemer) alternative variant bSep08, mRNA.
seemer	seemer.cSep08		1886	903	4	199	hexosaminidase containing (22.1 kD) (seemer) alternative variant cSep08, mRNA.
seemer	seemer.dSep08		3485	1110	4	172	hexosaminidase containing (seemer) alternative variant dSep08, mRNA.
seemer	seemer.eSep08		2354	638	6	150	hexosaminidase containing (seemer) alternative variant eSep08, mRNA.
seemer	seemer.fSep08		2500	369	4	122	hexosaminidase containing (seemer) alternative variant fSep08, mRNA.
seemer	seemer.gSep08		5502	1308	5	115	hexosaminidase containing CRA a (seemer) alternative variant gSep08, mRNA.
seemer	seemer.hSep08		3336	920	6	107	hexosaminidase containing (11.8 kD) (seemer) alternative variant hSep08, mRNA.
seenoy	seenoy.aSep08		31354	660		21	putative protein (2.4 kD) (seenoy) mRNA.
seepor	seepor.aSep08		2339	617		43	putative protein (5.2 kD) (seepor) mRNA.
seesa	seesa.aSep08		24095	664		97	putative protein (seesa) mRNA.
seeshee	seeshee.aSep08		3248	341		43	putative protein (4.8 kD) (seeshee) mRNA.
seetu	seetu.aSep08		897	355		117	otoferlin (seetu) mRNA.
seevar	seevar.aSep08		4385	669		54	putative protein (6.0 kD) (seevar) mRNA.
seewey	seewey.aSep08		19746	704		51	putative protein (seewey) mRNA.
Sel1.0	Sel1.0.aSep08		37471	1247	9	415	CRA a (Sel1.0) alternative variant aSep08, mRNA.
Sel1.0	Sel1.0.bSep08		17393	531	5	99	CRA a (Sel1.0) alternative variant bSep08, mRNA.
Sel1l	Sel1l.bSep08	314352	1496	398	3	122	sel1l (Sel1l) alternative variant bSep08, mRNA.
Sel1l2	Sel1l2.bSep08	311470	30860	792		264	sel-1 suppressor of 2 (Sel1l2) alternative variant bSep08, mRNA.
Sel1l2	Sel1l2.cSep08	311470	1039	325	2	82	sel-1 suppressor of 2 (Sel1l2) alternative variant cSep08, mRNA.
Selenbp1	Selenbp1.bSep08	140927	912	354	2	82	selenium binding protein 1 (Selenbp1) alternative variant bSep08, mRNA.
Selenbp1	Selenbp1.dSep08	140927	919	258	3	35	selenium binding protein 1 (Selenbp1) alternative variant dSep08, mRNA.
Seli	Seli.bSep08	362713	12694	2552	5	196	selenoprotein I (Seli) alternative variant bSep08, mRNA.
Selk	Selk.bSep08	290549	1937	1102	2	31	selenoprotein K (3.3 kD) (Selk) alternative variant bSep08, mRNA.
Selo	Selo.bSep08	315216	4853	1344	4	180	selenoprotein O (Selo) alternative variant bSep08, mRNA.
Selo	Selo.cSep08	315216	3904	873	6	123	selenoprotein O (Selo) alternative variant cSep08, mRNA.
Selo	Selo.dSep08	315216	2508	1960	2	66	selenoprotein O (7.9 kD) (Selo) alternative variant dSep08, mRNA.



Sels	Sels.bSep08	286900	960	426	2	71	selenoprotein S (Sels) alternative variant bSep08, mRNA.
Sema.0	Sema.0.aSep08		108985	813		271	sema 3C (Sema.0) mRNA.
Sema3b	Sema3b.bSep08	363142	2077	759	8	241	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B (Sema3b) alternative variant bSep08, mRNA.
Sema3c	Sema3c.aSep08	296787	62014	1136	2	378	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C (Sema3c) alternative variant aSep08, mRNA.
Sema3c	Sema3c.bSep08	296787	46346	730	2	242	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C (Sema3c) alternative variant bSep08, mRNA.
Sema3d	Sema3d.bSep08	246262	182172	2348	2	660	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3D (75.1 kD) (Sema3d) alternative variant bSep08, mRNA.
Sema3e	Sema3e.aSep08	296789	15677	640		210	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E (Sema3e) mRNA.
Sema3f	Sema3f.aSep08	315996	23724	2391	5	761	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3 F (Sema3f) alternative variant aSep08, mRNA.
Sema3f	Sema3f.bSep08	315996	2140	640	1	149	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3 F (Sema3f) alternative variant bSep08, mRNA.
Sema3g	Sema3g.aSep08	290562	2200	691		230	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3G (Sema3g) mRNA.
Sema4a	Sema4a.cSep08	310630	7584	611	6	203	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A (Sema4a) alternative variant cSep08, mRNA.
Sema4a	Sema4a.dSep08	310630	7713	570	6	189	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A (Sema4a) alternative variant dSep08, mRNA.
Sema4a	Sema4a.eSep08	310630	1566	405	4	113	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A (Sema4a) alternative variant eSep08, mRNA.
Sema4b	Sema4b.aSep08	293042	20997	1055		351	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B (Sema4b) mRNA.
Sema4c	Sema4c.aSep08	301346	718	395		131	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C (Sema4c) mRNA.
Sema4d	Sema4d.aSep08	306790	12552	1327		442	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D (Sema4d) alternative variant aSep08, mRNA.

Sema4d	Sema4d.bSep08	306790	10582	738		246	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D (Sema4d) alternative variant bSep08, mRNA.
Sema4f	Sema4f.bSep08	29745	3919	722	2	173	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F (Sema4f) alternative variant bSep08, mRNA.
Sema4g	Sema4g.aSep08	361764	2462	1167	9	389	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G (Sema4g) alternative variant aSep08, mRNA.
Sema4g	Sema4g.bSep08	361764	8073	1325	8	301	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G (Sema4g) alternative variant bSep08, mRNA.
Sema4g	Sema4g.cSep08	361764	1151	927	3	96	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G (10.2 kD) (Sema4g) alternative variant cSep08, mRNA.
Sema5b	Sema5b.bSep08	303901	4749	579	1	128	putative protein (Sema5b) alternative variant bSep08, mRNA.
Sema5b	Sema5b.bSep08	689558	4749	579	1	128	putative protein (Sema5b) alternative variant bSep08, mRNA.
Sema6a	Sema6a.bSep08	361324	33605	2658	7	560	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A (Sema6a) alternative variant bSep08, mRNA.
Sema6b	Sema6b.bSep08	84609	2070	774	6	258	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B (Sema6b) alternative variant bSep08, mRNA.
Sema6b	Sema6b.cSep08	84609	8642	447	4	148	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B (Sema6b) alternative variant cSep08, mRNA.
Sema6c	Sema6c.bSep08	29744	3271	1875	6	486	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant bSep08, mRNA.
Sema6c	Sema6c.cSep08	29744	2188	856	7	238	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant cSep08, mRNA.
Sema6c	Sema6c.dSep08	29744	7817	953	7	162	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (18.1 kD) (Sema6c) alternative variant dSep08, mRNA.
Sema6c	Sema6c.eSep08	29744	7575	663	6	139	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant eSep08, mRNA.

Sema6c	Sema6c.fSep08	29744	7547	878	8	130	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant fSep08, mRNA.
Sema6c	Sema6c.gSep08	29744	3857	433	4	124	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant gSep08, mRNA.
Sema6c	Sema6c.iSep08	29744	3879	407	3	64	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C (Sema6c) alternative variant iSep08, mRNA.
Sema6d	Sema6d.bSep08	311384	2006	342	2	114	sema domain transmembrane cytoplasmic 6D CRA b (Sema6d) alternative variant bSep08, mRNA.
Sema6d	Sema6d.cSep08	311384	44681	350	2	80	putative protein (Sema6d) alternative variant cSep08, mRNA.
Sema7a	Sema7a.aSep08	315711	1260	417		138	sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A (Sema7a) mRNA.
Senp1	Senp1.aSep08	300193	6691	338	3	112	SUMO1/sentrin specific peptidase 1 (Senp1) alternative variant aSep08, mRNA.
Senp1	Senp1.bSep08	300193	16066	761	3	59	SUMO1/sentrin specific peptidase 1 (Senp1) alternative variant bSep08, mRNA.
Senp2	Senp2.bSep08	78973	20854	910	9	186	SUMO/sentrin specific peptidase 2 (Senp2) alternative variant bSep08, mRNA.
Senp3	Senp3.bSep08	303245	4875	765	5	134	specific peptidase 3 CRA a (Senp3) alternative variant bSep08, mRNA.
Senp3	Senp3.cSep08	303245	2490	1882	3	55	specific peptidase 3 CRA a (6.5 kD) (Senp3) alternative variant cSep08, mRNA.
Senp3	Senp3.dSep08	303245	1964	1861	2	58	putative protein (6.2 kD) (Senp3) alternative variant dSep08, mRNA.
Senp5	Senp5.aSep08	303874	5804	3777		81	SUMO/sentrin specific protease 5 (Senp5) mRNA.
Senp6	Senp6.bSep08	300860	20542	1066	6	314	SUMO/sentrin specific peptidase 6 (Senp6) alternative variant bSep08, mRNA.
Senp6	Senp6.cSep08	300860	5957	744	3	172	SUMO/sentrin specific peptidase 6 (Senp6) alternative variant cSep08, mRNA.
Senp7	Senp7.aSep08	288167	55368	403		134	SUMO1/sentrin specific protease 7 (Senp7) mRNA.
Senp8	Senp8.aSep08	315723	13923	953	1	242	SUMO/sentrin specific peptidase 8 (Senp8) alternative variant aSep08, mRNA.
15-Sep	Sep15.aSep08	113922	32600	1735	7	214	selenoprotein (23.5 kD) (Sep15) alternative variant aSep08, mRNA.
15-Sep	Sep15.cSep08	113922	24550	736	4	124	selenoprotein (Sep15) alternative variant cSep08, mRNA.
15-Sep	Sep15.dSep08	113922	32033	844	4	106	selenoprotein (11.5 kD) (Sep15) alternative variant dSep08, mRNA.
15-Sep	Sep15.eSep08	113922	20199	393	3	92	selenoprotein (Sep15) alternative variant eSep08, mRNA.
15-Sep	Sep15.fSep08	113922	8276	965	2	71	selenoprotein (8.0 kD) (Sep15) alternative variant fSep08, mRNA.
15-Sep	Sep15.hSep08	113922	31682	735	6	113	selenoprotein (Sep15) alternative variant hSep08, mRNA.
Sephs1	Sephs1.bSep08	291314	18038	1110	7	270	selenophosphate synthetase 1 (30.2 kD) (Sephs1) alternative variant bSep08, mRNA.

Sephs1	Sephs1.cSep08	291314	20884	1251	6	263	selenophosphate synthetase 1 (Sephs1) alternative variant cSep08, mRNA.
Sephs1	Sephs1.dSep08	291314	13841	788	4	249	selenophosphate synthetase 1 (Sephs1) alternative variant dSep08, mRNA.
Sepp1	Sepp1.cSep08	29360	7280	905	4	126	selenoprotein P, plasma, 1 (14.9 kD) (Sepp1) alternative variant cSep08, mRNA.
1-Sep	Sept1.bSep08	293507	2169	648	5	215	septin 1 (Sept1) alternative variant bSep08, mRNA.
1-Sep	Sept1.cSep08	293507	3726	1902	10	163	septin 1 (18.3 kD) (Sept1) alternative variant cSep08, mRNA.
1-Sep	Sept1.dSep08	293507	3676	1971	9	191	septin 1 (21.5 kD) (Sept1) alternative variant dSep08, mRNA.
1-Sep	Sept1.eSep08	293507	2449	749	7	191	septin 1 (Sept1) alternative variant eSep08, mRNA.
1-Sep	Sept1.fSep08	293507	1807	734	5	165	septin 1 (Sept1) alternative variant fSep08, mRNA.
1-Sep	Sept1.gSep08	293507	674	570	2	45	septin 1 (5.3 kD) (Sept1) alternative variant gSep08, mRNA.
2-Sep	Sept2.bSep08	117515	26563	1087	10	345	septin 2 (Sept2) alternative variant bSep08, mRNA.
2-Sep	Sept2.cSep08	117515	6921	759	6	253	septin 2 (Sept2) alternative variant cSep08, mRNA.
2-Sep	Sept2.dSep08	117515	23302	775	8	215	septin 2 (Sept2) alternative variant dSep08, mRNA.
2-Sep	Sept2.eSep08	117515	20857	427	5	121	septin 2 (Sept2) alternative variant eSep08, mRNA.
2-Sep	Sept2.gSep08	117515	13185	995	3	73	septin 2 (Sept2) alternative variant gSep08, mRNA.
2-Sep	Sept2.hSep08	117515	19186	501	5	59	septin 2 (Sept2) alternative variant hSep08, mRNA.
3-Sep	Sept3.bSep08	56003	5614	458	4	118	septin 3 (Sept3) alternative variant bSep08, mRNA.
4-Sep	Sept4.aSep08	287606	12658	1333	11	360	septin 4 (41.9 kD) (Sept4) alternative variant aSep08, complete mRNA.
4-Sep	Sept4.cSep08	287606	11007	803	6	227	septin 4 (Sept4) alternative variant cSep08, mRNA.
4-Sep	Sept4.dSep08	287606	8031	691	6	207	septin 4 (Sept4) alternative variant dSep08, mRNA.
4-Sep	Sept4.eSep08	287606	11303	689	6	188	septin 4 CRA d (Sept4) alternative variant eSep08, mRNA.
4-Sep	Sept4.fSep08	287606	11674	668	7	179	septin 4 CRA a (Sept4) alternative variant fSep08, mRNA.
4-Sep	Sept4.hSep08	287606	11237	720	6	157	septin 4 CRA d (Sept4) alternative variant hSep08, mRNA.
4-Sep	Sept4.iSep08	287606	2021	686	4	114	septin 4 CRA l (Sept4) alternative variant iSep08, mRNA.
4-Sep	Sept4.jSep08	287606	1239	751	3	104	septin 4 (11.9 kD) (Sept4) alternative variant jSep08, mRNA.
4-Sep	Sept4.lSep08	287606	739	513	2	91	septin 4 CRA b (10.1 kD) (Sept4) alternative variant lSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.bSep08	116727	5839	756	8	252	septin 5 CRA g (Sept5andGp1bb) alternative variant bSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.bSep08	116728	5839	756	8	252	septin 5 CRA g (Sept5andGp1bb) alternative variant bSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.dSep08	116727	5970	1035	7	195	septin 5 CRA h (22.5 kD) (Sept5andGp1bb) alternative variant dSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.dSep08	116728	5970	1035	7	195	septin 5 CRA h (22.5 kD) (Sept5andGp1bb) alternative variant dSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.eSep08	116727	5368	575	6	191	septin 5 CRA d (Sept5andGp1bb) alternative variant eSep08, mRNA.

Sept5andGp1bb	Sept5andGp1bb.eSep08	116728	5368	575	6	191	septin 5 CRA d (Sept5andGp1bb) alternative variant eSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.fSep08	116727	508	412	2	65	septin 5 CRA f (Sept5andGp1bb) alternative variant fSep08, mRNA.
Sept5andGp1bb	Sept5andGp1bb.fSep08	116728	508	412	2	65	septin 5 CRA f (Sept5andGp1bb) alternative variant fSep08, mRNA.
6-Sep	Sept6.aSep08	691335	56118	1983	9	417	septin 6 (Sept6) alternative variant aSep08, mRNA.
6-Sep	Sept6.bSep08	691335	29800	1697	5	233	septin 6 (Sept6) alternative variant bSep08, mRNA.
6-Sep	Sept6.cSep08	691335	6897	731	1	38	septin 6 (Sept6) alternative variant cSep08, mRNA.
7-Sep	Sept7.bSep08	64551	30857	561	6	187	septin 7 (Sept7) alternative variant bSep08, mRNA.
7-Sep	Sept7.cSep08	64551	51656	727	8	187	septin 7 (Sept7) alternative variant cSep08, mRNA.
7-Sep	Sept7.dSep08	64551	51352	817	7	177	septin 7 (Sept7) alternative variant dSep08, mRNA.
8-Sep	Sept8.aSep08	303135	21764	1806	10	472	septin 8 (Sept8) alternative variant aSep08, mRNA.
8-Sep	Sept8.cSep08	303135	12349	1597	3	158	septin 8 (Sept8) alternative variant cSep08, mRNA.
8-Sep	Sept8.dSep08	303135	14124	412	4	137	septin 8 (Sept8) alternative variant dSep08, mRNA.
8-Sep	Sept8.eSep08	303135	2330	749	2	94	septin 8 (Sept8) alternative variant eSep08, mRNA.
9-Sep	Sept9.dSep08	83788	26292	725	6	209	septin 9 (Sept9) alternative variant dSep08, mRNA.
9-Sep	Sept9.fSep08	83788	5895	515	2	40	septin 9 (Sept9) alternative variant fSep08, mRNA.
10-Sep	Sept10.bSep08	309891	17425	1505	6	225	septin 10 (Sept10) alternative variant bSep08, mRNA.
10-Sep	Sept10.cSep08	309891	8606	886	2	72	septin 10 (Sept10) alternative variant cSep08, mRNA.
11-Sep	Sept11.aSep08	305227	83883	1510	10	475	septin 11 (Sept11) alternative variant aSep08, mRNA.
11-Sep	Sept11.bSep08	305227	84644	2203	9	468	septin 11 (Sept11) alternative variant bSep08, mRNA.
11-Sep	Sept11.dSep08	305227	84108	2376	10	431	septin 11 (49.7 kD) (Sept11) alternative variant dSep08, mRNA.
11-Sep	Sept11.fSep08	305227	4867	1453	1	56	septin 11 (Sept11) alternative variant fSep08, mRNA.
12-Sep	Sept12.aSep08	363542	9826	1295		381	septin 12 (Sept12) mRNA.
14-Sep	Sept14.aSep08	315702	3053	736	2	245	septin 14 (Sept14) alternative variant aSep08, mRNA.
14-Sep	Sept14.bSep08	315702	4097	1831	2	124	septin 14 (12.9 kD) (Sept14) alternative variant bSep08, mRNA.
Sepw1	Sepw1.aSep08	25545	5020	480	4	113	selenoprotein W, muscle 1 (12.3 kD) (Sepw1) alternative variant aSep08, complete mRNA.
Sepw1	Sepw1.bSep08	25545	3288	796	4	88	selenoprotein W, muscle 1 (9.6 kD) (Sepw1) alternative variant bSep08, mRNA.
Sepw1	Sepw1.cSep08	25545	5041	787	5	88	selenoprotein W, muscle 1 (9.6 kD) (Sepw1) alternative variant cSep08, complete mRNA.
Serac1	Serac1.aSep08	499015	7065	694		211	serine active site containing 1 (Serac1) mRNA.
Serbp1	Serbp1.bSep08	246303	17052	1586	8	413	serpine1 mRNA binding protein 1 (Serbp1) alternative variant bSep08, mRNA.
Serbp1	Serbp1.cSep08	246303	14536	1146	7	381	serpine1 mRNA binding protein 1 (Serbp1) alternative variant cSep08, mRNA.
Serbp1	Serbp1.dSep08	246303	16845	1289	7	377	serpine1 mRNA binding protein 1 (41.5 kD) (Serbp1) alternative variant dSep08, complete mRNA.
Serbp1	Serbp1.eSep08	246303	18587	3057	9	293	serpine1 mRNA binding protein 1 (Serbp1) alternative variant eSep08, mRNA.

Serbp1	Serbp1.fSep08	246303	5181	710	2	102	serpine1 mRNA binding protein 1 (Serbp1) alternative variant fSep08, mRNA.
serby	serby.aSep08		883	285		78	putative protein (serby) mRNA.
serchy	serchy.aSep08		3120	429		142	ryanodine receptor (serchy) mRNA.
serdar	serdar.aSep08		4050	274		80	putative protein (serdar) mRNA.
Serf1	Serf1.aSep08	502503	6048	476	3	92	small EDRK-rich factor 1 (Serf1) alternative variant aSep08, mRNA.
Serf2	Serf2.bSep08	502663	1318	583	3	153	small EDRK-rich factor 2 (Serf2) alternative variant bSep08, mRNA.
Serf2	Serf2.cSep08	502663	995	388	2	127	small EDRK-rich factor 2 (Serf2) alternative variant cSep08, mRNA.
Serf2	Serf2.eSep08	502663	1673	1139	2	39	small EDRK-rich factor 2 (4.5 kD) (Serf2) alternative variant eSep08, complete mRNA.
serflo	serflo.aSep08		3017	290		40	putative protein (serflo) mRNA.
serflu	serflu.aSep08		1701	348		116	leucine-rich repeat kinase 1 (serflu) mRNA.
sergar	sergar.aSep08		10546	267	3	83	putative protein, with a coiled coil domain (sergar) alternative variant aSep08, mRNA.
Sergef	Sergef.aSep08	365243	177623	718	4	178	secretion regulating guanine nucleotide exchange factor (18.9 kD) (Sergef) alternative variant aSep08, mRNA.
Serhl2	Serhl2.aSep08	500911	20444	1334	12	310	serine hydrolase-like 2 (35.5 kD) (Serhl2) alternative variant aSep08, mRNA.
Serhl2	Serhl2.bSep08	500911	17664	663	9	220	serine hydrolase-like 2 (Serhl2) alternative variant bSep08, mRNA.
Serinc1	Serinc1.aSep08	294421	18966	2792	2	453	serine incorporator 1 (50.6 kD) (Serinc1) alternative variant aSep08, mRNA.
Serinc1	Serinc1.bSep08	294421	3101	931	1	62	serine incorporator 1 (7.4 kD) (Serinc1) alternative variant bSep08, mRNA.
Serinc3	Serinc3.bSep08	296350	29073	2928	12	261	serine incorporator 3 (29.5 kD) (Serinc3) alternative variant bSep08, mRNA.
Serinc4	Serinc4.bSep08	311358	3481	881	2	263	serine incorporator 4 (29.2 kD) (Serinc4) alternative variant bSep08, mRNA.
serja	serja.aSep08		1378	271		35	putative protein (serja) mRNA.
serjey	serjey.aSep08		8200	416		138	centrosomal protein 135 CRA a (serjey) mRNA.
serkee	serkee.aSep08		25900	362		76	putative protein (serkee) mRNA.
serkler	serkler.aSep08		78366	427	3	118	putative protein (serkler) alternative variant aSep08, mRNA.
serkler	serkler.bSep08		20765	659	3	77	putative secreted or extracellular protein precursor (8.9 kD) (serkler) alternative variant bSep08, mRNA.
serkler	serkler.cSep08		49793	401	3	45	putative protein (serkler) alternative variant cSep08, mRNA.
serlo	serlo.aSep08		51932	573		81	furry homolog (serlo) mRNA.
sermee	sermee.aSep08		1348	399		133	adam metalloproteinase with thrombospondin type 1 motif 2 (sermee) mRNA.
sermer	sermer.aSep08		924	313		80	putative protein (8.7 kD) (sermer) mRNA.
sernoy	sernoy.aSep08		4648	770		64	putative protein (sernoy) mRNA.

Serp2	Serp2.bSep08	498546	24016	613	3	72	stress-associated endoplasmic reticulum protein family member 2 (Serp2) alternative variant bSep08, mRNA.
Serp2	Serp2.dSep08	498546	379	256	2	55	stress-associated endoplasmic reticulum protein family member 2 (Serp2) alternative variant dSep08, mRNA.
Serpina1	Serpina1.bSep08	24648	10086	1412	5	411	alpha-1-antiproteinase precursor (46.1 kD) (Serpina1) alternative variant bSep08, mRNA.
Serpina1	Serpina1.cSep08	24648	8276	775	2	258	alpha-1-antiproteinase (Serpina1) alternative variant cSep08, mRNA.
Serpina1	Serpina1.dSep08	24648	6278	684	2	213	alpha-1-antiproteinase precursor (23.9 kD) (Serpina1) alternative variant dSep08, mRNA.
Serpina1	Serpina1.eSep08	24648	4028	572	3	77	CRA b like (8.6 kD) (Serpina1) alternative variant eSep08, mRNA.
Serpina1	Serpina1.fSep08	24648	1274	468	2	32	CRA a like (Serpina1) alternative variant fSep08, mRNA.
Serpina3k	Serpina3k.bSep08	24794	4676	927	3	275	serine (or cysteine) peptidase inhibitor, clade A, member 3K (Serpina3k) alternative variant bSep08, mRNA.
Serpina3m	Serpina3m.aSep08	299276	5721	1311		321	serine (or cysteine) proteinase inhibitor, clade A, member 3M (36.1 kD) (Serpina3m) mRNA.
Serpina5	Serpina5.bSep08	65051	2087	828	2	273	serine (or cysteine) peptidase inhibitor, clade A, member 5 (Serpina5) alternative variant bSep08, mRNA.
Serpina6	Serpina6.bSep08	299270	1971	422	1	34	serine (or cysteine) peptidase inhibitor, clade A, member 6 (Serpina6) alternative variant bSep08, mRNA.
Serpina7	Serpina7.aSep08	81806	1142	430		98	serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 7 (Serpina7) mRNA.
Serpina10	Serpina10.bSep08	171154	2149	871	2	263	serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 10 (Serpina10) alternative variant bSep08, mRNA.
Serpina11	Serpina11.bSep08	362774	9424	1468	1	422	serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 11 (47.0 kD) (Serpina11) alternative variant bSep08, mRNA.
Serpina11	Serpina11.cSep08	362774	2935	543	1	150	serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 11 (Serpina11) alternative variant cSep08, mRNA.
Serpib1a	Serpib1a.bSep08	291091	4058	557	1	165	serine (or cysteine) proteinase inhibitor, clade B, member 1a (18.2 kD) (Serpib1a) alternative variant bSep08, complete mRNA.
Serpib3a	Serpib3a.aSep08	498209	5156	1617		395	serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 3A (Serpib3a) mRNA.
Serpib6a	Serpib6a.bSep08	291085	17971	835	6	245	serine (or cysteine) peptidase inhibitor, clade B, member 6a (Serpib6a) alternative variant bSep08, mRNA.
Serpib6a	Serpib6a.cSep08	291085	13471	762	5	181	serine (or cysteine) peptidase inhibitor, clade B, member 6a (Serpib6a) alternative variant cSep08, mRNA.
Serpib6b	Serpib6b.bSep08	364705	12423	778	5	259	serine (or cysteine) peptidase inhibitor, clade B, member 6b (Serpib6b) alternative variant bSep08, mRNA.
Serpib6b	Serpib6b.cSep08	364705	4043	736	1	44	serine (or cysteine) peptidase inhibitor, clade B, member 6b (Serpib6b) alternative variant cSep08, mRNA.

Serpib11	Serpib11.bSep08	304689	6798	811	1	165	serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 11 (Serpib11) alternative variant bSep08, mRNA.
Serpib12	Serpib12.aSep08	304692	21622	602		159	serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 12 (Serpib12) mRNA.
Serpinc1	Serpinc1.bSep08	304917	8246	995	3	295	serine (or cysteine) peptidase inhibitor, clade C (antithrombin), member 1 (Serpinc1) alternative variant bSep08, mRNA.
Serpinc1	Serpinc1.cSep08	304917	6424	728	1	212	serine (or cysteine) peptidase inhibitor, clade C (antithrombin), member 1 (23.6 kD) (Serpinc1) alternative variant cSep08, mRNA.
Serpind1	Serpind1.bSep08	79224	8517	1160		367	serine (or cysteine) peptidase inhibitor, clade D, member 1 (Serpind1) alternative variant bSep08, mRNA.
Serpine1	Serpine1.bSep08	24617	4971	750	2	123	serine (or cysteine) peptidase inhibitor, clade E, member 1 (Serpine1) alternative variant bSep08, mRNA.
Serpine2	Serpine2.bSep08	29366	52195	570	1	190	serine (or cysteine) peptidase inhibitor, clade E, member 2 (Serpine2) alternative variant bSep08, mRNA.
Serpinf1	Serpinf1.bSep08	287526	5778	1800	6	393	serine (or cysteine) peptidase inhibitor, clade F, member 1 (Serpinf1) alternative variant bSep08, mRNA.
Serpinf2	Serpinf2.bSep08	287527	3390	810	7	159	serine (or cysteine) peptidase inhibitor, clade F, member 2 (Serpinf2) alternative variant bSep08, mRNA.
Serpinf2	Serpinf2.cSep08	287527	930	808	1	131	serine (or cysteine) peptidase inhibitor, clade F, member 2 (Serpinf2) alternative variant cSep08, mRNA.
Serping1	Serping1.bSep08	295703	4718	912	3	282	serine (or cysteine) peptidase inhibitor, clade G, member 1 (Serping1) alternative variant bSep08, mRNA.
Serping1	Serping1.cSep08	295703	3554	702	2	213	serine (or cysteine) peptidase inhibitor, clade G, member 1 (Serping1) alternative variant cSep08, mRNA.
Serpinh1	Serpinh1.bSep08	29345	889	796	2	115	serine (or cysteine) peptidase inhibitor, clade H, member 1 (Serpinh1) alternative variant bSep08, mRNA.
Serpinh1	Serpinh1.cSep08	29345	4050	385	3	104	serine (or cysteine) peptidase inhibitor, clade H, member 1 (Serpinh1) alternative variant cSep08, mRNA.
Serpinh1	Serpinh1.dSep08	29345	3885	393	3	90	serine (or cysteine) peptidase inhibitor, clade H, member 1 (Serpinh1) alternative variant dSep08, mRNA.
Serpini1	Serpini1.bSep08	116459	51846	341	1	44	serine (or cysteine) peptidase inhibitor, clade I, member 1 (Serpini1) alternative variant bSep08, mRNA.
serpor	serpor.aSep08		6525	401		106	myosin Va (serpor) mRNA.
sersa	sersa.aSep08		2847	698		164	putative protein (18.3 kD) (sersa) mRNA.
sershee	sershee.aSep08		785	301		93	putative protein (sershee) mRNA.
SERTA.0	SERTA.0.aSep08		12060	743		151	SERTA (18.3 kD) (SERTA.0) mRNA.
Sertad2	Sertad2.bSep08	498423	16388	390	1	62	putative protein (Sertad2) alternative variant bSep08, mRNA.
Sertad4	Sertad4.cSep08	360899	8074	470	3	136	SERTA (Sertad4) alternative variant cSep08, mRNA.
Sertad4	Sertad4.dSep08	360899	5344	461	1	36	putative protein of mammalian origin (Sertad4) alternative variant dSep08, mRNA.
sertu	sertu.aSep08		37687	750		63	ras-related GTP-binding protein Rab10 like (sertu) mRNA.
servar	servar.aSep08		10215	436		120	paired box 7 (servar) mRNA.



serwey	serwey.aSep08		2989	370		62	putative protein (serwey) mRNA.
Sesn1	Sesn1.aSep08	294518	21175	2405	9	610	sestrin 1 (Sesn1) alternative variant aSep08, complete mRNA.
Sesn1	Sesn1.bSep08	294518	21147	1790	6	464	sestrin 1 (Sesn1) alternative variant bSep08, mRNA.
Sestd1	Sestd1.aSep08	295678	69053	2810	18	721	SEC14 and spectrin domains 1 (Sestd1) alternative variant aSep08, mRNA.
SET.1	SET.1.aSep08		4311	747		249	SET (SET.1) alternative variant aSep08, mRNA.
SET.1	SET.1.bSep08		2015	461		100	histone-lysine n-methyltransferase setd1b (SET.1) alternative variant bSep08, mRNA.
SET.2	SET.2.aSep08		656	379		126	mixed-lineage leukemia like (SET.2) mRNA.
SET.3	SET.3.aSep08		9157	2465	5	178	N-methyltransferase mll3 (SET.3) alternative variant aSep08, mRNA.
Setbp1	Setbp1.aSep08	291423	64219	1136		175	SET binding protein 1 (19.0 kD) (Setbp1) mRNA.
Setd2	Setd2.aSep08	316013	24745	4505	4	1501	putative protein of bilateral origin (Setd2) alternative variant aSep08, mRNA.
Setd2	Setd2.cSep08	316013	50618	3350	14	903	WW/Rsp5/WWP (Setd2) alternative variant cSep08, mRNA.
Setd2	Setd2.dSep08	316013	17822	841	4	147	putative protein of vertebrate origin (Setd2) alternative variant dSep08, mRNA.
Setd2	Setd2.gSep08	316013	16319	760	6	72	WW/Rsp5/WWP (Setd2) alternative variant gSep08, mRNA.
Setd3	Setd3.aSep08	299295	66871	2699	1	596	SET (67.4 kD) (Setd3) alternative variant aSep08, mRNA.
Setd3	Setd3.bSep08	299295	20929	404		81	putative protein of vertebrate origin (9.0 kD) (Setd3) alternative variant bSep08, mRNA.
Setd4	Setd4.bSep08	245975	4506	2221	3	270	putative protein of eukaryotic origin (30.5 kD) (Setd4) alternative variant bSep08, mRNA.
Setd4	Setd4.cSep08	245975	6365	1017	5	139	protein RDA279 CRA b (15.6 kD) (Setd4) alternative variant cSep08, mRNA.
Setd4	Setd4.eSep08	245975	12491	633	4	111	putative protein of mammalian origin (13.3 kD) (Setd4) alternative variant eSep08, mRNA.
Setd4	Setd4.fSep08	245975	3087	329	3	87	putative protein of vertebrate origin (Setd4) alternative variant fSep08, mRNA.
Setd4	Setd4.gSep08	245975	12680	785	4	44	putative protein (4.6 kD) (Setd4) alternative variant gSep08, mRNA.
Setd5	Setd5.bSep08	297514	7490	939	3	178	putative protein of vertebrate origin (Setd5) alternative variant bSep08, mRNA.
Setd5	Setd5.cSep08	297514	6327	743	7	117	putative protein of vertebrate origin (12.7 kD) (Setd5) alternative variant cSep08, mRNA.
Setd5	Setd5.iSep08	297514	5331	439	4	90	putative protein (Setd5) alternative variant iSep08, mRNA.
Setd5	Setd5.jSep08	297514	32131	437	3	33	putative protein (Setd5) alternative variant jSep08, mRNA.
Setd6	Setd6.bSep08	291844	943	707	2	156	putative protein of eukaryotic origin (Setd6) alternative variant bSep08, mRNA.
Setd6	Setd6.cSep08	291844	4637	448	2	103	putative protein of vertebrate origin (Setd6) alternative variant cSep08, mRNA.

Setdb1	Setdb1.aSep08	689873	3283	622	1	176	similar to Histone-lysine N-methyltransferase, H3 lysine-9 specific 4 (Histone H3-K9 methyltransferase 4) (H3-K9-HMTase 4) (SET domain bifurcated 1) (ERG-associated protein with SET domain) (ESET) and SET domain, bifurcated 1 (19.5 kD) (Setdb1) alternative variant aSep08, mRNA.
Setdb1	Setdb1.aSep08	689883	3283	622	1	176	similar to Histone-lysine N-methyltransferase, H3 lysine-9 specific 4 (Histone H3-K9 methyltransferase 4) (H3-K9-HMTase 4) (SET domain bifurcated 1) (ERG-associated protein with SET domain) (ESET) and SET domain, bifurcated 1 (19.5 kD) (Setdb1) alternative variant aSep08, mRNA.
Setdb1	Setdb1.cSep08	689873	2843	395	2	118	similar to Histone-lysine N-methyltransferase, H3 lysine-9 specific 4 (Histone H3-K9 methyltransferase 4) (H3-K9-HMTase 4) (SET domain bifurcated 1) (ERG-associated protein with SET domain) (ESET) and SET domain, bifurcated 1 (Setdb1) alternative variant cSep08, mRNA.
Setdb1	Setdb1.cSep08	689883	2843	395	2	118	similar to Histone-lysine N-methyltransferase, H3 lysine-9 specific 4 (Histone H3-K9 methyltransferase 4) (H3-K9-HMTase 4) (SET domain bifurcated 1) (ERG-associated protein with SET domain) (ESET) and SET domain, bifurcated 1 (Setdb1) alternative variant cSep08, mRNA.
Setmar	Setmar.aSep08	500281	12005	1557		315	SET domain and mariner transposase fusion gene (35.0 kD) (Setmar) mRNA.
Setx	Setx.aSep08	362096	4681	406		134	senataxin (Setx) mRNA.
seyby	seyby.aSep08		1953	1002		112	polyprotein (12.4 kD) (seyby) mRNA.
seychy	seychy.aSep08		3295	354		117	ryanodine receptor (seychy) mRNA.
seydar	seydar.aSep08		1596	194		64	putative protein (seydar) mRNA.
seyflo	seyflo.aSep08		15733	617		193	probable rna-binding protein 20 like (seyflo) mRNA.
seyflu	seyflu.aSep08		1107	214		21	putative protein (seyflu) mRNA.
seygar	seygar.aSep08		14200	542		37	putative protein (seygar) mRNA.
seyja	seyja.aSep08		7040	308		30	putative protein (seyja) mRNA.
seyjey	seyjey.aSep08		10625	1134		218	exocyst complex component 1 (seyjey) mRNA.
seykee	seykee.aSep08		23735	961		80	putative mitochondrial protein (9.1 kD) (seykee) mRNA.
seykler	seykler.aSep08		10975	322		46	putative protein (seykler) mRNA.
seylo	seylo.aSep08		50785	406		90	putative protein (seylo) mRNA.
seymee	seymee.aSep08		1573	537		98	putative protein (seymee) mRNA.
seymer	seymer.aSep08		853	525		77	putative protein (seymer) mRNA.
seynoy	seynoy.aSep08		5548	377		125	dedicator of cytokinesis 10 (seynoy) mRNA.
seypor	seypor.aSep08		8750	561		187	myosin Va (seypor) mRNA.
seysa	seysa.aSep08		2011	352		49	putative protein (seysa) mRNA.
seyshee	seyshee.aSep08		7099	1270	2	129	folliculin interacting protein (seyshee) alternative variant aSep08, mRNA.
seyshee	seyshee.bSep08		3874	437	1	69	folliculin interacting protein (seyshee) alternative variant bSep08, mRNA.

seyshee	seyshee.cSep08		5815	1779	3	64	folliculin interacting protein (seyshee) alternative variant cSep08, mRNA.
seytu	seytu.aSep08		14000	429		110	putative protein (seytu) mRNA.
seyvar	seyvar.aSep08		5795	947		34	putative protein (4.0 kD) (seyvar) mRNA.
seywey	seywey.aSep08		11397	613		45	putative protein (seywey) mRNA.
Sez6	Sez6.aSep08	192247	20683	1120		373	seizure related gene 6 (Sez6) mRNA.
Sez6l2	Sez6l2.bSep08	308988	1702	542	3	180	seizure related 6 homolog like 2 (Sez6l2) alternative variant bSep08, mRNA.
Sez6l2	Sez6l2.cSep08	308988	877	765	2	154	seizure related 6 homolog like 2 (Sez6l2) alternative variant cSep08, mRNA.
Sez6l2	Sez6l2.dSep08	308988	3181	915	4	131	seizure related 6 homolog like 2 (Sez6l2) alternative variant dSep08, mRNA.
Sf1	Sf1.cSep08	117855	2422	739	5	246	splicing factor 1 (Sf1) alternative variant cSep08, mRNA.
Sf1	Sf1.dSep08	117855	7621	682	3	102	splicing factor 1 (Sf1) alternative variant dSep08, mRNA.
Sf1	Sf1.eSep08	117855	783	668	2		
Sf3a1	Sf3a1.bSep08	305479	8165	1784	2	382	splicing factor 3a, subunit 1 (Sf3a1) alternative variant bSep08, mRNA.
Sf3a2	Sf3a2.bSep08	299620	1438	666	1	204	splicing factor 3a, subunit 2 (Sf3a2) alternative variant bSep08, mRNA.
Sf3a3	Sf3a3.bSep08	313583	11137	2146	9	243	splicing factor 3a, subunit 3 (Sf3a3) alternative variant bSep08, mRNA.
Sf3a3	Sf3a3.cSep08	313583	7577	566	7	45	splicing factor 3a, subunit 3 (5.1 kD) (Sf3a3) alternative variant cSep08, complete mRNA.
Sf3b1	Sf3b1.bSep08	84486	11420	1791	5	462	splicing factor 3b (Sf3b1) alternative variant bSep08, mRNA.
Sf3b1	Sf3b1.cSep08	84486	2327	1315	4	333	splicing factor 3b (Sf3b1) alternative variant cSep08, mRNA.
Sf3b1	Sf3b1.dSep08	84486	10943	608	4	159	splicing factor 3b CRA a (Sf3b1) alternative variant dSep08, mRNA.
Sf3b1	Sf3b1.eSep08	84486	1924	720	4	158	splicing factor 3b (Sf3b1) alternative variant eSep08, mRNA.
Sf3b1	Sf3b1.fSep08	84486	7578	1042	4	153	splicing factor 3b (Sf3b1) alternative variant fSep08, mRNA.
Sf3b1	Sf3b1.gSep08	84486	1738	602	3	128	splicing factor 3b (Sf3b1) alternative variant gSep08, mRNA.
Sf3b1	Sf3b1.hSep08	84486	708	343	3	55	putative protein (6.5 kD) (Sf3b1) alternative variant hSep08, mRNA.
Sf3b2	Sf3b2.bSep08	293671	6234	576	3	76	splicing factor 3b, subunit 2 (8.8 kD) (Sf3b2) alternative variant bSep08, mRNA.
Sf3b3	Sf3b3.bSep08	292019	4002	772	5	247	splicing factor 3b, subunit 3 (Sf3b3) alternative variant bSep08, mRNA.
Sf3b3	Sf3b3.cSep08	292019	3712	358	2	85	splicing factor 3b, subunit 3 (Sf3b3) alternative variant cSep08, mRNA.
Sf3b3	Sf3b3.dSep08	292019	5293	412	3	16	splicing factor 3b, subunit 3 (Sf3b3) alternative variant dSep08, mRNA.

Sf3b4	Sf3b4.aSep08	295270	4746	1791	2	541	splicing factor 3b, subunit 4 (Sf3b4) alternative variant aSep08, complete mRNA.
Sf3b4	Sf3b4.cSep08	295270	1353	862	2	171	splicing factor 3b, subunit 4 (18.5 kD) (Sf3b4) alternative variant cSep08, mRNA.
Sf4	Sf4.bSep08	290666	25175	1799	11	554	splicing factor 4 (Sf4) alternative variant bSep08, mRNA.
Sf4	Sf4.cSep08	290666	24125	788	7	262	splicing factor 4 (Sf4) alternative variant cSep08, mRNA.
Sf4	Sf4.dSep08	290666	7152	579	2	160	splicing factor 4 (Sf4) alternative variant dSep08, mRNA.
Sf4	Sf4.eSep08	290666	4600	638	2	121	splicing factor 4 CRA c (Sf4) alternative variant eSep08, mRNA.
Sf4	Sf4.fSep08	290666	752	400	2	86	splicing factor 4 (Sf4) alternative variant fSep08, mRNA.
Sfmbt1	Sfmbt1.bSep08	58967	4751	704	4	185	scm-like with four mbt domains 1 and similar to farnesyl diphosphate synthetase (Sfmbt1) alternative variant bSep08, mRNA.
Sfmbt1	Sfmbt1.bSep08	680626	4751	704	4	185	scm-like with four mbt domains 1 and similar to farnesyl diphosphate synthetase (Sfmbt1) alternative variant bSep08, mRNA.
Sfmbt1	Sfmbt1.cSep08	58967	30511	435	4	73	scm-like with four mbt domains 1 and similar to farnesyl diphosphate synthetase (Sfmbt1) alternative variant cSep08, mRNA.
Sfmbt1	Sfmbt1.cSep08	680626	30511	435	4	73	scm-like with four mbt domains 1 and similar to farnesyl diphosphate synthetase (Sfmbt1) alternative variant cSep08, mRNA.
Sfpi1	Sfpi1.bSep08	366126	31857	385	4	94	SFFV proviral integration 1 (Sfpi1) alternative variant bSep08, mRNA.
Sfpi1	Sfpi1.cSep08	366126	7300	167	2	31	SFFV proviral integration 1 (Sfpi1) alternative variant cSep08, mRNA.
Sfpq	Sfpq.bSep08	252855	10126	3902	10	607	splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (Sfpq) alternative variant bSep08, mRNA.
Sfpq	Sfpq.dSep08	252855	4282	3523	2	75	splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (8.3 kD) (Sfpq) alternative variant dSep08, mRNA.
Sfpq	Sfpq.eSep08	252855	3124	2082	3	75	splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (8.3 kD) (Sfpq) alternative variant eSep08, mRNA.
Sfpq	Sfpq.fSep08	252855	4821	850	4	43	splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (5.2 kD) (Sfpq) alternative variant fSep08, mRNA.
Sfpq	Sfpq.gSep08	252855	13335	667	6		
Sfrp1	Sfrp1.aSep08	84402	38338	3880		404	secreted frizzled-related protein 1 (Sfrp1) mRNA.
Sfrs1	Sfrs1.aSep08	689890	4178	3212	4	248	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (27.7 kD) (Sfrs1) alternative variant aSep08, mRNA.
Sfrs1	Sfrs1.aSep08	689898	4178	3212	4	248	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (27.7 kD) (Sfrs1) alternative variant aSep08, mRNA.

Sfrs1	Sfrs1.bSep08	689890	2294	1525	3	223	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant bSep08, mRNA.
Sfrs1	Sfrs1.bSep08	689898	2294	1525	3	223	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant bSep08, mRNA.
Sfrs1	Sfrs1.cSep08	689890	1094	675	2	161	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant cSep08, mRNA.
Sfrs1	Sfrs1.cSep08	689898	1094	675	2	161	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant cSep08, mRNA.
Sfrs1	Sfrs1.eSep08	689890	1481	594	2	53	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant eSep08, mRNA.
Sfrs1	Sfrs1.eSep08	689898	1481	594	2	53	splicing factor, arginine/serine-rich 1 and hypothetical protein LOC689898 (Sfrs1) alternative variant eSep08, mRNA.
Sfrs2	Sfrs2.cSep08	494445	1241	644	3	70	splicing factor, arginine/serine-rich 2 (SC-35) (Sfrs2) alternative variant cSep08, mRNA.
Sfrs2	Sfrs2.dSep08	494445	1589	1087	2	43	splicing factor, arginine/serine-rich 2 (SC-35) (4.9 kD) (Sfrs2) alternative variant dSep08, mRNA.
Sfrs2ip	Sfrs2ip.aSep08	312030	39552	649		216	splicing factor, arginine/serine-rich 2, interacting protein (Sfrs2ip) mRNA.
Sfrs3	Sfrs3.bSep08	361814	7967	1360	7	147	CRA c like (16.4 kD) (Sfrs3) alternative variant bSep08, complete mRNA.
Sfrs3	Sfrs3.cSep08	361814	5726	1069	3	129	splicing factor arginine serine-rich 3 (15.0 kD) (Sfrs3) alternative variant cSep08, complete mRNA.
Sfrs3	Sfrs3.eSep08	361814	882	795	2	55	CRA c like (Sfrs3) alternative variant eSep08, mRNA.
Sfrs3	Sfrs3.gSep08	361814	730	484	3	37	CRA e like (Sfrs3) alternative variant gSep08, mRNA.
Sfrs5	Sfrs5.aSep08	29667	4696	1470	8	269	splicing factor, arginine/serine-rich 5 (30.9 kD) (Sfrs5) alternative variant aSep08, complete mRNA.
Sfrs5	Sfrs5.cSep08	29667	3859	748	7	190	splicing factor, arginine/serine-rich 5 (22.0 kD) (Sfrs5) alternative variant cSep08, mRNA.
Sfrs5	Sfrs5.dSep08	29667	3804	818	7	183	splicing factor, arginine/serine-rich 5 (Sfrs5) alternative variant dSep08, mRNA.
Sfrs5	Sfrs5.eSep08	29667	2620	1449	6	138	splicing factor, arginine/serine-rich 5 (15.8 kD) (Sfrs5) alternative variant eSep08, mRNA.
Sfrs5	Sfrs5.fSep08	29667	2429	740	6	138	splicing factor, arginine/serine-rich 5 (15.8 kD) (Sfrs5) alternative variant fSep08, mRNA.
Sfrs5	Sfrs5.gSep08	29667	4535	2281	8	138	splicing factor, arginine/serine-rich 5 (15.8 kD) (Sfrs5) alternative variant gSep08, complete mRNA.
Sfrs5	Sfrs5.hSep08	29667	2973	1155	6	138	splicing factor, arginine/serine-rich 5 (15.8 kD) (Sfrs5) alternative variant hSep08, mRNA.
Sfrs5	Sfrs5.iSep08	29667	2185	758	5	124	splicing factor, arginine/serine-rich 5 (14.4 kD) (Sfrs5) alternative variant iSep08, mRNA.

Sfrs5	Sfrs5.jSep08	29667	2355	799	5	124	splicing factor, arginine/serine-rich 5 (14.4 kD) (Sfrs5) alternative variant jSep08, mRNA.
Sfrs5	Sfrs5.kSep08	29667	3052	1500	5	124	splicing factor, arginine/serine-rich 5 (14.4 kD) (Sfrs5) alternative variant kSep08, mRNA.
Sfrs5	Sfrs5.lSep08	29667	2284	912	4	119	splicing factor, arginine/serine-rich 5 (14.0 kD) (Sfrs5) alternative variant lSep08, complete mRNA.
Sfrs5	Sfrs5.nSep08	29667	1729	384	4	86	splicing factor, arginine/serine-rich 5 (Sfrs5) alternative variant nSep08, mRNA.
Sfrs6	Sfrs6.bSep08	362264	4615	4264	3	210	splicing factor, arginine/serine-rich 6 (24.1 kD) (Sfrs6) alternative variant bSep08, mRNA.
Sfrs6	Sfrs6.cSep08	362264	1256	408	3	94	splicing factor, arginine/serine-rich 6 (Sfrs6) alternative variant cSep08, mRNA.
Sfrs6	Sfrs6.dSep08	362264	1133	688	2	65	splicing factor, arginine/serine-rich 6 (Sfrs6) alternative variant dSep08, mRNA.
Sfrs7	Sfrs7.aSep08	362687	6077	1180	8	238	splicing factor arginine serine-rich 7 (27.4 kD) (Sfrs7) alternative variant aSep08, mRNA.
Sfrs7	Sfrs7.bSep08	362687	5784	854	8	227	splicing factor arginine serine-rich 7 (26.2 kD) (Sfrs7) alternative variant bSep08, mRNA.
Sfrs7	Sfrs7.cSep08	362687	5757	815	7	223	splicing factor arginine serine-rich 7 (25.6 kD) (Sfrs7) alternative variant cSep08, mRNA.
Sfrs7	Sfrs7.dSep08	362687	5768	1988	7	179	arginine serine-rich 7 splicing factor (Sfrs7) alternative variant dSep08, mRNA.
Sfrs7	Sfrs7.fSep08	362687	3544	632	5	117	splicing factor arginine serine-rich 7 (Sfrs7) alternative variant fSep08, mRNA.
Sfrs7	Sfrs7.gSep08	362687	4024	1259	6	113	putative mitochondrial protein (13.1 kD) (Sfrs7) alternative variant gSep08, mRNA.
Sfrs7	Sfrs7.hSep08	362687	422	341	2	93	putative mitochondrial protein (10.8 kD) (Sfrs7) alternative variant hSep08, mRNA.
Sfrs7	Sfrs7.iSep08	362687	3169	1898	3	74	CRA i like (Sfrs7) alternative variant iSep08, mRNA.
Sfrs8	Sfrs8.bSep08	304431	10200	928	6	309	splicing factor arginine serine-rich 8 (Sfrs8) alternative variant bSep08, mRNA.
Sfrs8	Sfrs8.dSep08	304431	5967	717	4	197	splicing factor arginine serine-rich 8 (Sfrs8) alternative variant dSep08, mRNA.
Sfrs8	Sfrs8.eSep08	304431	10754	819	4	134	splicing factor arginine serine-rich 8 (Sfrs8) alternative variant eSep08, mRNA.
Sfrs8	Sfrs8.fSep08	304431	2860	481	2	96	putative nuclear protein (10.7 kD) (Sfrs8) alternative variant fSep08, mRNA.
Sfrs8	Sfrs8.gSep08	304431	1337	744	2	77	putative mitochondrial protein (8.8 kD) (Sfrs8) alternative variant gSep08, mRNA.
Sfrs9	Sfrs9.bSep08	288701	5863	790	5	129	splicing factor arginine serine rich 9 CRA a (Sfrs9) alternative variant bSep08, mRNA.
Sfrs9	Sfrs9.dSep08	288701	4447	1783	3	89	putative protein (9.4 kD) (Sfrs9) alternative variant dSep08, mRNA.
Sfrs10	Sfrs10.bSep08	117259	5623	1539	6	147	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (Sfrs10) alternative variant bSep08, mRNA.

Sfrs10	Sfrs10.cSep08	117259	1017	386	2	128	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (Sfrs10) alternative variant cSep08, mRNA.
Sfrs10	Sfrs10.dSep08	117259	11487	737	4	84	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (Sfrs10) alternative variant dSep08, mRNA.
Sfrs11	Sfrs11.aSep08	502603	26559	1798	8	598	splicing factor arginine serine-rich 11 (Sfrs11) alternative variant aSep08, mRNA.
Sfrs11	Sfrs11.bSep08	502603	7740	3194	6	242	splicing factor arginine serine-rich 11 (29.3 kD) (Sfrs11) alternative variant bSep08, mRNA.
Sfrs11	Sfrs11.dSep08	502603	16964	643	6	213	splicing factor arginine serine-rich 11 CRA c (Sfrs11) alternative variant dSep08, mRNA.
Sfrs11	Sfrs11.eSep08	502603	4166	1043	5	211	splicing factor arginine serine-rich 11 (Sfrs11) alternative variant eSep08, mRNA.
Sfrs11	Sfrs11.fSep08	502603	10956	899	8	175	putative protein (Sfrs11) alternative variant fSep08, mRNA.
Sfrs11	Sfrs11.gSep08	502603	8324	1286	5	137	splicing factor arginine serine-rich 11 (14.5 kD) (Sfrs11) alternative variant gSep08, mRNA.
Sfrs11	Sfrs11.hSep08	502603	2263	1114	3	112	putative protein (Sfrs11) alternative variant hSep08, mRNA.
Sfrs12	Sfrs12.bSep08	56763	5576	2234	4	501	splicing factor arginine serine-rich 12 (Sfrs12) alternative variant bSep08, mRNA.
Sfrs12	Sfrs12.cSep08	56763	15788	3135	6	320	splicing factor arginine serine-rich 12 CRA b (38.4 kD) (Sfrs12) alternative variant cSep08, mRNA.
Sfrs12	Sfrs12.eSep08	56763	6183	586	4	83	putative secreted or extracellular protein precursor (8.7 kD) (Sfrs12) alternative variant eSep08, mRNA.
Sfrs12	Sfrs12.fSep08	56763	1385	244	2	78	splicing factor arginine serine-rich 12 (Sfrs12) alternative variant fSep08, mRNA.
Sfrs12	Sfrs12.gSep08	56763	17672	714	3	62	splicing factor arginine serine-rich 12 (Sfrs12) alternative variant gSep08, mRNA.
Sfrs12ip1	Sfrs12ip1.bSep08	361888	4660	724	1	116	SFRS12-interacting protein 1 (13.9 kD) (Sfrs12ip1) alternative variant bSep08, mRNA.
Sfrs14	Sfrs14.bSep08	361126	8921	1798	4	565	splicing factor, arginine/serine-rich 14 (Sfrs14) alternative variant bSep08, mRNA.
Sfrs14	Sfrs14.cSep08	361126	9624	884	3	294	splicing factor, arginine/serine-rich 14 (Sfrs14) alternative variant cSep08, mRNA.
Sfrs14	Sfrs14.dSep08	361126	2957	1058	4	98	splicing factor, arginine/serine-rich 14 (Sfrs14) alternative variant dSep08, mRNA.
Sfrs14	Sfrs14.fSep08	361126	2582	424	3	41	splicing factor, arginine/serine-rich 14 (4.4 kD) (Sfrs14) alternative variant fSep08, mRNA.
Sfrs15	Sfrs15.bSep08	245924	2517	413	4	137	splicing factor, arginine/serine-rich 15 (Sfrs15) alternative variant bSep08, mRNA.
Sfrs15	Sfrs15.cSep08	245924	4162	1831	3	74	splicing factor, arginine/serine-rich 15 (Sfrs15) alternative variant cSep08, mRNA.
Sfrs16	Sfrs16.bSep08	499390	33406	2483	21	471	splicing factor arginine serine-rich 16 CRA b (53.5 kD) (Sfrs16) alternative variant bSep08, mRNA.
Sfrs16	Sfrs16.cSep08	499390	8586	1084	4	250	reticuloendotheliosis viral oncogene (Sfrs16) alternative variant cSep08, mRNA.

Sfrs16	Sfrs16.dSep08	499390	4837	849	6	197	splicing factor arginine serine-rich 16 (Sfrs16) alternative variant dSep08, mRNA.
Sfrs16	Sfrs16.eSep08	499390	4063	716	5	131	splicing factor arginine serine-rich 16 (Sfrs16) alternative variant eSep08, mRNA.
Sfrs16	Sfrs16.fSep08	499390	1880	790	2	128	putative secreted or extracellular protein precursor (13.5 kD) (Sfrs16) alternative variant fSep08, complete mRNA.
Sfrs16	Sfrs16.gSep08	499390	1199	662	2	116	putative protein (11.7 kD) (Sfrs16) alternative variant gSep08, mRNA.
Sfrs16	Sfrs16.hSep08	499390	4979	1156	8	101	splicing factor arginine serine-rich 16 (12.7 kD) (Sfrs16) alternative variant hSep08, mRNA.
Sfrs18	Sfrs18.bSep08	297942	26896	4356	11	805	splicing factor, arginine/serine-rich 18 (92.4 kD) (Sfrs18) alternative variant bSep08, mRNA.
Sfrs18	Sfrs18.cSep08	297942	4388	2546	3	522	splicing factor, arginine/serine-rich 18 (Sfrs18) alternative variant cSep08, mRNA.
Sfrs18	Sfrs18.dSep08	297942	2107	620	2	123	splicing factor, arginine/serine-rich 18 (Sfrs18) alternative variant dSep08, mRNA.
Sft2d1	Sft2d1.dSep08	292305	16268	1365	9	77	SFT2-like (8.9 kD) (Sft2d1) alternative variant dSep08, mRNA.
Sftpa1	Sftpa1.aSep08	24773	3543	1650	5	280	surfactant, pulmonary-associated protein A1 (Sftpa1) alternative variant aSep08, mRNA.
Sftpa1	Sftpa1.bSep08	24773	2671	775	5	250	surfactant, pulmonary-associated protein A1 (Sftpa1) alternative variant bSep08, mRNA.
Sftpa1	Sftpa1.cSep08	24773	2632	950	5	228	surfactant, pulmonary-associated protein A1 (Sftpa1) alternative variant cSep08, mRNA.
Sftpa1	Sftpa1.eSep08	24773	1865	671	2	151	surfactant, pulmonary-associated protein A1 (Sftpa1) alternative variant eSep08, mRNA.
Sftpb	Sftpb.bSep08	192155	5315	839	8	278	surfactant associated protein B (Sftpb) alternative variant bSep08, mRNA.
Sftpb	Sftpb.cSep08	192155	2949	761	3	189	surfactant associated protein B (Sftpb) alternative variant cSep08, mRNA.
Sftpb	Sftpb.dSep08	192155	1884	712	5	189	surfactant associated protein B (Sftpb) alternative variant dSep08, mRNA.
Sftpc	Sftpc.bSep08	50683	2944	672	5	157	surfactant associated protein C (16.9 kD) (Sftpc) alternative variant bSep08, complete mRNA.
Sftpc	Sftpc.cSep08	50683	969	749	2	76	surfactant associated protein C (Sftpc) alternative variant cSep08, mRNA.
Sfxn1	Sfxn1.aSep08	364678	24455	893		297	sideroflexin 1 (Sfxn1) mRNA.
Sfxn2	Sfxn2.aSep08	294011	26339	2971	12	322	sideroflexin 2 CRA a (36.2 kD) (Sfxn2) alternative variant aSep08, complete mRNA.
Sfxn2	Sfxn2.bSep08	294011	3374	1154	5	238	sideroflexin 2 CRA a (Sfxn2) alternative variant bSep08, mRNA.
Sfxn2	Sfxn2.cSep08	294011	1360	674	2	162	sideroflexin 2 CRA a (17.4 kD) (Sfxn2) alternative variant cSep08, mRNA.
Sfxn2	Sfxn2.eSep08	294011	10907	732	5	127	sideroflexin 2 CRA a (14.3 kD) (Sfxn2) alternative variant eSep08, mRNA.
Sfxn3	Sfxn3.bSep08	65042	4497	752	7	250	sideroflexin 3 (Sfxn3) alternative variant bSep08, mRNA.
Sfxn3	Sfxn3.cSep08	65042	2408	393	4	107	sideroflexin 3 (Sfxn3) alternative variant cSep08, mRNA.



Sfxn4	Sfxn4.aSep08	361778	13600	686	5	141	sideroflexin 4 (Sfxn4) alternative variant aSep08, mRNA.
Sfxn4	Sfxn4.bSep08	361778	5962	877	3	112	sideroflexin 4 (13.0 kD) (Sfxn4) alternative variant bSep08, mRNA.
Sfxn5	Sfxn5.bSep08	261737	54568	3272	7	208	sideroflexin 5 (22.3 kD) (Sfxn5) alternative variant bSep08, mRNA.
Sfxn5	Sfxn5.dSep08	261737	44353	238	4	79	sideroflexin 5 (Sfxn5) alternative variant dSep08, mRNA.
Sgca	Sgca.bSep08	303468	10476	1306	10	331	sarcoglycan, alpha (dystrophin-associated glycoprotein) (37.0 kD) (Sgca) alternative variant bSep08, complete mRNA.
Sgcb	Sgcb.aSep08	680229	15011	3725	1	320	sarcoglycan, beta (dystrophin-associated glycoprotein) (35.0 kD) (Sgcb) alternative variant aSep08, complete mRNA.
Sgcb	Sgcb.bSep08	680229	8665	807	1	206	sarcoglycan, beta (dystrophin-associated glycoprotein) (Sgcb) alternative variant bSep08, mRNA.
Sgcd	Sgcd.cSep08	497892	101479	587		123	sarcoglycan, delta (dystrophin-associated glycoprotein) (Sgcd) alternative variant cSep08, mRNA.
Sgcg	Sgcg.bSep08	305941	11524	1214	5	161	sarcoglycan, gamma (dystrophin-associated glycoprotein) (17.2 kD) (Sgcg) alternative variant bSep08, mRNA.
Sgef	Sgef.aSep08	310460	49545	514		171	src homology 3 domain-containing guanine nucleotide exchange factor (Sgef) mRNA.
Sgip1	Sgip1.aSep08	313413	35115	1076	6	322	SH3-domain GRB2-like (endophilin) interacting protein 1 (Sgip1) alternative variant aSep08, mRNA.
Sgip1	Sgip1.bSep08	313413	106424	875	7	175	SH3-domain GRB2-like (endophilin) interacting protein 1 (19.3 kD) (Sgip1) alternative variant bSep08, mRNA.
Sgip1	Sgip1.cSep08	313413	106081	460	6	87	SH3-domain GRB2-like (endophilin) interacting protein 1 (Sgip1) alternative variant cSep08, mRNA.
Sgip1	Sgip1.dSep08	313413	8116	387	2	32	SH3-domain GRB2-like (endophilin) interacting protein 1 (Sgip1) alternative variant dSep08, mRNA.
Sgk1	Sgk1.bSep08	29517	2628	793	7	229	serum/glucocorticoid regulated kinase 1 (Sgk1) alternative variant bSep08, mRNA.
Sgk1	Sgk1.dSep08	29517	3476	399	4	90	serum/glucocorticoid regulated kinase 1 (Sgk1) alternative variant dSep08, mRNA.
Sgk2	Sgk2.aSep08	171497	4860	745		66	serum/glucocorticoid regulated kinase 2 (Sgk2) mRNA.
Sgms1	Sgms1.bSep08	353229	69936	3279	7	184	cholinephosphotransferase 1 (21.9 kD) (Sgms1) alternative variant bSep08, mRNA.
Sgms1	Sgms1.cSep08	353229	12544	391	4	119	cholinephosphotransferase 1 (Sgms1) alternative variant cSep08, mRNA.
Sgms1	Sgms1.dSep08	353229	4231	370	2	85	putative protein (Sgms1) alternative variant dSep08, mRNA.
Sgms1	Sgms1.eSep08	353229	225535	1220	6	169	cholinephosphotransferase 1 (Sgms1) alternative variant eSep08, mRNA.
Sgms1	Sgms1.fSep08	353229	5985	786	2	90	putative cytoplasmic protein (10.1 kD) (Sgms1) alternative variant fSep08, mRNA.
Sgms1	Sgms1.gSep08	353229	73619	743	4	103	cholinephosphotransferase 1 (Sgms1) alternative variant gSep08, mRNA.
Sgms1	Sgms1.hSep08	353229	165961	606	4	68	putative protein (Sgms1) alternative variant hSep08, mRNA.

Sgms1	Sgms1.iSep08	353229	46960	259	2	58	putative protein (Sgms1) alternative variant iSep08, mRNA.
Sgms2	Sgms2.bSep08	310849	3168	815	2	147	sphingomyelin synthase 2 (Sgms2) alternative variant bSep08, mRNA.
Sgms2	Sgms2.cSep08	310849	2567	727	3	51	sphingomyelin synthase 2 (Sgms2) alternative variant cSep08, mRNA.
Sgol1	Sgol1.aSep08	363174	3666	1495		220	shugoshin-like 1 ( <i>S. pombe</i> ) (Sgol1) mRNA.
Sgol2	Sgol2.aSep08	316425	2314	1047		348	shugoshin-like 2 ( <i>S. pombe</i> ) (Sgol2) mRNA.
Sgpp1	Sgpp1.aSep08	81536	20731	2887		336	sphingosine-1-phosphate phosphatase 1 (Sgpp1) mRNA.
Sgsm1	Sgsm1.bSep08	288743	21328	424	5	130	small G protein signaling modulator 1 (Sgsm1) alternative variant bSep08, mRNA.
Sgsm1	Sgsm1.cSep08	288743	9838	381	5	127	small G protein signaling modulator 1 (Sgsm1) alternative variant cSep08, mRNA.
Sgsm1	Sgsm1.dSep08	288743	7504	381	4	126	small G protein signaling modulator 1 (Sgsm1) alternative variant dSep08, mRNA.
Sgsm1	Sgsm1.eSep08	288743	588	391	2	45	small G protein signaling modulator 1 (4.9 kD) (Sgsm1) alternative variant eSep08, mRNA.
Sgsm2	Sgsm2.bSep08	303304	3474	1165	5	184	small G protein signaling modulator 2 (Sgsm2) alternative variant bSep08, mRNA.
Sgsm3	Sgsm3.bSep08	362963	3311	612	7	204	src homology-3 and variant SH3 (Sgsm3) alternative variant bSep08, mRNA.
Sgsm3	Sgsm3.cSep08	362963	2167	1668	5	127	RUN (14.6 kD) (Sgsm3) alternative variant cSep08, mRNA.
Sgsm3	Sgsm3.dSep08	362963	595	498	2	114	src homology-3 (Sgsm3) alternative variant dSep08, mRNA.
Sgta	Sgta.bSep08	64667	16662	1963	12	315	small glutamine-rich tetratricopeptide repeat (TPR)-containing, alpha (34.3 kD) (Sgta) alternative variant bSep08, complete mRNA.
Sgta	Sgta.cSep08	64667	4196	599	7	174	small glutamine-rich tetratricopeptide repeat (TPR)-containing, alpha (Sgta) alternative variant cSep08, mRNA.
Sgta	Sgta.eSep08	64667	9722	348	4	80	small glutamine-rich tetratricopeptide repeat (TPR)-containing, alpha (Sgta) alternative variant eSep08, mRNA.
Sgta	Sgta.fSep08	64667	438	361	2	62	small glutamine-rich tetratricopeptide repeat (TPR)-containing, alpha (Sgta) alternative variant fSep08, mRNA.
Sgtb	Sgtb.bSep08	294708	10885	3612	5	138	small glutamine-rich tetratricopeptide repeat (TPR)-containing, beta (Sgtb) alternative variant bSep08, mRNA.
SH2.0	SH2.0.aSep08		12177	1289		335	SH2 motif (37.2 kD) (SH2.0) mRNA.
Sh2b3	Sh2b3.bSep08	58838	3241	3053	2	235	SH2B adaptor protein 3 (Sh2b3) alternative variant bSep08, mRNA.
Sh2d1a	Sh2d1a.bSep08	501502	18278	613	4	81	SH2 domain protein 1A (9.1 kD) (Sh2d1a) alternative variant bSep08, mRNA.
Sh2d3c	Sh2d3c.bSep08	362111	29441	2495	12	371	SH2 motif (40.4 kD) (Sh2d3c) alternative variant bSep08, complete mRNA.
Sh2d3c	Sh2d3c.cSep08	362111	5288	1193	4	332	SH2 motif (Sh2d3c) alternative variant cSep08, mRNA.
Sh2d3c	Sh2d3c.dSep08	362111	3744	728	3	236	SH2 motif (Sh2d3c) alternative variant dSep08, mRNA.
Sh2d3c	Sh2d3c.eSep08	362111	5071	691	3	229	SH2 motif (Sh2d3c) alternative variant eSep08, mRNA.
Sh2d4b	Sh2d4b.aSep08	290612	17473	452		141	putative protein, with a coiled coil domain, of mammalian origin (Sh2d4b) mRNA.

Sh2d5	Sh2d5.aSep08	366489	8975	733		244	putative protein of metazoan origin (Sh2d5) mRNA.
Sh3bgr	Sh3bgr.bSep08	498066	24992	2072	8	204	SH3-binding domain glutamic acid-rich protein (22.1 kD) (Sh3bgr) alternative variant bSep08, mRNA.
Sh3bgr	Sh3bgr.cSep08	498066	23786	1010	7	203	SH3-binding domain glutamic acid-rich protein (22.2 kD) (Sh3bgr) alternative variant cSep08, mRNA.
Sh3bgr	Sh3bgr.dSep08	498066	7646	813	2	157	SH3-binding domain glutamic acid-rich protein (Sh3bgr) alternative variant dSep08, mRNA.
Sh3bgrl	Sh3bgrl.aSep08	302363	105957	2926		155	SH3-binding domain glutamic acid-rich protein like (Sh3bgrl) mRNA.
Sh3bgrl2	Sh3bgrl2.aSep08	501026	52461	3442		107	SH3 domain binding glutamic acid-rich protein like 2 (12.3 kD) (Sh3bgrl2) mRNA.
Sh3bp1	Sh3bp1.aSep08	300067	5449	1988		292	SH3-domain binding protein 1 (31.4 kD) (Sh3bp1) mRNA.
Sh3bp2	Sh3bp2.aSep08	305450	12791	2905	1	619	SH3-domain binding protein 2 (Sh3bp2) alternative variant aSep08, mRNA.
Sh3bp2	Sh3bp2.bSep08	305450	29226	415	1	138	SH3-domain binding protein 2 (Sh3bp2) alternative variant bSep08, mRNA.
Sh3bp4	Sh3bp4.bSep08	64634	67914	592	2	197	SH3-domain binding protein 4 (Sh3bp4) alternative variant bSep08, mRNA.
Sh3bp5l	Sh3bp5l.bSep08	690898	1063	601	1	58	SH3 binding domain protein 5 like (6.2 kD) (Sh3bp5l) alternative variant bSep08, mRNA.
Sh3bp5l	Sh3bp5l.cSep08	690898	770	458	1	42	SH3 binding domain protein 5 like (Sh3bp5l) alternative variant cSep08, mRNA.
Sh3d19	Sh3d19.aSep08	295171	19665	3496		430	SH3 domain protein D19 (Sh3d19) mRNA.
Sh3gl2	Sh3gl2.aSep08	116743	12046	1218		215	SH3-domain GRB2-like 2 (Sh3gl2) alternative variant aSep08, mRNA.
Sh3gl2	Sh3gl2.bSep08	116743	30302	462		154	SH3-domain GRB2-like 2 (Sh3gl2) alternative variant bSep08, mRNA.
Sh3gl3	Sh3gl3.aSep08	81921	129992	1526	4	347	SH3-domain GRB2-like 3 (39.2 kD) (Sh3gl3) alternative variant aSep08, complete mRNA.
Sh3gl3	Sh3gl3.cSep08	81921	107287	1099	3	246	SH3-domain GRB2-like 3 (Sh3gl3) alternative variant cSep08, mRNA.
Sh3gl3	Sh3gl3.dSep08	81921	106144	403	1	75	SH3-domain GRB2-like 3 (8.2 kD) (Sh3gl3) alternative variant dSep08, mRNA.
Sh3glb1	Sh3glb1.bSep08	292156	35217	1300	11	351	SH3-domain GRB2-like B1 (endophilin) (Sh3glb1) alternative variant bSep08, mRNA.
Sh3glb1	Sh3glb1.cSep08	292156	13624	668	5	117	SH3-domain GRB2-like B1 (endophilin) (Sh3glb1) alternative variant cSep08, mRNA.
Sh3glb2	Sh3glb2.aSep08	311848	13831	1018	10	323	endophilin B2 CRA a (Sh3glb2) alternative variant aSep08, mRNA.
Sh3glb2	Sh3glb2.bSep08	311848	13843	2502	9	207	endophilin B2 (23.2 kD) (Sh3glb2) alternative variant bSep08, mRNA.
Sh3glb2	Sh3glb2.cSep08	311848	10436	618	6	206	endophilin B2 CRA a (Sh3glb2) alternative variant cSep08, mRNA.
Sh3glb2	Sh3glb2.dSep08	311848	1306	737	3	192	endophilin B2 CRA a (Sh3glb2) alternative variant dSep08, mRNA.
Sh3glb2	Sh3glb2.eSep08	311848	8190	573	6	190	endophilin B2 CRA b (Sh3glb2) alternative variant eSep08, mRNA.

Sh3glb2	Sh3glb2.fSep08	311848	836	478	3	128	endophilin B2 (Sh3glb2) alternative variant fSep08, mRNA.
Sh3glb2	Sh3glb2.hSep08	311848	1470	1231	2	88	endophilin B2 (9.3 kD) (Sh3glb2) alternative variant hSep08, mRNA.
Sh3kbp1	Sh3kbp1.bSep08	84357	56454	813	6	242	kinase binding protein 1 like (Sh3kbp1) alternative variant bSep08, mRNA.
Sh3kbp1	Sh3kbp1.cSep08	84357	49747	710	6	236	kinase binding protein 1 like (Sh3kbp1) alternative variant cSep08, mRNA.
Sh3kbp1	Sh3kbp1.dSep08	84357	113284	614	5	204	kinase binding protein 1 like (Sh3kbp1) alternative variant dSep08, mRNA.
Sh3kbp1	Sh3kbp1.eSep08	84357	30509	884	5	182	kinase binding protein 1 like (19.4 kD) (Sh3kbp1) alternative variant eSep08, mRNA.
Sh3kbp1	Sh3kbp1.fSep08	84357	23613	790	4	180	kinase binding protein 1 like (Sh3kbp1) alternative variant fSep08, mRNA.
Sh3kbp1	Sh3kbp1.gSep08	84357	114885	374	4	79	kinase binding protein 1 like (8.7 kD) (Sh3kbp1) alternative variant gSep08, mRNA.
Sh3md4	Sh3md4.aSep08	294557	91845	768		256	SH3 multiple domains 4 (Sh3md4) mRNA.
Sh3pxd2a	Sh3pxd2a.bSep08	309460	10147	694	3	216	SH3 and PX domains 2A (Sh3pxd2a) alternative variant bSep08, mRNA.
Sh3pxd2a	Sh3pxd2a.cSep08	309460	5052	367	2	57	SH3 and PX domains 2A (Sh3pxd2a) alternative variant cSep08, mRNA.
Sh3rf2	Sh3rf2.bSep08	307472	12407	1015	1	295	putative mitochondrial protein of vertebrate origin (31.3 kD) (Sh3rf2) alternative variant bSep08, mRNA.
Sh3tc1	Sh3tc1.bSep08	305441	20506	649	5	215	SH3 domain and tetratricopeptide repeats 1 (Sh3tc1) alternative variant bSep08, mRNA.
Sh3tc2	Sh3tc2.aSep08	307393	8351	1468	5	271	SH3 domain and tetratricopeptide repeats 2 (Sh3tc2) alternative variant aSep08, mRNA.
Sh3yl1	Sh3yl1.bSep08	362724	20613	590	5	196	sh3 domain YSC-like 1 (Sh3yl1) alternative variant bSep08, mRNA.
Sh3yl1	Sh3yl1.cSep08	362724	18445	732	3	117	sh3 domain YSC-like 1 (13.1 kD) (Sh3yl1) alternative variant cSep08, mRNA.
Sh3yl1	Sh3yl1.dSep08	362724	30043	609	4	35	sh3 domain YSC-like 1 (Sh3yl1) alternative variant dSep08, mRNA.
Sh3yl1	Sh3yl1.eSep08	362724	20777	561	4	45	sh3 domain YSC-like 1 (Sh3yl1) alternative variant eSep08, mRNA.
SH3_1.0	SH3_1.0.aSep08		13971	1108		320	intersectin 1 (SH3_1.0) mRNA.
SH3_1.1	SH3_1.1.aSep08		18739	2319		294	intersectin 1 CRA a (SH3_1.1) alternative variant aSep08, mRNA.
SH3_1.3	SH3_1.3.aSep08		8035	2010		489	protein phosphatase 1 regulatory like (SH3_1.3) mRNA.
SH3_1.4	SH3_1.4.aSep08		8418	3235		465	dynamitin binding protein like (52.1 kD) (SH3_1.4) mRNA.
SH3_2.0	SH3_2.0.aSep08		15536	991	6	244	2 -binding protein like (SH3_2.0) alternative variant aSep08, mRNA.
SH3_2.0	SH3_2.0.bSep08		12824	1121	4	163	2 -binding protein like (SH3_2.0) alternative variant bSep08, mRNA.
SH3_2.0	SH3_2.0.cSep08		12777	601	5	148	2 -binding protein like (SH3_2.0) alternative variant cSep08, mRNA.
SH3_2.1	SH3_2.1.aSep08		57993	639		198	dedicator of 2 (SH3_2.1) mRNA.

SH3_2.2	SH3_2.2.aSep08		8206	1397	1	305	putative protein (SH3_2.2) alternative variant aSep08, mRNA.
SH3_2.2	SH3_2.2.bSep08		7816	1165	1	114	putative mitochondrial protein (12.8 kD) (SH3_2.2) alternative variant bSep08, mRNA.
SH3_2.2	SH3_2.2.cSep08		7479	168	1	54	cysteine rich domain (SH3_2.2) alternative variant cSep08, mRNA.
SH3_2.3	SH3_2.3.aSep08		7631	455		96	melanoma inhibitory activity 2 CRA b like (SH3_2.3) mRNA.
SH3_2.4	SH3_2.4.aSep08		6729	677		180	rho guanine nucleotide exchange factor 5 CRA a (SH3_2.4) alternative variant aSep08, mRNA.
SH3_2.5	SH3_2.5.aSep08		14054	420		127	domain-containing guanine exchange factor (SH3_2.5) mRNA.
SH3_2.6	SH3_2.6.aSep08		6280	470		156	src homology-3 and variant SH3 (SH3_2.6) mRNA.
SH3_2.7	SH3_2.7.aSep08		3577	332		110	intersectin 2 (SH3_2.7) mRNA.
shabor	shabor.aSep08		6700	703		233	axonemal dynein heavy Dnahc8 (shabor) mRNA.
shachy	shachy.aSep08		4614	721		114	putative protein (12.8 kD) (shachy) mRNA.
shadoy	shadoy.aSep08		30475	399	2	91	putative protein (shadoy) alternative variant aSep08, mRNA.
shaflu	shaflu.aSep08		1975	380		126	nucleoporin 98kDa CRA a (shaflu) mRNA.
shafly	shafly.aSep08		3841	426		38	putative protein (4.3 kD) (shafly) alternative variant aSep08, mRNA.
shagar	shagar.aSep08		19977	434		79	pericentriolar material 1 (shagar) mRNA.
shaja	shaja.aSep08		66734	409		38	putative protein (4.5 kD) (shaja) mRNA.
shajey	shajey.aSep08		6676	469		37	putative protein (4.1 kD) (shajey) mRNA.
shakee	shakee.aSep08		11723	531		176	tumor protein p53 binding 2 like (shakee) mRNA.
shakler	shakler.aSep08		13421	665		43	putative protein (shakler) mRNA.
shalo	shalo.aSep08		5259	733		45	putative protein (5.2 kD) (shalo) mRNA.
shamee	shamee.aSep08		1590	503		167	myosin heavy chain polypeptide 1 skeletal muscle adult (shamee) mRNA.
Shank1	Shank1.aSep08	78957	3544	1408		469	SH3/ankyrin domain gene 1 (Shank1) mRNA.
Shank3	Shank3.bSep08	59312	8172	308	1	102	SH3/ankyrin domain gene 3 (Shank3) alternative variant bSep08, mRNA.
shanoy	shanoy.aSep08		2849	740		24	putative protein (2.7 kD) (shanoy) mRNA.
shapor	shapor.aSep08		2102	193		63	topoisomerase II binding protein 1 like (shapor) mRNA.
sharbor	sharbor.bSep08		777	612	3	81	putative protein (sharbor) alternative variant bSep08, mRNA.
sharchy	sharchy.aSep08		1222	404	1	108	putative protein (sharchy) alternative variant aSep08, mRNA.
sharchy	sharchy.bSep08		11163	374	3	87	putative protein (sharchy) alternative variant bSep08, mRNA.
shardoy	shardoy.aSep08		1069	514		104	putative protein (shardoy) mRNA.
sharflu	sharflu.aSep08		2869	349		116	nucleoporin 98 (sharflu) mRNA.
sharfly	sharfly.aSep08		36043	749		50	atpase protein like (sharfly) mRNA.
shargar	shargar.aSep08		7656	489		61	putative protein (shargar) mRNA.

sharja	sharja.aSep08		2632	1132		54	putative protein (sharja) mRNA.
sharjey	sharjey.aSep08		4205	387		128	ellis van Creveld (sharjey) mRNA.
sharkee	sharkee.aSep08		3502	704		51	putative protein (5.6 kD) (sharkee) mRNA.
sharlo	sharlo.aSep08		16052	1204	6	338	autism susceptibility candidate 2 (sharlo) alternative variant aSep08, mRNA.
sharmee	sharmee.aSep08		403	298		24	putative protein (2.9 kD) (sharmee) mRNA.
sharnoy	sharnoy.bSep08		3139	363	2	35	putative protein (3.9 kD) (sharnoy) alternative variant bSep08, mRNA.
sharoy	sharoy.aSep08		1173	350		50	putative protein (5.9 kD) (sharoy) mRNA.
Sharpin	Sharpin.bSep08	81859	4277	1774	7	292	SHANK-associated RH domain interacting protein (Sharpin) alternative variant bSep08, mRNA.
Sharpin	Sharpin.cSep08	81859	830	579	2	191	SHANK-associated RH domain interacting protein (Sharpin) alternative variant cSep08, mRNA.
Sharpin	Sharpin.eSep08	81859	2303	335	3	111	SHANK-associated RH domain interacting protein (Sharpin) alternative variant eSep08, mRNA.
sharpor	sharpor.aSep08		16880	959		104	putative protein (11.5 kD) (sharpor) mRNA.
sharroy	sharroy.aSep08		11605	341			
sharsa	sharsa.aSep08		492	409		68	myeloid lymphoid mixed-lineage leukemia 2 CRA c like (sharsa) mRNA.
sharshee	sharshee.aSep08		15368	449		31	putative protein (3.8 kD) (sharshee) mRNA.
shartu	shartu.aSep08		6324	439		145	heat 5a (shartu) mRNA.
sharvo	sharvo.aSep08		856	250		82	putative protein (sharvo) mRNA.
sharwer	sharwer.aSep08		1704	337	1	63	putative protein (sharwer) alternative variant aSep08, mRNA.
sharwer	sharwer.bSep08		9653	597	4	33	putative protein (sharwer) alternative variant bSep08, mRNA.
sharwey	sharwey.aSep08		27016	788		203	lymphoid-restricted membrane protein like (sharwey) mRNA.
shasa	shasa.aSep08		498	227		62	putative protein (7.0 kD) (shasa) mRNA.
shashee	shashee.aSep08		3785	383		111	putative protein (shashee) mRNA.
shatu	shatu.bSep08		2011	568	2	49	putative protein (shatu) alternative variant bSep08, mRNA.
shavar	shavar.aSep08		3644	776		27	putative protein (shavar) mRNA.
shavo	shavo.aSep08		52215	316		80	putative protein (shavo) mRNA.
shawbor	shawbor.aSep08		886	237		78	putative protein (shawbor) mRNA.
shawchy	shawchy.aSep08		21529	651		216	CRA a (shawchy) mRNA.
shawdoy	shawdoy.aSep08		2652	1006		88	putative protein (shawdoy) alternative variant aSep08, mRNA.
shawdoy	shawdoy.bSep08		1954	309		88	putative protein (shawdoy) alternative variant bSep08, mRNA.
shawer	shawer.aSep08		62792	431		93	putative protein of vertebrate origin (shawer) mRNA.
shawey	shawey.aSep08		4863	753		136	organic anion transporter (shawey) mRNA.
shawflu	shawflu.aSep08		1151	556		30	putative protein (3.5 kD) (shawflu) mRNA.
shawfly	shawfly.aSep08		9708	830		36	putative protein (shawfly) mRNA.
shawgar	shawgar.aSep08		2507	722		83	putative protein (9.7 kD) (shawgar) mRNA.

shawja	shawja.aSep08		9427	429		143	homeobox containing 1 CRA a (shawja) mRNA.
shawjey	shawjey.aSep08		7573	746		97	janus kinase microtubule interacting protein 1 (shawjey) mRNA.
shawkee	shawkee.aSep08		5936	661		117	putative protein (shawkee) mRNA.
shawlo	shawlo.aSep08		76460	361		40	putative protein (shawlo) mRNA.
shawmee	shawmee.aSep08		1265	448		67	putative protein (7.5 kD) (shawmee) mRNA.
shawnoy	shawnoy.aSep08		375	261		43	putative protein (shawnoy) mRNA.
shawpor	shawpor.aSep08		5528	448		148	nephrocystin 3 (shawpor) mRNA.
shawroy	shawroy.bSep08		1417	271	2	41	putative protein (4.6 kD) (shawroy) alternative variant bSep08, mRNA.
shawsa	shawsa.aSep08		630	351		116	myeloid lymphoid mixed-lineage leukemia 2 like (shawsa) mRNA.
shawshee	shawshee.aSep08		26360	541	3	48	putative protein (shawshee) alternative variant aSep08, mRNA.
shawshee	shawshee.bSep08		944	317	2	27	putative protein (3.2 kD) (shawshee) alternative variant bSep08, mRNA.
shawtu	shawtu.aSep08		11984	552		113	heat 5A (12.8 kD) (shawtu) mRNA.
shawvo	shawvo.aSep08		307	205		20	putative protein (shawvo) mRNA.
shawwer	shawwer.aSep08		21531	2862	3	410	serine kinase (shawwer) alternative variant aSep08, mRNA.
shawwer	shawwer.bSep08		6337	522	2	91	serine kinase (shawwer) alternative variant bSep08, mRNA.
shawwey	shawwey.aSep08		1038	611		104	putative protein (shawwey) mRNA.
Shbg	Shbg.bSep08	24775	17269	1457	7	267	sex -binding like (29.1 kD) (Shbg) alternative variant bSep08, complete mRNA.
Shbg	Shbg.cSep08	24775	14978	761	4	155	sex hormone binding globulin like (17.2 kD) (Shbg) alternative variant cSep08, mRNA.
Shbg	Shbg.dSep08	24775	14951	729	4	155	sex hormone binding globulin like (17.2 kD) (Shbg) alternative variant dSep08, mRNA.
Shc1	Shc1.bSep08	85385	2128	956	4	248	src homology 2 domain-containing transforming protein C1 (Shc1) alternative variant bSep08, mRNA.
Shc1	Shc1.cSep08	85385	3815	2558	4	148	src homology 2 domain-containing transforming protein C1 (Shc1) alternative variant cSep08, mRNA.
Shc2	Shc2.bSep08	314612	1483	545	2	36	src homology 2 domain-containing transforming protein C2 (3.7 kD) (Shc2) alternative variant bSep08, mRNA.
Shc3	Shc3.bSep08	114858	90933	408		69	src homology 2 domain-containing transforming protein C3 (Shc3) alternative variant bSep08, mRNA.
Shcbp1	Shcbp1.aSep08	364648	29908	607	4	201	shc SH2-domain binding protein 1 (Shcbp1) alternative variant aSep08, mRNA.
Shcbp1	Shcbp1.bSep08	364648	3356	405	1	118	shc SH2-domain binding protein 1 (Shcbp1) alternative variant bSep08, mRNA.
Shd	Shd.bSep08	316507	1813	1017	3	154	src homology 2 domain-containing transforming protein D (17.4 kD) (Shd) alternative variant bSep08, mRNA.
She	She.aSep08	685088	14964	382		127	src homology 2 domain-containing transforming protein E (She) mRNA.
sheebor	sheebor.aSep08		22683	465		64	putative protein (sheebor) mRNA.

sheechy	sheechy.aSep08		3340	683		25	putative protein (sheechy) mRNA.
sheedoy	sheedoy.aSep08		47102	525	3	108	putative protein (sheedoy) alternative variant aSep08, mRNA.
sheedoy	sheedoy.bSep08		59140	1550	3	101	putative nuclear protein (11.1 kD) (sheedoy) alternative variant bSep08, mRNA.
sheedoy	sheedoy.cSep08		2448	524	2	65	CRA c like (7.4 kD) (sheedoy) alternative variant cSep08, mRNA.
sheeflu	sheeflu.aSep08		951	329		109	putative protein, with 2 coiled coil domains, of mammalian origin (sheeflu) mRNA.
sheefly	sheefly.aSep08		2643	440		146	nuclear receptor coactivator 7 CRA a (sheefly) mRNA.
sheegar	sheegar.aSep08		2492	351		116	putative protein of metazoan origin (sheegar) mRNA.
sheeja	sheeja.aSep08		65699	338		42	putative protein (4.5 kD) (sheeja) mRNA.
sheejey	sheejey.bSep08		3011	2687	1	70	putative protein (7.7 kD) (sheejey) alternative variant bSep08, mRNA.
sheekee	sheekee.aSep08		66650	541		107	putative protein (12.2 kD) (sheekee) mRNA.
sheelo	sheelo.aSep08		8219	479		159	2 -binding protein like (sheelo) mRNA.
sheemee	sheemee.bSep08		4467	419	3	36	putative protein (sheemee) alternative variant bSep08, mRNA.
sheenoy	sheenoy.aSep08		8411	816			
sheepor	sheepor.aSep08		2833	586		20	putative protein (sheepor) mRNA.
sheeroy	sheeroy.aSep08		2208	562		66	putative protein (7.2 kD) (sheeroy) mRNA.
sheesa	sheesa.aSep08		1163	424		141	myeloid lymphoid mixed-lineage leukemia 2 CRA a like (sheesa) mRNA.
sheeshee	sheeshee.aSep08		7900	303		94	putative protein (sheeshee) mRNA.
sheetu	sheetu.aSep08		638	194		51	neuronal PAS domain protein 3 (sheetu) mRNA.
sheevo	sheevo.aSep08		4232	431	6	79	putative protein (sheevo) alternative variant aSep08, mRNA.
sheewer	sheewer.aSep08		2836	1034	4	344	myeloid lymphoid mixed-lineage leukemia 5 like (sheewer) alternative variant aSep08, mRNA.
sheewer	sheewer.bSep08		871	206	2	68	myeloid lymphoid mixed-lineage leukemia 5 CRA d like (sheewer) alternative variant bSep08, mRNA.
sheewer	sheewer.cSep08		467	360	2	45	putative protein (5.4 kD) (sheewer) alternative variant cSep08, mRNA.
sheewey	sheewey.aSep08		2923	2878		51	putative protein (5.7 kD) (sheewey) mRNA.
sherbor	sherbor.aSep08		7982	737		245	protein particle complex (sherbor) mRNA.
sherchy	sherchy.aSep08		560	403		48	putative protein (5.3 kD) (sherchy) mRNA.
sherdoy	sherdoy.aSep08		7104	317	1	66	putative protein (sherdoy) alternative variant aSep08, mRNA.
sherdoy	sherdoy.bSep08		9906	322	2	35	putative protein (sherdoy) alternative variant bSep08, mRNA.
sherflu	sherflu.aSep08		69392	241		80	putative protein (sherflu) mRNA.
sherfly	sherfly.aSep08		31012	257		49	putative protein (sherfly) mRNA.
shergar	shergar.aSep08		9989	359		46	putative protein (shergar) mRNA.
sherja	sherja.aSep08		2025	710		64	putative protein (7.1 kD) (sherja) mRNA.



sherjey	sherjey.aSep08		14014	262		57	putative protein (sherjey) mRNA.
sherkee	sherkee.aSep08		7074	947	2	147	putative cytoplasmic protein (16.0 kD) (sherkee) alternative variant aSep08, mRNA.
sherlo	sherlo.aSep08		893	353		61	putative protein (6.7 kD) (sherlo) mRNA.
shermee	shermee.aSep08		1473	326		55	putative protein (shermee) mRNA.
shernoy	shernoy.aSep08		2247	848		145	helicase 1 (shernoy) mRNA.
sherpor	sherpor.aSep08		1234	501		85	putative protein (sherpor) mRNA.
sherroy	sherroy.aSep08		3526	1142		250	putative protein (sherroy) alternative variant aSep08, mRNA.
shersa	shersa.aSep08		271	182		60	putative protein (shersa) mRNA.
shershee	shershee.aSep08		632	338		41	putative protein (shershee) mRNA.
shertu	shertu.aSep08		515	230		68	putative protein (shertu) mRNA.
shervo	shervo.aSep08		1381	533		81	putative protein (9.4 kD) (shervo) mRNA.
sherwer	sherwer.aSep08		3541	613		60	ac2-154 like (sherwer) mRNA.
sherwey	sherwey.aSep08		31052	387		129	inositol 1 receptor 2 (sherwey) mRNA.
sheybor	sheybor.aSep08		3132	283		93	protein particle complex (sheybor) mRNA.
sheychy	sheychy.aSep08		32534	346		43	putative protein (sheychy) mRNA.
sheydoy	sheydoy.aSep08		11348	755		42	putative protein (sheydoy) mRNA.
sheyflu	sheyflu.aSep08		14187	814		271	suppression of tumorigenicity 5 like (sheyflu) mRNA.
sheyfly	sheyfly.aSep08		1823	468	2	109	putative protein (sheyfly) alternative variant aSep08, mRNA.
sheygar	sheygar.aSep08		17769	784	3	164	putative mitochondrial protein (18.9 kD) (sheygar) alternative variant aSep08, mRNA.
sheygar	sheygar.bSep08		21239	1355	2	164	putative mitochondrial protein (18.9 kD) (sheygar) alternative variant bSep08, mRNA.
sheyja	sheyja.aSep08		16760	312		104	F-box protein 16 (sheyja) mRNA.
sheyjey	sheyjey.aSep08		8173	705		140	regulator of G-protein signaling 12 CRA a (sheyjey) mRNA.
sheykee	sheykee.aSep08		1679	749		54	putative protein (sheykee) mRNA.
sheylo	sheylo.aSep08		1065	365		67	putative protein (7.8 kD) (sheylo) mRNA.
sheymee	sheymee.bSep08		28942	571	4	61	putative protein (sheymee) alternative variant bSep08, mRNA.
sheymee	sheymee.cSep08		19220	504	4	31	putative protein (3.3 kD) (sheymee) alternative variant cSep08, mRNA.
sheynoy	sheynoy.aSep08		6582	695		81	putative protein (sheynoy) mRNA.
sheypor	sheypor.aSep08		11171	600		44	ATPase (sheypor) mRNA.
sheyroy	sheyroy.aSep08		2383	1784		118	putative protein (sheyroy) mRNA.
sheysa	sheysa.aSep08		5381	784		187	myeloid lymphoid mixed-lineage leukemia 2 CRA c like (sheysa) mRNA.
sheyshee	sheyshee.aSep08		1416	881		48	putative protein (5.1 kD) (sheyshee) mRNA.
sheytu	sheytu.aSep08		5072	831		276	bromodomain adjacent zinc finger domain 1A (sheytu) mRNA.
sheyvo	sheyvo.aSep08		824	331		48	putative protein (5.7 kD) (sheyvo) mRNA.
sheywer	sheywer.aSep08		2634	255		74	putative protein (sheywer) mRNA.

sheywey	sheywey.aSep08		3127	658		219	inositol 1 receptor 2 (sheywey) mRNA.
Shf	Shf.aSep08	362205	6045	849	2	177	putative protein of eukaryotic origin (Shf) alternative variant aSep08, mRNA.
Shf	Shf.bSep08	362205	2138	707	1	130	putative protein of eukaryotic origin (Shf) alternative variant bSep08, mRNA.
Shh	Shh.bSep08	29499	1486	1086	2	292	sonic hedgehog (Shh) alternative variant bSep08, mRNA.
Shisa4	Shisa4.bSep08	360848	2437	382	3	120	shisa homolog 4 (Xenopus laevis) (Shisa4) alternative variant bSep08, mRNA.
Shkbp1	Shkbp1.aSep08	292735	13435	2322	18	711	sh3kbp1 binding protein 1 like (Shkbp1) alternative variant aSep08, mRNA.
Shkbp1	Shkbp1.bSep08	292735	4796	897	4	166	binding protein 1 like (Shkbp1) alternative variant bSep08, mRNA.
Shkbp1	Shkbp1.cSep08	292735	1250	1153	2	106	seta binding protein 1 like (11.1 kD) (Shkbp1) alternative variant cSep08, mRNA.
Shmt1	Shmt1.bSep08	287379	22341	2792	7	468	serine hydroxymethyltransferase 1 (soluble) (Shmt1) alternative variant bSep08, mRNA.
Shmt1	Shmt1.cSep08	287379	9123	1341	8	352	serine hydroxymethyltransferase 1 (soluble) (Shmt1) alternative variant cSep08, mRNA.
Shmt1	Shmt1.dSep08	287379	14147	897	7	229	serine hydroxymethyltransferase 1 (soluble) (Shmt1) alternative variant dSep08, mRNA.
Shmt1	Shmt1.eSep08	287379	13164	813	6	194	serine hydroxymethyltransferase 1 (soluble) (Shmt1) alternative variant eSep08, mRNA.
Shmt2	Shmt2.bSep08	299857	3176	842	7	265	serine hydroxymethyltransferase 2 (mitochondrial) (Shmt2) alternative variant bSep08, mRNA.
Shmt2	Shmt2.cSep08	299857	3543	1515	7	257	serine hydroxymethyltransferase 2 (mitochondrial) (Shmt2) alternative variant cSep08, mRNA.
Shmt2	Shmt2.dSep08	299857	2453	1340	6	256	serine hydroxymethyltransferase 2 (mitochondrial) (Shmt2) alternative variant dSep08, mRNA.
Shmt2	Shmt2.eSep08	299857	700	524	3	107	serine hydroxymethyltransferase 2 (mitochondrial) (Shmt2) alternative variant eSep08, mRNA.
shobor	shobor.aSep08		4328	622		39	putative protein (4.6 kD) (shobor) mRNA.
Shoc2	Shoc2.bSep08	309548	37756	342	1	14	putative protein (1.8 kD) (Shoc2) alternative variant bSep08, mRNA.
shochy	shochy.aSep08		41433	880		248	putative protein of metazoan origin (shochy) mRNA.
shodoy	shodoy.aSep08		1405	691		43	putative protein (5.1 kD) (shodoy) mRNA.
shoflu	shoflu.aSep08		2403	571	3	166	nucleoporin 98kDa (shoflu) alternative variant aSep08, mRNA.
shofly	shofly.aSep08		2756	656		86	putative protein (9.7 kD) (shofly) mRNA.
shogar	shogar.aSep08		10235	714		57	putative protein (6.8 kD) (shogar) mRNA.
shoja	shoja.aSep08		650	531		74	putative mitochondrial protein (8.0 kD) (shoja) mRNA.
shojey	shojey.aSep08		5655	442		41	putative protein (4.7 kD) (shojey) mRNA.
shokee	shokee.aSep08		1042	371		57	putative protein (shokee) mRNA.
sholo	sholo.aSep08		11238	598		140	autism susceptibility candidate 2 (sholo) mRNA.
shomee	shomee.aSep08		1467	800		116	putative protein (5.7 kD) (shomee) complete mRNA.
shonoy	shonoy.aSep08		5230	765		96	putative protein (10.6 kD) (shonoy) mRNA.

shopor	shopor.aSep08		11117	469		93	putative protein (shopor) mRNA.
shorbor	shorbor.aSep08		400	274		60	putative protein (6.6 kD) (shorbor) mRNA.
shorchy	shorchy.aSep08		3436	601		71	putative protein (shorchy) mRNA.
shordoy	shordoy.aSep08		539	373		124	CRA b (shordoy) mRNA.
shorflu	shorflu.aSep08		2863	1266		123	nuclear receptor interacting protein 3 (12.8 kD) (shorflu) mRNA.
shorfly	shorfly.aSep08		6050	1713		490	CRA b (shorfly) mRNA.
shorgar	shorgar.aSep08		2639	410		56	putative protein (6.4 kD) (shorgar) mRNA.
shorja	shorja.aSep08		7343	902		62	putative protein (6.9 kD) (shorja) mRNA.
shorjey	shorjey.aSep08		3540	920		113	putative membrane protein of metazoan origin (12.6 kD) (shorjey) mRNA.
shorkee	shorkee.aSep08		9650	1693	8	327	CRA a (shorkee) alternative variant aSep08, mRNA.
shorlo	shorlo.aSep08		12539	763		82	putative protein (shorlo) mRNA.
shormee	shormee.aSep08		1796	464		100	dynein axonemal heavy (shormee) mRNA.
shornoy	shornoy.aSep08		370	261		43	putative protein (shornoy) mRNA.
shoroy	shoroy.aSep08		3579	352		40	putative protein (shoroy) mRNA.
shorpor	shorpor.aSep08		9508	490		64	putative protein (shorpor) mRNA.
shorroy	shorroy.aSep08		2622	715	4	148	zinc finger RNA binding protein 2 like (shorroy) alternative variant aSep08, mRNA.
shorroy	shorroy.cSep08		2286	644	4	19	putative protein (2.3 kD) (shorroy) alternative variant cSep08, mRNA.
shorsa	shorsa.aSep08		23710	259	3	43	putative protein (shorsa) alternative variant aSep08, mRNA.
shorsa	shorsa.bSep08		7670	1055	2	81	putative protein (8.8 kD) (shorsa) alternative variant bSep08, mRNA.
shorshee	shorshee.aSep08		1023	930		106	putative protein (shorshee) mRNA.
shortu	shortu.aSep08		7762	494		164	bromodomain adjacent zinc finger domain 1A (shortu) mRNA.
shorvo	shorvo.aSep08		22052	482		160	aspartate-beta-hydroxylase CRA a (shorvo) mRNA.
shorwer	shorwer.aSep08		1380	254		35	putative protein (shorwer) mRNA.
shorwey	shorwey.aSep08		69098	398		79	inositol 1 receptor (8.7 kD) (shorwey) mRNA.
shosa	shosa.aSep08		1177	694		59	putative protein (shosa) mRNA.
shoshee	shoshee.aSep08		22487	401		56	putative protein (6.3 kD) (shoshee) mRNA.
shotu	shotu.aSep08		30351	351		26	putative protein (shotu) mRNA.
shovar	shovar.aSep08		2742	766		62	putative protein (7.0 kD) (shovar) mRNA.
shovo	shovo.aSep08		21962	788		74	putative cytoplasmic protein (8.8 kD) (shovo) mRNA.
shower	shower.aSep08		7166	684		227	pseudouridylate synthase 7 homolog (shower) mRNA.
showey	showey.aSep08		7359	650		88	putative protein (showey) mRNA.
Shox2	Shox2.bSep08	25546	8640	1564	1	235	short stature homeobox 2 (Shox2) alternative variant bSep08, mRNA.
shoybor	shoybor.aSep08		2840	1213		62	putative protein (shoybor) mRNA.
shoychy	shoychy.aSep08		10091	771		40	putative protein (4.1 kD) (shoychy) mRNA.
shoydoy	shoydoy.aSep08		9809	535		54	putative protein (shoydoy) mRNA.

shoyflu	shoyflu.aSep08		5923	307		49	wee 1 homolog like (shoyflu) mRNA.
shoyfly	shoyfly.aSep08		1173	262		87	solute carrier family 12 member 7 (shoyfly) mRNA.
shoygar	shoygar.aSep08		5411	413		75	putative protein (shoygar) mRNA.
shoyja	shoyja.aSep08		4616	635		96	putative protein (shoyja) mRNA.
shoyjey	shoyjey.aSep08		21569	362		47	putative protein (shoyjey) mRNA.
shoykee	shoykee.aSep08		1763	1700		61	putative protein (6.8 kD) (shoykee) mRNA.
shoylo	shoylo.aSep08		1083	411		68	putative protein (7.3 kD) (shoylo) mRNA.
shoymee	shoymee.aSep08		2380	669		223	putative protein of eukaryotic origin (shoymee) mRNA.
shoynoy	shoynoy.aSep08		370	261		43	putative protein (shoynoy) mRNA.
shoypor	shoypor.aSep08		6148	524		46	putative protein (shoypor) mRNA.
shoyroy	shoyroy.aSep08		1552	728		128	CRA b (shoyroy) mRNA.
shoysa	shoysa.aSep08		1265	927		262	putative protein of mammalian origin (shoysa) mRNA.
shoyshee	shoyshee.aSep08		32229	376		124	round spermatid basic protein 1 CRA c (shoyshee) mRNA.
shoytu	shoytu.aSep08		27537	306		102	bromodomain adjacent zinc finger domain 1A (shoytu) mRNA.
shoyvo	shoyvo.aSep08		487	324		45	putative protein (shoyvo) mRNA.
shoywer	shoywer.aSep08		52096	283		21	putative protein (2.5 kD) (shoywer) mRNA.
shoywey	shoywey.aSep08		5919	296		78	uncharacterized protein (shoywey) mRNA.
Shprh	Shprh.bSep08	308282	15579	724	7	241	SNF2 histone linker PHD RING helicase (Shprh) alternative variant bSep08, mRNA.
Shroom1	Shroom1.aSep08	287285	2286	1239		413	shroom family member 1 (Shroom1) mRNA.
Shroom2	Shroom2.bSep08	317435	16713	616	2	152	shroom family member 2 and hypothetical protein LOC685699 (Shroom2) alternative variant bSep08, mRNA.
Shroom2	Shroom2.bSep08	685699	16713	616	2	152	shroom family member 2 and hypothetical protein LOC685699 (Shroom2) alternative variant bSep08, mRNA.
Shroom2	Shroom2.cSep08	317435	16081	582	2	56	shroom family member 2 and hypothetical protein LOC685699 (Shroom2) alternative variant cSep08, mRNA.
Shroom2	Shroom2.cSep08	685699	16081	582	2	56	shroom family member 2 and hypothetical protein LOC685699 (Shroom2) alternative variant cSep08, mRNA.
Shroom3	Shroom3.aSep08	305230	12627	2049	4	340	shroom family member 3 (Shroom3) alternative variant aSep08, mRNA.
Shroom4	Shroom4.aSep08	317391	153799	1482		439	shroom family member 4 (Shroom4) mRNA.
shubor	shubor.aSep08		912	356		38	putative protein (shubor) mRNA.
shuchy	shuchy.aSep08		4446	418		138	putative protein of metazoan origin (shuchy) mRNA.
shudoy	shudoy.aSep08		5030	348		72	CRA b like (shudoy) mRNA.
shuflu	shuflu.aSep08		2856	1019		40	putative protein of mammalian origin (shuflu) mRNA.
shufly	shufly.aSep08		5438	730		75	putative mitochondrial protein (8.3 kD) (shufly) mRNA.
shugar	shugar.aSep08		37488	453		53	CRA b like (5.8 kD) (shugar) mRNA.
shuja	shuja.aSep08		9894	396		31	putative protein (shuja) mRNA.
shujey	shujey.aSep08		8336	423		15	putative protein (shujey) mRNA.
shukee	shukee.aSep08		20888	717	3	84	putative protein (9.4 kD) (shukee) alternative variant aSep08, mRNA.

shukee	shukee.bSep08		1167	292	2	82	putative protein (shukee) alternative variant bSep08, mRNA.
shulo	shulo.aSep08		8826	400		69	putative protein (8.1 kD) (shulo) mRNA.
shumee	shumee.aSep08		2581	440	2	116	myosin heavy chain (shumee) alternative variant aSep08, mRNA.
shumee	shumee.bSep08		2319	273	1	63	myosin heavy chain (shumee) alternative variant bSep08, mRNA.
shunoy	shunoy.aSep08		5567	670		42	putative protein (shunoy) mRNA.
shupor	shupor.aSep08		25264	664		142	transmembrane protein 108 (shupor) mRNA.
shuroy	shuroy.aSep08		4103	951		92	putative mitochondrial protein (10.5 kD) (shuroy) mRNA.
shusa	shusa.aSep08		4826	481		88	putative secreted or extracellular protein precursor (9.6 kD) (shusa) mRNA.
shushee	shushee.aSep08		1156	588		46	putative protein (5.3 kD) (shushee) mRNA.
shutu	shutu.aSep08		18829	411		136	protein kinase D2 (shutu) mRNA.
shuvar	shuvar.bSep08		1023	663	2	111	putative mitochondrial protein (12.1 kD) (shuvar) alternative variant bSep08, mRNA.
shuvo	shuvo.aSep08		12887	666		159	putative protein of metazoan origin (shuvo) mRNA.
shuwer	shuwer.aSep08		38955	974		319	CRA b (shuwer) alternative variant aSep08, mRNA.
shuwey	shuwey.aSep08		6876	493		35	putative protein (4.1 kD) (shuwey) mRNA.
shybor	shybor.aSep08		1245	485		67	putative protein (7.4 kD) (shybor) mRNA.
shychy	shychy.aSep08		2214	669	2	154	putative protein (shychy) alternative variant aSep08, mRNA.
shydoy	shydoy.aSep08		914	332		33	putative protein (3.5 kD) (shydoy) mRNA.
shyflu	shyflu.aSep08		7197	368		122	nucleoporin 98kDa (shyflu) mRNA.
shyfly	shyfly.aSep08		1495	258		85	putative protein (shyfly) mRNA.
shygar	shygar.aSep08		11623	602		40	putative protein (4.7 kD) (shygar) mRNA.
shyja	shyja.aSep08		15636	593	1	101	putative mitochondrial protein (11.3 kD) (shyja) alternative variant aSep08, mRNA.
shyja	shyja.bSep08		15746	656	2	57	putative protein (6.3 kD) (shyja) alternative variant bSep08, mRNA.
shyjej	shyjej.aSep08		1737	835		23	putative protein (2.6 kD) (shyjej) mRNA.
shykee	shykee.aSep08		4305	750		249	tumor protein p53 binding 2 like (shykee) mRNA.
shykler	shykler.aSep08		1716	725		61	putative protein (shykler) mRNA.
shymee	shymee.aSep08		1752	329		109	myosin heavy chain skeletal muscle (shymee) mRNA.
shynoy	shynoy.aSep08		895	659		64	putative protein (shynoy) mRNA.
shypor	shypor.aSep08		29523	663		220	phakinin (shypor) mRNA.
shyroj	shyroj.aSep08		8934	382		82	reverse transcriptase (shyroj) mRNA.
shysa	shysa.aSep08		762	408		8	putative protein (0.9 kD) (shysa) mRNA.
shyshee	shyshee.aSep08		637	550		88	zinc finger DHHC-type containing 3 (9.8 kD) (shyshee) mRNA.
shytu	shytu.aSep08		26558	402		133	kinase d1 (shytu) mRNA.
shyvar	shyvar.aSep08		1247	855	1		

shyvar	shyvar.bSep08		388	293	2	25	putative protein (shyvar) alternative variant bSep08, mRNA.
shyvo	shyvo.aSep08		39619	443	1	61	putative protein (shyvo) alternative variant aSep08, mRNA.
shyvo	shyvo.bSep08		39671	295		32	putative protein (shyvo) alternative variant bSep08, mRNA.
shywer	shywer.bSep08		3575	746		60	putative protein (6.0 kD) (shywer) alternative variant bSep08, mRNA.
shywey	shywey.aSep08		2822	893		50	putative protein (6.1 kD) (shywey) mRNA.
Si	Si.aSep08	497756	9428	709	2	235	sucrase-isomaltase (alpha-glucosidase) (Si) alternative variant aSep08, mRNA.
Si	Si.bSep08	497756	11572	883	1	143	sucrase-isomaltase (alpha-glucosidase) (Si) alternative variant bSep08, mRNA.
Siae	Siae.bSep08	363045	33452	1505	9	274	sialic acid acetyltransferase precursor (30.2 kD) (Siae) alternative variant bSep08, mRNA.
Siae	Siae.cSep08	363045	11115	407	3	107	sialic acid acetyltransferase (Siae) alternative variant cSep08, mRNA.
Siahbp1	Siahbp1.aSep08	84401	10938	1889	7	563	siah binding protein 1; FBP interacting repressor; pyrimidine tract binding splicing factor; Ro ribonucleoprotein-binding protein 1 (Siahbp1) alternative variant aSep08, complete mRNA.
Siahbp1	Siahbp1.bSep08	84401	9934	1415	3	319	siah binding protein 1; FBP interacting repressor; pyrimidine tract binding splicing factor; Ro ribonucleoprotein-binding protein 1 (34.4 kD) (Siahbp1) alternative variant bSep08, mRNA.
Siahbp1	Siahbp1.cSep08	84401	10179	1416	3	302	siah binding protein 1; FBP interacting repressor; pyrimidine tract binding splicing factor; Ro ribonucleoprotein-binding protein 1 (Siahbp1) alternative variant cSep08, mRNA.
Siahbp1	Siahbp1.dSep08	84401	9603	848	2	271	siah binding protein 1; FBP interacting repressor; pyrimidine tract binding splicing factor; Ro ribonucleoprotein-binding protein 1 (Siahbp1) alternative variant dSep08, mRNA.
Siahbp1	Siahbp1.eSep08	84401	9256	683	2	227	siah binding protein 1; FBP interacting repressor; pyrimidine tract binding splicing factor; Ro ribonucleoprotein-binding protein 1 (Siahbp1) alternative variant eSep08, mRNA.
Sidt1	Sidt1.aSep08	288109	71000	1249		296	SID1 transmembrane family, member 1 (Sidt1) mRNA.
Sidt2	Sidt2.bSep08	315617	6709	1941	14	270	SID1 transmembrane family member 2 (Sidt2) alternative variant bSep08, mRNA.
Sidt2	Sidt2.cSep08	315617	6096	748	10	249	SID1 transmembrane family member 2 (Sidt2) alternative variant cSep08, mRNA.
Sidt2	Sidt2.dSep08	315617	2564	811	4	229	SID1 transmembrane family member 2 (Sidt2) alternative variant dSep08, mRNA.
Sidt2	Sidt2.eSep08	315617	4551	649	9	216	SID1 transmembrane family member 2 (Sidt2) alternative variant eSep08, mRNA.
Sidt2	Sidt2.gSep08	315617	2559	433	6	143	SID1 transmembrane family member 2 (Sidt2) alternative variant gSep08, mRNA.

Sidt2	Sidt2.hSep08	315617	3705	1693	5	124	SID1 transmembrane family member 2 (14.5 kD) (Sidt2) alternative variant hSep08, mRNA.
Sigirr	Sigirr.bSep08	309106	1457	873	5	263	single immunoglobulin and toll-interleukin 1 receptor (TIR) domain (Sigirr) alternative variant bSep08, mRNA.
Sigirr	Sigirr.cSep08	309106	460	386	2	114	single immunoglobulin and toll-interleukin 1 receptor (TIR) domain (Sigirr) alternative variant cSep08, mRNA.
Siglec1	Siglec1.bSep08	311426	1277	637	1	35	sialic acid binding Ig-like lectin 1, sialoadhesin (Siglec1) alternative variant bSep08, mRNA.
Siglecg	Siglecg.aSep08	292844	7036	2053	4	444	sialic acid binding Ig-like lectin G (Siglecg) alternative variant aSep08, mRNA.
Siglech	Siglech.aSep08	361584	9165	1733		220	sialic acid binding Ig-like lectin H (Siglech) mRNA.
Sike	Sike.bSep08	362007	1974	871	2	123	suppressor of IKK epsilon (Sike) alternative variant bSep08, mRNA.
Sike	Sike.cSep08	362007	1845	357	3	113	suppressor of IKK epsilon (Sike) alternative variant cSep08, mRNA.
Sil1	Sil1.bSep08	291673	3222	709	1	118	endoplasmic reticulum chaperone SIL1 homolog (S. cerevisiae) (13.5 kD) (Sil1) alternative variant bSep08, mRNA.
Silv	Silv.bSep08	362818	3116	878	6	254	silver homolog (mouse) (Silv) alternative variant bSep08, mRNA.
Silv	Silv.cSep08	362818	6428	698	6	226	silver homolog (mouse) (Silv) alternative variant cSep08, mRNA.
Sim1	Sim1.bSep08	309888	5264	1697	3	115	single-minded homolog 1 (Drosophila) (13.3 kD) (Sim1) alternative variant bSep08, mRNA.
Sim2	Sim2.bSep08	304071	11593	784	5	133	single-minded homolog 2 (Drosophila) (Sim2) alternative variant bSep08, mRNA.
Sin3a	Sin3a.bSep08	363067	19742	2426	7	471	transcriptional regulator, SIN3A (yeast) (Sin3a) alternative variant bSep08, mRNA.
Sip1	Sip1.aSep08	84404	13762	1777	5	464	survival of motor neuron protein interacting protein 1 (Sip1) alternative variant aSep08, mRNA.
Sip1	Sip1.cSep08	84404	11665	918	9	197	survival of motor neuron protein interacting protein 1 (Sip1) alternative variant cSep08, mRNA.
Sip1	Sip1.dSep08	84404	6205	825	5	127	survival of motor neuron protein interacting protein 1 (14.3 kD) (Sip1) alternative variant dSep08, mRNA.
Sip1	Sip1.fSep08	84404	2539	725	2	63	survival of motor neuron protein interacting protein 1 (7.4 kD) (Sip1) alternative variant fSep08, mRNA.
Sip1	Sip1.hSep08	84404	949	672	2	46	survival of motor neuron protein interacting protein 1 (5.3 kD) (Sip1) alternative variant hSep08, mRNA.
Sipa1	Sipa1.bSep08	361710	1640	1309	5	198	signal-induced proliferation-associated gene 1 (Sipa1) alternative variant bSep08, mRNA.
Sipa1	Sipa1.cSep08	361710	998	829	3	116	signal-induced proliferation-associated gene 1 (Sipa1) alternative variant cSep08, mRNA.
Sipa1	Sipa1.dSep08	361710	1367	1108	4	106	signal-induced proliferation-associated gene 1 (Sipa1) alternative variant dSep08, mRNA.
Sipa111	Sipa111.bSep08	246212	56902	2500	13	770	signal-induced proliferation-associated 1 like CRA a (Sipa111) alternative variant bSep08, mRNA.

Sipa111	Sipa111.cSep08	246212	19821	1175	7	315	signal-induced proliferation-associated 1 like CRA a (Sipa111) alternative variant cSep08, mRNA.
Sipa111	Sipa111.dSep08	246212	29071	663	5	190	signal-induced proliferation-associated 1 like CRA b (Sipa111) alternative variant dSep08, mRNA.
Sipa111	Sipa111.eSep08	246212	167577	591	5	163	putative protein (Sipa111) alternative variant eSep08, mRNA.
Sipa111	Sipa111.fSep08	246212	6716	463	2	92	signal-induced proliferation-associated 1 like CRA a (Sipa111) alternative variant fSep08, mRNA.
Sipa112	Sipa112.bSep08	361442	72605	1783	9	545	signal-induced proliferation-associated 1 like 2 (Sipa112) alternative variant bSep08, mRNA.
Sipa112	Sipa112.cSep08	361442	25934	1518	8	321	signal-induced proliferation-associated 1 like 2 (Sipa112) alternative variant cSep08, mRNA.
Sirpa	Sirpa.bSep08	25528	37082	2515	8	513	signal-regulatory protein alpha (56.1 kD) (Sirpa) alternative variant bSep08, mRNA.
Sirpa	Sirpa.cSep08	25528	16865	394	2	91	signal-regulatory protein alpha (Sirpa) alternative variant cSep08, mRNA.
Sirt1	Sirt1.aSep08	309757	8188	2876		427	sirtuin 1 ((silent mating type information regulation 2, homolog) 1 ( <i>S. cerevisiae</i> ) (Sirt1) mRNA.
Sirt2	Sirt2.aSep08	361532	18707	2533	11	351	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (39.4 kD) (Sirt2) alternative variant aSep08, mRNA.
Sirt2	Sirt2.cSep08	361532	20561	968	9	322	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant cSep08, mRNA.
Sirt2	Sirt2.dSep08	361532	20207	826	9	274	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant dSep08, mRNA.
Sirt2	Sirt2.eSep08	361532	19903	1447	7	238	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant eSep08, mRNA.
Sirt2	Sirt2.fSep08	361532	11171	688	7	229	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant fSep08, mRNA.
Sirt2	Sirt2.gSep08	361532	16954	682	5	132	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant gSep08, mRNA.
Sirt2	Sirt2.hSep08	361532	12404	766	7	130	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant hSep08, mRNA.
Sirt2	Sirt2.iSep08	361532	19193	724	8	124	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (Sirt2) alternative variant iSep08, mRNA.
Sirt2	Sirt2.jSep08	361532	12204	657	6	52	sirtuin (silent mating type information regulation 2 homolog) 2 ( <i>S. cerevisiae</i> ) (5.9 kD) (Sirt2) alternative variant jSep08, mRNA.
Sirt3	Sirt3.bSep08	293615	22418	1254	6	276	sirtuin 3 (silent mating type information regulation 2, homolog) 3 ( <i>S. cerevisiae</i> ) (Sirt3) alternative variant bSep08, mRNA.
Sirt3	Sirt3.cSep08	293615	20984	753	6	250	sirtuin 3 (silent mating type information regulation 2, homolog) 3 ( <i>S. cerevisiae</i> ) (Sirt3) alternative variant cSep08, mRNA.
Sirt3	Sirt3.dSep08	293615	17224	781	5	156	sirtuin 3 (silent mating type information regulation 2, homolog) 3 ( <i>S. cerevisiae</i> ) (Sirt3) alternative variant dSep08, mRNA.



Sirt3	Sirt3.fSep08	293615	6774	404	3	63	sirtuin 3 (silent mating type information regulation 2, homolog) 3 ( <i>S. cerevisiae</i> ) (Sirt3) alternative variant fSep08, mRNA.
Sirt4	Sirt4.bSep08	304539	6762	1436	5	291	sirtuin 4 (silent mating type information regulation 2 homolog) 4 ( <i>S. cerevisiae</i> ) (33.0 kD) (Sirt4) alternative variant bSep08, mRNA.
Sirt5	Sirt5.bSep08	306840	20312	971	6	247	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (Sirt5) alternative variant bSep08, mRNA.
Sirt5	Sirt5.cSep08	306840	19506	827	5	238	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (Sirt5) alternative variant cSep08, mRNA.
Sirt5	Sirt5.dSep08	306840	19821	698	6	232	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (Sirt5) alternative variant dSep08, mRNA.
Sirt5	Sirt5.eSep08	306840	13831	702	5	216	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (23.9 kD) (Sirt5) alternative variant eSep08, mRNA.
Sirt5	Sirt5.fSep08	306840	20189	739	5	161	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (Sirt5) alternative variant fSep08, mRNA.
Sirt5	Sirt5.gSep08	306840	5211	310	2	87	sirtuin 5 (silent mating type information regulation 2 homolog) 5 ( <i>S. cerevisiae</i> ) (Sirt5) alternative variant gSep08, mRNA.
Sirt6	Sirt6.bSep08	299638	3648	588	6	139	sirtuin 6 (silent mating type information regulation 2, homolog) 6 ( <i>S. cerevisiae</i> ) (Sirt6) alternative variant bSep08, mRNA.
Sirt6	Sirt6.cSep08	299638	1174	356	4	118	sirtuin 6 (silent mating type information regulation 2, homolog) 6 ( <i>S. cerevisiae</i> ) (Sirt6) alternative variant cSep08, mRNA.
Sirt6	Sirt6.dSep08	299638	852	711	2	55	sirtuin 6 (silent mating type information regulation 2, homolog) 6 ( <i>S. cerevisiae</i> ) (Sirt6) alternative variant dSep08, mRNA.
Sirt7	Sirt7.bSep08	303745	6118	1228	9	180	sirtuin 7 CRA c (Sirt7) alternative variant bSep08, mRNA.
Sirt7	Sirt7.cSep08	303745	5106	1309	8	174	sirtuin 7 CRA c (19.0 kD) (Sirt7) alternative variant cSep08, mRNA.
Sirt7	Sirt7.dSep08	303745	2404	1125	3	110	sirtuin 7 (Sirt7) alternative variant dSep08, mRNA.
Sirt7	Sirt7.eSep08	303745	551	371	2	97	sirtuin 7 (Sirt7) alternative variant eSep08, mRNA.
SIS.0	SIS.0.aSep08		9610	4522	5	193	transaminase 1 (SIS.0) alternative variant aSep08, mRNA.
Siva1	Siva1.aSep08	362791	4379	717	2	177	SIVA1, apoptosis-inducing factor (18.9 kD) (Siva1) alternative variant aSep08, mRNA.
Six1	Six1.aSep08	114634	3212	1163	2	224	sine oculis-related homeobox 1 homolog ( <i>Drosophila</i> ) (Six1) alternative variant aSep08, mRNA.
Six6os1	Six6os1.aSep08	500673	8730	375		44	six6 opposite strand transcript 1 (5.3 kD) (Six6os1) mRNA.
skabor	skabor.aSep08		1183	298		93	putative protein (skabor) mRNA.
skachy	skachy.aSep08		1038	365		98	CRA a like (skachy) mRNA.

skadoy	skadoy.bSep08		16819	698	5	83	putative protein (skadoy) alternative variant bSep08, mRNA.
skafee	skafee.aSep08		2508	499		49	putative protein (skafee) mRNA.
skafly	skafly.aSep08		2489	590		78	putative mitochondrial protein (8.8 kD) (skafly) mRNA.
skafly	skafly.aSep08		2004	427		89	putative protein (skafly) mRNA.
skaja	skaja.aSep08		1425	384		65	putative protein (skaja) mRNA.
skajey	skajey.aSep08		17901	1730	6	177	CRA b (skajey) alternative variant aSep08, mRNA.
skalo	skalo.aSep08		6450	400		133	acetyl-Coenzyme a carboxylase (skalo) mRNA.
skamee	skamee.aSep08		3366	897		63	putative protein (skamee) mRNA.
Skap1	Skap1.bSep08	286975	27041	198	1	65	src family associated phosphoprotein 1 (Skap1) alternative variant bSep08, mRNA.
Skap2	Skap2.bSep08	155183	48434	819	5	118	src family associated phosphoprotein 2 (Skap2) alternative variant bSep08, mRNA.
Skap2	Skap2.cSep08	155183	40579	645	5	98	src family associated phosphoprotein 2 (11.0 kD) (Skap2) alternative variant cSep08, mRNA.
Skap2	Skap2.eSep08	155183	923	468	2	40	src family associated phosphoprotein 2 (4.5 kD) (Skap2) alternative variant eSep08, mRNA.
skapey	skapey.aSep08		2806	1132		86	putative protein (9.5 kD) (skapey) mRNA.
skapor	skapor.aSep08		8119	747		29	putative protein (skapor) mRNA.
skarbor	skarbor.aSep08		25165	1202		384	putative protein, with 3 coiled coil domains, of bilateral origin (skarbor) mRNA.
skarchy	skarchy.aSep08		5248	469		69	small adipocyte factor 1 like (skarchy) mRNA.
skardoy	skardoy.bSep08		2368	741		54	putative protein (6.3 kD) (skardoy) alternative variant bSep08, mRNA.
skarfee	skarfee.aSep08		1905	943		115	putative protein (12.6 kD) (skarfee) mRNA.
skarflu	skarflu.aSep08		8694	1103		139	envelope protein (skarflu) mRNA.
skarfly	skarfly.aSep08		32897	483		39	putative protein (4.5 kD) (skarfly) mRNA.
skarja	skarja.aSep08		53744	379		64	putative protein (6.8 kD) (skarja) mRNA.
skarjey	skarjey.aSep08		44116	489		64	CRA a like (skarjey) mRNA.
skarlo	skarlo.aSep08		1240	449		47	putative protein (5.5 kD) (skarlo) mRNA.
skarmee	skarmee.aSep08		1640	687		186	putative protein of mammalian origin (skarmee) alternative variant aSep08, mRNA.
skaroy	skaroy.aSep08		17196	697		33	putative protein (skaroy) mRNA.
skarpey	skarpey.aSep08		2287	183		61	repeat-containing protein (skarpey) mRNA.
skarpor	skarpor.aSep08		4177	403		34	putative protein (skarpor) mRNA.
skarroy	skarroy.aSep08		3786	438	1	42	putative protein (skarroy) alternative variant aSep08, mRNA.
skarroy	skarroy.bSep08		24189	381	2	47	putative protein (skarroy) alternative variant bSep08, mRNA.
skarshaw	skarshaw.aSep08		3381	623		74	putative mitochondrial protein (7.8 kD) (skarshaw) mRNA.
skarshee	skarshee.aSep08		4422	605		107	putative protein of mammalian origin (skarshee) mRNA.
skartu	skartu.aSep08		1437	570		80	gene model 1568 like (skartu) mRNA.
skarvo	skarvo.aSep08		4291	692		10	putative protein (1.1 kD) (skarvo) mRNA.

skarwer	skarwer.aSep08		832	710		40	putative protein (4.5 kD) (skarwer) mRNA.
skashaw	skashaw.aSep08		7258	1516	3	123	putative protein of metazoan origin (skashaw) alternative variant aSep08, mRNA.
skashee	skashee.aSep08		9763	255		20	putative protein (skashee) mRNA.
skatu	skatu.aSep08		4049	481		41	putative protein (4.6 kD) (skatu) mRNA.
skavo	skavo.aSep08		11393	371		86	putative protein (skavo) mRNA.
skawbor	skawbor.aSep08		15971	545		128	putative protein of metazoan origin (skawbor) mRNA.
skawchy	skawchy.aSep08		3141	326		58	putative protein (skawchy) mRNA.
skawdoy	skawdoy.aSep08		131119	377		39	putative protein (4.3 kD) (skawdoy) mRNA.
skawer	skawer.aSep08		4274	374		20	putative protein (2.3 kD) (skawer) mRNA.
skawfee	skawfee.bSep08		7807	784	5	53	putative protein (5.7 kD) (skawfee) alternative variant bSep08, mRNA.
skawflu	skawflu.bSep08		18049	375	3	37	putative protein (4.1 kD) (skawflu) alternative variant bSep08, mRNA.
skawfly	skawfly.aSep08		6678	330		33	putative protein (skawfly) mRNA.
skawja	skawja.aSep08		10417	1572	2	182	slain 1 (skawja) alternative variant aSep08, mRNA.
skawja	skawja.bSep08		31695	537	1	52	slain motif family member 1 (skawja) alternative variant bSep08, mRNA.
skawjey	skawjey.aSep08		3307	357		65	polyprotein (skawjey) mRNA.
skawlo	skawlo.aSep08		6683	252		83	phosphatidylinositol transfer protein beta (skawlo) mRNA.
skawmee	skawmee.aSep08		25771	508		88	neurofibromatosis 1 (skawmee) mRNA.
skawpey	skawpey.aSep08		10619	790	4	106	repeat-containing protein (skawpey) alternative variant aSep08, mRNA.
skawpey	skawpey.bSep08		5234	855	2	56	repeat-containing protein like (6.4 kD) (skawpey) alternative variant bSep08, mRNA.
skawpor	skawpor.aSep08		3584	457	2	73	putative protein (skawpor) alternative variant aSep08, mRNA.
skawroy	skawroy.aSep08		1717	398		132	utp20 processome component homolog CRA a (skawroy) mRNA.
skawshaw	skawshaw.aSep08		3781	237		76	putative protein (skawshaw) mRNA.
skawshee	skawshee.aSep08		6896	485		22	putative protein (2.4 kD) (skawshee) mRNA.
skawtu	skawtu.aSep08		38759	643	1	62	CRA b like (6.7 kD) (skawtu) alternative variant aSep08, mRNA.
skawtu	skawtu.bSep08		38229	528	2	108	putative protein (skawtu) alternative variant bSep08, mRNA.
skawtu	skawtu.cSep08		38181	356	1	87	putative protein (skawtu) alternative variant cSep08, mRNA.
skawvo	skawvo.aSep08		50878	575		191	ionotropic glutamate receptor (skawvo) mRNA.
skawwer	skawwer.aSep08		12929	800		145	CAMP protein like (skawwer) mRNA.
skeebor	skeebor.aSep08		3226	199		63	gag protein like (skeebor) mRNA.
skeechy	skeechy.aSep08		1231	357		72	putative protein (skeechy) mRNA.
skeedoy	skeedoy.aSep08		2832	400		71	putative mitochondrial protein (7.8 kD) (skeedoy) mRNA.
skeefee	skeefee.aSep08		733	498		27	putative protein (3.1 kD) (skeefee) mRNA.

skeeflu	skeeflu.bSep08		3970	426	2	59	putative protein (6.7 kD) (skeeflu) alternative variant bSep08, mRNA.
skeefly	skeefly.bSep08		1425	626	2	44	putative protein (skeefly) alternative variant bSep08, mRNA.
skeeja	skeeja.aSep08		7067	1573		53	putative protein (5.8 kD) (skeeja) mRNA.
skeejey	skeejey.aSep08		104118	397		34	putative protein (4.0 kD) (skeejey) mRNA.
skeelo	skeelo.aSep08		8009	392		130	tetratricopeptide repeat 28 (skeelo) mRNA.
skeemee	skeemee.aSep08		3590	347		115	neurofibromatosis 1 (skeemee) mRNA.
skeepey	skeepey.aSep08		17877	570	4	56	putative protein (6.5 kD) (skeepey) alternative variant aSep08, mRNA.
skeepey	skeepey.bSep08		9900	1294	3	40	putative protein (4.6 kD) (skeepey) alternative variant bSep08, mRNA.
skeepey	skeepey.cSep08		7453	1179	2	40	putative protein (4.6 kD) (skeepey) alternative variant cSep08, mRNA.
skeepey	skeepey.eSep08		19875	732	5	56	putative protein (6.5 kD) (skeepey) alternative variant eSep08, mRNA.
skeepor	skeepor.aSep08		2313	573		60	putative protein (skeepor) mRNA.
skeeroy	skeeroy.aSep08		7803	1055		63	putative protein (6.9 kD) (skeeroy) mRNA.
skeeshaw	skeeshaw.aSep08		1860	536		33	putative protein (3.3 kD) (skeeshaw) mRNA.
skeeshee	skeeshee.aSep08		2820	726	2	57	putative protein (skeeshee) alternative variant aSep08, mRNA.
skeeshee	skeeshee.bSep08		5657	386	2	50	putative protein (skeeshee) alternative variant bSep08, mRNA.
skeetu	skeetu.aSep08		5732	487		64	putative protein (7.3 kD) (skeetu) mRNA.
skeevo	skeevo.aSep08		5544	275		91	putative protein of mammalian origin (skeevo) mRNA.
skeewer	skeewer.bSep08		18465	379	3	40	putative protein (skeewer) alternative variant bSep08, mRNA.
skerbor	skerbor.aSep08		4155	686		85	putative protein of vertebrate origin (skerbor) mRNA.
skerchy	skerchy.aSep08		2479	536		65	putative protein (skerchy) mRNA.
skerdoy	skerdoy.aSep08		45513	712		137	putative mitochondrial protein (15.4 kD) (skerdoy) mRNA.
skerfee	skerfee.aSep08		3042	440		38	putative protein (skerfee) mRNA.
skerflu	skerflu.aSep08		804	538		97	putative protein (skerflu) mRNA.
skerfly	skerfly.aSep08		38218	921		47	putative protein (skerfly) mRNA.
skerja	skerja.aSep08		278648	737		100	putative cytoplasmic protein (11.6 kD) (skerja) mRNA.
skerjey	skerjey.aSep08		29978	594	4	43	putative protein (4.9 kD) (skerjey) alternative variant aSep08, mRNA.
skerjey	skerjey.cSep08		6910	490	4	40	putative protein (4.6 kD) (skerjey) alternative variant cSep08, mRNA.
skerlo	skerlo.aSep08		8079	280		75	putative protein (skerlo) mRNA.
skermee	skermee.aSep08		5878	481		52	neurofibromatosis 1 (skermee) mRNA.
skerpey	skerpey.aSep08		4285	348		116	putative protein (skerpey) mRNA.
skerpor	skerpor.bSep08		1731	588	2	30	putative protein (3.3 kD) (skerpor) alternative variant bSep08, mRNA.

skerroy	skerroy.cSep08		7410	381	3	49	putative protein (5.6 kD) (skerroy) alternative variant cSep08, mRNA.
skershaw	skershaw.aSep08		1307	341		34	putative protein (3.9 kD) (skershaw) mRNA.
skershee	skershee.aSep08		1270	307		43	putative protein (skershee) mRNA.
skertu	skertu.aSep08		113791	580	4	127	COX16 cytochrome c oxidase assembly homolog (14.6 kD) (skertu) alternative variant aSep08, mRNA.
skertu	skertu.bSep08		14716	2471	4	106	COX16 cytochrome c oxidase assembly homolog (12.3 kD) (skertu) alternative variant bSep08, mRNA.
skervo	skervo.aSep08		382	295		47	nipsnap homolog 3A (skervo) mRNA.
skerwer	skerwer.aSep08		60490	497		42	putative protein (4.7 kD) (skerwer) mRNA.
skeybor	skeybor.aSep08		11415	517		75	putative protein of vertebrate origin (skeybor) mRNA.
skeychy	skeychy.aSep08		3146	593		58	putative protein (6.9 kD) (skeychy) mRNA.
skeydoy	skeydoy.aSep08		2466	242		60	putative protein (skeydoy) mRNA.
skeyfee	skeyfee.aSep08		16546	599		48	kelch-like 3 (skeyfee) mRNA.
skeyflu	skeyflu.aSep08		2738	416		138	dedicator of cytokinesis 1 (skeyflu) mRNA.
skeyfly	skeyfly.aSep08		36240	799		49	putative protein (skeyfly) mRNA.
skeyja	skeyja.aSep08		71290	434	1	112	putative protein (12.5 kD) (skeyja) alternative variant aSep08, mRNA.
skeyja	skeyja.bSep08		72542	1031	2	83	putative protein (9.3 kD) (skeyja) alternative variant bSep08, mRNA.
skeyjey	skeyjey.aSep08		18944	1388		462	girdin (skeyjey) mRNA.
skeylo	skeylo.aSep08		2326	775		247	nucleolar complex associated 4 homolog (skeylo) mRNA.
skeymee	skeymee.aSep08		669	472		42	putative protein (4.3 kD) (skeymee) mRNA.
skeypey	skeypey.aSep08		7047	752		59	putative protein (6.5 kD) (skeypey) mRNA.
skeypor	skeypor.aSep08		634	505		63	putative protein (skeypor) mRNA.
skeyroy	skeyroy.aSep08		11435	430		71	putative protein (skeyroy) mRNA.
skeyshaw	skeyshaw.aSep08		16991	400		75	putative protein (skeyshaw) mRNA.
skeyshee	skeyshee.aSep08		3109	760		55	putative protein (skeyshee) mRNA.
skeytu	skeytu.aSep08		27172	767		43	putative protein (5.1 kD) (skeytu) mRNA.
skeyvo	skeyvo.aSep08		17722	785		57	putative protein (6.4 kD) (skeyvo) mRNA.
skeywer	skeywer.aSep08		23249	843		35	putative protein (4.0 kD) (skeywer) mRNA.
Skil	Skil.aSep08	114208	19445	768	4	256	SKI-like (Skil) alternative variant aSep08, mRNA.
Skil	Skil.bSep08	114208	22310	518	5	115	SKI-like (Skil) alternative variant bSep08, mRNA.
Skil	Skil.cSep08	114208	1936	383	2	33	SKI-like (Skil) alternative variant cSep08, mRNA.
Skip	Skip.bSep08	287533	5613	749	5	184	skeletal muscle and kidney enriched inositol phosphatase (Skip) alternative variant bSep08, mRNA.
Skip	Skip.cSep08	287533	6418	542	5	149	skeletal muscle and kidney enriched inositol phosphatase (16.7 kD) (Skip) alternative variant cSep08, mRNA.
Skip	Skip.dSep08	287533	2539	630	2	71	skeletal muscle and kidney enriched inositol phosphatase (Skip) alternative variant dSep08, mRNA.
Skip	Skip.eSep08	287533	2470	1929	4	97	skeletal muscle and kidney enriched inositol phosphatase (10.4 kD) (Skip) alternative variant eSep08, mRNA.

Skiv2l	Skiv2l.bSep08	294260	769	418	3	139	superkiller viralicidic activity 2-like (Skiv2l) alternative variant bSep08, mRNA.
Skiv2l2	Skiv2l2.aSep08	365668	23285	1423	13	473	superkiller viralicidic activity 2-like 2 (S. cerevisiae) (Skiv2l2) alternative variant aSep08, mRNA.
Skiv2l2	Skiv2l2.bSep08	365668	7380	856	4	285	superkiller viralicidic activity 2-like 2 (S. cerevisiae) (Skiv2l2) alternative variant bSep08, mRNA.
Skiv2l2	Skiv2l2.cSep08	365668	16917	1023	5	257	superkiller viralicidic activity 2-like 2 (S. cerevisiae) (Skiv2l2) alternative variant cSep08, mRNA.
skobor	skobor.aSep08		103686	2737	13	703	CRA a (80.9 kD) (skobor) alternative variant aSep08, complete mRNA.
skobor	skobor.bSep08		58255	697	5	231	putative protein, with a coiled coil domain, of vertebrate origin (skobor) alternative variant bSep08, mRNA.
skochoy	skochoy.aSep08		3704	354		117	protein CRA b (skochoy) mRNA.
skodoy	skodoy.aSep08		5906	1057		254	CRA a (skodoy) mRNA.
skofee	skofee.aSep08		8245	445		34	putative protein (3.9 kD) (skofee) mRNA.
skoflu	skoflu.aSep08		3360	528	2	43	putative protein (4.9 kD) (skoflu) alternative variant aSep08, mRNA.
skofly	skofly.aSep08		2078	398		48	putative protein (skofly) mRNA.
skoja	skoja.aSep08		937	324		55	CRA b like (skoja) mRNA.
skojey	skojey.aSep08		35340	323		57	putative protein (skojey) mRNA.
skolo	skolo.aSep08		1279	723		33	CRA a like (3.6 kD) (skolo) mRNA.
skomee	skomee.aSep08		10733	414		137	neurofibromin (skomee) mRNA.
skokey	skokey.aSep08		1446	708		58	putative protein (6.6 kD) (skokey) mRNA.
skopor	skopor.aSep08		75738	397		97	kinase 4 (skopor) mRNA.
skorbor	skorbor.aSep08		1344	710		39	putative protein (skorbor) mRNA.
skorchy	skorchy.aSep08		69154	702	2	53	putative protein (skorchy) alternative variant aSep08, mRNA.
skorchy	skorchy.bSep08		33526	616	1	55	putative protein (skorchy) alternative variant bSep08, mRNA.
skorchy	skorchy.cSep08		68976	468	1	61	putative protein (skorchy) alternative variant cSep08, mRNA.
skordoy	skordoy.aSep08		2966	231		22	putative protein (2.5 kD) (skordoy) mRNA.
skorfee	skorfee.aSep08		904	278		92	putative protein (skorfee) mRNA.
skorflu	skorflu.aSep08		22999	910		303	dedicator of cytokinesis 1 (skorflu) mRNA.
skorfly	skorfly.aSep08		4320	462	1	70	CRA b like (skorfly) alternative variant aSep08, mRNA.
skorfly	skorfly.bSep08		12255	381	2	71	putative protein (skorfly) alternative variant bSep08, mRNA.
skorja	skorja.aSep08		1121	439		80	transcription factor 1 (skorja) mRNA.
skorjey	skorjey.aSep08		113197	510		87	putative protein (skorjey) mRNA.
skorlo	skorlo.aSep08		2471	468		61	putative protein (7.2 kD) (skorlo) mRNA.
skormee	skormee.aSep08		17929	835		54	hepatocellular carcinoma-associated antigen 66 like (skormee) mRNA.
skoroy	skoroy.aSep08		36066	644	4	185	myosin binding protein C slow type like (skoroy) alternative variant aSep08, mRNA.

skoroy	skoroy.bSep08		36069	569	2	160	myosin binding protein C slow type like (skoroy) alternative variant bSep08, mRNA.
skoroy	skoroy.cSep08		35143	376	1	102	myosin binding protein C slow type like (skoroy) alternative variant cSep08, mRNA.
skorpey	skorpey.aSep08		2654	371		66	putative protein (skorpey) mRNA.
skorpor	skorpor.aSep08		103575	672		27	putative protein (3.1 kD) (skorpor) mRNA.
skorroy	skorroy.aSep08		1403	275		85	putative protein (skorroy) mRNA.
skorshaw	skorshaw.aSep08		12390	404		32	putative protein (skorshaw) mRNA.
skorshee	skorshee.aSep08		16317	698		82	putative nuclear protein (9.8 kD) (skorshee) mRNA.
skortu	skortu.aSep08		6808	581		118	putative protein (skortu) mRNA.
skorvo	skorvo.aSep08		59789	673		38	putative protein (4.5 kD) (skorvo) mRNA.
skorwer	skorwer.aSep08		1992	749	1	39	putative protein (4.7 kD) (skorwer) alternative variant aSep08, mRNA.
skorwer	skorwer.bSep08		3412	714	3	58	putative protein (6.6 kD) (skorwer) alternative variant bSep08, mRNA.
skoshaw	skoshaw.aSep08		2739	557		57	metaxin 3 (skoshaw) mRNA.
skoshee	skoshee.aSep08		6779	583		82	putative protein of vertebrate origin (skoshee) mRNA.
skotu	skotu.bSep08		2004	692	3	153	putative protein (skotu) alternative variant bSep08, mRNA.
skovo	skovo.aSep08		7900	1001		186	glutamate receptor ionotropic NMDA3A (skovo) mRNA.
skower	skower.aSep08		133690	577	5	84	CRA a like (9.7 kD) (skower) alternative variant aSep08, mRNA.
skower	skower.bSep08		1457	230	1	43	CRA d like (skower) alternative variant bSep08, mRNA.
skoybor	skoybor.aSep08		9733	850		283	RAN binding protein 2 like (skoybor) mRNA.
skoychy	skoychy.aSep08		2938	788		33	putative protein (3.8 kD) (skoychy) mRNA.
skoydoy	skoydoy.aSep08		2756	318		66	putative protein (7.3 kD) (skoydoy) mRNA.
skoyfee	skoyfee.aSep08		16168	557		158	transforming growth factor beta-induced 68kDa like (skoyfee) mRNA.
skoyflu	skoyflu.aSep08		37030	667		222	CRA f (skoyflu) mRNA.
skoyfly	skoyfly.aSep08		2581	427		22	putative protein (2.5 kD) (skoyfly) mRNA.
skoyja	skoyja.aSep08		3281	739		97	putative protein (skoyja) mRNA.
skoyjey	skoyjey.aSep08		3936	352		84	putative protein (9.5 kD) (skoyjey) mRNA.
skoylo	skoylo.aSep08		15549	461	3	153	CRA a like (skoylo) alternative variant aSep08, mRNA.
skoymee	skoymee.aSep08		2888	751		227	putative protein of eukaryotic origin (skoymee) mRNA.
skoypey	skoypey.aSep08		5913	288		22	putative protein (2.3 kD) (skoypey) mRNA.
skoypor	skoypor.aSep08		7086	649		63	putative protein (7.3 kD) (skoypor) mRNA.
skoyroy	skoyroy.aSep08		17098	432		73	putative protein (skoyroy) mRNA.
skoyshaw	skoyshaw.aSep08		24984	407		135	IQ motif containing GTPase activating protein 2 (skoyshaw) mRNA.
skoyshee	skoyshee.aSep08		879	755		101	putative protein (11.3 kD) (skoyshee) mRNA.
skoytu	skoytu.aSep08		28757	400		132	pecanex homolog CRA d (skoytu) mRNA.
skoyvo	skoyvo.aSep08		4206	505		38	putative protein (skoyvo) mRNA.
skoywer	skoywer.aSep08		12396	746	4	144	carboxypeptidase 3 cytosolic (16.6 kD) (skoywer) alternative variant aSep08, mRNA.

skoywer	skoywer.bSep08		2085	1277	2	88	carboxypeptidase 3 cytosolic (10.3 kD) (skoywer) alternative variant bSep08, mRNA.
Skp1a	Skp1a.aSep08	287280	14400	775	7	163	S-phase kinase-associated protein 1A (18.7 kD) (Skp1a) alternative variant aSep08, mRNA.
Skp1a	Skp1a.cSep08	287280	13189	828	5	149	S-phase kinase-associated protein 1A (Skp1a) alternative variant cSep08, mRNA.
Skp1a	Skp1a.dSep08	287280	11049	262	3	65	S-phase kinase-associated protein 1A (Skp1a) alternative variant dSep08, mRNA.
Skp1a	Skp1a.eSep08	287280	13126	780	5	63	S-phase kinase-associated protein 1A (6.9 kD) (Skp1a) alternative variant eSep08, mRNA.
Skp1a	Skp1a.fSep08	287280	2686	736	3	37	S-phase kinase-associated protein 1A (Skp1a) alternative variant fSep08, mRNA.
skubor	skubor.aSep08		35799	316		103	CUB (skubor) mRNA.
skuchy	skuchy.aSep08		12445	704		234	putative protein of eukaryotic origin (skuchy) mRNA.
skudoy	skudoy.aSep08		5950	733		244	CRA a (skudoy) mRNA.
skufee	skufee.aSep08		651	488		46	putative protein (4.8 kD) (skufee) mRNA.
skuflu	skuflu.aSep08		5269	350		81	putative protein (skuflu) mRNA.
skufly	skufly.aSep08		4010	657		81	putative protein (8.8 kD) (skufly) mRNA.
skuja	skuja.aSep08		17783	252		54	putative protein (skuja) mRNA.
skujey	skujey.aSep08		56215	400		90	putative cytoplasmic protein (10.1 kD) (skujey) mRNA.
skulo	skulo.aSep08		5465	1059		143	uncharacterized protein like (skulo) mRNA.
skumee	skumee.aSep08		5482	596		198	neurofibromin (skumee) mRNA.
skupey	skupey.aSep08		5740	1701		72	putative protein (skupey) alternative variant aSep08, mRNA.
skupor	skupor.aSep08		770	476		37	putative protein (skupor) mRNA.
skuroy	skuroy.bSep08		2104	587	2	37	putative protein (skuroy) alternative variant bSep08, mRNA.
skushaw	skushaw.aSep08		683	348		54	metaxin 1 (skushaw) mRNA.
skushee	skushee.aSep08		6157	720		70	putative protein (8.0 kD) (skushee) mRNA.
skutu	skutu.aSep08		1549	777		170	alpha-actinin like (skutu) alternative variant aSep08, mRNA.
skuvo	skuvo.aSep08		2567	467		134	procollagen type XV CRA a (skuvo) mRNA.
skuwer	skuwer.aSep08		11044	540		125	putative nuclear protein of mammalian origin (13.8 kD) (skuwer) mRNA.
skybor	skybor.bSep08		1665	553	2	65	putative protein (7.7 kD) (skybor) alternative variant bSep08, mRNA.
skychy	skychy.aSep08		11484	355		37	putative protein (skychy) mRNA.
skydoy	skydoy.aSep08		11241	800		266	CRA a (skydoy) mRNA.
skyfee	skyfee.aSep08		967	368		94	putative protein (skyfee) mRNA.
skyflu	skyflu.aSep08		2559	360		29	putative protein (skyflu) mRNA.
skyfly	skyfly.aSep08		2220	402		62	putative protein (skyfly) mRNA.
skyja	skyja.aSep08		3965	395		22	putative protein (2.6 kD) (skyja) mRNA.
skyjey	skyjey.aSep08		6864	356		11	putative protein (1.2 kD) (skyjey) mRNA.
skylo	skylo.aSep08		977	367		70	putative protein (skylo) mRNA.



skymee	skymee.aSep08		1259	379		33	nemo-like kinase (skymee) mRNA.
skypey	skypey.aSep08		4505	412		38	putative protein (skypey) mRNA.
skypor	skypor.aSep08		3399	783		40	putative protein (4.7 kD) (skypor) mRNA.
skyroy	skyroy.aSep08		33623	656		218	transferase (skyroy) mRNA.
skyshaw	skyshaw.aSep08		17625	476		158	thrombospondin 4 (skyshaw) mRNA.
skyshee	skyshee.aSep08		7397	291		24	putative protein (2.7 kD) (skyshee) mRNA.
skytu	skytu.aSep08		2339	726		38	putative protein (skytu) mRNA.
skyvo	skyvo.aSep08		41898	570		63	putative protein (7.4 kD) (skyvo) mRNA.
skywer	skywer.aSep08		3605	878		43	putative protein (4.8 kD) (skywer) mRNA.
Sla	Sla.bSep08	338477	29042	781	7	162	src-like adaptor (18.0 kD) (Sla) alternative variant bSep08, mRNA.
slabor	slabor.aSep08		612	367		121	RAN binding protein 2 like (slabor) mRNA.
slachy	slachy.aSep08		36554	562	1	42	putative protein (slachy) alternative variant aSep08, mRNA.
slachy	slachy.bSep08		46935	328	1	42	putative protein (5.0 kD) (slachy) alternative variant bSep08, mRNA.
sladoy	sladoy.aSep08		1181	364		86	putative protein (sladoy) mRNA.
slafee	slafee.aSep08		5122	523		97	putative protein (slafee) mRNA.
slaflu	slaflu.aSep08		22039	336	4	111	dedicator of cytokinesis 1 (slaflu) alternative variant aSep08, mRNA.
slafly	slafly.aSep08		9504	504		105	putative nuclear protein (12.1 kD) (slafly) mRNA.
Slain2	Slain2.bSep08	305310	77712	2466	10	469	slain motif family member 2 (Slain2) alternative variant bSep08, mRNA.
Slain2	Slain2.cSep08	305310	1267	780	2	185	slain motif family member 2 (Slain2) alternative variant cSep08, mRNA.
Slain2	Slain2.dSep08	305310	14611	707	4	184	slain motif family member 2 (Slain2) alternative variant dSep08, mRNA.
Slain2	Slain2.eSep08	305310	7542	283	3	69	putative protein (Slain2) alternative variant eSep08, mRNA.
slaja	slaja.aSep08		32386	1256	5	278	RNA binding motif protein 26 like (slaja) alternative variant aSep08, mRNA.
slaja	slaja.bSep08		376	254	1	84	RNA binding motif protein 26 like (slaja) alternative variant bSep08, mRNA.
slajey	slajey.aSep08		10353	195		43	putative protein (slajey) mRNA.
slalo	slalo.aSep08		40077	870		290	CRA a (slalo) mRNA.
slamee	slamee.aSep08		1554	882	3	193	putative protein, with a coiled coil domain, of metazoan origin (slamee) alternative variant aSep08, mRNA.
slamee	slamee.bSep08		1303	768	4	183	putative protein of eukaryotic origin (slamee) alternative variant bSep08, mRNA.
slamee	slamee.cSep08		1369	837	4	141	putative protein of eukaryotic origin (slamee) alternative variant cSep08, mRNA.
Slamf6	Slamf6.aSep08	498287	5131	752	6	160	SLAM family member 6 (Slamf6) alternative variant aSep08, mRNA.
Slamf6	Slamf6.bSep08	498287	3537	854	3	39	SLAM family member 6 (4.6 kD) (Slamf6) alternative variant bSep08, mRNA.
Slamf6	Slamf6.cSep08	498287	1556	729	2	36	SLAM family member 6 (4.2 kD) (Slamf6) alternative variant cSep08, mRNA.

Slamf7	Slamf7.aSep08	364049	3724	373		113	SLAM family member 7 (Slamf7) mRNA.
slapey	slapey.aSep08		4213	199		63	putative protein (slapey) mRNA.
slapor	slapor.aSep08		5268	530	3	150	C-type lectin domain family 3 member b (slapor) alternative variant aSep08, mRNA.
slarbor	slarbor.aSep08		7471	209		69	putative protein of metazoan origin (slarbor) mRNA.
slarchy	slarchy.aSep08		592	374		102	putative cytoplasmic protein (11.2 kD) (slarchy) mRNA.
slardoy	slardoy.aSep08		16769	912		66	putative protein (7.6 kD) (slardoy) mRNA.
slarfee	slarfee.aSep08		648	496		44	putative protein (4.9 kD) (slarfee) mRNA.
slarflu	slarflu.aSep08		3493	654		95	putative protein (slarflu) mRNA.
slarfly	slarfly.aSep08		1880	325		37	putative protein (4.2 kD) (slarfly) mRNA.
slarja	slarja.aSep08		58027	441	3	26	putative protein (slarja) alternative variant aSep08, mRNA.
slarja	slarja.bSep08		37223	301	2	23	putative protein (slarja) alternative variant bSep08, mRNA.
slarjey	slarjey.aSep08		2127	1199		55	CRA a like (6.8 kD) (slarjey) mRNA.
slarlo	slarlo.aSep08		752	258		79	putative protein (slarlo) mRNA.
slarmee	slarmee.aSep08		29277	385		61	RNA methyltransferase like 1 (slarmee) mRNA.
slaroy	slaroy.bSep08		4480	575	3	41	putative protein (4.8 kD) (slaroy) alternative variant bSep08, mRNA.
slarpey	slarpey.aSep08		760	227		75	intercellular adhesion molecule 5 Telencephalin (slarpey) mRNA.
slarpor	slarpor.aSep08		13564	769		197	complex D3 (slarpor) mRNA.
slarroy	slarroy.aSep08		1795	306		99	uncharacterized protein (slarroy) mRNA.
slarshaw	slarshaw.aSep08		1561	398		25	putative protein (3.0 kD) (slarshaw) mRNA.
slarshee	slarshee.aSep08		41936	797	4	116	tubulin tyrosine ligase-like family member 7 (slarshee) alternative variant aSep08, mRNA.
slarshee	slarshee.bSep08		41922	997	5	102	tubulin tyrosine ligase-like family member 7 (slarshee) alternative variant bSep08, mRNA.
slarshee	slarshee.cSep08		38767	338	1	34	tubulin tyrosine ligase-like family member 7 (slarshee) alternative variant cSep08, mRNA.
slartu	slartu.aSep08		22280	604		46	putative protein (5.0 kD) (slartu) mRNA.
slarvo	slarvo.aSep08		2331	795		52	putative protein (5.7 kD) (slarvo) mRNA.
slarwer	slarwer.aSep08		13126	396		131	putative protein of vertebrate origin (slarwer) mRNA.
slashaw	slashaw.aSep08		43222	885		294	IQ motif containing GTPase activating protein 2 (slashaw) mRNA.
slashee	slashee.aSep08		34794	367		117	collagen chain like (slashee) mRNA.
slatu	slatu.aSep08		15768	1062	8	353	pecanex homolog (slatu) alternative variant aSep08, mRNA.
slatu	slatu.bSep08		2251	382	1	35	putative protein (4.1 kD) (slatu) alternative variant bSep08, mRNA.
slavo	slavo.aSep08		8212	740		122	putative protein, with a transmembrane domain, of metazoan origin (slavo) mRNA.
slawbor	slawbor.aSep08		7706	334		40	putative protein (slawbor) mRNA.
slawchy	slawchy.aSep08		1695	288		90	putative protein (slawchy) mRNA.
slawdoy	slawdoy.aSep08		23429	345		80	putative protein (slawdoy) mRNA.

slawer	slawer.aSep08		1072	419		139	carboxypeptidase 3 cytosolic (slawer) mRNA.
slawfee	slawfee.aSep08		21593	794	3	50	putative protein (5.6 kD) (slawfee) alternative variant aSep08, mRNA.
slawflu	slawflu.aSep08		44490	709	1	236	putative protein (slawflu) alternative variant aSep08, mRNA.
slawflu	slawflu.bSep08		1779	694		168	putative protein (slawflu) alternative variant bSep08, mRNA.
slawfly	slawfly.aSep08		4716	385		67	putative protein (7.1 kD) (slawfly) mRNA.
slawja	slawja.aSep08		25918	282		93	putative protein (slawja) mRNA.
slawjey	slawjey.aSep08		625	238	2	79	putative protein (slawjey) alternative variant aSep08, mRNA.
slawlo	slawlo.aSep08		4036	746		61	putative protein (6.5 kD) (slawlo) mRNA.
slawmee	slawmee.aSep08		1902	1081		113	putative protein of metazoan origin (slawmee) mRNA.
slawpey	slawpey.aSep08		851	363		59	putative protein (slawpey) mRNA.
slawpor	slawpor.aSep08		450	343		38	putative protein (slawpor) mRNA.
slawroy	slawroy.bSep08		34439	522	5	38	CRA a like (4.2 kD) (slawroy) alternative variant bSep08, mRNA.
slawroy	slawroy.cSep08		34431	456	5	33	CRA b like (3.8 kD) (slawroy) alternative variant cSep08, mRNA.
slawshaw	slawshaw.aSep08		5093	556		56	G elongation factor mitochondrial 2 (6.5 kD) (slawshaw) mRNA.
slawshee	slawshee.aSep08		9417	437		145	tubulin polyglutamylase ttl17 (slawshee) mRNA.
slawtu	slawtu.aSep08		2884	199		63	gag protein like (slawtu) mRNA.
slawvo	slawvo.aSep08		37227	292		29	putative protein (slawvo) mRNA.
slawwer	slawwer.aSep08		56261	471		156	putative protein of mammalian origin (slawwer) mRNA.
Slbp	Slbp.bSep08	681062	3206	1141	2	115	stem-loop binding protein (Slbp) alternative variant bSep08, mRNA.
Slbp	Slbp.cSep08	681062	1457	762	1	58	stem-loop binding protein (6.5 kD) (Slbp) alternative variant cSep08, mRNA.
Slc1a1	Slc1a1.bSep08	25550	8122	1622	4	205	solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1 (21.7 kD) (Slc1a1) alternative variant bSep08, mRNA.
Slc1a2	Slc1a2.cSep08	29482	32707	885	4	222	solute carrier family 1 (glial high affinity glutamate transporter), member 2 (Slc1a2) alternative variant cSep08, mRNA.
Slc1a2	Slc1a2.eSep08	29482	1841	338	2	31	solute carrier family 1 (glial high affinity glutamate transporter), member 2 (3.3 kD) (Slc1a2) alternative variant eSep08, mRNA.
Slc1a3	Slc1a3.bSep08	29483	4407	423	1	124	solute carrier family 1 (glial high affinity glutamate transporter), member 3 (Slc1a3) alternative variant bSep08, mRNA.
Slc1a4	Slc1a4.bSep08	305540	10479	2349	2	259	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4 (27.4 kD) (Slc1a4) alternative variant bSep08, mRNA.

Slc1a6	Slc1a6.bSep08	84012	5215	506		117	solute carrier family 1 (high affinity aspartate/glutamate transporter), member 6 (Slc1a6) alternative variant bSep08, mRNA.
Slc2a4	Slc2a4.bSep08	25139	1041	710	4	236	solute carrier family 2 (facilitated glucose transporter), member 4 (Slc2a4) alternative variant bSep08, mRNA.
Slc2a5	Slc2a5.bSep08	65197	3037	672	3	145	solute carrier family 2 (facilitated glucose/fructose transporter), member 5 (Slc2a5) alternative variant bSep08, mRNA.
Slc2a5	Slc2a5.cSep08	65197	1928	1360	2	132	solute carrier family 2 (facilitated glucose/fructose transporter), member 5 (14.8 kD) (Slc2a5) alternative variant cSep08, mRNA.
Slc2a8	Slc2a8.bSep08	85256	2447	1385	5	288	solute carrier family 2, (facilitated glucose transporter) member 8 (Slc2a8) alternative variant bSep08, mRNA.
Slc2a8	Slc2a8.cSep08	85256	4390	1323	5	232	solute carrier family 2, (facilitated glucose transporter) member 8 (Slc2a8) alternative variant cSep08, mRNA.
Slc2a8	Slc2a8.dSep08	85256	7685	481	4	154	solute carrier family 2, (facilitated glucose transporter) member 8 (Slc2a8) alternative variant dSep08, mRNA.
Slc2a8	Slc2a8.eSep08	85256	1348	453	2	102	solute carrier family 2, (facilitated glucose transporter) member 8 (Slc2a8) alternative variant eSep08, mRNA.
Slc2a9	Slc2a9.aSep08	501925	48365	2436	6	229	solute carrier family 2 (facilitated glucose transporter), member 9 (Slc2a9) alternative variant aSep08, mRNA.
Slc2a9	Slc2a9.bSep08	501925	96567	363	3	71	solute carrier family 2 (facilitated glucose transporter), member 9 (Slc2a9) alternative variant bSep08, mRNA.
Slc2a13	Slc2a13.bSep08	171147	5352	1257	2	188	solute carrier family 2 (facilitated glucose transporter), member 13 (21.1 kD) (Slc2a13) alternative variant bSep08, mRNA.
Slc3a1	Slc3a1.bSep08	29484	3637	285	1	79	solute carrier family 3, member 1 (Slc3a1) alternative variant bSep08, mRNA.
Slc3a2	Slc3a2.bSep08	50567	5620	738	5	246	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (Slc3a2) alternative variant bSep08, mRNA.
Slc3a2	Slc3a2.cSep08	50567	3972	841	4	246	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (Slc3a2) alternative variant cSep08, mRNA.
Slc3a2	Slc3a2.eSep08	50567	892	779	2	92	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (Slc3a2) alternative variant eSep08, mRNA.
Slc3a2	Slc3a2.fSep08	50567	1414	1217	3	33	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (3.7 kD) (Slc3a2) alternative variant fSep08, mRNA.
Slc4a1	Slc4a1.bSep08	24779	11515	4497	10	850	solute carrier family 4 (anion exchanger), member 1 (94.4 kD) (Slc4a1) alternative variant bSep08, mRNA.
Slc4a1	Slc4a1.cSep08	24779	9988	2020	1	478	solute carrier family 4 (anion exchanger), member 1 (Slc4a1) alternative variant cSep08, mRNA.
Slc4a1ap	Slc4a1ap.aSep08	298805	27038	1752	13	432	solute carrier family 4 member 1 adaptor protein (Slc4a1ap) alternative variant aSep08, mRNA.

Slc4a1ap	Slc4a1ap.bSep08	298805	17809	1357	8	253	solute carrier family 4 member 1 adaptor protein CRA d (Slc4a1ap) alternative variant bSep08, mRNA.
Slc4a1ap	Slc4a1ap.dSep08	298805	2264	894	2	53	putative protein (Slc4a1ap) alternative variant dSep08, mRNA.
Slc4a2	Slc4a2.dSep08	24780	646	555	2	81	solute carrier family 4 (anion exchanger), member 2 (9.3 kD) (Slc4a2) alternative variant dSep08, mRNA.
Slc4a3	Slc4a3.bSep08	24781	2503	1416	5	267	solute carrier family 4 (anion exchanger), member 3 (29.6 kD) (Slc4a3) alternative variant bSep08, mRNA.
Slc4a3	Slc4a3.cSep08	24781	1674	772	4	219	solute carrier family 4 (anion exchanger), member 3 (Slc4a3) alternative variant cSep08, mRNA.
Slc4a4	Slc4a4.bSep08	84484	111820	598	1	199	solute carrier family 4 (anion exchanger), member 4 (Slc4a4) alternative variant bSep08, mRNA.
Slc4a5	Slc4a5.bSep08	297386	14790	757	5	252	solute carrier family 4, sodium bicarbonate cotransporter, member 5 (Slc4a5) alternative variant bSep08, mRNA.
Slc4a5	Slc4a5.cSep08	297386	21356	599	5	199	solute carrier family 4, sodium bicarbonate cotransporter, member 5 (Slc4a5) alternative variant cSep08, mRNA.
Slc4a5	Slc4a5.dSep08	297386	20397	402	4	133	solute carrier family 4, sodium bicarbonate cotransporter, member 5 (Slc4a5) alternative variant dSep08, mRNA.
Slc4a7	Slc4a7.bSep08	117955	10095	746	6	218	solute carrier family 4, sodium bicarbonate cotransporter, member 7 (Slc4a7) alternative variant bSep08, mRNA.
Slc4a8	Slc4a8.bSep08	315311	6920	418	4	139	solute carrier family 4 (anion exchanger), member 8 (Slc4a8) alternative variant bSep08, mRNA.
Slc4a8	Slc4a8.cSep08	315311	5831	1789	4	108	solute carrier family 4 (anion exchanger), member 8 (Slc4a8) alternative variant cSep08, mRNA.
Slc4a8	Slc4a8.dSep08	315311	3765	487	3	100	solute carrier family 4 (anion exchanger), member 8 (Slc4a8) alternative variant dSep08, mRNA.
Slc4a8	Slc4a8.fSep08	315311	2859	516	3	57	solute carrier family 4 (anion exchanger), member 8 (Slc4a8) alternative variant fSep08, mRNA.
Slc4a10	Slc4a10.bSep08	295645	183738	1716	5	391	solute carrier family 4, sodium bicarbonate transporter-like, member 10 (Slc4a10) alternative variant bSep08, mRNA.
Slc4a10	Slc4a10.cSep08	295645	21318	499	5	140	solute carrier family 4, sodium bicarbonate transporter-like, member 10 (Slc4a10) alternative variant cSep08, mRNA.
Slc4a10	Slc4a10.fSep08	295645	1840	300	2	23	solute carrier family 4, sodium bicarbonate transporter-like, member 10 (Slc4a10) alternative variant fSep08, mRNA.
Slc4a11	Slc4a11.bSep08	311423	1682	1401	4	132	solute carrier family 4, sodium bicarbonate transporter-like, member 11 (Slc4a11) alternative variant bSep08, mRNA.
Slc4a11	Slc4a11.cSep08	311423	822	665	3	125	solute carrier family 4, sodium bicarbonate transporter-like, member 11 (Slc4a11) alternative variant cSep08, mRNA.
Slc5a2	Slc5a2.bSep08	64522	2415	678	5	225	solute carrier family 5 (sodium/glucose cotransporter), member 2 (Slc5a2) alternative variant bSep08, mRNA.
Slc5a6	Slc5a6.bSep08	170551	11767	3287	16	634	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (68.6 kD) (Slc5a6) alternative variant bSep08, mRNA.
Slc5a6	Slc5a6.cSep08	170551	5853	1897	12	530	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (56.9 kD) (Slc5a6) alternative variant cSep08, mRNA.

Slc5a6	Slc5a6.dSep08	170551	2928	742	7	247	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (Slc5a6) alternative variant dSep08, mRNA.
Slc5a6	Slc5a6.eSep08	170551	1122	910	1	84	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (Slc5a6) alternative variant eSep08, mRNA.
Slc5a6	Slc5a6.fSep08	170551	4249	386	3	11	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (1.4 kD) (Slc5a6) alternative variant fSep08, mRNA.
Slc5a6	Slc5a6.gSep08	170551	4043	344	2	37	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (Slc5a6) alternative variant gSep08, mRNA.
Slc5a8	Slc5a8.aSep08	500820	40799	1759		563	solute carrier family 5 (iodide transporter), member 8 (Slc5a8) mRNA.
Slc5a9	Slc5a9.bSep08	366441	1040	450	4	149	solute carrier family 5 (sodium/glucose cotransporter), member 9 (Slc5a9) alternative variant bSep08, mRNA.
Slc5a9	Slc5a9.cSep08	366441	526	295	2	87	solute carrier family 5 (sodium/glucose cotransporter), member 9 (Slc5a9) alternative variant cSep08, mRNA.
Slc5a10	Slc5a10.bSep08	303205	47933	2214	15	540	solute carrier family 5 (sodium/glucose cotransporter), member 10 (59.1 kD) (Slc5a10) alternative variant bSep08, mRNA.
Slc5a10	Slc5a10.cSep08	303205	13271	796	7	262	solute carrier family 5 (sodium/glucose cotransporter), member 10 (Slc5a10) alternative variant cSep08, mRNA.
Slc5a10	Slc5a10.dSep08	303205	11892	734	7	230	solute carrier family 5 (sodium/glucose cotransporter), member 10 (24.4 kD) (Slc5a10) alternative variant dSep08, mRNA.
Slc5a10	Slc5a10.fSep08	303205	1142	310	2	97	solute carrier family 5 (sodium/glucose cotransporter), member 10 (Slc5a10) alternative variant fSep08, mRNA.
Slc5a11	Slc5a11.aSep08	252854	60262	2658	15	673	solute carrier family 5 (sodium/glucose cotransporter), member 11 (Slc5a11) alternative variant aSep08, mRNA.
Slc5a11	Slc5a11.bSep08	252854	18827	399		52	solute carrier family 5 (sodium/glucose cotransporter), member 11 (Slc5a11) alternative variant bSep08, mRNA.
Slc6a5	Slc6a5.aSep08	171148	1624	313		46	solute carrier family 6 (neurotransmitter transporter, glycine), member 5 (Slc6a5) mRNA.
Slc6a6	Slc6a6.bSep08	29464	41564	1436	6	235	solute carrier family 6 (neurotransmitter transporter, taurine), member 6 (Slc6a6) alternative variant bSep08, mRNA.
Slc6a9	Slc6a9.bSep08	116509	7992	602	4	183	solute carrier family 6 (neurotransmitter transporter, glycine), member 9 (Slc6a9) alternative variant bSep08, mRNA.
Slc6a9	Slc6a9.dSep08	116509	760	669	2	91	solute carrier family 6 (neurotransmitter transporter, glycine), member 9 (Slc6a9) alternative variant dSep08, mRNA.
Slc6a9	Slc6a9.eSep08	116509	2694	1290	3	91	solute carrier family 6 (neurotransmitter transporter, glycine), member 9 (10.0 kD) (Slc6a9) alternative variant eSep08, mRNA.

Slc6a9	Slc6a9.fSep08	116509	528	400	2	88	solute carrier family 6 (neurotransmitter transporter, glycine), member 9 (Slc6a9) alternative variant fSep08, mRNA.
Slc6a12	Slc6a12.bSep08	50676	11004	742	5	180	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12 (Slc6a12) alternative variant bSep08, mRNA.
Slc6a12	Slc6a12.cSep08	50676	6147	399	3	109	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12 (Slc6a12) alternative variant cSep08, mRNA.
Slc6a13	Slc6a13.bSep08	171163	22616	776	7	180	solute carrier family 6 (neurotransmitter transporter, GABA), member 13 (Slc6a13) alternative variant bSep08, mRNA.
Slc6a13	Slc6a13.cSep08	171163	2216	899	3	170	solute carrier family 6 (neurotransmitter transporter, GABA), member 13 (Slc6a13) alternative variant cSep08, mRNA.
Slc6a20	Slc6a20.cSep08	113918	3263	400	2	133	solute carrier family 6 (neurotransmitter transporter), member 20 (Slc6a20) alternative variant cSep08, mRNA.
Slc7a1	Slc7a1.aSep08	25648	10298	939		313	solute carrier family 7 (cationic amino acid transporter, y+ system), member 1 (Slc7a1) mRNA.
Slc7a2	Slc7a2.cSep08	64554	3721	1801	3	419	solute carrier family 7 (cationic amino acid transporter, y+ system), member 2 (Slc7a2) alternative variant cSep08, mRNA.
Slc7a4	Slc7a4.bSep08	303787	919	806	2	233	solute carrier family 7 (cationic amino acid transporter, y+ system), member 4 (Slc7a4) alternative variant bSep08, mRNA.
Slc7a4	Slc7a4.dSep08	303787	883	411	2	61	solute carrier family 7 (cationic amino acid transporter, y+ system), member 4 (6.5 kD) (Slc7a4) alternative variant dSep08, mRNA.
Slc7a6	Slc7a6.cSep08	307811	2684	731	3	43	solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 (Slc7a6) alternative variant cSep08, mRNA.
Slc7a7	Slc7a7.bSep08	83509	31450	687	3	228	solute carrier family 7 member (Slc7a7) alternative variant bSep08, mRNA.
Slc7a7	Slc7a7.cSep08	83509	38996	731	7	213	solute carrier family 7 member (Slc7a7) alternative variant cSep08, mRNA.
Slc7a7	Slc7a7.dSep08	83509	35492	1042	3	198	solute carrier family 7 member (Slc7a7) alternative variant dSep08, mRNA.
Slc7a7	Slc7a7.eSep08	83509	35679	682	3	169	solute carrier family 7 member (17.5 kD) (Slc7a7) alternative variant eSep08, mRNA.
Slc7a7	Slc7a7.fSep08	83509	41387	736	6	123	solute carrier family 7 member (Slc7a7) alternative variant fSep08, mRNA.
Slc7a7	Slc7a7.gSep08	83509	3798	294	3	82	putative protein (8.5 kD) (Slc7a7) alternative variant gSep08, mRNA.
Slc7a7	Slc7a7.hSep08	83509	39570	734	6	71	solute carrier family 7 member (Slc7a7) alternative variant hSep08, mRNA.

Slc7a8	Slc7a8.bSep08	84551	17978	491	3	130	solute carrier family 7 (cationic amino acid transporter, y+ system), member 8 (13.1 kD) (Slc7a8) alternative variant bSep08, mRNA.
Slc7a9	Slc7a9.bSep08	116726	4750	1037	3	213	solute carrier family 7 (cationic amino acid transporter, y+ system), member 9 (Slc7a9) alternative variant bSep08, mRNA.
Slc7a9	Slc7a9.cSep08	116726	5273	1134	4	191	solute carrier family 7 (cationic amino acid transporter, y+ system), member 9 (Slc7a9) alternative variant cSep08, mRNA.
Slc7a10	Slc7a10.bSep08	114518	510	423	2	91	solute carrier family 7 (cationic amino acid transporter, y+ system), member 10 (Slc7a10) alternative variant bSep08, mRNA.
Slc7a11	Slc7a11.bSep08	310392	28661	427	1	50	solute carrier family 7 (cationic amino acid transporter, y+ system), member 11 (Slc7a11) alternative variant bSep08, mRNA.
Slc7a15	Slc7a15.bSep08	298873	20073	1480	4	207	solute carrier family 7 (cationic amino acid transporter, y+ system), member 15 (23.1 kD) (Slc7a15) alternative variant bSep08, mRNA.
Slc7a15	Slc7a15.cSep08	298873	14297	426	2	91	solute carrier family 7 (cationic amino acid transporter, y+ system), member 15 (Slc7a15) alternative variant cSep08, mRNA.
Slc9a3r2	Slc9a3r2.bSep08	116501	4381	773	2	207	solute carrier family 9 (sodium/hydrogen exchanger), member 3 regulator 2 (Slc9a3r2) alternative variant bSep08, mRNA.
Slc9a5	Slc9a5.bSep08	192215	11588	1785	8	120	solute carrier family 9 (sodium/hydrogen exchanger), member 5 (13.6 kD) (Slc9a5) alternative variant bSep08, mRNA.
Slc9a5	Slc9a5.cSep08	192215	1050	565	2	75	solute carrier family 9 (sodium/hydrogen exchanger), member 5 (Slc9a5) alternative variant cSep08, mRNA.
Slc9a6	Slc9a6.aSep08	302863	18188	2920		182	solute carrier family 9 (sodium/hydrogen exchanger), member 6 (Slc9a6) mRNA.
Slc9a8	Slc9a8.aSep08	311651	24521	773		191	solute carrier family 9 (sodium/hydrogen exchanger), member 8 (Slc9a8) mRNA.
Slc10a3	Slc10a3.bSep08	501665	2478	566	1	81	solute carrier family 10 (sodium/bile acid cotransporter family), member 3 (8.4 kD) (Slc10a3) alternative variant bSep08, mRNA.
Slc11a1	Slc11a1.bSep08	316519	6757	1197	9	339	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1 (37.5 kD) (Slc11a1) alternative variant bSep08, mRNA.
Slc11a2	Slc11a2.aSep08	25715	16071	1251	9	297	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (Slc11a2) alternative variant aSep08, mRNA.
Slc11a2	Slc11a2.bSep08	25715	2371	447	2	71	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (Slc11a2) alternative variant bSep08, mRNA.
Slc12a1	Slc12a1.bSep08	25065	18855	682	6	227	solute carrier family 12, member 1 (Slc12a1) alternative variant bSep08, mRNA.



Slc12a1	Slc12a1.cSep08	25065	2294	1311	2	72	solute carrier family 12, member 1 (Slc12a1) alternative variant cSep08, mRNA.
Slc12a2	Slc12a2.bSep08	83629	17863	797		265	solute carrier family 12 (sodium/potassium/chloride transporters), member 2 (Slc12a2) alternative variant bSep08, mRNA.
Slc12a4	Slc12a4.bSep08	29501	3071	1190	6	236	solute carrier family 12, member 4 (Slc12a4) alternative variant bSep08, mRNA.
Slc12a4	Slc12a4.cSep08	29501	1340	535	4	177	solute carrier family 12, member 4 (Slc12a4) alternative variant cSep08, mRNA.
Slc12a5	Slc12a5.aSep08	171373	40120	5714	26	1158	solute carrier family 12, member 5 (Slc12a5) alternative variant aSep08, mRNA.
Slc12a5	Slc12a5.bSep08	171373	2818	460	4	152	solute carrier family 12, member 5 (Slc12a5) alternative variant bSep08, mRNA.
Slc12a5	Slc12a5.cSep08	171373	1010	510	2	78	solute carrier family 12, member 5 (Slc12a5) alternative variant cSep08, mRNA.
Slc12a7	Slc12a7.aSep08	308069	30935	1351	9	450	solute carrier family 12, member 7 (Slc12a7) alternative variant aSep08, mRNA.
Slc12a7	Slc12a7.bSep08	308069	12417	2556	8	284	solute carrier family 12, member 7 (Slc12a7) alternative variant bSep08, mRNA.
Slc12a7	Slc12a7.cSep08	308069	12405	2529	7	275	solute carrier family 12, member 7 (Slc12a7) alternative variant cSep08, mRNA.
Slc12a7	Slc12a7.dSep08	308069	53379	1474	7	266	solute carrier family 12, member 7 (Slc12a7) alternative variant dSep08, mRNA.
Slc12a7	Slc12a7.eSep08	308069	2272	380	3	81	solute carrier family 12, member 7 (Slc12a7) alternative variant eSep08, mRNA.
Slc12a8	Slc12a8.bSep08	266733	37095	550	3	59	solute carrier family 12 (potassium/chloride transporters), member 8 (Slc12a8) alternative variant bSep08, mRNA.
Slc13a4	Slc13a4.bSep08	503568	14496	1515	4	189	solute carrier family 13 (sodium/sulfate symporters), member 4 (Slc13a4) alternative variant bSep08, mRNA.
Slc13a4	Slc13a4.cSep08	503568	17248	373	3	124	solute carrier family 13 (sodium/sulfate symporters), member 4 (Slc13a4) alternative variant cSep08, mRNA.
Slc14a2	Slc14a2.dSep08	54302	8056	541	4	123	urea transporter (Slc14a2) alternative variant dSep08, mRNA.
Slc16a1	Slc16a1.bSep08	25027	13541	425	3	115	solute carrier family 16 (monocarboxylic acid transporters), member 1 (Slc16a1) alternative variant bSep08, mRNA.
Slc16a2	Slc16a2.aSep08	259248	6623	742	3	246	solute carrier family 16 (monocarboxylic acid transporters), member 2 (Slc16a2) alternative variant aSep08, mRNA.
Slc16a3	Slc16a3.bSep08	80878	2853	1024	4	235	solute carrier family 16 (monocarboxylic acid transporters), member 3 (25.5 kD) (Slc16a3) alternative variant bSep08, mRNA.
Slc16a3	Slc16a3.dSep08	80878	1697	368	3	95	solute carrier family 16 (monocarboxylic acid transporters), member 3 (Slc16a3) alternative variant dSep08, mRNA.
Slc16a3	Slc16a3.eSep08	80878	1303	340	2	74	solute carrier family 16 (monocarboxylic acid transporters), member 3 (Slc16a3) alternative variant eSep08, mRNA.
Slc16a4	Slc16a4.aSep08	295356	22467	1485	8	327	solute carrier family 16 (monocarboxylic acid transporters), member 4 (36.3 kD) (Slc16a4) alternative variant aSep08, mRNA.

Slc16a4	Slc16a4.bSep08	295356	8302	705	5	175	solute carrier family 16 (monocarboxylic acid transporters), member 4 (Slc16a4) alternative variant bSep08, mRNA.
Slc16a6	Slc16a6.bSep08	303772	17521	1461	6	416	solute carrier family 16 (monocarboxylic acid transporters), member 6 (Slc16a6) alternative variant bSep08, mRNA.
Slc16a6	Slc16a6.cSep08	303772	14247	738	4	153	solute carrier family 16 (monocarboxylic acid transporters), member 6 (Slc16a6) alternative variant cSep08, mRNA.
Slc16a6	Slc16a6.eSep08	303772	6633	418	2	42	solute carrier family 16 (monocarboxylic acid transporters), member 6 (4.9 kD) (Slc16a6) alternative variant eSep08, mRNA.
Slc16a7	Slc16a7.bSep08	29735	79810	705	5	162	solute carrier family 16 (monocarboxylic acid transporters), member 7 (Slc16a7) alternative variant bSep08, mRNA.
Slc16a7	Slc16a7.cSep08	29735	7097	396	2	55	solute carrier family 16 (monocarboxylic acid transporters), member 7 (Slc16a7) alternative variant cSep08, mRNA.
Slc16a11	Slc16a11.aSep08	287450	3829	2023	7	575	solute carrier family 16 (monocarboxylic acid transporters), member 11 (58.3 kD) (Slc16a11) alternative variant aSep08, mRNA.
Slc16a11	Slc16a11.cSep08	287450	2329	865	2	143	solute carrier family 16 (monocarboxylic acid transporters), member 11 (14.8 kD) (Slc16a11) alternative variant cSep08, mRNA.
Slc16a11	Slc16a11.dSep08	287450	1948	558	4	83	solute carrier family 16 (monocarboxylic acid transporters), member 11 (8.8 kD) (Slc16a11) alternative variant dSep08, mRNA.
Slc16a12	Slc16a12.aSep08	309525	2408	880		168	solute carrier family 16 (monocarboxylic acid transporters), member 12 (Slc16a12) mRNA.
Slc17a3	Slc17a3.aSep08	266730	21974	1536	10	487	solute carrier family 17 (sodium phosphate), member 3 (Slc17a3) alternative variant aSep08, mRNA.
Slc17a3	Slc17a3.bSep08	266730	11341	766	3	212	solute carrier family 17 (sodium phosphate), member 3 (Slc17a3) alternative variant bSep08, mRNA.
Slc17a3	Slc17a3.cSep08	266730	19497	1276	5	109	solute carrier family 17 (sodium phosphate), member 3 (Slc17a3) alternative variant cSep08, mRNA.
Slc17a3	Slc17a3.dSep08	266730	16276	1249	4	111	solute carrier family 17 (sodium phosphate), member 3 (12.1 kD) (Slc17a3) alternative variant dSep08, complete mRNA.
Slc17a4	Slc17a4.aSep08	679784	6568	1029		291	solute carrier family 17 (sodium phosphate), member 4 (31.8 kD) (Slc17a4) mRNA.
Slc17a7	Slc17a7.bSep08	116638	12010	2896	12	585	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 (Slc17a7) alternative variant bSep08, complete mRNA.
Slc17a7	Slc17a7.cSep08	116638	5424	600	1	180	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 (Slc17a7) alternative variant cSep08, mRNA.
Slc18a2	Slc18a2.aSep08	25549	7154	495		54	solute carrier family 18 (vesicular monoamine), member 2 (Slc18a2) mRNA.
Slc19a1	Slc19a1.cSep08	29723	7755	2086	4	278	solute carrier family 19 (sodium/hydrogen exchanger), member 1 (Slc19a1) alternative variant cSep08, mRNA.
Slc19a1	Slc19a1.dSep08	29723	6202	852	3	158	solute carrier family 19 (sodium/hydrogen exchanger), member 1 (Slc19a1) alternative variant dSep08, mRNA.

Slc20a1	Slc20a1.bSep08	81826	6006	872	4	241	solute carrier family 20 (phosphate transporter), member 1 (Slc20a1) alternative variant bSep08, mRNA.
Slc20a1	Slc20a1.cSep08	81826	2300	1617	2	99	solute carrier family 20 (phosphate transporter), member 1 (10.8 kD) (Slc20a1) alternative variant cSep08, mRNA.
Slc21a4	Slc21a4.bSep08	80899	23909	824	6	274	organic anion transporter K2 (Slc21a4) alternative variant bSep08, mRNA.
Slc21a4	Slc21a4.cSep08	80899	4960	736	4	207	organic anion transporter K5 (Slc21a4) alternative variant cSep08, mRNA.
Slc21a4	Slc21a4.dSep08	80899	30510	670	5	185	organic anion transporter K5 (Slc21a4) alternative variant dSep08, mRNA.
Slc21a4	Slc21a4.eSep08	80899	13723	775	5	184	organic anion transporter K6 (Slc21a4) alternative variant eSep08, mRNA.
Slc21a4	Slc21a4.fSep08	80899	5450	298	2	42	organic anion transporter K10 (Slc21a4) alternative variant fSep08, mRNA.
Slc22a1	Slc22a1.bSep08	24904	9081	553	4	184	solute carrier family 22 (organic cation transporter), member 1 (Slc22a1) alternative variant bSep08, mRNA.
Slc22a1	Slc22a1.cSep08	24904	4792	518	2	172	solute carrier family 22 (organic cation transporter), member 1 (Slc22a1) alternative variant cSep08, mRNA.
Slc22a1	Slc22a1.dSep08	24904	10875	773	3	154	solute carrier family 22 (organic cation transporter), member 1 (Slc22a1) alternative variant dSep08, mRNA.
Slc22a2	Slc22a2.bSep08	29503	14732	721	3	239	solute carrier family 22 (organic cation transporter), member 2 (Slc22a2) alternative variant bSep08, mRNA.
Slc22a2	Slc22a2.cSep08	29503	6132	386	3	123	solute carrier family 22 (organic cation transporter), member 2 (Slc22a2) alternative variant cSep08, mRNA.
Slc22a5	Slc22a5.bSep08	29726	27009	2778	1	469	solute carrier family 22 (organic cation transporter), member 5 (52.6 kD) (Slc22a5) alternative variant bSep08, complete mRNA.
Slc22a6	Slc22a6.bSep08	29509	3039	961	4	194	solute carrier family 22 member 6 (20.9 kD) (Slc22a6) alternative variant bSep08, complete mRNA.
Slc22a6	Slc22a6.cSep08	29509	2783	823	3	144	solute carrier family 22 member 6 (15.5 kD) (Slc22a6) alternative variant cSep08, mRNA.
Slc22a6	Slc22a6.dSep08	29509	1050	920	2	51	solute carrier family 22 member 6 (Slc22a6) alternative variant dSep08, mRNA.
Slc22a7	Slc22a7.bSep08	89776	4350	751	5	117	solute carrier family 22 (organic anion transporter), member 7 (Slc22a7) alternative variant bSep08, mRNA.
Slc22a8	Slc22a8.bSep08	83500	4236	757	6	236	solute carrier family 22 member 8 (Slc22a8) alternative variant bSep08, mRNA.
Slc22a8	Slc22a8.cSep08	83500	5045	406	4	134	solute carrier family 22 member 8 (Slc22a8) alternative variant cSep08, mRNA.
Slc22a8	Slc22a8.dSep08	83500	2750	464	3	126	putative protein (13.9 kD) (Slc22a8) alternative variant dSep08, mRNA.
Slc22a12	Slc22a12.bSep08	365398	4085	737	2	220	solute carrier family 22 (organic anion/cation transporter), member 12 (Slc22a12) alternative variant bSep08, mRNA.
Slc22a14	Slc22a14.aSep08	316061	13119	1777		529	solute carrier family 22 (organic cation transporter), member 14 (Slc22a14) alternative variant aSep08, mRNA.
Slc22a14	Slc22a14.bSep08	316061	4339	710		205	solute carrier family 22 (organic cation transporter), member 14 (Slc22a14) alternative variant bSep08, mRNA.

Slc22a15	Slc22a15.bSep08	310732	24614	1070	4	298	solute carrier family 22, member 15 (Slc22a15) alternative variant bSep08, mRNA.
Slc22a15	Slc22a15.cSep08	310732	1611	1311	2	51	solute carrier family 22, member 15 (5.8 kD) (Slc22a15) alternative variant cSep08, mRNA.
Slc22a17	Slc22a17.bSep08	305886	1261	1038	3	237	solute carrier family 22 member 17 CRA c (24.2 kD) (Slc22a17) alternative variant bSep08, complete mRNA.
Slc22a17	Slc22a17.cSep08	305886	784	647	2	132	solute carrier family 22 member 17 (Slc22a17) alternative variant cSep08, mRNA.
Slc22a17	Slc22a17.dSep08	305886	625	348	3	100	solute carrier family 22 member 17 (Slc22a17) alternative variant dSep08, mRNA.
Slc22a18	Slc22a18.bSep08	309131	16595	704	6	181	solute carrier family 22 (organic cation transporter), member 18 (Slc22a18) alternative variant bSep08, mRNA.
Slc22a18	Slc22a18.cSep08	309131	6272	443	4	147	solute carrier family 22 (organic cation transporter), member 18 (Slc22a18) alternative variant cSep08, mRNA.
Slc22a18	Slc22a18.dSep08	309131	1083	448	3	125	solute carrier family 22 (organic cation transporter), member 18 (Slc22a18) alternative variant dSep08, mRNA.
Slc22a23	Slc22a23.aSep08	64559	162464	2455		685	solute carrier family 22, member 23 (Slc22a23) mRNA.
Slc23a2	Slc23a2.bSep08	50622	4139	426	1	96	solute carrier family 23 (nucleobase transporters), member 2 (Slc23a2) alternative variant bSep08, mRNA.
Slc23a3	Slc23a3.bSep08	367298	3532	1347	2	214	solute carrier family 23 (nucleobase transporters), member 3 (23.3 kD) (Slc23a3) alternative variant bSep08, mRNA.
Slc24a3	Slc24a3.aSep08	85267	37820	1327		360	solute carrier family 24 (sodium/potassium/calcium exchanger), member 3 (Slc24a3) mRNA.
Slc24a4	Slc24a4.aSep08	314396	113005	2619	15	566	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4 (62.9 kD) (Slc24a4) alternative variant aSep08, mRNA.
Slc24a4	Slc24a4.cSep08	314396	1800	537	1	86	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4 (Slc24a4) alternative variant cSep08, mRNA.
Slc24a6	Slc24a6.bSep08	498185	6856	930	6	224	solute carrier family 24 (sodium/potassium/calcium exchanger), member 6 (Slc24a6) alternative variant bSep08, mRNA.
Slc25a1	Slc25a1.bSep08	29743	2156	839	8	275	solute carrier family 25 (mitochondrial carrier, citrate transporter), member 1 (Slc25a1) alternative variant bSep08, mRNA.
Slc25a1	Slc25a1.cSep08	29743	1280	580	5	193	solute carrier family 25 (mitochondrial carrier, citrate transporter), member 1 (Slc25a1) alternative variant cSep08, mRNA.
Slc25a1	Slc25a1.dSep08	29743	1649	825	6	153	solute carrier family 25 (mitochondrial carrier, citrate transporter), member 1 (Slc25a1) alternative variant dSep08, mRNA.
Slc25a1	Slc25a1.eSep08	29743	3014	2174	4	138	solute carrier family 25 (mitochondrial carrier, citrate transporter), member 1 (15.1 kD) (Slc25a1) alternative variant eSep08, complete mRNA.
Slc25a3	Slc25a3.aSep08	245959	7490	1377	8	357	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (39.6 kD) (Slc25a3) alternative variant aSep08, mRNA.

Slc25a3	Slc25a3.cSep08	245959	5443	760	4	184	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (Slc25a3) alternative variant cSep08, mRNA.
Slc25a3	Slc25a3.dSep08	245959	507	426	2	102	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (Slc25a3) alternative variant dSep08, mRNA.
Slc25a3	Slc25a3.eSep08	245959	4037	2785	3	94	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (Slc25a3) alternative variant eSep08, mRNA.
Slc25a3	Slc25a3.gSep08	245959	5564	806	6	89	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (9.6 kD) (Slc25a3) alternative variant gSep08, mRNA.
Slc25a3	Slc25a3.iSep08	245959	706	486	2	89	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3 (9.5 kD) (Slc25a3) alternative variant iSep08, mRNA.
Slc25a4	Slc25a4.cSep08	85333	847	364	2	63	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4 (Slc25a4) alternative variant cSep08, mRNA.
Slc25a5	Slc25a5.bSep08	25176	1777	780	2	218	solute carrier family 25 member 5 (23.8 kD) (Slc25a5) alternative variant bSep08, mRNA.
Slc25a5	Slc25a5.cSep08	25176	1300	964	2	98	solute carrier family 25 member 5 CRA a (11.1 kD) (Slc25a5) alternative variant cSep08, mRNA.
Slc25a10	Slc25a10.bSep08	170943	3340	722	9	222	solute carrier family 25 member 10 (Slc25a10) alternative variant bSep08, mRNA.
Slc25a10	Slc25a10.cSep08	170943	6707	721	9	221	solute carrier family 25 member 10 CRA a (24.2 kD) (Slc25a10) alternative variant cSep08, mRNA.
Slc25a10	Slc25a10.dSep08	170943	7499	2204	11	163	solute carrier family 25 member 10 CRA a (17.9 kD) (Slc25a10) alternative variant dSep08, complete mRNA.
Slc25a10	Slc25a10.eSep08	170943	3228	707	2	67	solute carrier family 25 member 10 (Slc25a10) alternative variant eSep08, mRNA.
Slc25a10	Slc25a10.fSep08	170943	1214	695	3	65	solute carrier family 25 member 10 CRA a (6.8 kD) (Slc25a10) alternative variant fSep08, mRNA.
Slc25a10	Slc25a10.gSep08	170943	4963	343	5	50	solute carrier family 25 member 10 (Slc25a10) alternative variant gSep08, mRNA.
Slc25a11	Slc25a11.bSep08	64201	1052	528	2	134	solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 (14.3 kD) (Slc25a11) alternative variant bSep08, mRNA.
Slc25a12	Slc25a12.aSep08	362145	50176	476	5	155	solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (Slc25a12) alternative variant aSep08, mRNA.
Slc25a12	Slc25a12.bSep08	362145	50172	472	5	155	solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (Slc25a12) alternative variant bSep08, mRNA.
Slc25a12	Slc25a12.cSep08	362145	12552	356	3	26	solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (3.2 kD) (Slc25a12) alternative variant cSep08, complete mRNA.
Slc25a12	Slc25a12.dSep08	362145	28758	360	3	66	solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (Slc25a12) alternative variant dSep08, mRNA.

Slc25a13	Slc25a13.aSep08	362322	155085	2942	17	642	solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 13 (70.7 kD) (Slc25a13) alternative variant aSep08, mRNA.
Slc25a13	Slc25a13.bSep08	362322	25119	777	8	259	solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 13 (Slc25a13) alternative variant bSep08, mRNA.
Slc25a13	Slc25a13.cSep08	362322	33495	886	6	144	solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 13 (15.5 kD) (Slc25a13) alternative variant cSep08, mRNA.
Slc25a14	Slc25a14.bSep08	85263	28734	669	7	204	solute carrier family 25 (mitochondrial carrier, brain), member 14 (Slc25a14) alternative variant bSep08, mRNA.
Slc25a14	Slc25a14.cSep08	85263	9827	543	4	124	solute carrier family 25 (mitochondrial carrier, brain), member 14 (Slc25a14) alternative variant cSep08, mRNA.
Slc25a14	Slc25a14.dSep08	85263	10012	428	6	83	solute carrier family 25 (mitochondrial carrier, brain), member 14 (Slc25a14) alternative variant dSep08, mRNA.
Slc25a15	Slc25a15.bSep08	306574	21965	1800	5	340	solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15 (Slc25a15) alternative variant bSep08, mRNA.
Slc25a15	Slc25a15.cSep08	306574	4568	725	1	165	solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15 (Slc25a15) alternative variant cSep08, mRNA.
Slc25a15	Slc25a15.dSep08	306574	15026	879	4	150	solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15 (Slc25a15) alternative variant dSep08, mRNA.
Slc25a16	Slc25a16.aSep08	361836	17253	2979		256	solute carrier family 25 (mitochondrial carrier, Graves disease autoantigen), member 16 (Slc25a16) alternative variant aSep08, mRNA.
Slc25a16	Slc25a16.bSep08	361836	28323	3182		234	solute carrier family 25 (mitochondrial carrier, Graves disease autoantigen), member 16 (26.1 kD) (Slc25a16) alternative variant bSep08, mRNA.
Slc25a17	Slc25a17.bSep08	300083	11175	889	5	195	solute carrier family 25 member 17 (22.1 kD) (Slc25a17) alternative variant bSep08, mRNA.
Slc25a17	Slc25a17.cSep08	300083	3235	615	3	71	solute carrier family 25 member 17 CRA a (Slc25a17) alternative variant cSep08, mRNA.
Slc25a19	Slc25a19.bSep08	303676	11440	706	1	185	solute carrier family 25 (mitochondrial thiamine pyrophosphate carrier), member 19 (Slc25a19) alternative variant bSep08, mRNA.
Slc25a19	Slc25a19.cSep08	303676	10072	747	2	163	solute carrier family 25 (mitochondrial thiamine pyrophosphate carrier), member 19 (Slc25a19) alternative variant cSep08, mRNA.
Slc25a20	Slc25a20.bSep08	117035	15879	820	1	153	solute carrier family 25 (mitochondrial carnitine/acylcarnitine translocase), member 20 (Slc25a20) alternative variant bSep08, mRNA.
Slc25a22	Slc25a22.aSep08	309111	4609	1607	10	323	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (34.7 kD) (Slc25a22) alternative variant aSep08, mRNA.

Slc25a22	Slc25a22.cSep08	309111	5767	707	7	188	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (Slc25a22) alternative variant cSep08, mRNA.
Slc25a22	Slc25a22.eSep08	309111	5193	753	5	98	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (10.7 kD) (Slc25a22) alternative variant eSep08, mRNA.
Slc25a22	Slc25a22.fSep08	309111	1616	450	4	74	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (Slc25a22) alternative variant fSep08, mRNA.
Slc25a22	Slc25a22.gSep08	309111	728	404	3	74	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (Slc25a22) alternative variant gSep08, mRNA.
Slc25a22	Slc25a22.hSep08	309111	953	381	4	71	solute carrier family 25 (mitochondrial carrier, glutamate), member 22 (Slc25a22) alternative variant hSep08, mRNA.
Slc25a23	Slc25a23.bSep08	301113	5004	577	2	155	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 23 (Slc25a23) alternative variant bSep08, mRNA.
Slc25a23	Slc25a23.cSep08	301113	1710	1103	2	123	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 23 (Slc25a23) alternative variant cSep08, mRNA.
Slc25a25	Slc25a25.bSep08	246771	2440	410	2	136	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25 (Slc25a25) alternative variant bSep08, mRNA.
Slc25a25	Slc25a25.cSep08	246771	3831	714	2	101	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25 (Slc25a25) alternative variant cSep08, mRNA.
Slc25a26	Slc25a26.aSep08	362403	96867	2862	10	274	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26 (29.1 kD) (Slc25a26) alternative variant aSep08, mRNA.
Slc25a26	Slc25a26.bSep08	362403	84793	800	8	217	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26 (Slc25a26) alternative variant bSep08, mRNA.
Slc25a26	Slc25a26.dSep08	362403	11306	571	3	94	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26 (Slc25a26) alternative variant dSep08, mRNA.
Slc25a26	Slc25a26.eSep08	362403	19501	477	3	73	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26 (Slc25a26) alternative variant eSep08, mRNA.
Slc25a27	Slc25a27.aSep08	85262	24405	1219		344	solute carrier family 25, member 27 (38.1 kD) (Slc25a27) mRNA.
Slc25a28	Slc25a28.bSep08	688811	11232	1489	4	211	solute carrier family 25, member 28 (Slc25a28) alternative variant bSep08, mRNA.
Slc25a28	Slc25a28.cSep08	688811	10603	1995	3	182	solute carrier family 25, member 28 (Slc25a28) alternative variant cSep08, mRNA.
Slc25a28	Slc25a28.dSep08	688811	7736	428	2	112	solute carrier family 25, member 28 (Slc25a28) alternative variant dSep08, mRNA.
Slc25a29	Slc25a29.bSep08	314441	989	710	1	97	solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine transporter), member 29 (Slc25a29) alternative variant bSep08, mRNA.
Slc25a30	Slc25a30.aSep08	361074	14900	750		203	solute carrier family 25, member 30 (Slc25a30) mRNA.

Slc25a31	Slc25a31.aSep08	689108	11636	443		147	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 31 (Slc25a31) mRNA.
Slc25a32	Slc25a32.aSep08	315023	17824	2316		316	solute carrier family 25, member 32 (35.0 kD) (Slc25a32) mRNA.
Slc25a35	Slc25a35.bSep08	497933	1187	877	1	76	solute carrier family 25, member 35 (Slc25a35) alternative variant bSep08, mRNA.
Slc25a36	Slc25a36.aSep08	501039	33851	1801	6	397	solute carrier family 25, member 36 (Slc25a36) alternative variant aSep08, mRNA.
Slc25a36	Slc25a36.bSep08	501039	33668	4497	7	190	solute carrier family 25, member 36 (21.4 kD) (Slc25a36) alternative variant bSep08, mRNA.
Slc25a36	Slc25a36.cSep08	501039	9768	2196	4	154	solute carrier family 25, member 36 (17.4 kD) (Slc25a36) alternative variant cSep08, mRNA.
Slc25a36	Slc25a36.dSep08	501039	9145	1291	5	154	solute carrier family 25, member 36 (17.4 kD) (Slc25a36) alternative variant dSep08, mRNA.
Slc25a37	Slc25a37.cSep08	306000	3002	750	2	187	solute carrier family 25, member 37 (21.2 kD) (Slc25a37) alternative variant cSep08, mRNA.
Slc25a38	Slc25a38.bSep08	301067	5375	552	4	151	solute carrier family 25, member 38 (Slc25a38) alternative variant bSep08, mRNA.
Slc25a38	Slc25a38.cSep08	301067	3973	271	2	68	solute carrier family 25, member 38 (Slc25a38) alternative variant cSep08, mRNA.
Slc25a38	Slc25a38.dSep08	301067	8251	698	6	89	solute carrier family 25, member 38 (Slc25a38) alternative variant dSep08, mRNA.
Slc25a39	Slc25a39.bSep08	360636	4313	2098	11	341	solute carrier family 25 member 39 (37.2 kD) (Slc25a39) alternative variant bSep08, complete mRNA.
Slc25a39	Slc25a39.cSep08	360636	4176	1133	10	306	solute carrier family 25 member 39 (Slc25a39) alternative variant cSep08, mRNA.
Slc25a39	Slc25a39.dSep08	360636	2927	943	8	270	solute carrier family 25 member 39 (Slc25a39) alternative variant dSep08, mRNA.
Slc25a39	Slc25a39.eSep08	360636	1899	704	5	159	solute carrier family 25 member 39 (17.4 kD) (Slc25a39) alternative variant eSep08, mRNA.
Slc25a39	Slc25a39.fSep08	360636	1136	678	4	144	solute carrier family 25 member 39 (Slc25a39) alternative variant fSep08, mRNA.
Slc25a39	Slc25a39.hSep08	360636	2258	696	6	110	solute carrier family 25 member 39 (Slc25a39) alternative variant hSep08, mRNA.
Slc25a39	Slc25a39.iSep08	360636	513	410	2	38	putative protein (Slc25a39) alternative variant iSep08, mRNA.
Slc25a40	Slc25a40.bSep08	296813	7105	609	1	102	solute carrier family 25, member 40 (Slc25a40) alternative variant bSep08, mRNA.
Slc25a42	Slc25a42.bSep08	689414	4259	2381	3	192	solute carrier family 25, member 42 (Slc25a42) alternative variant bSep08, mRNA.
Slc25a45	Slc25a45.aSep08	689625	4273	734	3	186	solute carrier family 25, member 45 (Slc25a45) alternative variant aSep08, mRNA.
Slc26a2	Slc26a2.bSep08	117267	9577	426	3	62	solute carrier family 26 (sulfate transporter), member 2 (Slc26a2) alternative variant bSep08, mRNA.
Slc26a3	Slc26a3.bSep08	114629	11956	327	1	74	solute carrier family 26, member 3 (Slc26a3) alternative variant bSep08, mRNA.



Slc26a4	Slc26a4.bSep08	29440	7641	789	1	212	solute carrier family 26, member 4 (Slc26a4) alternative variant bSep08, mRNA.
Slc26a8	Slc26a8.bSep08	309646	12785	1959	9	515	solute carrier family 26, member 8 (Slc26a8) alternative variant bSep08, mRNA.
Slc26a8	Slc26a8.cSep08	309646	46044	1782	1	456	solute carrier family 26, member 8 (Slc26a8) alternative variant cSep08, mRNA.
Slc26a11	Slc26a11.aSep08	360670	4739	367		122	solute carrier family 26, member 11 (Slc26a11) mRNA.
Slc27a1	Slc27a1.bSep08	94172	3646	1179	6	206	solute carrier family 27 (fatty acid transporter), member 1 (22.9 kD) (Slc27a1) alternative variant bSep08, mRNA.
Slc27a1	Slc27a1.cSep08	94172	9040	594	3	148	solute carrier family 27 (fatty acid transporter), member 1 (Slc27a1) alternative variant cSep08, mRNA.
Slc27a1	Slc27a1.dSep08	94172	7872	296	2	98	solute carrier family 27 (fatty acid transporter), member 1 (Slc27a1) alternative variant dSep08, mRNA.
Slc27a1	Slc27a1.eSep08	94172	893	401	2	75	solute carrier family 27 (fatty acid transporter), member 1 (Slc27a1) alternative variant eSep08, mRNA.
Slc27a2	Slc27a2.bSep08	65192	11130	2577	2	475	solute carrier family 27 (fatty acid transporter), member 2 (Slc27a2) alternative variant bSep08, mRNA.
Slc27a2	Slc27a2.cSep08	65192	17839	860	5	286	solute carrier family 27 (fatty acid transporter), member 2 (Slc27a2) alternative variant cSep08, mRNA.
Slc27a4	Slc27a4.aSep08	311839	12908	3053	13	643	solute carrier family 27 (fatty acid transporter), member 4 (72.2 kD) (Slc27a4) alternative variant aSep08, mRNA.
Slc28a1	Slc28a1.bSep08	116642	1364	990	1	69	solute carrier family 28 (sodium-coupled nucleoside transporter), member 1 (Slc28a1) alternative variant bSep08, mRNA.
Slc28a2	Slc28a2.bSep08	60423	12997	843	8	258	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2 (Slc28a2) alternative variant bSep08, mRNA.
Slc28a2	Slc28a2.cSep08	60423	8280	559	4	90	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2 (Slc28a2) alternative variant cSep08, mRNA.
Slc28a2	Slc28a2.eSep08	60423	728	174	2	19	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2 (Slc28a2) alternative variant eSep08, mRNA.
Slc29a1	Slc29a1.bSep08	63997	5995	798	7	158	solute carrier family 29 member 1 CRA a (Slc29a1) alternative variant bSep08, mRNA.
Slc29a1	Slc29a1.cSep08	63997	2813	954	3	151	solute carrier family 29 member 1 CRA a (16.5 kD) (Slc29a1) alternative variant cSep08, mRNA.
Slc29a1	Slc29a1.dSep08	63997	3982	452	5	150	solute carrier family 29 member 1 CRA c (Slc29a1) alternative variant dSep08, mRNA.
Slc29a1	Slc29a1.eSep08	63997	2900	741	2	114	equilibrative nucleoside transporter 1 (12.5 kD) (Slc29a1) alternative variant eSep08, mRNA.
Slc29a1	Slc29a1.gSep08	63997	3595	612	4	89	putative protein (9.6 kD) (Slc29a1) alternative variant gSep08, mRNA.
Slc29a1	Slc29a1.hSep08	63997	1186	887	2	81	solute carrier family 29 member 1 (9.0 kD) (Slc29a1) alternative variant hSep08, mRNA.
Slc29a1	Slc29a1.iSep08	63997	5606	750	7	75	equilibrative nucleoside transporter 1 (Slc29a1) alternative variant iSep08, mRNA.

Slc29a1	Slc29a1.jSep08	63997	5620	703	6	79	equilibrative nucleoside transporter 1 (Slc29a1) alternative variant jSep08, mRNA.
Slc29a1	Slc29a1.kSep08	63997	1038	687	2	87	putative protein (Slc29a1) alternative variant kSep08, mRNA.
Slc29a1	Slc29a1.lSep08	63997	5949	540	6	70	equilibrative nucleoside transporter 1 (Slc29a1) alternative variant lSep08, mRNA.
Slc30a2	Slc30a2.cSep08	25362	1018	733	2	139	solute carrier family 30 (zinc transporter), member 2 (Slc30a2) alternative variant cSep08, mRNA.
Slc30a2	Slc30a2.dSep08	25362	4680	672	3	106	solute carrier family 30 (zinc transporter), member 2 (Slc30a2) alternative variant dSep08, mRNA.
Slc30a2	Slc30a2.eSep08	25362	10562	1777	3	35	solute carrier family 30 (zinc transporter), member 2 (Slc30a2) alternative variant eSep08, mRNA.
Slc30a3	Slc30a3.bSep08	366568	2113	837	5	206	solute carrier family 30 (zinc transporter), member 3 (Slc30a3) alternative variant bSep08, mRNA.
Slc30a3	Slc30a3.cSep08	366568	19120	793	6	178	solute carrier family 30 (zinc transporter), member 3 (Slc30a3) alternative variant cSep08, mRNA.
Slc30a6	Slc30a6.aSep08	298786	28947	1584	2	475	solute carrier family 30 (zinc transporter), member 6 (Slc30a6) alternative variant aSep08, mRNA.
Slc30a7	Slc30a7.aSep08	310801	74607	2008		378	solute carrier family 30 (zinc transporter), member 7 (41.8 kD) (Slc30a7) mRNA.
Slc31a1	Slc31a1.bSep08	171135	26029	825	5	170	solute carrier family 31 (copper transporters), member 1 (19.2 kD) (Slc31a1) alternative variant bSep08, mRNA.
Slc31a2	Slc31a2.aSep08	298091	10510	1767	3	143	solute carrier family 31, member 2 (16.1 kD) (Slc31a2) alternative variant aSep08, mRNA.
Slc31a2	Slc31a2.bSep08	298091	10335	1335	1	143	solute carrier family 31, member 2 (Slc31a2) alternative variant bSep08, mRNA.
Slc33a1	Slc33a1.bSep08	64018	22091	1882	6	243	solute carrier family 33 (acetyl-CoA transporter), member 1 (26.9 kD) (Slc33a1) alternative variant bSep08, complete mRNA.
Slc33a1	Slc33a1.cSep08	64018	6556	561	4	186	solute carrier family 33 (acetyl-CoA transporter), member 1 (Slc33a1) alternative variant cSep08, mRNA.
Slc33a1	Slc33a1.dSep08	64018	2076	931	2	122	solute carrier family 33 (acetyl-CoA transporter), member 1 (13.4 kD) (Slc33a1) alternative variant dSep08, mRNA.
Slc34a1	Slc34a1.bSep08	25548	6198	2206	4	327	solute carrier family 34 member 1 CRA b (Slc34a1) alternative variant bSep08, mRNA.
Slc34a1	Slc34a1.cSep08	25548	2692	775	5	215	solute carrier family 34 member 1 CRA b (22.7 kD) (Slc34a1) alternative variant cSep08, mRNA.
Slc34a1	Slc34a1.dSep08	25548	9693	765	7	126	solute carrier family 34 member 1 (Slc34a1) alternative variant dSep08, mRNA.
Slc34a1	Slc34a1.eSep08	25548	897	791	2	122	solute carrier family 34 member 1 CRA d (Slc34a1) alternative variant eSep08, mRNA.
Slc35a2	Slc35a2.cSep08	100158233	1867	563	2	129	solute carrier family 35, member A2 (Slc35a2) alternative variant cSep08, mRNA.
Slc35a2	Slc35a2.dSep08	100158233	1780	478	2	103	solute carrier family 35, member A2 (Slc35a2) alternative variant dSep08, mRNA.
Slc35a4	Slc35a4.bSep08	257647	2355	781	2	103	putative cytoplasmic protein (11.3 kD) (Slc35a4) alternative variant bSep08, mRNA.

Slc35a4	Slc35a4.cSep08	257647	2160	487	1	70	putative protein (7.5 kD) (Slc35a4) alternative variant cSep08, mRNA.
Slc35a4	Slc35a4.dSep08	257647	1884	396	2	51	putative protein (Slc35a4) alternative variant dSep08, mRNA.
Slc35a5	Slc35a5.aSep08	498081	3091	1616	2	90	solute carrier family 35, member A5 (Slc35a5) alternative variant aSep08, mRNA.
Slc35b1	Slc35b1.bSep08	287642	4361	728	6	190	solute carrier family 35 member B1 (Slc35b1) alternative variant bSep08, mRNA.
Slc35b1	Slc35b1.cSep08	287642	2343	1670	2	163	solute carrier family 35 member B1 CRA b (18.3 kD) (Slc35b1) alternative variant cSep08, mRNA.
Slc35b1	Slc35b1.dSep08	287642	1456	268	3	88	solute carrier family 35 member B1 CRA c (Slc35b1) alternative variant dSep08, mRNA.
Slc35b1	Slc35b1.eSep08	287642	1928	905	3	64	solute carrier family 35 member B1 CRA c (7.0 kD) (Slc35b1) alternative variant eSep08, mRNA.
Slc35b3	Slc35b3.aSep08	306866	26763	1889	8	429	solute carrier family 35, member B3 (Slc35b3) alternative variant aSep08, mRNA.
Slc35b3	Slc35b3.bSep08	306866	6161	452	1	90	solute carrier family 35, member B3 (Slc35b3) alternative variant bSep08, mRNA.
Slc35b3	Slc35b3.dSep08	306866	2060	741	2	57	solute carrier family 35, member B3 (Slc35b3) alternative variant dSep08, mRNA.
Slc35b3	Slc35b3.eSep08	306866	10705	727	1	53	solute carrier family 35, member B3 (Slc35b3) alternative variant eSep08, mRNA.
Slc35b4	Slc35b4.bSep08	296969	5332	734	4	122	solute carrier family 35, member B4 (Slc35b4) alternative variant bSep08, mRNA.
Slc35b4	Slc35b4.cSep08	296969	1852	761	2	95	solute carrier family 35, member B4 (Slc35b4) alternative variant cSep08, mRNA.
Slc35b4	Slc35b4.dSep08	296969	1428	682	2	55	solute carrier family 35, member B4 (Slc35b4) alternative variant dSep08, mRNA.
Slc35c1	Slc35c1.bSep08	311204	1804	633	2	210	solute carrier family 35, member C1 (Slc35c1) alternative variant bSep08, mRNA.
Slc35c2	Slc35c2.bSep08	311637	11065	2047	11	364	solute carrier family 35 member C2 (40.3 kD) (Slc35c2) alternative variant bSep08, mRNA.
Slc35c2	Slc35c2.cSep08	311637	8919	1381	10	335	solute carrier family 35 member C2 (Slc35c2) alternative variant cSep08, mRNA.
Slc35c2	Slc35c2.dSep08	311637	6251	1875	7	200	putative protein (44.9 kD) (Slc35c2) alternative variant dSep08, mRNA.
Slc35c2	Slc35c2.eSep08	311637	10130	1202	9	197	solute carrier family 35 member C2 (Slc35c2) alternative variant eSep08, mRNA.
Slc35c2	Slc35c2.fSep08	311637	4774	726	6	130	solute carrier family 35 member C2 (Slc35c2) alternative variant fSep08, mRNA.
Slc35c2	Slc35c2.gSep08	311637	1420	717	2	119	solute carrier family 35 member C2 (Slc35c2) alternative variant gSep08, mRNA.
Slc35c2	Slc35c2.hSep08	311637	5499	816	5	112	solute carrier family 35 member C2 (12.8 kD) (Slc35c2) alternative variant hSep08, mRNA.
Slc35c2	Slc35c2.iSep08	311637	5593	663	6	63	putative protein (6.8 kD) (Slc35c2) alternative variant iSep08, mRNA.

Slc35d1	Slc35d1.bSep08	298280	50621	910	10	279	solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1 (Slc35d1) alternative variant bSep08, mRNA.
Slc35d1	Slc35d1.cSep08	298280	1294	339	1	62	solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1 (Slc35d1) alternative variant cSep08, mRNA.
Slc35d2	Slc35d2.aSep08	290959	10187	947		127	solute carrier family 35, member D2 (13.9 kD) (Slc35d2) mRNA.
Slc35e3	Slc35e3.aSep08	362883	13676	3432		313	solute carrier family 35, member E3 (34.8 kD) (Slc35e3) mRNA.
Slc35f2	Slc35f2.aSep08	300713	36528	747	5	249	solute carrier family 35, member F2 (Slc35f2) alternative variant aSep08, mRNA.
Slc35f4	Slc35f4.aSep08	305865	6765	1624		522	solute carrier family 35, member F4 (Slc35f4) alternative variant aSep08, mRNA.
Slc35f4	Slc35f4.bSep08	305865	6145	1004		154	solute carrier family 35, member F4 (Slc35f4) alternative variant bSep08, mRNA.
Slc35f5	Slc35f5.bSep08	288993	17529	741	8	178	solute carrier family 35, member F5 (20.0 kD) (Slc35f5) alternative variant bSep08, mRNA.
Slc36a1	Slc36a1.aSep08	155205	12499	1759	5	525	solute carrier family 36 (proton/amino acid symporter), member 1 (Slc36a1) alternative variant aSep08, mRNA.
Slc36a3	Slc36a3.bSep08	303148	11752	708	1	121	solute carrier family 36 (proton/amino acid symporter), member 3 (Slc36a3) alternative variant bSep08, mRNA.
Slc37a1	Slc37a1.bSep08	294321	14055	781	5	169	solute carrier family 37 (glycerol-3-phosphate transporter), member 1 (Slc37a1) alternative variant bSep08, mRNA.
Slc37a1	Slc37a1.cSep08	294321	11173	763	6	129	solute carrier family 37 (glycerol-3-phosphate transporter), member 1 (Slc37a1) alternative variant cSep08, mRNA.
Slc37a2	Slc37a2.aSep08	500973	7074	2004	10	243	solute carrier family 37 (glycerol-3-phosphate transporter), member 2 (Slc37a2) alternative variant aSep08, mRNA.
Slc37a3	Slc37a3.bSep08	312255	19995	393	4	130	solute carrier family 37 (glycerol-3-phosphate transporter), member 3 (Slc37a3) alternative variant bSep08, mRNA.
Slc37a3	Slc37a3.dSep08	312255	2301	759	2	40	solute carrier family 37 (glycerol-3-phosphate transporter), member 3 (4.8 kD) (Slc37a3) alternative variant dSep08, mRNA.
Slc37a4	Slc37a4.bSep08	29573	5452	1924	8	266	solute carrier family 37 member 4 (29.1 kD) (Slc37a4) alternative variant bSep08, mRNA.
Slc37a4	Slc37a4.cSep08	29573	1155	1056	2	58	solute carrier family 37 member 4 CRA b (6.4 kD) (Slc37a4) alternative variant cSep08, mRNA.
Slc38a1	Slc38a1.bSep08	170567	54417	1799	7	439	solute carrier family 38 member 1 CRA b (Slc38a1) alternative variant bSep08, mRNA.
Slc38a1	Slc38a1.cSep08	170567	6752	700	7	167	solute carrier family 38 member 1 CRA a (Slc38a1) alternative variant cSep08, mRNA.
Slc38a1	Slc38a1.eSep08	170567	26061	606	4	72	solute carrier family 38 member 1 CRA a (8.4 kD) (Slc38a1) alternative variant eSep08, mRNA.
Slc38a1	Slc38a1.fSep08	170567	1744	417	3	50	solute carrier family 38 member 1 CRA c (Slc38a1) alternative variant fSep08, mRNA.
Slc38a2	Slc38a2.bSep08	29642	2543	707	5	116	solute carrier family 38, member 2 (Slc38a2) alternative variant bSep08, mRNA.

Slc38a2	Slc38a2.cSep08	29642	968	431	3	104	solute carrier family 38, member 2 (Slc38a2) alternative variant cSep08, mRNA.
Slc38a3	Slc38a3.bSep08	252919	3316	964	5	187	solute carrier family 38 member 3 CRA a (20.9 kD) (Slc38a3) alternative variant bSep08, mRNA.
Slc38a3	Slc38a3.cSep08	252919	3099	836	5	178	solute carrier family 38 member 3 CRA a (Slc38a3) alternative variant cSep08, mRNA.
Slc38a3	Slc38a3.dSep08	252919	3293	1143	4	110	solute carrier family 38 member 3 CRA a (12.2 kD) (Slc38a3) alternative variant dSep08, mRNA.
Slc38a4	Slc38a4.bSep08	170573	4052	741	6	246	solute carrier family 38, member 4 (Slc38a4) alternative variant bSep08, mRNA.
Slc38a4	Slc38a4.cSep08	170573	29598	418	5	123	solute carrier family 38, member 4 (Slc38a4) alternative variant cSep08, mRNA.
Slc38a4	Slc38a4.dSep08	170573	24527	433	5	114	solute carrier family 38, member 4 (Slc38a4) alternative variant dSep08, mRNA.
Slc38a4	Slc38a4.eSep08	170573	39845	415	4	104	solute carrier family 38, member 4 (Slc38a4) alternative variant eSep08, mRNA.
Slc38a4	Slc38a4.fSep08	170573	30472	531	2	30	solute carrier family 38, member 4 (3.5 kD) (Slc38a4) alternative variant fSep08, mRNA.
Slc38a4	Slc38a4.gSep08	170573	39203	391	5	53	solute carrier family 38, member 4 (Slc38a4) alternative variant gSep08, mRNA.
Slc38a5	Slc38a5.bSep08	192208	4124	754	1	251	solute carrier family 38, member 5 (Slc38a5) alternative variant bSep08, mRNA.
Slc38a6	Slc38a6.bSep08	299139	8714	1781	5	159	solute carrier family 38, member 6 (Slc38a6) alternative variant bSep08, mRNA.
Slc38a6	Slc38a6.cSep08	299139	45921	586	9	119	solute carrier family 38, member 6 (12.8 kD) (Slc38a6) alternative variant cSep08, mRNA.
Slc38a6	Slc38a6.fSep08	299139	1901	413	2	33	solute carrier family 38, member 6 (Slc38a6) alternative variant fSep08, mRNA.
Slc38a7	Slc38a7.bSep08	291840	4780	420	1	59	solute carrier family 38, member 7 (Slc38a7) alternative variant bSep08, mRNA.
Slc38a9	Slc38a9.bSep08	310091	7413	954	4	111	solute carrier family 38, member 9 (12.7 kD) (Slc38a9) alternative variant bSep08, mRNA.
Slc38a10	Slc38a10.aSep08	303740	7618	3040	4	645	solute carrier family 38, member 10 (Slc38a10) alternative variant aSep08, mRNA.
Slc38a10	Slc38a10.bSep08	303740	5631	1855	2	428	solute carrier family 38, member 10 (Slc38a10) alternative variant bSep08, mRNA.
Slc38a11	Slc38a11.aSep08	362141	16219	787	6	237	solute carrier family 38, member 11 (Slc38a11) alternative variant aSep08, mRNA.
Slc38a11	Slc38a11.bSep08	362141	2685	338	1	65	solute carrier family 38, member 11 (Slc38a11) alternative variant bSep08, mRNA.
Slc39a3	Slc39a3.aSep08	314637	5932	1538	1	317	solute carrier family 39 (zinc transporter), member 3 (34.0 kD) (Slc39a3) alternative variant aSep08, mRNA.
Slc39a4	Slc39a4.bSep08	300051	934	846	2	100	solute carrier family 39 (zinc transporter), member 4 (Slc39a4) alternative variant bSep08, mRNA.
Slc39a5	Slc39a5.bSep08	362812	4682	1052	5	299	solute carrier family 39 (metal ion transporter), member 5 (Slc39a5) alternative variant bSep08, mRNA.

Slc39a6	Slc39a6.bSep08	291733	11548	1776	4	585	solute carrier family 39 (metal ion transporter), member 6 (Slc39a6) alternative variant bSep08, mRNA.
Slc39a6	Slc39a6.cSep08	291733	4254	428	3	65	solute carrier family 39 (metal ion transporter), member 6 (Slc39a6) alternative variant cSep08, mRNA.
Slc39a9	Slc39a9.bSep08	314275	12805	674	5	224	solute carrier family 39 (zinc transporter), member 9 (Slc39a9) alternative variant bSep08, mRNA.
Slc39a10_predicted	Slc39a10_predicted.bSep08	363229	6919	603	3	191	solute carrier family 39 (zinc transporter), member 10 (predicted) (Slc39a10_predicted) alternative variant bSep08, mRNA.
Slc39a11	Slc39a11.bSep08	287796	266557	702	5	210	solute carrier family 39 (metal ion transporter), member 11 (Slc39a11) alternative variant bSep08, mRNA.
Slc39a11	Slc39a11.cSep08	287796	208955	864	6	207	solute carrier family 39 (metal ion transporter), member 11 (21.9 kD) (Slc39a11) alternative variant cSep08, mRNA.
Slc39a11	Slc39a11.dSep08	287796	129866	835	3	82	solute carrier family 39 (metal ion transporter), member 11 (Slc39a11) alternative variant dSep08, mRNA.
Slc39a11	Slc39a11.eSep08	287796	5827	342	2	43	solute carrier family 39 (metal ion transporter), member 11 (Slc39a11) alternative variant eSep08, mRNA.
Slc39a13	Slc39a13.bSep08	295928	2056	971	5	144	solute carrier family 39 (metal ion transporter), member 13 (Slc39a13) alternative variant bSep08, mRNA.
Slc39a13	Slc39a13.cSep08	295928	1318	858	4	113	solute carrier family 39 (metal ion transporter), member 13 (12.3 kD) (Slc39a13) alternative variant cSep08, mRNA.
Slc39a14	Slc39a14.bSep08	306009	6073	410	4	136	solute carrier family 39 (zinc transporter), member 14 (Slc39a14) alternative variant bSep08, mRNA.
Slc39a14	Slc39a14.cSep08	306009	6584	391	3	129	solute carrier family 39 (zinc transporter), member 14 (Slc39a14) alternative variant cSep08, mRNA.
Slc41a2	Slc41a2.bSep08	362861	46286	751	4	175	solute carrier family 41, member 2 (Slc41a2) alternative variant bSep08, mRNA.
Slc41a2	Slc41a2.cSep08	362861	19710	385	3	107	solute carrier family 41, member 2 (Slc41a2) alternative variant cSep08, mRNA.
Slc41a2	Slc41a2.eSep08	362861	14194	683	3	69	solute carrier family 41, member 2 (Slc41a2) alternative variant eSep08, mRNA.
Slc43a1	Slc43a1.bSep08	311168	33498	933	8	264	solute carrier family 43, member 1 (Slc43a1) alternative variant bSep08, mRNA.
Slc43a1	Slc43a1.cSep08	311168	4430	1041	4	156	solute carrier family 43, member 1 (17.5 kD) (Slc43a1) alternative variant cSep08, mRNA.
Slc43a2	Slc43a2.bSep08	287532	4765	382	3	127	solute carrier family 43 member 2 CRA c (Slc43a2) alternative variant bSep08, mRNA.
Slc43a2	Slc43a2.cSep08	287532	13618	1103	4	94	solute carrier family 43 member 2 CRA b (10.8 kD) (Slc43a2) alternative variant cSep08, mRNA.
Slc43a2	Slc43a2.dSep08	287532	10264	535	3	92	putative protein (10.1 kD) (Slc43a2) alternative variant dSep08, mRNA.
Slc43a3	Slc43a3.bSep08	311170	14224	831	7	277	solute carrier family 43, member 3 (Slc43a3) alternative variant bSep08, mRNA.
Slc43a3	Slc43a3.cSep08	311170	1562	394	3	131	solute carrier family 43, member 3 (Slc43a3) alternative variant cSep08, mRNA.
Slc44a2	Slc44a2.bSep08	363024	6708	2083	7	347	solute carrier family 44 member 2 (Slc44a2) alternative variant bSep08, mRNA.

Slc44a2	Slc44a2.cSep08	363024	6724	1403	8	270	solute carrier family 44 member 2 CRA b (Slc44a2) alternative variant cSep08, mRNA.
Slc44a2	Slc44a2.dSep08	363024	24935	1362	9	252	solute carrier family 44 member 2 (Slc44a2) alternative variant dSep08, mRNA.
Slc44a2	Slc44a2.eSep08	363024	4432	734	5	229	solute carrier family 44 member 2 (Slc44a2) alternative variant eSep08, mRNA.
Slc44a2	Slc44a2.fSep08	363024	24487	875	8	199	solute carrier family 44 member 2 (Slc44a2) alternative variant fSep08, mRNA.
Slc44a2	Slc44a2.gSep08	363024	6153	990	9	117	solute carrier family 44 member 2 (Slc44a2) alternative variant gSep08, mRNA.
Slc44a3	Slc44a3.bSep08	295417	22389	799	7	266	solute carrier family 44, member 3 (Slc44a3) alternative variant bSep08, mRNA.
Slc44a3	Slc44a3.cSep08	295417	1467	467	2	44	solute carrier family 44, member 3 (Slc44a3) alternative variant cSep08, mRNA.
Slc44a3	Slc44a3.dSep08	295417	30396	726	4	6	solute carrier family 44, member 3 (0.7 kD) (Slc44a3) alternative variant dSep08, mRNA.
Slc44a4	Slc44a4.aSep08	294255	16161	2297		709	solute carrier family 44, member 4 (Slc44a4) mRNA.
Slc45a1	Slc45a1.cSep08	246258	1002	461	2	82	solute carrier family 45, member 1 (Slc45a1) alternative variant cSep08, mRNA.
Slc45a3	Slc45a3.aSep08	304785	20551	3163	1	564	solute carrier family 45, member 3 (60.7 kD) (Slc45a3) alternative variant aSep08, complete mRNA.
Slc45a3	Slc45a3.bSep08	304785	15533	779	1	158	solute carrier family 45, member 3 (Slc45a3) alternative variant bSep08, mRNA.
Slc45a4	Slc45a4.aSep08	315054	59820	1323		204	solute carrier family 45, member 4 (Slc45a4) mRNA.
Slc46a3	Slc46a3.bSep08	288454	5063	223	1	74	solute carrier family 46, member 3 (Slc46a3) alternative variant bSep08, mRNA.
Slc47a2	Slc47a2.aSep08	497921	17920	740		134	solute carrier family 47, member 2 (Slc47a2) mRNA.
Slco1a4	Slco1a4.cSep08	170698	5475	842	2	56	solute carrier organic anion transporter family, member 1a4 (Slco1a4) alternative variant cSep08, mRNA.
Slco1a5andSlco1a1	Slco1a5andSlco1a1.bSep08	50572	1805	650	1	49	solute carrier organic anion transporter family, member 1a1 and solute carrier organic anion transporter family, member 1a5 (Slco1a5andSlco1a1) alternative variant bSep08, mRNA.
Slco1a5andSlco1a1	Slco1a5andSlco1a1.bSep08	80900	1805	650	1	49	solute carrier organic anion transporter family, member 1a1 and solute carrier organic anion transporter family, member 1a5 (Slco1a5andSlco1a1) alternative variant bSep08, mRNA.
Slco1a6	Slco1a6.bSep08	84608	25325	848	2	159	solute carrier organic anion transporter family, member 1a6 (Slco1a6) alternative variant bSep08, mRNA.
Slco1c1	Slco1c1.bSep08	84511	5962	608	4	167	solute carrier organic anion transporter family, member 1c1 (Slco1c1) alternative variant bSep08, mRNA.
Slco2b1	Slco2b1.bSep08	140860	12553	857	2	175	solute carrier organic anion transporter family, member 2b1 (Slco2b1) alternative variant bSep08, mRNA.
Slco3a1	Slco3a1.bSep08	140915	53354	1657	7	358	solute carrier organic anion transporter family, member 3a1 (Slco3a1) alternative variant bSep08, mRNA.
Slco3a1	Slco3a1.cSep08	140915	55007	585	2	194	solute carrier organic anion transporter family, member 3a1 (Slco3a1) alternative variant cSep08, mRNA.

Slco4a1	Slco4a1.bSep08	171144	981	734	1	176	solute carrier organic anion transporter family, member 4a1 (Slco4a1) alternative variant bSep08, mRNA.
Slco6b1	Slco6b1.bSep08	170925	26766	913	2	144	solute carrier organic anion transporter family, member 6b1 (Slco6b1) alternative variant bSep08, mRNA.
sleebor	sleebor.aSep08		1159	642		126	type X alpha 1 (sleebor) mRNA.
sleechy	sleechy.aSep08		1322	439		60	putative protein (7.2 kD) (sleechy) mRNA.
sleedoy	sleedoy.bSep08		1261	1151		96	putative protein (sleedoy) alternative variant bSep08, mRNA.
sleefee	sleefee.aSep08		2131	284		56	putative protein (sleefee) mRNA.
sleeflu	sleeflu.aSep08		38032	952	2	103	CRA a (sleeflu) alternative variant aSep08, mRNA.
sleeflu	sleeflu.bSep08		11842	713	1	62	CRA a (sleeflu) alternative variant bSep08, mRNA.
sleefly	sleefly.aSep08		726	492		79	putative nuclear protein (7.9 kD) (sleefly) mRNA.
sleeja	sleeja.aSep08		41825	369		53	putative protein (6.0 kD) (sleeja) mRNA.
sleelo	sleelo.aSep08		22188	615		72	putative protein (sleelo) mRNA.
sleemee	sleemee.aSep08		420	315	2	54	putative protein (sleemee) alternative variant aSep08, mRNA.
sleepy	sleepy.aSep08		445	351		71	tyrosine kinase 2 (sleepy) mRNA.
sleeroy	sleeroy.aSep08		4620	306		56	putative protein (sleeroy) mRNA.
sleeshaw	sleeshaw.aSep08		1751	1611	2	39	putative protein (4.5 kD) (sleeshaw) alternative variant aSep08, mRNA.
sleeshee	sleeshee.aSep08		5238	344		99	tubulin polyglutamylase tll17 like (sleeshee) mRNA.
sleetu	sleetu.aSep08		6933	375		48	putative protein (sleetu) mRNA.
sleevo	sleevo.aSep08		6258	626		208	proteasome-associated protein Ecm29 homolog (sleevo) mRNA.
sleewer	sleewer.aSep08		9811	523		79	CRA a like (sleewer) mRNA.
slerbor	slerbor.aSep08		2117	311		35	putative protein (slerbor) mRNA.
slerchy	slerchy.aSep08		3203	295		35	putative protein (slerchy) mRNA.
slerdoy	slerdoy.aSep08		4630	427	3	141	CRA c (slerdoy) alternative variant aSep08, mRNA.
slerdoy	slerdoy.bSep08		1840	302	1	64	CRA c (slerdoy) alternative variant bSep08, mRNA.
slerfee	slerfee.aSep08		1293	519		26	putative protein (2.9 kD) (slerfee) mRNA.
slerflu	slerflu.aSep08		13783	279		45	putative protein (slerflu) mRNA.
slerfly	slerfly.aSep08		3895	577		166	upf0470 protein homolog (slerfly) mRNA.
slerja	slerja.aSep08		2624	441		82	putative protein (slerja) mRNA.
slerlo	slerlo.aSep08		14270	738	1	91	putative protein of mammalian origin (slerlo) alternative variant aSep08, mRNA.
slerlo	slerlo.bSep08		738	649	1	108	putative protein (slerlo) alternative variant bSep08, mRNA.
slermee	slermee.aSep08		4307	434		144	myosin ID (slermee) mRNA.
slerpey	slerpey.aSep08		6956	1327		437	tyrosine kinase 2 (slerpey) mRNA.
slerroy	slerroy.aSep08		35444	575		138	putative protein of metazoan origin (slerroy) mRNA.
slershaw	slershaw.aSep08		1147	247		64	putative protein (slershaw) mRNA.
slershee	slershee.aSep08		1125	293		76	putative protein (8.2 kD) (slershee) mRNA.
slertu	slertu.aSep08		4903	944	4	290	numb homolog (slertu) alternative variant aSep08, mRNA.
slertu	slertu.bSep08		5444	2272	2	246	numb (slertu) alternative variant bSep08, mRNA.



slervo	slervo.aSep08		7917	558		185	proteasome-associated protein ecm29 homolog (slervo) mRNA.
slerwer	slerwer.aSep08		13410	810		269	putative protein of vertebrate origin (slerwer) mRNA.
sleybor	sleybor.aSep08		1545	721		37	putative protein (sleybor) mRNA.
sleychy	sleychy.aSep08		1314	896		84	putative protein (sleychy) mRNA.
sleydoy	sleydoy.aSep08		3689	210		24	putative protein (sleydoy) mRNA.
sleyfee	sleyfee.aSep08		2290	271		49	putative protein (sleyfee) mRNA.
sleyflu	sleyflu.aSep08		3478	586	2	74	putative protein (sleyflu) alternative variant aSep08, mRNA.
sleyflu	sleyflu.cSep08		8762	1126	3	36	putative protein (4.1 kD) (sleyflu) alternative variant cSep08, mRNA.
sleyfly	sleyfly.aSep08		2745	851		272	CRA a (sleyfly) mRNA.
sleyja	sleyja.aSep08		1724	217		32	putative protein (3.8 kD) (sleyja) mRNA.
sleylo	sleylo.aSep08		915	406		99	putative protein (sleylo) mRNA.
sleymee	sleymee.aSep08		87581	773	4	206	myosin ID CRA b (sleymee) alternative variant aSep08, mRNA.
sleymee	sleymee.bSep08		1123	610	1	124	putative protein of vertebrate origin (14.2 kD) (sleymee) alternative variant bSep08, mRNA.
sleypey	sleypey.aSep08		5027	285		85	putative protein (sleypey) mRNA.
sleyroy	sleyroy.aSep08		1801	649		53	putative protein (sleyroy) mRNA.
sleyshaw	sleyshaw.aSep08		6835	636		39	putative protein (4.4 kD) (sleyshaw) mRNA.
sleyshee	sleyshee.aSep08		1834	567		84	putative protein (sleyshee) mRNA.
sleytu	sleytu.aSep08		9188	631		51	putative protein (5.7 kD) (sleytu) mRNA.
sleyvo	sleyvo.aSep08		45028	829		275	novel protein containing 10 HEAT domains (sleyvo) alternative variant aSep08, mRNA.
sleyvo	sleyvo.bSep08		33206	287	1	58	novel protein containing 10 HEAT domains (sleyvo) alternative variant bSep08, mRNA.
sleywer	sleywer.bSep08		4357	744	3	30	putative protein (sleywer) alternative variant bSep08, mRNA.
sleywer	sleywer.cSep08		1258	428	3	49	CRA b like (sleywer) alternative variant cSep08, mRNA.
Slfn2	Slfn2.bSep08	303380	5477	736	1	80	schlafen 2 (9.3 kD) (Slfn2) alternative variant bSep08, mRNA.
Slfn5	Slfn5.aSep08	303377	4184	587		195	schlafen 5 (Slfn5) mRNA.
Slfn8	Slfn8.bSep08	303378	13735	1511	2	410	schlafen 8 (46.2 kD) (Slfn8) alternative variant bSep08, mRNA.
Slfnl1	Slfnl1.bSep08	500540	1329	721	2	208	schlafen-like 1 (Slfnl1) alternative variant bSep08, mRNA.
Slfnl1	Slfnl1.cSep08	500540	931	325	2	58	schlafen-like 1 (Slfnl1) alternative variant cSep08, mRNA.
Slit1	Slit1.bSep08	65047	6985	2061	5	283	slit homolog 1 (Drosophila) (Slit1) alternative variant bSep08, mRNA.
Slit2	Slit2.aSep08	360272	38925	1674	8	339	slit homolog 2 (Drosophila) (36.9 kD) (Slit2) alternative variant aSep08, mRNA.
Slit3	Slit3.bSep08	83467	25768	398	3	132	slit homolog 3 (Drosophila) (Slit3) alternative variant bSep08, mRNA.
Slitrk3	Slitrk3.aSep08	310519	4927	3318	1	979	SLIT and NTRK-like family, member 3 (109.2 kD) (Slitrk3) alternative variant aSep08, mRNA.

Slk	Slk.bSep08	54308	22480	4284	9	375	STE20-like kinase (yeast) (Slk) alternative variant bSep08, mRNA.
Slk	Slk.dSep08	54308	1308	861	2	53	STE20-like kinase (yeast) (6.2 kD) (Slk) alternative variant dSep08, mRNA.
Slmap	Slmap.bSep08	290533	116679	5085	20	749	sarcolemma associated protein (86.0 kD) (Slmap) alternative variant bSep08, mRNA.
Slmo1	Slmo1.bSep08	690253	11589	705	6	168	slowmo homolog 1 (Drosophila) (Slmo1) alternative variant bSep08, mRNA.
slobor	slobor.aSep08		28471	591	2	43	putative protein (slobor) alternative variant aSep08, mRNA.
slobor	slobor.bSep08		27991	578	5	73	putative cytoplasmic protein (8.2 kD) (slobor) alternative variant bSep08, mRNA.
slochy	slochy.aSep08		49867	556		49	putative protein (5.7 kD) (slochy) mRNA.
slodoy	slodoy.aSep08		682	487		113	putative protein (slodoy) mRNA.
slofee	slofee.aSep08		562	404		102	putative protein (slofee) mRNA.
sloflu	sloflu.aSep08		26364	493		49	putative protein (5.3 kD) (sloflu) mRNA.
slofly	slofly.aSep08		931	547		26	putative protein (2.9 kD) (slofly) mRNA.
sloja	sloja.aSep08		69608	314		60	putative protein (sloja) mRNA.
slojey	slojey.aSep08		15378	497		60	putative protein (slojey) mRNA.
slolo	slolo.aSep08		32435	1420	4	54	CRA a like (6.3 kD) (slolo) alternative variant aSep08, mRNA.
slolo	slolo.cSep08		27377	910	4	79	CRA c like (slolo) alternative variant cSep08, mRNA.
slolo	slolo.dSep08		25591	865	4	59	putative protein (6.8 kD) (slolo) alternative variant dSep08, mRNA.
slolo	slolo.eSep08		1828	487	2	57	putative protein (slolo) alternative variant eSep08, mRNA.
slomee	slomee.aSep08		2301	96		31	active -related (slomee) mRNA.
slopey	slopey.aSep08		2647	736		138	putative protein (slopey) mRNA.
slopor	slopor.aSep08		13833	481		55	putative protein (slopor) mRNA.
slorbor	slorbor.aSep08		7604	876	5	39	putative protein (4.6 kD) (slorbor) alternative variant aSep08, mRNA.
slorbor	slorbor.bSep08		680	442	1	39	putative protein (4.6 kD) (slorbor) alternative variant bSep08, mRNA.
slorchy	slorchy.aSep08		5783	593		55	putative protein (6.3 kD) (slorchy) mRNA.
slordoy	slordoy.aSep08		4320	444	3	147	solute carrier family 14 member 1 CRA a (slordoy) alternative variant aSep08, mRNA.
slorfee	slorfee.aSep08		1071	483		64	putative protein (slorfee) mRNA.
slorflu	slorflu.aSep08		1089	288		30	putative protein (slorflu) mRNA.
slorfly	slorfly.aSep08		3049	257	2	66	putative protein (slorfly) alternative variant aSep08, mRNA.
slorja	slorja.aSep08	306163	14082	4572		126	glypican 6 (slorja) mRNA.
slormee	slormee.aSep08		16305	1787		595	putative protein (slormee) mRNA.
sloroy	sloroy.aSep08		31843	545		181	putative protein of metazoan origin (sloroy) mRNA.
slorpey	slorpey.aSep08		12996	707		55	putative protein (slorpey) mRNA.
slorroy	slorroy.aSep08		308	217		15	putative protein (slorroy) mRNA.
slorshaw	slorshaw.aSep08		4984	736		43	putative protein (4.9 kD) (slorshaw) mRNA.

slorshee	slorshee.aSep08		6102	675		39	putative protein (slorshee) mRNA.
slortu	slortu.dSep08		5480	607	3	37	putative protein (slortu) alternative variant dSep08, mRNA.
slorvo	slorvo.aSep08		13741	1300		326	CRA a (slorvo) mRNA.
slorwer	slorwer.aSep08		8951	493		149	v-raf murine sarcoma viral oncogene homolog B1 (slorwer) mRNA.
sloshaw	sloshaw.aSep08		6817	427		141	putative protein of metazoan origin (sloshaw) mRNA.
sloshee	sloshee.aSep08		463	374		124	putative protein (sloshee) mRNA.
slotu	slotu.aSep08		9501	357	2	33	putative protein (3.5 kD) (slotu) alternative variant aSep08, mRNA.
slotu	slotu.bSep08		9569	826	2	74	putative protein (slotu) alternative variant bSep08, mRNA.
slovo	slovo.aSep08		1493	390		38	putative protein (4.5 kD) (slovo) mRNA.
slower	slower.aSep08		28949	891		69	putative protein (7.9 kD) (slower) mRNA.
sloybor	sloybor.aSep08		4460	391		129	laminin (sloybor) mRNA.
sloycha	sloycha.aSep08		586	485		77	putative protein of mammalian origin (sloycha) mRNA.
sloychy	sloychy.aSep08		9348	552		59	putative protein (sloychy) mRNA.
sloydoy	sloydoy.aSep08		1334	288		37	putative protein (sloydoy) mRNA.
sloyfee	sloyfee.aSep08		1221	724		51	putative protein (5.4 kD) (sloyfee) mRNA.
sloyflu	sloyflu.aSep08		2509	500		37	putative protein (sloyflu) mRNA.
sloyfly	sloyfly.aSep08		2686	759		123	putative protein of mammalian origin (sloyfly) mRNA.
sloyja	sloyja.aSep08		4554	496		164	ATP-binding cassette sub-family C member 4 like (sloyja) mRNA.
sloymee	sloymee.aSep08		1098	502		35	putative protein (4.0 kD) (sloymee) mRNA.
sloypey	sloypey.aSep08		7015	684		109	phosphodiesterase 4A (11.6 kD) (sloypey) mRNA.
sloyroy	sloyroy.aSep08		24304	255		47	putative protein (sloyroy) mRNA.
sloyshaw	sloyshaw.aSep08		907	638		43	putative protein (sloyshaw) mRNA.
sloyshee	sloyshee.aSep08		6572	792		42	putative protein (5.0 kD) (sloyshee) mRNA.
sloytu	sloytu.aSep08		1774	504		35	putative protein (sloytu) mRNA.
sloyvo	sloyvo.aSep08		8407	809		269	fk506 binding protein 15 133kDa like (sloyvo) mRNA.
sloywer	sloywer.aSep08		56723	354	1	118	v-raf murine sarcoma viral oncogene homolog B1 (sloywer) alternative variant aSep08, mRNA.
sloywer	sloywer.bSep08		56715	395	1	80	serine threonine kinase (sloywer) alternative variant bSep08, mRNA.
slubor	slubor.aSep08		4010	554		82	protein particle complex (slubor) mRNA.
sluchy	sluchy.aSep08		4090	317	2	57	putative protein (sluchy) alternative variant aSep08, mRNA.
sluchy	sluchy.bSep08		644	326	1	54	putative protein (sluchy) alternative variant bSep08, mRNA.
sludoy	sludoy.aSep08		51986	408		91	putative protein (sludoy) mRNA.
slufee	slufee.aSep08		3816	794		27	putative protein (slufee) mRNA.
sluflu	sluflu.aSep08		1422	584	2	194	ki-67 (sluflu) alternative variant aSep08, mRNA.
sluflu	sluflu.aSep08		10138	453		37	putative protein (4.2 kD) (sluflu) mRNA.
sluja	sluja.aSep08		3778	383		90	putative protein (sluja) mRNA.
slujey	slujey.aSep08		11436	474		157	heparanase (slujey) mRNA.
slulo	slulo.aSep08		1582	557		66	putative protein (7.3 kD) (slulo) mRNA.

slumee	slumee.aSep08		929	654		77	putative protein (8.8 kD) (slumee) mRNA.
slupey	slupey.aSep08		21274	647		39	putative protein (slupey) mRNA.
slupor	slupor.aSep08		2550	719		138	solute carrier family 6 member 20B (slupor) mRNA.
sluroy	sluroy.aSep08		40748	294		41	putative protein of mammalian origin (sluroy) mRNA.
slushaw	slushaw.aSep08		11760	689		229	3-hydroxy-3-methylglutaryl-Coenzyme A reductase (slushaw) mRNA.
slushee	slushee.aSep08		488	265		46	putative protein (slushee) mRNA.
slutu	slutu.aSep08		36786	864		134	putative protein (15.2 kD) (slutu) mRNA.
sluvo	sluvo.aSep08		6463	821	3	48	putative protein (5.5 kD) (sluvo) alternative variant aSep08, mRNA.
sluvo	sluvo.bSep08		6443	774	3	48	putative protein (5.5 kD) (sluvo) alternative variant bSep08, mRNA.
sluvo	sluvo.cSep08		1396	318	2	23	putative protein (2.5 kD) (sluvo) alternative variant cSep08, mRNA.
sluwer	sluwer.aSep08		21752	527		84	putative protein (sluwer) mRNA.
slybor	slybor.aSep08		2577	407		135	RAN binding protein 2 like (slybor) mRNA.
slychy	slychy.aSep08		2948	373		86	putative protein (slychy) mRNA.
slydoy	slydoy.aSep08		10851	381	3	126	putative protein of metazoan origin (slydoy) alternative variant aSep08, mRNA.
slyfee	slyfee.aSep08		3282	431		64	F-box leucine-rich repeat protein 21 (slyfee) mRNA.
slyflu	slyflu.aSep08		3987	751		109	putative protein (11.9 kD) (slyflu) mRNA.
slyfly	slyfly.aSep08		1475	408		135	zinc finger protein 580 (slyfly) mRNA.
slyja	slyja.aSep08		13313	698		36	putative protein (slyja) mRNA.
slyjey	slyjey.aSep08		547	393		44	putative protein (slyjey) mRNA.
slylo	slylo.aSep08		6100	396		39	putative protein (4.6 kD) (slylo) mRNA.
slymee	slymee.aSep08		13208	374		50	putative protein (5.4 kD) (slymee) mRNA.
slypey	slypey.aSep08		9080	418		117	putative protein (slypey) mRNA.
slypor	slypor.aSep08		8312	417		65	putative protein (7.2 kD) (slypor) mRNA.
slyroy	slyroy.aSep08		1548	415	2	114	CRA b like (slyroy) alternative variant aSep08, mRNA.
slyroy	slyroy.bSep08		1839	386	2	37	CRA b like (slyroy) alternative variant bSep08, mRNA.
slyshaw	slyshaw.aSep08		2673	350		70	putative protein (slyshaw) mRNA.
slyshee	slyshee.aSep08		650	558		41	putative protein (4.8 kD) (slyshee) mRNA.
slytu	slytu.aSep08		1436	379		76	putative nuclear protein (8.2 kD) (slytu) mRNA.
slyvo	slyvo.aSep08		902	797		57	putative protein (slyvo) mRNA.
slywer	slywer.aSep08		1392	552		93	putative mitochondrial protein (10.0 kD) (slywer) mRNA.
smabor	smabor.aSep08		2866	450		65	putative protein (7.2 kD) (smabor) mRNA.
smacha	smacha.aSep08		2971	371		85	GTPase activating protein testicular GAP1 like (smacha) mRNA.
smachy	smachy.bSep08		2129	631	4	101	CRA a (10.9 kD) (smachy) alternative variant bSep08, mRNA.
Smad1	Smad1.bSep08	25671	5294	501	2	166	MAD homolog 1 (Drosophila) (Smad1) alternative variant bSep08, mRNA.

Smad2	Smad2.bSep08	29357	61943	2597	11	467	MAD homolog 2 (Drosophila) (52.2 kD) (Smad2) alternative variant bSep08, mRNA.
Smad2	Smad2.cSep08	29357	47632	755	5	212	MAD homolog 2 (Drosophila) (Smad2) alternative variant cSep08, mRNA.
Smad2	Smad2.dSep08	29357	2574	765	2	67	MAD homolog 2 (Drosophila) (Smad2) alternative variant dSep08, mRNA.
Smad3	Smad3.bSep08	25631	2385	1511	2	121	MAD homolog 3 (Drosophila) (13.5 kD) (Smad3) alternative variant bSep08, mRNA.
Smad4	Smad4.bSep08	50554	43836	1779	2	298	MAD homolog 4 (Drosophila) (Smad4) alternative variant bSep08, mRNA.
Smad4	Smad4.cSep08	50554	12892	703	1	197	MAD homolog 4 (Drosophila) (Smad4) alternative variant cSep08, mRNA.
Smad4	Smad4.dSep08	50554	4727	278	2	92	MAD homolog 4 (Drosophila) (Smad4) alternative variant dSep08, mRNA.
Smad5	Smad5.bSep08	59328	5611	941	1	313	MAD homolog 5 (Drosophila) (Smad5) alternative variant bSep08, mRNA.
Smad6	Smad6.bSep08	367100	1531	1448	2	208	MAD homolog 6 (Drosophila) (Smad6) alternative variant bSep08, mRNA.
Smad7	Smad7.bSep08	81516	24461	790	4	245	MAD homolog 7 (Drosophila) (Smad7) alternative variant bSep08, mRNA.
Smad7	Smad7.cSep08	81516	25208	2298	3	235	MAD homolog 7 (Drosophila) (Smad7) alternative variant cSep08, mRNA.
smadoy	smadoy.aSep08		8737	273		43	putative protein (smadoy) mRNA.
smafee	smafee.aSep08		12941	706		33	putative protein (smafee) mRNA.
smaflu	smaflu.aSep08		7017	581		118	putative protein of vertebrate origin (smaflu) mRNA.
smafly	smafly.aSep08		2375	736		123	putative protein of mammalian origin (smafly) mRNA.
Smagp	Smagp.bSep08	300236	35238	736	1	39	small cell adhesion glycoprotein and hypothetical protein LOC685015 (4.1 kD) (Smagp) alternative variant bSep08, mRNA.
Smagp	Smagp.bSep08	685015	35238	736	1	39	small cell adhesion glycoprotein and hypothetical protein LOC685015 (4.1 kD) (Smagp) alternative variant bSep08, mRNA.
smaja	smaja.aSep08		1283	738		55	putative protein (6.1 kD) (smaja) mRNA.
smamee	smamee.aSep08		4474	486		161	schlafen 5 CRA b (smamee) mRNA.
Smap2	Smap2.bSep08	298500	9160	1523	6	356	stromal membrane-associated GTPase-activating protein 2 and hypothetical protein LOC680690 (Smap2) alternative variant bSep08, mRNA.
Smap2	Smap2.bSep08	680690	9160	1523	6	356	stromal membrane-associated GTPase-activating protein 2 and hypothetical protein LOC680690 (Smap2) alternative variant bSep08, mRNA.
smapey	smapey.aSep08		443	363		54	putative protein (smapey) mRNA.
smarbor	smarbor.aSep08		2568	340		55	putative protein (6.2 kD) (smarbor) mRNA.
Smarca1	Smarca1.aSep08	317575	13990	568	3	102	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 (Smarca1) alternative variant aSep08, mRNA.

Smarca2	Smarca2.aSep08	361745	104115	3896	23	954	SWI SNF-related matrix-associated actin-dependent regulator of chromatin a2 (Smarca2) alternative variant aSep08, mRNA.
Smarca2	Smarca2.bSep08	361745	61381	1055	7	351	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 2 like (Smarca2) alternative variant bSep08, mRNA.
Smarca2	Smarca2.cSep08	361745	58595	752	5	250	SWI SNF-related matrix-associated actin-dependent regulator of chromatin a2 (Smarca2) alternative variant cSep08, mRNA.
Smarca2	Smarca2.dSep08	361745	46324	839	7	220	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 2 (Smarca2) alternative variant dSep08, mRNA.
Smarca2	Smarca2.eSep08	361745	23250	1781	4	159	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 2 (Smarca2) alternative variant eSep08, mRNA.
Smarca2	Smarca2.fSep08	361745	4578	384	3	92	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 2 (Smarca2) alternative variant fSep08, mRNA.
Smarca2	Smarca2.gSep08	361745	9361	459	3	72	SWI SNF-related matrix-associated actin-dependent regulator of chromatin a2 like (Smarca2) alternative variant gSep08, mRNA.
Smarca2	Smarca2.kSep08	361745	3420	390	3	47	putative protein (Smarca2) alternative variant kSep08, mRNA.
Smarca4	Smarca4.aSep08	171379	91244	5578	34	1616	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (181.7 kD) (Smarca4) alternative variant aSep08, mRNA.
Smarca4	Smarca4.bSep08	171379	26819	1863	10	486	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant bSep08, mRNA.
Smarca4	Smarca4.cSep08	171379	23372	1251	8	343	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant cSep08, mRNA.
Smarca4	Smarca4.dSep08	171379	13705	895	7	297	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant dSep08, mRNA.
Smarca4	Smarca4.eSep08	171379	13050	564	3	187	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant eSep08, mRNA.
Smarca4	Smarca4.fSep08	171379	31929	536	5	173	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant fSep08, mRNA.
Smarca4	Smarca4.gSep08	171379	1341	363	2	63	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (Smarca4) alternative variant gSep08, mRNA.
Smarca5	Smarca5.bSep08	307766	3268	384	1	56	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 (6.2 kD) (Smarca5) alternative variant bSep08, mRNA.

Smarcad1	Smarcad1.bSep08	312398	1524	701	3	123	SWI SNF-related matrix-associated actin-dependent regulator of chromatin subfamily a containing dead H box 1 (Smarcad1) alternative variant bSep08, mRNA.
Smarcad1	Smarcad1.cSep08	312398	1679	771	2	84	SWI SNF-related matrix-associated actin-dependent regulator of chromatin subfamily a containing dead H box 1 (Smarcad1) alternative variant cSep08, mRNA.
Smarcal1	Smarcal1.bSep08	316477	46888	3055	18	910	hepa-related protein HARP (101.2 kD) (Smarcal1) alternative variant bSep08, mRNA.
Smarcal1	Smarcal1.bSep08	690314	46888	3055	18	910	hepa-related protein HARP (101.2 kD) (Smarcal1) alternative variant bSep08, mRNA.
Smarcal1	Smarcal1.cSep08	316477	4934	559	4	119	hepa-related protein HARP (13.3 kD) (Smarcal1) alternative variant cSep08, mRNA.
Smarcal1	Smarcal1.cSep08	690314	4934	559	4	119	hepa-related protein HARP (13.3 kD) (Smarcal1) alternative variant cSep08, mRNA.
Smarcal1	Smarcal1.eSep08	316477	2270	235	3	39	putative protein (4.4 kD) (Smarcal1) alternative variant eSep08, mRNA.
Smarcal1	Smarcal1.eSep08	690314	2270	235	3	39	putative protein (4.4 kD) (Smarcal1) alternative variant eSep08, mRNA.
Smarcb1	Smarcb1.bSep08	361825	15030	1164	1	323	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1 (Smarcb1) alternative variant bSep08, mRNA.
Smarcc1	Smarcc1.bSep08	301020	27509	1551	5	244	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1 (Smarcc1) alternative variant bSep08, mRNA.
Smarcc1	Smarcc1.cSep08	301020	17012	720	7	196	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1 (Smarcc1) alternative variant cSep08, mRNA.
Smarcc1	Smarcc1.dSep08	301020	26836	617	3	126	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1 (Smarcc1) alternative variant dSep08, mRNA.
Smarcc1	Smarcc1.eSep08	301020	17359	403	3	119	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1 (Smarcc1) alternative variant eSep08, mRNA.
Smarcd1	Smarcd1.bSep08	363002	1574	415	1	73	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1 (Smarcd1) alternative variant bSep08, mRNA.
Smarcd2	Smarcd2.bSep08	83833	3271	1510	7	450	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2 (50.3 kD) (Smarcd2) alternative variant bSep08, mRNA.
Smarcd2	Smarcd2.cSep08	83833	2188	1606	2	207	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2 (Smarcd2) alternative variant cSep08, mRNA.
Smarcd3	Smarcd3.bSep08	296732	11513	863	8	287	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 (Smarcd3) alternative variant bSep08, mRNA.

Smarcd3	Smarcd3.cSep08	296732	6829	897	7	259	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 (Smarcd3) alternative variant cSep08, mRNA.
Smarce1	Smarce1.bSep08	303518	16913	1323	7	194	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (22.8 kD) (Smarce1) alternative variant bSep08, mRNA.
Smarce1	Smarce1.cSep08	303518	17057	776	5	192	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (Smarce1) alternative variant cSep08, mRNA.
Smarce1	Smarce1.dSep08	303518	17166	1079	8	171	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (20.2 kD) (Smarce1) alternative variant dSep08, mRNA.
Smarce1	Smarce1.eSep08	303518	6771	995	2	154	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (17.9 kD) (Smarce1) alternative variant eSep08, mRNA.
Smarce1	Smarce1.fSep08	303518	16419	777	6	145	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (Smarce1) alternative variant fSep08, mRNA.
smarcha	smarcha.aSep08		16443	1355		85	putative protein (smarcha) mRNA.
smarchy	smarchy.aSep08		513	447		45	putative protein (5.4 kD) (smarchy) mRNA.
smardoy	smardoy.aSep08		4642	199		63	putative protein (smardoy) mRNA.
smarfee	smarfee.aSep08		1239	636		89	putative protein (9.9 kD) (smarfee) mRNA.
smarflu	smarflu.aSep08		12694	1337	4	331	mitochondrial GTPase 1 (37.0 kD) (smarflu) alternative variant aSep08, complete mRNA.
smarflu	smarflu.bSep08		11911	847	4	277	mitochondrial GTPase 1 (smarflu) alternative variant bSep08, mRNA.
smarflu	smarflu.cSep08		926	618	1	94	mitochondrial GTPase 1 homolog (smarflu) alternative variant cSep08, mRNA.
smarflu	smarflu.dSep08		4707	725	2	69	mitochondrial GTPase 1 (smarflu) alternative variant dSep08, mRNA.
smarfly	smarfly.bSep08		904	319	2	23	putative protein (2.8 kD) (smarfly) alternative variant bSep08, mRNA.
smarja	smarja.aSep08		23992	289		49	putative protein (smarja) mRNA.
smarmee	smarmee.aSep08		3296	774	2	79	ac1576 like (smarmee) alternative variant aSep08, mRNA.
smarmee	smarmee.bSep08		12553	713	2	38	putative protein (4.2 kD) (smarmee) alternative variant bSep08, mRNA.
smarnor	smarnor.aSep08		9747	734		244	shugoshin-like 1 (smarnor) mRNA.
smaroy	smaroy.aSep08		63859	765		152	putative nuclear protein (16.5 kD) (smaroy) mRNA.
smarpey	smarpey.aSep08		1907	758	4	166	hepatocellular carcinoma-associated td26 like (smarpey) alternative variant aSep08, mRNA.
smarpey	smarpey.cSep08		846	738	2	71	putative protein (8.6 kD) (smarpey) alternative variant cSep08, mRNA.
smarroy	smarroy.aSep08		26633	1400	4	74	CRA b like (smarroy) alternative variant aSep08, mRNA.
smarroy	smarroy.bSep08		25017	485	2	53	putative protein (smarroy) alternative variant bSep08, mRNA.



smarshaw	smarshaw.aSep08		2011	315	2	98	putative protein (smarshaw) alternative variant aSep08, mRNA.
smarshaw	smarshaw.bSep08		21066	772	2	35	putative protein (4.3 kD) (smarshaw) alternative variant bSep08, mRNA.
smarshee	smarshee.bSep08		3211	439	4	22	putative protein (2.7 kD) (smarshee) alternative variant bSep08, mRNA.
smartu	smartu.aSep08		923	731		45	putative protein (smartu) mRNA.
smarvo	smarvo.aSep08		229970	1780		540	astrotactin 2 (smarvo) alternative variant aSep08, mRNA.
smarvo	smarvo.bSep08		13603	451		95	astrotactin 2 CRA b (smarvo) alternative variant bSep08, mRNA.
smarwer	smarwer.aSep08		1734	298		99	maltase-glucoamylase (smarwer) mRNA.
smashaw	smashaw.aSep08		7966	955		37	putative protein (smashaw) mRNA.
smashee	smashee.aSep08		1174	262		24	putative protein (2.6 kD) (smashee) mRNA.
smatu	smatu.aSep08		5890	396		51	CRA a (smatu) mRNA.
smavo	smavo.aSep08		3840	836		136	fk506 binding protein 15 like (smavo) mRNA.
smawbor	smawbor.aSep08		1648	370		123	putative protein, with a coiled coil domain, of metazoan origin (smawbor) mRNA.
smawcha	smawcha.aSep08		24775	780		260	calmodulin regulated spectrin-associated protein 1 CRA b (smawcha) alternative variant aSep08, mRNA.
smawchy	smawchy.bSep08		1472	360	2	29	putative protein (smawchy) alternative variant bSep08, mRNA.
smawdoy	smawdoy.aSep08		4100	766		136	putative protein (14.7 kD) (smawdoy) mRNA.
smawer	smawer.aSep08		945	472		66	putative protein (7.9 kD) (smawer) mRNA.
smawfee	smawfee.aSep08		6504	306		57	putative protein of mammalian origin (smawfee) mRNA.
smawflu	smawflu.aSep08		1043	625		107	CRA a (smawflu) mRNA.
smawfly	smawfly.aSep08		1788	714		121	putative protein of vertebrate origin (smawfly) mRNA.
smawja	smawja.aSep08		6802	232		52	putative protein (smawja) mRNA.
smawmee	smawmee.aSep08		9502	381		75	putative protein (smawmee) mRNA.
smawnor	smawnor.aSep08		13212	309		37	putative protein (smawnor) mRNA.
smawpey	smawpey.aSep08		5615	812		270	dedicator of cytokinesis (smawpey) mRNA.
smawroy	smawroy.aSep08		586	489		67	putative protein (smawroy) mRNA.
smawshaw	smawshaw.aSep08		683	396		26	putative protein (smawshaw) mRNA.
smawshee	smawshee.aSep08		848	442	1	121	putative protein (smawshee) alternative variant aSep08, mRNA.
smawshee	smawshee.bSep08		831	421	1	114	putative protein (smawshee) alternative variant bSep08, mRNA.
smawshee	smawshee.cSep08		4766	386	1	49	putative protein (smawshee) alternative variant cSep08, mRNA.
smawtu	smawtu.aSep08		21040	1320		119	putative protein of mammalian origin (smawtu) mRNA.
smawvo	smawvo.aSep08		29350	209		69	astrotactin (smawvo) mRNA.
smawwer	smawwer.aSep08		2400	377		125	maltase-glucoamylase (smawwer) mRNA.
Smc1a	Smc1a.cSep08	63996	1146	583	3	92	putative protein of ancient origin (Smc1a) alternative variant cSep08, mRNA.

Smc1a	Smc1a.eSep08	63996	9299	547	4	37	putative protein (Smc1a) alternative variant eSep08, mRNA.
Smc2	Smc2.bSep08	362519	6480	419	1	139	putative protein, with a coiled coil domain, of ancient origin (Smc2) alternative variant bSep08, mRNA.
Smc3	Smc3.aSep08	29486	22373	2563		731	SMC protein, N-terminal and SMCs flexible hinge (Smc3) alternative variant aSep08, mRNA.
Smc3	Smc3.bSep08	29486	33593	2190		680	SMC protein, N-terminal and SMCs flexible hinge (Smc3) alternative variant bSep08, mRNA.
Smc3	Smc3.cSep08	29486	8170	2032		503	SMC protein, N-terminal (57.9 kD) (Smc3) alternative variant cSep08, mRNA.
Smc4	Smc4.bSep08	295107	13876	1506	9	465	SMC protein, N-terminal (Smc4) alternative variant bSep08, mRNA.
Smc4	Smc4.cSep08	295107	4094	547	3	117	SMC protein, N-terminal (Smc4) alternative variant cSep08, mRNA.
Smc4	Smc4.dSep08	295107	1157	590	2	71	putative cytoplasmic protein of ancient origin (8.1 kD) (Smc4) alternative variant dSep08, mRNA.
Smc5	Smc5.bSep08	293967	31827	1367	10	455	putative protein, with 3 coiled coil domains, of eukaryotic origin (Smc5) alternative variant bSep08, mRNA.
Smc5	Smc5.cSep08	293967	10300	980	6	198	putative protein, with a coiled coil domain, of ancient origin (Smc5) alternative variant cSep08, mRNA.
Smc5	Smc5.dSep08	293967	13032	552	5	184	putative protein, with 2 coiled coil domains, of eukaryotic origin (Smc5) alternative variant dSep08, mRNA.
Smc6l1	Smc6l1.bSep08	313961	11758	2715	4	134	putative protein of ancient origin (15.3 kD) (Smc6l1) alternative variant bSep08, mRNA.
Smcr7	Smcr7.aSep08	497916	4944	2451		494	putative protein of vertebrate origin (Smcr7) mRNA.
smeebor	smeebor.aSep08		5016	394		130	putative protein of metazoan origin (smeebor) mRNA.
smeecha	smeecha.aSep08		12975	463		90	putative protein (smeecha) mRNA.
smeechy	smeechy.aSep08		1985	409	2	72	putative protein (smeechy) alternative variant aSep08, mRNA.
smeechy	smeechy.bSep08		3880	829	3	90	putative protein (10.1 kD) (smeechy) alternative variant bSep08, mRNA.
smeechy	smeechy.cSep08		1312	608	2	67	putative protein (smeechy) alternative variant cSep08, mRNA.
smeedoy	smeedoy.aSep08		1668	570		66	putative protein (7.3 kD) (smeedoy) mRNA.
smeefee	smeefee.aSep08		3757	172		27	putative protein (smeefee) mRNA.
smeeflu	smeeflu.aSep08		1941	1547	4	132	ATH1 acid trehalase-like 1 (smeeflu) alternative variant aSep08, mRNA.
smeeflu	smeeflu.bSep08		399	319	1	105	ATH1 acid trehalase-like 1 CRA b (smeeflu) alternative variant bSep08, mRNA.
smeefly	smeefly.aSep08		9592	380		46	putative protein (5.3 kD) (smeefly) mRNA.
smeeja	smeeja.aSep08		3467	585		90	putative protein (smeeja) mRNA.
smee mee	smee mee.aSep08		1448	440	2	31	CRA b like (3.3 kD) (smee mee) alternative variant aSep08, mRNA.
smee mee	smee mee.bSep08		10141	1443	3	97	CRA b like (10.5 kD) (smee mee) alternative variant bSep08, mRNA.

smeemee	smeemee.dSep08		10176	471	3	60	putative protein (smeemee) alternative variant dSep08, mRNA.
smeenor	smeenor.aSep08		6212	722	4	125	sulfotransferase K2 (smeenor) alternative variant aSep08, mRNA.
smeeroy	smeeroy.aSep08		2647	569		179	putative protein, with a coiled coil domain (smeeroy) mRNA.
smeeshaw	smeeshaw.aSep08		4003	285		21	putative protein (smeeshaw) mRNA.
smeeshee	smeeshee.aSep08		4128	711		179	CRA b (smeeshee) mRNA.
smeetu	smeetu.aSep08		17220	437	3	145	CRA a (smeetu) alternative variant aSep08, mRNA.
smeevo	smeevo.aSep08		177953	571		189	astrotactin 2 (smeevo) mRNA.
smeewer	smeewer.aSep08		1734	298		99	maltase-glucoamylase (smeewer) mRNA.
Smek1	Smek1.aSep08	314388	42214	3989	12	833	smek homolog (Smek1) alternative variant aSep08, mRNA.
Smek1	Smek1.aSep08	690008	42214	3989	12	833	smek homolog (Smek1) alternative variant aSep08, mRNA.
Smek1	Smek1.bSep08	314388	8668	1028	7	342	smek homolog 1 (Smek1) alternative variant bSep08, mRNA.
Smek1	Smek1.bSep08	690008	8668	1028	7	342	smek homolog 1 (Smek1) alternative variant bSep08, mRNA.
Smek1	Smek1.dSep08	314388	2321	1269	2	106	smek 1 (10.8 kD) (Smek1) alternative variant dSep08, mRNA.
Smek1	Smek1.dSep08	690008	2321	1269	2	106	smek 1 (10.8 kD) (Smek1) alternative variant dSep08, mRNA.
Smek2	Smek2.bSep08	360993	3846	429	2	91	SMEK homolog 2, suppressor of mek1 (Dictyostelium) (Smek2) alternative variant bSep08, mRNA.
Smek2	Smek2.cSep08	360993	1911	824	2	74	SMEK homolog 2, suppressor of mek1 (Dictyostelium) (8.4 kD) (Smek2) alternative variant cSep08, mRNA.
smerbor	smerbor.aSep08		737	596		41	CRA c like (smerbor) mRNA.
smercha	smercha.bSep08		747	570	2	57	inositol e (smercha) alternative variant bSep08, mRNA.
smerchy	smerchy.aSep08		3524	393		130	nuclear receptor coactivator 3 (smerchy) mRNA.
smerdoy	smerdoy.aSep08		967	748		31	putative protein (smerdoy) mRNA.
smerfee	smerfee.aSep08		40927	797		47	putative protein (5.7 kD) (smerfee) mRNA.
smerflu	smerflu.aSep08		1940	437		52	putative protein (smerflu) mRNA.
smerfly	smerfly.aSep08		3767	1840		122	putative protein of mammalian origin (smerfly) mRNA.
smerja	smerja.aSep08		2201	721		58	putative protein (6.5 kD) (smerja) mRNA.
smermee	smermee.aSep08		4520	846		61	putative protein (smermee) mRNA.
smernor	smernor.aSep08		1143	1068		47	putative protein (4.8 kD) (smernor) mRNA.
smerpey	smerpey.aSep08		10513	401		75	putative protein (smerpey) mRNA.
smerroy	smerroy.aSep08		1089	510		21	putative protein (2.6 kD) (smerroy) mRNA.
smershaw	smershaw.aSep08		4453	384		72	putative protein (6.1 kD) (smershaw) mRNA.
smershee	smershee.aSep08		3769	447		32	putative protein (smershee) mRNA.
smertu	smertu.aSep08		19638	340		112	CRA b like (smertu) mRNA.
smervo	smervo.aSep08		7877	377		64	cdk5 regulatory associated protein 2 CRA b (smervo) mRNA.
smerwer	smerwer.aSep08		10403	506		168	maltase-glucoamylase (smerwer) mRNA.

smeybor	smeybor.aSep08		2376	455		118	putative protein of metazoan origin (smeybor) mRNA.
smeycha	smeycha.aSep08		2726	529		175	putative protein of mammalian origin (smeycha) mRNA.
smeychy	smeychy.aSep08		1966	703		75	putative protein (8.2 kD) (smeychy) mRNA.
smeydoy	smeydoy.aSep08		5735	709		58	putative protein (6.5 kD) (smeydoy) mRNA.
smeyfee	smeyfee.aSep08		5237	370		88	CRA b like (10.0 kD) (smeyfee) mRNA.
smeyflu	smeyflu.aSep08		1029	938		33	anoctamin 9 like (smeyflu) mRNA.
smeyfly	smeyfly.aSep08		3723	586	5	195	glioma tumor suppressor candidate region gene 1 like (smeyfly) alternative variant aSep08, mRNA.
smeyja	smeyja.aSep08		14926	435		69	putative protein (smeyja) mRNA.
smeymee	smeymee.bSep08		1108	642	2	62	putative protein (6.9 kD) (smeymee) alternative variant bSep08, mRNA.
smeynor	smeynor.aSep08		24694	849	3	52	CRA b like (5.8 kD) (smeynor) mRNA.
smeypey	smeypey.aSep08		5177	713	1	66	putative protein (7.6 kD) (smeypey) alternative variant aSep08, mRNA.
smeypey	smeypey.bSep08		7222	395	1	43	putative protein (4.7 kD) (smeypey) alternative variant bSep08, mRNA.
smeyroy	smeyroy.aSep08		1801	425		70	centrosomal protein 290kDa CRA b (smeyroy) mRNA.
smeyshaw	smeyshaw.aSep08		3862	435		58	putative protein (6.2 kD) (smeyshaw) mRNA.
smeyshee	smeyshee.aSep08		2141	335	2	76	CRA b (smeyshee) alternative variant aSep08, mRNA.
smeytu	smeytu.aSep08		1444	680		56	putative protein (6.1 kD) (smeytu) mRNA.
smeyvo	smeyvo.aSep08		34847	1140		379	cdk5 regulatory associated protein 2 (smeyvo) mRNA.
smeywer	smeywer.aSep08		2078	364		121	maltase-glucoamylase (smeywer) mRNA.
Smg5	Smg5.aSep08	681012	27112	4420	22	1064	smg-5 homolog nonsense mediated mRNA decay factor (Smg5) alternative variant aSep08, mRNA.
Smg5	Smg5.cSep08	681012	6957	415	4	112	smg-5 homolog nonsense mediated mRNA decay factor (Smg5) alternative variant cSep08, mRNA.
Smg7	Smg7.aSep08	360855	17339	1783		402	smg-7 homolog (Smg7) alternative variant aSep08, mRNA.
Smg7	Smg7.bSep08	360855	4121	995		331	smg-7 homolog (Smg7) alternative variant bSep08, mRNA.
Smndc1	Smndc1.cSep08	287768	10012	1136	5	199	putative protein, with a coiled coil domain, of eukaryotic origin (22.5 kD) (Smndc1) alternative variant cSep08, mRNA.
Smndc1	Smndc1.dSep08	287768	9163	797	3	193	putative protein, with 2 coiled coil domains, of eukaryotic origin (Smndc1) alternative variant dSep08, mRNA.
Smndc1	Smndc1.eSep08	287768	8906	853	4	159	putative protein, with a coiled coil domain, of eukaryotic origin (18.1 kD) (Smndc1) alternative variant eSep08, mRNA.
Smndc1	Smndc1.fSep08	287768	8343	584	3	121	putative protein, with a coiled coil domain, of fungal and metazoan origin (13.5 kD) (Smndc1) alternative variant fSep08, mRNA.
Smndc1	Smndc1.gSep08	287768	9913	864	4	121	putative protein, with a coiled coil domain, of fungal and metazoan origin (13.5 kD) (Smndc1) alternative variant gSep08, complete mRNA.
Smndc1	Smndc1.hSep08	287768	6978	575	2	79	putative protein (Smndc1) alternative variant hSep08, mRNA.

Smo	Smo.bSep08	25273	3386	2062	2	265	smoothened homolog (Drosophila) (29.3 kD) (Smo) alternative variant bSep08, mRNA.
smobor	smobor.aSep08		19934	627		59	putative protein (6.4 kD) (smobor) mRNA.
Smoc1	Smoc1.bSep08	314280	4906	338	3	111	SPARC related modular calcium binding 1 (Smoc1) alternative variant bSep08, mRNA.
Smoc1	Smoc1.cSep08	314280	1379	987	2	86	SPARC related modular calcium binding 1 (9.4 kD) (Smoc1) alternative variant cSep08, mRNA.
Smoc1	Smoc1.dSep08	314280	5405	783	2	51	SPARC related modular calcium binding 1 (Smoc1) alternative variant dSep08, mRNA.
Smoc2	Smoc2.bSep08	292401	1410	874	1	107	SPARC related modular calcium binding 2 (Smoc2) alternative variant bSep08, mRNA.
smocha	smocha.aSep08		3156	314	1	68	putative protein (smocha) alternative variant aSep08, mRNA.
smocha	smocha.bSep08		825	215	1	67	putative protein (smocha) alternative variant bSep08, mRNA.
smochy	smochy.bSep08		1499	400	2	39	putative protein (4.7 kD) (smochy) alternative variant bSep08, mRNA.
smodoy	smodoy.aSep08		4328	199		63	gag protein like (smodoy) mRNA.
smofee	smofee.bSep08		1670	546	2	73	putative protein (smofee) alternative variant bSep08, mRNA.
smoflu	smoflu.aSep08		3558	474		103	putative protein (smoflu) mRNA.
smofly	smofly.aSep08		5382	365		72	putative cytoplasmic protein (8.1 kD) (smofly) mRNA.
smoja	smoja.aSep08		1444	346		67	UDP-glucose ceramide glucosyltransferase-like 2 (smoja) mRNA.
smomee	smomee.aSep08		784	644		149	putative protein (16.4 kD) (smomee) mRNA.
smopey	smopey.aSep08		6080	407		135	dedicator of cytokinesis (smopey) mRNA.
smorbor	smorbor.aSep08		8196	663		160	armadillo repeat containing 2 like (smorbor) mRNA.
smorcha	smorcha.aSep08		3737	651		86	putative protein of mammalian origin (smorcha) mRNA.
smorchy	smorchy.aSep08		25937	573		40	putative protein (4.2 kD) (smorchy) mRNA.
smordoy	smordoy.aSep08		1575	313		22	putative protein (smordoy) mRNA.
smorfee	smorfee.aSep08		2023	482		41	putative protein (4.6 kD) (smorfee) mRNA.
smorflu	smorflu.aSep08		8238	394	4	131	PHD ring finger domains 1 like (smorflu) alternative variant aSep08, mRNA.
smorflu	smorflu.bSep08		8641	485	4	128	PHD ring finger domains 1 like (smorflu) alternative variant bSep08, mRNA.
smorfly	smorfly.aSep08		4204	656	4	69	putative protein (smorfly) alternative variant aSep08, mRNA.
smorja	smorja.aSep08		986	487		47	putative protein (5.3 kD) (smorja) mRNA.
smormee	smormee.aSep08		2255	257		67	putative protein (7.4 kD) (smormee) mRNA.
smornor	smornor.aSep08		144287	1049		349	raft-linking protein CRA a (smornor) mRNA.
smoroy	smoroy.aSep08		3416	853		69	putative protein (smoroy) mRNA.
smorpey	smorpey.aSep08		5773	578		192	anillin (smorpey) mRNA.
smorroy	smorroy.aSep08		2472	302		100	centrosomal protein 290kDa CRA b (smorroy) mRNA.
smorshaw	smorshaw.aSep08		8397	365		86	cd180 antigen CRA a like (smorshaw) mRNA.

smorshee	smorshee.aSep08		24036	408		56	putative protein (smorshee) mRNA.
smortu	smortu.aSep08		53744	1733		75	putative protein (smortu) mRNA.
smorvo	smorvo.aSep08		10655	595		198	cdk5 regulatory associated protein 2 (smorvo) mRNA.
smorwer	smorwer.aSep08		28919	1521		224	maltase-glucoamylase (smorwer) mRNA.
smoshaw	smoshaw.aSep08		1841	670		42	putative protein (4.9 kD) (smoshaw) mRNA.
smoshee	smoshee.aSep08		1370	636		59	putative protein (6.7 kD) (smoshee) mRNA.
smotu	smotu.aSep08		975	515		110	putative protein (smotu) mRNA.
smovo	smovo.aSep08		3085	498		90	putative protein of mammalian origin (smovo) mRNA.
smower	smower.aSep08		3237	390		129	maltase-glucoamylase CRA a (smower) mRNA.
Smox	Smox.bSep08	308652	31732	1980	6	514	spermine oxidase (57.1 kD) (Smox) alternative variant bSep08, mRNA.
Smox	Smox.cSep08	308652	34383	1799	7	468	spermine oxidase (Smox) alternative variant cSep08, mRNA.
Smox	Smox.dSep08	308652	34385	1691	6	465	spermine oxidase (Smox) alternative variant dSep08, mRNA.
Smox	Smox.fSep08	308652	29309	392	3	71	spermine oxidase (7.3 kD) (Smox) alternative variant fSep08, mRNA.
smoybor	smoybor.aSep08		10834	692		62	putative protein (7.4 kD) (smoybor) mRNA.
smoycha	smoycha.bSep08		863	538	2	40	putative protein (4.3 kD) (smoycha) alternative variant bSep08, mRNA.
smoychy	smoychy.aSep08		9392	2534		355	phosphatidylinositol 3 4 5-trisphosphate-dependent Rac exchanger 1 (smoychy) mRNA.
smoydoy	smoydoy.aSep08		2724	674	4	68	putative protein (smoydoy) alternative variant aSep08, mRNA.
smoyfee	smoyfee.aSep08		517	399		35	putative protein (4.1 kD) (smoyfee) mRNA.
smoyflu	smoyflu.aSep08		552	285		94	kinase 2 (smoyflu) mRNA.
smoyfly	smoyfly.aSep08		851	468		155	glioma tumor suppressor candidate region gene 1 like (smoyfly) mRNA.
smoyja	smoyja.aSep08		19150	312		27	putative protein (3.1 kD) (smoyja) mRNA.
smoylor	smoylor.aSep08		136496	782		105	putative protein (smoylor) alternative variant aSep08, mRNA.
smoylor	smoylor.bSep08		136392	783	1	46	putative protein (5.2 kD) (smoylor) alternative variant bSep08, mRNA.
smoymee	smoymee.aSep08		158818	1567	16	478	breast carcinoma amplified sequence 3 like (smoymee) alternative variant aSep08, mRNA.
smoymee	smoymee.bSep08		51676	1027	7	303	breast carcinoma amplified sequence 3 like (smoymee) alternative variant bSep08, mRNA.
smoymee	smoymee.cSep08		19803	306	5	101	breast carcinoma amplified sequence 3 like (smoymee) alternative variant cSep08, mRNA.
smoymee	smoymee.dSep08		12672	335	3	92	breast carcinoma amplified sequence 3 CRA c like (smoymee) alternative variant dSep08, mRNA.
smoymee	smoymee.eSep08		1812	414	3	49	putative protein (smoymee) alternative variant eSep08, mRNA.
smoynor	smoynor.aSep08		3841	405		71	kinesin family member 6 (smoynor) mRNA.
smoypey	smoypey.aSep08		31038	781	4	210	CRA a (smoypey) alternative variant aSep08, mRNA.

smoypey	smoypey.bSep08		14214	959	3	176	putative protein of mammalian origin (smoypey) alternative variant bSep08, mRNA.
smoypey	smoypey.cSep08		11309	352	2	117	CRA b like (smoypey) alternative variant cSep08, mRNA.
smoyroy	smoyroy.aSep08		2668	549		182	centrosomal protein 290kDa CRA a (smoyroy) mRNA.
smoyshaw	smoyshaw.aSep08		3439	681		226	microtubule-associated serine threonine-protein kinase 4 (smoyshaw) mRNA.
smoyshee	smoyshee.aSep08		20049	508		20	putative protein (smoyshee) mRNA.
smoytu	smoytu.aSep08		1094	863		43	echinoderm microtubule associated protein like 5 CRA b (smoytu) mRNA.
smoyvo	smoyvo.aSep08		1587	891		38	putative protein (4.0 kD) (smoyvo) mRNA.
smoywer	smoywer.aSep08		582	486		106	T-cell receptor like precursor (12.1 kD) (smoywer) mRNA.
Smpd1	Smpd1.bSep08	308909	2643	1537	4	479	sphingomyelin phosphodiesterase 1, acid lysosomal (Smpd1) alternative variant bSep08, mRNA.
Smpd1	Smpd1.cSep08	308909	1394	763	4	108	sphingomyelin phosphodiesterase 1, acid lysosomal (Smpd1) alternative variant cSep08, mRNA.
Smpd2	Smpd2.bSep08	83537	1319	687	1	56	sphingomyelin phosphodiesterase 2, neutral (Smpd2) alternative variant bSep08, mRNA.
Smpd4	Smpd4.aSep08	303790	23739	3444	9	860	sphingomyelin phosphodiesterase 4 (Smpd4) alternative variant aSep08, mRNA.
Smpd4	Smpd4.bSep08	303790	12547	598	1	199	sphingomyelin phosphodiesterase 4 (Smpd4) alternative variant bSep08, mRNA.
Smpd4	Smpd4.cSep08	303790	11616	635	2	156	sphingomyelin phosphodiesterase 4 (17.5 kD) (Smpd4) alternative variant cSep08, mRNA.
Smpdl3a	Smpdl3a.bSep08	294422	3922	1584	2	229	sphingomyelin phosphodiesterase, acid-like 3A (26.1 kD) (Smpdl3a) alternative variant bSep08, mRNA.
Smpdl3bandXkr8	Smpdl3bandXkr8.cSep08	313033	27311	1092	4	160	sphingomyelin phosphodiesterase acid-like 3B (18.3 kD) (Smpdl3bandXkr8) alternative variant cSep08, mRNA.
Smpdl3bandXkr8	Smpdl3bandXkr8.cSep08	362619	27311	1092	4	160	sphingomyelin phosphodiesterase acid-like 3B (18.3 kD) (Smpdl3bandXkr8) alternative variant cSep08, mRNA.
Smpx	Smpx.aSep08	84416	56203	890	2	85	small muscle protein, X-linked (9.1 kD) (Smpx) alternative variant aSep08, mRNA.
Smpx	Smpx.cSep08	84416	59583	520	4	85	small muscle protein, X-linked (9.1 kD) (Smpx) alternative variant cSep08, mRNA.
Smr.0	Smr.0.aSep08		5799	392		87	smr protein/MutS2 C-terminal (Smr.0) mRNA.
Smtn	Smtn.bSep08	289734	5470	682	4	175	smoothelin (Smtn) alternative variant bSep08, mRNA.
Smtn	Smtn.cSep08	289734	2035	699	2	92	putative protein (10.8 kD) (Smtn) alternative variant cSep08, mRNA.
Smtn	Smtn.eSep08	289734	1091	587	2	72	smoothelin (Smtn) alternative variant eSep08, mRNA.
Smtnl1	Smtnl1.aSep08	311167	6145	913		229	smoothelin-like 1 (Smtnl1) mRNA.
Smtnl2	Smtnl2.aSep08	679629	12833	1835		276	smoothelin-like 2 (Smtnl2) mRNA.
Smu1	Smu1.bSep08	117541	4689	705	3	151	smu-1 suppressor of mec-8 unc-52 homolog (Smu1) alternative variant bSep08, mRNA.
Smu1	Smu1.cSep08	117541	4662	420	3	140	smu-1 suppressor of mec-8 unc-52 homolog CRA b (Smu1) alternative variant cSep08, mRNA.
smubor	smubor.aSep08		1675	272		82	putative protein (smubor) mRNA.

smucha	smucha.aSep08		952	635	2	147	putative protein (16.2 kD) (smucha) mRNA.
smuchy	smuchy.aSep08		2215	900		299	CRA a (smuchy) mRNA.
smudoy	smudoy.aSep08		14508	720		93	putative protein (smudoy) mRNA.
smufee	smufee.aSep08		3475	1899		60	sideroflexin 1 (6.6 kD) (smufee) mRNA.
smuflu	smuflu.aSep08		2468	345		58	A disintegrin metallopeptidase domain 8 like (smuflu) mRNA.
smufly	smufly.aSep08		8231	262		39	putative protein (smufly) mRNA.
Smug1	Smug1.bSep08	315344	481	277	2	69	single-strand selective monofunctional uracil DNA glycosylase (Smug1) alternative variant bSep08, mRNA.
smuja	smuja.aSep08		24929	708		143	UDP-glucose ceramide glucosyltransferase-like (smuja) alternative variant aSep08, mRNA.
smuja	smuja.bSep08		9716	308		47	UDP-glucose ceramide glucosyltransferase-like 2 (smuja) alternative variant bSep08, mRNA.
smumee	smumee.bSep08		3050	962	3	91	putative protein (smumee) alternative variant bSep08, mRNA.
smumee	smumee.cSep08		3065	937	2	95	putative protein (smumee) alternative variant cSep08, mRNA.
smupey	smupey.aSep08		31580	1593	13	530	dicator of cytokinesis 6 CRA d (smupey) alternative variant aSep08, mRNA.
smuroy	smuroy.aSep08		14468	269		31	putative protein (smuroy) mRNA.
smushaw	smushaw.aSep08		828	718	2	74	putative protein (8.2 kD) (smushaw) alternative variant aSep08, mRNA.
smushee	smushee.aSep08		19260	712		101	putative nuclear protein (11.7 kD) (smushee) mRNA.
smutu	smutu.aSep08		1896	433	1	59	uncharacterized protein like (smutu) alternative variant aSep08, mRNA.
smutu	smutu.bSep08		1829	272		46	putative protein of mammalian origin (smutu) alternative variant bSep08, mRNA.
smuvo	smuvo.aSep08		4021	689		56	putative protein (smuvo) mRNA.
smuwer	smuwer.aSep08		32007	635		46	putative protein (smuwer) mRNA.
smybor	smybor.bSep08		1426	605	2	58	putative protein (6.3 kD) (smybor) alternative variant bSep08, mRNA.
smybor	smybor.cSep08		795	544	2	58	putative protein (6.3 kD) (smybor) alternative variant cSep08, mRNA.
smychy	smychy.aSep08		1380	388		46	putative protein (smychy) mRNA.
Smyd1	Smyd1.bSep08	297333	31502	541	3	103	putative protein of eukaryotic origin (Smyd1) alternative variant bSep08, mRNA.
Smyd2	Smyd2.aSep08	289372	41327	1578	12	444	zinc finger, MYND-type (Smyd2) alternative variant aSep08, mRNA.
Smyd2	Smyd2.bSep08	289372	3346	634	3	130	putative protein of eukaryotic origin (Smyd2) alternative variant bSep08, mRNA.
Smyd2	Smyd2.cSep08	289372	2044	662	2	107	putative protein of vertebrate origin (11.8 kD) (Smyd2) alternative variant cSep08, mRNA.
Smyd2	Smyd2.dSep08	289372	3262	1072	3	71	putative endoplasmic reticulum protein, with a transmembrane domain (7.7 kD) (Smyd2) alternative variant dSep08, mRNA.



Smyd5	Smyd5.aSep08	312503	10325	911	10	303	smyd family member 5 (Smyd5) alternative variant aSep08, mRNA.
Smyd5	Smyd5.bSep08	312503	723	246	2	39	smyd family member 5 (Smyd5) alternative variant bSep08, mRNA.
smydoy	smydoy.aSep08		968	304		35	putative protein (4.1 kD) (smydoy) mRNA.
smyfee	smyfee.aSep08		9551	1470		188	CRA b (smyfee) mRNA.
smyflu	smyflu.aSep08		959	701		47	putative protein (smyflu) mRNA.
smyfly	smyfly.aSep08		29012	724		52	putative protein (6.1 kD) (smyfly) mRNA.
smyja	smyja.aSep08		10347	921	3	69	putative protein (7.8 kD) (smyja) alternative variant aSep08, mRNA.
smymee	smymee.aSep08		4591	1002	3	53	CRA a like (6.1 kD) (smymee) alternative variant aSep08, mRNA.
smymee	smymee.bSep08		3902	967	2	34	CRA a like (smymee) alternative variant bSep08, mRNA.
smypey	smypey.aSep08		1494	739		61	putative protein (smypey) mRNA.
smyroy	smyroy.aSep08		7411	487		70	putative protein (7.6 kD) (smyroy) mRNA.
smyshaw	smyshaw.aSep08		11418	438		13	putative protein (smyshaw) mRNA.
smyshee	smyshee.aSep08		558	417		23	putative protein (2.5 kD) (smyshee) mRNA.
smytu	smytu.aSep08		30908	752		42	putative protein (smytu) mRNA.
smyvo	smyvo.aSep08		1827	269		47	putative protein (smyvo) mRNA.
smywer	smywer.aSep08		3089	1067		154	putative protein of vertebrate origin (17.7 kD) (smywer) mRNA.
snabor	snabor.aSep08		957	316		78	putative protein (snabor) mRNA.
snacha	snacha.aSep08		2879	931	2	66	CRA b like (7.3 kD) (snacha) alternative variant aSep08, complete mRNA.
snacha	snacha.bSep08		2835	883	2	66	CRA b like (7.3 kD) (snacha) alternative variant bSep08, mRNA.
snacha	snacha.cSep08		2370	813	1	40	CRA a like (4.3 kD) (snacha) alternative variant cSep08, complete mRNA.
snachy	snachy.aSep08		3862	758		223	phosphatidylinositol 3 4 5-trisphosphate-dependent Rac exchanger 1 (snachy) mRNA.
snadoy	snadoy.aSep08		1151	331		80	putative protein (snadoy) mRNA.
snafee	snafee.bSep08		1127	694	2	42	putative protein (snafee) alternative variant bSep08, mRNA.
snafly	snafly.aSep08		9856	380		44	zinc finger protein 541 like (snafly) mRNA.
snaja	snaja.aSep08		2833	553	2	60	putative protein (6.9 kD) (snaja) alternative variant aSep08, mRNA.
snalor	snalor.aSep08		1449	599		86	putative cytoplasmic protein (13.2 kD) (snalor) mRNA.
snamee	snamee.bSep08		23048	546	3	102	putative cytoplasmic protein (11.2 kD) (snamee) alternative variant bSep08, mRNA.
snamee	snamee.cSep08		5685	284	2	52	putative protein (snamee) alternative variant cSep08, mRNA.
snanor	snanor.aSep08		10500	407		59	putative protein (snanor) mRNA.
Snap25	Snap25.aSep08	25012	34363	1825	5	182	synaptosomal-associated protein 25 (Snap25) alternative variant aSep08, mRNA.

Snap25	Snap25.cSep08	25012	86945	988	5	175	synaptosomal-associated protein 25 (19.7 kD) (Snap25) alternative variant cSep08, mRNA.
Snap25	Snap25.dSep08	25012	69487	2760	1	144	synaptosomal-associated protein 25 (Snap25) alternative variant dSep08, mRNA.
Snap91	Snap91.aSep08	65178	24174	1206	11	401	synaptosomal-associated protein 91 (Snap91) alternative variant aSep08, mRNA.
Snap91	Snap91.bSep08	65178	25229	2346	11	314	synaptosomal-associated protein 91 (Snap91) alternative variant bSep08, mRNA.
Snap91	Snap91.cSep08	65178	27907	729	8	242	synaptosomal-associated protein 91 (Snap91) alternative variant cSep08, mRNA.
Snap91	Snap91.dSep08	65178	6765	389	4	122	synaptosomal-associated protein 91 (Snap91) alternative variant dSep08, mRNA.
Snap91	Snap91.eSep08	65178	13725	403	6	121	synaptosomal-associated protein 91 (Snap91) alternative variant eSep08, mRNA.
Snapap	Snapap.aSep08	295217	1876	825	4	136	SNAP-associated protein (14.9 kD) (Snapap) alternative variant aSep08, complete mRNA.
Snapap	Snapap.cSep08	295217	1461	672	3	65	SNAP-associated protein (Snapap) alternative variant cSep08, mRNA.
Snapap	Snapap.dSep08	295217	1431	614	3	37	SNAP-associated protein (4.7 kD) (Snapap) alternative variant dSep08, mRNA.
Snapc1	Snapc1.bSep08	314228	10661	743	4	138	small nuclear RNA activating complex, polypeptide 1 (Snapc1) alternative variant bSep08, mRNA.
Snapc2	Snapc2.aSep08	304204	2414	970	4	252	small nuclear RNA activating complex, polypeptide 2 (Snapc2) alternative variant aSep08, mRNA.
Snapc2	Snapc2.bSep08	304204	1969	853	4	152	small nuclear RNA activating complex, polypeptide 2 (15.7 kD) (Snapc2) alternative variant bSep08, mRNA.
Snapc3	Snapc3.bSep08	362537	21515	640	5	213	small nuclear RNA activating complex polypeptide 3 CRA c (Snapc3) alternative variant bSep08, mRNA.
Snapc3	Snapc3.cSep08	362537	23258	889	6	166	small nuclear RNA activating complex polypeptide 3 CRA b (Snapc3) alternative variant cSep08, mRNA.
Snapc3	Snapc3.dSep08	362537	22634	1675	4	67	putative protein (7.9 kD) (Snapc3) alternative variant dSep08, mRNA.
Snapc3	Snapc3.eSep08	362537	17166	425	3	54	putative protein (Snapc3) alternative variant eSep08, mRNA.
Snapc3	Snapc3.gSep08	362537	1666	729	2	48	putative protein (5.7 kD) (Snapc3) alternative variant gSep08, mRNA.
Snapc4	Snapc4.bSep08	362088	1588	752	2	216	small nuclear RNA activating complex, polypeptide 4 (Snapc4) alternative variant bSep08, mRNA.
Snapc4	Snapc4.cSep08	362088	1122	505	2	46	small nuclear RNA activating complex, polypeptide 4 (Snapc4) alternative variant cSep08, mRNA.
Snapc4	Snapc4.dSep08	362088	12607	1800	5	41	small nuclear RNA activating complex, polypeptide 4 (Snapc4) alternative variant dSep08, mRNA.
Snapc4	Snapc4.eSep08	362088	1799	418	3	50	small nuclear RNA activating complex, polypeptide 4 (Snapc4) alternative variant eSep08, mRNA.
Snapc4	Snapc4.gSep08	362088	2090	397	2	47	small nuclear RNA activating complex, polypeptide 4 (5.2 kD) (Snapc4) alternative variant gSep08, mRNA.
snapey	snapey.aSep08		998	676		87	putative protein (9.4 kD) (snapey) mRNA.

snarbor	snarbor.aSep08		399	284	1	54	putative protein (snarbor) alternative variant aSep08, mRNA.
snarbor	snarbor.bSep08		304	185	1	23	putative protein (snarbor) alternative variant bSep08, mRNA.
snarcha	snarcha.aSep08		1044	424		124	putative protein (snarcha) mRNA.
snarchy	snarchy.aSep08		1548	415		71	putative protein (snarchy) mRNA.
snardoy	snardoy.aSep08		1292	342		51	putative protein (snardoy) mRNA.
SNARE.0	SNARE.0.aSep08		5254	726	3	95	syntaxin (SNARE.0) alternative variant aSep08, mRNA.
SNARE.0	SNARE.0.cSep08		1541	712	2	60	syntaxin (6.7 kD) (SNARE.0) alternative variant cSep08, mRNA.
snarfee	snarfee.aSep08		15643	710	3	67	putative protein (7.4 kD) (snarfee) alternative variant aSep08, mRNA.
snarfee	snarfee.bSep08		16821	319	1	69	putative protein (snarfee) alternative variant bSep08, mRNA.
snarflu	snarflu.aSep08		59252	260		46	putative protein (snarflu) mRNA.
snarfly	snarfly.aSep08		1657	390		68	carcinoembryonic antigen-related like (snarfly) mRNA.
snarja	snarja.aSep08		1819	576		147	putative protein (snarja) mRNA.
snarlor	snarlor.aSep08		94167	390		53	putative protein (snarlor) mRNA.
snarmee	snarmee.aSep08		8110	746		248	integrator complex (snarmee) mRNA.
snarnor	snarnor.aSep08		7857	777		17	putative protein (snarnor) mRNA.
snaroy	snaroy.aSep08		972	370		123	centrosomal protein 290kDa CRA a (snaroy) mRNA.
snarpey	snarpey.aSep08		426	262		49	putative protein (5.4 kD) (snarpey) mRNA.
snarroy	snarroy.aSep08		762	355		31	putative protein (snarroy) mRNA.
snarshaw	snarshaw.aSep08		14216	385	4	128	tripartite motif protein 23 CRA b (snarshaw) mRNA.
snarshee	snarshee.aSep08	362068	3477	399		132	monogenic audiogenic seizure susceptibility 1 homolog (snarshee) mRNA.
snartu	snartu.aSep08		2021	746		75	putative protein (8.4 kD) (snartu) mRNA.
snarvo	snarvo.aSep08		7317	706		51	putative protein (6.0 kD) (snarvo) mRNA.
snarwer	snarwer.aSep08		1390	681		85	putative protein (9.3 kD) (snarwer) mRNA.
snashaw	snashaw.aSep08		16494	299		43	putative protein (snashaw) mRNA.
snashee	snashee.aSep08		46164	570		138	uncharacterized protein (snashee) mRNA.
snatu	snatu.aSep08		1440	392		130	echinoderm microtubule associated protein like 5 CRA b (snatu) mRNA.
snavo	snavo.aSep08		2719	369		122	transducin-like enhancer 1 (snavo) mRNA.
snawbor	snawbor.bSep08		28412	764	2	77	putative protein (8.3 kD) (snawbor) alternative variant bSep08, mRNA.
snawcha	snawcha.aSep08		1580	1473		90	putative protein (snawcha) mRNA.
snawchy	snawchy.aSep08		4926	320		61	putative protein (snawchy) mRNA.
snawdoy	snawdoy.aSep08		1176	740		97	putative protein (11.1 kD) (snawdoy) mRNA.
snawer	snawer.aSep08		8198	593		104	putative protein (snawer) mRNA.
snawfee	snawfee.aSep08		13457	741		64	putative protein (7.1 kD) (snawfee) mRNA.
snawflu	snawflu.aSep08		11580	335	2	31	putative protein (snawflu) mRNA.
snawfly	snawfly.aSep08		1323	1259		79	putative protein (snawfly) mRNA.

snawja	snawja.aSep08		33274	266		60	putative protein (snawja) mRNA.
snawlor	snawlor.aSep08		5796	1768	5	6	putative protein (snawlor) alternative variant aSep08, mRNA.
snawnor	snawnor.aSep08		805	286		92	putative protein (snawnor) mRNA.
snawpey	snawpey.aSep08		3478	329		40	putative protein (4.5 kD) (snawpey) mRNA.
snawroy	snawroy.aSep08		1015	617		55	putative protein (6.5 kD) (snawroy) mRNA.
snawshaw	snawshaw.aSep08		4953	434		144	kinesin heavy chain member 2 (snawshaw) mRNA.
snawshee	snawshee.aSep08		12017	444		37	putative protein (snawshee) mRNA.
snawtu	snawtu.aSep08		5069	376		68	putative protein (snawtu) mRNA.
snawvo	snawvo.aSep08		19631	392		35	putative protein (3.9 kD) (snawvo) mRNA.
snawwer	snawwer.aSep08		26055	1950		104	putative mitochondrial protein (11.9 kD) (snawwer) mRNA.
Snca	Snca.bSep08	29219	65819	378	1	67	synuclein, alpha (Snca) alternative variant bSep08, mRNA.
Sncaip	Sncaip.bSep08	307309	966	489	2	60	synuclein, alpha interacting protein (synphilin) (Sncaip) alternative variant bSep08, mRNA.
Sncaip	Sncaip.cSep08	307309	8451	543	2	41	synuclein, alpha interacting protein (synphilin) (Sncaip) alternative variant cSep08, mRNA.
Sncb	Sncb.bSep08	113893	8011	714	3	116	synuclein, beta (Sncb) alternative variant bSep08, mRNA.
Sncb	Sncb.cSep08	113893	1149	601	3	23	synuclein, beta (2.8 kD) (Sncb) alternative variant cSep08, mRNA.
Sncb	Sncb.dSep08	113893	966	308	2	56	synuclein, beta (Sncb) alternative variant dSep08, mRNA.
Sncg	Sncg.bSep08	64347	4377	620	6	148	synuclein, gamma (15.6 kD) (Sncg) alternative variant bSep08, mRNA.
Snd1	Snd1.bSep08	64635	54842	1642	9	310	maternal tudor protein (35.2 kD) (Snd1) alternative variant bSep08, mRNA.
Snd1	Snd1.cSep08	64635	217006	684	6	227	staphylococcus nuclease (SNase-like) (Snd1) alternative variant cSep08, mRNA.
Snd1	Snd1.dSep08	64635	18875	411	4	110	putative protein of eukaryotic origin (Snd1) alternative variant dSep08, mRNA.
Snd1	Snd1.eSep08	64635	40851	326	3	108	putative protein of eukaryotic origin (Snd1) alternative variant eSep08, mRNA.
Snd1	Snd1.fSep08	64635	646	538	2	80	putative protein of metazoan origin (9.2 kD) (Snd1) alternative variant fSep08, mRNA.
Snd1	Snd1.gSep08	64635	692	294	3	31	putative protein of metazoan origin (Snd1) alternative variant gSep08, mRNA.
Sned1	Sned1.aSep08	316638	10117	610		201	sushi, nidogen and EGF-like domains 1 (Sned1) mRNA.
sneebor	sneebor.aSep08		720	308		46	putative protein (5.0 kD) (sneebor) mRNA.
sneecha	sneecha.aSep08		3879	552		116	putative protein (sneecha) mRNA.
sneechy	sneechy.aSep08		6433	381		70	putative protein (sneechy) mRNA.
sneedoy	sneedoy.aSep08		2958	602		63	putative protein (7.3 kD) (sneedoy) mRNA.
sneefee	sneefee.aSep08		6325	744		103	putative protein (11.1 kD) (sneefee) mRNA.
sneeflu	sneeflu.aSep08		6295	525		105	putative protein (sneeflu) mRNA.
sneefly	sneefly.aSep08		1580	736		133	putative protein of mammalian origin (sneefly) mRNA.
sneeja	sneeja.aSep08		561	402		32	T cell receptor (sneeja) mRNA.
sneelor	sneelor.aSep08		8565	980		80	putative protein (sneelor) mRNA.

sneemee	sneemee.aSep08		500	402		115	putative protein (sneemee) mRNA.
sneenor	sneenor.aSep08		1043	674		62	putative protein (sneenor) mRNA.
sneepey	sneepey.aSep08		4194	419		121	putative protein (sneepey) mRNA.
sneeroy	sneeroy.aSep08		766	580		33	putative protein (3.8 kD) (sneeroy) mRNA.
sneeshaw	sneeshaw.aSep08		34366	286		95	kinesin-like protein kif2a (sneeshaw) mRNA.
sneeshee	sneeshee.aSep08		2586	351		116	glutamyl aminopeptidase (sneeshee) mRNA.
sneetu	sneetu.aSep08		13341	399		67	putative protein (sneetu) mRNA.
sneevo	sneevo.aSep08		102373	640		71	putative protein (sneevo) mRNA.
sneewer	sneewer.aSep08		35365	367		96	putative protein of mammalian origin (sneewer) mRNA.
snerbor	snerbor.aSep08		2281	608		36	putative protein (3.9 kD) (snerbor) mRNA.
snercha	snercha.aSep08		1404	1277		16	putative protein (1.9 kD) (snercha) mRNA.
snerchy	snerchy.aSep08		686	407		90	putative protein (snerchy) mRNA.
snerdoy	snerdoy.aSep08		1682	1307		114	putative protein (12.5 kD) (snerdoy) mRNA.
snerfee	snerfee.aSep08		6394	785		112	putative protein (snerfee) mRNA.
snerflu	snerflu.aSep08		4895	666	2	144	CRA b like (15.8 kD) (snerflu) alternative variant aSep08, mRNA.
snerflu	snerflu.cSep08		27775	780	5	66	CRA b like (7.2 kD) (snerflu) alternative variant cSep08, mRNA.
snerflu	snerflu.dSep08		15511	222	3	22	putative protein (2.4 kD) (snerflu) alternative variant dSep08, mRNA.
snerfly	snerfly.aSep08		5086	348		116	kinesin light CRA a (snerfly) mRNA.
snerja	snerja.aSep08		3187	199		63	gag protein like (snerja) mRNA.
snerlor	snerlor.aSep08		1317	703		61	putative protein of mammalian origin (7.0 kD) (snerlor) mRNA.
snermee	snermee.aSep08		6972	754	2	74	putative protein (snermee) alternative variant aSep08, mRNA.
snernor	snernor.bSep08		2609	657	2	70	putative protein of mammalian origin (7.9 kD) (snernor) alternative variant bSep08, mRNA.
snerpey	snerpey.aSep08		10629	346		39	immunoglobulin superfamily member 9B (snerpey) mRNA.
snerroy	snerroy.aSep08		7564	385		52	putative protein (snerroy) mRNA.
snershaw	snershaw.aSep08		9380	711		112	putative protein (12.6 kD) (snershaw) mRNA.
snershee	snershee.aSep08		8968	254		82	phosphatidylinositol glycan anchor biosynthesis class (snershee) mRNA.
snertu	snertu.aSep08		22351	335		111	putative protein of bilateral origin (snertu) mRNA.
snervo	snervo.aSep08		2009	391		22	putative protein (snervo) mRNA.
snerwer	snerwer.aSep08		56504	2639		183	contactin associated protein-like 2 CRA b (20.0 kD) (snerwer) mRNA.
sneybor	sneybor.bSep08		3768	1103	2	30	putative protein (3.4 kD) (sneybor) alternative variant bSep08, mRNA.
sneycha	sneycha.aSep08		1374	526		67	microtubule-associated protein 1 light beta CRA a (sneycha) mRNA.
sneychy	sneychy.aSep08		16076	1076		65	putative protein (7.1 kD) (sneychy) alternative variant aSep08, mRNA.
sneydoy	sneydoy.aSep08		16958	1209	2	284	rotatin CRA e (sneydoy) alternative variant aSep08, mRNA.

sneydoy	sneydoy.bSep08		13481	672	1	178	rotatin (sneydoy) alternative variant bSep08, mRNA.
sneyfee	sneyfee.aSep08		5539	412		62	isoleucyl-tRNA synthetase (sneyfee) mRNA.
sneyflu	sneyflu.aSep08		2319	319		102	putative protein (sneyflu) mRNA.
sneyfly	sneyfly.aSep08		17249	631		209	MAP microtubule affinity-regulating kinase (sneyfly) mRNA.
sneylor	sneylor.bSep08		4200	689	2	71	putative mitochondrial protein (7.6 kD) (sneylor) alternative variant bSep08, mRNA.
sneymee	sneymee.aSep08		1655	592	4	197	benzodiazapine receptor associated protein 1 CRA a (sneymee) alternative variant aSep08, mRNA.
sneynor	sneynor.aSep08		1184	439	2	131	putative protein (sneynor) alternative variant aSep08, mRNA.
sneypey	sneypey.aSep08		6982	1554		123	putative protein of mammalian origin (sneypey) mRNA.
sneyroy	sneyroy.aSep08		9194	185		57	putative protein (sneyroy) mRNA.
sneyshaw	sneyshaw.aSep08		79805	552		56	putative protein (6.5 kD) (sneyshaw) mRNA.
sneytu	sneytu.aSep08		7461	534		61	putative protein (sneytu) mRNA.
sneyvo	sneyvo.aSep08		5191	321		72	putative protein (sneyvo) mRNA.
sneywer	sneywer.aSep08		4492	837		83	putative protein (6.4 kD) (sneywer) mRNA.
Snf1lk	Snf1lk.bSep08	59329	1545	627	1	208	SNF1-like kinase (Snf1lk) alternative variant bSep08, mRNA.
SNF2_N.0	SNF2_N.0.aSep08		14979	1113		301	transcriptional regulator atrx (SNF2_N.0) mRNA.
SNF2_N.1	SNF2_N.1.aSep08		24816	1294	1	359	stretch responsive protein 278 CRA b (SNF2_N.1) alternative variant aSep08, mRNA.
SNF2_N.1	SNF2_N.1.bSep08		17594	883	1	229	stretch responsive protein 278 CRA b (SNF2_N.1) alternative variant bSep08, mRNA.
SNF2_N.2	SNF2_N.2.aSep08		9924	793	4	264	CRA b (SNF2_N.2) alternative variant aSep08, mRNA.
SNF2_N.3	SNF2_N.3.aSep08		29960	3140	19	748	rad54 (84.9 kD) (SNF2_N.3) alternative variant aSep08, mRNA.
SNF2_N.3	SNF2_N.3.bSep08		8350	755	6	225	rad54 (SNF2_N.3) alternative variant bSep08, mRNA.
SNF2_N.3	SNF2_N.3.cSep08		1190	406	4	75	rad54 (SNF2_N.3) alternative variant cSep08, mRNA.
SNF2_N.4	SNF2_N.4.aSep08		6612	734		244	SNF2-related (SNF2_N.4) mRNA.
Snf7.0	Snf7.0.aSep08		37464	1135	4	286	chromatin modifying protein 4b (Snf7.0) alternative variant aSep08, mRNA.
Snf8	Snf8.bSep08	287645	11961	672	6	127	SNF8, ESCRT-II complex subunit, homolog (S. cerevisiae) (Snf8) alternative variant bSep08, mRNA.
Snf8	Snf8.cSep08	287645	10371	757	8	122	SNF8, ESCRT-II complex subunit, homolog (S. cerevisiae) (Snf8) alternative variant cSep08, mRNA.
Snf8	Snf8.dSep08	287645	1446	661	2	47	SNF8, ESCRT-II complex subunit, homolog (S. cerevisiae) (4.3 kD) (Snf8) alternative variant dSep08, mRNA.
Snhg8	Snhg8.aSep08	361111	3889	445		143	small nucleolar RNA host gene (non-protein coding) 8 (Snhg8) mRNA.
Snip	Snip.bSep08	56029	4951	1091	5	331	SNAP25-interacting protein (Snip) alternative variant bSep08, mRNA.
Snip	Snip.cSep08	56029	10917	404	4	134	SNAP25-interacting protein (Snip) alternative variant cSep08, mRNA.
Snip	Snip.dSep08	56029	10577	399	4	133	SNAP25-interacting protein (Snip) alternative variant dSep08, mRNA.

Snip1	Snip1.bSep08	313588	2074	1093	2	243	smad nuclear interacting protein 1 (28.3 kD) (Snip1) alternative variant bSep08, mRNA.
Snip1	Snip1.cSep08	313588	5128	403	2	118	smad nuclear interacting protein 1 (Snip1) alternative variant cSep08, mRNA.
snobor	snobor.aSep08		2875	925		188	putative protein, with a coiled coil domain (snobor) mRNA.
snocha	snocha.aSep08		1957	413		119	putative protein (snocha) mRNA.
snochy	snochy.aSep08		2183	1697	2	84	solute carrier family 9 member 8 CRA a (snochy) alternative variant aSep08, mRNA.
snodoy	snodoy.aSep08		12090	577		104	putative secreted or extracellular protein precursor (10.9 kD) (snodoy) mRNA.
snofee	snofee.aSep08		53262	1799	2	472	putative protein (snofee) alternative variant aSep08, mRNA.
snofee	snofee.bSep08		12887	868	1	109	putative cytoplasmic protein (12.5 kD) (snofee) alternative variant bSep08, mRNA.
snofee	snofee.eSep08		64181	665	1	37	putative protein (4.3 kD) (snofee) alternative variant eSep08, mRNA.
snoflu	snoflu.aSep08		2071	1099	4	109	ASM15 like (11.8 kD) (snoflu) alternative variant aSep08, mRNA.
snoflu	snoflu.dSep08		2672	2320	5	59	putative protein (6.3 kD) (snoflu) alternative variant dSep08, complete mRNA.
snofly	snofly.aSep08		10104	690		100	putative nuclear protein (11.3 kD) (snofly) mRNA.
snoja	snoja.aSep08		13634	1611		82	putative protein (9.7 kD) (snoja) mRNA.
snolor	snolor.aSep08		7546	382		127	roundabout homolog 2 (snolor) mRNA.
snomee	snomee.aSep08		2557	418		139	integrator complex CRA b (snomee) mRNA.
snonor	snonor.aSep08		1077	580		39	putative protein (4.5 kD) (snonor) mRNA.
snopey	snopey.aSep08		5898	546		44	putative protein (5.0 kD) (snopey) mRNA.
snorbor	snorbor.aSep08		17923	1375		63	putative protein (7.5 kD) (snorbor) mRNA.
snorcha	snorcha.aSep08		763	492		76	putative protein (snorcha) mRNA.
snorchy	snorchy.aSep08		727	343		45	putative protein (snorchy) mRNA.
snorfee	snorfee.aSep08		6586	326		102	putative protein (snorfee) mRNA.
snorflu	snorflu.aSep08		23028	1797		15	putative protein (snorflu) alternative variant aSep08, mRNA.
snorflu	snorflu.bSep08		9847	407		48	putative protein (snorflu) alternative variant bSep08, mRNA.
snorfly	snorfly.aSep08		11095	2306		264	exocyst complex component 3-like protein 2 (snorfly) mRNA.
snorlor	snorlor.aSep08		43852	736		59	putative protein (6.5 kD) (snorlor) mRNA.
snormee	snormee.aSep08		763	412		25	putative protein (2.5 kD) (snormee) mRNA.
snornor	snornor.aSep08		8473	1027		258	ubiquitin ligase E3 alpha-II (snornor) mRNA.
snoroy	snoroy.aSep08		1587	799	2	102	centrosomal protein of like (snoroy) alternative variant aSep08, mRNA.
snorpey	snorpey.aSep08		23797	669		62	putative protein (7.5 kD) (snorpey) mRNA.
snorrooy	snorrooy.aSep08		197482	1246		43	putative protein (5.2 kD) (snorrooy) mRNA.
snorshaw	snorshaw.aSep08		1003	748		45	putative protein (5.1 kD) (snorshaw) mRNA.

snorshee	snorshee.aSep08		1094	834		84	putative protein of mammalian origin (snorshee) mRNA.
snortu	snortu.aSep08		3364	721		75	putative protein (8.5 kD) (snortu) mRNA.
snorvo	snorvo.aSep08		59545	713		74	putative secreted or extracellular protein precursor (8.1 kD) (snorvo) mRNA.
snorwer	snorwer.aSep08		3493	869		289	CRA a (snorwer) mRNA.
snoshaw	snoshaw.aSep08		4199	2722		60	erbB2 interacting protein (snoshaw) mRNA.
snoshee	snoshee.aSep08		2400	820		84	putative protein (snoshee) mRNA.
snotu	snotu.aSep08		2671	480		68	putative protein (snotu) mRNA.
snovo	snovo.aSep08		6848	691		32	putative protein (3.7 kD) (snovo) mRNA.
snower	snower.aSep08		38780	982	2	297	CRA d (snower) alternative variant aSep08, mRNA.
snower	snower.bSep08		6663	775	1	258	CRA d (snower) alternative variant bSep08, mRNA.
snoybor	snoybor.aSep08		2117	882		67	putative protein (7.1 kD) (snoybor) mRNA.
snoycha	snoycha.aSep08		1770	482		104	transcription factor (snoycha) mRNA.
snoychy	snoychy.aSep08		1820	588		88	putative protein (10.2 kD) (snoychy) mRNA.
snoyfee	snoyfee.aSep08		176045	825	5	275	centromere protein P (snoyfee) alternative variant aSep08, mRNA.
snoyfee	snoyfee.bSep08		17859	559	1	81	centromere protein P (9.2 kD) (snoyfee) alternative variant bSep08, mRNA.
snoyflu	snoyflu.aSep08		1899	391		102	transient receptor potential cation channel subfamily M member 5 (snoyflu) mRNA.
snoylor	snoylor.aSep08		12642	514		55	putative protein (snoylor) mRNA.
snoymee	snoymee.aSep08		700	370		52	putative protein (snoymee) mRNA.
snoynor	snoynor.aSep08		8478	819	5	272	ubiquitin ligase E3 alpha-II (snoynor) alternative variant aSep08, mRNA.
snoynor	snoynor.bSep08		3307	640	1	110	ubiquitin ligase E3 alpha-II (snoynor) alternative variant bSep08, mRNA.
snoypey	snoypey.aSep08		1808	484		140	putative protein (snoypey) mRNA.
snoyroy	snoyroy.aSep08		5791	338		112	neuron navigator 3 CRA b (snoyroy) mRNA.
snoyshaw	snoyshaw.aSep08		12930	593		38	putative protein (4.4 kD) (snoyshaw) mRNA.
snoytu	snoytu.aSep08		27333	1462	2	306	putative protein of ancient origin (snoytu) alternative variant aSep08, mRNA.
snoyvo	snoyvo.aSep08		13438	542		112	putative protein, with a coiled coil domain (snoyvo) mRNA.
snoywer	snoywer.aSep08		6878	713		38	putative protein (snoywer) mRNA.
Snrp70	Snrp70.aSep08	361574	20433	1578	10	451	small nuclear ribonucleoprotein (52.1 kD) (Snrp70) alternative variant aSep08, complete mRNA.
Snrp70	Snrp70.bSep08	361574	7197	3639	4	352	small nuclear ribonucleoprotein polypeptide (39.3 kD) (Snrp70) alternative variant bSep08, mRNA.
Snrp70	Snrp70.cSep08	361574	19660	1065	10	308	small nuclear ribonucleoprotein (Snrp70) alternative variant cSep08, mRNA.
Snrp70	Snrp70.fSep08	361574	3739	370	3	63	putative protein (Snrp70) alternative variant fSep08, mRNA.
Snrpa	Snrpa.bSep08	292729	6894	883	6	228	small nuclear ribonucleoprotein polypeptide A (Snrpa) alternative variant bSep08, mRNA.



Snrpb	Snrpb.bSep08	171365	6234	773	6	140	small nuclear ribonucleoprotein polypeptides B and B1 (Snrpb) alternative variant bSep08, mRNA.
Snrpb2	Snrpb2.bSep08	362223	9474	1041	4	218	u2 small nuclear ribonucleoprotein B (24.6 kD) (Snrpb2) alternative variant bSep08, mRNA.
Snrpb2	Snrpb2.cSep08	362223	6011	452	1	37	u2 small nuclear ribonucleoprotein B (Snrpb2) alternative variant cSep08, mRNA.
Snrpc	Snrpc.aSep08	361808	10590	725	1	172	u1 small nuclear ribonucleoprotein C (Snrpc) alternative variant aSep08, mRNA.
Snrpc	Snrpc.bSep08	361808	10524	769	1	118	u1 small nuclear ribonucleoprotein C (12.3 kD) (Snrpc) alternative variant bSep08, mRNA.
Snta1	Snta1.aSep08	362242	30564	2102	4	535	syntrophin, acidic 1 (Snta1) alternative variant aSep08, mRNA.
Snta1	Snta1.bSep08	362242	4101	643	1	214	syntrophin, acidic 1 (Snta1) alternative variant bSep08, mRNA.
Sntg1	Sntg1.aSep08	500394	54298	464		83	syntrophin, gamma 1 (Sntg1) mRNA.
snuor	snuor.aSep08		2388	1086		279	putative protein of mammalian origin (snuor) mRNA.
snucha	snucha.aSep08		1201	590		47	putative protein (5.3 kD) (snucha) mRNA.
snuchy	snuchy.aSep08		11357	692	4	230	solute carrier family 9 member 8 CRA a (snuchy) alternative variant aSep08, mRNA.
snuchy	snuchy.bSep08		11920	433	5	144	solute carrier family 9 member 8 CRA a (snuchy) alternative variant bSep08, mRNA.
snuchy	snuchy.cSep08		754	302	2	100	solute carrier family 9 member 8 CRA c (snuchy) alternative variant cSep08, mRNA.
snudoy	snudoy.aSep08		43756	381		34	putative protein (3.6 kD) (snudoy) mRNA.
snufee	snufee.aSep08		4128	750		46	putative protein (5.4 kD) (snufee) mRNA.
snufly	snufly.aSep08		10921	150		29	putative protein (snufly) mRNA.
snufly	snufly.aSep08		1839	485		161	dapper homolog 3 (snufly) mRNA.
snuja	snuja.aSep08		423	299		40	putative protein (4.5 kD) (snuja) mRNA.
snulor	snulor.aSep08		3649	702		110	putative protein (snulor) mRNA.
snumee	snumee.aSep08		3577	1624		151	integrator complex (snumee) mRNA.
snunor	snunor.aSep08		4551	419		104	putative protein (snunor) mRNA.
snupey	snupey.aSep08		4640	537		45	putative protein (snupey) mRNA.
Snupn	Snupn.bSep08	316108	27788	2175	7	202	snurportin 1 (23.2 kD) (Snupn) alternative variant bSep08, mRNA.
Snupn	Snupn.cSep08	316108	12333	391	4	67	snurportin 1 (Snupn) alternative variant cSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.bSep08	81781	19773	1304	9	221	small nuclear (23.1 kD) (SnurfandSnrpn) alternative variant bSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.bSep08	113938	19773	1304	9	221	small nuclear (23.1 kD) (SnurfandSnrpn) alternative variant bSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.cSep08	81781	19546	921	7	140	small nuclear ribonucleoprotein (SnurfandSnrpn) alternative variant cSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.cSep08	113938	19546	921	7	140	small nuclear ribonucleoprotein (SnurfandSnrpn) alternative variant cSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.dSep08	81781	15879	694	5	90	SNRPN upstream reading frame (10.7 kD) (SnurfandSnrpn) alternative variant dSep08, mRNA.

SnurfandSnrpn	SnurfandSnrpn.dSep08	113938	15879	694	5	90	SNRPN upstream reading frame (10.7 kD) (SnurfandSnrpn) alternative variant dSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.eSep08	81781	19783	437	4	77	SNRPN upstream reading frame (9.1 kD) (SnurfandSnrpn) alternative variant eSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.eSep08	113938	19783	437	4	77	SNRPN upstream reading frame (9.1 kD) (SnurfandSnrpn) alternative variant eSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.gSep08	81781	15898	848	6	90	SNRPN upstream reading frame (10.7 kD) (SnurfandSnrpn) alternative variant gSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.gSep08	113938	15898	848	6	90	SNRPN upstream reading frame (10.7 kD) (SnurfandSnrpn) alternative variant gSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.hSep08	81781	18475	804	7	71	SNRPN upstream reading frame (8.4 kD) (SnurfandSnrpn) alternative variant hSep08, complete mRNA.
SnurfandSnrpn	SnurfandSnrpn.hSep08	113938	18475	804	7	71	SNRPN upstream reading frame (8.4 kD) (SnurfandSnrpn) alternative variant hSep08, complete mRNA.
SnurfandSnrpn	SnurfandSnrpn.iSep08	81781	6455	726	3	59	putative protein (SnurfandSnrpn) alternative variant iSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.iSep08	113938	6455	726	3	59	putative protein (SnurfandSnrpn) alternative variant iSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.jSep08	81781	9135	666	3	46	putative protein (5.4 kD) (SnurfandSnrpn) alternative variant jSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.jSep08	113938	9135	666	3	46	putative protein (5.4 kD) (SnurfandSnrpn) alternative variant jSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.kSep08	81781	17410	592	6	70	SNRPN upstream reading frame (SnurfandSnrpn) alternative variant kSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.kSep08	113938	17410	592	6	70	SNRPN upstream reading frame (SnurfandSnrpn) alternative variant kSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.lSep08	81781	135526	585	7	86	SNRPN upstream reading frame (10.2 kD) (SnurfandSnrpn) alternative variant lSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.lSep08	113938	135526	585	7	86	SNRPN upstream reading frame (10.2 kD) (SnurfandSnrpn) alternative variant lSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.mSep08	81781	16766	547	5	63	putative protein (SnurfandSnrpn) alternative variant mSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.mSep08	113938	16766	547	5	63	putative protein (SnurfandSnrpn) alternative variant mSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.oSep08	81781	15593	342	4	24	putative protein (SnurfandSnrpn) alternative variant oSep08, mRNA.
SnurfandSnrpn	SnurfandSnrpn.oSep08	113938	15593	342	4	24	putative protein (SnurfandSnrpn) alternative variant oSep08, mRNA.
snuroy	snuroy.aSep08		5677	527		175	centrosomal protein 290kDa CRA b (snuroy) mRNA.
snushaw	snushaw.aSep08		2645	1774		13	putative protein (1.6 kD) (snushaw) mRNA.
snushee	snushee.aSep08		70905	1783			
snutu	snutu.aSep08		5187	313		99	CRA b like (snutu) mRNA.
snuvo	snuvo.aSep08		30690	575		191	protein tyrosine phosphatase receptor type D (snuvo) mRNA.
snuwer	snuwer.aSep08		1869	350		55	putative protein (snuwer) mRNA.
Snx1	Snx1.bSep08	84471	6979	833	6	220	sorting nexin 1 (Snx1) alternative variant bSep08, mRNA.

Snx1	Snx1.cSep08	84471	32126	696	7	195	sorting nexin 1 (Snx1) alternative variant cSep08, mRNA.
Snx1	Snx1.dSep08	84471	2461	481	5	101	sorting nexin 1 (Snx1) alternative variant dSep08, mRNA.
Snx2	Snx2.bSep08	291464	22312	1748	10	445	sorting nexin 2 (Snx2) alternative variant bSep08, mRNA.
Snx2	Snx2.cSep08	291464	10556	604	7	172	sorting nexin 2 CRA c (Snx2) alternative variant cSep08, mRNA.
Snx2	Snx2.dSep08	291464	9740	911	6	157	sorting nexin 2 (Snx2) alternative variant dSep08, mRNA.
Snx2	Snx2.eSep08	291464	11577	793	6	132	sorting nexin 2 (15.6 kD) (Snx2) alternative variant eSep08, mRNA.
Snx2	Snx2.fSep08	291464	7708	442	3	81	sorting nexin 2 (Snx2) alternative variant fSep08, mRNA.
Snx4	Snx4.bSep08	360725	10271	1561	5	172	sorting nexin 4 (Snx4) alternative variant bSep08, mRNA.
Snx4	Snx4.cSep08	360725	14565	1048	7	163	sorting nexin 4 (Snx4) alternative variant cSep08, mRNA.
Snx5	Snx5.aSep08	296199	19297	2036	14	404	sorting nexin 5 (46.8 kD) (Snx5) alternative variant aSep08, mRNA.
Snx5	Snx5.bSep08	296199	14657	1084	8	264	sorting nexin 5 CRA b (30.3 kD) (Snx5) alternative variant bSep08, mRNA.
Snx5	Snx5.cSep08	296199	14242	757	7	240	sorting nexin 5 CRA b (27.7 kD) (Snx5) alternative variant cSep08, mRNA.
Snx5	Snx5.eSep08	296199	1921	702	2	91	sorting nexin 5 (Snx5) alternative variant eSep08, mRNA.
Snx5	Snx5.gSep08	296199	691	597	2	55	putative protein (6.5 kD) (Snx5) alternative variant gSep08, mRNA.
Snx6	Snx6.bSep08	362738	4509	727	2	9	sorting nexin 6 (1.1 kD) (Snx6) alternative variant bSep08, complete mRNA.
Snx10	Snx10.bSep08	297096	45561	2548	7	201	sorting nexin 10 (23.6 kD) (Snx10) alternative variant bSep08, mRNA.
Snx10	Snx10.cSep08	297096	8642	561	3	117	sorting nexin 10 (13.2 kD) (Snx10) alternative variant cSep08, mRNA.
Snx10	Snx10.dSep08	297096	19191	221	3	20	sorting nexin 10 (Snx10) alternative variant dSep08, mRNA.
Snx11	Snx11.aSep08	303493	8651	725	6	220	sorting nexin 11 (Snx11) alternative variant aSep08, mRNA.
Snx11	Snx11.bSep08	303493	6756	692	6	133	sorting nexin 11 (Snx11) alternative variant bSep08, mRNA.
Snx11	Snx11.dSep08	303493	3199	407	1	45	sorting nexin 11 (Snx11) alternative variant dSep08, mRNA.
Snx12	Snx12.aSep08	363478	126524	1823	4	448	sorting nexin 12 (Snx12) alternative variant aSep08, mRNA.
Snx12	Snx12.cSep08	363478	3505	671	2	101	sorting nexin 12 (Snx12) alternative variant cSep08, mRNA.
Snx12	Snx12.dSep08	363478	7730	705	2	58	sorting nexin 12 (Snx12) alternative variant dSep08, mRNA.
Snx13	Snx13.bSep08	362731	11399	611	4	147	sorting nexin 13 (Snx13) alternative variant bSep08, mRNA.
Snx14	Snx14.bSep08	315871	64848	1928	18	642	sorting nexin 14 (Snx14) alternative variant bSep08, mRNA.
Snx14	Snx14.cSep08	315871	4539	661	8	219	sorting nexin 14 (Snx14) alternative variant cSep08, mRNA.

Snx14	Snx14.dSep08	315871	18065	472	5	157	sorting nexin 14 (Snx14) alternative variant dSep08, mRNA.
Snx14	Snx14.eSep08	315871	10163	759	3	119	sorting nexin 14 (13.4 kD) (Snx14) alternative variant eSep08, mRNA.
Snx14	Snx14.fSep08	315871	3763	579	2	21	sorting nexin 14 (2.4 kD) (Snx14) alternative variant fSep08, mRNA.
Snx15	Snx15.bSep08	293691	2491	839	4	182	sorting nexin 15 (20.4 kD) (Snx15) alternative variant bSep08, mRNA.
Snx15	Snx15.cSep08	293691	1124	755	3	107	sorting nexin 15 (Snx15) alternative variant cSep08, mRNA.
Snx16	Snx16.aSep08	64088	20537	1787	1	499	sorting nexin 16 (Snx16) alternative variant aSep08, mRNA.
Snx17	Snx17.bSep08	298836	1936	1159	6	212	sorting nexin 17 (Snx17) alternative variant bSep08, mRNA.
Snx17	Snx17.cSep08	298836	2055	515	4	150	sorting nexin 17 (Snx17) alternative variant cSep08, mRNA.
Snx20	Snx20.aSep08	307742	6130	1414	4	313	sorting nexin 20 (35.7 kD) (Snx20) alternative variant aSep08, mRNA.
Snx22	Snx22.bSep08	300796	1198	415		73	sorting nexin 22 (Snx22) alternative variant bSep08, mRNA.
Snx24	Snx24.aSep08	361328	157988	2147	7	103	sorting nexin 24 (12.5 kD) (Snx24) alternative variant aSep08, mRNA.
Snx25	Snx25.aSep08	306471	103010	2885	3	635	sorting nexin 25 (73.0 kD) (Snx25) alternative variant aSep08, mRNA.
Snx27	Snx27.bSep08	260323	19406	1865	1	176	sorting nexin family member 27 (Snx27) alternative variant bSep08, mRNA.
Snx30	Snx30.bSep08	298033	4816	501		40	sorting nexin family member 30 (Snx30) alternative variant bSep08, mRNA.
Snx32	Snx32.aSep08	361708	17452	1630	13	285	sorting nexin 32 (33.2 kD) (Snx32) alternative variant aSep08, complete mRNA.
Snx32	Snx32.bSep08	361708	15367	807	7	233	sorting nexin 32 (Snx32) alternative variant bSep08, mRNA.
Snx32	Snx32.cSep08	361708	15313	729	7	216	sorting nexin 32 (Snx32) alternative variant cSep08, mRNA.
Snx32	Snx32.dSep08	361708	1402	500	5	133	sorting nexin 32 (Snx32) alternative variant dSep08, mRNA.
snybor	snybor.aSep08		879	292		27	putative protein (snybor) mRNA.
snycha	snycha.aSep08		611	408		49	putative protein (5.5 kD) (snycha) complete mRNA.
snychy	snychy.aSep08		2917	556	5	85	CRA a like (9.5 kD) (snychy) alternative variant aSep08, mRNA.
snychy	snychy.bSep08		2588	487	5	85	CRA a like (9.5 kD) (snychy) alternative variant bSep08, mRNA.
snychy	snychy.cSep08		1008	753	2	69	CRA a like (7.7 kD) (snychy) alternative variant cSep08, mRNA.
snydoy	snydoy.aSep08		1217	285		94	putative protein of vertebrate origin (snydoy) mRNA.
snyfee	snyfee.aSep08		951	496		80	putative protein (snyfee) mRNA.

snyflu	snyflu.aSep08		7345	484		41	putative protein (snyflu) mRNA.
snyfly	snyfly.aSep08		579	404		134	zinc finger CCCH-type containing 4 like (snyfly) mRNA.
snyja	snyja.aSep08		1402	547		33	putative protein (3.8 kD) (snyja) mRNA.
snylor	snylor.aSep08		2812	245		65	putative protein (snylor) mRNA.
snymee	snymee.aSep08		8361	418		111	BRCA1 interacting protein C-terminal helicase 1 like (snymee) mRNA.
snynor	snynor.aSep08		7890	610		46	putative protein (snynor) mRNA.
snypey	snypey.aSep08		6367	440		20	putative protein (snypey) mRNA.
snyroy	snyroy.aSep08		10980	482		160	centrosomal protein 290kDa CRA a (snyroy) mRNA.
snyshaw	snyshaw.aSep08		2189	386		19	putative protein (snyshaw) mRNA.
snyshee	snyshee.aSep08		652	608		26	putative protein (snyshee) mRNA.
snytu	snytu.aSep08		373	308		68	putative protein (snytu) mRNA.
snyvo	snyvo.aSep08		10515	577		100	putative protein of eukaryotic origin (snyvo) mRNA.
snywer	snywer.aSep08		1196	565		187	eph receptor B6 (snywer) mRNA.
soby	soby.aSep08		8542	397		72	WD repeat (soby) mRNA.
sochy	sochy.aSep08		11274	699		204	kinesin 18A (sochy) alternative variant aSep08, mRNA.
sochy	sochy.bSep08		2859	499		134	kinesin family member 18B (sochy) alternative variant bSep08, mRNA.
Socs5	Socs5.bSep08	500616	5098	660	2	40	suppressor of cytokine signaling 5 and hypothetical protein LOC681150 (Socs5) alternative variant bSep08, mRNA.
Socs5	Socs5.bSep08	681150	5098	660	2	40	suppressor of cytokine signaling 5 and hypothetical protein LOC681150 (Socs5) alternative variant bSep08, mRNA.
Socs7	Socs7.aSep08	287659	26023	989		126	suppressor of cytokine signaling 7 (Socs7) mRNA.
SOCS_box.1	SOCS_box.1.aSep08	360206	4983	964		293	ankyrin and SOCS protein, C-terminal (SOCS_box.1) mRNA.
Sod2	Sod2.bSep08	24787	5867	598	5	185	superoxide dismutase 2, mitochondrial (Sod2) alternative variant bSep08, mRNA.
sodar	sodar.aSep08		35408	446		46	putative protein (5.1 kD) (sodar) mRNA.
sofer	sofer.aSep08		37988	360	2	120	type II (sofer) alternative variant aSep08, mRNA.
sofer	sofer.bSep08		19696	264	1	47	type II (sofer) alternative variant bSep08, mRNA.
soflo	soflo.aSep08		17350	762		93	putative nuclear protein (10.2 kD) (soflo) mRNA.
soflu	soflu.aSep08		1032	678		61	putative protein (soflu) mRNA.
sogar	sogar.aSep08		1566	206		68	SH2 domain-containing protein (sogar) mRNA.
Sohlh1	Sohlh1.aSep08	362085	1535	418		139	spermatogenesis and oogenesis specific basic helix-loop-helix 1 (Sohlh1) mRNA.
soja	soja.aSep08		88936	737	2	46	putative protein (5.5 kD) (soja) alternative variant aSep08, mRNA.
soja	soja.bSep08		76245	726	2	31	putative protein (3.6 kD) (soja) alternative variant bSep08, mRNA.
sojey	sojey.aSep08		4788	835		77	aasdh protein (8.4 kD) (sojey) mRNA.
sokee	sokee.aSep08		5537	865		288	putative protein of vertebrate origin (sokee) mRNA.
sokler	sokler.aSep08		1977	308		102	signal transducing adaptor family member 2 (sokler) mRNA.

Solh	Solh.bSep08	303000	623	537	2	178	small optic lobes homolog (Drosophila) (Solh) alternative variant bSep08, mRNA.
solo	solo.aSep08		15496	946		314	furry homolog (solo) mRNA.
somee	somee.aSep08		1123	1073		47	putative protein (4.8 kD) (somee) mRNA.
somer	somer.aSep08		514	360		120	CRA c like (somer) mRNA.
Son	Son.aSep08	304092	6086	2280	5	576	SON DNA binding protein CRA f like (Son) alternative variant aSep08, mRNA.
Son	Son.bSep08	304092	19579	2584	10	560	SON DNA binding protein like (Son) alternative variant bSep08, mRNA.
Son	Son.cSep08	304092	1802	1515	3	277	SON protein (Son) alternative variant cSep08, mRNA.
Son	Son.dSep08	304092	16652	1261	10	252	SON protein (Son) alternative variant dSep08, mRNA.
Son	Son.eSep08	304092	3872	2118	3	202	SON protein (Son) alternative variant eSep08, mRNA.
Son	Son.fSep08	304092	4833	2411	4	152	SON DNA binding protein like (16.8 kD) (Son) alternative variant fSep08, mRNA.
Son	Son.gSep08	304092	6516	1169	7	152	SON DNA binding protein like (16.8 kD) (Son) alternative variant gSep08, mRNA.
Son	Son.hSep08	304092	3427	1058	4	119	SON protein (13.7 kD) (Son) alternative variant hSep08, mRNA.
Son	Son.iSep08	304092	4517	1523	5	119	SON protein (13.7 kD) (Son) alternative variant iSep08, mRNA.
sonoy	sonoy.aSep08		3931	346		68	putative protein (sonoy) mRNA.
sopor	sopor.aSep08		4506	581	2	96	putative protein (sopor) alternative variant aSep08, mRNA.
Sorbs1	Sorbs1.aSep08	686098	148646	5521	25	788	sorbin-like and src homology-3 and variant SH3 (88.6 kD) (Sorbs1) alternative variant aSep08, mRNA.
Sorbs1	Sorbs1.bSep08	686098	184628	1342	16	385	sorbin-like (Sorbs1) alternative variant bSep08, mRNA.
Sorbs1	Sorbs1.cSep08	686098	35451	792	9	264	putative protein, with a coiled coil domain, of vertebrate origin (Sorbs1) alternative variant cSep08, mRNA.
Sorbs1	Sorbs1.dSep08	686098	22811	1015	6	196	src homology-3 and variant SH3 (Sorbs1) alternative variant dSep08, mRNA.
Sorbs1	Sorbs1.eSep08	686098	19264	465	5	155	putative protein of vertebrate origin (Sorbs1) alternative variant eSep08, mRNA.
Sorbs1	Sorbs1.fSep08	686098	5942	837	3	136	src homology-3 and variant SH3 (Sorbs1) alternative variant fSep08, mRNA.
Sorbs1	Sorbs1.gSep08	686098	13236	375	4	124	putative protein of vertebrate origin (Sorbs1) alternative variant gSep08, mRNA.
Sorbs1	Sorbs1.hSep08	686098	558	423	2	101	putative protein of metazoan origin (Sorbs1) alternative variant hSep08, mRNA.
Sorbs1	Sorbs1.iSep08	686098	11980	296	4	98	putative protein of vertebrate origin (Sorbs1) alternative variant iSep08, mRNA.
Sorbs1	Sorbs1.jSep08	686098	3621	294	2	97	putative protein of vertebrate origin (Sorbs1) alternative variant jSep08, mRNA.
Sorbs1	Sorbs1.kSep08	686098	7223	257	4	85	putative protein of vertebrate origin (Sorbs1) alternative variant kSep08, mRNA.
Sorbs1	Sorbs1.lSep08	686098	8904	345	5	65	putative protein of vertebrate origin (Sorbs1) alternative variant lSep08, mRNA.

Sorbs3	Sorbs3.bSep08	282843	13019	2851	11	562	src homology-3 and variant SH3 (Sorbs3) alternative variant bSep08, mRNA.
Sorbs3	Sorbs3.cSep08	282843	4992	830	9	217	putative nuclear protein of mammalian origin (24.2 kD) (Sorbs3) alternative variant cSep08, mRNA.
Sorbs3	Sorbs3.dSep08	282843	11771	580	6	193	putative protein of mammalian origin (Sorbs3) alternative variant dSep08, mRNA.
Sorbs3	Sorbs3.eSep08	282843	7447	436	4	145	sorbin-like (Sorbs3) alternative variant eSep08, mRNA.
Sorbs3	Sorbs3.fSep08	282843	3141	1119	2	133	putative protein (Sorbs3) alternative variant fSep08, mRNA.
Sorbs3	Sorbs3.gSep08	282843	2870	629	3	128	src homology-3 and variant SH3 (Sorbs3) alternative variant gSep08, mRNA.
Sorbs3	Sorbs3.hSep08	282843	1993	325	3	108	putative protein of mammalian origin (Sorbs3) alternative variant hSep08, mRNA.
Sorbs3	Sorbs3.iSep08	282843	814	389	4	52	putative protein of mammalian origin (Sorbs3) alternative variant iSep08, mRNA.
Sorbs3	Sorbs3.jSep08	282843	766	390	3	35	putative protein of mammalian origin (Sorbs3) alternative variant jSep08, mRNA.
sorby	sorby.aSep08		4075	227		62	putative protein (7.0 kD) (sorby) mRNA.
sorchy	sorchy.aSep08		1669	268		17	putative protein (2.2 kD) (sorchy) mRNA.
Sorcs2	Sorcs2.bSep08	305438	6655	702	5	182	vps10 domain receptor protein sorcs 2 (Sorcs2) alternative variant bSep08, mRNA.
Sorcs2	Sorcs2.cSep08	305438	5752	696	4	105	vps10 domain receptor protein sorcs 2 (Sorcs2) alternative variant cSep08, mRNA.
Sorcs2	Sorcs2.dSep08	305438	2250	314	2	36	vps10 domain receptor protein sorcs 2 (Sorcs2) alternative variant dSep08, mRNA.
Sord	Sord.bSep08	24788	25593	802	2	221	sorbitol dehydrogenase (Sord) alternative variant bSep08, mRNA.
Sord	Sord.cSep08	24788	25408	724	2	164	sorbitol dehydrogenase (Sord) alternative variant cSep08, mRNA.
sordar	sordar.aSep08		1027	375		72	putative protein (8.1 kD) (sordar) mRNA.
sorflo	sorflo.aSep08		13987	642	4	139	CRA a like (sorflo) alternative variant aSep08, mRNA.
sorflo	sorflo.bSep08		1345	672	2	70	CRA a like (8.1 kD) (sorflo) alternative variant bSep08, mRNA.
sorflo	sorflo.cSep08		14897	1095	3	27	putative protein (3.1 kD) (sorflo) alternative variant cSep08, mRNA.
sorflo	sorflo.dSep08		14294	900	4	70	CRA a like (8.1 kD) (sorflo) alternative variant dSep08, mRNA.
sorflo	sorflo.eSep08		16611	440	3	44	putative protein (sorflo) alternative variant eSep08, mRNA.
sorflu	sorflu.aSep08		957	433		92	putative protein (sorflu) mRNA.
sorgar	sorgar.aSep08		1329	901	2	72	putative protein (7.9 kD) (sorgar) alternative variant aSep08, mRNA.
sorja	sorja.bSep08		4926	819	2	52	putative protein of vertebrate origin (sorja) alternative variant bSep08, mRNA.
sorjey	sorjey.aSep08		25483	831		172	putative protein of metazoan origin (20.2 kD) (sorjey) mRNA.
sorkee	sorkee.aSep08		2529	443		147	flavin containing monooxygenase 3 (sorkee) mRNA.

sorkler	sorkler.aSep08		3075	1375		71	putative cytoplasmic protein (8.3 kD) (sorkler) mRNA.
Sor1	Sor1.aSep08	300652	20241	1957	13	622	sortilin-related receptor, LDLR class A repeats-containing (Sor1) alternative variant aSep08, mRNA.
Sor1	Sor1.bSep08	300652	26991	2187	17	523	sortilin-related receptor, LDLR class A repeats-containing (Sor1) alternative variant bSep08, mRNA.
Sor1	Sor1.cSep08	300652	6353	1001	3	182	sortilin-related receptor, LDLR class A repeats-containing (Sor1) alternative variant cSep08, mRNA.
Sor1	Sor1.dSep08	300652	2792	806	1	137	sortilin-related receptor, LDLR class A repeats-containing (Sor1) alternative variant dSep08, mRNA.
sorlo	sorlo.aSep08		2689	520		48	putative protein (5.4 kD) (sorlo) mRNA.
sormee	sormee.aSep08		4650	295		48	putative protein (5.6 kD) (sormee) mRNA.
sornoy	sornoy.aSep08		21427	1117		372	dedicator of cytokinesis 10 (sornoy) mRNA.
sorpor	sorpor.aSep08		8440	759		251	myosin VC CRA b (sorpor) mRNA.
sorsa	sorsa.aSep08		1813	269		43	putative protein (sorsa) mRNA.
sorshee	sorshee.aSep08		7454	369		122	folliculin-interacting protein (sorshee) mRNA.
Sort1	Sort1.aSep08	83576	21629	2087	11	425	sortilin 1 (Sort1) alternative variant aSep08, mRNA.
Sort1	Sort1.bSep08	83576	6353	775	7	257	sortilin 1 (Sort1) alternative variant bSep08, mRNA.
Sort1	Sort1.cSep08	83576	6620	635	4	211	sortilin 1 (Sort1) alternative variant cSep08, mRNA.
sortu	sortu.aSep08		5798	478		79	putative protein (sortu) mRNA.
sorvar	sorvar.aSep08		12726	1398		416	ATPase type 13A2 (sorvar) mRNA.
sorwey	sorwey.aSep08		3429	352		33	LYST-interacting protein 8 like (sorwey) mRNA.
Sos1	Sos1.bSep08	313845	16286	681	6	110	son of sevenless homolog 1 (Drosophila) (Sos1) alternative variant bSep08, mRNA.
sosa	sosa.aSep08		550	291		60	putative protein (sosa) mRNA.
soshee	soshee.aSep08		2219	376		125	sucrase-isomaltase (soshee) mRNA.
Sostdc1	Sostdc1.bSep08	266803	3534	835	2	157	sclerostin (Sostdc1) alternative variant bSep08, mRNA.
sotu	sotu.aSep08		2839	210		63	putative protein (sotu) mRNA.
sovar	sovar.aSep08		6342	374		103	putative protein (sovar) mRNA.
sowey	sowey.aSep08		1598	344		114	murinoglobulin (sowey) mRNA.
Sox5	Sox5.aSep08	140587	134222	639		198	SRY-box containing gene 5 (Sox5) mRNA.
Sox15	Sox15.aSep08	363632	1674	1102	2	142	SRY-box containing gene 15 (Sox15) alternative variant aSep08, mRNA.
Sox17	Sox17.aSep08	312936	4998	2544	4	423	SRY-box containing gene 17 (45.0 kD) (Sox17) alternative variant aSep08, mRNA.
Sox17	Sox17.cSep08	312936	1143	706	2	172	SRY-box containing gene 17 (Sox17) alternative variant cSep08, mRNA.
Sox17	Sox17.dSep08	312936	1377	1212	2	42	SRY-box containing gene 17 (4.7 kD) (Sox17) alternative variant dSep08, mRNA.
Sox17	Sox17.eSep08	312936	519	350	2	32	SRY-box containing gene 17 (Sox17) alternative variant eSep08, mRNA.
Sox30	Sox30.aSep08	689918	14718	746		232	SRY (sex determining region Y)-box 30 (Sox30) mRNA.
soyby	soyby.aSep08		965	544		28	putative protein (3.3 kD) (soyby) mRNA.
soychy	soychy.aSep08		2821	2514		530	rho GTPase activating protein 11A (soychy) mRNA.



soydar	soydar.aSep08		2981	264		34	putative protein (3.8 kD) (soydar) mRNA.
soyflo	soyflo.aSep08		30277	968	4	239	acyl-CoA synthetase long-chain family member 5 (soyflo) alternative variant aSep08, mRNA.
soyflo	soyflo.bSep08		14741	751	4	193	acyl-CoA synthetase long-chain family member 5 (soyflo) alternative variant bSep08, mRNA.
soyflo	soyflo.cSep08		27474	1067	3	168	acyl-CoA synthetase long-chain family member 5 (soyflo) alternative variant cSep08, mRNA.
soyflo	soyflo.dSep08		13678	704	4	154	acyl-CoA synthetase long-chain family member 5 (soyflo) alternative variant dSep08, mRNA.
soyflo	soyflo.eSep08		25912	726	2	130	acyl-CoA synthetase long-chain family member 5 (soyflo) alternative variant eSep08, mRNA.
soyflu	soyflu.aSep08		1355	637		81	putative protein (9.0 kD) (soyflu) mRNA.
soygar	soygar.aSep08		1685	687		229	transmembrane protein 16h (soygar) mRNA.
soyja	soyja.aSep08		5453	317		34	putative protein (soyja) mRNA.
soyjoy	soyjoy.aSep08		62711	678		62	putative protein (7.2 kD) (soyjoy) mRNA.
soykee	soykee.aSep08		4124	882		49	putative protein (5.4 kD) (soykee) mRNA.
soykler	soykler.aSep08		10352	322		46	putative protein (soykler) mRNA.
soylo	soylo.aSep08		1644	329		82	putative protein (soylo) mRNA.
soymee	soymee.aSep08		872	276		91	transcription factor 7 (soymee) mRNA.
soynoy	soynoy.aSep08		2565	484		72	putative protein (soynoy) mRNA.
soypor	soypor.aSep08		1798	817		138	myosin VC CRA c (16.0 kD) (soypor) mRNA.
soysa	soysa.aSep08		4091	810		186	CAP-binding protein complex interacting 1 like (soysa) mRNA.
soyshee	soyshee.aSep08		4141	720		42	putative protein (4.5 kD) (soyshee) mRNA.
soytu	soytu.aSep08		23582	590	4	185	CRA b (soytu) alternative variant aSep08, mRNA.
soyvar	soyvar.aSep08		1398	697		119	putative protein (soyvar) mRNA.
soywey	soywey.aSep08		1602	725	3	126	CRA c (13.2 kD) (soywey) alternative variant aSep08, mRNA.
soywey	soywey.bSep08		1483	803	3	86	gene rich cluster C10 CRA b like (9.1 kD) (soywey) alternative variant bSep08, mRNA.
Sp1	Sp1.bSep08	24790	1353	1087	2	203	trans-acting transcription factor 1 (Sp1) alternative variant bSep08, mRNA.
Sp2	Sp2.aSep08	303499	24972	3385	8	623	sp2 transcription factor (Sp2) alternative variant aSep08, mRNA.
Sp3	Sp3.aSep08	367846	33263	1219		240	trans-acting transcription factor 3 (26.5 kD) (Sp3) mRNA.
Sp100	Sp100.aSep08	363269	35270	1428	14	447	nuclear antigen Sp100 (Sp100) alternative variant aSep08, mRNA.
Sp100	Sp100.bSep08	363269	42849	2649	15	410	nuclear antigen Sp100 (Sp100) alternative variant bSep08, mRNA.
Sp100	Sp100.cSep08	363269	23607	958	8	188	nuclear antigen Sp100 (Sp100) alternative variant cSep08, mRNA.
Sp100	Sp100.dSep08	363269	13226	863	5	166	nuclear antigen Sp100 (Sp100) alternative variant dSep08, mRNA.
Sp100	Sp100.eSep08	363269	9918	560	6	156	nuclear antigen Sp100 (Sp100) alternative variant eSep08, mRNA.

Sp100	Sp100.fSep08	363269	4781	641	3	131	nuclear antigen Sp100 (Sp100) alternative variant fSep08, mRNA.
Sp100	Sp100.gSep08	363269	12230	661	5	123	nuclear antigen Sp100 (Sp100) alternative variant gSep08, mRNA.
Sp100	Sp100.hSep08	363269	3068	348	2	61	nuclear antigen Sp100 (Sp100) alternative variant hSep08, mRNA.
Sp110	Sp110.bSep08	301570	19582	1353	4	395	SP110 nuclear body protein (Sp110) alternative variant bSep08, mRNA.
Sp110	Sp110.cSep08	301570	13613	820	1	273	SP110 nuclear body protein (Sp110) alternative variant cSep08, mRNA.
Spa17	Spa17.bSep08	85244	10056	1428	1	132	sperm autoantigenic protein 17 (Spa17) alternative variant bSep08, mRNA.
spabor	spabor.bSep08		5133	709		47	putative protein (5.0 kD) (spabor) alternative variant bSep08, mRNA.
Spaca1	Spaca1.aSep08	500432	5021	414		110	sperm acrosome associated 1 (Spaca1) mRNA.
Spaca3	Spaca3.aSep08	287557	7476	756	1	163	sperm acrosome associated 3 (18.3 kD) (Spaca3) alternative variant aSep08, mRNA.
spacha	spacha.aSep08		4427	350		116	senataxin (spacha) mRNA.
spachy	spachy.aSep08		12187	495		61	putative protein (spachy) mRNA.
spafee	spafee.aSep08		1727	267		88	putative protein (spafee) mRNA.
spafly	spafly.aSep08		1279	284		57	putative protein (spafly) mRNA.
spafly	spafly.aSep08		160283	497		165	protein phosphatase 1 regulatory like (spafly) mRNA.
Spag4	Spag4.bSep08	83623	1109	878	2	108	sperm associated antigen 4 (Spag4) alternative variant bSep08, mRNA.
Spag4	Spag4.cSep08	83623	804	702	2	36	sperm associated antigen 4 (4.3 kD) (Spag4) alternative variant cSep08, mRNA.
Spag5	Spag5.bSep08	252918	6788	1302	2	227	sperm associated antigen 5 (Spag5) alternative variant bSep08, mRNA.
Spag8	Spag8.aSep08	362508	2574	1909		467	sperm associated antigen 8 (51.0 kD) (Spag8) mRNA.
Spag9	Spag9.aSep08	360600	64892	3282	25	1038	sperm associated antigen 9 (Spag9) alternative variant aSep08, mRNA.
Spag9	Spag9.cSep08	360600	20869	2951	9	408	sperm associated antigen 9 (Spag9) alternative variant cSep08, mRNA.
Spag9	Spag9.dSep08	360600	37761	610	6	203	sperm associated antigen 9 (Spag9) alternative variant dSep08, mRNA.
Spag9	Spag9.eSep08	360600	13321	3798	5	197	sperm associated antigen 9 (Spag9) alternative variant eSep08, mRNA.
Spag9	Spag9.fSep08	360600	36859	437	4	145	sperm associated antigen 9 (Spag9) alternative variant fSep08, mRNA.
Spag16	Spag16.aSep08	501158	133421	743	5	206	sperm associated antigen 16 (Spag16) alternative variant aSep08, mRNA.
spalor	spalor.aSep08		31328	326		50	putative protein (spalor) mRNA.
spamee	spamee.aSep08		4885	676	3	52	putative protein (5.6 kD) (spamee) alternative variant aSep08, mRNA.
spamee	spamee.bSep08		2678	369	1	37	putative protein (spamee) alternative variant bSep08, mRNA.

spanor	spanor.aSep08		16303	2103	13	432	ubiquitin ligase E3 alpha-II (spanor) alternative variant aSep08, mRNA.
spapey	spapey.aSep08		8638	521		90	putative protein (10.2 kD) (spapey) mRNA.
sparbor	sparbor.aSep08		15358	291		97	cell division cycle 40 homolog (sparbor) mRNA.
Sparc	Sparc.aSep08	24791	20070	963	9	320	secreted acidic cysteine rich glycoprotein (Sparc) alternative variant aSep08, mRNA.
Sparc	Sparc.cSep08	24791	20024	899	9	260	secreted acidic cysteine rich glycoprotein (Sparc) alternative variant cSep08, mRNA.
sparcha	sparcha.aSep08		4686	337		71	putative protein (sparcha) mRNA.
sparchy	sparchy.aSep08		1275	323		70	putative protein (sparchy) mRNA.
Sparcl1	Sparcl1.bSep08	25434	18292	760	1	253	SPARC-like 1 (mast9, hevin) (Sparcl1) alternative variant bSep08, mRNA.
sparfee	sparfee.aSep08		35242	384		63	putative protein (sparfee) mRNA.
sparflu	sparflu.aSep08		551	255		36	putative protein (sparflu) mRNA.
sparfly	sparfly.aSep08		348	255		36	putative protein (sparfly) mRNA.
sparlor	sparlor.aSep08		1183	1053	2	57	putative protein (6.4 kD) (sparlor) alternative variant aSep08, mRNA.
sparmee	sparmee.aSep08		1755	272		90	tripartite motif protein 25 CRA a (sparmee) mRNA.
sparnor	sparnor.aSep08		1613	491		163	parkin-like cytoplasmic protein (sparnor) mRNA.
sparoy	sparoy.aSep08		6044	517		172	neuron navigator 3 CRA b (sparoy) mRNA.
sparpey	sparpey.aSep08		687	200		33	putative protein (sparpey) mRNA.
sparroy	sparroy.aSep08		3645	296		32	putative protein (sparroy) mRNA.
sparshaw	sparshaw.aSep08		3022	797		50	putative protein (5.6 kD) (sparshaw) mRNA.
spartu	spartu.aSep08		2652	767		117	thyroid hormone receptor interactor 11 (spartu) mRNA.
sparvo	sparvo.aSep08		757	388		27	putative protein (sparvo) mRNA.
sparwer	sparwer.aSep08		1298	1245		96	putative protein (sparwer) mRNA.
spashaw	spashaw.aSep08		5815	3156		74	putative protein, with a coiled coil domain (8.6 kD) (spashaw) mRNA.
Spast	Spast.aSep08	362700	58690	4489	15	362	spastin (40.4 kD) (Spast) alternative variant aSep08, mRNA.
Spast	Spast.cSep08	362700	2348	390	1	28	spastin (3.3 kD) (Spast) alternative variant cSep08, mRNA.
Spata1	Spata1.bSep08	362056	12716	682	5	182	spermatogenesis associated 1 (20.5 kD) (Spata1) alternative variant bSep08, mRNA.
Spata1	Spata1.cSep08	362056	10219	670	4	161	spermatogenesis associated 1 (Spata1) alternative variant cSep08, mRNA.
Spata1	Spata1.dSep08	362056	18944	701	6	129	spermatogenesis associated 1 CRA a (Spata1) alternative variant dSep08, mRNA.
Spata1	Spata1.eSep08	362056	1509	924	2	126	spermatogenesis associated 1 (14.8 kD) (Spata1) alternative variant eSep08, mRNA.
Spata2	Spata2.bSep08	114210	10272	3928	3	511	spermatogenesis associated 2 (57.5 kD) (Spata2) alternative variant bSep08, mRNA.
Spata2L	Spata2L.bSep08	498963	4727	3412	1	109	spermatogenesis associated 2-like (12.3 kD) (Spata2L) alternative variant bSep08, mRNA.
Spata3	Spata3.aSep08	363270	4814	892	2	241	spermatogenesis associated 3 (Spata3) alternative variant aSep08, mRNA.

Spata5	Spata5.bSep08	361935	172135	1785	1	538	spermatogenesis associated 5 (Spata5) alternative variant bSep08, mRNA.
Spata5	Spata5.cSep08	361935	140698	717	1	182	spermatogenesis associated 5 (Spata5) alternative variant cSep08, mRNA.
Spata6	Spata6.bSep08	171413	21279	580	1	108	spermatogenesis associated 6 (Spata6) alternative variant bSep08, mRNA.
Spata7	Spata7.aSep08	192225	45592	2008	12	621	spermatogenesis associated 7 (Spata7) alternative variant aSep08, mRNA.
Spata7	Spata7.bSep08	192225	35085	747	6	249	spermatogenesis associated 7 (Spata7) alternative variant bSep08, mRNA.
Spata7	Spata7.cSep08	192225	36681	742	5	235	spermatogenesis associated 7 (Spata7) alternative variant cSep08, mRNA.
Spata7	Spata7.dSep08	192225	35157	727	5	225	spermatogenesis associated 7 (Spata7) alternative variant dSep08, mRNA.
Spata7	Spata7.eSep08	192225	26406	510	6	138	spermatogenesis associated 7 (Spata7) alternative variant eSep08, mRNA.
Spata7	Spata7.fSep08	192225	34925	734	7	129	spermatogenesis associated 7 (14.6 kD) (Spata7) alternative variant fSep08, complete mRNA.
Spata7	Spata7.gSep08	192225	39821	1048	7	72	spermatogenesis associated 7 (8.3 kD) (Spata7) alternative variant gSep08, mRNA.
Spata9	Spata9.bSep08	294594	21895	596	1	136	spermatogenesis associated 9 (Spata9) alternative variant bSep08, mRNA.
Spata13	Spata13.aSep08	305938	17085	748		223	spermatogenesis associated 13 (Spata13) mRNA.
Spata16	Spata16.aSep08	294932	41641	300		70	spermatogenesis associated 16 (Spata16) mRNA.
Spata17	Spata17.aSep08	498305	120099	728	7	242	spermatogenesis associated 17 (Spata17) alternative variant aSep08, mRNA.
Spata17	Spata17.bSep08	498305	67450	738	4	157	spermatogenesis associated 17 (Spata17) alternative variant bSep08, mRNA.
Spata18	Spata18.bSep08	289586	26684	1788	7	499	spermatogenesis associated 18 (Spata18) alternative variant bSep08, mRNA.
Spata18	Spata18.cSep08	289586	11335	773	6	197	spermatogenesis associated 18 (Spata18) alternative variant cSep08, mRNA.
Spata18	Spata18.dSep08	289586	9966	662	6	160	spermatogenesis associated 18 (Spata18) alternative variant dSep08, mRNA.
Spata18	Spata18.fSep08	289586	14814	716	7	116	spermatogenesis associated 18 (Spata18) alternative variant fSep08, mRNA.
Spata18	Spata18.gSep08	289586	6774	770	3	87	spermatogenesis associated 18 (Spata18) alternative variant gSep08, mRNA.
Spata20	Spata20.bSep08	360604	1186	755	1	190	spermatogenesis associated 20 (Spata20) alternative variant bSep08, mRNA.
Spata21	Spata21.bSep08	366491	9019	801	1	171	spermatogenesis associated 21 (Spata21) alternative variant bSep08, mRNA.
Spats1	Spats1.bSep08	301255	7539	652	4	121	spermatogenesis associated, serine-rich 1 (Spats1) alternative variant bSep08, mRNA.
Spats2	Spats2.aSep08	300221	73601	3067	6	561	spermatogenesis associated, serine-rich 2 (60.5 kD) (Spats2) alternative variant aSep08, mRNA.
spatu	spatu.aSep08		3997	384		98	putative protein (spatu) mRNA.

spavo	spavo.aSep08		5822	708		45	putative protein (5.1 kD) (spavo) mRNA.
spawcha	spawcha.aSep08		11825	411		97	putative protein (spawcha) mRNA.
spawchy	spawchy.aSep08		4043	495		99	putative protein (11.0 kD) (spawchy) mRNA.
spawer	spawer.aSep08		4993	683		41	CRA a like (4.5 kD) (spawer) mRNA.
spawfee	spawfee.bSep08		2575	350		116	PHD finger protein 2 (spawfee) alternative variant bSep08, mRNA.
spawflu	spawflu.aSep08		1885	216		47	putative protein (spawflu) mRNA.
spawfly	spawfly.aSep08		2699	486		138	putative protein (spawfly) mRNA.
spawlor	spawlor.aSep08		1470	809		64	putative protein (spawlor) mRNA.
spawmee	spawmee.aSep08		10896	809		52	putative protein (5.9 kD) (spawmee) mRNA.
spawnor	spawnor.aSep08		1258	541		157	cullin 7 (spawnor) mRNA.
spawpey	spawpey.aSep08		3575	505		67	putative protein (7.3 kD) (spawpey) mRNA.
spawroy	spawroy.aSep08		18581	1592		125	putative mitochondrial protein (14.0 kD) (spawroy) mRNA.
spawshaw	spawshaw.aSep08		38358	479		53	putative protein (spawshaw) mRNA.
spawtu	spawtu.aSep08		11450	212		70	thyroid hormone receptor interactor 11 (spawtu) mRNA.
spawvo	spawvo.aSep08		12094	686		228	putative protein of metazoan origin (spawvo) mRNA.
spawwer	spawwer.aSep08		883	383		84	putative protein (10.1 kD) (spawwer) mRNA.
SPC22.0	SPC22.0.aSep08		2187	1941		252	putative protein (SPC22.0) mRNA.
Spc24	Spc24.aSep08	363028	4783	701	4	167	SPC24, NDC80 kinetochore complex component, homolog ( <i>S. cerevisiae</i> ) (Spc24) alternative variant aSep08, mRNA.
Spc24	Spc24.cSep08	363028	1545	482	3	54	SPC24, NDC80 kinetochore complex component, homolog ( <i>S. cerevisiae</i> ) (Spc24) alternative variant cSep08, mRNA.
Spc25	Spc25.bSep08	295661	12664	683	1	213	SPC25, NDC80 kinetochore complex component, homolog ( <i>S. cerevisiae</i> ) (Spc25) alternative variant bSep08, mRNA.
Spcs1	Spcs1.bSep08	290555	1383	515	4	82	signal peptidase complex subunit 1 homolog ( <i>S. cerevisiae</i> ) (Spcs1) alternative variant bSep08, mRNA.
Spcs2	Spcs2.aSep08	293142	19435	2965	5	226	signal peptidase complex subunit 2 homolog ( <i>S. cerevisiae</i> ) (25.0 kD) (Spcs2) alternative variant aSep08, complete mRNA.
Spdef	Spdef.bSep08	689210	14245	1801	1	325	sterile alpha motif/pointed and ets (36.3 kD) (Spdef) alternative variant bSep08, complete mRNA.
Spdya	Spdya.bSep08	192209	40746	1794	6	311	speedy homolog A ( <i>Drosophila</i> ) (36.1 kD) (Spdya) alternative variant bSep08, complete mRNA.
Spdya	Spdya.cSep08	192209	39829	706	6	235	speedy homolog A ( <i>Drosophila</i> ) (Spdya) alternative variant cSep08, mRNA.
Spdya	Spdya.dSep08	192209	44095	761	5	96	speedy homolog A ( <i>Drosophila</i> ) (11.7 kD) (Spdya) alternative variant dSep08, mRNA.
Specc1	Specc1.bSep08	303208	115733	545	4	161	sperm antigen with calponin homology and coiled-coil domains 1 (Specc1) alternative variant bSep08, mRNA.
Specc1l	Specc1l.cSep08	361828	6068	343	3	114	SPECC1-like (Specc1l) alternative variant cSep08, mRNA.
Specc1l	Specc1l.dSep08	361828	8581	2914	5	109	SPECC1-like (12.4 kD) (Specc1l) alternative variant dSep08, mRNA.
Spectrin.0	Spectrin.0.aSep08		48765	1468		489	dystrophin Dp427l (Spectrin.0) mRNA.
Spectrin.1	Spectrin.1.aSep08		61604	619		206	dystrophin CRA c (Spectrin.1) mRNA.

Spectrin.2	Spectrin.2.aSep08		50419	1020	5	339	spectrin repeat containing nuclear envelope (Spectrin.2) alternative variant aSep08, mRNA.
Spectrin.3	Spectrin.3.aSep08		3101	857		285	microtubule-actin crosslinking factor 1 (Spectrin.3) mRNA.
Spectrin.4	Spectrin.4.aSep08		28522	914		304	microtubule-actin crosslinking factor 1 (Spectrin.4) mRNA.
Spectrin.6	Spectrin.6.aSep08		12153	1146	7	333	utrophin (Spectrin.6) alternative variant aSep08, mRNA.
Spectrin.7	Spectrin.7.aSep08		3974	952		317	spectrin beta 4 CRA b (Spectrin.7) mRNA.
speecha	speecha.aSep08		7628	410	2	136	tbc1 domain family member 13 (speecha) alternative variant aSep08, mRNA.
speecha	speecha.bSep08		6948	583	1	40	tbc1 domain family member 13 (speecha) alternative variant bSep08, mRNA.
speechy	speechy.aSep08		7701	279		48	putative protein (speechy) mRNA.
speefee	speefee.aSep08		3719	2417		190	PHD finger protein 2 (speefee) mRNA.
speeflu	speeflu.aSep08		59403	671			
speefly	speefly.aSep08		3753	1328		60	putative protein of mammalian origin (6.5 kD) (speefly) mRNA.
speelor	speelor.aSep08		7964	1784	3	471	T-cell lymphoma invasion metastasis 1 like (speelor) alternative variant aSep08, mRNA.
speelor	speelor.bSep08		10212	1311	4	285	T-cell lymphoma invasion metastasis 1 like (speelor) alternative variant bSep08, mRNA.
speemee	speemee.bSep08		730	593	2	73	putative protein (speemee) alternative variant bSep08, mRNA.
speenor	speenor.aSep08		620	367		122	putative protein of vertebrate origin (speenor) mRNA.
speepey	speepey.aSep08		11134	347		75	putative protein of mammalian origin (speepey) mRNA.
speeroy	speeroy.aSep08		1628	709		72	putative protein (7.7 kD) (speeroy) mRNA.
speeshaw	speeshaw.aSep08		520	442		147	mesoderm induction early response 1 family member 3 CRA c (speeshaw) mRNA.
speetu	speetu.aSep08		2481	346		104	putative protein (speetu) mRNA.
speevo	speevo.aSep08		8071	359		119	CRA a like (speevo) mRNA.
speewer	speewer.aSep08		2168	251		31	putative protein (3.7 kD) (speewer) mRNA.
Spef2	Spef2.bSep08	64555	37178	1526	9	494	sperm flagellar 2 (Spef2) alternative variant bSep08, mRNA.
Spef2	Spef2.cSep08	64555	18128	396	3	131	sperm flagellar 2 (Spef2) alternative variant cSep08, mRNA.
Speg	Speg.cSep08	363256	13078	1350	5	187	SPEG complex locus (Speg) alternative variant cSep08, mRNA.
Speg	Speg.dSep08	363256	1371	538	4	167	SPEG complex locus (Speg) alternative variant dSep08, mRNA.
Speg	Speg.eSep08	363256	3970	817	5	113	SPEG complex locus (12.7 kD) (Speg) alternative variant eSep08, mRNA.
Speg	Speg.fSep08	363256	1884	333	4	111	SPEG complex locus (Speg) alternative variant fSep08, mRNA.
Speg	Speg.gSep08	363256	876	369	3	82	SPEG complex locus (Speg) alternative variant gSep08, mRNA.
Speg	Speg.hSep08	363256	1117	392	2	82	SPEG complex locus (7.0 kD) (Speg) alternative variant hSep08, mRNA.

Speg	Speg.iSep08	363256	587	450	2	36	SPEG complex locus (4.1 kD) (Speg) alternative variant iSep08, mRNA.
spercha	spercha.aSep08		21818	825		273	nucleoporin (spercha) mRNA.
sperchy	sperchy.aSep08		1758	499		36	putative protein (3.7 kD) (sperchy) mRNA.
sperfee	sperfee.aSep08		3634	621		125	putative protein (13.8 kD) (sperfee) mRNA.
sperflu	sperflu.aSep08		2490	783		51	putative protein (sperflu) mRNA.
sperfly	sperfly.aSep08		5140	285		40	putative protein (4.3 kD) (sperfly) mRNA.
sperlor	sperlor.aSep08		2690	1624	2	27	putative protein (9.1 kD) (sperlor) alternative variant aSep08, mRNA.
sperlor	sperlor.cSep08		2684	779	3	27	putative protein (9.1 kD) (sperlor) alternative variant cSep08, mRNA.
spermee	spermee.aSep08		24757	824		80	putative protein (spermee) mRNA.
sperpey	sperpey.aSep08		2110	490	3	73	putative protein of mammalian origin (sperpey) alternative variant aSep08, mRNA.
sperpey	sperpey.bSep08		800	353	2	42	putative protein (sperpey) alternative variant bSep08, mRNA.
sperroy	sperroy.aSep08		2396	708		46	putative protein (sperroy) mRNA.
spershaw	spershaw.aSep08		506	354	2	60	putative protein (7.0 kD) (spershaw) alternative variant aSep08, mRNA.
Spert	Spert.bSep08	498572	8794	2206	3	372	spermatid associated (43.1 kD) (Spert) alternative variant bSep08, mRNA.
Spert	Spert.cSep08	498572	6442	742	2	247	spermatid associated (Spert) alternative variant cSep08, mRNA.
spertu	spertu.aSep08		11795	1938	3	97	uncharacterized protein (11.2 kD) (spertu) alternative variant aSep08, mRNA.
spervo	spervo.aSep08		31741	1780	11	485	CRA d (spervo) alternative variant aSep08, mRNA.
spervo	spervo.bSep08		11544	858	6	285	CRA d (spervo) alternative variant bSep08, mRNA.
spervo	spervo.cSep08		24007	782	6	123	CRA d like (spervo) alternative variant cSep08, mRNA.
spervo	spervo.dSep08		4244	411	3	86	CRA d like (spervo) alternative variant dSep08, mRNA.
sperwer	sperwer.aSep08		7520	466	3	97	putative protein (sperwer) alternative variant aSep08, mRNA.
sperwer	sperwer.bSep08		9695	847	4	79	putative protein (sperwer) alternative variant bSep08, mRNA.
Spesp1	Spesp1.bSep08	501010	14489	735	1	178	sperm equatorial segment protein 1 (Spesp1) alternative variant bSep08, mRNA.
Spetex-2AandSpetex-2B	Spetex-2AandSpetex-2B.aSep08	364261	20931	747		106	spetex-2A protein and Spetex-2B protein and similar to Spetex-2F protein (Spetex-2AandSpetex-2B) mRNA.
Spetex-2AandSpetex-2B	Spetex-2AandSpetex-2B.aSep08	497084	20931	747		106	spetex-2A protein and Spetex-2B protein and similar to Spetex-2F protein (Spetex-2AandSpetex-2B) mRNA.
Spetex-2AandSpetex-2B	Spetex-2AandSpetex-2B.aSep08	680542	20931	747		106	spetex-2A protein and Spetex-2B protein and similar to Spetex-2F protein (Spetex-2AandSpetex-2B) mRNA.
Spetex-2H	Spetex-2H.bSep08	361008	6252	375	1	124	spetex-2H protein (Spetex-2H) alternative variant bSep08, mRNA.

speycha	speycha.aSep08		3024	564		188	nucleoporin (speycha) mRNA.
speychy	speychy.aSep08		2201	778		78	putative protein (speychy) mRNA.
speyfee	speyfee.aSep08		3856	242		72	putative cytoplasmic protein (7.9 kD) (speyfee) mRNA.
speyflu	speyflu.aSep08		6652	384		34	putative protein (4.0 kD) (speyflu) mRNA.
speyfly	speyfly.aSep08		5280	576	2	68	CRA a like (speyfly) alternative variant aSep08, mRNA.
speyfly	speyfly.bSep08		3098	353	1	44	putative protein (speyfly) alternative variant bSep08, mRNA.
speylor	speylor.aSep08		7224	870		132	nucleolar protein 1 (speylor) mRNA.
speymee	speymee.aSep08		28021	492		50	putative protein (speymee) mRNA.
speynor	speynor.aSep08		876	573		118	peroxisomal biogenesis factor 6 (speynor) mRNA.
speypey	speypey.aSep08		2254	925		106	putative protein of mammalian origin (speypey) mRNA.
speyroy	speyroy.aSep08		15156	564		187	proline serine-rich coiled-coil 2 (speyroy) mRNA.
speyshaw	speyshaw.aSep08		6986	328		71	putative protein (speyshaw) mRNA.
speytu	speytu.aSep08		4683	393		131	uncharacterized protein (speytu) mRNA.
speyvo	speyvo.aSep08		12981	415		58	putative protein (speyvo) mRNA.
speywer	speywer.aSep08		7002	274		54	putative protein (speywer) mRNA.
Spg3a	Spg3a.bSep08	362750	43599	865	7	196	spastic paraplegia 3A homolog (human) (Spg3a) alternative variant bSep08, mRNA.
Spg7	Spg7.bSep08	353231	792	386	2	102	spastic paraplegia 7 homolog (human) (Spg7) alternative variant bSep08, mRNA.
Spg21	Spg21.aSep08	300791	27788	1962	9	308	spastic paraplegia 21 homolog (human) (35.0 kD) (Spg21) alternative variant aSep08, complete mRNA.
Sphk1	Sphk1.aSep08	170897	2906	1692	6	474	sphingosine kinase 1 (Sphk1) alternative variant aSep08, mRNA.
Sphk1	Sphk1.cSep08	170897	2787	786	5	195	sphingosine kinase 1 (Sphk1) alternative variant cSep08, mRNA.
Sphk1	Sphk1.dSep08	170897	2556	703	3	127	sphingosine kinase 1 (13.4 kD) (Sphk1) alternative variant dSep08, mRNA.
Sphk2	Sphk2.bSep08	308589	5510	818	3	185	sphingosine kinase 2 (Sphk2) alternative variant bSep08, mRNA.
Sphk2	Sphk2.dSep08	308589	565	483	2	67	sphingosine kinase 2 (Sphk2) alternative variant dSep08, mRNA.
Sphkap	Sphkap.bSep08	316561	10409	1593	2	498	protein sphkap (Sphkap) alternative variant bSep08, mRNA.
Sphkap	Sphkap.cSep08	316561	159521	1432	6	242	putative protein of vertebrate origin (Sphkap) alternative variant cSep08, mRNA.
Sphkap	Sphkap.dSep08	316561	5192	1073	3	79	sphkap protein (Sphkap) alternative variant dSep08, mRNA.
Spin-Ssty.3	Spin-Ssty.3.aSep08		5761	731	3	125	putative mitochondrial protein of eukaryotic origin (14.3 kD) (Spin-Ssty.3) alternative variant aSep08, mRNA.
Spin1	Spin1.bSep08	361217	43159	637	2	132	spindlin 1 and hypothetical protein LOC689247 (Spin1) alternative variant bSep08, mRNA.
Spin1	Spin1.bSep08	689247	43159	637	2	132	spindlin 1 and hypothetical protein LOC689247 (Spin1) alternative variant bSep08, mRNA.



Spin1	Spin1.cSep08	361217	48074	652	2	129	spindlin 1 and hypothetical protein LOC689247 (Spin1) alternative variant cSep08, mRNA.
Spin1	Spin1.cSep08	689247	48074	652	2	129	spindlin 1 and hypothetical protein LOC689247 (Spin1) alternative variant cSep08, mRNA.
Spin2	Spin2.aSep08	317395	16717	714	3	144	spindlin family, member 2 (Spin2) alternative variant aSep08, mRNA.
Spin2	Spin2.bSep08	317395	5247	521	3	62	spindlin family, member 2 (Spin2) alternative variant bSep08, mRNA.
Spin2	Spin2.cSep08	317395	10296	688	4	56	spindlin family, member 2 (6.3 kD) (Spin2) alternative variant cSep08, mRNA.
Spin2	Spin2.dSep08	317395	16663	738	4	43	spindlin family, member 2 (4.9 kD) (Spin2) alternative variant dSep08, mRNA.
Spink5	Spink5.aSep08	361319	49259	2789		836	serine peptidase inhibitor, Kazal type 5 (Spink5) mRNA.
Spink8	Spink8.bSep08	301016	11307	580	3	101	serine peptidase inhibitor, Kazal type 8 (11.4 kD) (Spink8) alternative variant bSep08, mRNA.
Spink8	Spink8.cSep08	301016	11531	2003	2	81	serine peptidase inhibitor, Kazal type 8 (Spink8) alternative variant cSep08, mRNA.
Spinlw1	Spinlw1.bSep08	685161	6736	431	2	55	serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin) (Spinlw1) alternative variant bSep08, mRNA.
Spinlw1	Spinlw1.cSep08	685161	790	351	1	36	serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin) (4.1 kD) (Spinlw1) alternative variant cSep08, mRNA.
Spint1	Spint1.bSep08	311331	12036	1773	3	451	serine protease inhibitor, Kunitz type 1 (50.4 kD) (Spint1) alternative variant bSep08, mRNA.
Spint1	Spint1.cSep08	311331	3061	615	2	171	serine protease inhibitor, Kunitz type 1 (Spint1) alternative variant cSep08, mRNA.
Spint2	Spint2.cSep08	292770	20793	732	5	222	serine protease inhibitor, Kunitz type 2 (Spint2) alternative variant cSep08, mRNA.
Spint2	Spint2.dSep08	292770	20171	743	4	73	serine protease inhibitor, Kunitz type 2 (Spint2) alternative variant dSep08, mRNA.
Spint2	Spint2.eSep08	292770	1636	1016	2	93	serine protease inhibitor, Kunitz type 2 (Spint2) alternative variant eSep08, mRNA.
Spire1	Spire1.bSep08	307348	26736	657	7	219	spire homolog 1 (Drosophila) (Spire1) alternative variant bSep08, mRNA.
Spire1	Spire1.cSep08	307348	7037	1026	5	159	spire homolog 1 (Drosophila) (Spire1) alternative variant cSep08, mRNA.
Spna2	Spna2.bSep08	64159	5550	1660	11	430	spectrin alpha 2 (Spna2) alternative variant bSep08, mRNA.
Spna2	Spna2.cSep08	64159	9644	1224	10	408	non-erythrocytic spectrin alpha (Spna2) alternative variant cSep08, mRNA.
Spna2	Spna2.dSep08	64159	3704	899	7	273	alpha spectrin (Spna2) alternative variant dSep08, mRNA.
Spna2	Spna2.eSep08	64159	1254	732	4	162	alpha spectrin (Spna2) alternative variant eSep08, mRNA.
Spna2	Spna2.fSep08	64159	5142	379	3	107	alpha spectrin (Spna2) alternative variant fSep08, mRNA.
Spna2	Spna2.gSep08	64159	2310	846	3	93	non-erythrocytic spectrin alpha (Spna2) alternative variant gSep08, mRNA.

Spna2	Spna2.hSep08	64159	855	775	2	79	spectrin alpha precursor (9.2 kD) (Spna2) alternative variant hSep08, mRNA.
Spnb1	Spnb1.bSep08	314251	7410	1784	5	367	spectrin beta 1 and hypothetical protein LOC679700 (Spnb1) alternative variant bSep08, mRNA.
Spnb1	Spnb1.bSep08	679700	7410	1784	5	367	spectrin beta 1 and hypothetical protein LOC679700 (Spnb1) alternative variant bSep08, mRNA.
Spnb1	Spnb1.cSep08	314251	898	670	2	79	spectrin beta 1 and hypothetical protein LOC679700 (Spnb1) alternative variant cSep08, mRNA.
Spnb1	Spnb1.cSep08	679700	898	670	2	79	spectrin beta 1 and hypothetical protein LOC679700 (Spnb1) alternative variant cSep08, mRNA.
Spnb2	Spnb2.bSep08	305614	17412	1110	7	370	spectrin beta (Spnb2) alternative variant bSep08, mRNA.
Spnb2	Spnb2.cSep08	305614	15083	998	7	332	beta spectrin (Spnb2) alternative variant cSep08, mRNA.
Spnb2	Spnb2.dSep08	305614	5170	673	4	224	spectrin beta (Spnb2) alternative variant dSep08, mRNA.
Spnb2	Spnb2.eSep08	305614	6687	3205	4	224	spectrin beta (24.7 kD) (Spnb2) alternative variant eSep08, mRNA.
Spnb2	Spnb2.fSep08	305614	8485	682	5	176	spectrin beta (Spnb2) alternative variant fSep08, mRNA.
Spnb2	Spnb2.gSep08	305614	2185	1639	3	84	spectrin beta 2 CRA b (Spnb2) alternative variant gSep08, mRNA.
Spnb2	Spnb2.hSep08	305614	3942	3178	2	80	spectrin beta (8.6 kD) (Spnb2) alternative variant hSep08, mRNA.
Spnb3	Spnb3.bSep08	29211	1214	842	2	280	spectrin beta 3 (Spnb3) alternative variant bSep08, mRNA.
Spnb4	Spnb4.aSep08	308458	16435	752		250	spectrin beta 4 (Spnb4) mRNA.
Spo11	Spo11.bSep08	366261	6611	666	8	221	SPO11 meiotic protein covalently bound to DSB homolog ( <i>S. cerevisiae</i> ) (Spo11) alternative variant bSep08, mRNA.
spobor	spobor.aSep08		1521	394	2	35	putative protein (spobor) alternative variant aSep08, mRNA.
spobor	spobor.bSep08		466	383	1	33	putative protein (spobor) alternative variant bSep08, mRNA.
SPOC.0	SPOC.0.aSep08		8897	455		151	death (SPOC.0) mRNA.
spocha	spocha.aSep08		1494	496	2	77	CRA a like (spocha) alternative variant aSep08, mRNA.
spochy	spochy.aSep08		13982	334		25	putative protein (spochy) mRNA.
Spock1	Spock1.aSep08	306759	503589	1790		431	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1 (Spock1) alternative variant aSep08, mRNA.
Spock1	Spock1.bSep08	306759	11785	1132		203	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1 (Spock1) alternative variant bSep08, mRNA.
Spock1	Spock1.cSep08	306759	947	343		114	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1 (Spock1) alternative variant cSep08, mRNA.
Spock2	Spock2.aSep08	361840	26098	3626	2	385	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2 (43.0 kD) (Spock2) alternative variant aSep08, mRNA.
Spock3	Spock3.bSep08	306404	106845	555	1	139	sparc/osteonectin, cwcv and kazal-like domains proteoglycan 3 (Spock3) alternative variant bSep08, mRNA.

spofee	spofee.aSep08		10129	901	5	299	WNK lysine deficient protein kinase 2 (spofee) alternative variant aSep08, mRNA.
spoflu	spoflu.aSep08		392	286		44	putative protein (4.8 kD) (spoflu) mRNA.
spofly	spofly.aSep08		5288	2135	2	299	excision repair cross-complementing rodent deficiency complementation group 2 (spofly) alternative variant aSep08, mRNA.
spofly	spofly.bSep08		821	532	1	136	excision repair cross-complementing rodent deficiency complementation group 2 (spofly) alternative variant bSep08, mRNA.
spolor	spolor.aSep08		1598	380		69	putative protein (7.7 kD) (spolor) mRNA.
spomee	spomee.aSep08		1180	267		59	putative protein (spomee) mRNA.
Spon1	Spon1.bSep08	64456	11550	3555	9	506	spondin 1, extracellular matrix protein (Spon1) alternative variant bSep08, mRNA.
Spon1	Spon1.cSep08	64456	52627	334	3	111	spondin 1, extracellular matrix protein (Spon1) alternative variant cSep08, mRNA.
sponsor	sponsor.aSep08		787	409		136	parkin-like cytoplasmic protein (sponsor) mRNA.
Spop	Spop.bSep08	287643	67630	771	6	152	speckle-type POZ protein (Spop) alternative variant bSep08, mRNA.
Spop	Spop.cSep08	287643	67363	397	5	81	speckle-type POZ protein (Spop) alternative variant cSep08, mRNA.
Spop	Spop.dSep08	287643	4345	390	2	73	speckle-type POZ protein (Spop) alternative variant dSep08, mRNA.
Spop	Spop.fSep08	287643	67236	300	3	36	speckle-type POZ protein (3.7 kD) (Spop) alternative variant fSep08, complete mRNA.
spopey	spopey.aSep08		5556	470		69	putative protein (7.2 kD) (spopey) mRNA.
sporcha	sporcha.aSep08		5545	379		40	putative protein (sporcha) mRNA.
sporchy	sporchy.aSep08		691	261		46	putative protein (sporchy) mRNA.
sporfee	sporfee.aSep08		1685	1161		84	putative protein (8.7 kD) (sporfee) mRNA.
sporflu	sporflu.aSep08		1414	553		95	putative protein (sporflu) mRNA.
sporfly	sporfly.aSep08		849	377		71	immunity-related GTPase family Q CRA a (sporfly) mRNA.
sporlor	sporlor.aSep08		997	754		111	putative protein (sporlor) mRNA.
spormee	spormee.aSep08		24205	1784		71	putative protein (7.8 kD) (spormee) alternative variant aSep08, mRNA.
spormee	spormee.bSep08		23503	314	1	71	putative protein (7.8 kD) (spormee) alternative variant bSep08, mRNA.
spormee	spormee.cSep08		5743	791	2	53	putative protein (spormee) alternative variant cSep08, mRNA.
spornor	spornor.aSep08		12874	831	3	276	zinc finger protein 318 (spornor) alternative variant aSep08, mRNA.
spornor	spornor.bSep08		12506	311	2	103	zinc finger protein 318 CRA b (spornor) alternative variant bSep08, mRNA.
sporoy	sporoy.aSep08		4663	409		136	oxysterol-binding protein-like protein 8 (sporoy) mRNA.
sporpey	sporpey.aSep08		657	232		72	putative protein of mammalian origin (sporpey) mRNA.
sporroy	sporroy.aSep08		785	371		30	putative protein (3.4 kD) (sporroy) mRNA.

sporshaw	sporshaw.aSep08		2190	649	2	142	putative protein (sporshaw) alternative variant aSep08, mRNA.
sportu	sportu.aSep08		883	252		84	putative protein of vertebrate origin (sportu) mRNA.
sporvo	sporvo.aSep08		1822	575		110	putative protein (12.3 kD) (sporvo) mRNA.
sporwer	sporwer.aSep08		12717	944		32	putative protein of mammalian origin (sporwer) mRNA.
sposhaw	sposhaw.aSep08		7824	784		66	putative protein (7.3 kD) (sposhaw) mRNA.
spotu	spotu.aSep08		16062	394		131	CRA b (spotu) mRNA.
spovo	spovo.aSep08		47068	413		95	putative protein (10.5 kD) (spovo) mRNA.
spower	spower.aSep08		18741	528		128	oxysterol-binding protein-like protein 3 (spower) mRNA.
spoycha	spoycha.aSep08		1230	522		81	putative protein (8.6 kD) (spoycha) mRNA.
spoychy	spoychy.aSep08		12169	381		54	putative protein (5.9 kD) (spoychy) mRNA.
spoyfee	spoyfee.bSep08		26518	390	2	49	putative protein (spoyfee) alternative variant bSep08, mRNA.
spoyflu	spoyflu.aSep08		5652	410		106	putative cytoplasmic protein (11.3 kD) (spoyflu) mRNA.
spoyfly	spoyfly.aSep08		2117	751		250	immunity-related GTPase family Q CRA a (spoyfly) mRNA.
spoylor	spoylor.aSep08		4676	441		89	synaptojanin 1 CRA b (spoylor) mRNA.
spoymee	spoymee.aSep08		2491	329		109	putative protein (spoymee) mRNA.
spoynor	spoynor.aSep08		9882	603		119	putative protein (spoynor) mRNA.
spoyroy	spoyroy.aSep08		16995	717		67	putative protein (7.4 kD) (spoyroy) mRNA.
spoyshaw	spoyshaw.aSep08		90603	683		69	putative protein (7.7 kD) (spoyshaw) mRNA.
spoytu	spoytu.aSep08		1506	278		46	putative protein (spoytu) mRNA.
spoyvo	spoyvo.aSep08		2978	1714		67	putative protein (spoyvo) alternative variant aSep08, mRNA.
spoywer	spoywer.aSep08		4191	854		50	putative protein (5.5 kD) (spoywer) mRNA.
Spp2	Spp2.aSep08	94168	19461	818	2	232	secreted phosphoprotein 2 (Spp2) alternative variant aSep08, mRNA.
Spp2	Spp2.cSep08	94168	19398	647	1	176	secreted phosphoprotein 2 (Spp2) alternative variant cSep08, mRNA.
Sppl2b	Sppl2b.bSep08	362828	2265	794	5	211	signal peptide peptidase-like 2B (Sppl2b) alternative variant bSep08, mRNA.
Sppl2b	Sppl2b.cSep08	362828	6977	611	6	200	signal peptide peptidase-like 2B (Sppl2b) alternative variant cSep08, mRNA.
Sppl2b	Sppl2b.eSep08	362828	11394	563	6	116	signal peptide peptidase-like 2B (Sppl2b) alternative variant eSep08, mRNA.
Sppl2b	Sppl2b.fSep08	362828	2030	341	3	113	signal peptide peptidase-like 2B (Sppl2b) alternative variant fSep08, mRNA.
Sppl2b	Sppl2b.gSep08	362828	456	365	2	49	signal peptide peptidase-like 2B (Sppl2b) alternative variant gSep08, mRNA.
Sppl3	Sppl3.aSep08	360822	85699	2836	11	448	signal peptide peptidase 3 (Sppl3) alternative variant aSep08, mRNA.
Sppl3	Sppl3.cSep08	360822	34993	770	5	138	signal peptide peptidase 3 (15.0 kD) (Sppl3) alternative variant cSep08, mRNA.
Sppl3	Sppl3.dSep08	360822	8852	1290	5	119	signal peptide peptidase 3 (13.2 kD) (Sppl3) alternative variant dSep08, mRNA.

Sppl3	Sppl3.eSep08	360822	1437	725	2	66	signal peptide peptidase 3 (10.6 kD) (Sppl3) alternative variant eSep08, mRNA.
Spr	Spr.bSep08	29270	1349	704	1	36	sepiapterin reductase (4.2 kD) (Spr) alternative variant bSep08, mRNA.
Spred2	Spred2.aSep08	305539	24441	1786	6	456	2 sprouty-related 1 domain (Spred2) alternative variant aSep08, mRNA.
Spred3	Spred3.aSep08	308478	8808	1775		419	sprouty (43.5 kD) (Spred3) mRNA.
SPRY.0	SPRY.0.aSep08		954	396		132	midline 2 (SPRY.0) mRNA.
SPRY.1	SPRY.1.aSep08		1368	958		212	tripartite motif-containing 50 (SPRY.1) mRNA.
SPRY.2	SPRY.2.aSep08		1385	788		262	tripartite motif-containing 65 (SPRY.2) mRNA.
SPRY.5	SPRY.5.aSep08		13428	1089	4	341	probable E3 ubiquitin-protein ligase herc1 (SPRY.5) alternative variant aSep08, mRNA.
SPRY.5	SPRY.5.bSep08		4433	367	1	122	probable E3 ubiquitin-protein ligase herc1 like (SPRY.5) alternative variant bSep08, mRNA.
SPRY.6	SPRY.6.aSep08		27916	4569	39	1312	ring finger protein 123 CRA a (148.3 kD) (SPRY.6) alternative variant aSep08, mRNA.
SPRY.6	SPRY.6.bSep08		9219	1089	10	303	ring finger protein 123 (33.0 kD) (SPRY.6) alternative variant bSep08, mRNA.
SPRY.6	SPRY.6.cSep08		1923	682	5	127	ring finger protein 123 (SPRY.6) alternative variant cSep08, mRNA.
SPRY.6	SPRY.6.dSep08		1271	736	2	96	CRA d like (SPRY.6) alternative variant dSep08, mRNA.
SPRY.6	SPRY.6.eSep08		7436	787	5	90	ring finger protein 123 (SPRY.6) alternative variant eSep08, mRNA.
SPRY.6	SPRY.6.fSep08		7766	450	4	72	ring finger protein 123 (SPRY.6) alternative variant fSep08, mRNA.
SPRY.7	SPRY.7.bSep08		1901	445	4	115	tripartite motif-containing 6 (SPRY.7) alternative variant bSep08, mRNA.
SPRY.8	SPRY.8.aSep08		11883	2749		124	nuclear ribonucleoprotein U-like 2 (SPRY.8) mRNA.
Spry3	Spry3.bSep08	498159	9138	3259	2	281	sprouty homolog 3 (Drosophila) (30.4 kD) (Spry3) alternative variant bSep08, mRNA.
Spryd3	Spryd3.aSep08	315327	17202	2470	9	461	spry 3 (Spryd3) alternative variant aSep08, mRNA.
Spryd3	Spryd3.bSep08	315327	3943	455	1	135	putative protein of eukaryotic origin (Spryd3) alternative variant bSep08, mRNA.
Spsb1	Spsb1.aSep08	313722	60930	2965	2	273	splA/ryanodine receptor domain and SOCS box containing 1 (30.8 kD) (Spsb1) alternative variant aSep08, complete mRNA.
Spsb2	Spsb2.aSep08	297592	1690	1496	1	276	splA/ryanodine receptor domain and SOCS box containing 2 (29.9 kD) (Spsb2) alternative variant aSep08, mRNA.
Spsb3	Spsb3.bSep08	302981	4571	776	5	240	splA ryanodine receptor domain socs box containing 3 (Spsb3) alternative variant bSep08, mRNA.
Spsb3	Spsb3.cSep08	302981	4507	1057	5	233	splA ryanodine receptor domain socs box containing 3 (Spsb3) alternative variant cSep08, mRNA.
Spsb3	Spsb3.dSep08	302981	4488	799	6	224	splA ryanodine receptor domain socs box containing 3 (Spsb3) alternative variant dSep08, mRNA.
Spsb3	Spsb3.eSep08	302981	4317	674	3	181	splA ryanodine receptor domain socs box containing 3 (19.8 kD) (Spsb3) alternative variant eSep08, mRNA.

Spsb3	Spsb3.fSep08	302981	3766	712	2	83	putative mitochondrial protein (9.2 kD) (Spsb3) alternative variant fSep08, mRNA.
Spsb4	Spsb4.bSep08	300950	71711	545		86	spiA/ryanodine receptor domain and SOCS box containing 4 (Spsb4) alternative variant bSep08, mRNA.
Sptbn5	Sptbn5.aSep08	296090	1543	767		255	spectrin, beta, non-erythrocytic 5 (Sptbn5) mRNA.
Sptlc1	Sptlc1.bSep08	361213	882	316	1	43	serine palmitoyltransferase, long chain base subunit 1 (Sptlc1) alternative variant bSep08, mRNA.
spubor	spubor.aSep08		3898	342		113	putative protein of metazoan origin (spubor) mRNA.
spucha	spucha.aSep08		5169	350	1	114	rap guanine nucleotide exchange factor 1 (spucha) alternative variant aSep08, mRNA.
spucha	spucha.bSep08		5120	317	1	105	rap guanine nucleotide exchange factor 1 (spucha) alternative variant bSep08, mRNA.
spucha	spucha.cSep08		5203	270		89	rap guanine nucleotide exchange factor 1 (spucha) alternative variant cSep08, mRNA.
spuchy	spuchy.aSep08		10141	779		36	putative protein (spuchy) mRNA.
spufee	spufee.aSep08		368	284		90	putative protein (spufee) mRNA.
spuflu	spuflu.aSep08		5072	437		48	putative protein (spuflu) mRNA.
spuflly	spuflly.aSep08		3388	372		123	excision repair cross-complementing rodent deficiency complementation group 2 CRA b (spuflly) mRNA.
spulor	spulor.aSep08		1610	472		126	putative protein (spulor) mRNA.
spumee	spumee.aSep08		10305	326		42	putative protein (spumee) mRNA.
spunor	spunor.aSep08		4283	446		142	putative protein (spunor) mRNA.
spupey	spupey.aSep08		38653	587		46	putative protein (spupey) mRNA.
spuroy	spuroy.aSep08		1188	491	2	50	putative protein (spuroy) alternative variant aSep08, mRNA.
spuroy	spuroy.bSep08		1216	485	2	64	putative protein (spuroy) alternative variant bSep08, mRNA.
spushaw	spushaw.aSep08		12403	791		35	putative protein (4.1 kD) (spushaw) mRNA.
sputu	sputu.aSep08		4702	417	2	138	CRA b (sputu) alternative variant aSep08, mRNA.
sputu	sputu.bSep08		2646	375	1	79	putative protein (sputu) alternative variant bSep08, mRNA.
spuvo	spuvo.aSep08		917	530		51	putative protein (spuvo) mRNA.
spuwer	spuwer.aSep08		4599	568		100	oxysterol-binding protein-like protein 3 (spuwer) mRNA.
spybor	spybor.aSep08		445	260		35	putative protein (spybor) mRNA.
spycha	spycha.aSep08		6755	604		92	putative protein (10.1 kD) (spycha) mRNA.
spychy	spychy.aSep08		2518	311		48	putative protein (spychy) mRNA.
spyfee	spyfee.aSep08		12300	670		222	kinase (spyfee) mRNA.
spyflu	spyflu.aSep08		42751	770	2	54	putative protein (spyflu) mRNA.
spyfly	spyfly.aSep08		445	365		121	protein phosphatase 1 regulatory like (spyfly) mRNA.
spylor	spylor.aSep08		4479	709		86	putative protein (spylor) mRNA.
spymee	spymee.aSep08		4032	542	1	53	putative protein (5.9 kD) (spymee) alternative variant aSep08, mRNA.
spymee	spymee.bSep08		4136	1003	1	59	putative protein (spymee) alternative variant bSep08, mRNA.
spynor	spynor.aSep08		1551	808		268	putative protein (spynor) mRNA.

spypey	spypey.aSep08		12401	4932		1591	rho GTPase-activating protein (spypey) mRNA.
spyroy	spyroy.aSep08		6988	541		115	neuron navigator 3 (spyroy) mRNA.
spyshaw	spyshaw.aSep08		2778	543		74	putative membrane protein of eukaryotic origin (8.4 kD) (spyshaw) mRNA.
spytu	spytu.aSep08		2709	741		247	CRA a (spytu) mRNA.
spyvo	spyvo.aSep08		4601	813		76	putative protein (spyvo) mRNA.
spywer	spywer.aSep08		26665	602		194	deafness autosomal dominant 5 (spywer) mRNA.
Sqrdl	Sqrdl.aSep08	362292	7290	481	2	153	sulfide quinone reductase-like (yeast) (Sqrdl) alternative variant aSep08, mRNA.
Sqstm1	Sqstm1.aSep08	113894	11296	2058	8	472	sequestosome 1 (Sqstm1) alternative variant aSep08, mRNA.
Sqstm1	Sqstm1.cSep08	113894	9080	1170	8	384	sequestosome 1 (Sqstm1) alternative variant cSep08, mRNA.
Sqstm1	Sqstm1.dSep08	113894	10835	1300	8	374	sequestosome 1 (Sqstm1) alternative variant dSep08, mRNA.
Sqstm1	Sqstm1.eSep08	113894	3345	1146	3	312	sequestosome 1 (Sqstm1) alternative variant eSep08, complete mRNA.
Sqstm1	Sqstm1.fSep08	113894	3197	896	4	278	sequestosome 1 (Sqstm1) alternative variant fSep08, complete mRNA.
Sqstm1	Sqstm1.gSep08	113894	6552	760	5	253	sequestosome 1 CRA c (Sqstm1) alternative variant gSep08, mRNA.
Sqstm1	Sqstm1.iSep08	113894	2667	811	4	216	sequestosome 1 CRA b (Sqstm1) alternative variant iSep08, mRNA.
Sqstm1	Sqstm1.jSep08	113894	10610	1127	7	188	sequestosome 1 CRA b (Sqstm1) alternative variant jSep08, mRNA.
Sqstm1	Sqstm1.kSep08	113894	3050	1418	3	130	sequestosome 1 (14.1 kD) (Sqstm1) alternative variant kSep08, mRNA.
Sqstm1	Sqstm1.lSep08	113894	2493	747	3	92	sequestosome 1 (10.2 kD) (Sqstm1) alternative variant lSep08, mRNA.
Sqstm1	Sqstm1.oSep08	113894	562	384	2		
Srbd1	Srbd1.aSep08	301665	200314	2788	12	735	s1 RNA binding domain 1 (Srbd1) alternative variant aSep08, mRNA.
Srbd1	Srbd1.bSep08	301665	32138	1309	2	258	s1 RNA binding domain 1 (Srbd1) alternative variant bSep08, mRNA.
Srbd1	Srbd1.cSep08	301665	7926	812	1	174	s1 RNA binding domain 1 (Srbd1) alternative variant cSep08, mRNA.
Src	Src.bSep08	83805	24944	790	6	181	rous sarcoma oncogene (Src) alternative variant bSep08, mRNA.
SRCR.0	SRCR.0.aSep08		7839	882		293	deleted in malignant brain tumors 1 like (SRCR.0) mRNA.
Srcrb4d	Srcrb4d.aSep08	304401	2046	395	3	131	CRA c (Srcrb4d) alternative variant aSep08, mRNA.
Srcrb4d	Srcrb4d.bSep08	304401	2557	1129	3	113	putative protein (Srcrb4d) alternative variant bSep08, mRNA.
Srcrb4d	Srcrb4d.cSep08	304401	8368	365	3	101	CRA c like (Srcrb4d) alternative variant cSep08, mRNA.
Srd5a3	Srd5a3.aSep08	305291	15584	2102	1	348	steroid 5 alpha-reductase 3 (Srd5a3) alternative variant aSep08, mRNA.

Sreb1	Sreb1.aSep08	78968	22525	4876	17	1144	sterol regulatory element binding transcription factor 1 (Sreb1) alternative variant aSep08, mRNA.
Sreb1	Sreb1.bSep08	78968	1071	981	1	127	sterol regulatory element binding transcription factor 1 (Sreb1) alternative variant bSep08, mRNA.
Sreb2	Sreb2.bSep08	300095	20443	783	1	116	sterol regulatory element binding factor 2 (12.5 kD) (Sreb2) alternative variant bSep08, mRNA.
Srf	Srf.bSep08	501099	4211	577		192	serum response factor (Srf) alternative variant bSep08, mRNA.
Srfbp1	Srfbp1.bSep08	291469	13939	1978	7	417	serum response factor binding protein 1 (46.6 kD) (Srfbp1) alternative variant bSep08, mRNA.
Srgap1	Srgap1.aSep08	314903	9663	374		124	SLIT-ROBO Rho GTPase activating protein 1 (Srgap1) alternative variant aSep08, mRNA.
Srgap3	Srgap3.aSep08	500287	10242	708		229	SLIT-ROBO Rho GTPase activating protein 3 (Srgap3) mRNA.
Srl	Srl.aSep08	302948	41018	805	5	268	sarcalumenin (Srl) alternative variant aSep08, mRNA.
Srl	Srl.bSep08	302948	25832	379	1	126	sarcalumenin (Srl) alternative variant bSep08, mRNA.
Srm	Srm.bSep08	84596	1306	646	4	129	spermidine synthase (Srm) alternative variant bSep08, mRNA.
Srm	Srm.cSep08	84596	1394	1152	3	110	spermidine synthase (12.7 kD) (Srm) alternative variant cSep08, mRNA.
Srp14	Srp14.bSep08	296076	3180	467	4	112	signal recognition particle 14 (Srp14) alternative variant bSep08, mRNA.
Srp14	Srp14.cSep08	296076	3155	299	5	96	signal recognition particle 14 (Srp14) alternative variant cSep08, mRNA.
Srp14	Srp14.dSep08	296076	3173	841	4	92	signal recognition particle 14 (10.3 kD) (Srp14) alternative variant dSep08, mRNA.
Srp19	Srp19.bSep08	291685	2164	738	3	45	putative protein (5.2 kD) (Srp19) alternative variant bSep08, mRNA.
Srp19	Srp19.cSep08	291685	4296	443	4	43	signal recognition particle (4.9 kD) (Srp19) alternative variant cSep08, mRNA.
Srp68	Srp68.bSep08	363707	1458	780	3	222	putative protein (Srp68) alternative variant bSep08, mRNA.
Srp68	Srp68.cSep08	363707	6326	797	5	115	signal recognition particle 68kDa CRA a (Srp68) alternative variant cSep08, mRNA.
Srp68	Srp68.dSep08	363707	6447	373	4	98	signal recognition particle 68kDa CRA a (Srp68) alternative variant dSep08, mRNA.
Srp68	Srp68.eSep08	363707	1060	748	2	89	putative protein (Srp68) alternative variant eSep08, mRNA.
Srp72	Srp72.aSep08	498351	12434	2473	9	307	signal recognition particle 72 (Srp72) alternative variant aSep08, mRNA.
Srp72	Srp72.bSep08	498351	2059	867	3	137	signal recognition particle 72 (Srp72) alternative variant bSep08, mRNA.
Srp72	Srp72.cSep08	498351	1410	815	2	124	signal recognition particle 72 (Srp72) alternative variant cSep08, mRNA.
Srp72	Srp72.dSep08	498351	915	233	2	54	signal recognition particle 72 (Srp72) alternative variant dSep08, mRNA.
Srp1	Srp1.aSep08	361811	43377	4204	1	517	serine/arginine-rich protein specific kinase 1 (Srp1) alternative variant aSep08, mRNA.



Srpk1	Srpk1.bSep08	361811	10474	792		171	serine/arginine-rich protein specific kinase 1 (Srpk1) alternative variant bSep08, mRNA.
Srpk2	Srpk2.aSep08	296753	100069	1006	9	271	serine/arginine-rich protein specific kinase 2 (Srpk2) alternative variant aSep08, mRNA.
Srpk2	Srpk2.bSep08	296753	160490	789	6	197	serine/arginine-rich protein specific kinase 2 (Srpk2) alternative variant bSep08, mRNA.
Srpk3	Srpk3.bSep08	293854	2600	630	7	210	serine/arginine-rich protein specific kinase 3 (Srpk3) alternative variant bSep08, mRNA.
Srpr	Srpr.bSep08	315548	1654	865	6	288	signal recognition particle receptor ('docking protein') (Srpr) alternative variant bSep08, mRNA.
Srpr	Srpr.cSep08	315548	2876	1798	6	281	signal recognition particle receptor ('docking protein') (Srpr) alternative variant cSep08, mRNA.
Srpr	Srpr.dSep08	315548	1046	739	3	135	signal recognition particle receptor ('docking protein') (14.5 kD) (Srpr) alternative variant dSep08, mRNA.
Srprb	Srprb.bSep08	300965	2889	386	3	112	signal recognition particle receptor, B subunit (Srprb) alternative variant bSep08, mRNA.
Srpx	Srpx.bSep08	64316	58480	695		231	sushi-repeat-containing protein (Srpx) alternative variant bSep08, mRNA.
Srr	Srr.bSep08	303306	16200	786	6	166	serine racemase (Srr) alternative variant bSep08, mRNA.
Srrd	Srrd.aSep08	288717	5983	1484	7	291	SRR1 (Srrd) alternative variant aSep08, mRNA.
Srrd	Srrd.bSep08	288717	4572	761	5	167	SRR1 (Srrd) alternative variant bSep08, mRNA.
Srrm1	Srrm1.bSep08	313620	8874	955	7	318	serine arginine repetitive matrix 1 (Srrm1) alternative variant bSep08, mRNA.
Srrm1	Srrm1.cSep08	313620	6300	801	6	267	serine arginine repetitive matrix 1 (Srrm1) alternative variant cSep08, mRNA.
Srrm1	Srrm1.dSep08	313620	19370	656	4	218	serine arginine repetitive matrix 1 (Srrm1) alternative variant dSep08, mRNA.
Srrm1	Srrm1.eSep08	313620	3784	789	4	187	serine arginine repetitive matrix 1 (Srrm1) alternative variant eSep08, mRNA.
Srrm1	Srrm1.fSep08	313620	2444	445	2	102	putative protein (Srrm1) alternative variant fSep08, mRNA.
Srrm1	Srrm1.gSep08	313620	741	397	2	84	serine arginine repetitive matrix 1 (Srrm1) alternative variant gSep08, mRNA.
Srrm1	Srrm1.hSep08	313620	1879	1233	3	72	serine arginine repetitive matrix 1 (Srrm1) alternative variant hSep08, mRNA.
Srrm2	Srrm2.aSep08	302969	5481	4439	5	1114	serine arginine repetitive matrix 2 (Srrm2) alternative variant aSep08, mRNA.
Srrm2	Srrm2.bSep08	302969	5025	3011	5	837	serine arginine repetitive matrix 2 (Srrm2) alternative variant bSep08, mRNA.
Srrm2	Srrm2.cSep08	302969	2215	1914	2	302	putative nuclear protein of mammalian origin (31.4 kD) (Srrm2) alternative variant cSep08, mRNA.
Srrm2	Srrm2.dSep08	302969	2225	2015	3	227	putative protein of mammalian origin (Srrm2) alternative variant dSep08, mRNA.
Srrm2	Srrm2.eSep08	302969	1422	410	3	136	serine arginine repetitive matrix 2 CRA f (Srrm2) alternative variant eSep08, mRNA.
Srrm2	Srrm2.fSep08	302969	595	371	2	80	putative protein of mammalian origin (8.7 kD) (Srrm2) alternative variant fSep08, mRNA.

Ss18	Ss18.bSep08	361295	25401	1131	6	217	putative protein of vertebrate origin (Ss18) alternative variant bSep08, mRNA.
Ss18	Ss18.cSep08	361295	25342	967	5	159	putative protein (Ss18) alternative variant cSep08, mRNA.
Ss18	Ss18.dSep08	361295	98001	693	4	143	putative protein of vertebrate origin (Ss18) alternative variant dSep08, mRNA.
Ss18	Ss18.eSep08	361295	62018	1051	7	79	synovial sarcoma (8.8 kD) (Ss18) alternative variant eSep08, complete mRNA.
Ssb	Ssb.bSep08	81783	5853	719	6	185	sjogren syndrome antigen B like (21.5 kD) (Ssb) alternative variant bSep08, mRNA.
Ssb	Ssb.cSep08	81783	5849	789	5	152	sjogren syndrome antigen B like (17.8 kD) (Ssb) alternative variant cSep08, mRNA.
Ssb	Ssb.dSep08	81783	5290	740	4	134	sjogren syndrome antigen B like (15.8 kD) (Ssb) alternative variant dSep08, mRNA.
Ssb	Ssb.eSep08	81783	1105	686	3	125	sjogren syndrome antigen B like (Ssb) alternative variant eSep08, mRNA.
Ssb	Ssb.fSep08	81783	1221	727	2	113	sjogren syndrome antigen B like (12.9 kD) (Ssb) alternative variant fSep08, mRNA.
Ssbp1	Ssbp1.aSep08	54304	10006	642	6	148	single-stranded DNA binding protein 1 (17.2 kD) (Ssbp1) alternative variant aSep08, mRNA.
Ssbp1	Ssbp1.bSep08	54304	10042	605	6	148	single-stranded DNA binding protein 1 (17.2 kD) (Ssbp1) alternative variant bSep08, mRNA.
Ssbp1	Ssbp1.cSep08	54304	10008	1186	5	148	single-stranded DNA binding protein 1 (17.2 kD) (Ssbp1) alternative variant cSep08, mRNA.
Ssbp1	Ssbp1.eSep08	54304	10043	693	6	144	single-stranded DNA binding protein 1 (16.7 kD) (Ssbp1) alternative variant eSep08, mRNA.
Ssbp1	Ssbp1.fSep08	54304	3735	643	1	57	single-stranded DNA binding protein 1 (6.6 kD) (Ssbp1) alternative variant fSep08, mRNA.
Ssbp2	Ssbp2.aSep08	361877	203803	840	14	280	single-stranded DNA binding protein 2 like (Ssbp2) alternative variant aSep08, mRNA.
Ssbp2	Ssbp2.bSep08	361877	252013	816	11	247	single-stranded DNA binding protein 2 CRA a like (Ssbp2) alternative variant bSep08, mRNA.
Ssbp2	Ssbp2.cSep08	361877	62601	603	8	140	single-stranded DNA binding protein 2 like (Ssbp2) alternative variant cSep08, mRNA.
Ssbp2	Ssbp2.dSep08	361877	26064	504	3	90	single-stranded DNA binding protein 2 CRA a like (Ssbp2) alternative variant dSep08, mRNA.
Ssbp3	Ssbp3.aSep08	84354	94019	1343	14	412	single stranded DNA binding protein 3 (Ssbp3) alternative variant aSep08, mRNA.
Ssbp3	Ssbp3.bSep08	84354	140954	3078	17	361	single stranded DNA binding protein 3 (37.7 kD) (Ssbp3) alternative variant bSep08, mRNA.
Ssbp3	Ssbp3.cSep08	84354	129637	882	12	294	single stranded DNA binding protein 3 (Ssbp3) alternative variant cSep08, mRNA.
Ssbp4	Ssbp4.aSep08	364534	10376	1261	17	382	single stranded DNA binding protein 4 (Ssbp4) alternative variant aSep08, mRNA.
Ssbp4	Ssbp4.cSep08	364534	917	388	6	129	single stranded DNA binding protein 4 (Ssbp4) alternative variant cSep08, mRNA.
Ssbp4	Ssbp4.fSep08	364534	441	314	2	45	single stranded DNA binding protein 4 (Ssbp4) alternative variant fSep08, mRNA.

Ssfa2	Ssfa2.bSep08	311146	9881	491	4	116	sperm specific antigen 2 (Ssfa2) alternative variant bSep08, mRNA.
Ssh1	Ssh1.bSep08	304580	15517	1779	6	539	slingshot homolog 1 (Drosophila) (Ssh1) alternative variant bSep08, mRNA.
Ssh3	Ssh3.bSep08	365396	5888	1408	7	265	slingshot homolog 3 (Drosophila) (Ssh3) alternative variant bSep08, mRNA.
Ssh3	Ssh3.cSep08	365396	6759	2468	4	172	slingshot homolog 3 (Drosophila) (20.0 kD) (Ssh3) alternative variant cSep08, mRNA.
Ssna1	Ssna1.aSep08	311802	1044	681	2	139	sjogren's syndrome nuclear autoantigen 1 (15.6 kD) (Ssna1) alternative variant aSep08, mRNA.
Ssna1	Ssna1.bSep08	311802	1454	918	3	119	sjogren's syndrome nuclear autoantigen 1 (13.6 kD) (Ssna1) alternative variant bSep08, mRNA.
Ssr1	Ssr1.bSep08	361233	5882	421	4	65	signal sequence receptor, alpha (Ssr1) alternative variant bSep08, mRNA.
Ssr3	Ssr3.aSep08	81784	3540	2569		67	signal sequence receptor, gamma (Ssr3) mRNA.
Ssr4	Ssr4.bSep08	29435	4064	921	6	214	signal sequence receptor, delta (Ssr4) alternative variant bSep08, mRNA.
Ssr4	Ssr4.cSep08	29435	3786	571	5	148	signal sequence receptor, delta (Ssr4) alternative variant cSep08, complete mRNA.
Ssr4	Ssr4.dSep08	29435	2434	1303	3	144	signal sequence receptor, delta (Ssr4) alternative variant dSep08, mRNA.
Ssrp1	Ssrp1.bSep08	81785	3247	1033	6	344	structure specific recognition protein 1 (Ssrp1) alternative variant bSep08, mRNA.
Ssrp1	Ssrp1.cSep08	81785	3250	950	6	316	structure specific recognition protein 1 (Ssrp1) alternative variant cSep08, mRNA.
Ssrp1	Ssrp1.dSep08	81785	5416	1696	7	303	structure specific recognition protein 1 (34.2 kD) (Ssrp1) alternative variant dSep08, mRNA.
Ssrp1	Ssrp1.eSep08	81785	2546	721	5	240	structure specific recognition protein 1 (Ssrp1) alternative variant eSep08, mRNA.
Ssrp1	Ssrp1.fSep08	81785	2871	1290	4	136	structure specific recognition protein 1 (Ssrp1) alternative variant fSep08, mRNA.
Ssrp1	Ssrp1.gSep08	81785	2470	559	5	132	structure specific recognition protein 1 (Ssrp1) alternative variant gSep08, mRNA.
Ssrp1	Ssrp1.hSep08	81785	1225	353	3	117	structure specific recognition protein 1 (Ssrp1) alternative variant hSep08, mRNA.
Ssrp1	Ssrp1.iSep08	81785	3219	1467	3	106	structure specific recognition protein 1 (12.0 kD) (Ssrp1) alternative variant iSep08, mRNA.
Sssca1	Sssca1.bSep08	689397	1178	453	2	89	sjogren's syndrome/scleroderma autoantigen 1 homolog (human) (Sssca1) alternative variant bSep08, mRNA.
Ssu72	Ssu72.bSep08	298681	14091	365	2	111	ssu72 RNA polymerase II CTD phosphatase homolog (yeast) (Ssu72) alternative variant bSep08, mRNA.
Ssx2ip	Ssx2ip.bSep08	308023	12011	699	6	232	synovial sarcoma, X breakpoint 2 interacting protein (Ssx2ip) alternative variant bSep08, mRNA.
Ssx2ip	Ssx2ip.cSep08	308023	18083	954	6	205	synovial sarcoma, X breakpoint 2 interacting protein (Ssx2ip) alternative variant cSep08, mRNA.
Ssx2ip	Ssx2ip.dSep08	308023	16830	702	5	179	synovial sarcoma, X breakpoint 2 interacting protein (Ssx2ip) alternative variant dSep08, mRNA.

Ssx2ip	Ssx2ip.eSep08	308023	11774	358	2	119	synovial sarcoma, X breakpoint 2 interacting protein (Ssx2ip) alternative variant eSep08, mRNA.
Ssx2ip	Ssx2ip.fSep08	308023	12526	723	5	102	synovial sarcoma, X breakpoint 2 interacting protein (Ssx2ip) alternative variant fSep08, mRNA.
Ssx2ip	Ssx2ip.gSep08	308023	2916	1787	2	95	synovial sarcoma, X breakpoint 2 interacting protein (10.1 kD) (Ssx2ip) alternative variant gSep08, mRNA.
St3gal1	St3gal1.bSep08	362924	1407	384	1	120	ST3 beta-galactoside alpha-2,3-sialyltransferase 1 (St3gal1) alternative variant bSep08, mRNA.
St3gal2	St3gal2.aSep08	64442	16005	3355		417	ST3 beta-galactoside alpha-2,3-sialyltransferase 2 (St3gal2) mRNA.
St3gal4	St3gal4.aSep08	363040	48778	2095	12	420	ST3 beta-galactoside alpha-2,3-sialyltransferase 4 (St3gal4) alternative variant aSep08, mRNA.
St3gal4	St3gal4.bSep08	363040	42152	743	8	169	ST3 beta-galactoside alpha-2,3-sialyltransferase 4 (St3gal4) alternative variant bSep08, mRNA.
St3gal4	St3gal4.cSep08	363040	40658	788	3	118	ST3 beta-galactoside alpha-2,3-sialyltransferase 4 (St3gal4) alternative variant cSep08, mRNA.
St3gal4	St3gal4.dSep08	363040	1022	727	2	92	ST3 beta-galactoside alpha-2,3-sialyltransferase 4 (10.3 kD) (St3gal4) alternative variant dSep08, mRNA.
St3gal4	St3gal4.fSep08	363040	6289	580	4	9	ST3 beta-galactoside alpha-2,3-sialyltransferase 4 (1.0 kD) (St3gal4) alternative variant fSep08, mRNA.
St3gal5	St3gal5.bSep08	83505	48376	762	1	246	ST3 beta-galactoside alpha-2,3-sialyltransferase 5 (St3gal5) alternative variant bSep08, mRNA.
St3gal5	St3gal5.cSep08	83505	48426	707	2	235	ST3 beta-galactoside alpha-2,3-sialyltransferase 5 (St3gal5) alternative variant cSep08, mRNA.
St3gal5	St3gal5.dSep08	83505	52122	737	2	167	ST3 beta-galactoside alpha-2,3-sialyltransferase 5 (St3gal5) alternative variant dSep08, mRNA.
St5	St5.aSep08	308944	33602	3165		825	suppression of tumorigenicity 5 (St5) mRNA.
St6gal1	St6gal1.cSep08	25197	6567	788	3	184	beta galactoside alpha 2,6 sialyltransferase 1 (21.9 kD) (St6gal1) alternative variant cSep08, mRNA.
St6gal1	St6gal1.dSep08	25197	2700	584	1	107	beta galactoside alpha 2,6 sialyltransferase 1 (12.4 kD) (St6gal1) alternative variant dSep08, mRNA.
St6gal1	St6gal1.eSep08	25197	3580	417	2	104	beta galactoside alpha 2,6 sialyltransferase 1 (St6gal1) alternative variant eSep08, mRNA.
St6gal2	St6gal2.aSep08	301155	105155	2343		525	beta galactoside alpha 2,6 sialyltransferase 2 (59.9 kD) (St6gal2) mRNA.
St6galnac2	St6galnac2.cSep08	303692	8838	709	5	138	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 2 (St6galnac2) alternative variant cSep08, mRNA.
St6galnac2	St6galnac2.dSep08	303692	3324	748	3	123	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 2 (St6galnac2) alternative variant dSep08, mRNA.
St6galnac2	St6galnac2.eSep08	303692	2031	1295	2	75	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 2 (8.8 kD) (St6galnac2) alternative variant eSep08, mRNA.
ST7	ST7.aSep08	296911	127288	2088	17	413	suppression of tumorigenicity 7 (47.2 kD) (ST7) alternative variant aSep08, mRNA.

ST7	ST7.bSep08	296911	247416	1282	9	285	suppression of tumorigenicity 7 (32.7 kD) (ST7) alternative variant bSep08, mRNA.
ST7	ST7.cSep08	296911	39115	677	7	225	suppression of tumorigenicity 7 (ST7) alternative variant cSep08, mRNA.
ST7	ST7.dSep08	296911	3485	829	2	77	suppression of tumorigenicity 7 (ST7) alternative variant dSep08, mRNA.
St7l	St7l.bSep08	295344	14313	468	2	146	suppression of tumorigenicity 7-like (St7l) alternative variant bSep08, mRNA.
St7l	St7l.cSep08	295344	24156	692	5	101	suppression of tumorigenicity 7-like (St7l) alternative variant cSep08, mRNA.
St8sia3	St8sia3.bSep08	25547	3427	804	3	239	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 3 (St8sia3) alternative variant bSep08, mRNA.
St8sia5	St8sia5.aSep08	364901	61632	2175	2	558	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5 (St8sia5) alternative variant aSep08, mRNA.
St13	St13.bSep08	81800	30089	3219	2	334	suppression of tumorigenicity 13 (37.1 kD) (St13) alternative variant bSep08, mRNA.
St13	St13.cSep08	81800	15730	748	1	154	suppression of tumorigenicity 13 (St13) alternative variant cSep08, mRNA.
St18	St18.bSep08	266680	3863	1411	2	91	suppression of tumorigenicity 18 (10.2 kD) (St18) alternative variant bSep08, mRNA.
Stac	Stac.aSep08	363152	52787	471	1	157	src homology three (SH3) and cysteine rich domain (Stac) alternative variant aSep08, mRNA.
Stac	Stac.bSep08	363152	52774	425	1	141	src homology three (SH3) and cysteine rich domain (Stac) alternative variant bSep08, mRNA.
Stac3	Stac3.bSep08	362895	5854	1140	7	307	SH3 and cysteine rich domain 3 (Stac3) alternative variant bSep08, mRNA.
Stac3	Stac3.cSep08	362895	856	410	4	90	SH3 and cysteine rich domain 3 (Stac3) alternative variant cSep08, mRNA.
stacha	stacha.aSep08		3307	326		108	putative protein (stacha) mRNA.
stachy	stachy.aSep08		543	314		33	putative protein (stachy) mRNA.
stafee	stafee.aSep08		1793	678		40	putative protein (4.3 kD) (stafee) mRNA.
staflu	staflu.aSep08		973	485		87	putative protein (staflu) mRNA.
stafly	stafly.bSep08		563	365	2	60	putative protein (stafly) alternative variant bSep08, mRNA.
Stag1	Stag1.bSep08	315958	48308	1186	8	394	stromal antigen 1 (Stag1) alternative variant bSep08, mRNA.
Stag1	Stag1.cSep08	315958	10624	483	2	126	stromal antigen 1 (Stag1) alternative variant cSep08, mRNA.
Stag2	Stag2.aSep08	313304	27484	3495	1	451	stromal antigen 2 (52.0 kD) (Stag2) alternative variant aSep08, mRNA.
Stag2	Stag2.bSep08	313304	12061	662		192	stromal antigen 2 (Stag2) alternative variant bSep08, mRNA.
Stag3	Stag3.bSep08	114522	6812	1208		402	stromal antigen 3 (Stag3) alternative variant bSep08, mRNA.
Stag3	Stag3.cSep08	114522	2380	508		54	stromal antigen 3 (Stag3) alternative variant cSep08, mRNA.

stalor	stalor.aSep08		20997	540	2	180	synaptojanin 1 CRA e (stalor) alternative variant aSep08, mRNA.
stalor	stalor.bSep08		20980	410	2	136	synaptojanin 1 CRA e (stalor) alternative variant bSep08, mRNA.
stalor	stalor.cSep08		20864	542	2	119	synaptojanin 1 CRA e (stalor) alternative variant cSep08, mRNA.
stalor	stalor.dSep08		18826	528	2	95	synaptojanin (stalor) alternative variant dSep08, mRNA.
Stam	Stam.bSep08	498798	28092	745	6	168	signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (Stam) alternative variant bSep08, mRNA.
Stampb	Stampb.aSep08	171565	24506	1796	2	505	stam binding protein (Stampb) alternative variant aSep08, complete mRNA.
Stampb	Stampb.cSep08	171565	13770	1106	7	247	stam binding protein (Stampb) alternative variant cSep08, mRNA.
Stampb	Stampb.dSep08	171565	11542	645	3	145	stam binding protein (Stampb) alternative variant dSep08, mRNA.
Stampb	Stampb.eSep08	171565	11437	744	6	83	stam binding protein (Stampb) alternative variant eSep08, mRNA.
stamee	stamee.aSep08		940	468		42	putative protein (4.6 kD) (stamee) mRNA.
stanor	stanor.aSep08		1108	687		32	putative protein (3.7 kD) (stanor) mRNA.
Stap1	Stap1.aSep08	305269	29699	1801	2	426	signal transducing adaptor family member 1 (Stap1) alternative variant aSep08, complete mRNA.
Stap1	Stap1.cSep08	305269	23300	777	3	239	signal transducing adaptor family member 1 (Stap1) alternative variant cSep08, mRNA.
Stap2	Stap2.aSep08	363334	2360	701	1	156	signal transducing adaptor family member 2 (Stap2) alternative variant aSep08, mRNA.
Stap2	Stap2.bSep08	363334	2488	633	1	135	signal transducing adaptor family member 2 (Stap2) alternative variant bSep08, mRNA.
stapey	stapey.aSep08		1703	264		55	putative protein (stapey) mRNA.
starcha	starcha.aSep08		846	599	2	61	putative protein (starcha) alternative variant aSep08, mRNA.
starchy	starchy.aSep08		2604	527		41	putative protein (4.5 kD) (starchy) mRNA.
Stard3	Stard3.bSep08	363675	18262	784	5	210	putative endoplasmic reticulum protein, with at least 2 transmembrane domains, of metazoan origin (23.1 kD) (Stard3) alternative variant bSep08, mRNA.
Stard3	Stard3.cSep08	363675	1150	754	2	149	putative protein (Stard3) alternative variant cSep08, mRNA.
Stard3	Stard3.dSep08	363675	18066	575	5	121	putative endoplasmic reticulum protein, with a transmembrane domain, of bilateral origin (13.7 kD) (Stard3) alternative variant dSep08, mRNA.
Stard3	Stard3.eSep08	363675	13855	594	2	108	putative membrane protein of bilateral origin (11.6 kD) (Stard3) alternative variant eSep08, mRNA.
Stard3nl	Stard3nl.bSep08	291182	6167	1086	1	158	STAR3 N-terminal like (Stard3nl) alternative variant bSep08, mRNA.
Stard5	Stard5.aSep08	502348	10000	1335	5	231	StAR-related lipid transfer 5 (Stard5) mRNA.
Stard6	Stard6.bSep08	291527	8675	506	2	76	hypothetical protein LOC680108 (Stard6) alternative variant bSep08, mRNA.

Stard6	Stard6.bSep08	680108	8675	506	2	76	hypothetical protein LOC680108 (Stard6) alternative variant bSep08, mRNA.
Stard7	Stard7.aSep08	296128	28650	3058	8	373	lipid-binding START precursor (43.0 kD) (Stard7) alternative variant aSep08, mRNA.
Stard7	Stard7.cSep08	296128	19972	836	5	138	putative protein, with a coiled coil domain, of metazoan origin (15.8 kD) (Stard7) alternative variant cSep08, mRNA.
Stard10	Stard10.aSep08	293150	26259	2200	6	372	lipid-binding START (Stard10) alternative variant aSep08, mRNA.
Stard10	Stard10.bSep08	293150	27167	1059	5	353	putative protein of eukaryotic origin (Stard10) alternative variant bSep08, mRNA.
Stard10	Stard10.dSep08	293150	26493	809	5	226	putative protein of eukaryotic origin (Stard10) alternative variant dSep08, mRNA.
Stard10	Stard10.eSep08	293150	24805	1101	5	222	lipid-binding START (Stard10) alternative variant eSep08, mRNA.
Stard10	Stard10.fSep08	293150	23624	744	5	165	putative protein of eukaryotic origin (Stard10) alternative variant fSep08, mRNA.
Stard10	Stard10.gSep08	293150	1243	632	1	128	putative protein of eukaryotic origin (Stard10) alternative variant gSep08, mRNA.
Stard13	Stard13.bSep08	498130	45935	606	2	113	StAR-related lipid transfer 13 (13.0 kD) (Stard13) alternative variant bSep08, mRNA.
starfee	starfee.aSep08		6614	745		106	putative protein (starfee) alternative variant aSep08, mRNA.
starflu	starflu.aSep08		736	497		112	potassium channel (starflu) mRNA.
starfly	starfly.aSep08		2705	459		42	putative protein (4.8 kD) (starfly) mRNA.
starlor	starlor.aSep08		19383	759		79	putative cytoplasmic protein (8.9 kD) (starlor) mRNA.
starmee	starmee.aSep08		820	278		40	putative protein (4.4 kD) (starmee) mRNA.
starnor	starnor.aSep08		2191	408		70	putative protein (starnor) mRNA.
staroy	staroy.aSep08		800	367		58	putative protein of mammalian origin (staroy) mRNA.
starpey	starpey.aSep08		14746	376		55	putative protein (starpey) mRNA.
starroy	starroy.aSep08		8214	1182		214	CRA d like (23.7 kD) (starroy) mRNA.
starshaw	starshaw.aSep08		20448	1131		59	putative protein (7.2 kD) (starshaw) alternative variant aSep08, mRNA.
starshaw	starshaw.bSep08		19650	332		25	putative protein (2.8 kD) (starshaw) alternative variant bSep08, mRNA.
startu	startu.aSep08		50239	702	3	144	putative secreted or extracellular protein precursor (16.0 kD) (startu) alternative variant aSep08, mRNA.
startu	startu.bSep08		19017	498	3	73	putative protein (startu) alternative variant bSep08, mRNA.
starvo	starvo.aSep08		3970	1637	5	91	putative protein (10.1 kD) (starvo) alternative variant aSep08, mRNA.
starvo	starvo.bSep08		1921	732	2	88	putative protein (starvo) alternative variant bSep08, mRNA.
starvo	starvo.cSep08		3277	1790	3	25	putative protein (starvo) alternative variant cSep08, mRNA.
starvo	starvo.dSep08		1662	1157	4	128	putative protein (starvo) alternative variant dSep08, mRNA.
starvo	starvo.eSep08		3184	721	6	91	putative protein (10.1 kD) (starvo) alternative variant eSep08, mRNA.

starwer	starwer.aSep08		5704	993		208	putative mitochondrial protein (22.4 kD) (starwer) mRNA.
stashaw	stashaw.aSep08		15363	732		94	putative protein of vertebrate origin (stashaw) mRNA.
Stat1	Stat1.cSep08	25124	15810	771	7	180	signal transducer and activator of transcription 1 (Stat1) alternative variant cSep08, mRNA.
Stat1	Stat1.dSep08	25124	4323	785	4	82	signal transducer and activator of transcription 1 (Stat1) alternative variant dSep08, mRNA.
Stat1	Stat1.eSep08	25124	3600	377	2	41	signal transducer and activator of transcription 1 (Stat1) alternative variant eSep08, mRNA.
Stat2	Stat2.aSep08	288774	16918	4084	24	842	signal transducer and activator of transcription 2 (96.9 kD) (Stat2) alternative variant aSep08, mRNA.
Stat2	Stat2.bSep08	288774	1277	1080	2	117	signal transducer and activator of transcription 2 (12.9 kD) (Stat2) alternative variant bSep08, mRNA.
Stat3	Stat3.bSep08	25125	4805	397	5	132	signal transducer and activator of transcription 3 (Stat3) alternative variant bSep08, mRNA.
Stat3	Stat3.cSep08	25125	4679	912	4	110	signal transducer and activator of transcription 3 (Stat3) alternative variant cSep08, mRNA.
Stat3	Stat3.dSep08	25125	1003	710	3	69	signal transducer and activator of transcription 3 (Stat3) alternative variant dSep08, mRNA.
Stat3	Stat3.eSep08	25125	4743	673	4	38	signal transducer and activator of transcription 3 (Stat3) alternative variant eSep08, mRNA.
Stat5a	Stat5a.aSep08	24918	5552	1928	7	246	signal transducer and activator of transcription 5A (Stat5a) alternative variant aSep08, mRNA.
Stat6	Stat6.bSep08	362896	2496	376	5	125	signal transducer and activator of transcription 6 (Stat6) alternative variant bSep08, mRNA.
statu	statu.aSep08		5569	404		55	putative protein (statu) mRNA.
STAT_alpha.0	STAT_alpha.0.aSep08		10471	608		202	activator of transcription signal transducer 5 (STAT_alpha.0) mRNA.
Stau1	Stau1.cSep08	84496	4954	877	5	292	stau RNA binding protein homolog 1 (Drosophila) (Stau1) alternative variant cSep08, mRNA.
Stau1	Stau1.dSep08	84496	3152	696	6	216	stau RNA binding protein homolog 1 (Drosophila) (Stau1) alternative variant dSep08, mRNA.
stavo	stavo.aSep08		3160	608		69	putative protein (7.7 kD) (stavo) mRNA.
stawcha	stawcha.aSep08		4337	819		215	nucleoporin 214kDa (stawcha) mRNA.
stawchy	stawchy.aSep08		851	595	1	39	putative protein (stawchy) alternative variant aSep08, mRNA.
stawchy	stawchy.bSep08		39510	419	2	35	putative protein (stawchy) alternative variant bSep08, mRNA.
stawer	stawer.aSep08		71692	463		26	putative protein (stawer) mRNA.
stawfee	stawfee.aSep08		33545	712	2	104	putative nuclear protein (11.4 kD) (stawfee) alternative variant aSep08, mRNA.
stawfee	stawfee.bSep08		13196	1035	4	11	putative protein (stawfee) alternative variant bSep08, mRNA.
stawfee	stawfee.cSep08		1369	757	2	49	putative protein (5.5 kD) (stawfee) alternative variant cSep08, mRNA.
stawfee	stawfee.dSep08		1388	745	2	49	putative protein (5.5 kD) (stawfee) alternative variant dSep08, mRNA.



stawflu	stawflu.bSep08		736	354	2	64	putative protein (stawflu) alternative variant bSep08, mRNA.
stawflu	stawflu.cSep08		482	344	2	18	putative protein (2.0 kD) (stawflu) alternative variant cSep08, mRNA.
stawfly	stawfly.aSep08		3209	703		40	putative protein (4.0 kD) (stawfly) mRNA.
stawlor	stawlor.aSep08		10711	1917	4	88	putative protein (9.7 kD) (stawlor) alternative variant aSep08, mRNA.
stawmee	stawmee.aSep08		8562	509	4	144	putative protein (stawmee) alternative variant aSep08, mRNA.
stawmee	stawmee.bSep08		3672	984	3	65	putative protein (7.5 kD) (stawmee) alternative variant bSep08, mRNA.
stawmee	stawmee.dSep08		3346	372	2	51	putative protein (stawmee) alternative variant dSep08, mRNA.
stawnor	stawnor.aSep08		1464	321		33	putative protein (3.6 kD) (stawnor) mRNA.
stawpey	stawpey.aSep08		4815	828		33	putative protein (stawpey) mRNA.
stawroy	stawroy.aSep08		7615	662		92	putative cytoplasmic protein (10.0 kD) (stawroy) mRNA.
stawshaw	stawshaw.aSep08		2024	396		92	putative protein (stawshaw) mRNA.
stawtu	stawtu.aSep08		7710	400		87	putative protein (stawtu) mRNA.
stawvo	stawvo.aSep08		2918	418		51	putative protein (5.8 kD) (stawvo) mRNA.
stawwer	stawwer.aSep08		19290	1775		526	putative protein of eukaryotic origin (stawwer) alternative variant aSep08, mRNA.
Steap2	Steap2.cSep08	312052	10478	646	3	12	six transmembrane epithelial antigen of the prostate 2 (1.3 kD) (Steap2) alternative variant cSep08, mRNA.
Steap2	Steap2.dSep08	312052	1865	501	2	31	six transmembrane epithelial antigen of the prostate 2 (Steap2) alternative variant dSep08, mRNA.
Steap3	Steap3.bSep08	170824	25401	912	1	282	STEAP family member 3 (Steap3) alternative variant bSep08, mRNA.
Steap3	Steap3.cSep08	170824	18736	735	1	220	STEAP family member 3 (Steap3) alternative variant cSep08, mRNA.
Steap4	Steap4.bSep08	499991	3012	928	3	309	STEAP family member 4 (Steap4) alternative variant bSep08, mRNA.
steecha	steecha.aSep08		2797	228		75	nucleoporin 214kDa CRA b (steecha) mRNA.
steechy	steechy.aSep08		1943	587		59	putative protein of mammalian origin (6.5 kD) (steechy) mRNA.
steefee	steefee.aSep08		9104	777		113	putative nuclear protein (13.2 kD) (steefee) mRNA.
steeflu	steeflu.aSep08		1468	541		54	putative protein (5.8 kD) (steeflu) mRNA.
steefly	steefly.aSep08		1563	235		56	putative protein (steefly) mRNA.
steelor	steelor.aSep08		6180	416		27	putative protein (3.1 kD) (steelor) mRNA.
steemee	steemee.aSep08		14617	1505	4	223	putative protein (24.5 kD) (steemee) alternative variant aSep08, mRNA.
steemee	steemee.cSep08		1453	1078	2	75	putative protein (9.2 kD) (steemee) alternative variant cSep08, mRNA.
steenor	steenor.aSep08		673	356		43	putative protein (steenor) mRNA.
steepcy	steepcy.aSep08		4473	1216	3	104	BCSC-1 like (steepcy) alternative variant aSep08, mRNA.

steepcy	steepcy.bSep08		1028	516	1	59	BCSC-1 like (6.9 kD) (steepcy) alternative variant bSep08, mRNA.
steeroy	steeroy.aSep08		844	741		59	putative protein (6.8 kD) (steeroy) mRNA.
steeshaw	steeshaw.aSep08		5881	537		100	putative protein (steeshaw) mRNA.
steetu	steetu.aSep08		1631	492	2	54	putative protein (6.0 kD) (steetu) alternative variant aSep08, mRNA.
steevo	steevo.aSep08		5075	738		68	putative protein (7.5 kD) (steevo) mRNA.
steewer	steewer.aSep08		15147	1037		299	CRA d (steewer) mRNA.
stercha	stercha.aSep08		1424	402		133	nucleoporin 214 like (stercha) mRNA.
sterchy	sterchy.aSep08		1369	619		108	CRA d like (11.3 kD) (sterchy) mRNA.
sterfee	sterfee.aSep08		1280	518		39	putative protein (4.5 kD) (sterfee) mRNA.
sterflu	sterflu.aSep08		3284	1000		85	calpain 1 (sterflu) mRNA.
sterfly	sterfly.aSep08		1621	1533		208	putative protein of vertebrate origin (sterfly) mRNA.
sterlor	sterlor.aSep08		66857	298		79	intersectin 1 ITSN-1 (sterlor) mRNA.
stermee	stermee.aSep08		2705	783	4	75	putative protein (7.9 kD) (stermee) alternative variant aSep08, mRNA.
stermee	stermee.bSep08		2123	641	3	75	putative protein (7.9 kD) (stermee) alternative variant bSep08, mRNA.
sternor	sternor.aSep08		2249	335	2	65	putative protein (sternor) alternative variant aSep08, mRNA.
sterpey	sterpey.aSep08		3328	398		132	putative protein of eukaryotic origin (sterpey) mRNA.
sterroy	sterroy.aSep08		475	292		83	putative protein (sterroy) mRNA.
stershaw	stershaw.bSep08		1376	611	3	45	putative protein (5.3 kD) (stershaw) alternative variant bSep08, mRNA.
stertu	stertu.bSep08		2340	1292	2	75	putative protein (stertu) alternative variant bSep08, mRNA.
stertu	stertu.cSep08		1194	422	2	28	putative protein (stertu) alternative variant cSep08, mRNA.
stervo	stervo.aSep08		15774	1168		40	putative protein (4.7 kD) (stervo) mRNA.
sterwer	sterwer.aSep08		7070	700	6	232	putative protein of metazoan origin (sterwer) alternative variant aSep08, mRNA.
steycha	steycha.aSep08		12546	938		312	nucleoporin 214kDa (steycha) mRNA.
steychy	steychy.aSep08		57455	738		79	putative cytoplasmic protein (9.0 kD) (steychy) mRNA.
steyfee	steyfee.aSep08		3264	513		61	putative protein (6.8 kD) (steyfee) mRNA.
steyflu	steyflu.aSep08		967	514		36	putative protein (steyflu) mRNA.
steyfly	steyfly.aSep08		1515	506		80	cytochrome p450 (steyfly) mRNA.
steylor	steylor.aSep08		1191	306		31	putative protein (3.6 kD) (steylor) mRNA.
steymee	steymee.aSep08		13155	361		110	putative protein (steymee) mRNA.
steynor	steynor.aSep08		23443	915		51	putative protein (6.0 kD) (steynor) mRNA.
steypey	steypey.aSep08		3330	937		99	putative nuclear protein (11.1 kD) (steypey) mRNA.
steyroy	steyroy.aSep08		49170	501	1	60	CRA a like (6.9 kD) (steyroy) alternative variant aSep08, mRNA.
steyroy	steyroy.bSep08		60010	477	1	77	CRA b like (8.4 kD) (steyroy) alternative variant bSep08, mRNA.
steyslaw	steyslaw.aSep08		4659	704		76	putative protein (8.1 kD) (steyslaw) mRNA.

steytu	steytu.aSep08		27319	793		160	putative protein of vertebrate origin (steytu) mRNA.
steyvo	steyvo.aSep08		6413	358		33	putative protein (steyvo) mRNA.
steywer	steywer.aSep08		8701	653		164	family with sequence similarity 13 member A1 (steywer) mRNA.
Stfa2l2	Stfa2l2.aSep08	288070	2544	356		93	stefin A2-like 2 (Stfa2l2) mRNA.
Stfa2l3	Stfa2l3.aSep08	498087	2305	330		85	stefin A2-like 3 (Stfa2l3) mRNA.
Stil	Stil.aSep08	313506	4801	410		136	scl/Tal1 interrupting locus (Stil) mRNA.
Stim1	Stim1.bSep08	361618	154222	2523	7	413	stromal interaction molecule 1 (48.1 kD) (Stim1) alternative variant bSep08, mRNA.
Stim1	Stim1.cSep08	361618	9022	669	4	223	stromal interaction molecule 1 (Stim1) alternative variant cSep08, mRNA.
Stim2	Stim2.bSep08	117087	1093	643	2	43	stromal interaction molecule 2 (Stim2) alternative variant bSep08, mRNA.
Stip1	Stip1.bSep08	192277	1251	392	1	97	stress-induced phosphoprotein 1 (Stip1) alternative variant bSep08, mRNA.
Stk10	Stk10.aSep08	29398	9162	1196		275	serine/threonine kinase 10 (Stk10) alternative variant aSep08, mRNA.
Stk10	Stk10.bSep08	29398	23675	965	1	271	serine/threonine kinase 10 (Stk10) alternative variant bSep08, mRNA.
Stk10	Stk10.cSep08	29398	21779	773	2	257	serine/threonine kinase 10 (Stk10) alternative variant cSep08, mRNA.
Stk11	Stk11.bSep08	314621	2538	745	6	231	serine/threonine kinase 11 (Stk11) alternative variant bSep08, mRNA.
Stk11ip	Stk11ip.bSep08	301535	2355	1148	3	154	serine/threonine kinase 11 interacting protein (Stk11ip) alternative variant bSep08, mRNA.
Stk11ip	Stk11ip.cSep08	301535	767	344	4	114	serine/threonine kinase 11 interacting protein (Stk11ip) alternative variant cSep08, mRNA.
Stk16	Stk16.bSep08	286927	9278	1909	8	273	serine threonine kinase 16 (30.6 kD) (Stk16) alternative variant bSep08, mRNA.
Stk16	Stk16.cSep08	286927	2076	896	5	205	serine threonine kinase 16 (23.4 kD) (Stk16) alternative variant cSep08, mRNA.
Stk16	Stk16.eSep08	286927	922	736	2	129	putative protein human specific (Stk16) alternative variant eSep08, mRNA.
Stk16	Stk16.fSep08	286927	936	827	2	78	serine threonine kinase 16 (8.5 kD) (Stk16) alternative variant fSep08, mRNA.
Stk19	Stk19.bSep08	361800	7310	734	1	158	serine/threonine kinase 19 (17.7 kD) (Stk19) alternative variant bSep08, mRNA.
Stk22s1	Stk22s1.bSep08	292890	9127	719	6	220	serine/threonine kinase 22 substrate 1 (Stk22s1) alternative variant bSep08, mRNA.
Stk22s1	Stk22s1.cSep08	292890	3492	801	4	152	serine/threonine kinase 22 substrate 1 (15.0 kD) (Stk22s1) alternative variant cSep08, mRNA.
Stk22s1	Stk22s1.dSep08	292890	1482	434	3	112	serine/threonine kinase 22 substrate 1 (Stk22s1) alternative variant dSep08, mRNA.
Stk22s1	Stk22s1.eSep08	292890	824	689	2	108	serine/threonine kinase 22 substrate 1 (11.0 kD) (Stk22s1) alternative variant eSep08, mRNA.

Stk24	Stk24.bSep08	361092	15430	3929	2	207	serine/threonine kinase 24 (STE20 homolog, yeast) (23.1 kD) (Stk24) alternative variant bSep08, mRNA.
Stk25	Stk25.bSep08	373542	9234	2198	6	154	kinase (17.6 kD) (Stk25) alternative variant bSep08, mRNA.
Stk25	Stk25.cSep08	373542	1361	941	3	88	kinase (9.9 kD) (Stk25) alternative variant cSep08, mRNA.
Stk25	Stk25.eSep08	373542	2593	912	3	61	putative protein (6.7 kD) (Stk25) alternative variant eSep08, mRNA.
Stk32a	Stk32a.aSep08	364858	55419	645		134	serine/threonine kinase 32A (Stk32a) mRNA.
Stk33	Stk33.aSep08	690861	986	218		72	serine/threonine kinase 33 (Stk33) mRNA.
Stk36	Stk36.aSep08	301516	3706	2010		393	serine/threonine kinase 36 (fused homolog, Drosophila) (Stk36) mRNA.
Stk38	Stk38.aSep08	361813	10034	2084	7	318	serine/threonine kinase 38 (37.0 kD) (Stk38) alternative variant aSep08, mRNA.
Stk38	Stk38.bSep08	361813	2346	1124	2	209	serine/threonine kinase 38 (23.8 kD) (Stk38) alternative variant bSep08, mRNA.
Stk39	Stk39.bSep08	54348	162515	1525	5	168	serine/threonine kinase 39, STE20/SPS1 homolog (yeast) (17.4 kD) (Stk39) alternative variant bSep08, mRNA.
Stk40	Stk40.bSep08	360230	4561	997	2	107	serine/threonine kinase 40 (Stk40) alternative variant bSep08, mRNA.
Stmn1	Stmn1.aSep08	29332	5750	1020	5	149	stathmin 1 (17.3 kD) (Stmn1) alternative variant aSep08, mRNA.
Stmn1	Stmn1.cSep08	29332	3017	872	3	92	stathmin 1 (10.8 kD) (Stmn1) alternative variant cSep08, mRNA.
Stmn3	Stmn3.aSep08	29246	7982	790	2	213	stathmin-like 3 (Stmn3) alternative variant aSep08, mRNA.
Stmn3	Stmn3.bSep08	29246	8106	1015	3	180	stathmin-like 3 (21.0 kD) (Stmn3) alternative variant bSep08, complete mRNA.
Stmn3	Stmn3.dSep08	29246	7317	591	1	168	stathmin-like 3 (Stmn3) alternative variant dSep08, mRNA.
Stmn4	Stmn4.aSep08	79423	18656	1328	7	216	stathmin-like 4 (25.4 kD) (Stmn4) alternative variant aSep08, mRNA.
Stmn4	Stmn4.bSep08	79423	18445	1036	6	189	stathmin-like 4 (22.1 kD) (Stmn4) alternative variant bSep08, mRNA.
Stmn4	Stmn4.cSep08	79423	18181	769	6	188	stathmin-like 4 (22.0 kD) (Stmn4) alternative variant cSep08, mRNA.
Stmn4	Stmn4.dSep08	79423	13952	610	5	99	stathmin-like 4 (Stmn4) alternative variant dSep08, mRNA.
stocha	stocha.aSep08		1385	1065		86	putative protein (stocha) mRNA.
stochy	stochy.aSep08		2407	405		37	putative protein (stochy) mRNA.
stofee	stofee.aSep08		5405	856		60	putative protein (6.5 kD) (stofee) mRNA.
stoflu	stoflu.aSep08		998	458		61	putative protein (stoflu) mRNA.
stofly	stofly.aSep08		5655	658	4	171	CRA c (stofly) alternative variant aSep08, mRNA.
stofly	stofly.bSep08		1990	663	2	113	CRA c (stofly) alternative variant bSep08, mRNA.
stofly	stofly.dSep08		1503	392	2	71	CRA d like (stofly) alternative variant dSep08, mRNA.
stolor	stolor.aSep08		596	397		39	putative protein (stolor) mRNA.
stomee	stomee.aSep08		82071	365		94	carbonic anhydrase (stomee) mRNA.
Stoml1	Stoml1.aSep08	300748	4137	1467	4	248	stomatin-like 1 (Stoml1) alternative variant aSep08, mRNA.

Stoml2	Stoml2.bSep08	298203	2360	854	8	233	stomatin (Epb7.2)-like 2 (Stoml2) alternative variant bSep08, mRNA.
Stoml2	Stoml2.cSep08	298203	1849	1256	5	91	stomatin (Epb7.2)-like 2 (Stoml2) alternative variant cSep08, mRNA.
Stoml2	Stoml2.dSep08	298203	1303	438	2	46	stomatin (Epb7.2)-like 2 (4.8 kD) (Stoml2) alternative variant dSep08, mRNA.
Ston2	Ston2.aSep08	314349	44889	928		187	stonin 2 (20.4 kD) (Ston2) mRNA.
stonor	stonor.aSep08		6802	746		83	putative protein (9.2 kD) (stonor) mRNA.
stopey	stopey.aSep08		8618	3214		324	CRA b (37.3 kD) (stopey) mRNA.
storcha	storcha.aSep08		23598	1014	8	337	putative protein (storcha) alternative variant aSep08, mRNA.
storchy	storchy.aSep08		2642	224		64	putative protein (storchy) mRNA.
storfee	storfee.aSep08		29707	298		18	putative protein (storfee) mRNA.
storflu	storflu.aSep08		14029	1131	3	55	putative protein (6.2 kD) (storflu) alternative variant aSep08, mRNA.
storfly	storfly.aSep08		1175	370		70	putative protein (storfly) alternative variant aSep08, mRNA.
storlor	storlor.aSep08		52873	782	2	160	mitochondrial ribosomal protein S6 (storlor) alternative variant aSep08, mRNA.
storlor	storlor.bSep08		18192	387	1	50	mitochondrial ribosomal protein S6 like (storlor) alternative variant bSep08, mRNA.
stormee	stormee.aSep08		2629	566		63	putative protein (7.1 kD) (stormee) mRNA.
stornor	stornor.aSep08		728	529		53	putative protein (5.9 kD) (stornor) mRNA.
storoy	storoy.aSep08		850	334		21	putative protein (storoy) mRNA.
storpey	storpey.aSep08		9332	404		134	putative protein of bilateral origin (storpey) mRNA.
storroy	storroy.aSep08		14050	739		55	putative protein (storroy) mRNA.
storshaw	storshaw.bSep08		4504	1184	4	65	putative protein (7.3 kD) (storshaw) alternative variant bSep08, mRNA.
storshaw	storshaw.cSep08		4437	506	3	40	putative protein (storshaw) alternative variant cSep08, mRNA.
stortu	stortu.aSep08		741	580		89	putative protein (10.2 kD) (stortu) mRNA.
storvo	storvo.aSep08		9517	668		222	InaD-like (storvo) mRNA.
storwer	storwer.aSep08		580	422		82	putative protein (storwer) mRNA.
stoshaw	stoshaw.aSep08		13422	736		59	putative protein (stoshaw) mRNA.
stotu	stotu.aSep08		1895	558		185	dicer1 (stotu) mRNA.
stovo	stovo.aSep08		7326	966	1	99	putative cytoplasmic protein (10.5 kD) (stovo) alternative variant aSep08, mRNA.
stovo	stovo.bSep08		7960	690	2	68	putative protein (stovo) alternative variant bSep08, mRNA.
stower	stower.bSep08		16336	1776	4	76	putative protein (8.2 kD) (stower) alternative variant bSep08, mRNA.
stoycha	stoycha.aSep08		926	452		93	putative protein (stoycha) mRNA.
stoychy	stoychy.aSep08		476	434		70	putative protein (stoychy) mRNA.
stoyfee	stoyfee.aSep08		7994	899		151	putative protein (stoyfee) mRNA.
stoyflu	stoyflu.aSep08		3958	683		141	putative protein (stoyflu) mRNA.

stoyfly	stoyfly.aSep08		5036	552		123	latent transforming growth factor beta binding protein 4 like (stoyfly) mRNA.
stoylor	stoylor.aSep08		2967	239		35	putative protein (stoylor) mRNA.
stoymee	stoymee.aSep08		3700	485	4	161	CRA a like (stoymee) alternative variant aSep08, mRNA.
stoymee	stoymee.bSep08		4024	604	2	110	CRA a like (stoymee) alternative variant bSep08, mRNA.
stoymee	stoymee.cSep08		4074	531	2	46	CRA a like (5.1 kD) (stoymee) alternative variant cSep08, mRNA.
stoynor	stoynor.aSep08		6600	296		33	putative protein (stoynor) mRNA.
stoypey	stoypey.aSep08		3729	474		90	putative protein (9.9 kD) (stoypey) mRNA.
stoyroy	stoyroy.aSep08		793	259		34	putative protein (stoyroy) mRNA.
stoyshaw	stoyshaw.aSep08		3534	647		83	putative nuclear protein (9.6 kD) (stoyshaw) mRNA.
stoytu	stoytu.aSep08		6110	677	3	106	putative protein (stoytu) alternative variant aSep08, mRNA.
stoyvo	stoyvo.aSep08		2096	299		99	dedicator of cytokinesis 7 (stoyvo) mRNA.
stoywer	stoywer.aSep08		4379	802		54	putative protein (stoywer) mRNA.
Stra6	Stra6.cSep08	363071	5738	410	5	98	stimulated by retinoic acid gene 6 (Stra6) alternative variant cSep08, mRNA.
Strap	Strap.bSep08	297699	2810	404	3	87	serine/threonine kinase receptor associated protein (Strap) alternative variant bSep08, mRNA.
Strbp	Strbp.bSep08	84476	21986	547	3	121	spermatid perinuclear RNA binding protein (Strbp) alternative variant bSep08, mRNA.
Strbp	Strbp.cSep08	84476	27997	737	5	109	spermatid perinuclear RNA binding protein (11.8 kD) (Strbp) alternative variant cSep08, mRNA.
Strbp	Strbp.dSep08	84476	11706	694	2	85	spermatid perinuclear RNA binding protein (Strbp) alternative variant dSep08, mRNA.
Strn3	Strn3.bSep08	114520	42470	1852	8	445	striatin, calmodulin binding protein 3 (Strn3) alternative variant bSep08, mRNA.
Strn3	Strn3.cSep08	114520	1125	1045	2	83	striatin, calmodulin binding protein 3 (9.0 kD) (Strn3) alternative variant cSep08, mRNA.
Strn4	Strn4.bSep08	308392	19641	2735	17	655	zinedin (Strn4) alternative variant bSep08, mRNA.
Strn4	Strn4.cSep08	308392	8587	2773	8	374	zinedin (40.3 kD) (Strn4) alternative variant cSep08, mRNA.
Strn4	Strn4.dSep08	308392	8062	1143	4	279	zinedin (29.4 kD) (Strn4) alternative variant dSep08, mRNA.
Strn4	Strn4.eSep08	308392	2224	979	3	124	zinedin (Strn4) alternative variant eSep08, mRNA.
Strn4	Strn4.gSep08	308392	3670	237	2	38	zinedin (Strn4) alternative variant gSep08, mRNA.
Strumpellin.0	Strumpellin.0.aSep08		28081	5502	20	776	CRA a (Strumpellin.0) alternative variant aSep08, mRNA.
Sts	Sts.bSep08	24800	4031	1185	6	190	steroid sulfatase (Sts) alternative variant bSep08, mRNA.
Sts	Sts.cSep08	24800	1601	748	2	174	steroid sulfatase (Sts) alternative variant cSep08, mRNA.
Sts	Sts.dSep08	24800	1789	766	3	151	steroid sulfatase (Sts) alternative variant dSep08, mRNA.
Stub1	Stub1.bSep08	287155	2852	2183	4	224	STIP1 homology and U-Box containing protein 1 (25.4 kD) (Stub1) alternative variant bSep08, mRNA.
Stub1	Stub1.cSep08	287155	1771	1519	4	192	STIP1 homology and U-Box containing protein 1 (21.0 kD) (Stub1) alternative variant cSep08, mRNA.
Stub1	Stub1.dSep08	287155	1284	608	4	191	STIP1 homology and U-Box containing protein 1 (Stub1) alternative variant dSep08, mRNA.

Stub1	Stub1.eSep08	287155	768	685	2	131	STIP1 homology and U-Box containing protein 1 (Stub1) alternative variant eSep08, mRNA.
stucha	stucha.aSep08		2899	772		37	putative protein (stucha) mRNA.
stuchy	stuchy.aSep08		2165	1062		90	putative protein (10.3 kD) (stuchy) alternative variant aSep08, mRNA.
stufee	stufee.aSep08		836	749		40	putative protein (stufee) mRNA.
stufly	stufly.aSep08		885	380		126	pecanex-like 3 CRA c (stufly) mRNA.
stufly	stufly.aSep08		112960	744	7	95	putative protein (10.9 kD) (stufly) alternative variant aSep08, mRNA.
stufly	stufly.bSep08		61488	355	3	46	putative protein (stufly) alternative variant bSep08, mRNA.
stulor	stulor.aSep08		4858	274		43	putative protein (stulor) mRNA.
stumee	stumee.aSep08		18062	349		65	putative protein (7.2 kD) (stumee) mRNA.
stunor	stunor.aSep08		12490	677		155	transmembrane protein 63b (stunor) mRNA.
stupey	stupey.aSep08		1660	748		151	hepatocyte cell adhesion molecule (stupey) mRNA.
sturoy	sturoy.aSep08		897	371		20	putative protein (2.2 kD) (sturoy) mRNA.
stushaw	stushaw.aSep08		9790	780	6	232	DEAH box polypeptide 29 (stushaw) alternative variant aSep08, mRNA.
stutu	stutu.aSep08		689	354		60	putative protein (6.6 kD) (stutu) mRNA.
stuvo	stuvo.aSep08		5196	693		60	putative protein (7.1 kD) (stuvo) mRNA.
stuwcr	stuwcr.aSep08		1710	1365		199	gag-pro-pol polyprotein (stuwcr) mRNA.
Stx1a	Stx1a.cSep08	116470	1541	712	2	60	syntaxin 1A (brain) (6.7 kD) (Stx1a) alternative variant cSep08, mRNA.
Stx2	Stx2.bSep08	25130	32915	1019	11	259	epimorphin (29.9 kD) (Stx2) alternative variant bSep08, mRNA.
Stx2	Stx2.cSep08	25130	6238	642	7	106	epimorphin CRA e (12.3 kD) (Stx2) alternative variant cSep08, mRNA.
Stx2	Stx2.eSep08	25130	10917	262	3	79	putative protein (Stx2) alternative variant eSep08, mRNA.
Stx2	Stx2.fSep08	25130	6510	1378	3	66	epimorphin CRA e (Stx2) alternative variant fSep08, mRNA.
Stx3	Stx3.bSep08	81802	12053	648	4	109	syntaxin 3 (Stx3) alternative variant bSep08, mRNA.
Stx3	Stx3.cSep08	81802	44569	1041	2	65	syntaxin 3 (Stx3) alternative variant cSep08, mRNA.
Stx4a	Stx4a.aSep08	81803	7455	1238	10	348	syntaxin 4A (placental) (Stx4a) alternative variant aSep08, mRNA.
Stx4a	Stx4a.cSep08	81803	5804	1336	6	192	syntaxin 4A (placental) (22.4 kD) (Stx4a) alternative variant cSep08, mRNA.
Stx4a	Stx4a.eSep08	81803	6031	1381	7	141	syntaxin 4A (placental) (16.5 kD) (Stx4a) alternative variant eSep08, mRNA.
Stx4a	Stx4a.fSep08	81803	5102	778	7	133	syntaxin 4A (placental) (15.4 kD) (Stx4a) alternative variant fSep08, mRNA.
Stx4a	Stx4a.gSep08	81803	1522	1025	3	71	syntaxin 4A (placental) (Stx4a) alternative variant gSep08, mRNA.
Stx4a	Stx4a.hSep08	81803	1834	547	3	69	syntaxin 4A (placental) (Stx4a) alternative variant hSep08, mRNA.
Stx5a	Stx5a.bSep08	65134	1477	793	4	118	syntaxin 5a (Stx5a) alternative variant bSep08, mRNA.
Stx5a	Stx5a.cSep08	65134	4102	638	1	108	syntaxin 5a (Stx5a) alternative variant cSep08, mRNA.

Stx7	Stx7.bSep08	60466	36213	815	6	222	syntaxin 7 (Stx7) alternative variant bSep08, mRNA.
Stx7	Stx7.cSep08	60466	28413	303	2	77	syntaxin 7 (Stx7) alternative variant cSep08, mRNA.
Stx8	Stx8.aSep08	59074	111484	1779	3	574	syntaxin 8 (Stx8) alternative variant aSep08, mRNA.
Stx8	Stx8.cSep08	59074	225232	606	3	165	syntaxin 8 (Stx8) alternative variant cSep08, mRNA.
Stx8	Stx8.dSep08	59074	146031	375	2	54	syntaxin 8 (Stx8) alternative variant dSep08, mRNA.
Stx12	Stx12.aSep08	65033	28561	2632	8	274	syntaxin 12 (31.2 kD) (Stx12) alternative variant aSep08, mRNA.
Stx12	Stx12.bSep08	65033	2574	591	1	38	syntaxin 12 (4.6 kD) (Stx12) alternative variant bSep08, mRNA.
Stx16	Stx16.aSep08	362283	23928	1752	7	328	syntaxin 16 (37.3 kD) (Stx16) alternative variant aSep08, mRNA.
Stx16	Stx16.bSep08	362283	8736	3279	6	221	syntaxin 16 (Stx16) alternative variant bSep08, mRNA.
Stx16	Stx16.dSep08	362283	6767	1403	7	139	syntaxin 16 (16.0 kD) (Stx16) alternative variant dSep08, mRNA.
Stx18	Stx18.bSep08	360953	25850	1342	9	252	syntaxin 18 (Stx18) alternative variant bSep08, mRNA.
Stx18	Stx18.cSep08	360953	1513	683	2	102	syntaxin 18 (11.5 kD) (Stx18) alternative variant cSep08, mRNA.
Stxbp1	Stxbp1.bSep08	25558	7326	1482	5	174	syntaxin binding protein 1 (Stxbp1) alternative variant bSep08, mRNA.
Stxbp1	Stxbp1.cSep08	25558	8181	744	4	111	syntaxin binding protein 1 (Stxbp1) alternative variant cSep08, mRNA.
Stxbp2	Stxbp2.aSep08	81804	9694	2000	17	543	syntaxin binding protein 2 (Stxbp2) alternative variant aSep08, mRNA.
Stxbp2	Stxbp2.bSep08	81804	3751	720	7	230	syntaxin binding protein 2 (Stxbp2) alternative variant bSep08, mRNA.
Stxbp2	Stxbp2.cSep08	81804	7210	667	4	200	syntaxin binding protein 2 (Stxbp2) alternative variant cSep08, mRNA.
Stxbp2	Stxbp2.dSep08	81804	3538	425	6	110	syntaxin binding protein 2 (Stxbp2) alternative variant dSep08, mRNA.
Stxbp2	Stxbp2.eSep08	81804	14936	908	3	86	syntaxin binding protein 2 (9.5 kD) (Stxbp2) alternative variant eSep08, mRNA.
Stxbp3	Stxbp3.bSep08	114095	3371	479	4	118	syntaxin binding protein 3 like (Stxbp3) alternative variant bSep08, mRNA.
Stxbp3	Stxbp3.cSep08	114095	10135	671	2	37	syntaxin binding protein 3 like (4.3 kD) (Stxbp3) alternative variant cSep08, mRNA.
Stxbp4	Stxbp4.bSep08	303443	39295	334	4	111	syntaxin binding protein 4 (Stxbp4) alternative variant bSep08, mRNA.
Stxbp5	Stxbp5.dSep08	81022	10636	2472	5	291	syntaxin binding protein 5 like (32.1 kD) (Stxbp5) alternative variant dSep08, mRNA.
Stxbp5	Stxbp5.eSep08	81022	18518	680	5	204	syntaxin binding protein 5 like (Stxbp5) alternative variant eSep08, mRNA.
Stxbp5	Stxbp5.fSep08	81022	7695	774	2	52	syntaxin binding protein 5 like (Stxbp5) alternative variant fSep08, mRNA.
Stxbp6	Stxbp6.aSep08	362734	240663	1437		308	syntaxin binding protein 6 (amisyn) (Stxbp6) alternative variant aSep08, mRNA.



Stxbp6	Stxbp6.bSep08	362734	200336	551	1	150	syntaxin binding protein 6 (amisyn) (Stxbp6) alternative variant bSep08, mRNA.
stycha	stycha.aSep08		5562	643		65	putative protein (stycha) mRNA.
stychy	stychy.aSep08		3199	464		99	putative protein (11.2 kD) (stychy) mRNA.
styfee	styfee.aSep08		89484	438		63	jumonji AT rich interactive domain 2 CRA b (styfee) mRNA.
styflu	styflu.aSep08		581	432		54	putative protein (styflu) mRNA.
styfly	styfly.aSep08		492	408		116	multiple -like domain (styfly) mRNA.
stylor	stylor.aSep08		724	556		92	putative protein (stylor) mRNA.
stymee	stymee.aSep08		47537	269		23	putative protein (stymee) mRNA.
stynor	stynor.aSep08		17608	326	4	62	putative protein (stynor) alternative variant aSep08, mRNA.
stynor	stynor.bSep08		18956	360	3	62	tight junction associated protein 1 (stynor) alternative variant bSep08, mRNA.
stynor	stynor.cSep08		18570	529	3	39	putative protein (4.3 kD) (stynor) alternative variant cSep08, mRNA.
stynor	stynor.dSep08		19496	480	5	57	putative protein (stynor) alternative variant dSep08, mRNA.
stypey	stypey.aSep08		9409	251		69	pbx knotted 1 homeobox 2 (stypey) mRNA.
styroy	styroy.aSep08		22811	668	7	222	putative protein of mammalian origin (styroy) alternative variant aSep08, mRNA.
styshaw	styshaw.aSep08		4024	1728			
stytu	stytu.aSep08		946	330		75	putative protein (stytu) mRNA.
styvo	styvo.aSep08		7632	274		90	myb-like SWIRM MPN domains 1 (styvo) mRNA.
stywer	stywer.aSep08		470	352		30	putative protein (stywer) mRNA.
Styxl1	Styxl1.aSep08	360792	32717	1788	8	451	serine/threonine/tyrosine interacting-like 1 (Styxl1) alternative variant aSep08, mRNA.
Styxl1	Styxl1.cSep08	360792	20783	637	6	164	serine/threonine/tyrosine interacting-like 1 (19.4 kD) (Styxl1) alternative variant cSep08, mRNA.
Styxl1	Styxl1.dSep08	360792	20154	577	5	132	serine/threonine/tyrosine interacting-like 1 (Styxl1) alternative variant dSep08, mRNA.
Sub1	Sub1.bSep08	192269	14767	3354	5	127	SUB1 homolog (S. cerevisiae) (14.4 kD) (Sub1) alternative variant bSep08, complete mRNA.
suby	suby.aSep08		4906	415		138	putative protein of eukaryotic origin (suby) mRNA.
suchy	suchy.aSep08		43143	851		48	putative protein (5.2 kD) (suchy) mRNA.
Sucla2	Sucla2.bSep08	361071	27563	1524	7	236	succinate-Coenzyme A ligase, ADP-forming, beta subunit (25.6 kD) (Sucla2) alternative variant bSep08, mRNA.
Suclg1	Suclg1.bSep08	114597	8855	503	4	167	succinate-CoA ligase, GDP-forming, alpha subunit (Suclg1) alternative variant bSep08, mRNA.
Suclg1	Suclg1.cSep08	114597	1764	427	2	76	succinate-CoA ligase, GDP-forming, alpha subunit (Suclg1) alternative variant cSep08, mRNA.
Suclg1	Suclg1.dSep08	114597	8999	621	3	60	succinate-CoA ligase, GDP-forming, alpha subunit (6.5 kD) (Suclg1) alternative variant dSep08, complete mRNA.
Suclg2	Suclg2.aSep08	362404	158722	1895	8	323	succinyl-CoA synthetase (Suclg2) alternative variant aSep08, mRNA.
Suclg2	Suclg2.bSep08	362404	166727	856	7	214	succinyl-CoA synthetase (Suclg2) alternative variant bSep08, mRNA.

Suc1g2	Suc1g2.dSep08	362404	156425	921	6	116	succinyl-CoA synthetase (Suc1g2) alternative variant dSep08, mRNA.
Suc1g2	Suc1g2.eSep08	362404	10977	800	2	110	succinyl-CoA synthetase (Suc1g2) alternative variant eSep08, mRNA.
Suc1g2	Suc1g2.fSep08	362404	120363	875	3	103	succinyl-CoA synthetase (Suc1g2) alternative variant fSep08, mRNA.
Suc1g2	Suc1g2.gSep08	362404	25828	634	3	79	succinyl-CoA synthetase (Suc1g2) alternative variant gSep08, mRNA.
sudar	sudar.aSep08		1597	1401		129	CRA b like (14.7 kD) (sudar) mRNA.
Suds3	Suds3.aSep08	360819	5629	746	3	132	suppressor of defective silencing 3 homolog ( <i>S. cerevisiae</i> ) (14.9 kD) (Suds3) alternative variant aSep08, mRNA.
Suds3	Suds3.bSep08	360819	7373	1530	6	122	suppressor of defective silencing 3 homolog ( <i>S. cerevisiae</i> ) (Suds3) alternative variant bSep08, mRNA.
Suds3	Suds3.cSep08	360819	6671	917	5	108	suppressor of defective silencing 3 homolog ( <i>S. cerevisiae</i> ) (Suds3) alternative variant cSep08, mRNA.
Suds3	Suds3.dSep08	360819	2364	777	2	142	suppressor of defective silencing 3 homolog ( <i>S. cerevisiae</i> ) (15.7 kD) (Suds3) alternative variant dSep08, mRNA.
sufer	sufer.aSep08		10119	538		179	G protein coupled receptor 158 (sufer) mRNA.
suflo	suflo.aSep08		84008	280		23	putative protein (suflo) mRNA.
suflu	suflu.aSep08		16145	529		138	putative protein (suflu) mRNA.
Sufu	Sufu.bSep08	361769	40402	2984	9	294	suppressor of fused homolog ( <i>Drosophila</i> ) (Sufu) alternative variant bSep08, mRNA.
Sufu	Sufu.cSep08	361769	3586	582	3	89	suppressor of fused homolog ( <i>Drosophila</i> ) (Sufu) alternative variant cSep08, mRNA.
Sufu	Sufu.dSep08	361769	28446	736	2	81	suppressor of fused homolog ( <i>Drosophila</i> ) (Sufu) alternative variant dSep08, mRNA.
sugar	sugar.aSep08		141788	712		33	putative protein (3.8 kD) (sugar) mRNA.
Sugt1	Sugt1.bSep08	290408	30769	742	11	240	SGT1, suppressor of G2 allele of SKP1 ( <i>S. cerevisiae</i> ) (Sugt1) alternative variant bSep08, mRNA.
Sugt1	Sugt1.cSep08	290408	3977	472	2	148	SGT1, suppressor of G2 allele of SKP1 ( <i>S. cerevisiae</i> ) (Sugt1) alternative variant cSep08, mRNA.
suja	suja.aSep08		632	429		57	putative protein (suja) mRNA.
sujey	sujey.aSep08		1673	520		11	putative protein (sujey) mRNA.
sukee	sukee.aSep08		4889	1143	7	380	putative protein of vertebrate origin (sukee) alternative variant aSep08, mRNA.
sukee	sukee.bSep08		5716	2172	5	132	putative protein of vertebrate origin (sukee) alternative variant bSep08, mRNA.
sukler	sukler.aSep08		3951	1941		464	chromatin assembly factor 1 (sukler) mRNA.
Sulf1	Sulf1.bSep08	171396	40060	2023	3	465	sulfatase 1 (Sulf1) alternative variant bSep08, mRNA.
Sulf1	Sulf1.cSep08	171396	2451	718	1	183	sulfatase 1 (Sulf1) alternative variant cSep08, mRNA.
Sulf2	Sulf2.bSep08	311642	101162	3643	21	918	sulfatase 2 (Sulf2) alternative variant bSep08, mRNA.
Sulf2	Sulf2.cSep08	311642	63981	1785	3	548	sulfatase 2 (Sulf2) alternative variant cSep08, mRNA.
Sulf2	Sulf2.dSep08	311642	3868	396	4	132	sulfatase 2 (Sulf2) alternative variant dSep08, mRNA.
Sulf2	Sulf2.eSep08	311642	16273	299	2	61	sulfatase 2 (Sulf2) alternative variant eSep08, mRNA.

Sulfate_transp.0	Sulfate_transp.0.aSep08		9542	1874	16	552	solute carrier family 26 member 6 CRA a (Sulfate_transp.0) alternative variant aSep08, mRNA.
Sulfate_transp.0	Sulfate_transp.0.bSep08		835	448	2	144	transmembrane protein 89 (Sulfate_transp.0) alternative variant bSep08, mRNA.
Sulfate_transp.0	Sulfate_transp.0.cSep08		2459	532	6	110	solute carrier family 26 member 6 CRA e (Sulfate_transp.0) alternative variant cSep08, mRNA.
Sulfate_transp.0	Sulfate_transp.0.dSep08		1611	529	4	91	solute carrier family 26 member 6 CRA a (Sulfate_transp.0) alternative variant dSep08, mRNA.
Sulfotransfer_1.0	Sulfotransfer_1.0.aSep08		14311	790		159	hydroxysteroid sulfotransferase precursor (18.7 kD) (Sulfotransfer_1.0) mRNA.
sulo	sulo.aSep08		4194	572		190	furry homolog (sulo) mRNA.
Sult1a1	Sult1a1.bSep08	83783	2408	792	1	201	sulfotransferase family 1A, phenol-preferring, member 1 (22.3 kD) (Sult1a1) alternative variant bSep08, mRNA.
Sult1c2	Sult1c2.cSep08	171072	21926	802	3	234	sulfotransferase family, cytosolic, 1C, member 2 (Sult1c2) alternative variant cSep08, mRNA.
Sult1c2	Sult1c2.dSep08	171072	21150	669	2	167	sulfotransferase family, cytosolic, 1C, member 2 (Sult1c2) alternative variant dSep08, mRNA.
Sult1c2a	Sult1c2a.aSep08	316153	4368	2025	4	172	sulfotransferase family, cytosolic, 1C, member 2a (Sult1c2a) alternative variant aSep08, mRNA.
Sult1e1andste2	Sult1e1andste2.bSep08	25355	13539	880	1	211	estrogen sulfotransferase (25.1 kD) (Sult1e1andste2) alternative variant bSep08, complete mRNA.
Sult1e1andste2	Sult1e1andste2.bSep08	360268	13539	880	1	211	estrogen sulfotransferase (25.1 kD) (Sult1e1andste2) alternative variant bSep08, complete mRNA.
Sult2a2	Sult2a2.aSep08	361510	75416	1025	6	284	sulfotransferase family 2A, dehydroepiandrosterone (DHEA)-preferring, member 2 (33.2 kD) (Sult2a2) alternative variant aSep08, mRNA.
Sult2a2	Sult2a2.bSep08	361510	61378	635	5	193	sulfotransferase family 2A, dehydroepiandrosterone (DHEA)-preferring, member 2 (Sult2a2) alternative variant bSep08, mRNA.
Sult2b1	Sult2b1.aSep08	292915	29713	1801	2	594	sulfotransferase family, cytosolic, 2B, member 1 (Sult2b1) alternative variant aSep08, mRNA.
Sult2b1	Sult2b1.bSep08	292915	13040	1777	1	586	sulfotransferase family, cytosolic, 2B, member 1 (Sult2b1) alternative variant bSep08, mRNA.
Sult2b1	Sult2b1.dSep08	292915	8284	538	2	179	sulfotransferase family, cytosolic, 2B, member 1 (Sult2b1) alternative variant dSep08, mRNA.
Sult5a1	Sult5a1.aSep08	292077	5732	881	6	256	sulfotransferase family 5A member 1 (Sult5a1) alternative variant aSep08, mRNA.
Sult5a1	Sult5a1.bSep08	292077	15273	945	6	154	sulfotransferase family 5A member 1 like (Sult5a1) alternative variant bSep08, mRNA.
Sult5a1	Sult5a1.dSep08	292077	1531	824	2	65	sulfotransferase family 5A member 1 CRA b (7.5 kD) (Sult5a1) alternative variant dSep08, mRNA.
Sult5a1	Sult5a1.eSep08	292077	26158	770	5	9	putative protein (Sult5a1) alternative variant eSep08, mRNA.
Sult5a1	Sult5a1.fSep08	292077	1108	543	2	50	putative protein (5.7 kD) (Sult5a1) alternative variant fSep08, mRNA.
Sult6b1	Sult6b1.aSep08	503103	4546	490		95	sulfotransferase family, cytosolic, 6B, member 1 (Sult6b1) mRNA.

sumee	sumee.aSep08		11582	732	2	21	putative protein (2.5 kD) (sumee) alternative variant aSep08, mRNA.
sumer	sumer.bSep08		993	828	2	83	putative cytoplasmic protein (9.4 kD) (sumer) alternative variant bSep08, mRNA.
Sumf2	Sumf2.bSep08	360800	22929	680	6	226	sulfatase modifying factor 2 (Sumf2) alternative variant bSep08, mRNA.
Sumf2	Sumf2.cSep08	360800	22729	702	5	192	sulfatase modifying factor 2 (Sumf2) alternative variant cSep08, mRNA.
Sumf2	Sumf2.dSep08	360800	14982	386	4	125	sulfatase modifying factor 2 (Sumf2) alternative variant dSep08, mRNA.
Sumf2	Sumf2.eSep08	360800	3410	1486	3	71	sulfatase modifying factor 2 (7.7 kD) (Sumf2) alternative variant eSep08, mRNA.
Sumo1	Sumo1.bSep08	301442	29793	769	4	62	SMT3 suppressor of mif two 3 homolog 1 (yeast) (7.2 kD) (Sumo1) alternative variant bSep08, mRNA.
Sumo1	Sumo1.cSep08	301442	29390	740	4	62	SMT3 suppressor of mif two 3 homolog 1 (yeast) (7.2 kD) (Sumo1) alternative variant cSep08, mRNA.
Sumo1	Sumo1.dSep08	301442	29258	734	5	61	SMT3 suppressor of mif two 3 homolog 1 (yeast) (Sumo1) alternative variant dSep08, mRNA.
Sumo1	Sumo1.eSep08	301442	29700	673	4	62	SMT3 suppressor of mif two 3 homolog 1 (yeast) (7.2 kD) (Sumo1) alternative variant eSep08, mRNA.
Sumo3	Sumo3.bSep08	499417	7998	910	3	75	SMT3 suppressor of mif two 3 homolog 3 ( <i>S. cerevisiae</i> ) (8.6 kD) (Sumo3) alternative variant bSep08, complete mRNA.
Sumo3	Sumo3.cSep08	499417	2649	752	2	57	SMT3 suppressor of mif two 3 homolog 3 ( <i>S. cerevisiae</i> ) (6.5 kD) (Sumo3) alternative variant cSep08, mRNA.
Sumo3	Sumo3.dSep08	499417	2744	674	3	57	SMT3 suppressor of mif two 3 homolog 3 ( <i>S. cerevisiae</i> ) (6.5 kD) (Sumo3) alternative variant dSep08, mRNA.
sunoy	sunoy.aSep08		72068	473		34	putative protein (sunoy) mRNA.
Suox	Suox.bSep08	81805	1014	417	2	82	sulfite oxidase (Suox) alternative variant bSep08, mRNA.
supor	supor.aSep08		22619	819	7	273	testis expressed 9 (supor) alternative variant aSep08, mRNA.
supor	supor.bSep08		36663	1805	11	188	testis expressed 9 (22.2 kD) (supor) alternative variant bSep08, mRNA.
supor	supor.cSep08		18090	1172	4	150	testis expressed 9 (supor) alternative variant cSep08, mRNA.
supor	supor.dSep08		15635	634	6	112	putative nuclear protein of vertebrate origin (12.3 kD) (supor) alternative variant dSep08, mRNA.
Supt3h	Supt3h.aSep08	685697	340000	2933	1	391	suppressor of Ty 3 homolog ( <i>S. cerevisiae</i> ) (Supt3h) alternative variant aSep08, mRNA.
Supt3h	Supt3h.bSep08	685697	263490	823	1	193	suppressor of Ty 3 homolog ( <i>S. cerevisiae</i> ) (Supt3h) alternative variant bSep08, mRNA.
Supt4h1	Supt4h1.aSep08	287608	6195	1054	5	117	suppressor of Ty 4 homolog 1 ( <i>S. cerevisiae</i> ) (13.2 kD) (Supt4h1) alternative variant aSep08, complete mRNA.
Supt4h1	Supt4h1.bSep08	287608	5510	449	6	89	suppressor of Ty 4 homolog 1 ( <i>S. cerevisiae</i> ) (10.1 kD) (Supt4h1) alternative variant bSep08, complete mRNA.
Supt4h1	Supt4h1.dSep08	287608	5520	455	6	85	suppressor of Ty 4 homolog 1 ( <i>S. cerevisiae</i> ) (9.6 kD) (Supt4h1) alternative variant dSep08, complete mRNA.

Supt5h	Supt5h.aSep08	308472	29223	4443	28	556	suppressor of Ty 5 homolog (62.9 kD) (Supt5h) alternative variant aSep08, mRNA.
Supt5h	Supt5h.bSep08	308472	7052	1436	13	478	suppressor of Ty 5 homolog (Supt5h) alternative variant bSep08, mRNA.
Supt5h	Supt5h.cSep08	308472	7276	2103	13	328	suppressor of Ty 5 homolog (35.3 kD) (Supt5h) alternative variant cSep08, mRNA.
Supt5h	Supt5h.dSep08	308472	15130	906	10	262	suppressor of Ty 5 homolog (Supt5h) alternative variant dSep08, mRNA.
Supt5h	Supt5h.eSep08	308472	6651	756	9	252	suppressor of Ty 5 homolog (Supt5h) alternative variant eSep08, mRNA.
Supt5h	Supt5h.fSep08	308472	12990	704	10	207	suppressor of Ty 5 homolog (Supt5h) alternative variant fSep08, mRNA.
Supt5h	Supt5h.gSep08	308472	1494	363	3	121	suppressor of Ty 5 homolog CRA b (Supt5h) alternative variant gSep08, mRNA.
Supt5h	Supt5h.iSep08	308472	1846	1768	2	78	suppressor of Ty 5 homolog CRA b (8.7 kD) (Supt5h) alternative variant iSep08, mRNA.
Supt5h	Supt5h.jSep08	308472	1112	534	3	74	putative protein (Supt5h) alternative variant jSep08, mRNA.
Supt5h	Supt5h.kSep08	308472	9476	264	3	65	putative protein (Supt5h) alternative variant kSep08, mRNA.
Supt6h	Supt6h.aSep08	303281	14767	3332	16	832	suppressor of Ty 6 homolog CRA c (Supt6h) alternative variant aSep08, mRNA.
Supt6h	Supt6h.bSep08	303281	3814	740	5	187	suppressor of Ty 6 homolog CRA c (Supt6h) alternative variant bSep08, mRNA.
Supt6h	Supt6h.cSep08	303281	1363	708	3	159	suppressor of Ty 6 homolog (Supt6h) alternative variant cSep08, mRNA.
Supt6h	Supt6h.dSep08	303281	4542	1327	5	153	suppressor of Ty 6 homolog CRA c (Supt6h) alternative variant dSep08, mRNA.
Supt6h	Supt6h.eSep08	303281	1115	410	2	136	suppressor of Ty 6 homolog CRA c (Supt6h) alternative variant eSep08, mRNA.
Supt6h	Supt6h.fSep08	303281	1242	753	2	113	suppressor of Ty 6 homolog CRA c (Supt6h) alternative variant fSep08, mRNA.
Supt7l	Supt7l.aSep08	313905	10751	1777	1	438	suppressor of Ty 7 ( <i>S. cerevisiae</i> )-like (Supt7l) alternative variant aSep08, mRNA.
Supt16h	Supt16h.bSep08	305851	5377	3229	4	142	suppressor of Ty 16 homolog ( <i>S. cerevisiae</i> ) (16.2 kD) (Supt16h) alternative variant bSep08, mRNA.
Surf1	Surf1.bSep08	64463	2078	731	6	201	surfeit 1 (22.9 kD) (Surf1) alternative variant bSep08, mRNA.
Surf1	Surf1.cSep08	64463	2028	492	5	148	surfeit 1 (Surf1) alternative variant cSep08, mRNA.
Surf1	Surf1.dSep08	64463	487	340	2	72	surfeit 1 (Surf1) alternative variant dSep08, mRNA.
Surf1	Surf1.eSep08	64463	2817	2348	4	56	surfeit 1 (Surf1) alternative variant eSep08, mRNA.
Surf2	Surf2.bSep08	619345	3812	1347	5	203	surfeit gene 2 (23.8 kD) (Surf2) alternative variant bSep08, mRNA.
Surf2	Surf2.cSep08	619345	3715	2540	4	110	surfeit gene 2 (12.7 kD) (Surf2) alternative variant cSep08, mRNA.
Surf2	Surf2.dSep08	619345	1265	811	2	92	surfeit gene 2 (10.8 kD) (Surf2) alternative variant dSep08, mRNA.

susa	susa.aSep08		889	677		152	putative protein of mammalian origin (susa) mRNA.
Susd2	Susd2.bSep08	294335	7352	3008	13	767	AMOP and von Willebrand factor, type D and sushi/SCR/CCP (84.7 kD) (Susd2) alternative variant bSep08, complete mRNA.
Susd2	Susd2.cSep08	294335	922	727	2	230	putative protein of ancient origin (Susd2) alternative variant cSep08, mRNA.
Susd2	Susd2.dSep08	294335	474	382	1	123	putative protein of ancient origin (Susd2) alternative variant dSep08, mRNA.
sushee	sushee.aSep08		4247	420		139	sucrase-isomaltase (sushee) mRNA.
Sushi.0	Sushi.0.aSep08		1926	553		170	complement component 2 CRA d (Sushi.0) mRNA.
Sushi.1	Sushi.1.aSep08		14281	1084	4	361	cfh protein (Sushi.1) alternative variant aSep08, mRNA.
Sushi.1	Sushi.1.bSep08		14054	680	3	226	complement factor H-related protein B (Sushi.1) alternative variant bSep08, mRNA.
Sushi.1	Sushi.1.cSep08		11300	1065	2	31	cfh protein like (Sushi.1) alternative variant cSep08, mRNA.
Sushi.2	Sushi.2.aSep08		16553	617		129	complement component factor H (Sushi.2) mRNA.
Sushi.3	Sushi.3.aSep08		4685	2175	9	396	seizure related 6 (Sushi.3) alternative variant aSep08, mRNA.
Sushi.3	Sushi.3.bSep08		2379	738	6	245	seizure related 6 (Sushi.3) alternative variant bSep08, mRNA.
Sushi.3	Sushi.3.cSep08		573	463	2	85	seizure related 6 (Sushi.3) alternative variant cSep08, mRNA.
Sushi.3	Sushi.3.dSep08		782	697	2	74	seizure related 6 precursor (8.0 kD) (Sushi.3) alternative variant dSep08, mRNA.
Sushi.4	Sushi.4.aSep08		26286	1989		588	polydom (Sushi.4) alternative variant aSep08, mRNA.
sutu	sutu.bSep08		2875	563		39	putative protein (4.7 kD) (sutu) alternative variant bSep08, mRNA.
Suv39h1	Suv39h1.bSep08	302553	12743	2687	6	412	suppressor of variegation 3-9 homolog 1 (Drosophila) (47.8 kD) (Suv39h1) alternative variant bSep08, complete mRNA.
Suv39h1	Suv39h1.cSep08	302553	1586	855	2	91	suppressor of variegation 3-9 homolog 1 (Drosophila) (Suv39h1) alternative variant cSep08, mRNA.
Suv39h1	Suv39h1.eSep08	302553	481	142	2	17	suppressor of variegation 3-9 homolog 1 (Drosophila) (Suv39h1) alternative variant eSep08, mRNA.
Suv39h2	Suv39h2.bSep08	364785	10814	879		217	suppressor of variegation 3-9 homolog 2 (Drosophila) (Suv39h2) alternative variant bSep08, mRNA.
Suv420h1	Suv420h1.cSep08	361688	656	575	2	66	suppressor of variegation 4-20 homolog 1 (Drosophila) (Suv420h1) alternative variant cSep08, mRNA.
Suv420h2	Suv420h2.bSep08	308345	2074	1840	3	269	suppressor of variegation 4-20 homolog 2 (Drosophila) (30.7 kD) (Suv420h2) alternative variant bSep08, mRNA.
Suv420h2	Suv420h2.cSep08	308345	2553	411	4	121	suppressor of variegation 4-20 homolog 2 (Drosophila) (Suv420h2) alternative variant cSep08, mRNA.
suvar	suvar.aSep08		619	264		87	heparan sulfate proteoglycan (suvar) mRNA.
suwey	suwey.aSep08		2588	1074		216	murinoglobulin (suwey) mRNA.
Sv2a	Sv2a.bSep08	117559	2460	378	5	86	synaptic vesicle glycoprotein 2a (Sv2a) alternative variant bSep08, mRNA.

Sv2b	Sv2b.bSep08	117556	14853	534	4	126	synaptic vesicle glycoprotein 2 (Sv2b) alternative variant bSep08, mRNA.
Sv2b	Sv2b.cSep08	117556	98782	619	2	85	putative protein (Sv2b) alternative variant cSep08, mRNA.
Sv2b	Sv2b.dSep08	117556	99090	682	2	83	synaptic vesicle glycoprotein 2b CRA b (Sv2b) alternative variant dSep08, mRNA.
Svep1	Svep1.aSep08	685899	41668	3130		1042	polydom (Svep1) mRNA.
Svil	Svil.aSep08	361256	122063	677	6	103	supervillin (Svil) alternative variant aSep08, mRNA.
Svil	Svil.bSep08	361256	76212	314	5	65	supervillin (Svil) alternative variant bSep08, mRNA.
Svil	Svil.cSep08	361256	49571	517	4	65	supervillin (Svil) alternative variant cSep08, mRNA.
Svil	Svil.dSep08	361256	38088	557	2	42	supervillin (Svil) alternative variant dSep08, mRNA.
Svs3	Svs3.bSep08	192239	51908	304	2	37	seminal vesicle secretion 3 (4.1 kD) (Svs3) alternative variant bSep08, mRNA.
Svs4	Svs4.bSep08	24802	1557	376	1	49	seminal vesicle secretory protein 4 (5.5 kD) (Svs4) alternative variant bSep08, mRNA.
Svs6	Svs6.aSep08	362267	1663	494	1	99	seminal vesicle secretory protein 6 (11.4 kD) (Svs6) alternative variant aSep08, complete mRNA.
swacha	swacha.aSep08		1553	693		95	putative protein (swacha) mRNA.
swachy	swachy.aSep08		4833	497		93	putative protein (swachy) mRNA.
swafee	swafee.aSep08		1378	542		53	CRA b like (swafee) mRNA.
swafly	swafly.aSep08		3090	722		85	spectrin beta 4 CRA b (swafly) mRNA.
swafly	swafly.bSep08		498	365	2	110	putative protein (swafly) alternative variant bSep08, mRNA.
swafly	swafly.aSep08		3090	722		85	spectrin beta 4 CRA b (swafly) mRNA.
swalor	swalor.bSep08		827	377	2	39	putative protein (swalor) alternative variant bSep08, mRNA.
swamee	swamee.aSep08		4071	716		68	putative protein (7.5 kD) (swamee) mRNA.
swanor	swanor.aSep08		1923	480		52	putative protein (5.8 kD) (swanor) mRNA.
swapey	swapey.aSep08		21264	769		68	putative protein (7.5 kD) (swapey) mRNA.
swarcha	swarcha.aSep08		554	323		88	uncharacterized protein homolog like (swarcha) mRNA.
swarchy	swarchy.aSep08		2190	738		34	putative protein (3.8 kD) (swarchy) mRNA.
swarfee	swarfee.aSep08		6172	759		123	putative protein of mammalian origin (swarfee) mRNA.
swarflu	swarflu.aSep08		2684	418	4	67	putative protein of mammalian origin (swarflu) alternative variant aSep08, mRNA.
swarflu	swarflu.bSep08		911	269	2	44	putative protein of mammalian origin (swarflu) alternative variant bSep08, mRNA.
swarfly	swarfly.aSep08		2993	1309	4	119	polyprotein (swarfly) alternative variant aSep08, mRNA.
swarfly	swarfly.bSep08		854	668	2	89	polyprotein -pol (9.7 kD) (swarfly) alternative variant bSep08, mRNA.
swarlor	swarlor.aSep08		900	316		26	putative protein (swarlor) mRNA.
swarmee	swarmee.aSep08		1572	321		59	putative protein of mammalian origin (7.1 kD) (swarmee) mRNA.
swarnor	swarnor.aSep08		12578	1800		53	putative protein (swarnor) mRNA.
swaroy	swaroy.aSep08		2031	276		30	putative protein (3.6 kD) (swaroy) mRNA.
swarpey	swarpey.aSep08		10724	816	2	50	putative protein (5.7 kD) (swarpey) alternative variant aSep08, mRNA.

swarpey	swarpey.bSep08		3692	378	2	50	putative protein (5.7 kD) (swarpey) alternative variant bSep08, mRNA.
swarroy	swarroy.aSep08		33324	416		121	putative protein (swarroy) mRNA.
swarshaw	swarshaw.aSep08		3438	2098	2	77	putative mitochondrial protein (8.4 kD) (swarshaw) alternative variant aSep08, mRNA.
swartu	swartu.aSep08		1227	625		136	putative protein (swartu) mRNA.
swarvo	swarvo.aSep08		7649	621		147	EF-hand calcium binding domain 7 like (swarvo) mRNA.
swarwer	swarwer.aSep08		1176	1013		42	putative protein (swarwer) mRNA.
swashaw	swashaw.aSep08		24590	294		55	putative protein (swashaw) mRNA.
swatu	swatu.aSep08		44485	374	4	103	ATG2 autophagy related 2 homolog b (swatu) alternative variant aSep08, mRNA.
swavo	swavo.aSep08		10050	414		137	dedicator of cytokinesis 7 (swavo) mRNA.
swawcha	swawcha.aSep08		1534	1091	3	99	putative mitochondrial protein (11.4 kD) (swawcha) alternative variant aSep08, mRNA.
swawchy	swawchy.aSep08		2198	990		330	laminin (swawchy) mRNA.
swawer	swawer.aSep08		27236	354		34	putative protein (swawer) mRNA.
swawfee	swawfee.aSep08		1932	531		176	desmoplakin CRA b (swawfee) mRNA.
swawflu	swawflu.aSep08		3017	478		42	CRA b like (4.6 kD) (swawflu) mRNA.
swawfly	swawfly.aSep08		2420	759		53	putative protein (5.6 kD) (swawfly) mRNA.
swawlor	swawlor.bSep08		1025	324	2	55	putative protein (6.0 kD) (swawlor) alternative variant bSep08, mRNA.
swawmee	swawmee.aSep08		747	272		39	putative protein (swawmee) mRNA.
swawnor	swawnor.bSep08		4051	961	2	109	putative endoplasmic reticulum protein (12.3 kD) (swawnor) alternative variant bSep08, mRNA.
swawpey	swawpey.aSep08		34865	731		243	sortilin-related receptor L A repeats-containing (swawpey) mRNA.
swawroy	swawroy.bSep08		10308	394	2	48	putative protein (swawroy) alternative variant bSep08, mRNA.
swawshaw	swawshaw.aSep08		3087	638		47	putative protein (5.8 kD) (swawshaw) mRNA.
swawtu	swawtu.aSep08		887	315		52	domain-containing protein 85c like (swawtu) mRNA.
swawvo	swawvo.aSep08		810	751		39	putative protein (4.3 kD) (swawvo) mRNA.
swawwer	swawwer.aSep08		7996	1137	1	76	CRA a like (8.3 kD) (swawwer) alternative variant aSep08, mRNA.
swawwer	swawwer.bSep08		3970	1102	1	76	CRA a like (8.3 kD) (swawwer) alternative variant bSep08, mRNA.
sweecha	sweecha.aSep08		985	563		129	putative protein (14.1 kD) (sweecha) mRNA.
sweechy	sweechy.aSep08		16054	759		35	putative protein (sweechy) mRNA.
sweefee	sweefee.aSep08		1743	420		139	desmoplakin CRA b (sweefee) mRNA.
sweeflu	sweeflu.aSep08		7216	502		9	putative protein (1.0 kD) (sweeflu) mRNA.
sweefly	sweefly.aSep08		2449	1925		543	putative protein of bilateral origin (sweefly) mRNA.
sweelor	sweelor.bSep08		1023	324	2	55	putative protein (6.0 kD) (sweelor) alternative variant bSep08, mRNA.
sweemee	sweemee.aSep08		34756	812		117	putative protein (sweemee) mRNA.
sweenor	sweenor.aSep08		10213	787		202	CRA b (23.2 kD) (sweenor) mRNA.



sweepy	sweepy.aSep08		1705	754		59	putative protein (6.3 kD) (sweepy) mRNA.
sweeroy	sweeroy.aSep08		12731	258		57	putative protein (sweeroy) mRNA.
sweeshaw	sweeshaw.aSep08		3688	116		38	rapamycin-insensitive companion of mTOR CRA b (sweeshaw) mRNA.
sweetu	sweetu.aSep08		8018	527		175	protein-like 1 (sweetu) mRNA.
sweevo	sweevo.aSep08		5752	349		116	putative protein of ancient origin (sweevo) mRNA.
sweewer	sweewer.aSep08		53750	423		19	putative protein (sweewer) mRNA.
swercha	swercha.aSep08		20944	1463	8	359	GTPase activating rap RanGAP domain-like 3 (38.5 kD) (swercha) alternative variant aSep08, mRNA.
swerchy	swerchy.bSep08		1861	289		53	putative protein (swerchy) alternative variant bSep08, mRNA.
swerfee	swerfee.aSep08		7865	969		322	desmoplakin CRA b (swerfee) mRNA.
swerflu	swerflu.aSep08		618	538		72	putative protein (swerflu) mRNA.
swerfly	swerfly.aSep08		2890	1783		533	putative protein (swerfly) alternative variant aSep08, mRNA.
swerlor	swerlor.aSep08		6043	529		56	putative protein (5.7 kD) (swerlor) mRNA.
swermee	swermee.aSep08		17136	538		157	putative protein (swermee) mRNA.
swernor	swernor.aSep08		3701	571		190	methylmalonyl-CoA mutase (swernor) mRNA.
swerpey	swerpey.aSep08		1683	419		38	putative protein (4.5 kD) (swerpey) mRNA.
swerroy	swerroy.aSep08		47655	663		50	putative protein (swerroy) mRNA.
swershaw	swershaw.aSep08		8388	444		64	putative protein (swershaw) mRNA.
swertu	swertu.aSep08		25420	401		97	putative protein (swertu) mRNA.
swervo	swervo.aSep08		11646	637		211	putative protein of ancient origin (swervo) mRNA.
swerwer	swerwer.aSep08		1676	688		83	putative protein (swerwer) mRNA.
sweycha	sweycha.aSep08		56793	583	3	143	GTPase activating RanGAP domain-like 3 (sweycha) alternative variant aSep08, mRNA.
sweycha	sweycha.bSep08		33861	320	3	106	GTPase activating RanGAP domain-like 3 (sweycha) alternative variant bSep08, mRNA.
sweychy	sweychy.aSep08		1005	866		159	putative protein (sweychy) mRNA.
sweyfee	sweyfee.aSep08		649	498		36	putative protein (4.0 kD) (sweyfee) mRNA.
sweyflu	sweyflu.aSep08		1053	574	2	41	putative protein (5.0 kD) (sweyflu) alternative variant aSep08, mRNA.
sweyflu	sweyflu.bSep08		526	418	1	8	putative protein (1.0 kD) (sweyflu) alternative variant bSep08, mRNA.
sweyfly	sweyfly.aSep08		1859	620		74	putative protein (sweyfly) mRNA.
sweylor	sweylor.aSep08		14194	717		125	carbonyl reductase (sweylor) mRNA.
sweymee	sweymee.aSep08		7733	295	2	65	putative protein (7.5 kD) (sweymee) alternative variant aSep08, mRNA.
sweynor	sweynor.aSep08		2653	265		88	methylmalonyl-CoA mutase (sweynor) mRNA.
sweypey	sweypey.aSep08		2258	527	1	57	putative protein (sweypey) alternative variant aSep08, mRNA.
sweypey	sweypey.bSep08		1166	519	1	7	putative protein (sweypey) alternative variant bSep08, mRNA.

sweyroy	sweyroy.aSep08		14518	341		113	slit-robo Rho GTPase activating protein 3 (sweyroy) mRNA.
sweyshaw	sweyshaw.aSep08		3150	1230		64	putative protein (sweyshaw) mRNA.
sweytu	sweytu.aSep08		1791	515	3	105	putative protein (sweytu) alternative variant aSep08, mRNA.
sweyvo	sweyvo.aSep08		11635	929		309	putative protein of metazoan origin (sweyvo) mRNA.
sweywer	sweywer.aSep08		980	301		99	putative protein (sweywer) mRNA.
Swi3.0	Swi3.0.aSep08		16558	1719		240	timeless interacting protein (26.9 kD) (Swi3.0) mRNA.
SWIM.0	SWIM.0.aSep08		792	330		110	finger swim domain-containing protein (SWIM.0) mRNA.
swocha	swocha.aSep08		4060	731		243	putative protein of mammalian origin (swocha) mRNA.
swochy	swochy.aSep08		20439	411		84	putative protein (swochy) mRNA.
swofee	swofee.aSep08		5454	653		19	putative protein (swofee) mRNA.
swoflu	swoflu.aSep08		1629	935		268	putative protein, with 2 coiled coil domains, of bilateral origin (swoflu) mRNA.
swofly	swofly.aSep08		391	284		67	putative protein (swofly) mRNA.
swolor	swolor.aSep08		5162	723		118	putative protein (13.2 kD) (swolor) mRNA.
swomee	swomee.aSep08		3990	741		111	putative protein (swomee) mRNA.
swonor	swonor.aSep08		5851	437		84	putative protein (10.0 kD) (swonor) mRNA.
swopey	swopey.aSep08		34613	1261	2	35	putative protein (4.0 kD) (swopey) alternative variant aSep08, mRNA.
sworcha	sworcha.aSep08		2202	919		37	putative protein (4.2 kD) (sworcha) mRNA.
sworchy	sworchy.aSep08		1573	748		103	putative protein of mammalian origin (sworchy) mRNA.
sworfee	sworfee.aSep08		5294	620	2	42	putative protein (sworfee) alternative variant aSep08, mRNA.
sworfee	sworfee.bSep08		5605	1347	4	31	putative protein (3.5 kD) (sworfee) alternative variant bSep08, mRNA.
sworflu	sworflu.aSep08		49082	723		49	putative protein (5.5 kD) (sworflu) mRNA.
sworfly	sworfly.aSep08		760	348		53	putative protein (sworfly) mRNA.
sworlor	sworlor.aSep08		6399	1777		592	dopey family member 2 (sworlor) mRNA.
swormee	swormee.aSep08		6967	660	5	39	putative protein (4.5 kD) (swormee) alternative variant aSep08, mRNA.
swormee	swormee.bSep08		705	460	2	59	putative protein (swormee) alternative variant bSep08, mRNA.
swornor	swornor.aSep08		2654	265		88	methylmalonyl-CoA mutase (swornor) mRNA.
sworoy	sworoy.aSep08		16448	365		52	putative protein (6.1 kD) (sworoy) mRNA.
sworpey	sworpey.aSep08		7891	412		34	putative protein (sworpey) mRNA.
sworroy	sworroy.aSep08		26293	710		32	putative protein (sworroy) mRNA.
sworshaw	sworshaw.aSep08		31965	648		49	putative protein (sworshaw) mRNA.
swortu	swortu.aSep08	691052	1294	915		263	putative protein of vertebrate origin (swortu) mRNA.
sworvo	sworvo.aSep08		3902	505		44	putative protein (sworvo) mRNA.
sworwer	sworwer.aSep08		14227	655		31	putative protein (3.5 kD) (sworwer) alternative variant aSep08, mRNA.
swoshaw	swoshaw.aSep08		5953	635		211	CRA a (swoshaw) mRNA.

swotu	swotu.aSep08		3026	450	2	44	putative protein (4.9 kD) (swotu) alternative variant aSep08, mRNA.
swotu	swotu.bSep08		5079	341	1	44	putative protein (swotu) alternative variant bSep08, mRNA.
swovo	swovo.aSep08		16934	682		227	EF-hand calcium binding domain 7 like (swovo) mRNA.
swower	swower.aSep08		37533	540		43	putative protein (4.7 kD) (swower) mRNA.
swoycha	swoycha.aSep08		22560	530		94	GTPase activating protein vps9 domains 1 (swoycha) mRNA.
swoychy	swoychy.aSep08		2833	641	6	138	collagen type XX alpha 1 (swoychy) alternative variant aSep08, mRNA.
swoyfee	swoyfee.aSep08		7625	603		67	putative protein (7.2 kD) (swoyfee) mRNA.
swoyflu	swoyflu.aSep08		2672	788		56	putative protein (swoyflu) mRNA.
swoyfly	swoyfly.aSep08		1230	730	1	99	putative protein (swoyfly) alternative variant aSep08, mRNA.
swoyfly	swoyfly.bSep08		2876	694	3	79	putative protein (swoyfly) alternative variant bSep08, mRNA.
swoylor	swoylor.aSep08		63127	698	2	131	putative protein (swoylor) alternative variant aSep08, mRNA.
swoylor	swoylor.bSep08		4916	405	1	30	putative protein (swoylor) alternative variant bSep08, mRNA.
swoymee	swoymee.aSep08		13921	318		73	putative cytoplasmic protein (8.2 kD) (swoymee) mRNA.
swoynor	swoynor.aSep08		3701	571		190	methylmalonyl-CoA mutase (swoynor) mRNA.
swoypey	swoypey.aSep08		5589	640		212	rho guanine nucleotide exchange factor 12 (swoypey) mRNA.
swoyroy	swoyroy.aSep08		5228	614	1	45	putative protein (swoyroy) alternative variant aSep08, mRNA.
swoyroy	swoyroy.bSep08		9939	341	1	45	putative protein (5.2 kD) (swoyroy) alternative variant bSep08, mRNA.
swoyshaw	swoyshaw.aSep08		1284	1067		56	CRA a like (swoyshaw) alternative variant aSep08, mRNA.
swoyshaw	swoyshaw.bSep08		729	512		59	CRA a like (swoyshaw) alternative variant bSep08, mRNA.
swoytu	swoytu.aSep08		4948	818		94	putative mitochondrial protein (10.2 kD) (swoytu) mRNA.
swoyvo	swoyvo.aSep08		837	213		16	putative protein (swoyvo) mRNA.
swoywer	swoywer.aSep08		1494	495		119	gag protein (swoywer) mRNA.
swucha	swucha.aSep08		12892	760	10	252	golgi autoantigen golgin subfamily a like (swucha) alternative variant aSep08, mRNA.
swucha	swucha.bSep08		11398	407	5	135	golgi autoantigen golgin subfamily a like (swucha) alternative variant bSep08, mRNA.
swucha	swucha.cSep08		553	454	1	46	golgi autoantigen golgin subfamily a like (swucha) alternative variant cSep08, mRNA.
swuchy	swuchy.aSep08		986	817			
swufee	swufee.aSep08		6380	964		35	putative protein (4.2 kD) (swufee) mRNA.
swuflu	swuflu.aSep08		2407	696		210	CRA c like (swuflu) mRNA.
swuflu	swuflu.aSep08		2125	283		93	beta spectrin (swuflu) mRNA.
swuly	swuly.aSep08		2125	283		93	beta spectrin (swuly) mRNA.
swulor	swulor.aSep08		22788	701	2	67	putative protein (7.5 kD) (swulor) alternative variant aSep08, mRNA.

swulor	swulor.bSep08		10219	617	1	39	putative protein (4.4 kD) (swulor) alternative variant bSep08, mRNA.
swumee	swumee.aSep08		3976	925		83	putative protein (9.0 kD) (swumee) mRNA.
swunor	swunor.aSep08		8307	715		50	putative protein (swunor) mRNA.
swupey	swupey.bSep08		2435	592	2	32	putative protein (3.6 kD) (swupey) alternative variant bSep08, mRNA.
swupey	swupey.cSep08		115316	511	4	37	putative protein (swupey) alternative variant cSep08, mRNA.
swupey	swupey.dSep08		61418	353	3	46	putative protein (5.5 kD) (swupey) alternative variant dSep08, mRNA.
swuroy	swuroy.aSep08		1404	492	2	44	putative protein (5.0 kD) (swuroy) alternative variant aSep08, mRNA.
swuroy	swuroy.bSep08		1078	396	1	26	putative protein (2.9 kD) (swuroy) alternative variant bSep08, mRNA.
swutu	swutu.aSep08		6977	266		88	ATG2 autophagy related 2 homolog B (swutu) mRNA.
swuvo	swuvo.aSep08		1340	503		47	putative protein (5.8 kD) (swuvo) mRNA.
swuwer	swuwer.aSep08		21289	416		49	putative protein (swuwer) mRNA.
swycha	swycha.aSep08		1222	851		90	protein-O-mannosyltransferase 1 (swycha) mRNA.
swychy	swychy.aSep08		716	513		31	putative protein (swychy) mRNA.
swyfee	swyfee.bSep08		2962	322	3	38	putative protein (swyfee) alternative variant bSep08, mRNA.
swyflu	swyflu.aSep08		1539	710		46	putative protein (swyflu) mRNA.
swyfly	swyfly.aSep08		1231	315		104	spectrin (swyfly) mRNA.
swylor	swylor.aSep08		1373	602		38	putative protein (4.3 kD) (swylor) mRNA.
swymee	swymee.aSep08		8088	611			
swynor	swynor.aSep08		6674	285		15	putative protein (swynor) mRNA.
swypey	swypey.aSep08		3601	624		142	putative protein (swypey) mRNA.
swyro	swyro.aSep08		6201	1525	1	95	hclb mouse DNA Helicase B like (swyro) alternative variant aSep08, mRNA.
swyro	swyro.bSep08		970	714	1	61	putative protein (swyro) alternative variant bSep08, mRNA.
swyshaw	swyshaw.aSep08		18103	421		44	putative protein (swyshaw) mRNA.
swytu	swytu.aSep08		1240	284		94	ATG2 autophagy related 2 homolog B (swytu) mRNA.
swyvo	swyvo.aSep08		7191	776		84	putative membrane protein (9.2 kD) (swyvo) mRNA.
Syap1	Syap1.aSep08	302678	34918	2679	9	360	synapse associated protein 1 (40.8 kD) (Syap1) alternative variant aSep08, mRNA.
Syap1	Syap1.bSep08	302678	32007	732	6	232	synapse associated protein 1 (Syap1) alternative variant bSep08, mRNA.
syby	syby.bSep08		4789	597		50	putative protein (syby) alternative variant bSep08, mRNA.
Syce2	Syce2.aSep08	364976	1419	393		56	synaptonemal complex central element protein 2 (Syce2) mRNA.
sychy	sychy.bSep08		1329	359	2	16	putative protein (2.0 kD) (sychy) alternative variant bSep08, mRNA.
sychy	sychy.cSep08		1249	314	3	16	putative protein (2.0 kD) (sychy) alternative variant cSep08, mRNA.

Sycn	Sycn.bSep08	245917	1388	519	2	122	syncollin (Sycn) alternative variant bSep08, mRNA.
sydar	sydar.aSep08		933	235		42	putative protein (sydar) mRNA.
Syde1	Syde1.aSep08	362842	3293	1291	6	430	synapse defective 1 Rho GTPase homolog (Syde1) alternative variant aSep08, mRNA.
Syde1	Syde1.bSep08	362842	602	393	2	86	synapse defective 1 Rho GTPase homolog CRA a (Syde1) alternative variant bSep08, mRNA.
Syde2	Syde2.aSep08	308021	13840	779		242	RhoGAP (Syde2) mRNA.
Syf2	Syf2.bSep08	170933	4163	747	4	152	SYF2 homolog, RNA splicing factor ( <i>S. cerevisiae</i> ) (17.5 kD) (Syf2) alternative variant bSep08, mRNA.
Syf2	Syf2.dSep08	170933	2593	865	2	77	SYF2 homolog, RNA splicing factor ( <i>S. cerevisiae</i> ) (8.2 kD) (Syf2) alternative variant dSep08, mRNA.
Syf2	Syf2.fSep08	170933	2055	548	3	35	SYF2 homolog, RNA splicing factor ( <i>S. cerevisiae</i> ) (Syf2) alternative variant fSep08, mRNA.
syfer	syfer.aSep08		2083	624		58	rho GTPase activating protein 21 (syfer) mRNA.
syflo	syflo.aSep08		1947	612		63	putative protein (syflo) mRNA.
syflu	syflu.aSep08		2270	392		86	putative nuclear protein (9.6 kD) (syflu) mRNA.
sygar	sygar.aSep08		2712	338		89	CRA a like (sygar) mRNA.
syja	syja.aSep08		7904	822		53	putative protein (syja) mRNA.
Syja_N.0	Syja_N.0.aSep08		7319	420		139	synaptojanin 1 CRA e (Syja_N.0) mRNA.
syjey	syjey.aSep08		13746	769		56	putative protein (6.3 kD) (syjey) mRNA.
Syk	Syk.bSep08	25155	8266	389	1	129	spleen tyrosine kinase (Syk) alternative variant bSep08, mRNA.
Syk	Syk.cSep08	25155	16318	638	3	114	spleen tyrosine kinase (Syk) alternative variant cSep08, mRNA.
sykee	sykee.aSep08		2152	789		211	putative protein (sykee) mRNA.
sykler	sykler.aSep08		2177	684	2	71	chromatin assembly factor 1 like (sykler) alternative variant aSep08, mRNA.
sylo	sylo.aSep08		8482	747		248	furry homolog CRA a (sylo) mRNA.
symee	symee.aSep08		1164	415		36	putative protein (symee) mRNA.
symer	symer.aSep08		16765	724		167	solute carrier family 38 member 10 (symer) mRNA.
Sympk	Sympk.aSep08	292683	17157	2585	17	861	symplekin (Sympk) alternative variant aSep08, mRNA.
Sympk	Sympk.bSep08	292683	2657	1416	5	333	symplekin (36.4 kD) (Sympk) alternative variant bSep08, mRNA.
Sympk	Sympk.cSep08	292683	17465	787	6	222	symplekin (Sympk) alternative variant cSep08, mRNA.
Sympk	Sympk.dSep08	292683	9076	419	6	139	symplekin (Sympk) alternative variant dSep08, mRNA.
Syn1	Syn1.bSep08	24949	2517	1574	2	186	synapsin I (Syn1) alternative variant bSep08, mRNA.
Syn1	Syn1.cSep08	24949	37492	679	3	95	synapsin I (Syn1) alternative variant cSep08, mRNA.
Syn3	Syn3.bSep08	29130	9272	560	4	165	synapsin III (Syn3) alternative variant bSep08, mRNA.
Sync	Sync.bSep08	297884	14629	1780	4	103	similar to 60S ribosomal protein L23a and syncoilin (Sync) alternative variant bSep08, mRNA.
Sync	Sync.bSep08	362606	14629	1780	4	103	similar to 60S ribosomal protein L23a and syncoilin (Sync) alternative variant bSep08, mRNA.
Syncrip	Syncrip.aSep08	363113	31945	3528	12	542	synaptotagmin binding, cytoplasmic RNA interacting protein (Syncrip) alternative variant aSep08, mRNA.

Syne1	Syne1.bSep08	499010	28554	2861	11	631	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant bSep08, mRNA.
Syne1	Syne1.cSep08	499010	31245	994	7	174	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant cSep08, mRNA.
Syne1	Syne1.dSep08	499010	4630	760	3	137	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant dSep08, mRNA.
Syne1	Syne1.eSep08	499010	15735	393	4	131	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant eSep08, mRNA.
Syne1	Syne1.gSep08	499010	2347	472	2	114	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant gSep08, mRNA.
Syne1	Syne1.hSep08	499010	11184	951	3	86	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant hSep08, mRNA.
Syne1	Syne1.jSep08	499010	11837	180	3	60	spectrin repeat containing, nuclear envelope 1 (Syne1) alternative variant jSep08, mRNA.
Syne2	Syne2.aSep08	299151	18681	1146		381	synaptic nuclear envelope 2 (Syne2) mRNA.
Syngr1	Syngr1.bSep08	29205	20372	574	2	191	synaptogyrin 1 (Syngr1) alternative variant bSep08, mRNA.
Syngr1	Syngr1.cSep08	29205	20507	759	2	186	synaptogyrin 1 (Syngr1) alternative variant cSep08, mRNA.
Syngr3	Syngr3.aSep08	302975	4750	1989	3	271	synaptogyrin 3 (Syngr3) alternative variant aSep08, mRNA.
Syngr3	Syngr3.bSep08	302975	1011	471	1	153	synaptogyrin 3 (Syngr3) alternative variant bSep08, mRNA.
Synj1	Synj1.aSep08	85238	26731	4203	4	694	synaptojanin 1 (Synj1) alternative variant aSep08, mRNA.
Synj1	Synj1.bSep08	85238	19082	1239	1	269	synaptojanin 1 (Synj1) alternative variant bSep08, mRNA.
Synj1	Synj1.cSep08	85238	13138	737	2	179	synaptojanin 1 (Synj1) alternative variant cSep08, mRNA.
Synj2	Synj2.dSep08	84018	4719	532	3	177	synaptojanin 2 (Synj2) alternative variant dSep08, mRNA.
Synj2	Synj2.eSep08	84018	3217	381	3	120	synaptojanin 2 (Synj2) alternative variant eSep08, mRNA.
synoy	synoy.aSep08		350	257		41	putative protein (synoy) mRNA.
Synpo	Synpo.aSep08	60324	53399	3093	4	985	synaptopodin (105.4 kD) (Synpo) alternative variant aSep08, mRNA.
Synpo2	Synpo2.aSep08	499702	41841	906	2	301	synaptopodin 2 (Synpo2) alternative variant aSep08, mRNA.
Synpo2l	Synpo2l.aSep08	305675	3085	1191		396	synaptopodin 2-like (Synpo2l) mRNA.
Synpr	Synpr.bSep08	66030	322006	611	5	153	synaptoporin (Synpr) alternative variant bSep08, mRNA.
Synpr	Synpr.cSep08	66030	164248	570	4	139	synaptoporin (Synpr) alternative variant cSep08, mRNA.
Synpr	Synpr.dSep08	66030	124700	447	3	88	synaptoporin (Synpr) alternative variant dSep08, mRNA.
Sypl	Sypl.aSep08	366595	19073	1001	4	259	synaptophysin-like protein (28.1 kD) (Sypl) alternative variant aSep08, complete mRNA.
Sypl	Sypl.bSep08	366595	17530	750	6	250	synaptophysin-like protein (Sypl) alternative variant bSep08, mRNA.
Sypl	Sypl.dSep08	366595	20229	2017	5	162	synaptophysin-like protein (18.2 kD) (Sypl) alternative variant dSep08, complete mRNA.
Sypl	Sypl.eSep08	366595	14751	467	3	145	synaptophysin-like protein (Sypl) alternative variant eSep08, mRNA.
Sypl	Sypl.fSep08	366595	3773	730	3	122	synaptophysin-like protein (Sypl) alternative variant fSep08, mRNA.
sypor	sypor.aSep08		6663	396	1	120	suppressor of hairy wing homolog 4 (sypor) alternative variant aSep08, mRNA.

sypor	sypor.bSep08		20531	372	2	97	suppressor of hairy wing homolog 4 (sypor) alternative variant bSep08, mRNA.
sysa	sysa.aSep08		677	388		103	putative protein (sysa) mRNA.
syshee	syshee.aSep08		29265	1225		80	CRA a like (8.9 kD) (syshee) mRNA.
Syt1	Syt1.bSep08	25716	14111	897	4	191	synaptotagmin I (Syt1) alternative variant bSep08, mRNA.
Syt3	Syt3.bSep08	25731	3880	1762	4	550	synaptotagmin III (Syt3) alternative variant bSep08, mRNA.
Syt3	Syt3.cSep08	25731	3960	380	2	117	synaptotagmin III (Syt3) alternative variant cSep08, mRNA.
Syt3	Syt3.dSep08	25731	5085	437	3	95	synaptotagmin III (Syt3) alternative variant dSep08, mRNA.
Syt6	Syt6.bSep08	60565	50815	1973	5	467	synaptotagmin VI (52.5 kD) (Syt6) alternative variant bSep08, complete mRNA.
Syt6	Syt6.cSep08	60565	10983	473	2	141	synaptotagmin VI (Syt6) alternative variant cSep08, mRNA.
Syt6	Syt6.dSep08	60565	12703	1674	5	122	synaptotagmin VI (Syt6) alternative variant dSep08, mRNA.
Syt7	Syt7.bSep08	59267	31846	394	4	131	synaptotagmin VII (Syt7) alternative variant bSep08, mRNA.
Syt7	Syt7.dSep08	59267	32900	928	4	116	synaptotagmin VII (13.0 kD) (Syt7) alternative variant dSep08, mRNA.
Syt7	Syt7.eSep08	59267	6020	644	6	91	synaptotagmin VII (Syt7) alternative variant eSep08, mRNA.
Syt8	Syt8.bSep08	60566	1050	624		131	synaptotagmin VIII (Syt8) alternative variant bSep08, mRNA.
Syt11	Syt11.bSep08	60568	13281	963	1	224	synaptotagmin XI (Syt11) alternative variant bSep08, mRNA.
Syt16	Syt16.aSep08	299142	15359	395		119	synaptotagmin XVI (Syt16) mRNA.
Syt17	Syt17.aSep08	192189	52407	612		193	synaptotagmin XVII (Syt17) mRNA.
Sytl1	Sytl1.bSep08	297872	3074	732	5	218	synaptotagmin-like 1 (Sytl1) alternative variant bSep08, mRNA.
Sytl1	Sytl1.cSep08	297872	599	524	2	92	synaptotagmin-like 1 (9.9 kD) (Sytl1) alternative variant cSep08, mRNA.
Sytl2	Sytl2.bSep08	361604	25174	2911	10	376	synaptotagmin-like 2 (42.7 kD) (Sytl2) alternative variant bSep08, mRNA.
Sytl2	Sytl2.cSep08	361604	7700	718	4	199	synaptotagmin-like 2 (Sytl2) alternative variant cSep08, mRNA.
Sytl2	Sytl2.dSep08	361604	13307	585	4	90	synaptotagmin-like 2 (Sytl2) alternative variant dSep08, mRNA.
Sytl3	Sytl3.bSep08	499017	1761	645	2	150	synaptotagmin-like 3 (Sytl3) alternative variant bSep08, mRNA.
sytu	sytu.aSep08		1157	869		33	putative protein (3.8 kD) (sytu) mRNA.
syvar	syvar.aSep08		33975	414		120	heparan sulfate proteoglycan (syvar) mRNA.
Syvn1	Syvn1.bSep08	361712	3522	2221	9	397	synovial apoptosis inhibitor 1, synoviolin (Syvn1) alternative variant bSep08, mRNA.
Syvn1	Syvn1.cSep08	361712	1636	1063	6	279	synovial apoptosis inhibitor 1, synoviolin (29.7 kD) (Syvn1) alternative variant cSep08, mRNA.
Syvn1	Syvn1.dSep08	361712	2131	1381	4	208	synovial apoptosis inhibitor 1, synoviolin (Syvn1) alternative variant dSep08, mRNA.

Syvn1	Syvn1.eSep08	361712	1248	444	5	147	synovial apoptosis inhibitor 1, synoviolin (Syvn1) alternative variant eSep08, mRNA.
Syvn1	Syvn1.fSep08	361712	1393	386	3	120	synovial apoptosis inhibitor 1, synoviolin (Syvn1) alternative variant fSep08, mRNA.
sywey	sywey.aSep08		1992	133		44	murinoglobulin 2 like (sywey) mRNA.
T2	T2.aSep08	681288	5573	388	3	129	brachyury 2 (T2) alternative variant aSep08, mRNA.
T2	T2.bSep08	681288	910	506	2	54	brachyury 2 (T2) alternative variant bSep08, mRNA.
taby	taby.aSep08		2828	728		44	putative protein of vertebrate origin (taby) mRNA.
Tacc2	Tacc2.cSep08	309025	85309	1623	4	540	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant cSep08, mRNA.
Tacc2	Tacc2.dSep08	309025	20232	1554	9	295	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant dSep08, mRNA.
Tacc2	Tacc2.eSep08	309025	9826	731	6	243	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant eSep08, mRNA.
Tacc2	Tacc2.fSep08	309025	23318	652	8	217	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant fSep08, mRNA.
Tacc2	Tacc2.gSep08	309025	15752	495	5	164	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant gSep08, mRNA.
Tacc2	Tacc2.iSep08	309025	425	327	2	41	transforming, acidic coiled-coil containing protein 2 (Tacc2) alternative variant iSep08, mRNA.
Tacc3	Tacc3.aSep08	360962	14799	2442	15	632	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant aSep08, mRNA.
Tacc3	Tacc3.bSep08	360962	7459	891	6	292	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant bSep08, mRNA.
Tacc3	Tacc3.cSep08	360962	1871	818	3	272	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant cSep08, mRNA.
Tacc3	Tacc3.dSep08	360962	6621	810	6	263	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant dSep08, mRNA.
Tacc3	Tacc3.eSep08	360962	2467	1431	5	123	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant eSep08, mRNA.
Tacc3	Tacc3.fSep08	360962	1369	793	3	104	transforming, acidic coiled-coil containing protein 3 (Tacc3) alternative variant fSep08, mRNA.
tachy	tachy.aSep08		2371	350		49	putative protein (5.1 kD) (tachy) mRNA.
Tada1l	Tada1l.bSep08	360874	14440	1812	7	248	transcriptional adaptor 1 (HFI1 homolog, yeast) like (28.0 kD) (Tada1l) alternative variant bSep08, mRNA.
Tada1l	Tada1l.cSep08	360874	4286	1491	3	174	transcriptional adaptor 1 (HFI1 homolog, yeast) like (Tada1l) alternative variant cSep08, mRNA.
Tada2l	Tada2l.bSep08	360581	24322	731	8	98	transcriptional adaptor 2 (ADA2 homolog, yeast)-like (12.0 kD) (Tada2l) alternative variant bSep08, mRNA.
Tada3l	Tada3l.aSep08	362414	7446	1838	7	181	transcriptional adaptor 3 -like CRA c (20.5 kD) (Tada3l) alternative variant aSep08, mRNA.
Tada3l	Tada3l.bSep08	362414	2889	850	4	135	transcriptional adaptor 3 -like CRA c (Tada3l) alternative variant bSep08, mRNA.
Tada3l	Tada3l.cSep08	362414	4498	551	2	74	transcriptional adaptor 3 -like CRA d (Tada3l) alternative variant cSep08, mRNA.



Tada3l	Tada3l.eSep08	362414	2597	644	3	37	transcriptional adaptor 3 -like CRA d (Tada3l) alternative variant eSep08, mRNA.
tadar	tadar.aSep08		2475	223		73	hydrocephalus inducing (tadar) mRNA.
Taf1	Taf1.aSep08	317256	19472	735	1	214	TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf1) alternative variant aSep08, mRNA.
Taf1	Taf1.bSep08	317256	17663	579	1	193	TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf1) alternative variant bSep08, mRNA.
Taf1a	Taf1a.bSep08	360893	4808	471		8	TATA box binding protein (Tbp)-associated factor, RNA polymerase I, A (1.0 kD) (Taf1a) alternative variant bSep08, mRNA.
Taf1b	Taf1b.aSep08	690450	27815	673		224	TATA box binding protein (Tbp)-associated factor, RNA polymerase I, B (Taf1b) mRNA.
Taf1c	Taf1c.bSep08	361420	2205	639	3	160	CRA b (17.5 kD) (Taf1c) alternative variant bSep08, mRNA.
Taf1c	Taf1c.cSep08	361420	2186	494	4	129	CRA c (Taf1c) alternative variant cSep08, mRNA.
Taf1c	Taf1c.dSep08	361420	2204	470	5	108	tata box binding protein -associated factor RNA polymerase I C 110kDa like (Taf1c) alternative variant dSep08, mRNA.
Taf1c	Taf1c.fSep08	361420	494	415	2	77	putative protein (Taf1c) alternative variant fSep08, mRNA.
Taf2	Taf2.aSep08	170844	21942	1475		412	TAF2 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf2) mRNA.
TAF4.0	TAF4.0.aSep08		51558	1904		149	associated factor (TAF4.0) mRNA.
TAF4.1	TAF4.1.aSep08		11601	1770	6	169	CRA a (TAF4.1) alternative variant aSep08, mRNA.
TAF4.1	TAF4.1.bSep08		1924	536	3	24	putative protein (TAF4.1) alternative variant bSep08, mRNA.
Taf5	Taf5.bSep08	294018	3264	805	1	134	TAF5 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf5) alternative variant bSep08, mRNA.
Taf5l	Taf5l.aSep08	307927	17587	2782	1	609	TAF5-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor (Taf5l) alternative variant aSep08, mRNA.
Taf5l	Taf5l.bSep08	307927	17022	984	1	327	TAF5-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor (Taf5l) alternative variant bSep08, mRNA.
Taf6	Taf6.bSep08	288533	2369	1031	4	261	TAF6 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf6) alternative variant bSep08, mRNA.
Taf6	Taf6.cSep08	288533	1508	734	5	187	TAF6 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf6) alternative variant cSep08, mRNA.
Taf6l	Taf6l.bSep08	309194	12652	2332	5	623	TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor (68.0 kD) (Taf6l) alternative variant bSep08, mRNA.
Taf6l	Taf6l.cSep08	309194	7654	1057	3	283	TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor (31.6 kD) (Taf6l) alternative variant cSep08, mRNA.
Taf7l	Taf7l.aSep08	363493	4258	1099		209	TAF7-like RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf7l) mRNA.
Taf8	Taf8.cSep08	316216	4456	371	2	76	TAF8 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf8) alternative variant cSep08, mRNA.

Taf9	Taf9.cSep08	373541	3319	801	3	54	TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf9) alternative variant cSep08, mRNA.
Taf9	Taf9.fSep08	373541	1005	273	2	23	TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf9) alternative variant fSep08, mRNA.
Taf9b	Taf9b.bSep08	171152	8250	760	1	201	TAF9B RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf9b) alternative variant bSep08, mRNA.
Taf10	Taf10.bSep08	293345	1118	715	2	74	TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf10) alternative variant bSep08, mRNA.
Taf11	Taf11.bSep08	309638	4188	991	4	144	putative protein (Taf11) alternative variant bSep08, mRNA.
Taf11	Taf11.dSep08	309638	1762	566	2	28	putative protein (Taf11) alternative variant dSep08, mRNA.
Taf13	Taf13.aSep08	310784	11374	1363	3	124	TAF13 RNA polymerase II, TATA box binding protein (TBP)-associated factor (14.3 kD) (Taf13) alternative variant aSep08, complete mRNA.
Taf13	Taf13.bSep08	310784	10767	658	2	89	TAF13 RNA polymerase II, TATA box binding protein (TBP)-associated factor (10.6 kD) (Taf13) alternative variant bSep08, mRNA.
Taf15	Taf15.aSep08	287571	31981	2070	16	583	TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf15) alternative variant aSep08, mRNA.
Taf15	Taf15.cSep08	287571	3245	818	3	130	TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor (Taf15) alternative variant cSep08, mRNA.
taflo	taflo.aSep08		6730	465		28	putative protein (taflo) mRNA.
taflu	taflu.aSep08		1150	601			
tagar	tagar.aSep08		3641	421		95	putative protein (tagar) mRNA.
Tagln2	Tagln2.aSep08	304983	5176	1405	5	207	transgelin 2 (23.3 kD) (Tagln2) alternative variant aSep08, mRNA.
Tagln2	Tagln2.bSep08	304983	6014	935	5	199	transgelin 2 (22.4 kD) (Tagln2) alternative variant bSep08, mRNA.
Tagln2	Tagln2.dSep08	304983	1588	749	4	170	transgelin 2 (Tagln2) alternative variant dSep08, mRNA.
Tagln2	Tagln2.eSep08	304983	6146	823	4	153	transgelin 2 (17.4 kD) (Tagln2) alternative variant eSep08, mRNA.
Tagln2	Tagln2.fSep08	304983	3868	365	2	121	transgelin 2 (Tagln2) alternative variant fSep08, mRNA.
Tagln3	Tagln3.aSep08	63837	13035	740	3	246	transgelin 3 (Tagln3) alternative variant aSep08, mRNA.
Tagln3	Tagln3.dSep08	63837	12116	802	2	153	transgelin 3 (Tagln3) alternative variant dSep08, mRNA.
taja	taja.aSep08		1458	414		61	putative protein (taja) mRNA.
tajey	tajey.aSep08		28254	1106		32	putative protein (tajey) mRNA.
takee	takee.aSep08		5308	411	1	136	ezrin-binding PACE-1 like (takee) alternative variant aSep08, mRNA.
takee	takee.bSep08		5238	556	1	110	ezrin-binding PACE-1 like (takee) alternative variant bSep08, mRNA.
takler	takler.aSep08		460	359		32	putative protein (3.8 kD) (takler) mRNA.

Tal1	Tal1.cSep08	313507	3474	1081	3	7	T-cell acute lymphocytic leukemia 1 (Tal1) alternative variant cSep08, mRNA.
Tal1	Tal1.dSep08	313507	3410	215	2	54	T-cell acute lymphocytic leukemia 1 (Tal1) alternative variant dSep08, mRNA.
Taldo1	Taldo1.bSep08	83688	9481	1024	3	108	transaldolase 1 (12.1 kD) (Taldo1) alternative variant bSep08, mRNA.
Talin_middle.0	Talin_middle.0.aSep08		13257	2295	16	665	talin (Talin_middle.0) alternative variant aSep08, mRNA.
Talin_middle.0	Talin_middle.0.bSep08		2786	446	1	70	putative protein (Talin_middle.0) alternative variant bSep08, mRNA.
talo	talo.aSep08		2650	438		48	putative protein (5.3 kD) (talo) mRNA.
tamee	tamee.aSep08		92097	720		66	putative protein (tamee) mRNA.
Tank	Tank.aSep08	252961	79239	2034	8	413	TRAF family member-associated Nf-kappa B activator (Tank) alternative variant aSep08, mRNA.
Tank	Tank.bSep08	252961	60602	680	5	226	TRAF family member-associated Nf-kappa B activator (Tank) alternative variant bSep08, mRNA.
Tank	Tank.cSep08	252961	55154	755	6	187	TRAF family member-associated Nf-kappa B activator (Tank) alternative variant cSep08, mRNA.
Tank	Tank.dSep08	252961	18393	503	4	113	TRAF family member-associated Nf-kappa B activator (Tank) alternative variant dSep08, mRNA.
tanoy	tanoy.aSep08		3994	396		131	dedicator of cytokinesis 10 (tanoy) mRNA.
Taok1	Taok1.bSep08	286993	26769	2895	10	704	TAO kinase 1 (Taok1) alternative variant bSep08, mRNA.
Taok2	Taok2.bSep08	64666	15094	3888	18	1056	tao kinase 2 (Taok2) alternative variant bSep08, mRNA.
Taok2	Taok2.cSep08	64666	1685	1605	2	153	kinase 2 (Taok2) alternative variant cSep08, mRNA.
Taok2	Taok2.dSep08	64666	1694	765	4	114	TAO kinase (Taok2) alternative variant dSep08, mRNA.
Taok3	Taok3.bSep08	304530	31116	685	9	228	TAO kinase 3 (Taok3) alternative variant bSep08, mRNA.
Tap1	Tap1.dSep08	24811	588	336	2	81	transporter 1, ATP-binding cassette, sub-family B (MDR/TAP) (Tap1) alternative variant dSep08, mRNA.
Tap2	Tap2.bSep08	24812	4669	2148	5	315	putative protein (21.6 kD) (Tap2) alternative variant bSep08, mRNA.
Tap2	Tap2.cSep08	24812	1277	599	2	194	transporter 2 (Tap2) alternative variant cSep08, mRNA.
Tap2	Tap2.dSep08	24812	1403	1144	3	169	tap2 (Tap2) alternative variant dSep08, mRNA.
Tap2	Tap2.eSep08	24812	2993	712	3	123	putative protein (Tap2) alternative variant eSep08, mRNA.
Tap2	Tap2.fSep08	24812	2046	1337	2	78	putative mitochondrial protein (9.2 kD) (Tap2) alternative variant fSep08, mRNA.
tapor	tapor.aSep08		1418	394		74	putative protein of mammalian origin (tapor) mRNA.
Tapt1	Tapt1.aSep08	305386	15235	607	4	202	transmembrane anterior posterior transformation 1 (Tapt1) alternative variant aSep08, mRNA.
Tapt1	Tapt1.bSep08	305386	22795	345	2	114	transmembrane anterior posterior transformation 1 (Tapt1) alternative variant bSep08, mRNA.
Tarbp2	Tarbp2.aSep08	363006	1757	1339		146	TAR (HIV) RNA binding protein 2 (15.6 kD) (Tarbp2) mRNA.
tarby	tarby.bSep08		2272	494	4	62	putative protein (tarby) alternative variant bSep08, mRNA.
tarby	tarby.cSep08		21182	546	7	36	putative protein (tarby) alternative variant cSep08, mRNA.
tarby	tarby.dSep08		1860	341	4	49	putative protein (tarby) alternative variant dSep08, mRNA.
tarchy	tarchy.aSep08		1173	722		53	putative protein (6.1 kD) (tarchy) mRNA.

tardar	tardar.aSep08		2291	2099		276	putative protein (tardar) mRNA.
Tardbp	Tardbp.aSep08	298648	10880	2174	5	414	TAR DNA binding protein (44.5 kD) (Tardbp) alternative variant aSep08, complete mRNA.
Tardbp	Tardbp.bSep08	298648	9642	1348	5	298	TAR DNA binding protein (33.6 kD) (Tardbp) alternative variant bSep08, mRNA.
Tardbp	Tardbp.dSep08	298648	4228	655	3	162	TAR DNA binding protein (Tardbp) alternative variant dSep08, mRNA.
Tardbp	Tardbp.eSep08	298648	4229	647	3	145	TAR DNA binding protein (Tardbp) alternative variant eSep08, mRNA.
Tardbp	Tardbp.fSep08	298648	4225	684	3	137	TAR DNA binding protein (Tardbp) alternative variant fSep08, mRNA.
tarflo	tarflo.aSep08		2260	475		35	putative protein (3.9 kD) (tarflo) mRNA.
tarflu	tarflu.aSep08		1838	619		52	putative protein (5.8 kD) (tarflu) mRNA.
targar	targar.aSep08		2576	514		171	receptor beta (targar) mRNA.
tarja	tarja.aSep08		3351	607		94	putative protein (tarja) mRNA.
tarjey	tarjey.aSep08		17715	942		272	platelet-derived growth factor receptor (tarjey) mRNA.
tarkee	tarkee.aSep08		2339	574		59	putative protein (7.0 kD) (tarkee) mRNA.
tarkler	tarkler.aSep08		5419	484		34	putative protein (tarkler) mRNA.
tarlo	tarlo.aSep08		4975	387		66	putative protein (tarlo) mRNA.
tarmee	tarmee.aSep08		5967	386		128	shroom family member 1 CRA b (tarmee) mRNA.
tarnoy	tarnoy.aSep08		4992	697		56	putative protein (6.4 kD) (tarnoy) mRNA.
tarpor	tarpor.aSep08		14503	723		43	putative protein of eukaryotic origin (tarpor) mRNA.
Tars	Tars.aSep08	294810	18847	2601	19	762	threonyl-tRNA synthetase (Tars) alternative variant aSep08, mRNA.
Tars	Tars.bSep08	294810	7150	698	6	216	threonyl-tRNA synthetase (Tars) alternative variant bSep08, mRNA.
Tars2	Tars2.bSep08	310672	9706	730	7	168	threonyl-tRNA synthetase 2 mitochondrial (Tars2) alternative variant bSep08, mRNA.
Tars2	Tars2.cSep08	310672	3833	464	3	154	threonyl-tRNA synthetase mitochondrial (Tars2) alternative variant cSep08, mRNA.
Tars2	Tars2.dSep08	310672	7394	743	6	131	threonyl-tRNA synthetase-like 1 (Tars2) alternative variant dSep08, mRNA.
Tars2	Tars2.eSep08	310672	4928	645	6	124	threonyl-tRNA synthetase-like (13.9 kD) (Tars2) alternative variant eSep08, mRNA.
tarsa	tarsa.bSep08		1235	636	2	62	putative protein (6.5 kD) (tarsa) alternative variant bSep08, mRNA.
tarshee	tarshee.aSep08		6489	735		90	putative protein (tarshee) mRNA.
Tarsl2	Tarsl2.bSep08	308701	19870	715	1	238	threonyl-tRNA synthetase-like 2 (Tarsl2) alternative variant bSep08, mRNA.
tartu	tartu.aSep08		8294	337		98	putative protein of mammalian origin (tartu) mRNA.
tarvar	tarvar.aSep08		1309	276		60	putative protein (tarvar) mRNA.
tarwey	tarwey.aSep08		755	405		134	nucleolar protein 1 120kDa (tarwey) mRNA.
tasa	tasa.aSep08		2980	720		98	putative protein (10.7 kD) (tasa) mRNA.
tashee	tashee.aSep08		89777	959	8	298	CRA a (tashee) alternative variant aSep08, mRNA.

tashee	tashee.bSep08		37178	705	3	40	putative protein (tashee) alternative variant bSep08, mRNA.
tashee	tashee.cSep08		30126	672	2	16	putative protein (tashee) alternative variant cSep08, mRNA.
tashee	tashee.dSep08		30124	670	2	15	putative protein (tashee) alternative variant dSep08, mRNA.
Tat	Tat.bSep08	24813	1550	1160	2	86	tyrosine aminotransferase (Tat) alternative variant bSep08, mRNA.
Tatdn2	Tatdn2.bSep08	500295	3353	882	1	145	TatD-related deoxyribonuclease (Tatdn2) alternative variant bSep08, mRNA.
Tatdn2	Tatdn2.cSep08	500295	9994	2191	4	116	TatD-related deoxyribonuclease (Tatdn2) alternative variant cSep08, mRNA.
tatu	tatu.aSep08		6462	248		82	CRA b like (tatu) mRNA.
tavar	tavar.aSep08		6700	551		47	putative protein (5.5 kD) (tavar) mRNA.
tawby	tawby.aSep08		3419	755		102	putative protein (tawby) mRNA.
tawchy	tawchy.aSep08		3346	662		59	putative protein (tawchy) mRNA.
tawdar	tawdar.aSep08		19334	393		99	WD repeat domain 59 (tawdar) mRNA.
tawey	tawey.aSep08		19048	566		64	putative protein (6.8 kD) (tawey) mRNA.
tawflo	tawflo.aSep08		760	468		121	putative protein (tawflo) mRNA.
tawflu	tawflu.aSep08		1733	377		55	putative protein (tawflu) mRNA.
tawgar	tawgar.aSep08		7435	957		58	putative protein (tawgar) mRNA.
tawja	tawja.aSep08		6074	818		272	CRA a (tawja) mRNA.
tawjey	tawjey.aSep08		6026	365		118	putative protein (tawjey) mRNA.
tawkee	tawkee.aSep08		2832	248		70	soluble adenylyl cyclase (tawkee) mRNA.
tawkler	tawkler.aSep08		2062	600		95	putative protein of mammalian origin (tawkler) mRNA.
tawlo	tawlo.aSep08		3480	2134		52	solute carrier family 7 member (tawlo) mRNA.
tawmee	tawmee.bSep08		746	442	2	111	putative protein (tawmee) alternative variant bSep08, mRNA.
tawnoy	tawnoy.bSep08		10223	461	1	48	putative protein (5.4 kD) (tawnoy) alternative variant bSep08, mRNA.
tawnoy	tawnoy.cSep08		3606	389	1	50	putative protein (tawnoy) alternative variant cSep08, mRNA.
tawpor	tawpor.aSep08		40206	423		141	filamin A interacting protein 1 CRA b (tawpor) mRNA.
tawsa	tawsa.aSep08		2472	501	1	35	putative protein (tawsa) alternative variant aSep08, mRNA.
tawsa	tawsa.bSep08		2086	403	1	36	putative protein (tawsa) alternative variant bSep08, mRNA.
tawshee	tawshee.aSep08		1280	597	2	77	putative protein of mammalian origin (tawshee) alternative variant aSep08, mRNA.
tawtu	tawtu.aSep08		6299	1374		289	putative protein (tawtu) alternative variant aSep08, mRNA.
tawvar	tawvar.aSep08		4586	690		84	putative cytoplasmic protein (9.0 kD) (tawvar) mRNA.
tawwey	tawwey.aSep08		4449	387		67	condensin complex (tawwey) mRNA.
Tax1bp1	Tax1bp1.bSep08	246244	30275	1467	11	488	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant bSep08, mRNA.
Tax1bp1	Tax1bp1.cSep08	246244	26221	1054	9	350	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant cSep08, mRNA.

Tax1bp1	Tax1bp1.dSep08	246244	26229	1021	7	267	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant dSep08, mRNA.
Tax1bp1	Tax1bp1.eSep08	246244	24731	756	8	252	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant eSep08, mRNA.
Tax1bp1	Tax1bp1.fSep08	246244	23182	721	5	206	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant fSep08, mRNA.
Tax1bp1	Tax1bp1.gSep08	246244	7678	407	3	135	tax1 (human T-cell leukemia virus type I) binding protein 1 (Tax1bp1) alternative variant gSep08, mRNA.
Tax1bp3	Tax1bp3.aSep08	360564	4822	1535	4	124	tax1 (human T-cell leukemia virus type I) binding protein 3 (13.7 kD) (Tax1bp3) alternative variant aSep08, mRNA.
Taz	Taz.bSep08	363521	8682	916	9	194	tafazzin (22.4 kD) (Taz) alternative variant bSep08, mRNA.
Taz	Taz.cSep08	363521	5860	700	6	186	tafazzin (Taz) alternative variant cSep08, mRNA.
Taz	Taz.dSep08	363521	4885	606	2	110	tafazzin (Taz) alternative variant dSep08, mRNA.
Taz	Taz.eSep08	363521	7188	1781	7	99	tafazzin (Taz) alternative variant eSep08, mRNA.
TB.0	TB.0.aSep08		10622	2998	12	832	latent transforming growth factor beta binding protein 3 like (TB.0) alternative variant aSep08, mRNA.
TB.0	TB.0.bSep08		553	424	2	111	latent transforming growth factor beta binding protein 3 like (TB.0) alternative variant bSep08, mRNA.
TB.0	TB.0.cSep08		1988	1365	1	108	latent transforming growth factor beta binding protein 3 like (TB.0) alternative variant cSep08, mRNA.
TBC.0	TBC.0.aSep08		21039	1229	2	362	growth hormone regulated TBC protein 1 (TBC.0) alternative variant aSep08, mRNA.
TBC.0	TBC.0.bSep08		18494	748		248	growth hormone regulated tbc protein 1 (TBC.0) alternative variant bSep08, mRNA.
Tbc1d1	Tbc1d1.aSep08	360937	60561	1767	8	555	TBC1 domain family, member 1 (Tbc1d1) alternative variant aSep08, mRNA.
Tbc1d1	Tbc1d1.bSep08	360937	16785	630	1	157	TBC1 domain family, member 1 (Tbc1d1) alternative variant bSep08, mRNA.
Tbc1d2	Tbc1d2.bSep08	313234	47413	4193	13	704	TBC1 domain family, member 2 (80.7 kD) (Tbc1d2) alternative variant bSep08, mRNA.
Tbc1d2	Tbc1d2.cSep08	313234	929	395	2	38	TBC1 domain family, member 2 (Tbc1d2) alternative variant cSep08, mRNA.
Tbc1d2b	Tbc1d2b.bSep08	315880	18998	376	2	125	TBC1 domain family, member 2B (Tbc1d2b) alternative variant bSep08, mRNA.
Tbc1d5	Tbc1d5.bSep08	501088	379377	1790	10	414	TBC1 domain family, member 5 (Tbc1d5) alternative variant bSep08, mRNA.
Tbc1d5	Tbc1d5.cSep08	501088	307768	1575	13	332	TBC1 domain family, member 5 (38.9 kD) (Tbc1d5) alternative variant cSep08, mRNA.
Tbc1d5	Tbc1d5.dSep08	501088	73388	316	3	105	TBC1 domain family, member 5 (Tbc1d5) alternative variant dSep08, mRNA.
Tbc1d7	Tbc1d7.bSep08	361227	17551	1572	6	204	TBC1 domain family, member 7 (23.8 kD) (Tbc1d7) alternative variant bSep08, mRNA.
Tbc1d9b	Tbc1d9b.aSep08	360520	17242	2900	14	762	TBC1 domain family, member 9B (Tbc1d9b) alternative variant aSep08, mRNA.
Tbc1d9b	Tbc1d9b.bSep08	360520	21526	3001	11	692	TBC1 domain family, member 9B (Tbc1d9b) alternative variant bSep08, mRNA.

Tbc1d9b	Tbc1d9b.cSep08	360520	7166	583	6	194	TBC1 domain family, member 9B (Tbc1d9b) alternative variant cSep08, mRNA.
Tbc1d9b	Tbc1d9b.dSep08	360520	3218	572	3	149	TBC1 domain family, member 9B (Tbc1d9b) alternative variant dSep08, mRNA.
Tbc1d10a	Tbc1d10a.bSep08	360968	1379	749	2	176	TBC1 domain family, member 10a (Tbc1d10a) alternative variant bSep08, mRNA.
Tbc1d10b	Tbc1d10b.aSep08	365372	10918	2989	3	717	TBC1 domain family, member 10b (Tbc1d10b) alternative variant aSep08, mRNA.
Tbc1d10b	Tbc1d10b.bSep08	365372	5669	749		249	TBC1 domain family, member 10b (Tbc1d10b) alternative variant bSep08, mRNA.
Tbc1d13	Tbc1d13.aSep08	499768	5356	2824		201	TBC1 domain family, member 13 (Tbc1d13) mRNA.
Tbc1d15	Tbc1d15.aSep08	366896	59149	3682	17	687	tbc1 domain family member 15 (Tbc1d15) alternative variant aSep08, mRNA.
Tbc1d15	Tbc1d15.bSep08	366896	13953	1242	10	307	tbc1 domain family member 15 (36.4 kD) (Tbc1d15) alternative variant bSep08, mRNA.
Tbc1d15	Tbc1d15.cSep08	366896	5981	660	3	97	tbc1 domain family member 15 (Tbc1d15) alternative variant cSep08, mRNA.
Tbc1d15	Tbc1d15.dSep08	366896	1932	516	3	79	tbc1 domain family member 15 (Tbc1d15) alternative variant dSep08, mRNA.
Tbc1d20	Tbc1d20.bSep08	362237	13003	684	5	227	TBC1 domain family, member 20 (Tbc1d20) alternative variant bSep08, mRNA.
Tbc1d20	Tbc1d20.cSep08	362237	12653	596	5	126	TBC1 domain family, member 20 (Tbc1d20) alternative variant cSep08, mRNA.
Tbc1d22b	Tbc1d22b.bSep08	502414	13845	1100		340	TBC1 domain family, member 22B (Tbc1d22b) alternative variant bSep08, mRNA.
Tbc1d23	Tbc1d23.aSep08	304019	58662	3657	18	684	TBC1 domain family, member 23 (76.6 kD) (Tbc1d23) alternative variant aSep08, mRNA.
Tbc1d24	Tbc1d24.bSep08	287110	6646	1784	6	136	TBC1 domain family, member 24 (Tbc1d24) alternative variant bSep08, mRNA.
Tbc1d24	Tbc1d24.cSep08	287110	5519	761	2	45	TBC1 domain family, member 24 (Tbc1d24) alternative variant cSep08, mRNA.
Tbcb	Tbcb.cSep08	292777	1781	599	3	161	tubulin folding cofactor B (Tbcb) alternative variant cSep08, mRNA.
Tbcb	Tbcb.dSep08	292777	2868	1284	4	120	tubulin folding cofactor B (13.3 kD) (Tbcb) alternative variant dSep08, mRNA.
Tbccd1	Tbccd1.bSep08	303830	29017	2546	8	503	putative protein of eukaryotic origin (57.5 kD) (Tbccd1) alternative variant bSep08, mRNA.
Tbccd1	Tbccd1.cSep08	303830	20059	903	2	188	putative protein of metazoan origin (21.8 kD) (Tbccd1) alternative variant cSep08, mRNA.
Tbccd1	Tbccd1.dSep08	303830	13917	368	1	72	putative protein (Tbccd1) alternative variant dSep08, mRNA.
Tbcd	Tbcd.aSep08	360683	21974	729		243	tubulin-specific chaperone d (Tbcd) mRNA.
Tbce	Tbce.bSep08	361255	37749	759	7	155	tubulin-specific chaperone e (Tbce) alternative variant bSep08, mRNA.
Tbce	Tbce.cSep08	361255	8402	1462	9	152	tubulin-specific chaperone e CRA e (17.4 kD) (Tbce) alternative variant cSep08, mRNA.

Tbce	Tbce.dSep08	361255	3818	1108	4	93	tubulin-specific chaperone e (10.4 kD) (Tbce) alternative variant dSep08, mRNA.
Tbce	Tbce.eSep08	361255	3649	727	4	88	tubulin-specific chaperone e (10.2 kD) (Tbce) alternative variant eSep08, mRNA.
Tbce	Tbce.gSep08	361255	784	608	2	59	tubulin-specific chaperone e CRA e (6.8 kD) (Tbce) alternative variant gSep08, mRNA.
Tbk1	Tbk1.bSep08	299827	8202	1383	9	225	TANK-binding kinase 1 like (Tbk1) alternative variant bSep08, mRNA.
Tbk1	Tbk1.cSep08	299827	8796	674	4	184	TANK-binding kinase 1 like (Tbk1) alternative variant cSep08, mRNA.
Tbk1	Tbk1.dSep08	299827	1828	1216	3	94	TANK-binding kinase 1 like (11.5 kD) (Tbk1) alternative variant dSep08, mRNA.
Tbl1xr1	Tbl1xr1.bSep08	365755	42391	205		68	transducin (beta)-like 1X-linked receptor 1 (Tbl1xr1) alternative variant bSep08, mRNA.
Tbl3	Tbl3.bSep08	287120	1288	771	5	218	transducin (beta)-like 3 (Tbl3) alternative variant bSep08, mRNA.
Tbl3	Tbl3.cSep08	287120	1491	704	4	155	transducin (beta)-like 3 (16.9 kD) (Tbl3) alternative variant cSep08, mRNA.
Tbp	Tbp.aSep08	117526	15668	1270	7	335	TATA box binding protein (36.5 kD) (Tbp) alternative variant aSep08, mRNA.
TBP-binding.0	TBP-binding.0.aSep08		5846	478		159	taf1 RNA polymerase II TATA box binding protein - associated factor like (TBP-binding.0) mRNA.
Tbpl1	Tbpl1.aSep08	689030	26263	2834	7	218	TATA box binding protein-like 1 (24.2 kD) (Tbpl1) alternative variant aSep08, complete mRNA.
Tbpl1	Tbpl1.cSep08	689030	822	729	2	145	TATA box binding protein-like 1 (16.3 kD) (Tbpl1) alternative variant cSep08, mRNA.
Tbpl1	Tbpl1.dSep08	689030	24699	1044	7	130	TATA box binding protein-like 1 (14.5 kD) (Tbpl1) alternative variant dSep08, complete mRNA.
Tbpl1	Tbpl1.eSep08	689030	776	649	2	60	TATA box binding protein-like 1 (Tbpl1) alternative variant eSep08, mRNA.
Tbr1	Tbr1.aSep08	680427	2061	912		303	T-box brain gene 1 (Tbr1) mRNA.
Tbrg1	Tbrg1.bSep08	300521	1406	807	2	124	transforming growth factor beta 1 (Tbrg1) alternative variant bSep08, mRNA.
Tbrg1	Tbrg1.cSep08	300521	1099	358	2	95	transforming growth factor beta 1 (Tbrg1) alternative variant cSep08, mRNA.
Tbrg1	Tbrg1.dSep08	300521	3905	532	4	43	putative protein (5.1 kD) (Tbrg1) alternative variant dSep08, complete mRNA.
Tbrg4	Tbrg4.bSep08	360977	2998	574	4	190	transforming growth factor beta regulated gene 4 (Tbrg4) alternative variant bSep08, mRNA.
Tbrg4	Tbrg4.cSep08	360977	1775	788	4	162	transforming growth factor beta regulated gene 4 (Tbrg4) alternative variant cSep08, mRNA.
Tbrg4	Tbrg4.dSep08	360977	3079	381	3	100	transforming growth factor beta regulated gene 4 (Tbrg4) alternative variant dSep08, mRNA.
Tbx2	Tbx2.bSep08	303398	6333	2627	5	485	T-box 2 (50.7 kD) (Tbx2) alternative variant bSep08, mRNA.
Tbx3	Tbx3.bSep08	353305	9816	2368	6	363	T-box 3 (Tbx3) alternative variant bSep08, mRNA.
Tbx3	Tbx3.cSep08	353305	4501	2417	3	260	T-box 3 (Tbx3) alternative variant cSep08, mRNA.



Tbx15	Tbx15.aSep08	295315	47939	1783	3	363	T-box 15 (Tbx15) alternative variant aSep08, mRNA.
Tbx18	Tbx18.bSep08	315870	2076	486	2	161	T-box18 (Tbx18) alternative variant bSep08, mRNA.
Tbx19	Tbx19.bSep08	304935	6139	357	1	84	T-box 19 (Tbx19) alternative variant bSep08, mRNA.
Tbx22	Tbx22.bSep08	302369	5241	787	1	62	T-box 22 (Tbx22) alternative variant bSep08, mRNA.
Tc2n	Tc2n.bSep08	500707	29586	585	5	195	tandem C2 domains, nuclear (Tc2n) alternative variant bSep08, mRNA.
Tc2n	Tc2n.cSep08	500707	13895	414	1	89	tandem C2 domains, nuclear (Tc2n) alternative variant cSep08, mRNA.
Tc2n	Tc2n.dSep08	500707	22567	1006	2	83	tandem C2 domains, nuclear (Tc2n) alternative variant dSep08, mRNA.
Tc2n	Tc2n.eSep08	500707	6779	366	1	77	tandem C2 domains, nuclear (Tc2n) alternative variant eSep08, mRNA.
Tcam1	Tcam1.bSep08	59305	12806	2775	3	319	testicular cell adhesion molecule 1 (35.3 kD) (Tcam1) alternative variant bSep08, mRNA.
Tcea1	Tcea1.aSep08	362479	37256	2364	10	330	transcription elongation factor A (SII) 1 (Tcea1) alternative variant aSep08, mRNA.
Tcea1	Tcea1.bSep08	362479	32949	943	8	269	transcription elongation factor A (SII) 1 (Tcea1) alternative variant bSep08, mRNA.
Tcea1	Tcea1.cSep08	362479	26588	1469	10	263	transcription elongation factor A (SII) 1 (Tcea1) alternative variant cSep08, mRNA.
Tcea1	Tcea1.dSep08	362479	15978	1143	8	256	transcription elongation factor A (SII) 1 (Tcea1) alternative variant dSep08, mRNA.
Tcea1	Tcea1.eSep08	362479	19298	623	2	41	transcription elongation factor A (SII) 1 (Tcea1) alternative variant eSep08, mRNA.
Tcea2	Tcea2.aSep08	29575	4602	822	7	215	transcription elongation factor A (SII), 2 (Tcea2) alternative variant aSep08, mRNA.
Tcea2	Tcea2.bSep08	29575	4130	399	4	132	transcription elongation factor A (SII), 2 (Tcea2) alternative variant bSep08, mRNA.
Tcea2	Tcea2.cSep08	29575	854	779	2	110	transcription elongation factor A (SII), 2 (Tcea2) alternative variant cSep08, mRNA.
Tceal1	Tceal1.bSep08	302593	1097	741	1	164	transcription elongation factor A (SII)-like 1 (Tceal1) alternative variant bSep08, mRNA.
Tceal3	Tceal3.aSep08	501628	1963	1094		64	transcription elongation factor A (SII)-like 3 (Tceal3) mRNA.
Tceb1	Tceb1.bSep08	64525	15096	762	2	112	transcription elongation factor B (SIII), polypeptide 1 (12.5 kD) (Tceb1) alternative variant bSep08, mRNA.
Tceb1	Tceb1.cSep08	64525	15170	1004	2	112	transcription elongation factor B (SIII), polypeptide 1 (12.5 kD) (Tceb1) alternative variant cSep08, mRNA.
Tceb1	Tceb1.dSep08	64525	14132	704	2	33	transcription elongation factor B (SIII), polypeptide 1 (Tceb1) alternative variant dSep08, mRNA.
Tceb3	Tceb3.bSep08	25562	1469	930	2	82	transcription elongation factor B (SIII), polypeptide 3 (Tceb3) alternative variant bSep08, mRNA.
Tcerg1	Tcerg1.bSep08	307474	60935	4100	22	1077	transcription elongation regulator 1 (121.7 kD) (Tcerg1) alternative variant bSep08, mRNA.
Tcerg1	Tcerg1.cSep08	307474	20102	2207	10	378	transcription elongation regulator 1 CRA c (45.6 kD) (Tcerg1) alternative variant cSep08, mRNA.

Tcerg1	Tcerg1.dSep08	307474	21524	1490	12	279	transcription elongation regulator 1 CRA c (33.0 kD) (Tcerg1) alternative variant dSep08, mRNA.
Tcerg1	Tcerg1.eSep08	307474	9165	768	4	232	transcription elongation regulator 1 (Tcerg1) alternative variant eSep08, mRNA.
Tcerg1	Tcerg1.fSep08	307474	5300	927	4	149	transcription elongation regulator 1 (Tcerg1) alternative variant fSep08, mRNA.
Tcerg1	Tcerg1.gSep08	307474	8250	1905	3	78	transcription elongation regulator 1 CRA a (Tcerg1) alternative variant gSep08, mRNA.
Tcf3	Tcf3.bSep08	312451	3687	820	3	160	transcription factor 3 (17.9 kD) (Tcf3) alternative variant bSep08, mRNA.
Tcf4	Tcf4.bSep08	84382	10003	3774	5	221	transcription factor 4 CRA c (Tcf4) alternative variant bSep08, mRNA.
Tcf4	Tcf4.cSep08	84382	57159	556	5	166	transcription factor 4 CRA d (Tcf4) alternative variant cSep08, mRNA.
Tcf4	Tcf4.dSep08	84382	47236	544	4	163	transcription factor 4 CRA a (Tcf4) alternative variant dSep08, mRNA.
Tcf4	Tcf4.eSep08	84382	6974	604	4	158	transcription factor 4 CRA b (Tcf4) alternative variant eSep08, mRNA.
Tcf4	Tcf4.fSep08	84382	1742	409	3	116	transcription factor 4 CRA a (Tcf4) alternative variant fSep08, mRNA.
Tcf4	Tcf4.hSep08	84382	2506	571	2	54	putative protein, with a coiled coil domain (5.9 kD) (Tcf4) alternative variant hSep08, mRNA.
Tcf7	Tcf7.aSep08	363595	9262	2243		214	transcription factor 7, T-cell specific (Tcf7) mRNA.
Tcf12	Tcf12.bSep08	25720	41170	1797	12	512	transcription factor 12 (Tcf12) alternative variant bSep08, mRNA.
Tcf12	Tcf12.cSep08	25720	28531	1097	9	365	transcription factor 12 (Tcf12) alternative variant cSep08, mRNA.
Tcf20	Tcf20.bSep08	366964	30228	514	1	55	transcription factor 20 (Tcf20) alternative variant bSep08, mRNA.
Tcf25	Tcf25.aSep08	292082	34494	5302	18	780	transcription factor 25 (basic helix-loop-helix) (Tcf25) alternative variant aSep08, mRNA.
Tcf25	Tcf25.bSep08	292082	17053	829	7	263	transcription factor 25 (basic helix-loop-helix) (Tcf25) alternative variant bSep08, mRNA.
Tcf25	Tcf25.cSep08	292082	16960	2276	9	244	transcription factor 25 (basic helix-loop-helix) (27.8 kD) (Tcf25) alternative variant cSep08, mRNA.
Tcf25	Tcf25.eSep08	292082	2401	534	3	109	transcription factor 25 (basic helix-loop-helix) (Tcf25) alternative variant eSep08, mRNA.
Tcf25	Tcf25.gSep08	292082	4400	465	4	69	transcription factor 25 (basic helix-loop-helix) (Tcf25) alternative variant gSep08, mRNA.
Tcf25	Tcf25.hSep08	292082	5743	1076	4	46	transcription factor 25 (basic helix-loop-helix) (5.3 kD) (Tcf25) alternative variant hSep08, mRNA.
Tcf25	Tcf25.iSep08	292082	3042	713	2	15	transcription factor 25 (basic helix-loop-helix) (Tcf25) alternative variant iSep08, mRNA.
Tcfap2b	Tcfap2b.bSep08	301285	20405	1249	1	365	transcription factor AP-2 beta (Tcfap2b) alternative variant bSep08, mRNA.
Tcfap2c	Tcfap2c.bSep08	362280	7504	872	4	290	transcription factor AP-2 gamma (Tcfap2c) alternative variant bSep08, mRNA.

Tcfap2c	Tcfap2c.cSep08	362280	3938	765	5	238	transcription factor AP-2 gamma (Tcfap2c) alternative variant cSep08, mRNA.
Tcfap2c	Tcfap2c.dSep08	362280	932	814	2	112	putative mitochondrial protein (12.6 kD) (Tcfap2c) alternative variant dSep08, mRNA.
Tcfap2c	Tcfap2c.fSep08	362280	5858	507	2	80	putative protein (Tcfap2c) alternative variant fSep08, mRNA.
Tcfap4	Tcfap4.bSep08	360482	11154	393	4	64	transcription factor AP4 (Tcfap4) alternative variant bSep08, mRNA.
Tcfcp2	Tcfcp2.bSep08	315309	9716	775	7	209	transcription factor CP2 (Tcfcp2) alternative variant bSep08, mRNA.
Tcfcp2	Tcfcp2.cSep08	315309	19238	491	5	129	transcription factor CP2 (Tcfcp2) alternative variant cSep08, mRNA.
Tcfcp2	Tcfcp2.dSep08	315309	8235	810	5	127	transcription factor CP2 (Tcfcp2) alternative variant dSep08, mRNA.
Tcfe2a	Tcfe2a.bSep08	171046	22256	2505	18	405	transcription factor E2a (42.5 kD) (Tcfe2a) alternative variant bSep08, mRNA.
Tcfe2a	Tcfe2a.cSep08	171046	6823	719	5	239	transcription factor E2a (Tcfe2a) alternative variant cSep08, mRNA.
Tcfe2a	Tcfe2a.dSep08	171046	3004	551	4	183	transcription factor E2a (Tcfe2a) alternative variant dSep08, mRNA.
Tcfe3	Tcfe3.aSep08	317376	9098	2669		437	transcription factor E3 (Tcfe3) mRNA.
Tcfef	Tcfef.bSep08	316214	52039	987	6	236	transcription factor EB (Tcfef) alternative variant bSep08, mRNA.
Tcfef	Tcfef.cSep08	316214	2685	1137	2	189	transcription factor EB (16.5 kD) (Tcfef) alternative variant cSep08, mRNA.
Tcfef	Tcfef.dSep08	316214	5302	308	2	102	transcription factor EB (Tcfef) alternative variant dSep08, mRNA.
Tchp	Tchp.aSep08	304547	4188	1547		128	trichoplein, keratin filament binding (Tchp) mRNA.
Tcirg1	Tcirg1.aSep08	293650	11870	2570	19	499	T-cell immune regulator 1 like (55.9 kD) (Tcirg1) alternative variant aSep08, complete mRNA.
Tcirg1	Tcirg1.bSep08	293650	3203	1398	10	444	T-cell immune regulator 1 like (Tcirg1) alternative variant bSep08, mRNA.
Tcirg1	Tcirg1.cSep08	293650	5070	1345	9	227	T-cell immune regulator 1 like (Tcirg1) alternative variant cSep08, mRNA.
Tcirg1	Tcirg1.dSep08	293650	3958	717	6	211	T-cell immune regulator 1 like (Tcirg1) alternative variant dSep08, mRNA.
Tcirg1	Tcirg1.eSep08	293650	953	567	4	162	T-cell immune regulator 1 like (Tcirg1) alternative variant eSep08, mRNA.
Tcirg1	Tcirg1.fSep08	293650	1172	831	4	160	T-cell immune regulator 1 like (Tcirg1) alternative variant fSep08, mRNA.
Tcirg1	Tcirg1.gSep08	293650	735	653	2	129	CRA a (13.8 kD) (Tcirg1) alternative variant gSep08, mRNA.
Tcirg1	Tcirg1.hSep08	293650	1370	498	3	102	vacuolar proton (Tcirg1) alternative variant hSep08, mRNA.
Tcn2	Tcn2.bSep08	64365	8121	910	6	253	transcobalamin 2 (Tcn2) alternative variant bSep08, mRNA.
Tcn2	Tcn2.cSep08	64365	6856	775	5	195	transcobalamin 2 (Tcn2) alternative variant cSep08, mRNA.

Tcn2	Tcn2.dSep08	64365	6795	711	5	178	transcobalamin 2 (Tcn2) alternative variant dSep08, mRNA.
Tcn2	Tcn2.eSep08	64365	5180	541	4	111	transcobalamin 2 (12.6 kD) (Tcn2) alternative variant eSep08, complete mRNA.
Tcn2	Tcn2.fSep08	64365	4600	394	3	84	transcobalamin 2 (Tcn2) alternative variant fSep08, mRNA.
Tcof1	Tcof1.aSep08	291571	21074	3709	20	1084	treacher Collins Franceschetti syndrome 1, homolog (Tcof1) alternative variant aSep08, mRNA.
Tcof1	Tcof1.cSep08	291571	4604	725	5	241	treacher Collins Franceschetti syndrome 1, homolog (Tcof1) alternative variant cSep08, mRNA.
Tcp1	Tcp1.bSep08	24818	3477	1507	4	273	t-complex protein 1 (Tcp1) alternative variant bSep08, mRNA.
Tcp1	Tcp1.cSep08	24818	2588	285	3	33	t-complex protein 1 (Tcp1) alternative variant cSep08, mRNA.
Tcp10b	Tcp10b.bSep08	308169	23883	1912	7	389	t-complex protein 10b (Tcp10b) alternative variant bSep08, mRNA.
Tcp10b	Tcp10b.cSep08	308169	11796	796	2	265	t-complex protein 10b (Tcp10b) alternative variant cSep08, mRNA.
Tcp10b	Tcp10b.dSep08	308169	13006	693	3	158	t-complex protein 10b (Tcp10b) alternative variant dSep08, mRNA.
Tcp11	Tcp11.bSep08	309641	6892	932	2	310	t-complex protein 11 (Tcp11) alternative variant bSep08, mRNA.
Tcp11	Tcp11.cSep08	309641	8729	736	2	244	t-complex protein 11 (Tcp11) alternative variant cSep08, mRNA.
Tcp111	Tcp111.bSep08	499846	9891	483	1	98	t-complex 11 like 1 (Tcp111) alternative variant bSep08, mRNA.
Tcp112	Tcp112.bSep08	314683	16436	799	6	228	t-complex 11 (mouse) like 2 (Tcp112) alternative variant bSep08, mRNA.
Tcp112	Tcp112.cSep08	314683	8081	168	2	56	t-complex 11 (mouse) like 2 (Tcp112) alternative variant cSep08, mRNA.
Tcrb	Tcrb.aSep08	24820	424251	1408	5	204	T-cell receptor like (Tcrb) alternative variant aSep08, mRNA.
Tcrb	Tcrb.bSep08	24820	441032	1184	7	201	T-cell receptor like (22.4 kD) (Tcrb) alternative variant bSep08, mRNA.
Tcrb	Tcrb.cSep08	24820	5779	724	5	184	T-cell receptor like (Tcrb) alternative variant cSep08, mRNA.
Tcrb	Tcrb.dSep08	24820	3953	428	2	142	T-cell receptor like (Tcrb) alternative variant dSep08, mRNA.
Tcrb	Tcrb.eSep08	24820	508	396		131	tcrb protein (Tcrb) alternative variant eSep08, mRNA.
Tcrb	Tcrb.fSep08	24820	538537	765	4	113	T-cell receptor like (12.8 kD) (Tcrb) alternative variant fSep08, mRNA.
Tcrb	Tcrb.hSep08	24820	1856	694	2	26	putative protein (Tcrb) alternative variant hSep08, mRNA.
Tcta	Tcta.aSep08	306587	3404	1974	2	199	T-cell leukemia translocation altered gene (Tcta) alternative variant aSep08, mRNA.
Tctex-1.0	Tctex-1.0.aSep08		2792	1226		74	tctex-1 (8.5 kD) (Tctex-1.0) mRNA.
Tctex1d1	Tctex1d1.aSep08	362553	13265	816		55	putative protein (6.3 kD) (Tctex1d1) mRNA.
Tctn3	Tctn3.aSep08	309486	6967	1647		372	tectonic family member 3 (Tctn3) mRNA.

Tdg	Tdg.aSep08	114521	19625	2937	10	410	thymine-DNA glycosylase (45.2 kD) (Tdg) alternative variant aSep08, mRNA.
Tdg	Tdg.bSep08	114521	14314	675	5	176	thymine-DNA glycosylase (Tdg) alternative variant bSep08, mRNA.
Tdrd1	Tdrd1.bSep08	292129	3477	772	1	109	putative protein of vertebrate origin (12.2 kD) (Tdrd1) alternative variant bSep08, mRNA.
Tdrd3	Tdrd3.bSep08	306066	63385	700	6	233	putative protein of eukaryotic origin (Tdrd3) alternative variant bSep08, mRNA.
Tdrd3	Tdrd3.cSep08	306066	37870	1780	4	37	putative protein (Tdrd3) alternative variant cSep08, mRNA.
Tdrd3	Tdrd3.dSep08	306066	6809	1394	2	39	putative protein (4.0 kD) (Tdrd3) alternative variant dSep08, mRNA.
Tdrd7	Tdrd7.bSep08	85425	32249	737	5	245	putative protein of metazoan origin (Tdrd7) alternative variant bSep08, mRNA.
Tdrd7	Tdrd7.cSep08	85425	25271	1131	5	201	putative protein of metazoan origin (Tdrd7) alternative variant cSep08, mRNA.
Tdrd9	Tdrd9.aSep08	299343	15390	751		250	atp-dependent rna helicase tdrd9 (Tdrd9) mRNA.
Tdrd12	Tdrd12.aSep08	292813	11492	581		132	tudor domain-containing protein 12 (Tdrd12) mRNA.
Tdrkh	Tdrkh.aSep08	310652	7165	1585	12	499	CRA a (Tdrkh) alternative variant aSep08, mRNA.
Tdrkh	Tdrkh.bSep08	310652	17061	1244	6	296	CRA a (32.4 kD) (Tdrkh) alternative variant bSep08, complete mRNA.
Tdrkh	Tdrkh.cSep08	310652	1919	494	3	164	CRA a (Tdrkh) alternative variant cSep08, mRNA.
Tdrkh	Tdrkh.dSep08	310652	14762	610	3	80	tdrkh protein (9.1 kD) (Tdrkh) alternative variant dSep08, complete mRNA.
Tead1	Tead1.aSep08	361630	75612	374		124	TEA domain family member 1 (Tead1) mRNA.
Tead2	Tead2.bSep08	308582	16815	2093	12	466	TEA domain family member 2 (Tead2) alternative variant bSep08, mRNA.
Tead2	Tead2.cSep08	308582	1673	1149	2	103	TEA domain family member 2 (12.1 kD) (Tead2) alternative variant cSep08, mRNA.
Tead2	Tead2.dSep08	308582	4171	705	3	45	TEA domain family member 2 (Tead2) alternative variant dSep08, mRNA.
Tead3	Tead3.bSep08	294299	1282	1089	3	109	TEA domain family member 3 (12.7 kD) (Tead3) alternative variant bSep08, mRNA.
Tec	Tec.bSep08	84492	4670	462	3	139	cytoplasmic tyrosine kinase, Dscr28C related (Drosophila) (Tec) alternative variant bSep08, mRNA.
Tecta	Tecta.cSep08	300653	5063	695	3	78	tectorin alpha (9.1 kD) (Tecta) alternative variant cSep08, mRNA.
teeby	teeby.aSep08		2375	278		92	nik related kinase (teeby) mRNA.
teechy	teechy.aSep08		9576	930	4	309	eukaryotic translation initiation factor 2 alpha kinase 4 CRA a (teechy) mRNA.
teedar	teedar.aSep08		9299	692		48	putative protein (5.8 kD) (teedar) mRNA.
teeflo	teeflo.aSep08		5181	596		198	CTCL tumor antigen like (teeflo) mRNA.
teeflu	teeflu.aSep08		19478	1785		75	putative protein (teeflu) mRNA.
teegar	teegar.aSep08		815	479		97	putative protein (teegar) mRNA.
teeja	teeja.aSep08		6014	334		110	retinitis pigmentosa GTPase regulator interacting protein 1 like (teeja) mRNA.

teejey	teejey.aSep08		3137	622		37	putative protein (4.3 kD) (teejey) mRNA.
teekee	teekee.aSep08		3719	742		163	oct-1 (teekee) mRNA.
teekler	teekler.aSep08		32289	553		93	putative protein (teekler) mRNA.
teelo	teelo.aSep08		694	598		35	putative protein (4.0 kD) (teelo) mRNA.
teemee	teemee.aSep08		7563	850		263	kinesin family member 3A (teemee) mRNA.
teenoy	teenoy.aSep08		2473	223		22	putative protein (teenoy) mRNA.
teepor	teepor.aSep08		4687	358	1	52	myosin VI (teepor) alternative variant aSep08, mRNA.
teepor	teepor.bSep08		7855	323	2	59	myosin VI (teepor) alternative variant bSep08, mRNA.
teesa	teesa.aSep08		21997	679		32	putative protein (3.6 kD) (teesa) mRNA.
teeshee	teeshee.aSep08		1614	287		44	putative protein (teeshee) mRNA.
teetu	teetu.aSep08		16215	294		28	putative protein (teetu) mRNA.
teevar	teevar.aSep08		13842	488		73	ac1576 like (teevar) mRNA.
teewey	teewey.aSep08		2489	572		64	putative protein (7.3 kD) (teewey) mRNA.
Tef	Tef.bSep08	29362	20667	613	2	204	thyrotroph embryonic factor (Tef) alternative variant bSep08, mRNA.
Tef	Tef.cSep08	29362	1715	1255	1	101	thyrotroph embryonic factor (Tef) alternative variant cSep08, mRNA.
Tegt	Tegt.bSep08	24822	16260	2406	10	237	testis enhanced gene transcript (26.5 kD) (Tegt) alternative variant bSep08, complete mRNA.
Tegt	Tegt.cSep08	24822	13442	773	9	229	testis enhanced gene transcript (Tegt) alternative variant cSep08, mRNA.
Tegt	Tegt.dSep08	24822	12485	782	8	204	testis enhanced gene transcript (Tegt) alternative variant dSep08, mRNA.
Tegt	Tegt.eSep08	24822	15774	656	5	122	testis enhanced gene transcript (13.3 kD) (Tegt) alternative variant eSep08, mRNA.
Tek	Tek.bSep08	89804	3336	371	1	92	endothelial-specific receptor tyrosine kinase (Tek) alternative variant bSep08, mRNA.
Tekt2	Tekt2.bSep08	298532	1333	719	1	122	tektin 2 (Tekt2) alternative variant bSep08, mRNA.
Tekt3	Tekt3.bSep08	287392	7585	508	1	129	tektin 3 (Tekt3) alternative variant bSep08, mRNA.
Telo2	Telo2.aSep08	302986	15213	3210		834	TEL2, telomere maintenance 2, homolog ( <i>S. cerevisiae</i> ) (93.1 kD) (Telo2) mRNA.
Tenc1	Tenc1.bSep08	315326	1696	769	6	255	putative protein of eukaryotic origin (Tenc1) alternative variant bSep08, mRNA.
Tenc1	Tenc1.cSep08	315326	1719	1015	5	157	SH2 motif (Tenc1) alternative variant cSep08, mRNA.
Tenc1	Tenc1.dSep08	315326	1281	314	5	104	putative protein of eukaryotic origin (Tenc1) alternative variant dSep08, mRNA.
Tenc1	Tenc1.eSep08	315326	950	295	4	98	SH2 motif (Tenc1) alternative variant eSep08, mRNA.
Tep1	Tep1.aSep08	64523	14845	2448	17	777	telomerase associated protein 1 (Tep1) alternative variant aSep08, mRNA.
Tep1	Tep1.bSep08	64523	2113	414	2	138	telomerase associated protein 1 (Tep1) alternative variant bSep08, mRNA.
Tep1	Tep1.cSep08	64523	1023	420	4	107	telomerase associated protein 1 (Tep1) alternative variant cSep08, mRNA.
Tepp	Tepp.bSep08	291850	1280	375	1	74	testis/prostate/placenta-expressed protein (8.5 kD) (Tepp) alternative variant bSep08, mRNA.

terby	terby.aSep08		2675	830		79	nik related kinase (terby) mRNA.
terdar	terdar.aSep08		4384	707		132	CRA b like (terdar) mRNA.
Terf1	Terf1.bSep08	297758	12214	358	4	119	telomeric repeat binding factor 1 (Terf1) alternative variant bSep08, mRNA.
Terf1	Terf1.cSep08	297758	1739	792	2	30	telomeric repeat binding factor 1 (3.5 kD) (Terf1) alternative variant cSep08, mRNA.
Terf2	Terf2.bSep08	361403	20745	2475	7	478	telomeric repeat binding factor 2 (Terf2) alternative variant bSep08, mRNA.
Terf2ip	Terf2ip.cSep08	307861	5419	1605	3	74	telomeric repeat binding factor 2, interacting protein (Terf2ip) alternative variant cSep08, mRNA.
Terf2ip	Terf2ip.dSep08	307861	5401	1584	3	68	telomeric repeat binding factor 2, interacting protein (Terf2ip) alternative variant dSep08, mRNA.
terflo	terflo.aSep08		8655	392		68	oocyte-testis gene 1 like (terflo) mRNA.
terflu	terflu.aSep08		4739	506		71	putative protein (terflu) mRNA.
tergar	tergar.aSep08		1802	885	4	208	regulator of nonsense transcripts 1 (tergar) alternative variant aSep08, mRNA.
tergar	tergar.bSep08		501	211	1	65	putative protein (tergar) alternative variant bSep08, mRNA.
terja	terja.aSep08		6828	618		60	putative protein (6.5 kD) (terja) mRNA.
terjey	terjey.aSep08		3745	265		34	putative protein (terjey) mRNA.
terkee	terkee.aSep08		5600	1209		38	putative protein (4.5 kD) (terkee) mRNA.
terkler	terkler.aSep08		689	566		42	putative protein (4.8 kD) (terkler) mRNA.
terlo	terlo.aSep08		2119	1042		17	putative protein (2.0 kD) (terlo) mRNA.
termee	termee.aSep08		10492	460		46	CRA c like (5.0 kD) (termee) mRNA.
ternoy	ternoy.aSep08		7466	696		231	thyroid hormone receptor interactor 12 CRA f (ternoy) mRNA.
terpor	terpor.aSep08		7608	419		139	myosin VI (terpor) mRNA.
tersa	tersa.aSep08		702	250		41	putative protein (4.4 kD) (tersa) mRNA.
tershee	tershee.aSep08		49536	668	2	184	putative protein of vertebrate origin (tershee) alternative variant aSep08, mRNA.
tershee	tershee.bSep08		50204	451	2	129	CRA b like (tershee) alternative variant bSep08, mRNA.
Tert	Tert.bSep08	301965	1818	540	2	107	telomerase reverse transcriptase (12.0 kD) (Tert) alternative variant bSep08, mRNA.
tertu	tertu.aSep08		8672	669		190	WD repeat domain 35 (tertu) mRNA.
tervar	tervar.aSep08		912	814		123	putative protein (14.0 kD) (tervar) mRNA.
terwey	terwey.aSep08		5466	406		135	von Willebrand factor (terwey) mRNA.
Tesb	Tesb.bSep08	414826	14539	1500	6	273	tesb pseudogene (31.2 kD) (Tesb) alternative variant bSep08, mRNA.
Tesb	Tesb.dSep08	414826	5256	678	4	107	putative nuclear protein (11.7 kD) (Tesb) alternative variant dSep08, mRNA.
Tesb	Tesb.eSep08	414826	9307	319	4	76	putative protein (Tesb) alternative variant eSep08, mRNA.
Tesb	Tesb.fSep08	414826	3603	645	4	72	tesb pseudogene like (8.4 kD) (Tesb) alternative variant fSep08, mRNA.
Tesb	Tesb.gSep08	414826	14974	751	10	66	testis specific basic protein (Tesb) alternative variant gSep08, mRNA.

Tesc	Tesc.aSep08	288689	33946	899	8	244	tescalcin (Tesc) alternative variant aSep08, mRNA.
Tesc	Tesc.cSep08	288689	5896	540	5	105	tescalcin (12.1 kD) (Tesc) alternative variant cSep08, mRNA.
Tesk2	Tesk2.bSep08	170908	32770	752	3	110	testis-specific kinase 2 (12.1 kD) (Tesk2) alternative variant bSep08, mRNA.
Tessp2	Tessp2.bSep08	301027	2822	864	1	185	testis serine protease 2 (Tessp2) alternative variant bSep08, mRNA.
Tet1	Tet1.bSep08	309902	5237	423	5	140	tet oncogene 1 (Tet1) alternative variant bSep08, mRNA.
Tex2	Tex2.aSep08	303611	63963	5194	11	1061	testis expressed 2 (Tex2) alternative variant aSep08, mRNA.
Tex2	Tex2.bSep08	303611	8430	1011	3	138	testis expressed 2 (16.1 kD) (Tex2) alternative variant bSep08, mRNA.
Tex2	Tex2.cSep08	303611	2865	388	2	82	putative protein of metazoan origin (Tex2) alternative variant cSep08, mRNA.
Tex10	Tex10.bSep08	298065	18093	870	5	252	hypothetical protein LOC680809 (Tex10) alternative variant bSep08, mRNA.
Tex10	Tex10.bSep08	680809	18093	870	5	252	hypothetical protein LOC680809 (Tex10) alternative variant bSep08, mRNA.
Tex10	Tex10.cSep08	298065	4363	498	4	69	hypothetical protein LOC680809 (Tex10) alternative variant cSep08, mRNA.
Tex10	Tex10.cSep08	680809	4363	498	4	69	hypothetical protein LOC680809 (Tex10) alternative variant cSep08, mRNA.
Tex11	Tex11.aSep08	501588	78261	785		261	putative protein of metazoan origin (Tex11) mRNA.
Tex12	Tex12.aSep08	690393	3638	399		95	putative protein, with a coiled coil domain, of mammalian origin (Tex12) mRNA.
Tex13	Tex13.aSep08	680865	2755	611		203	putative protein, with a coiled coil domain, of mammalian origin (Tex13) mRNA.
Tex14	Tex14.aSep08	287603	2957	447		46	testis expressed 14 (Tex14) mRNA.
Tex264	Tex264.bSep08	300988	26650	765	5	214	testis expressed 264 (23.7 kD) (Tex264) alternative variant bSep08, mRNA.
Tex264	Tex264.cSep08	300988	23316	751	5	187	testis expressed 264 (Tex264) alternative variant cSep08, mRNA.
Tex264	Tex264.dSep08	300988	22950	974	5	184	testis expressed 264 (Tex264) alternative variant dSep08, mRNA.
Tex264	Tex264.eSep08	300988	26454	722	5	133	testis expressed 264 (Tex264) alternative variant eSep08, mRNA.
Tex264	Tex264.fSep08	300988	22424	392	3	82	testis expressed 264 (Tex264) alternative variant fSep08, mRNA.
Tex264	Tex264.hSep08	300988	2523	402	2	58	putative protein (Tex264) alternative variant hSep08, mRNA.
teyby	teyby.aSep08		1917	418		139	putative protein of mammalian origin (teyby) mRNA.
teychy	teychy.aSep08		1400	375		89	putative protein, with a coiled coil domain, of vertebrate origin (teychy) mRNA.
teydar	teydar.aSep08		791	178		47	putative protein (teydar) mRNA.
teyflo	teyflo.aSep08		1953	334		110	actin filament associated Protein 1-like 2 (teyflo) mRNA.
teyflu	teyflu.aSep08		1634	217		72	phosphodiesterase 8A CRA b (teyflu) mRNA.



teygar	teygar.aSep08		3100	1773		96	putative nuclear protein (10.2 kD) (teygar) mRNA.
teyja	teyja.aSep08		41941	818		21	putative protein (2.3 kD) (teyja) mRNA.
teyjey	teyjey.bSep08		982	564	2	39	putative protein (teyjey) alternative variant bSep08, mRNA.
teykee	teykee.aSep08		6773	651		217	dual specificity phosphatase 27 (teykee) mRNA.
teykler	teykler.aSep08		4367	484		34	putative protein (teykler) mRNA.
teylo	teylo.aSep08		1953	771		61	putative protein (6.8 kD) (teylo) mRNA.
teymee	teymee.aSep08		1553	500		81	putative protein (teymee) mRNA.
teynoy	teynoy.aSep08		26307	617	6	205	thyroid hormone receptor interactor 12 CRA c (teynoy) alternative variant aSep08, mRNA.
teypor	teypor.aSep08		23684	2578	4	336	myosin VI (teypor) alternative variant aSep08, mRNA.
teypor	teypor.bSep08		18296	611	3	203	myosin VI (teypor) alternative variant bSep08, mRNA.
teypor	teypor.cSep08		8443	419	2	139	myosin VI (teypor) alternative variant cSep08, mRNA.
teysa	teysa.aSep08		1576	400		67	death-inducing-protein CRA c (teysa) mRNA.
teyshee	teyshee.aSep08		17265	691		230	putative protein of vertebrate origin (teyshee) mRNA.
teytu	teytu.aSep08		4634	347		115	WD repeat domain 35 (teytu) mRNA.
teyvar	teyvar.aSep08		1320	418		80	putative protein (teyvar) mRNA.
teywey	teywey.aSep08		2423	177		58	von Willebrand factor (teywey) mRNA.
Tfb1m	Tfb1m.bSep08	308140	40078	754	6	107	transcription factor B1, mitochondrial (11.8 kD) (Tfb1m) alternative variant bSep08, mRNA.
Tfb2m	Tfb2m.bSep08	289307	5246	1159	5	134	transcription factor B2 mitochondrial (15.9 kD) (Tfb2m) alternative variant bSep08, mRNA.
Tfb2m	Tfb2m.cSep08	289307	4641	546	3	104	transcription factor B2 mitochondrial (Tfb2m) alternative variant cSep08, mRNA.
Tfb2m	Tfb2m.dSep08	289307	5103	773	2	87	transcription factor B2 mitochondrial (Tfb2m) alternative variant dSep08, mRNA.
Tfdp2	Tfdp2.aSep08	300947	69707	2372	8	385	transcription factor Dp 2 (42.9 kD) (Tfdp2) alternative variant aSep08, mRNA.
Tfdp2	Tfdp2.cSep08	300947	77397	646	3	142	transcription factor Dp 2 (Tfdp2) alternative variant cSep08, mRNA.
Tfg	Tfg.bSep08	360709	21834	1564	7	372	trk-fused gene (Tfg) alternative variant bSep08, mRNA.
Tfg	Tfg.cSep08	360709	15149	971	5	285	trk-fused gene (30.2 kD) (Tfg) alternative variant cSep08, mRNA.
Tfg	Tfg.dSep08	360709	13957	937	5	267	trk-fused gene (Tfg) alternative variant dSep08, mRNA.
Tfg	Tfg.fSep08	360709	1311	598	2	140	trk-fused gene (Tfg) alternative variant fSep08, mRNA.
TFIIS_M.0	TFIIS_M.0.aSep08		13662	980	5	210	death (TFIIS_M.0) alternative variant aSep08, mRNA.
TFIIS_M.0	TFIIS_M.0.bSep08		756	550	2	183	death (TFIIS_M.0) alternative variant bSep08, mRNA.
Tfpi	Tfpi.aSep08	29436	43041	1302	9	306	tissue factor pathway inhibitor precursor (34.9 kD) (Tfpi) alternative variant aSep08, complete mRNA.
Tfpi	Tfpi.cSep08	29436	35802	3039	8	232	tissue factor pathway inhibitor precursor (26.5 kD) (Tfpi) alternative variant cSep08, complete mRNA.
Tfpi	Tfpi.dSep08	29436	41413	852	8	229	tissue factor pathway inhibitor (Tfpi) alternative variant dSep08, mRNA.
Tfpi	Tfpi.eSep08	29436	26647	723	5	131	putative protein (Tfpi) alternative variant eSep08, mRNA.

Tfpi2	Tfpi2.bSep08	286926	3340	620	3	166	tissue factor pathway inhibitor 2 (Tfpi2) alternative variant bSep08, mRNA.
Tfpt	Tfpt.aSep08	85423	9450	797	6	253	TCF3 (E2A) fusion partner (Tfpt) alternative variant aSep08, mRNA.
Tfpt	Tfpt.cSep08	85423	8848	601	4	189	TCF3 (E2A) fusion partner (Tfpt) alternative variant cSep08, mRNA.
Tfpt	Tfpt.dSep08	85423	8970	824	5	179	TCF3 (E2A) fusion partner (Tfpt) alternative variant dSep08, mRNA.
Tfrc	Tfrc.aSep08	64678	21839	2140		563	transferrin receptor (Tfrc) alternative variant aSep08, mRNA.
Tfrc	Tfrc.bSep08	64678	10609	3475	2	368	transferrin receptor (Tfrc) alternative variant bSep08, mRNA.
Tfrc	Tfrc.cSep08	64678	1451	443	1	55	transferrin receptor (6.5 kD) (Tfrc) alternative variant cSep08, mRNA.
Tg	Tg.bSep08	24826	30140	1048	1	349	thyroglobulin (Tg) alternative variant bSep08, mRNA.
Tgds	Tgds.aSep08	306164	21059	1808	11	354	TDP-glucose 4,6-dehydratase (Tgds) alternative variant aSep08, mRNA.
Tgds	Tgds.cSep08	306164	976	311	1	74	TDP-glucose 4,6-dehydratase (Tgds) alternative variant cSep08, mRNA.
Tgfb1i1	Tgfb1i1.aSep08	84574	6021	2091	11	473	transforming growth factor beta 1 induced transcript 1 (51.4 kD) (Tgfb1i1) alternative variant aSep08, mRNA.
Tgfb1i1	Tgfb1i1.bSep08	84574	5178	818	7	272	transforming growth factor beta 1 induced transcript 1 (Tgfb1i1) alternative variant bSep08, mRNA.
Tgfb1i1	Tgfb1i1.cSep08	84574	2456	746	7	237	transforming growth factor beta 1 induced transcript 1 (Tgfb1i1) alternative variant cSep08, mRNA.
Tgfb1i1	Tgfb1i1.dSep08	84574	5033	678	7	225	transforming growth factor beta 1 induced transcript 1 (Tgfb1i1) alternative variant dSep08, mRNA.
Tgfb1i1	Tgfb1i1.fSep08	84574	2869	340	2	113	transforming growth factor beta 1 induced transcript 1 (Tgfb1i1) alternative variant fSep08, mRNA.
Tgfb1i1	Tgfb1i1.gSep08	84574	2039	434	6	106	transforming growth factor beta 1 induced transcript 1 (Tgfb1i1) alternative variant gSep08, mRNA.
Tgfb2	Tgfb2.bSep08	81809	95816	1569	6	362	transforming growth factor, beta 2 (Tgfb2) alternative variant bSep08, mRNA.
Tgfb2	Tgfb2.dSep08	81809	67179	374	3	56	transforming growth factor, beta 2 (Tgfb2) alternative variant dSep08, mRNA.
Tgfb3	Tgfb3.bSep08	25717	10400	399	4	109	transforming growth factor, beta 3 (Tgfb3) alternative variant bSep08, mRNA.
Tgfb3	Tgfb3.cSep08	25717	2952	406	3	75	transforming growth factor, beta 3 (Tgfb3) alternative variant cSep08, mRNA.
Tgfb1	Tgfb1.aSep08	116487	11820	2368	11	562	transforming growth factor, beta induced (Tgfb1) alternative variant aSep08, mRNA.
Tgfb1	Tgfb1.bSep08	116487	7476	1374	8	253	transforming growth factor, beta induced (Tgfb1) alternative variant bSep08, mRNA.
Tgfb3	Tgfb3.bSep08	29610	10195	866	6	288	transforming growth factor, beta receptor III (Tgfb3) alternative variant bSep08, mRNA.
Tgfb3	Tgfb3.cSep08	29610	45719	686	5	174	transforming growth factor, beta receptor III (Tgfb3) alternative variant cSep08, mRNA.

TGF_beta.1	TGF_beta.1.aSep08		10419	750	2	249	myostatin (TGF_beta.1) alternative variant aSep08, mRNA.
TGF_beta.1	TGF_beta.1.bSep08		4805	434	1	144	growth differentiation (TGF_beta.1) alternative variant bSep08, mRNA.
TGF_beta.2	TGF_beta.2.aSep08		15908	620	3	178	bone morphogenetic protein 8 (TGF_beta.2) alternative variant aSep08, mRNA.
TGF_beta.2	TGF_beta.2.bSep08		2516	817	1	112	bone morphogenetic protein 8 (TGF_beta.2) alternative variant bSep08, mRNA.
Tgif1	Tgif1.aSep08	316742	82362	1419	3	313	TG interacting factor 1 (Tgif1) alternative variant aSep08, mRNA.
Tgif1	Tgif1.cSep08	316742	6592	972	3	222	TG interacting factor 1 (Tgif1) alternative variant cSep08, mRNA.
Tgif1	Tgif1.dSep08	316742	5136	715	3	198	TG interacting factor 1 (Tgif1) alternative variant dSep08, mRNA.
Tgif1	Tgif1.eSep08	316742	1977	1110	2	168	TG interacting factor 1 (Tgif1) alternative variant eSep08, mRNA.
Tgm1	Tgm1.aSep08	60335	13433	2700	15	843	transglutaminase 1, K polypeptide (Tgm1) alternative variant aSep08, mRNA.
Tgm1	Tgm1.cSep08	60335	3686	803	5	198	transglutaminase 1, K polypeptide (Tgm1) alternative variant cSep08, mRNA.
Tgm1	Tgm1.dSep08	60335	4449	345	3	105	transglutaminase 1, K polypeptide (Tgm1) alternative variant dSep08, mRNA.
Tgm2	Tgm2.bSep08	56083	6862	719	3	191	transglutaminase 2, C polypeptide (Tgm2) alternative variant bSep08, mRNA.
Tgm2	Tgm2.cSep08	56083	7619	354	3	83	transglutaminase 2, C polypeptide (Tgm2) alternative variant cSep08, mRNA.
Tgm2	Tgm2.dSep08	56083	1517	675		80	transglutaminase 2, C polypeptide (Tgm2) alternative variant dSep08, mRNA.
Tgm2	Tgm2.fSep08	56083	2945	410	3	63	transglutaminase 2, C polypeptide (Tgm2) alternative variant fSep08, mRNA.
Tgm4	Tgm4.bSep08	64679	23850	737	7	245	transglutaminase 4 (prostate) (Tgm4) alternative variant bSep08, mRNA.
Tgm4	Tgm4.cSep08	64679	38203	2764	5	242	transglutaminase 4 (prostate) (27.0 kD) (Tgm4) alternative variant cSep08, complete mRNA.
Tgm4	Tgm4.dSep08	64679	3908	633	5	210	transglutaminase 4 (prostate) (Tgm4) alternative variant dSep08, mRNA.
Tgm4	Tgm4.eSep08	64679	22522	799	7	210	transglutaminase 4 (prostate) (Tgm4) alternative variant eSep08, mRNA.
Tgm4	Tgm4.fSep08	64679	23823	598	6	194	transglutaminase 4 (prostate) (Tgm4) alternative variant fSep08, mRNA.
Tgm4	Tgm4.gSep08	64679	22679	766	7	89	transglutaminase 4 (prostate) (10.1 kD) (Tgm4) alternative variant gSep08, mRNA.
Tgm5	Tgm5.aSep08	691929	2274	1081		262	transglutaminase 5 (Tgm5) mRNA.
Th	Th.bSep08	25085	870	257	1	85	tyrosine hydroxylase (Th) alternative variant bSep08, mRNA.
Thada	Thada.aSep08	313865	99873	3548	10	573	thyroid adenoma associated (Thada) mRNA.
Thap3	Thap3.cSep08	362667	2325	642	2	83	zinc finger, C2CH-type (Thap3) alternative variant cSep08, mRNA.

Thap4	Thap4.bSep08	363291	25242	729	4	187	thap 4 (Thap4) alternative variant bSep08, mRNA.
Thap4	Thap4.cSep08	363291	25235	722	4	180	thap 4 (Thap4) alternative variant cSep08, mRNA.
Thap4	Thap4.dSep08	363291	32337	409	2	136	putative protein, with a coiled coil domain, of ancient origin (Thap4) alternative variant dSep08, mRNA.
Thap4	Thap4.eSep08	363291	25005	388	3	119	putative protein of ancient origin (Thap4) alternative variant eSep08, mRNA.
Thap6	Thap6.bSep08	305244	12139	2661	4	175	putative nuclear protein, with a coiled coil domain, of vertebrate origin (20.2 kD) (Thap6) alternative variant bSep08, mRNA.
Thap6	Thap6.cSep08	305244	17879	437	4	82	putative cytoplasmic protein of vertebrate origin (9.4 kD) (Thap6) alternative variant cSep08, mRNA.
Thap7	Thap7.aSep08	287944	2944	1612	3	356	zinc finger, C2CH-type (39.7 kD) (Thap7) alternative variant aSep08, mRNA.
Thap7	Thap7.cSep08	287944	2224	595	4	197	zinc finger, C2CH-type (Thap7) alternative variant cSep08, mRNA.
Thap7	Thap7.dSep08	287944	1853	1293	3	139	putative protein human specific (Thap7) alternative variant dSep08, mRNA.
Thbs2	Thbs2.aSep08	292406	6200	2709		240	thrombospondin 2 (Thbs2) mRNA.
Thbs4	Thbs4.aSep08	29220	21192	1723		573	thrombospondin 4 (Thbs4) mRNA.
Theg	Theg.aSep08	299599	9373	1209	4	352	theg homolog (Theg) alternative variant aSep08, mRNA.
Theg	Theg.cSep08	299599	6639	632	1	159	putative protein of metazoan origin (Theg) alternative variant cSep08, mRNA.
Them4	Them4.bSep08	361992	14616	720	1	117	thioesterase superfamily member 4 (Them4) alternative variant bSep08, mRNA.
Thg1l	Thg1l.bSep08	303067	5443	951	5	269	tRNA-histidine guanylyltransferase 1-like ( <i>S. cerevisiae</i> ) (31.4 kD) (Thg1l) alternative variant bSep08, mRNA.
Thiol-ester_cl.0	Thiol-ester_cl.0.aSep08		3067	380		126	putative protein of ancient origin (Thiol-ester_cl.0) mRNA.
Thiolase_C.0	Thiolase_C.0.aSep08		15508	639		161	acetyl-Coenzyme a acetyltransferase 2 (Thiolase_C.0) mRNA.
Thiolase_C.1	Thiolase_C.1.aSep08		724	405		76	acetyl-Coenzyme a acetyltransferase 2 (Thiolase_C.1) mRNA.
Thioredoxin.0	Thioredoxin.0.aSep08		28125	2613	10	434	thioredoxin domain containing protein (Thioredoxin.0) alternative variant aSep08, mRNA.
Thioredoxin.0	Thioredoxin.0.bSep08		1840	618	3	47	putative protein of fungal and metazoan origin (5.2 kD) (Thioredoxin.0) alternative variant bSep08, mRNA.
Thnsl2	Thnsl2.bSep08	297332	970	616	2	125	threonine synthase-like 2 (bacterial) (Thnsl2) alternative variant bSep08, mRNA.
Thoc1	Thoc1.aSep08	291797	35527	3243	21	657	THO complex 1 (75.4 kD) (Thoc1) alternative variant aSep08, mRNA.
Thoc1	Thoc1.cSep08	291797	14208	778	7	153	THO complex 1 (Thoc1) alternative variant cSep08, mRNA.
Thoc1	Thoc1.dSep08	291797	6917	698	3	85	THO complex 1 (10.0 kD) (Thoc1) alternative variant dSep08, mRNA.
Thoc1	Thoc1.eSep08	291797	1145	791	2	49	THO complex 1 (5.3 kD) (Thoc1) alternative variant eSep08, mRNA.

Thoc2	Thoc2.aSep08	313308	32757	3063	13	453	THO complex 2 (Thoc2) alternative variant aSep08, mRNA.
Thoc2	Thoc2.bSep08	313308	15050	768	3	225	THO complex 2 (Thoc2) alternative variant bSep08, mRNA.
Thoc4	Thoc4.aSep08	690585	4074	1532	1	259	THO complex 4 (Thoc4) alternative variant aSep08, mRNA.
Thoc5	Thoc5.bSep08	360972	15530	941	10	313	THO complex 5 (Thoc5) alternative variant bSep08, mRNA.
Thoc5	Thoc5.cSep08	360972	8000	708	7	236	THO complex 5 (Thoc5) alternative variant cSep08, mRNA.
Thoc5	Thoc5.dSep08	360972	3699	648	4	151	THO complex 5 (17.9 kD) (Thoc5) alternative variant dSep08, mRNA.
Thoc7	Thoc7.aSep08	305714	15897	1324	7	152	THO complex 7 homolog (Drosophila) (17.8 kD) (Thoc7) alternative variant aSep08, mRNA.
Thoc7	Thoc7.bSep08	305714	767	210	1	69	THO complex 7 homolog (Drosophila) (Thoc7) alternative variant bSep08, mRNA.
Thop1	Thop1.bSep08	64517	3455	691	5	188	thimet oligopeptidase 1 (Thop1) alternative variant bSep08, mRNA.
Thop1	Thop1.cSep08	64517	7774	750	5	186	thimet oligopeptidase 1 (Thop1) alternative variant cSep08, mRNA.
Thop1	Thop1.dSep08	64517	649	374	3	112	thimet oligopeptidase 1 (Thop1) alternative variant dSep08, mRNA.
Thpo	Thpo.bSep08	81811	814	698	2	122	thrombopoietin (Thpo) alternative variant bSep08, mRNA.
Thpo	Thpo.cSep08	81811	4257	728	5	106	thrombopoietin (Thpo) alternative variant cSep08, mRNA.
Thra	Thra.bSep08	81812	1518	1033	2	144	thyroid hormone receptor alpha (Thra) alternative variant bSep08, mRNA.
Thrap3	Thrap3.bSep08	313591	14443	3706	8	463	thyroid hormone receptor associated protein 3 (53.7 kD) (Thrap3) alternative variant bSep08, mRNA.
Thrap3	Thrap3.cSep08	313591	8940	1596	5	201	thyroid hormone receptor associated protein 3 (Thrap3) alternative variant cSep08, mRNA.
Thrsp	Thrsp.bSep08	25357	4169	696	2	89	thyroid hormone responsive (10.2 kD) (Thrsp) alternative variant bSep08, complete mRNA.
Thsd1	Thsd1.bSep08	364630	18746	2647	1	797	thrombospondin, type I, domain 1 (88.7 kD) (Thsd1) alternative variant bSep08, mRNA.
Thsd7a	Thsd7a.aSep08	500032	22870	536	3	178	thrombospondin type-1 domain-containing protein 7A (Thsd7a) alternative variant aSep08, mRNA.
Thsd7a	Thsd7a.bSep08	500032	120266	517	2	171	thrombospondin type-1 domain-containing protein 7A (Thsd7a) alternative variant bSep08, mRNA.
Thumpd1	Thumpd1.bSep08	309041	1730	734	2	210	THUMP (Thumpd1) alternative variant bSep08, mRNA.
Thumpd1	Thumpd1.cSep08	309041	1773	777		153	putative protein of eukaryotic origin (Thumpd1) alternative variant cSep08, mRNA.
Thumpd1	Thumpd1.dSep08	309041	3551	454	3	35	putative protein (Thumpd1) alternative variant dSep08, mRNA.
Thumpd2	Thumpd2.bSep08	313851	39685	1241	7	359	putative RNA methylase (Thumpd2) alternative variant bSep08, mRNA.
Thumpd2	Thumpd2.cSep08	313851	2068	807	1	200	thump domain-containing protein 2 (20.9 kD) (Thumpd2) alternative variant cSep08, mRNA.

Thumpd3	Thumpd3.aSep08	500288	24297	3290	10	525	THUMP and putative RNA methylase (Thumpd3) alternative variant aSep08, mRNA.
Thumpd3	Thumpd3.bSep08	500288	14503	1160	5	314	THUMP (35.3 kD) (Thumpd3) alternative variant bSep08, mRNA.
Thumpd3	Thumpd3.cSep08	500288	14280	914	5	288	putative protein of ancient origin (Thumpd3) alternative variant cSep08, mRNA.
Thumpd3	Thumpd3.dSep08	500288	2857	893	4	65	putative protein of metazoan origin (Thumpd3) alternative variant dSep08, mRNA.
Thy1	Thy1.aSep08	24832	5076	1985	4	222	thymus cell antigen 1, theta (24.6 kD) (Thy1) alternative variant aSep08, mRNA.
Thy1	Thy1.bSep08	24832	4108	736	5	217	thymus cell antigen 1, theta (Thy1) alternative variant bSep08, mRNA.
Thy1	Thy1.dSep08	24832	4251	784	4	146	thymus cell antigen 1, theta (16.5 kD) (Thy1) alternative variant dSep08, mRNA.
Thy1	Thy1.eSep08	24832	4287	739	3	146	thymus cell antigen 1, theta (16.5 kD) (Thy1) alternative variant eSep08, mRNA.
Thy1	Thy1.fSep08	24832	4149	654	4	146	thymus cell antigen 1, theta (16.5 kD) (Thy1) alternative variant fSep08, mRNA.
Thy1	Thy1.gSep08	24832	1452	790	2	85	thymus cell antigen 1, theta (Thy1) alternative variant gSep08, mRNA.
Thy1	Thy1.hSep08	24832	4435	625	3		
Thymosin.0	Thymosin.0.aSep08		2054	330		45	thymosin -like (5.4 kD) (Thymosin.0) mRNA.
Thyn1	Thyn1.bSep08	300470	1945	514	4	140	thymocyte nuclear protein 1 (Thyn1) alternative variant bSep08, mRNA.
Thyn1	Thyn1.cSep08	300470	3888	710	6	121	thymocyte nuclear protein 1 (14.2 kD) (Thyn1) alternative variant cSep08, mRNA.
Thyn1	Thyn1.eSep08	300470	992	347	2	68	thymocyte nuclear protein 1 (Thyn1) alternative variant eSep08, mRNA.
Tia1	Tia1.bSep08	312510	19372	3114	12	353	cytotoxic granule-associated RNA binding protein 1 (Tia1) alternative variant bSep08, mRNA.
Tia1	Tia1.dSep08	312510	16177	740	4	97	cytotoxic granule-associated RNA binding protein 1 (10.9 kD) (Tia1) alternative variant dSep08, mRNA.
Tia1	Tia1.eSep08	312510	10066	394	2	85	cytotoxic granule-associated RNA binding protein 1 (Tia1) alternative variant eSep08, mRNA.
Tia1	Tia1.fSep08	312510	880	444	2	84	cytotoxic granule-associated RNA binding protein 1 (9.7 kD) (Tia1) alternative variant fSep08, mRNA.
Tia1	Tia1.hSep08	312510	16256	737	6	54	cytotoxic granule-associated RNA binding protein 1 (6.1 kD) (Tia1) alternative variant hSep08, mRNA.
Tia1	Tia1.jSep08	312510	2190	570	5	66	cytotoxic granule-associated RNA binding protein 1 (Tia1) alternative variant jSep08, mRNA.
Tial1	Tial1.bSep08	361655	21278	5342	11	419	TIA-1 precursor (Tial1) alternative variant bSep08, mRNA.
Tial1	Tial1.cSep08	361655	2798	957	5	318	cytotoxic granule-associated RNA binding protein-like 1 (Tial1) alternative variant cSep08, mRNA.
Tial1	Tial1.dSep08	361655	6462	1789	8	189	cytotoxic granule-associated RNA binding protein-like 1 (21.1 kD) (Tial1) alternative variant dSep08, mRNA.
Tial1	Tial1.eSep08	361655	2750	733	5	133	cytotoxic granule-associated RNA binding protein-like 1 (Tial1) alternative variant eSep08, mRNA.

Tial1	Tial1.fSep08	361655	27027	870	5	127	cytotoxic granule-associated RNA binding protein-like 1 (Tial1) alternative variant fSep08, mRNA.
Tial1	Tial1.gSep08	361655	985	875	2	59	cytotoxic granule-associated RNA binding protein-like 1 (Tial1) alternative variant gSep08, mRNA.
Tiam1	Tiam1.aSep08	304109	34252	532		176	T-cell lymphoma invasion and metastasis 1 (Tiam1) mRNA.
Tie1	Tie1.bSep08	89806	3195	535	3	178	tyrosine kinase with immunoglobulin-like and EGF-like domains 1 (Tie1) alternative variant bSep08, mRNA.
Tie1	Tie1.cSep08	89806	729	389	3	129	tyrosine kinase with immunoglobulin-like and EGF-like domains 1 (Tie1) alternative variant cSep08, mRNA.
Tie1	Tie1.dSep08	89806	781	575	3	66	tyrosine kinase with immunoglobulin-like and EGF-like domains 1 (Tie1) alternative variant dSep08, mRNA.
Tifa	Tifa.aSep08	310877	8224	1729	1	181	TRAF-interacting protein with forkhead-associated domain (21.0 kD) (Tifa) alternative variant aSep08, mRNA.
Tifab	Tifab.cSep08	364674	1546	400	2	26	TRAF-interacting protein with forkhead-associated domain, family member B (Tifab) alternative variant cSep08, mRNA.
TIL.0	TIL.0.aSep08		9296	433		143	willebrand factor (TIL.0) mRNA.
TIL.1	TIL.1.aSep08		2193	581		193	protease inhibitor I8, cysteine-rich trypsin inhibitor-like (TIL.1) mRNA.
Timd2	Timd2.bSep08	287222	37147	1195	3	283	TIM2 (Timd2) alternative variant bSep08, mRNA.
Timeless	Timeless.bSep08	83508	17403	2491	13	552	timeless homolog (Drosophila) (64.2 kD) (Timeless) alternative variant bSep08, mRNA.
Timeless	Timeless.cSep08	83508	8074	2526	12	421	timeless homolog (Drosophila) (Timeless) alternative variant cSep08, mRNA.
Timeless	Timeless.dSep08	83508	2051	1056	5	182	timeless homolog (Drosophila) (Timeless) alternative variant dSep08, mRNA.
Timeless	Timeless.eSep08	83508	2066	1068	5	180	timeless homolog (Drosophila) (Timeless) alternative variant eSep08, mRNA.
Timm8a2	Timm8a2.aSep08	680794	5372	615	2	80	translocase of inner mitochondrial membrane 8 homolog a2 (yeast) (Timm8a2) alternative variant aSep08, mRNA.
Timm9	Timm9.aSep08	171139	12795	700	3	107	translocase of inner mitochondrial membrane 9 homolog CRA a (Timm9) alternative variant aSep08, mRNA.
Timm9	Timm9.bSep08	171139	12810	876	5	89	translocase inner membrane (10.4 kD) (Timm9) alternative variant bSep08, complete mRNA.
Timm9	Timm9.cSep08	171139	1774	160	2	51	putative protein (Timm9) alternative variant cSep08, mRNA.
Timm9	Timm9.eSep08	171139	2311	471	2	31	putative protein (Timm9) alternative variant eSep08, mRNA.
Timm10	Timm10.bSep08	64464	3497	940	2	90	translocase of inner mitochondrial membrane 10 homolog (yeast) (10.3 kD) (Timm10) alternative variant bSep08, mRNA.
Timm10	Timm10.cSep08	64464	3503	653	2	90	translocase of inner mitochondrial membrane 10 homolog (yeast) (10.3 kD) (Timm10) alternative variant cSep08, mRNA.
Timm17a	Timm17a.bSep08	54311	11913	1059	4	161	translocase of inner mitochondrial membrane 17a (16.9 kD) (Timm17a) alternative variant bSep08, complete mRNA.

Timm17a	Timm17a.cSep08	54311	5861	1708	1	115	translocase of inner mitochondrial membrane 17a (12.5 kD) (Timm17a) alternative variant cSep08, mRNA.
Timm22	Timm22.aSep08	79463	7743	1785	4	190	translocase of inner mitochondrial membrane 22 homolog (yeast) (Timm22) alternative variant aSep08, mRNA.
Timm22	Timm22.bSep08	79463	2782	477	2	158	translocase of inner mitochondrial membrane 22 homolog (yeast) (Timm22) alternative variant bSep08, mRNA.
Timm23	Timm23.bSep08	54312	10810	509	1	115	translocase of inner mitochondrial membrane 23 homolog (yeast) (11.9 kD) (Timm23) alternative variant bSep08, mRNA.
Timm50	Timm50.bSep08	685725	1786	307	3	102	translocase of inner mitochondrial membrane 50 homolog ( <i>S. cerevisiae</i> ) (Timm50) alternative variant bSep08, mRNA.
Timp1	Timp1.bSep08	116510	4680	818	2	217	tissue inhibitor of metalloproteinase 1 (23.8 kD) (Timp1) alternative variant bSep08, mRNA.
Timp1	Timp1.cSep08	116510	3371	324		98	tissue inhibitor of metalloproteinase 1 (Timp1) alternative variant cSep08, mRNA.
Timp2	Timp2.cSep08	29543	765	638	2	27	tissue inhibitor of metalloproteinase 2 (Timp2) alternative variant cSep08, mRNA.
Timp3	Timp3.bSep08	25358	5416	4007	1	143	tissue inhibitor of metalloproteinase 3 (Timp3) alternative variant bSep08, mRNA.
Tinagl1	Tinagl1.bSep08	94174	9998	1937	12	467	tubulointerstitial nephritis antigen-like 1 (52.8 kD) (Tinagl1) alternative variant bSep08, complete mRNA.
Tinagl1	Tinagl1.cSep08	94174	7477	775	6	233	tubulointerstitial nephritis antigen-like 1 (Tinagl1) alternative variant cSep08, mRNA.
Tinagl1	Tinagl1.dSep08	94174	6079	377	4	125	tubulointerstitial nephritis antigen-like 1 (Tinagl1) alternative variant dSep08, mRNA.
Titin_Z.0	Titin_Z.0.aSep08		2074	277		92	titin N2-B (Titin_Z.0) mRNA.
Tjap1	Tjap1.aSep08	316233	2657	2016	1	389	tight junction associated protein 1 (41.9 kD) (Tjap1) alternative variant aSep08, mRNA.
Tjap1	Tjap1.bSep08	316233	2845	1028	3	127	tight junction associated protein 1 (13.9 kD) (Tjap1) alternative variant bSep08, mRNA.
Tjp1	Tjp1.bSep08	292994	5629	852	2	210	tight junction protein 1 (Tjp1) alternative variant bSep08, mRNA.
Tjp1	Tjp1.cSep08	292994	5938	1122	2	195	tight junction protein 1 (Tjp1) alternative variant cSep08, mRNA.
Tjp2	Tjp2.bSep08	115769	94291	3087	20	938	tight junction protein 2 (Tjp2) alternative variant bSep08, mRNA.
Tjp2	Tjp2.cSep08	115769	44091	787	5	213	tight junction protein 2 (Tjp2) alternative variant cSep08, mRNA.
Tjp2	Tjp2.dSep08	115769	5100	576	4	192	tight junction protein 2 (Tjp2) alternative variant dSep08, mRNA.
Tjp2	Tjp2.eSep08	115769	3609	1391	3	165	tight junction protein 2 (Tjp2) alternative variant eSep08, mRNA.
Tjp2	Tjp2.fSep08	115769	14168	313	2	85	tight junction protein 2 (Tjp2) alternative variant fSep08, mRNA.
Tjp2	Tjp2.gSep08	115769	737	222	2	70	tight junction protein 2 (Tjp2) alternative variant gSep08, mRNA.



Tjp3	Tjp3.bSep08	314640	1615	967	3	171	tight junction protein 3 (Tjp3) alternative variant bSep08, mRNA.
Tjp3	Tjp3.cSep08	314640	4018	741	6	170	tight junction protein 3 (Tjp3) alternative variant cSep08, mRNA.
Tk2	Tk2.cSep08	291824	6569	402	4	33	thymidine kinase 2, mitochondrial (Tk2) alternative variant cSep08, mRNA.
Tkt	Tkt.bSep08	64524	17931	893	6	249	transketolase (Tkt) alternative variant bSep08, mRNA.
Tkt	Tkt.cSep08	64524	1477	739	3	181	transketolase (Tkt) alternative variant cSep08, mRNA.
Tkt	Tkt.dSep08	64524	13329	711	5	162	transketolase (Tkt) alternative variant dSep08, mRNA.
Tkt	Tkt.eSep08	64524	1881	921	3	109	transketolase (Tkt) alternative variant eSep08, mRNA.
Tkt	Tkt.fSep08	64524	2624	1145	3	99	transketolase (10.8 kD) (Tkt) alternative variant fSep08, mRNA.
Tktl1	Tktl1.bSep08	689374	18323	704	1	64	transketolase-like 1 (Tktl1) alternative variant bSep08, mRNA.
Tlcd2	Tlcd2.aSep08	497955	818	403		116	putative protein of metazoan origin (Tlcd2) mRNA.
TLD.0	TLD.0.aSep08		11221	2569		157	nuclear receptor coactivator 7 CRA c (18.0 kD) (TLD.0) mRNA.
Tle1	Tle1.aSep08	362533	16342	854	4	284	transducin-like enhancer of split 1, homolog of Drosophila E(spl) (Tle1) alternative variant aSep08, mRNA.
Tle1	Tle1.bSep08	362533	4912	1551	3	191	transducin-like enhancer of split 1, homolog of Drosophila E(spl) (21.3 kD) (Tle1) alternative variant bSep08, mRNA.
Tle2	Tle2.bSep08	299636	2246	1225	4	297	transducin-like enhancer of split 2, homolog of Drosophila E(spl) (32.3 kD) (Tle2) alternative variant bSep08, mRNA.
Tle2	Tle2.cSep08	299636	4572	437	6	145	transducin-like enhancer of split 2, homolog of Drosophila E(spl) (Tle2) alternative variant cSep08, mRNA.
Tle2	Tle2.dSep08	299636	766	404	3	134	transducin-like enhancer of split 2, homolog of Drosophila E(spl) (Tle2) alternative variant dSep08, mRNA.
Tle2	Tle2.hSep08	299636	525	413	2	80	transducin-like enhancer of split 2, homolog of Drosophila E(spl) (Tle2) alternative variant hSep08, mRNA.
Tle3	Tle3.aSep08	84424	44137	3621	20	610	transducin-like enhancer 3 (Tle3) alternative variant aSep08, mRNA.
Tle3	Tle3.cSep08	84424	4008	862	5	287	transducin-like enhancer 3 (Tle3) alternative variant cSep08, mRNA.
Tle3	Tle3.eSep08	84424	2483	271	3	42	transducin-like enhancer of split 3 (Tle3) alternative variant eSep08, mRNA.
Tle4	Tle4.aSep08	25565	20833	3304	7	571	transducin-like enhancer of split 4, homolog of Drosophila E(spl) (Tle4) alternative variant aSep08, mRNA.
Tle4	Tle4.bSep08	25565	1196	309	1	102	transducin-like enhancer of split 4, homolog of Drosophila E(spl) (Tle4) alternative variant bSep08, mRNA.
TLE_N.0	TLE_N.0.aSep08		44265	1307	1	236	transducin-like enhancer 1 (TLE_N.0) alternative variant aSep08, mRNA.
TLE_N.0	TLE_N.0.bSep08		44289	1361	1	163	transducin-like enhancer 1 (TLE_N.0) alternative variant bSep08, mRNA.
Tlk1	Tlk1.bSep08	311118	28351	2374	10	435	tousled-like kinase 1 (Tlk1) alternative variant bSep08, mRNA.

Tlk1	Tlk1.cSep08	311118	51039	411	3	137	tousled-like kinase 1 (Tlk1) alternative variant cSep08, mRNA.
Tlk2	Tlk2.aSep08	303592	91479	3424	22	837	tousled-like kinase 2 (Arabidopsis) and hypothetical protein LOC688398 (Tlk2) alternative variant aSep08, mRNA.
Tlk2	Tlk2.aSep08	688398	91479	3424	22	837	tousled-like kinase 2 (Arabidopsis) and hypothetical protein LOC688398 (Tlk2) alternative variant aSep08, mRNA.
Tlk2	Tlk2.bSep08	303592	22114	1145	6	258	tousled-like kinase 2 (Arabidopsis) and hypothetical protein LOC688398 (Tlk2) alternative variant bSep08, mRNA.
Tlk2	Tlk2.bSep08	688398	22114	1145	6	258	tousled-like kinase 2 (Arabidopsis) and hypothetical protein LOC688398 (Tlk2) alternative variant bSep08, mRNA.
Tln1	Tln1.aSep08	313494	9886	4563	24	1100	talin 1 (Tln1) alternative variant aSep08, mRNA.
Tln1	Tln1.bSep08	313494	532	450	2	150	talin 1 (Tln1) alternative variant bSep08, mRNA.
Tln1	Tln1.cSep08	313494	14184	654	5	136	talin 1 (Tln1) alternative variant cSep08, mRNA.
Tln1	Tln1.dSep08	313494	11265	482	3	69	talin 1 (Tln1) alternative variant dSep08, mRNA.
Tlr6	Tlr6.aSep08	305353	6525	2527	1	806	toll-like receptor 6 (92.6 kD) (Tlr6) alternative variant aSep08, mRNA.
Tlr7	Tlr7.aSep08	317468	26617	4060	2	1068	toll-like receptor 7 (123.4 kD) (Tlr7) alternative variant aSep08, mRNA.
Tlx3	Tlx3.aSep08	497881	1484	562	1	150	T-cell leukemia, homeobox 3 (Tlx3) alternative variant aSep08, mRNA.
Tlx3	Tlx3.bSep08	497881	1328	368	1	122	T-cell leukemia, homeobox 3 (Tlx3) alternative variant bSep08, mRNA.
Tm2d1	Tm2d1.aSep08	362545	40932	1290	5	208	hypothetical protein LOC685326 (22.2 kD) (Tm2d1) alternative variant aSep08, mRNA.
Tm2d1	Tm2d1.aSep08	685326	40932	1290	5	208	hypothetical protein LOC685326 (22.2 kD) (Tm2d1) alternative variant aSep08, mRNA.
Tm2d1	Tm2d1.bSep08	362545	40972	1000	4	141	hypothetical protein LOC685326 (14.9 kD) (Tm2d1) alternative variant bSep08, mRNA.
Tm2d1	Tm2d1.bSep08	685326	40972	1000	4	141	hypothetical protein LOC685326 (14.9 kD) (Tm2d1) alternative variant bSep08, mRNA.
Tm2d1	Tm2d1.cSep08	362545	20236	623	1	81	hypothetical protein LOC685326 (9.1 kD) (Tm2d1) alternative variant cSep08, mRNA.
Tm2d1	Tm2d1.cSep08	685326	20236	623	1	81	hypothetical protein LOC685326 (9.1 kD) (Tm2d1) alternative variant cSep08, mRNA.
Tm2d1	Tm2d1.dSep08	362545	11026	499	3	27	hypothetical protein LOC685326 (3.1 kD) (Tm2d1) alternative variant dSep08, mRNA.
Tm2d1	Tm2d1.dSep08	685326	11026	499	3	27	hypothetical protein LOC685326 (3.1 kD) (Tm2d1) alternative variant dSep08, mRNA.
Tm2d2	Tm2d2.bSep08	290833	2581	498	3	143	domain-containing protein 2 precursor (15.2 kD) (Tm2d2) alternative variant bSep08, mRNA.
Tm2d3	Tm2d3.bSep08	292995	11936	828	2	227	TM2 (Tm2d3) alternative variant bSep08, mRNA.
Tm2d3	Tm2d3.cSep08	292995	10673	1141	1	183	TM2 (20.3 kD) (Tm2d3) alternative variant cSep08, mRNA.
Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.cSep08	288044	14463	639	5	143	tctex1 domain-containing protein 2 (Tm4sf19andTctex1d2) alternative variant cSep08, mRNA.
Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.cSep08	498095	14463	639	5	143	tctex1 domain-containing protein 2 (Tm4sf19andTctex1d2) alternative variant cSep08, mRNA.

Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.dSep08	288044	3549	314	2	95	putative protein of metazoan origin (Tm4sf19andTctex1d2) alternative variant dSep08, mRNA.
Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.dSep08	498095	3549	314	2	95	putative protein of metazoan origin (Tm4sf19andTctex1d2) alternative variant dSep08, mRNA.
Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.eSep08	288044	1691	905	2	36	tctex1 domain-containing protein 2 like (4.2 kD) (Tm4sf19andTctex1d2) alternative variant eSep08, mRNA.
Tm4sf19andTctex1d2	Tm4sf19andTctex1d2.eSep08	498095	1691	905	2	36	tctex1 domain-containing protein 2 like (4.2 kD) (Tm4sf19andTctex1d2) alternative variant eSep08, mRNA.
Tm6sf1	Tm6sf1.bSep08	361600	18639	1204	8	266	transmembrane 6 superfamily member 1 (Tm6sf1) alternative variant bSep08, mRNA.
Tm6sf2	Tm6sf2.bSep08	689029	1414	752	1	149	transmembrane 6 superfamily member 2 (15.5 kD) (Tm6sf2) alternative variant bSep08, mRNA.
Tm7sf2	Tm7sf2.bSep08	293688	3267	660	6	219	transmembrane 7 superfamily member 2 (Tm7sf2) alternative variant bSep08, mRNA.
Tm9sf1	Tm9sf1.bSep08	361043	7309	1886	6	628	transmembrane 9 superfamily member 1 (Tm9sf1) alternative variant bSep08, mRNA.
Tm9sf1	Tm9sf1.cSep08	361043	4976	1186	3	280	transmembrane 9 superfamily member 1 (31.7 kD) (Tm9sf1) alternative variant cSep08, mRNA.
Tm9sf1	Tm9sf1.dSep08	361043	2132	955	3	264	transmembrane 9 superfamily member 1 (Tm9sf1) alternative variant dSep08, mRNA.
Tm9sf3	Tm9sf3.aSep08	309475	53077	6396	15	587	transmembrane 9 superfamily member 3 (67.5 kD) (Tm9sf3) alternative variant aSep08, mRNA.
Tm9sf3	Tm9sf3.bSep08	309475	21839	786	8	247	transmembrane 9 superfamily member 3 (Tm9sf3) alternative variant bSep08, mRNA.
Tm9sf4	Tm9sf4.bSep08	296279	34076	691	5	160	transmembrane 9 superfamily protein member 4 (Tm9sf4) alternative variant bSep08, mRNA.
Tmbim1	Tmbim1.bSep08	316516	1722	644	3	111	transmembrane BAX inhibitor motif containing 1 (Tmbim1) alternative variant bSep08, mRNA.
Tmbim1	Tmbim1.cSep08	316516	5600	713	8	109	transmembrane BAX inhibitor motif containing 1 (Tmbim1) alternative variant cSep08, mRNA.
Tmbim1	Tmbim1.dSep08	316516	2556	839	4	78	transmembrane BAX inhibitor motif containing 1 (8.4 kD) (Tmbim1) alternative variant dSep08, mRNA.
Tmbim4	Tmbim4.bSep08	362884	2931	1313	3	139	transmembrane BAX inhibitor motif containing 4 (15.8 kD) (Tmbim4) alternative variant bSep08, mRNA.
Tmbim4	Tmbim4.cSep08	362884	5970	575	5	133	transmembrane BAX inhibitor motif containing 4 (14.8 kD) (Tmbim4) alternative variant cSep08, mRNA.
Tmc3	Tmc3.aSep08	293065	12431	703		195	transmembrane channel-like gene family 3 (Tmc3) mRNA.
Tmc5	Tmc5.bSep08	365360	8455	398	2	74	transmembrane channel-like gene family 5 (Tmc5) alternative variant bSep08, mRNA.
Tmcc1	Tmcc1.aSep08	312654	28791	1410		220	transmembrane and coiled coil domains 1 (Tmcc1) mRNA.
Tmcc2	Tmcc2.aSep08	305095	13276	2752	4	470	transmembrane and coiled-coil domains 2 (51.9 kD) (Tmcc2) alternative variant aSep08, mRNA.
Tmcc3	Tmcc3.bSep08	314751	25198	749	2	126	transmembrane and coiled coil domains 3 (Tmcc3) alternative variant bSep08, mRNA.
Tmcc3	Tmcc3.cSep08	314751	26622	395	1	75	transmembrane and coiled coil domains 3 (Tmcc3) alternative variant cSep08, mRNA.

Tmco3	Tmco3.aSep08	306607	37860	3103	15	678	transmembrane and coiled-coil domains 3 (75.7 kD) (Tmco3) alternative variant aSep08, mRNA.
Tmco3	Tmco3.bSep08	306607	22570	723	6	241	transmembrane and coiled-coil domains 3 (Tmco3) alternative variant bSep08, mRNA.
Tmco3	Tmco3.eSep08	306607	4255	671	4	81	transmembrane and coiled-coil domains 3 (Tmco3) alternative variant eSep08, mRNA.
Tmco4	Tmco4.aSep08	500573	39214	916		142	transmembrane and coiled-coil domains 4 and hypothetical protein LOC690449 (Tmco4) mRNA.
Tmco4	Tmco4.aSep08	690449	39214	916		142	transmembrane and coiled-coil domains 4 and hypothetical protein LOC690449 (Tmco4) mRNA.
Tmed1	Tmed1.bSep08	315461	2727	1620	2	82	putative protein, with a coiled coil domain, of bilateral origin (9.6 kD) (Tmed1) alternative variant bSep08, complete mRNA.
Tmed2	Tmed2.aSep08	65165	7808	749	5	208	transmembrane emp24 domain trafficking protein 2 (23.4 kD) (Tmed2) alternative variant aSep08, mRNA.
Tmed4	Tmed4.aSep08	305502	4181	1607	5	227	emp24/gp25L/p24 (26.1 kD) (Tmed4) alternative variant aSep08, complete mRNA.
Tmed4	Tmed4.bSep08	305502	4560	2099	5	223	emp24/gp25L/p24 (24.9 kD) (Tmed4) alternative variant bSep08, complete mRNA.
Tmed10	Tmed10.bSep08	84599	12091	3154	3	107	transmembrane emp24-like trafficking protein 10 (yeast) (Tmed10) alternative variant bSep08, mRNA.
Tmeff1	Tmeff1.bSep08	63845	23085	613	5	150	transmembrane protein with EGF-like and two follistatin-like domains 1 (Tmeff1) alternative variant bSep08, mRNA.
Tmeff2	Tmeff2.bSep08	363228	55947	816	2	195	transmembrane protein with EGF-like and two follistatin-like domains 2 (Tmeff2) alternative variant bSep08, mRNA.
Tmem1	Tmem1.aSep08	309678	16677	4505	2	604	transmembrane protein 1 (Tmem1) alternative variant aSep08, mRNA.
Tmem1	Tmem1.bSep08	309678	4665	784	1	163	transmembrane protein 1 (Tmem1) alternative variant bSep08, mRNA.
Tmem2	Tmem2.bSep08	309400	15018	738	8	238	transmembrane protein 2 (Tmem2) alternative variant bSep08, mRNA.
Tmem2	Tmem2.cSep08	309400	12777	619	6	206	transmembrane protein 2 (Tmem2) alternative variant cSep08, mRNA.
Tmem2	Tmem2.dSep08	309400	9826	1104	3	99	transmembrane protein 2 (11.0 kD) (Tmem2) alternative variant dSep08, mRNA.
Tmem5	Tmem5.bSep08	299841	11810	817	4	188	transmembrane protein 5 (Tmem5) alternative variant bSep08, mRNA.
Tmem5	Tmem5.cSep08	299841	9642	774	4	165	transmembrane protein 5 (Tmem5) alternative variant cSep08, mRNA.
Tmem8	Tmem8.cSep08	303004	3011	729	1	96	transmembrane protein 8 (five membrane-spanning domains) (Tmem8) alternative variant cSep08, mRNA.
Tmem9	Tmem9.bSep08	289046	17293	1583	6	183	transmembrane protein 9 (20.6 kD) (Tmem9) alternative variant bSep08, mRNA.
Tmem9	Tmem9.cSep08	289046	17357	1463	6	183	transmembrane protein 9 (20.6 kD) (Tmem9) alternative variant cSep08, complete mRNA.
Tmem9	Tmem9.dSep08	289046	7581	772	2	52	transmembrane protein 9 (Tmem9) alternative variant dSep08, mRNA.

Tmem9b	Tmem9b.aSep08	293415	20018	3037	2	199	TMEM9 domain family, member B (22.6 kD) (Tmem9b) alternative variant aSep08, mRNA.
Tmem11	Tmem11.bSep08	303196	14049	722	2	133	transmembrane protein 11 (Tmem11) alternative variant bSep08, mRNA.
Tmem14a	Tmem14a.bSep08	363206	11947	469		44	transmembrane protein 14A (Tmem14a) alternative variant bSep08, mRNA.
Tmem14c	Tmem14c.aSep08	171432	6027	776	4	114	transmembrane protein 14C (11.7 kD) (Tmem14c) alternative variant aSep08, mRNA.
Tmem14c	Tmem14c.cSep08	171432	2048	433	2	63	transmembrane protein 14C (Tmem14c) alternative variant cSep08, mRNA.
Tmem16a	Tmem16a.bSep08	309135	35854	1134	13	286	transmembrane protein 16A (Tmem16a) alternative variant bSep08, mRNA.
Tmem16a	Tmem16a.cSep08	309135	9386	398	2	132	transmembrane protein 16A (Tmem16a) alternative variant cSep08, mRNA.
Tmem16c	Tmem16c.aSep08	311287	52174	560		186	transmembrane protein 16C (Tmem16c) mRNA.
Tmem16d	Tmem16d.bSep08	299714	78728	647	8	215	transmembrane protein 16D (eight membrane-spanning domains) (Tmem16d) alternative variant bSep08, mRNA.
Tmem16d	Tmem16d.cSep08	299714	9849	568	5	189	transmembrane protein 16D (eight membrane-spanning domains) (Tmem16d) alternative variant cSep08, mRNA.
Tmem16d	Tmem16d.dSep08	299714	160000	574	3	117	transmembrane protein 16D (eight membrane-spanning domains) (Tmem16d) alternative variant dSep08, mRNA.
Tmem16f	Tmem16f.bSep08	315272	17207	1026	6	341	transmembrane protein 16F (Tmem16f) alternative variant bSep08, mRNA.
Tmem16f	Tmem16f.cSep08	315272	33976	951	7	317	transmembrane protein 16F (Tmem16f) alternative variant cSep08, mRNA.
Tmem16f	Tmem16f.dSep08	315272	10087	375	3	124	transmembrane protein 16F (Tmem16f) alternative variant dSep08, mRNA.
Tmem16f	Tmem16f.eSep08	315272	8729	3052	2	109	transmembrane protein 16F (Tmem16f) alternative variant eSep08, mRNA.
Tmem16k	Tmem16k.aSep08	301111	117339	2650	2	723	transmembrane protein 16K (Tmem16k) alternative variant aSep08, mRNA.
Tmem16k	Tmem16k.bSep08	301111	17713	742		247	transmembrane protein 16K (Tmem16k) alternative variant bSep08, mRNA.
Tmem17	Tmem17.cSep08	360985	35266	748	2	94	transmembrane protein 17 (9.9 kD) (Tmem17) alternative variant cSep08, mRNA.
Tmem17	Tmem17.dSep08	360985	7657	272	2	57	transmembrane protein 17 (Tmem17) alternative variant dSep08, mRNA.
Tmem18	Tmem18.aSep08	362722	6744	3570	4	148	transmembrane protein 18 (17.3 kD) (Tmem18) alternative variant aSep08, mRNA.
Tmem18	Tmem18.dSep08	362722	1078	333	2	42	transmembrane protein 18 (Tmem18) alternative variant dSep08, mRNA.
Tmem19	Tmem19.bSep08	299800	20098	1007	2	270	transmembrane protein 19 (Tmem19) alternative variant bSep08, mRNA.
Tmem24	Tmem24.bSep08	300666	4851	1602	6	273	transmembrane protein 24 (Tmem24) alternative variant bSep08, mRNA.
Tmem24	Tmem24.cSep08	300666	2821	856	2	166	transmembrane protein 24 (Tmem24) alternative variant cSep08, mRNA.

Tmem24	Tmem24.dSep08	300666	925	839	2	125	transmembrane protein 24 (Tmem24) alternative variant dSep08, mRNA.
Tmem25	Tmem25.bSep08	689172	2066	1839	1	70	transmembrane protein 25 (7.8 kD) (Tmem25) alternative variant bSep08, mRNA.
Tmem28l	Tmem28l.aSep08	688841	1069	681		186	transmembrane protein 28-like (Tmem28l) mRNA.
Tmem30a	Tmem30a.aSep08	300857	24489	4346	2	364	transmembrane protein 30A (41.1 kD) (Tmem30a) alternative variant aSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.dSep08	59303	12275	527	6	175	solute carrier family 30 member 9 CRA a (Tmem33andSlc30a9) alternative variant dSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.dSep08	498358	12275	527	6	175	solute carrier family 30 member 9 CRA a (Tmem33andSlc30a9) alternative variant dSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.eSep08	59303	15193	462	7	146	solute carrier family 30 member 9 CRA a (Tmem33andSlc30a9) alternative variant eSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.eSep08	498358	15193	462	7	146	solute carrier family 30 member 9 CRA a (Tmem33andSlc30a9) alternative variant eSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.fSep08	59303	20359	542	2	50	transmembrane protein 33 (Tmem33andSlc30a9) alternative variant fSep08, mRNA.
Tmem33andSlc30a9	Tmem33andSlc30a9.fSep08	498358	20359	542	2	50	transmembrane protein 33 (Tmem33andSlc30a9) alternative variant fSep08, mRNA.
Tmem34	Tmem34.bSep08	291946	7330	902	7	235	putative protein, with a transmembrane domain, of eukaryotic origin (Tmem34) alternative variant bSep08, mRNA.
Tmem34	Tmem34.cSep08	291946	983	537	2	91	putative protein of metazoan origin (Tmem34) alternative variant cSep08, mRNA.
Tmem38a	Tmem38a.bSep08	306327	16054	2715	4	198	transmembrane protein 38a (21.4 kD) (Tmem38a) alternative variant bSep08, mRNA.
Tmem38a	Tmem38a.cSep08	306327	8216	522	2	58	transmembrane protein 38a (Tmem38a) alternative variant cSep08, mRNA.
Tmem39a	Tmem39a.bSep08	288092	4529	732	2	214	transmembrane protein 39a (Tmem39a) alternative variant bSep08, mRNA.
Tmem39a	Tmem39a.cSep08	288092	18550	695	4	186	transmembrane protein 39a (Tmem39a) alternative variant cSep08, mRNA.
Tmem39a	Tmem39a.dSep08	288092	4440	578	3	106	transmembrane protein 39a (12.3 kD) (Tmem39a) alternative variant dSep08, mRNA.
Tmem39b	Tmem39b.bSep08	362608	2119	580	2	120	transmembrane protein 39b (Tmem39b) alternative variant bSep08, mRNA.
Tmem40	Tmem40.aSep08	680858	10676	340	6	112	transmembrane protein 40 (Tmem40) alternative variant aSep08, mRNA.
Tmem41b	Tmem41b.bSep08	361626	14338	2306	1	206	transmembrane protein 41B (23.3 kD) (Tmem41b) alternative variant bSep08, mRNA.
Tmem42	Tmem42.aSep08	363171	3511	2319	3	125	transmembrane protein 42 (13.6 kD) (Tmem42) alternative variant aSep08, complete mRNA.
Tmem42	Tmem42.cSep08	363171	1560	936	2	60	transmembrane protein 42 (Tmem42) alternative variant cSep08, mRNA.
Tmem43	Tmem43.bSep08	362401	2028	469	1	61	transmembrane protein 43 (Tmem43) alternative variant bSep08, mRNA.
Tmem44	Tmem44.aSep08	288028	15162	933		311	transmembrane protein 44 (Tmem44) mRNA.

Tmem45a	Tmem45a.aSep08	680866	85840	1581		273	transmembrane protein 45A (31.2 kD) (Tmem45a) mRNA.
Tmem48	Tmem48.bSep08	362557	16809	366		122	transmembrane protein 48 (Tmem48) alternative variant bSep08, mRNA.
Tmem49	Tmem49.bSep08	192129	27590	639	6	182	transmembrane protein 49 (Tmem49) alternative variant bSep08, mRNA.
Tmem49	Tmem49.cSep08	192129	37927	1616	6	173	transmembrane protein 49 (Tmem49) alternative variant cSep08, mRNA.
Tmem49	Tmem49.dSep08	192129	17784	1144	3	87	transmembrane protein 49 (9.9 kD) (Tmem49) alternative variant dSep08, mRNA.
Tmem49	Tmem49.eSep08	192129	1931	898	2	61	transmembrane protein 49 (Tmem49) alternative variant eSep08, mRNA.
Tmem50b	Tmem50b.aSep08	360698	31240	695	5	158	transmembrane protein 50B CRA a (17.9 kD) (Tmem50b) alternative variant aSep08, mRNA.
Tmem50b	Tmem50b.cSep08	360698	28428	593	4	139	transmembrane protein 50B CRA a (Tmem50b) alternative variant cSep08, mRNA.
Tmem50b	Tmem50b.dSep08	360698	21769	400	2	74	putative protein (Tmem50b) alternative variant dSep08, mRNA.
Tmem50b	Tmem50b.eSep08	360698	22000	375	2	69	putative protein (Tmem50b) alternative variant eSep08, mRNA.
Tmem50b	Tmem50b.fSep08	360698	22017	356	2	45	transmembrane protein 50B CRA a (Tmem50b) alternative variant fSep08, mRNA.
Tmem53	Tmem53.aSep08	313529	15410	872		275	transmembrane protein 53 (Tmem53) mRNA.
Tmem55a	Tmem55a.bSep08	362490	25375	783	1	184	transmembrane protein 55A (Tmem55a) alternative variant bSep08, mRNA.
Tmem55b	Tmem55b.bSep08	364298	1964	742	5	230	transmembrane protein 55B (Tmem55b) alternative variant bSep08, mRNA.
Tmem55b	Tmem55b.cSep08	364298	1667	691	4	150	transmembrane protein 55B (Tmem55b) alternative variant cSep08, mRNA.
Tmem55b	Tmem55b.dSep08	364298	2015	908	2	115	transmembrane protein 55B (13.0 kD) (Tmem55b) alternative variant dSep08, mRNA.
Tmem55b	Tmem55b.eSep08	364298	2246	891	3	115	transmembrane protein 55B (13.0 kD) (Tmem55b) alternative variant eSep08, mRNA.
Tmem57	Tmem57.aSep08	313618	57039	1938		645	transmembrane protein 57 (Tmem57) mRNA.
Tmem59l	Tmem59l.aSep08	306349	3666	1412	8	330	transmembrane protein 59-like (Tmem59l) alternative variant aSep08, mRNA.
Tmem59l	Tmem59l.bSep08	306349	2994	873	7	290	transmembrane protein 59-like (Tmem59l) alternative variant bSep08, mRNA.
Tmem59l	Tmem59l.cSep08	306349	2539	555	5	129	transmembrane protein 59-like (Tmem59l) alternative variant cSep08, mRNA.
Tmem59l	Tmem59l.dSep08	306349	674	599	2	110	transmembrane protein 59-like (Tmem59l) alternative variant dSep08, mRNA.
Tmem60	Tmem60.aSep08	296761	4851	1365	2	76	transmembrane protein 60 (8.1 kD) (Tmem60) alternative variant aSep08, mRNA.
Tmem62	Tmem62.aSep08	311350	22169	1966		425	transmembrane protein 62 (Tmem62) mRNA.
Tmem63a	Tmem63a.bSep08	289318	13209	742	1	138	transmembrane protein 63a (Tmem63a) alternative variant bSep08, mRNA.

Tmem63b	Tmem63b.aSep08	363197	12086	1768	15	584	transmembrane protein 63b (Tmem63b) alternative variant aSep08, mRNA.
Tmem63b	Tmem63b.bSep08	363197	2794	1145	7	249	transmembrane protein 63b (29.1 kD) (Tmem63b) alternative variant bSep08, mRNA.
Tmem67	Tmem67.bSep08	313067	11744	490	4	137	transmembrane protein 67 (Tmem67) alternative variant bSep08, mRNA.
Tmem67	Tmem67.cSep08	313067	4275	792	4	120	transmembrane protein 67 CRA b (Tmem67) alternative variant cSep08, mRNA.
Tmem67	Tmem67.dSep08	313067	1494	730	2	78	transmembrane protein 67 CRA b (9.1 kD) (Tmem67) alternative variant dSep08, mRNA.
Tmem68	Tmem68.bSep08	312946	17619	683	4	227	transmembrane protein 68 (Tmem68) alternative variant bSep08, mRNA.
Tmem68	Tmem68.cSep08	312946	8994	867	5	97	transmembrane protein 68 (11.3 kD) (Tmem68) alternative variant cSep08, mRNA.
Tmem71	Tmem71.aSep08	690657	20632	406		72	transmembrane protein 71 (Tmem71) alternative variant aSep08, mRNA.
Tmem77	Tmem77.bSep08	362011	26793	930	9	227	transmembrane protein 77 (Tmem77) alternative variant bSep08, mRNA.
Tmem77	Tmem77.cSep08	362011	25353	855	7	200	transmembrane protein 77 (Tmem77) alternative variant cSep08, mRNA.
Tmem77	Tmem77.dSep08	362011	25469	774	7	195	transmembrane protein 77 (Tmem77) alternative variant dSep08, mRNA.
Tmem77	Tmem77.eSep08	362011	9156	481	4	133	transmembrane protein 77 (Tmem77) alternative variant eSep08, mRNA.
Tmem77	Tmem77.fSep08	362011	17888	599	5	130	transmembrane protein 77 (14.6 kD) (Tmem77) alternative variant fSep08, mRNA.
Tmem77	Tmem77.gSep08	362011	28710	1723	5	53	transmembrane protein 77 (6.2 kD) (Tmem77) alternative variant gSep08, complete mRNA.
Tmem80	Tmem80.aSep08	309109	7519	796	4	172	transmembrane protein 80 (Tmem80) alternative variant aSep08, mRNA.
Tmem80	Tmem80.cSep08	309109	4928	829	1	77	transmembrane protein 80 (Tmem80) alternative variant cSep08, mRNA.
Tmem82	Tmem82.aSep08	298605	3934	1798	5	548	transmembrane protein 82 (Tmem82) alternative variant aSep08, mRNA.
Tmem82	Tmem82.bSep08	298605	2036	878	3	249	transmembrane protein 82 (Tmem82) alternative variant bSep08, mRNA.
Tmem82	Tmem82.dSep08	298605	430	322	2	46	transmembrane protein 82 (Tmem82) alternative variant dSep08, mRNA.
Tmem85	Tmem85.bSep08	296049	3752	487	4	138	transmembrane protein 85 (Tmem85) alternative variant bSep08, mRNA.
Tmem85	Tmem85.cSep08	296049	920	618	2	89	transmembrane protein 85 (9.4 kD) (Tmem85) alternative variant cSep08, mRNA.
Tmem85	Tmem85.dSep08	296049	4510	1789	3	89	transmembrane protein 85 (9.4 kD) (Tmem85) alternative variant dSep08, mRNA.
Tmem86a	Tmem86a.aSep08	308602	3062	442	1	147	transmembrane protein 86A (Tmem86a) alternative variant aSep08, mRNA.



Tmem86a	Tmem86a.bSep08	308602	12771	544	1	109	transmembrane protein 86A (Tmem86a) alternative variant bSep08, mRNA.
Tmem86b	Tmem86b.aSep08	690610	2926	2100	1	233	transmembrane protein 86B (25.9 kD) (Tmem86b) alternative variant aSep08, mRNA.
Tmem87a	Tmem87a.aSep08	366170	38839	1869	16	498	transmembrane protein 87A (Tmem87a) alternative variant aSep08, mRNA.
Tmem87a	Tmem87a.bSep08	366170	19715	1116	9	251	transmembrane protein 87A (Tmem87a) alternative variant bSep08, mRNA.
Tmem87a	Tmem87a.cSep08	366170	18273	543	7	139	transmembrane protein 87A (Tmem87a) alternative variant cSep08, mRNA.
Tmem87a	Tmem87a.dSep08	366170	3412	1264	2	12	transmembrane protein 87A (Tmem87a) alternative variant dSep08, mRNA.
Tmem87b	Tmem87b.aSep08	362212	12724	3328		130	transmembrane protein 87B (Tmem87b) mRNA.
Tmem93	Tmem93.bSep08	287477	845	650	2	154	transmembrane protein 93 (Tmem93) alternative variant bSep08, mRNA.
Tmem97	Tmem97.bSep08	303330	1497	485	1	161	transmembrane protein 97 (Tmem97) alternative variant bSep08, mRNA.
Tmem98	Tmem98.cSep08	303356	2377	1076	2	81	transmembrane protein 98 (8.8 kD) (Tmem98) alternative variant cSep08, mRNA.
Tmem100	Tmem100.aSep08	497979	4327	2756	2	134	transmembrane protein 100 (14.4 kD) (Tmem100) alternative variant aSep08, mRNA.
Tmem100	Tmem100.dSep08	497979	2721	739	3	134	transmembrane protein 100 (14.4 kD) (Tmem100) alternative variant dSep08, mRNA.
Tmem101	Tmem101.aSep08	303564	3753	1694		257	transmembrane protein 101 (28.8 kD) (Tmem101) complete mRNA.
Tmem102	Tmem102.aSep08	497937	1537	1436		469	transmembrane protein 102 (Tmem102) mRNA.
Tmem104	Tmem104.aSep08	303670	16026	617		151	transmembrane protein 104 (Tmem104) mRNA.
Tmem106a	Tmem106a.bSep08	287722	5997	826	7	203	transmembrane protein 106A (Tmem106a) alternative variant bSep08, mRNA.
Tmem106a	Tmem106a.cSep08	287722	609	558	2	65	transmembrane protein 106A (7.2 kD) (Tmem106a) alternative variant cSep08, mRNA.
Tmem106b	Tmem106b.bSep08	312132	2238	562	3	63	transmembrane protein 106B (Tmem106b) alternative variant bSep08, mRNA.
Tmem106c	Tmem106c.aSep08	315286	5279	1584	4	260	transmembrane protein 106C (28.9 kD) (Tmem106c) alternative variant aSep08, mRNA.
Tmem106c	Tmem106c.cSep08	315286	2918	709	2	161	transmembrane protein 106C (Tmem106c) alternative variant cSep08, mRNA.
Tmem108	Tmem108.aSep08	300967	23852	1392	3	324	transmembrane protein 108 (Tmem108) alternative variant aSep08, mRNA.
Tmem108	Tmem108.bSep08	300967	247258	626	5	52	transmembrane protein 108 (Tmem108) alternative variant bSep08, mRNA.
Tmem108	Tmem108.cSep08	300967	247323	396	3	70	transmembrane protein 108 (Tmem108) alternative variant cSep08, mRNA.
Tmem109	Tmem109.aSep08	361732	9536	947	4	243	transmembrane protein 109 (26.2 kD) (Tmem109) alternative variant aSep08, mRNA.
Tmem109	Tmem109.cSep08	361732	5774	437	1	64	transmembrane protein 109 (Tmem109) alternative variant cSep08, mRNA.

Tmem109	Tmem109.dSep08	361732	7009	354	2	51	transmembrane protein 109 (Tmem109) alternative variant dSep08, mRNA.
Tmem110	Tmem110.bSep08	361110	9515	725	2	175	transmembrane protein 110 (Tmem110) alternative variant bSep08, mRNA.
Tmem116	Tmem116.aSep08	690442	33313	1330	10	352	transmembrane protein 116 CRA a (Tmem116) alternative variant aSep08, mRNA.
Tmem116	Tmem116.aSep08	690455	33313	1330	10	352	transmembrane protein 116 CRA a (Tmem116) alternative variant aSep08, mRNA.
Tmem116	Tmem116.bSep08	690442	42571	750	7	212	transmembrane protein 116 CRA c (Tmem116) alternative variant bSep08, mRNA.
Tmem116	Tmem116.bSep08	690455	42571	750	7	212	transmembrane protein 116 CRA c (Tmem116) alternative variant bSep08, mRNA.
Tmem116	Tmem116.cSep08	690442	1865	1137	2	87	transmembrane protein 116 CRA a precursor (9.5 kD) (Tmem116) alternative variant cSep08, mRNA.
Tmem116	Tmem116.cSep08	690455	1865	1137	2	87	transmembrane protein 116 CRA a precursor (9.5 kD) (Tmem116) alternative variant cSep08, mRNA.
Tmem116	Tmem116.dSep08	690442	13012	350	3	55	transmembrane protein 116 CRA b (Tmem116) alternative variant dSep08, mRNA.
Tmem116	Tmem116.dSep08	690455	13012	350	3	55	transmembrane protein 116 CRA b (Tmem116) alternative variant dSep08, mRNA.
Tmem120a	Tmem120a.bSep08	288591	6821	614	4	186	transmembrane protein 120A (Tmem120a) alternative variant bSep08, mRNA.
Tmem121	Tmem121.aSep08	691678	2435	414		138	transmembrane protein 121 (Tmem121) mRNA.
Tmem123	Tmem123.aSep08	363013	26866	876	1	226	transmembrane protein 123 (Tmem123) alternative variant aSep08, mRNA.
Tmem123	Tmem123.bSep08	363013	26827	831	1	200	transmembrane protein 123 (Tmem123) alternative variant bSep08, mRNA.
Tmem123	Tmem123.cSep08	363013	3662	2794	1	141	transmembrane protein 123 (14.7 kD) (Tmem123) alternative variant cSep08, mRNA.
Tmem126a	Tmem126a.aSep08	293113	6160	854	5	196	transmembrane protein 126A (21.7 kD) (Tmem126a) alternative variant aSep08, complete mRNA.
Tmem126a	Tmem126a.cSep08	293113	4729	739	3	122	transmembrane protein 126A (13.7 kD) (Tmem126a) alternative variant cSep08, mRNA.
Tmem126a	Tmem126a.dSep08	293113	1014	466	2	100	transmembrane protein 126A (11.1 kD) (Tmem126a) alternative variant dSep08, mRNA.
Tmem126b	Tmem126b.bSep08	293114	11941	916	4	143	transmembrane protein 126B (15.9 kD) (Tmem126b) alternative variant bSep08, complete mRNA.
Tmem126b	Tmem126b.cSep08	293114	11943	690	4	80	transmembrane protein 126B (8.8 kD) (Tmem126b) alternative variant cSep08, complete mRNA.
Tmem126b	Tmem126b.dSep08	293114	7367	1327	1	53	transmembrane protein 126B (6.1 kD) (Tmem126b) alternative variant dSep08, mRNA.
Tmem128	Tmem128.bSep08	360952	7247	930	4	125	transmembrane protein 128 (14.6 kD) (Tmem128) alternative variant bSep08, mRNA.
Tmem128	Tmem128.cSep08	360952	5153	1253	3	107	transmembrane protein 128 (12.1 kD) (Tmem128) alternative variant cSep08, mRNA.
Tmem128	Tmem128.dSep08	360952	2743	393	2	65	transmembrane protein 128 (Tmem128) alternative variant dSep08, mRNA.

Tmem130	Tmem130.aSep08	304280	23889	2758	8	461	transmembrane protein 130 (Tmem130) alternative variant aSep08, mRNA.
Tmem131	Tmem131.aSep08	316335	104947	963		321	transmembrane protein 131 (Tmem131) mRNA.
Tmem132a	Tmem132a.bSep08	338474	11657	1834	4	488	transmembrane protein 132A (Tmem132a) alternative variant bSep08, complete mRNA.
Tmem132a	Tmem132a.cSep08	338474	3239	1037	5	345	transmembrane protein 132A (Tmem132a) alternative variant cSep08, mRNA.
Tmem132a	Tmem132a.dSep08	338474	2298	859	3	221	transmembrane protein 132A (24.1 kD) (Tmem132a) alternative variant dSep08, mRNA.
Tmem132d	Tmem132d.cSep08	288750	2457	398	3	62	transmembrane protein 132D (Tmem132d) alternative variant cSep08, mRNA.
Tmem134	Tmem134.cSep08	361695	1754	889	2	125	putative protein (13.5 kD) (Tmem134) alternative variant cSep08, mRNA.
Tmem134	Tmem134.dSep08	361695	4388	695	6	111	transmembrane protein 134 (12.4 kD) (Tmem134) alternative variant dSep08, mRNA.
Tmem138	Tmem138.aSep08	361728	6547	886	5	240	transmembrane protein 138 (Tmem138) alternative variant aSep08, mRNA.
Tmem138	Tmem138.cSep08	361728	6285	662	5	80	transmembrane protein 138 like (Tmem138) alternative variant cSep08, mRNA.
Tmem138	Tmem138.dSep08	361728	2392	412	3	69	transmembrane protein 138 (Tmem138) alternative variant dSep08, mRNA.
Tmem138	Tmem138.eSep08	361728	5744	748	3	65	transmembrane protein 138 (Tmem138) alternative variant eSep08, mRNA.
Tmem138	Tmem138.fSep08	361728	6318	762	5	63	transmembrane protein 138 (7.8 kD) (Tmem138) alternative variant fSep08, mRNA.
Tmem138	Tmem138.gSep08	361728	6092	647	5	58	transmembrane protein 138 (Tmem138) alternative variant gSep08, mRNA.
Tmem141	Tmem141.aSep08	499755	2525	1636	4	128	transmembrane protein 141 (14.9 kD) (Tmem141) alternative variant aSep08, mRNA.
Tmem141	Tmem141.cSep08	499755	1619	625	3	93	transmembrane protein 141 (11.0 kD) (Tmem141) alternative variant cSep08, mRNA.
Tmem141	Tmem141.dSep08	499755	1957	1006	4	77	transmembrane protein 141 (8.7 kD) (Tmem141) alternative variant dSep08, mRNA.
Tmem143	Tmem143.bSep08	308593	10434	804	5	254	transmembrane protein 143 (Tmem143) alternative variant bSep08, mRNA.
Tmem143	Tmem143.cSep08	308593	7530	621	3	207	transmembrane protein 143 (Tmem143) alternative variant cSep08, mRNA.
Tmem144	Tmem144.bSep08	361968	9539	461	4	106	transmembrane protein 144 (Tmem144) alternative variant bSep08, mRNA.
Tmem144	Tmem144.cSep08	361968	4416	461	3	95	transmembrane protein 144 (Tmem144) alternative variant cSep08, mRNA.
Tmem146	Tmem146.aSep08	680264	36498	2368	22	725	transmembrane protein 146 (Tmem146) alternative variant aSep08, mRNA.
Tmem146	Tmem146.bSep08	680264	25003	388	5	87	transmembrane protein 146 (Tmem146) alternative variant bSep08, mRNA.
Tmem147	Tmem147.bSep08	292792	466	380	2	111	transmembrane protein 147 (Tmem147) alternative variant bSep08, mRNA.

Tmem147	Tmem147.cSep08	292792	487	415	2	79	transmembrane protein 147 (8.9 kD) (Tmem147) alternative variant cSep08, mRNA.
Tmem147	Tmem147.dSep08	292792	1543	308	3	77	transmembrane protein 147 (Tmem147) alternative variant dSep08, mRNA.
Tmem150	Tmem150.bSep08	245966	5007	2408	8	271	transmembrane protein 150 (29.0 kD) (Tmem150) alternative variant bSep08, complete mRNA.
Tmem150	Tmem150.dSep08	245966	3179	743	7	100	transmembrane protein 150 (10.7 kD) (Tmem150) alternative variant dSep08, mRNA.
Tmem150	Tmem150.eSep08	245966	861	572	2	69	transmembrane protein 150 (Tmem150) alternative variant eSep08, mRNA.
Tmem151a	Tmem151a.bSep08	309158	4592	740	2	191	transmembrane protein 151A (Tmem151a) alternative variant bSep08, mRNA.
Tmem151b	Tmem151b.aSep08	301253	1275	563		187	transmembrane protein 151B (Tmem151b) mRNA.
Tmem154	Tmem154.bSep08	361972	37306	697	5	179	transmembrane protein 154 (Tmem154) alternative variant bSep08, mRNA.
Tmem154	Tmem154.cSep08	361972	20129	425	1	138	transmembrane protein 154 (Tmem154) alternative variant cSep08, mRNA.
Tmem156	Tmem156.bSep08	498365	9516	594	2	69	transmembrane protein 156 (Tmem156) alternative variant bSep08, mRNA.
Tmem161b	Tmem161b.aSep08	309953	68882	1050	7	143	transmembrane protein 161B (16.8 kD) (Tmem161b) alternative variant aSep08, mRNA.
Tmem161b	Tmem161b.bSep08	309953	47732	375	3	116	transmembrane protein 161B (Tmem161b) alternative variant bSep08, mRNA.
Tmem161b	Tmem161b.cSep08	309953	67530	754	6	115	transmembrane protein 161B (13.6 kD) (Tmem161b) alternative variant cSep08, mRNA.
Tmem161b	Tmem161b.dSep08	309953	67559	758	6	48	transmembrane protein 161B (Tmem161b) alternative variant dSep08, mRNA.
Tmem163	Tmem163.bSep08	360839	157943	541	5	156	transmembrane protein 163 (Tmem163) alternative variant bSep08, mRNA.
Tmem163	Tmem163.cSep08	360839	5911	685	3	124	transmembrane protein 163 (Tmem163) alternative variant cSep08, mRNA.
Tmem165	Tmem165.bSep08	364137	747	392	1	116	transmembrane protein 165 (Tmem165) alternative variant bSep08, mRNA.
Tmem168	Tmem168.aSep08	312135	23129	3904	4	697	transmembrane protein 168 (79.8 kD) (Tmem168) alternative variant aSep08, mRNA.
Tmem169	Tmem169.bSep08	690294	23686	2499	1	297	transmembrane protein 169 (33.4 kD) (Tmem169) alternative variant bSep08, mRNA.
Tmem169	Tmem169.cSep08	690294	18830	343	1	84	transmembrane protein 169 (Tmem169) alternative variant cSep08, mRNA.
Tmem171	Tmem171.bSep08	293634	7136	697	3	202	transmembrane protein 171 (Tmem171) alternative variant bSep08, mRNA.
Tmem175	Tmem175.cSep08	305623	16170	1657	10	199	transmembrane protein 175 (22.4 kD) (Tmem175) alternative variant cSep08, complete mRNA.
Tmem175	Tmem175.dSep08	305623	4709	584	7	158	transmembrane protein 175 (Tmem175) alternative variant dSep08, mRNA.
Tmem175	Tmem175.fSep08	305623	11556	1052	4	86	transmembrane protein 175 (9.1 kD) (Tmem175) alternative variant fSep08, mRNA.

Tmem175	Tmem175.gSep08	305623	1242	455	2	61	transmembrane protein 175 (Tmem175) alternative variant gSep08, mRNA.
Tmem176b	Tmem176b.aSep08	171411	7220	1262	7	274	transmembrane protein 176B (29.7 kD) (Tmem176b) alternative variant aSep08, mRNA.
Tmem176b	Tmem176b.bSep08	171411	7429	1112	8	263	transmembrane protein 176B (28.5 kD) (Tmem176b) alternative variant bSep08, complete mRNA.
Tmem176b	Tmem176b.dSep08	171411	7188	805	7	258	transmembrane protein 176B (Tmem176b) alternative variant dSep08, mRNA.
Tmem176b	Tmem176b.eSep08	171411	7246	866	7	247	transmembrane protein 176B (Tmem176b) alternative variant eSep08, mRNA.
Tmem176b	Tmem176b.fSep08	171411	6669	1248	6	214	transmembrane protein 176B (Tmem176b) alternative variant fSep08, mRNA.
Tmem176b	Tmem176b.gSep08	171411	836	480	2	61	transmembrane protein 176B (Tmem176b) alternative variant gSep08, mRNA.
Tmem179b	Tmem179b.aSep08	690263	1948	837	5	195	transmembrane protein 179B (Tmem179b) alternative variant aSep08, mRNA.
Tmem179b	Tmem179b.cSep08	690263	1731	633	5	147	transmembrane protein 179B (16.5 kD) (Tmem179b) alternative variant cSep08, mRNA.
Tmem179b	Tmem179b.dSep08	690263	1314	1143	2	103	transmembrane protein 179B (11.5 kD) (Tmem179b) alternative variant dSep08, mRNA.
Tmem183a	Tmem183a.bSep08	289034	10719	790	6	244	transmembrane protein 183A (Tmem183a) alternative variant bSep08, mRNA.
Tmem183a	Tmem183a.cSep08	289034	8321	653	5	213	transmembrane protein 183A (Tmem183a) alternative variant cSep08, mRNA.
Tmem183a	Tmem183a.dSep08	289034	6614	2417	4	157	transmembrane protein 183A (Tmem183a) alternative variant dSep08, mRNA.
Tmem183a	Tmem183a.eSep08	289034	7247	466	3	119	transmembrane protein 183A (Tmem183a) alternative variant eSep08, mRNA.
Tmem184a	Tmem184a.bSep08	304325	6367	738	6	242	transmembrane protein 184A (Tmem184a) alternative variant bSep08, mRNA.
Tmem184a	Tmem184a.cSep08	304325	1219	403	2	89	transmembrane protein 184A (Tmem184a) alternative variant cSep08, mRNA.
Tmem184b	Tmem184b.aSep08	362959	43715	3324	9	463	transmembrane protein 184B (Tmem184b) alternative variant aSep08, mRNA.
Tmem184b	Tmem184b.bSep08	362959	3388	506	4	166	transmembrane protein 184B (Tmem184b) alternative variant bSep08, mRNA.
Tmem184b	Tmem184b.eSep08	362959	20440	386	2	45	transmembrane protein 184B (5.3 kD) (Tmem184b) alternative variant eSep08, mRNA.
Tmem188	Tmem188.aSep08	291914	14972	1806	5	126	transmembrane protein 188 (14.3 kD) (Tmem188) alternative variant aSep08, complete mRNA.
Tmem188	Tmem188.cSep08	291914	13996	755	4	101	transmembrane protein 188 (11.3 kD) (Tmem188) alternative variant cSep08, mRNA.
Tmem188	Tmem188.dSep08	291914	14015	772	4	95	transmembrane protein 188 (10.7 kD) (Tmem188) alternative variant dSep08, mRNA.
Tmem188	Tmem188.eSep08	291914	4858	717	1	68	transmembrane protein 188 (Tmem188) alternative variant eSep08, mRNA.

Tmem188	Tmem188.fSep08	291914	13820	505	3	23	transmembrane protein 188 (Tmem188) alternative variant fSep08, mRNA.
Tmem192	Tmem192.bSep08	361137	13161	822	5	202	transmembrane protein 192 (Tmem192) alternative variant bSep08, mRNA.
Tmem199	Tmem199.bSep08	303332	4144	1059	5	176	transmembrane protein 199 (19.9 kD) (Tmem199) alternative variant bSep08, mRNA.
Tmem200a	Tmem200a.bSep08	498987	797	261	2	46	transmembrane protein 200A (Tmem200a) alternative variant bSep08, mRNA.
Tmem204	Tmem204.bSep08	287129	3247	783	2	98	putative protein (Tmem204) alternative variant bSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.bSep08	140665	12010	1159	6	223	family rab3 member RAS oncogene (24.8 kD) (Tmem205andRab3d) alternative variant bSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.bSep08	300441	12010	1159	6	223	family rab3 member RAS oncogene (24.8 kD) (Tmem205andRab3d) alternative variant bSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.cSep08	140665	5594	688	4	216	transmembrane protein 205 (Tmem205andRab3d) alternative variant cSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.cSep08	300441	5594	688	4	216	transmembrane protein 205 (Tmem205andRab3d) alternative variant cSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.dSep08	140665	7534	732	5	211	GTP-binding protein like (Tmem205andRab3d) alternative variant dSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.dSep08	300441	7534	732	5	211	GTP-binding protein like (Tmem205andRab3d) alternative variant dSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.eSep08	140665	5571	728	5	189	transmembrane protein 205 (21.2 kD) (Tmem205andRab3d) alternative variant eSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.eSep08	300441	5571	728	5	189	transmembrane protein 205 (21.2 kD) (Tmem205andRab3d) alternative variant eSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.fSep08	140665	5566	649	4	189	transmembrane protein 205 (21.2 kD) (Tmem205andRab3d) alternative variant fSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.fSep08	300441	5566	649	4	189	transmembrane protein 205 (21.2 kD) (Tmem205andRab3d) alternative variant fSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.hSep08	140665	4676	617	3	159	transmembrane protein 205 (17.9 kD) (Tmem205andRab3d) alternative variant hSep08, mRNA.
Tmem205andRab3d	Tmem205andRab3d.hSep08	300441	4676	617	3	159	transmembrane protein 205 (17.9 kD) (Tmem205andRab3d) alternative variant hSep08, mRNA.
Tmem206	Tmem206.bSep08	305070	1720	722	2	87	transmembrane protein 206 (10.2 kD) (Tmem206) alternative variant bSep08, mRNA.
Tmem206	Tmem206.cSep08	305070	16184	328	3	76	transmembrane protein 206 (Tmem206) alternative variant cSep08, mRNA.
Tmem208	Tmem208.bSep08	291963	912	585	2	144	transmembrane protein 208 (Tmem208) alternative variant bSep08, mRNA.
Tmem208	Tmem208.cSep08	291963	2552	811	3	114	transmembrane protein 208 (Tmem208) alternative variant cSep08, mRNA.
Tmem208	Tmem208.dSep08	291963	2006	577	2	108	transmembrane protein 208 (Tmem208) alternative variant dSep08, mRNA.
Tmem208	Tmem208.eSep08	291963	2682	805	4	100	transmembrane protein 208 (11.0 kD) (Tmem208) alternative variant eSep08, mRNA.

Tmem209	Tmem209.bSep08	312200	6949	690	6	213	transmembrane protein 209 (Tmem209) alternative variant bSep08, mRNA.
Tmem209	Tmem209.cSep08	312200	4322	922	5	212	transmembrane protein 209 (23.1 kD) (Tmem209) alternative variant cSep08, mRNA.
Tmem209	Tmem209.dSep08	312200	7403	780	3	55	transmembrane protein 209 (Tmem209) alternative variant dSep08, mRNA.
Tmem209	Tmem209.eSep08	312200	1699	296	2	37	transmembrane protein 209 (Tmem209) alternative variant eSep08, mRNA.
Tmem211	Tmem211.aSep08	679783	855	650		50	transmembrane protein 211 (Tmem211) mRNA.
Tmf1	Tmf1.bSep08	114206	8227	1801	7	415	TATA element modulatory factor 1 (Tmf1) alternative variant bSep08, mRNA.
Tmf1	Tmf1.cSep08	114206	5622	3683	4	130	TATA element modulatory factor 1 (15.1 kD) (Tmf1) alternative variant cSep08, mRNA.
Tmf1	Tmf1.dSep08	114206	514	294	2	72	TATA element modulatory factor 1 (Tmf1) alternative variant dSep08, mRNA.
Tmie	Tmie.bSep08	501061	4488	681	2	75	putative protein (8.5 kD) (Tmie) alternative variant bSep08, mRNA.
Tmigd1	Tmigd1.aSep08	363654	5778	243		79	putative protein of vertebrate origin (Tmigd1) mRNA.
Tmlhe	Tmlhe.bSep08	170898	26851	714	4	237	trimethyllysine hydroxylase, epsilon (Tmlhe) alternative variant bSep08, mRNA.
Tmlhe	Tmlhe.cSep08	170898	12491	784	4	219	trimethyllysine hydroxylase, epsilon (Tmlhe) alternative variant cSep08, mRNA.
Tmod1	Tmod1.aSep08	25566	67366	1785	8	540	tropomodulin 1 (Tmod1) alternative variant aSep08, mRNA.
Tmod1	Tmod1.cSep08	25566	50900	602	5	155	tropomodulin 1 (Tmod1) alternative variant cSep08, mRNA.
Tmod1	Tmod1.dSep08	25566	37086	1372	3	72	tropomodulin 1 (Tmod1) alternative variant dSep08, mRNA.
Tmod2	Tmod2.bSep08	58814	26131	925	6	174	tropomodulin 2 (Tmod2) alternative variant bSep08, mRNA.
Tmod2	Tmod2.cSep08	58814	24177	615	5	147	tropomodulin 2 (Tmod2) alternative variant cSep08, mRNA.
Tmod2	Tmod2.dSep08	58814	5677	534	4	99	tropomodulin 2 (Tmod2) alternative variant dSep08, mRNA.
Tmod3	Tmod3.bSep08	300838	17903	3337	8	213	tropomodulin 3 (23.8 kD) (Tmod3) alternative variant bSep08, mRNA.
Tmod3	Tmod3.cSep08	300838	3401	705	3	124	tropomodulin 3 (Tmod3) alternative variant cSep08, mRNA.
Tmod3	Tmod3.dSep08	300838	5658	595	3	21	tropomodulin 3 (2.5 kD) (Tmod3) alternative variant dSep08, mRNA.
Tmod4	Tmod4.bSep08	295261	3197	765	7	184	tropomodulin 4 (Tmod4) alternative variant bSep08, mRNA.
Tmod4	Tmod4.cSep08	295261	3135	633	6	176	tropomodulin 4 (Tmod4) alternative variant cSep08, mRNA.
Tmpo	Tmpo.aSep08	25359	12435	3733	4	692	thymopoietin (74.9 kD) (Tmpo) alternative variant aSep08, mRNA.
Tmpo	Tmpo.cSep08	25359	24923	1939	7	416	thymopoietin (Tmpo) alternative variant cSep08, mRNA.
Tmpo	Tmpo.dSep08	25359	18609	727	7	241	thymopoietin (Tmpo) alternative variant dSep08, mRNA.
Tmpo	Tmpo.eSep08	25359	9810	891	4	174	thymopoietin (Tmpo) alternative variant eSep08, mRNA.
Tmpo	Tmpo.gSep08	25359	6134	271	3	90	thymopoietin (Tmpo) alternative variant gSep08, mRNA.

Tmprss2	Tmprss2.bSep08	156435	28874	818	7	215	transmembrane protease, serine 2 (Tmprss2) alternative variant bSep08, mRNA.
Tmprss4	Tmprss4.aSep08	367074	2769	1202		101	transmembrane protease, serine 4 (Tmprss4) mRNA.
Tmprss5	Tmprss5.bSep08	266681	595	311	2	103	transmembrane protease, serine 5 (spinesin) (Tmprss5) alternative variant bSep08, mRNA.
Tmprss6	Tmprss6.bSep08	315388	3543	707	4	223	transmembrane serine protease 6 (Tmprss6) alternative variant bSep08, mRNA.
Tmprss6	Tmprss6.cSep08	315388	604	399	2	99	transmembrane serine protease 6 (Tmprss6) alternative variant cSep08, mRNA.
Tmprss9	Tmprss9.aSep08	314636	1465	569		137	transmembrane protease, serine 9 (Tmprss9) mRNA.
Tmsb10	Tmsb10.aSep08	50665	1042	486	3	75	thymosin, beta 10 (Tmsb10) alternative variant aSep08, mRNA.
Tmtc1	Tmtc1.aSep08	362465	13625	357		119	transmembrane and tetratricopeptide repeat containing 1 (Tmtc1) mRNA.
Tmtc2	Tmtc2.aSep08	299762	52149	415		138	transmembrane and tetratricopeptide repeat containing 2 (Tmtc2) mRNA.
Tmtc4	Tmtc4.bSep08	290501	45268	2003	14	606	transmembrane and tetratricopeptide repeat containing 4 (67.7 kD) (Tmtc4) alternative variant bSep08, mRNA.
Tmtc4	Tmtc4.eSep08	290501	946	349	2	36	transmembrane and tetratricopeptide repeat containing 4 (Tmtc4) alternative variant eSep08, mRNA.
Tmub2	Tmub2.bSep08	303567	3186	784	3	260	putative protein of bilateral origin (Tmub2) alternative variant bSep08, mRNA.
Tmub2	Tmub2.cSep08	303567	3169	822	4	229	putative protein of bilateral origin (Tmub2) alternative variant cSep08, mRNA.
Tnfaip1	Tnfaip1.bSep08	287543	2418	452	2	28	tumor necrosis factor, alpha-induced protein 1 (endothelial) (Tnfaip1) alternative variant bSep08, mRNA.
Tnfaip2	Tnfaip2.aSep08	299339	6171	1425	6	253	tumor necrosis factor, alpha-induced protein 2 (Tnfaip2) alternative variant aSep08, mRNA.
Tnfaip2	Tnfaip2.bSep08	299339	10275	267	2	68	tumor necrosis factor, alpha-induced protein 2 (Tnfaip2) alternative variant bSep08, mRNA.
Tnfaip2	Tnfaip2.dSep08	299339	1882	273	3	48	tumor necrosis factor, alpha-induced protein 2 (Tnfaip2) alternative variant dSep08, mRNA.
Tnfaip8	Tnfaip8.aSep08	307428	63037	735	2	141	tumor necrosis factor, alpha-induced protein 8 (Tnfaip8) alternative variant aSep08, mRNA.
Tnfaip8l1	Tnfaip8l1.aSep08	301131	8572	372	1	114	tumor necrosis factor, alpha-induced protein 8-like 1 (Tnfaip8l1) alternative variant aSep08, mRNA.
Tnfaip8l1	Tnfaip8l1.bSep08	301131	9001	799	1	99	tumor necrosis factor, alpha-induced protein 8-like 1 (Tnfaip8l1) alternative variant bSep08, mRNA.
Tnfrsf1a	Tnfrsf1a.aSep08	25625	12940	2527	1	531	tumor necrosis factor receptor superfamily, member 1a (Tnfrsf1a) alternative variant aSep08, mRNA.
Tnfrsf1a	Tnfrsf1a.bSep08	25625	11701	1230	2	306	tumor necrosis factor receptor superfamily, member 1a (Tnfrsf1a) alternative variant bSep08, mRNA.
Tnfrsf9	Tnfrsf9.bSep08	500590	10279	473	2	98	tumor necrosis factor receptor superfamily, member 9 (Tnfrsf9) alternative variant bSep08, mRNA.
Tnfrsf9	Tnfrsf9.cSep08	500590	4356	330	1	54	tumor necrosis factor receptor superfamily, member 9 (Tnfrsf9) alternative variant cSep08, mRNA.



Tnfrsf10b	Tnfrsf10b.bSep08	364420	17170	927		219	tumor necrosis factor receptor superfamily, member 10b (Tnfrsf10b) alternative variant bSep08, mRNA.
Tnfrsf11a	Tnfrsf11a.aSep08	498206	12060	716		238	tumor necrosis factor receptor superfamily, member 11a (Tnfrsf11a) mRNA.
Tnfrsf13c	Tnfrsf13c.aSep08	500910	1380	910		125	tumor necrosis factor receptor superfamily, member 13c (Tnfrsf13c) mRNA.
Tnfrsf14	Tnfrsf14.cSep08	366518	4363	617	5	118	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator) (Tnfrsf14) alternative variant cSep08, mRNA.
Tnfrsf14	Tnfrsf14.eSep08	366518	939	777	2	74	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator) (Tnfrsf14) alternative variant eSep08, mRNA.
Tnfrsf22	Tnfrsf22.aSep08	686008	4502	641		132	tumor necrosis factor receptor superfamily, member 22 (14.4 kD) (Tnfrsf22) mRNA.
Tnfrsf25	Tnfrsf25.aSep08	500592	2800	1174	7	303	tumor necrosis factor receptor superfamily, member 25 (Tnfrsf25) alternative variant aSep08, mRNA.
Tnfrsf25	Tnfrsf25.bSep08	500592	2713	723	7	157	tumor necrosis factor receptor superfamily, member 25 (Tnfrsf25) alternative variant bSep08, mRNA.
Tnfrsf25	Tnfrsf25.cSep08	500592	2252	592	6	126	tumor necrosis factor receptor superfamily, member 25 (Tnfrsf25) alternative variant cSep08, mRNA.
Tnfrsf25	Tnfrsf25.dSep08	500592	904	701	2	72	tumor necrosis factor receptor superfamily, member 25 (Tnfrsf25) alternative variant dSep08, mRNA.
Tnfrsf26	Tnfrsf26.aSep08	361685	20454	2804	1	211	tumor necrosis factor receptor superfamily, member 26 (Tnfrsf26) alternative variant aSep08, mRNA.
Tnfrsf26	Tnfrsf26.bSep08	361685	18436	738	1	186	tumor necrosis factor receptor superfamily, member 26 (Tnfrsf26) alternative variant bSep08, mRNA.
Tnfsf12	Tnfsf12.aSep08	360548	8673	916	1	275	tumor necrosis factor ligand superfamily member 12 (Tnfsf12) alternative variant aSep08, mRNA.
Tnfsf13	Tnfsf13.aSep08	287437	3547	2052	6	241	tumor necrosis factor (ligand) superfamily, member 13 (26.7 kD) (Tnfsf13) alternative variant aSep08, mRNA.
Tnfsf13	Tnfsf13.cSep08	287437	1796	717	3	118	tumor necrosis factor (ligand) superfamily, member 13 (13.3 kD) (Tnfsf13) alternative variant cSep08, mRNA.
Tnfsf13	Tnfsf13.dSep08	287437	1460	714	2	106	tumor necrosis factor (ligand) superfamily, member 13 (Tnfsf13) alternative variant dSep08, mRNA.
Tnip1	Tnip1.bSep08	363599	13198	1833	11	382	TNFAIP3 interacting protein 1 (Tnip1) alternative variant bSep08, mRNA.
Tnip1	Tnip1.cSep08	363599	30533	622	5	206	TNFAIP3 interacting protein 1 (Tnip1) alternative variant cSep08, mRNA.
Tnip1	Tnip1.dSep08	363599	15891	260	3	62	TNFAIP3 interacting protein 1 (Tnip1) alternative variant dSep08, mRNA.
Tnk2	Tnk2.bSep08	303882	11767	2461	7	565	tyrosine kinase, non-receptor, 2 (Tnk2) alternative variant bSep08, mRNA.
Tnk2	Tnk2.dSep08	303882	2392	418	3	85	tyrosine kinase, non-receptor, 2 (Tnk2) alternative variant dSep08, mRNA.
Tnks	Tnks.bSep08	290794	46388	1783	8	474	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase (Tnks) alternative variant bSep08, mRNA.

Tnks	Tnks.cSep08	290794	18060	872	3	174	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase (Tnks) alternative variant cSep08, mRNA.
Tnks	Tnks.dSep08	290794	4071	348	3	116	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase (Tnks) alternative variant dSep08, mRNA.
Tnks	Tnks.eSep08	290794	3169	414	4	95	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase (Tnks) alternative variant eSep08, mRNA.
Tnks1bp1	Tnks1bp1.aSep08	295707	10264	2627		721	tankyrase 1 binding protein 1 (Tnks1bp1) mRNA.
Tnks2	Tnks2.bSep08	309512	1529	514	2	48	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2 (Tnks2) alternative variant bSep08, mRNA.
Tnks2	Tnks2.cSep08	309512	1851	577	2	48	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2 (Tnks2) alternative variant cSep08, mRNA.
Tnn	Tnn.bSep08	304913	9300	379	3	126	tenascin N (Tnn) alternative variant bSep08, mRNA.
Tnnc2	Tnnc2.bSep08	296369	2938	720	2	145	troponin C2, fast (16.4 kD) (Tnnc2) alternative variant bSep08, mRNA.
Tnni1	Tnni1.aSep08	29388	12230	846	3	171	troponin I, skeletal, slow 1 (19.8 kD) (Tnni1) alternative variant aSep08, mRNA.
Tnni1	Tnni1.bSep08	29388	12128	822	4	171	troponin I, skeletal, slow 1 (19.8 kD) (Tnni1) alternative variant bSep08, mRNA.
Tnni1	Tnni1.dSep08	29388	12383	1029	4	171	troponin I, skeletal, slow 1 (19.8 kD) (Tnni1) alternative variant dSep08, complete mRNA.
Tnni1	Tnni1.eSep08	29388	10234	645	3	136	troponin I, skeletal, slow 1 (Tnni1) alternative variant eSep08, mRNA.
Tnni1	Tnni1.fSep08	29388	10106	759	3	104	troponin I, skeletal, slow 1 (Tnni1) alternative variant fSep08, mRNA.
Tnni2	Tnni2.aSep08	29389	2009	757	1	161	troponin I type 2 (skeletal, fast) (18.8 kD) (Tnni2) alternative variant aSep08, mRNA.
Tnni3	Tnni3.cSep08	29248	1469	748	4	82	troponin I type 3 (cardiac) (8.9 kD) (Tnni3) alternative variant cSep08, mRNA.
Tnni3	Tnni3.dSep08	29248	2190	1761	3	82	troponin I type 3 (cardiac) (8.9 kD) (Tnni3) alternative variant dSep08, mRNA.
Tnni3	Tnni3.eSep08	29248	1037	777	2	46	troponin I type 3 (cardiac) (Tnni3) alternative variant eSep08, mRNA.
Tnni3	Tnni3.fSep08	29248	1813	500	3	62	troponin I type 3 (cardiac) (Tnni3) alternative variant fSep08, mRNA.
Tnni3k	Tnni3k.bSep08	295531	14022	535	1	102	TNNI3 interacting kinase (Tnni3k) alternative variant bSep08, mRNA.
Tnnt1	Tnnt1.bSep08	171409	8557	739	12	232	troponin T1, skeletal, slow (Tnnt1) alternative variant bSep08, mRNA.
Tnnt2	Tnnt2.aSep08	24837	17675	1027	14	305	troponin T2 cardiac (Tnnt2) alternative variant aSep08, mRNA.
Tnnt2	Tnnt2.bSep08	24837	13356	1785	13	295	troponin T2 cardiac CRA a (34.6 kD) (Tnnt2) alternative variant bSep08, mRNA.
Tnnt2	Tnnt2.cSep08	24837	9359	1306	11	268	troponin T2 cardiac (Tnnt2) alternative variant cSep08, mRNA.
Tnnt2	Tnnt2.eSep08	24837	8353	1359	5	93	putative protein (Tnnt2) alternative variant eSep08, mRNA.
Tnnt2	Tnnt2.fSep08	24837	11661	894	5	89	putative protein (Tnnt2) alternative variant fSep08, mRNA.

Tnnt2	Tnnt2.gSep08	24837	1910	497	5	81	troponin T (9.8 kD) (Tnnt2) alternative variant gSep08, mRNA.
Tnnt2	Tnnt2.hSep08	24837	990	716	2	49	putative protein (5.9 kD) (Tnnt2) alternative variant hSep08, mRNA.
Tnnt3	Tnnt3.aSep08	24838	16943	1022	17	259	troponin T3, skeletal, fast (30.7 kD) (Tnnt3) alternative variant aSep08, complete mRNA.
Tnnt3	Tnnt3.bSep08	24838	16944	990	15	248	troponin T3, skeletal, fast (29.3 kD) (Tnnt3) alternative variant bSep08, complete mRNA.
Tnnt3	Tnnt3.cSep08	24838	12561	309	9	103	troponin T3, skeletal, fast (Tnnt3) alternative variant cSep08, mRNA.
Tnnt3	Tnnt3.dSep08	24838	4333	413	4	50	troponin T3, skeletal, fast (Tnnt3) alternative variant dSep08, mRNA.
Tnnt3	Tnnt3.eSep08	24838	3897	609	6	32	troponin T3, skeletal, fast (3.8 kD) (Tnnt3) alternative variant eSep08, mRNA.
Tnpo1	Tnpo1.aSep08	309126	42686	2380	3	780	transportin 1 (Tnpo1) alternative variant aSep08, mRNA.
Tnpo1	Tnpo1.bSep08	309126	16045	1783		352	transportin 1 (Tnpo1) alternative variant bSep08, mRNA.
Tnpo1	Tnpo1.cSep08	309126	4085	624	1	94	transportin 1 (Tnpo1) alternative variant cSep08, mRNA.
Tnpo2	Tnpo2.bSep08	304670	7385	548	5	182	transportin 2 (importin 3, karyopherin beta 2b) (Tnpo2) alternative variant bSep08, mRNA.
Tnpo2	Tnpo2.cSep08	304670	3647	382	4	88	transportin 2 (importin 3, karyopherin beta 2b) (Tnpo2) alternative variant cSep08, mRNA.
Tnpo2	Tnpo2.dSep08	304670	1183	943	4	83	transportin 2 (importin 3, karyopherin beta 2b) (Tnpo2) alternative variant dSep08, mRNA.
Tnpo3	Tnpo3.bSep08	296954	20314	420	3	50	transportin 3 (Tnpo3) alternative variant bSep08, mRNA.
Tnr	Tnr.bSep08	25567	17166	445	4	148	tenascin R (Tnr) alternative variant bSep08, mRNA.
Tnr	Tnr.cSep08	25567	868	492	2	134	tenascin R (Tnr) alternative variant cSep08, mRNA.
Tnr	Tnr.dSep08	25567	4958	352	4	117	tenascin R (Tnr) alternative variant dSep08, mRNA.
Tnrc4	Tnrc4.bSep08	499669	3326	742	8	246	trinucleotide repeat containing 4 (Tnrc4) alternative variant bSep08, mRNA.
Tnrc4	Tnrc4.cSep08	499669	2909	1190	6	241	trinucleotide repeat containing 4 (Tnrc4) alternative variant cSep08, mRNA.
Tnrc4	Tnrc4.dSep08	499669	3663	1336	3	80	trinucleotide repeat containing 4 (Tnrc4) alternative variant dSep08, mRNA.
Tnrc6a	Tnrc6a.aSep08	308971	17916	4544	15	720	trinucleotide repeat containing 6a (Tnrc6a) alternative variant aSep08, mRNA.
Tnrc6a	Tnrc6a.bSep08	308971	3535	366	3	122	trinucleotide repeat containing 6a (Tnrc6a) alternative variant bSep08, mRNA.
Tnrc6a	Tnrc6a.cSep08	308971	1311	376	2	98	trinucleotide repeat containing 6a (Tnrc6a) alternative variant cSep08, mRNA.
Tnrc6b	Tnrc6b.dSep08	192178	144938	360	5	44	trinucleotide repeat containing 6B (Tnrc6b) alternative variant dSep08, mRNA.
Tns1	Tns1.bSep08	301509	1648	459	2	46	tensin 1 (5.4 kD) (Tns1) alternative variant bSep08, mRNA.
Tnxb	Tnxb.aSep08	415089	17988	4570	5	1037	tenascin XB (Tnxb) alternative variant aSep08, mRNA.
Tnxb	Tnxb.bSep08	415089	9635	3691	1	721	tenascin XB (Tnxb) alternative variant bSep08, mRNA.
Tnxb	Tnxb.cSep08	415089	2133	1356	1	387	tenascin XB (Tnxb) alternative variant cSep08, mRNA.

Tnxb	Tnxb.dSep08	415089	2489	980		304	tenascin XB (Tnxb) alternative variant dSep08, mRNA.
Tnxb	Tnxb.eSep08	415089	1366	803	2	204	tenascin XB (Tnxb) alternative variant eSep08, mRNA.
toby	toby.aSep08		1249	384		41	putative protein (toby) mRNA.
tochy	tochy.aSep08		1652	640		44	putative protein (5.0 kD) (tochy) mRNA.
todar	todar.aSep08		7921	606		202	hydrocephalus inducing (todar) mRNA.
Toe1	Toe1.aSep08	298443	3475	2027	7	426	target of EGR1, member 1 (nuclear) (Toe1) alternative variant aSep08, mRNA.
Toe1	Toe1.cSep08	298443	1952	1605	2	113	target of EGR1, member 1 (nuclear) (Toe1) alternative variant cSep08, mRNA.
Toe1	Toe1.dSep08	298443	1099	409	3	82	target of EGR1, member 1 (nuclear) (Toe1) alternative variant dSep08, mRNA.
Toe1	Toe1.eSep08	298443	1782	773	4	113	target of EGR1, member 1 (nuclear) (Toe1) alternative variant eSep08, mRNA.
Toe1	Toe1.fSep08	298443	1100	620	3	59	target of EGR1, member 1 (nuclear) (Toe1) alternative variant fSep08, mRNA.
toflo	toflo.aSep08		2949	842		61	putative protein (6.7 kD) (toflo) mRNA.
toflu	toflu.aSep08		1447	814		51	putative protein (5.7 kD) (toflu) mRNA.
togar	togar.aSep08		1255	490		105	unc-13 homolog A CRA a (12.1 kD) (togar) mRNA.
toja	toja.aSep08		2994	522		36	putative protein (4.4 kD) (toja) mRNA.
tojey	tojey.aSep08		8855	746	5	248	platelet-derived growth factor receptor (tojey) alternative variant aSep08, mRNA.
tojey	tojey.bSep08		1066	423	1	97	platelet-derived growth factor receptor (tojey) alternative variant bSep08, mRNA.
tokee	tokee.aSep08		25795	341		87	putative protein (tokee) mRNA.
tokler	tokler.aSep08		692	569		42	putative protein (4.8 kD) (tokler) mRNA.
Tollip	Tollip.bSep08	361677	772	678	2	116	putative endoplasmic reticulum protein, with a transmembrane domain (12.5 kD) (Tollip) alternative variant bSep08, mRNA.
tolo	tolo.aSep08		5632	279		34	putative protein (tolo) mRNA.
Tom1l2	Tom1l2.bSep08	360537	86997	592	2	197	target of myb1-like 2 (chicken) (Tom1l2) alternative variant bSep08, mRNA.
Tom1l2	Tom1l2.cSep08	360537	31432	385	2	127	target of myb1-like 2 (chicken) (Tom1l2) alternative variant cSep08, mRNA.
tomee	tomee.aSep08		13319	577		103	putative protein (tomee) mRNA.
Tomm7	Tomm7.aSep08	685620	6842	377	3	55	translocase of outer mitochondrial membrane 7 homolog (yeast) (6.2 kD) (Tomm7) alternative variant aSep08, complete mRNA.
Tomm20	Tomm20.aSep08	266601	12331	1006	1	149	translocase of outer mitochondrial membrane 20 homolog (yeast) (16.8 kD) (Tomm20) alternative variant aSep08, complete mRNA.
Tomm34	Tomm34.bSep08	311621	4471	613	2	123	translocase of outer mitochondrial membrane 34 (Tomm34) alternative variant bSep08, mRNA.
Tomm40	Tomm40.aSep08	308416	12098	1558	10	361	translocase of outer mitochondrial membrane 40 homolog (yeast) (37.9 kD) (Tomm40) alternative variant aSep08, complete mRNA.

Tomm40	Tomm40.cSep08	308416	10015	743	7	238	translocase of outer mitochondrial membrane 40 homolog (yeast) (Tomm40) alternative variant cSep08, mRNA.
Tomm40	Tomm40.dSep08	308416	2241	927	4	114	translocase of outer mitochondrial membrane 40 homolog (yeast) (12.5 kD) (Tomm40) alternative variant dSep08, mRNA.
tonoy	tonoy.aSep08		2858	795	2	39	putative protein (4.4 kD) (tonoy) alternative variant aSep08, mRNA.
Top1	Top1.aSep08	64550	62267	1362		305	topoisomerase (DNA) I (36.8 kD) (Top1) mRNA.
Top1mt	Top1mt.bSep08	300029	10487	687	4	171	DNA topoisomerase 1, mitochondrial (Top1mt) alternative variant bSep08, mRNA.
Top1mt	Top1mt.cSep08	300029	9919	814	3	164	DNA topoisomerase 1, mitochondrial (Top1mt) alternative variant cSep08, mRNA.
Top1mt	Top1mt.dSep08	300029	10208	694	4	144	DNA topoisomerase 1, mitochondrial (Top1mt) alternative variant dSep08, mRNA.
Top2a	Top2a.aSep08	360243	17446	3610	20	915	topoisomerase (DNA) II alpha (Top2a) alternative variant aSep08, mRNA.
Top2a	Top2a.bSep08	360243	8670	594	5	132	topoisomerase (DNA) II alpha (Top2a) alternative variant bSep08, mRNA.
Top2a	Top2a.cSep08	360243	3016	1051	5	118	topoisomerase (DNA) II alpha (Top2a) alternative variant cSep08, mRNA.
Top3b	Top3b.bSep08	287930	4012	749	6	204	topoisomerase (DNA) III beta (Top3b) alternative variant bSep08, mRNA.
Top3b	Top3b.cSep08	287930	18277	749	6	193	topoisomerase (DNA) III beta (Top3b) alternative variant cSep08, mRNA.
Top3b	Top3b.dSep08	287930	5329	2817	4	178	topoisomerase (DNA) III beta (20.5 kD) (Top3b) alternative variant dSep08, mRNA.
Topbp1	Topbp1.aSep08	315969	14750	2322	12	635	topoisomerase (DNA) II binding protein 1 (Topbp1) alternative variant aSep08, mRNA.
Topbp1	Topbp1.bSep08	315969	14643	1816	9	397	topoisomerase (DNA) II binding protein 1 (44.5 kD) (Topbp1) alternative variant bSep08, mRNA.
Topbp1	Topbp1.cSep08	315969	1986	669	3	101	topoisomerase (DNA) II binding protein 1 (Topbp1) alternative variant cSep08, mRNA.
Topoisom_I.0	Topoisom_I.0.aSep08		11558	2329	1	311	topoisomerase I (Topoisom_I.0) alternative variant aSep08, mRNA.
topor	topor.aSep08		16648	693		73	putative protein (8.4 kD) (topor) mRNA.
Tor1aip1	Tor1aip1.bSep08	246314	26862	2064	9	396	torsin A interacting protein 1 (44.4 kD) (Tor1aip1) alternative variant bSep08, mRNA.
Tor1aip1	Tor1aip1.cSep08	246314	27106	934	10	255	torsin A interacting protein 1 (Tor1aip1) alternative variant cSep08, mRNA.
Tor1aip2	Tor1aip2.bSep08	304881	11572	2353	3	121	putative protein (13.6 kD) (Tor1aip2) alternative variant bSep08, complete mRNA.
Tor1aip2	Tor1aip2.cSep08	304881	11158	1892	3	120	putative protein (Tor1aip2) alternative variant cSep08, mRNA.
Tor1aip2	Tor1aip2.dSep08	304881	10230	904	2	89	putative protein (9.9 kD) (Tor1aip2) alternative variant dSep08, mRNA.
Tor1aip2	Tor1aip2.eSep08	304881	19506	808	5	68	torsin A interacting protein 2 (Tor1aip2) alternative variant eSep08, mRNA.

Tor1b	Tor1b.bSep08	311854	3638	819	3	163	torsin family 1 member B (Tor1b) alternative variant bSep08, mRNA.
Tor1b	Tor1b.cSep08	311854	1712	1030	2	158	torsin family 1 member B CRA a (18.2 kD) (Tor1b) alternative variant cSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.cSep08	362112	4771	1932	5	293	torsin family 2 member A (Tor2aandPtrh1) alternative variant cSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.cSep08	362113	4771	1932	5	293	torsin family 2 member A (Tor2aandPtrh1) alternative variant cSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.dSep08	362112	1785	934	5	208	peptidyl-tRNA hydrolase 1 homolog (Tor2aandPtrh1) alternative variant dSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.dSep08	362113	1785	934	5	208	peptidyl-tRNA hydrolase 1 homolog (Tor2aandPtrh1) alternative variant dSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.eSep08	362112	2458	727	3	168	family 2 member a (Tor2aandPtrh1) alternative variant eSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.eSep08	362113	2458	727	3	168	family 2 member a (Tor2aandPtrh1) alternative variant eSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.fSep08	362112	6846	659	2	72	peptidyl-tRNA hydrolase 1 homolog like (Tor2aandPtrh1) alternative variant fSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.fSep08	362113	6846	659	2	72	peptidyl-tRNA hydrolase 1 homolog like (Tor2aandPtrh1) alternative variant fSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.gSep08	362112	668	390	3	56	peptidyl-tRNA hydrolase 1 homolog (Tor2aandPtrh1) alternative variant gSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.gSep08	362113	668	390	3	56	peptidyl-tRNA hydrolase 1 homolog (Tor2aandPtrh1) alternative variant gSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.hSep08	362112	923	537	2	37	family 2 member A (Tor2aandPtrh1) alternative variant hSep08, mRNA.
Tor2aandPtrh1	Tor2aandPtrh1.hSep08	362113	923	537	2	37	family 2 member A (Tor2aandPtrh1) alternative variant hSep08, mRNA.
torby	torby.aSep08		12317	744		247	microrchidia 4 CRA a (torby) mRNA.
torchy	torchy.aSep08		1679	311		103	ino80 complex homolog 1 (torchy) mRNA.
tordar	tordar.aSep08		85671	419		44	putative protein (tordar) mRNA.
torflo	torflo.aSep08		13443	732		83	putative protein (9.6 kD) (torflo) mRNA.
torflu	torflu.aSep08		12020	920	1	244	adaptor-related protein complex 3 beta CRA b (torflu) alternative variant aSep08, mRNA.
torflu	torflu.bSep08		15948	773	1	153	adaptor-related protein complex 3 beta CRA c (torflu) alternative variant bSep08, mRNA.
torgar	torgar.bSep08		5524	184	2	46	putative protein (torgar) alternative variant bSep08, mRNA.
torja	torja.aSep08		398	251		83	T cell receptor (torja) mRNA.
torjey	torjey.aSep08		4577	3702		94	ubiquitin specific peptidase 46 CRA a (torjey) mRNA.
torkee	torkee.aSep08		37145	793	4	137	putative protein (14.8 kD) (torkee) alternative variant aSep08, mRNA.
torkee	torkee.bSep08		22110	760	2	38	putative protein (torkee) alternative variant bSep08, mRNA.
torkler	torkler.aSep08		2062	600		95	putative protein of mammalian origin (torkler) mRNA.
torlo	torlo.bSep08		932	652	2	20	putative protein (torlo) alternative variant bSep08, mRNA.
tormee	tormee.aSep08		18103	467		112	CRA b like (tormee) mRNA.

tornoy	tornoy.aSep08		50425	782	2	201	thyroid hormone receptor interactor 12 CRA c (tornoy) alternative variant aSep08, mRNA.
tornoy	tornoy.bSep08		1544	281	1	93	thyroid hormone receptor interactor 12 (tornoy) alternative variant bSep08, mRNA.
torpor	torpor.aSep08		6418	418		102	interphotoreceptor matrix proteoglycan 1 (torpor) mRNA.
torsa	torsa.aSep08		3375	2441		83	death-inducing-protein CRA e (torsa) mRNA.
torshee	torshee.aSep08		29790	439		91	putative protein (torshee) mRNA.
tortu	tortu.aSep08		4270	350		116	WD repeat domain 35 (tortu) mRNA.
torvar	torvar.aSep08		4800	218		72	procollagen-lysine 1 (torvar) mRNA.
torwey	torwey.aSep08		3088	576		146	von Willebrand factor (torwey) mRNA.
tosa	tosa.aSep08		2088	475		102	putative protein (tosa) mRNA.
toshee	toshee.aSep08		13280	386		92	putative protein (toshee) mRNA.
totu	totu.aSep08		113069	735		156	putative protein (totu) mRNA.
tovar	tovar.aSep08		1690	325		108	putative protein, with a coiled coil domain, of mammalian origin (tovar) mRNA.
towey	towey.aSep08		2255	493		164	nucleolar protein 1 (towey) mRNA.
Tox	Tox.bSep08	362481	2244	2054	2	73	thymocyte selection-associated high mobility group box (Tox) alternative variant bSep08, mRNA.
Tox3	Tox3.bSep08	291908	1963	1707	2	172	TOX high mobility group box family member 3 (Tox3) alternative variant bSep08, mRNA.
Tox3	Tox3.cSep08	291908	83682	475	3	135	TOX high mobility group box family member 3 (Tox3) alternative variant cSep08, mRNA.
toyby	toyby.aSep08		15195	344		42	putative protein (4.6 kD) (toyby) mRNA.
toychy	toychy.aSep08		5120	394		131	ino80 complex homolog 1 (toychy) mRNA.
toydar	toydar.aSep08		1023	473		21	putative protein (toydar) mRNA.
toydoy	toydoy.aSep08		1466	740	3	79	putative protein (toydoy) alternative variant aSep08, mRNA.
toydoy	toydoy.bSep08		1645	796	2	41	putative protein (4.9 kD) (toydoy) alternative variant bSep08, mRNA.
toydoy	toydoy.cSep08		1536	455	3	30	putative protein (3.3 kD) (toydoy) alternative variant cSep08, mRNA.
toyflo	toyflo.aSep08		4914	1784		15	putative protein (toyflo) mRNA.
toyflu	toyflu.aSep08		1361	460		128	CRA b (toyflu) mRNA.
toygar	toygar.bSep08		1847	298	2	47	putative protein (toygar) alternative variant bSep08, mRNA.
toyja	toyja.aSep08		4592	883		108	putative nuclear protein (11.7 kD) (toyja) alternative variant aSep08, mRNA.
toyjey	toyjey.aSep08		24889	758	4	49	putative protein (toyjey) alternative variant aSep08, mRNA.
toyjey	toyjey.bSep08		4173	710	3	80	putative cytoplasmic protein (8.9 kD) (toyjey) alternative variant bSep08, mRNA.
toykee	toykee.aSep08		6979	703		82	putative protein (toykee) mRNA.
toykler	toykler.aSep08		3556	805		124	putative protein, with a coiled coil domain, of mammalian origin (toykler) mRNA.
toylo	toylo.aSep08		7427	776		258	poly-specific ribonuclease PAN3b (toylo) mRNA.
toymee	toymee.aSep08		10431	762		45	putative protein (5.4 kD) (toymee) mRNA.

toynoy	toynoy.aSep08		7207	665		81	putative protein (8.7 kD) (toynoy) mRNA.
toypor	toypor.aSep08		28031	702		75	putative protein (toypor) mRNA.
toysa	toysa.aSep08		18966	363		83	tbc1 domain family member 22a (toysa) mRNA.
toyshee	toyshee.aSep08		606	249		44	putative protein (toyshee) mRNA.
toytu	toytu.aSep08		3767	863		58	putative protein (5.9 kD) (toytu) mRNA.
toyvar	toyvar.aSep08		2770	856		195	5 10-methylenetetrahydrofolate reductase CRA a (toyvar) mRNA.
toywey	toywey.aSep08		15203	1105		158	transmembrane protein 16B (toywey) mRNA.
Tpbg	Tpbg.bSep08	83684	1397	721	2	148	trophoblast glycoprotein (Tpbg) alternative variant bSep08, mRNA.
Tpbg	Tpbg.cSep08	83684	442	347	2	25	trophoblast glycoprotein (Tpbg) alternative variant cSep08, mRNA.
Tpbpa	Tpbpa.bSep08	64509	1303	605	4	107	trophoblast specific protein alpha (Tpbpa) alternative variant bSep08, mRNA.
Tpcn1	Tpcn1.aSep08	246215	16267	3520	18	492	two pore channel 1 (Tpcn1) alternative variant aSep08, mRNA.
Tpcn2	Tpcn2.bSep08	309139	29694	3008	25	309	two pore segment channel 2 (35.0 kD) (Tpcn2) alternative variant bSep08, mRNA.
Tpcn2	Tpcn2.cSep08	309139	2917	842	7	155	two pore segment channel 2 (Tpcn2) alternative variant cSep08, mRNA.
Tpcn2	Tpcn2.dSep08	309139	1242	880	2	52	two pore segment channel 2 (5.5 kD) (Tpcn2) alternative variant dSep08, mRNA.
Tpd52	Tpd52.bSep08	294900	80384	2356	8	208	tumor protein D52 (22.6 kD) (Tpd52) alternative variant bSep08, complete mRNA.
Tpd52	Tpd52.cSep08	294900	78978	881	6	190	tumor protein D52 (Tpd52) alternative variant cSep08, mRNA.
Tpd52	Tpd52.dSep08	294900	12932	432	5	111	tumor protein D52 (Tpd52) alternative variant dSep08, mRNA.
Tpd52l1	Tpd52l1.cSep08	689256	13134	788	4	99	tumor protein D52-like 1 (Tpd52l1) alternative variant cSep08, mRNA.
Tpd52l2	Tpd52l2.aSep08	296480	18391	978	7	245	tumor protein D52-like 2 (Tpd52l2) alternative variant aSep08, mRNA.
Tpd52l2	Tpd52l2.bSep08	296480	18332	919	9	235	tumor protein D52-like 2 (Tpd52l2) alternative variant bSep08, mRNA.
Tpd52l2	Tpd52l2.dSep08	296480	18233	760	6	216	tumor protein D52-like 2 (Tpd52l2) alternative variant dSep08, mRNA.
Tpd52l2	Tpd52l2.eSep08	296480	14821	577	6	191	tumor protein D52-like 2 (Tpd52l2) alternative variant eSep08, mRNA.
Tph1	Tph1.bSep08	24848	3315	382	3	127	tryptophan hydroxylase 1 (Tph1) alternative variant bSep08, mRNA.
Tph1	Tph1.dSep08	24848	493	344	2	28	tryptophan hydroxylase 1 (Tph1) alternative variant dSep08, mRNA.
Tph2	Tph2.bSep08	317675	13571	368	1	57	tryptophan hydroxylase 2 (Tph2) alternative variant bSep08, mRNA.
Tpi1	Tpi1.bSep08	24849	1909	728	2	177	triosephosphate isomerase 1 (Tpi1) alternative variant bSep08, mRNA.



Tpi1	Tpi1.cSep08	24849	2051	781	3	160	triosephosphate isomerase 1 (17.3 kD) (Tpi1) alternative variant cSep08, mRNA.
Tpk1	Tpk1.aSep08	680668	201500	2136		108	thiamine pyrophosphokinase (11.8 kD) (Tpk1) mRNA.
Tpm1	Tpm1.dSep08	24851	20917	931	10	284	tropomyosin (32.6 kD) (Tpm1) alternative variant dSep08, mRNA.
Tpm1	Tpm1.eSep08	24851	19495	992	9	283	tropomyosin (Tpm1) alternative variant eSep08, mRNA.
Tpm1	Tpm1.jSep08	24851	18711	744	8	206	tropomyosin (24.0 kD) (Tpm1) alternative variant jSep08, mRNA.
Tpm1	Tpm1.kSep08	24851	11173	698	4	152	tropomyosin (17.4 kD) (Tpm1) alternative variant kSep08, mRNA.
Tpm1	Tpm1.lSep08	24851	1209	438	2	127	tropomyosin 1 (Tpm1) alternative variant lSep08, mRNA.
Tpm1	Tpm1.mSep08	24851	2676	1564	2	110	tropomyosin (12.4 kD) (Tpm1) alternative variant mSep08, complete mRNA.
Tpm1	Tpm1.nSep08	24851	3418	1680	3	70	tropomyosin (Tpm1) alternative variant nSep08, mRNA.
Tpm2	Tpm2.aSep08	500450	9018	1992	9	284	tropomyosin 2 (Tpm2) alternative variant aSep08, mRNA.
Tpm2	Tpm2.bSep08	500450	8471	1031	8	235	tropomyosin 2 (27.2 kD) (Tpm2) alternative variant bSep08, mRNA.
Tpm2	Tpm2.cSep08	500450	7157	908	9	165	tropomyosin 2 (Tpm2) alternative variant cSep08, mRNA.
Tpm2	Tpm2.dSep08	500450	5645	2086	8	158	tropomyosin 2 (18.4 kD) (Tpm2) alternative variant dSep08, mRNA.
Tpm3	Tpm3.aSep08	117557	26546	1046	10	285	tropomyosin 3, gamma (33.2 kD) (Tpm3) alternative variant aSep08, mRNA.
Tpm3	Tpm3.bSep08	117557	27751	2172	9	285	tropomyosin 3, gamma (33.1 kD) (Tpm3) alternative variant bSep08, mRNA.
Tpm3	Tpm3.eSep08	117557	13235	631	8	172	tropomyosin 3, gamma (Tpm3) alternative variant eSep08, mRNA.
Tpm3	Tpm3.fSep08	117557	8034	572	4	152	tropomyosin 3, gamma (17.6 kD) (Tpm3) alternative variant fSep08, mRNA.
Tpm3	Tpm3.iSep08	117557	10533	633	5	101	tropomyosin 3, gamma (Tpm3) alternative variant iSep08, mRNA.
Tpm3	Tpm3.kSep08	117557	10295	432	3	59	tropomyosin 3, gamma (Tpm3) alternative variant kSep08, mRNA.
Tpm3	Tpm3.lSep08	117557	7984	418	2	10	tropomyosin 3, gamma (Tpm3) alternative variant lSep08, mRNA.
Tpm3	Tpm3.mSep08	117557	1219	367	2	44	tropomyosin 3, gamma (Tpm3) alternative variant mSep08, mRNA.
Tpm4	Tpm4.bSep08	24852	7614	957	4	151	tropomyosin 4 (Tpm4) alternative variant bSep08, mRNA.
Tpm4	Tpm4.cSep08	24852	5549	1379	3	101	tropomyosin 4 (Tpm4) alternative variant cSep08, mRNA.
Tpmt	Tpmt.aSep08	690050	16129	1778	1	546	thiopurine methyltransferase (Tpmt) alternative variant aSep08, mRNA.
Tpmt	Tpmt.bSep08	690050	18532	1068	6	240	thiopurine methyltransferase (27.7 kD) (Tpmt) alternative variant bSep08, mRNA.
Tpo1	Tpo1.bSep08	170907	78821	784	2	260	developmentally regulated protein TPO1 (Tpo1) alternative variant bSep08, mRNA.
Tpo1	Tpo1.cSep08	170907	80261	767	2	195	developmentally regulated protein TPO1 (Tpo1) alternative variant cSep08, mRNA.

Tpp1	Tpp1.bSep08	83534	824	596	2	83	tripeptidyl peptidase I (Tpp1) alternative variant bSep08, mRNA.
Tpp2	Tpp2.bSep08	81815	20659	1160	9	386	tripeptidyl peptidase II (Tpp2) alternative variant bSep08, mRNA.
Tpp2	Tpp2.cSep08	81815	14111	1065	5	208	tripeptidyl peptidase II (Tpp2) alternative variant cSep08, mRNA.
Tpp2	Tpp2.dSep08	81815	5553	843	2	55	tripeptidyl peptidase II (6.3 kD) (Tpp2) alternative variant dSep08, mRNA.
Tppp	Tppp.bSep08	361466	6174	959	2	152	tubulin polymerization promoting protein (Tppp) alternative variant bSep08, mRNA.
Tppp3	Tppp3.aSep08	291966	3474	775	1	207	tubulin polymerization-promoting protein family member 3 (Tppp3) alternative variant aSep08, mRNA.
Tpr	Tpr.bSep08	304862	10858	1784	7	594	translocated promoter region (Tpr) alternative variant bSep08, mRNA.
Tpr	Tpr.cSep08	304862	10374	868	6	289	translocated promoter region (Tpr) alternative variant cSep08, mRNA.
Tpr	Tpr.dSep08	304862	1394	413	2	101	translocated promoter region (Tpr) alternative variant dSep08, mRNA.
Tpr	Tpr.eSep08	304862	1603	454	3	11	translocated promoter region (Tpr) alternative variant eSep08, mRNA.
Tpr	Tpr.fSep08	304862	1283	179	2	7	translocated promoter region (Tpr) alternative variant fSep08, mRNA.
Tprkb	Tprkb.bSep08	297411	14894	2100	3	175	tp53rk binding protein (19.5 kD) (Tprkb) alternative variant bSep08, complete mRNA.
Tprkb	Tprkb.cSep08	297411	12540	600	3	155	tp53rk binding protein (Tprkb) alternative variant cSep08, mRNA.
Tprkb	Tprkb.dSep08	297411	13595	728	3	106	tp53rk binding protein (12.0 kD) (Tprkb) alternative variant dSep08, mRNA.
TPR_1.0	TPR_1.0.aSep08		68713	388		128	ubiquitously transcribed tetratricopeptide repeat (TPR_1.0) mRNA.
TPR_1.1	TPR_1.1.aSep08		9558	439		146	RAN binding protein 2 like (TPR_1.1) mRNA.
TPR_1.2	TPR_1.2.aSep08		10846	831	5	249	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (TPR_1.2) alternative variant aSep08, mRNA.
TPR_1.2	TPR_1.2.bSep08		2780	2021	3	101	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (11.8 kD) (TPR_1.2) alternative variant bSep08, mRNA.
TPR_1.5	TPR_1.5.aSep08		5791	902		300	tetratricopeptide TPR 1 and sel1-like and tetratricopeptide TPR 2 and tetratricopeptide TPR 3 (TPR_1.5) mRNA.
TPR_1.6	TPR_1.6.aSep08		4301	578		192	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (TPR_1.6) mRNA.
TPR_2.0	TPR_2.0.aSep08		9796	840		275	signal recognition particle (TPR_2.0) mRNA.
TPR_2.1	TPR_2.1.aSep08		105814	403		134	transmembrane tetratricopeptide repeat containing 2 (TPR_2.1) mRNA.
Tpst2	Tpst2.bSep08	288719	7998	1148	4	331	tyrosylprotein sulfotransferase 2 (Tpst2) alternative variant bSep08, mRNA.
Tpst2	Tpst2.cSep08	288719	2857	1452	2	91	tyrosylprotein sulfotransferase 2 (Tpst2) alternative variant cSep08, mRNA.

Tpx2	Tpx2.bSep08	311546	29051	815	7	230	TPX2, microtubule-associated protein homolog ( <i>Xenopus laevis</i> ) (Tpx2) alternative variant bSep08, mRNA.
Tpx2	Tpx2.cSep08	311546	7524	1045	6	198	TPX2, microtubule-associated protein homolog ( <i>Xenopus laevis</i> ) (Tpx2) alternative variant cSep08, mRNA.
Tpx2	Tpx2.dSep08	311546	22822	377	4	63	TPX2, microtubule-associated protein homolog ( <i>Xenopus laevis</i> ) (Tpx2) alternative variant dSep08, mRNA.
Tpx2	Tpx2.eSep08	311546	22759	362	4	49	TPX2, microtubule-associated protein homolog ( <i>Xenopus laevis</i> ) (Tpx2) alternative variant eSep08, mRNA.
Tra1	Tra1.aSep08	362862	10993	2315	14	667	heat shock protein 90kDa beta member 1 (Tra1) alternative variant aSep08, mRNA.
Tra1	Tra1.cSep08	362862	3252	801	5	179	shock protein beta member 1 (Tra1) alternative variant cSep08, mRNA.
Tra1	Tra1.dSep08	362862	4246	937	3	124	shock protein beta member 1 (Tra1) alternative variant dSep08, mRNA.
Tra1	Tra1.eSep08	362862	1977	883	4	116	shock protein beta member 1 (Tra1) alternative variant eSep08, mRNA.
Tra1	Tra1.fSep08	362862	795	705	2	100	shock protein (11.6 kD) (Tra1) alternative variant fSep08, mRNA.
Tra2a	Tra2a.aSep08	500116	18915	1859	8	282	transformer 2 alpha homolog ( <i>Drosophila</i> ) (32.6 kD) (Tra2a) alternative variant aSep08, mRNA.
Tra2a	Tra2a.dSep08	500116	3632	683	3	103	transformer 2 alpha homolog ( <i>Drosophila</i> ) (Tra2a) alternative variant dSep08, mRNA.
Tra2a	Tra2a.eSep08	500116	10873	328	3	72	transformer 2 alpha homolog ( <i>Drosophila</i> ) (Tra2a) alternative variant eSep08, mRNA.
Tra2a	Tra2a.gSep08	500116	2755	452	3	78	transformer 2 alpha homolog ( <i>Drosophila</i> ) (Tra2a) alternative variant gSep08, mRNA.
Trabd	Trabd.bSep08	300142	9376	1406	9	324	TraB determinant (Trabd) alternative variant bSep08, mRNA.
Trabd	Trabd.cSep08	300142	3798	722	6	174	TraB determinant (Trabd) alternative variant cSep08, mRNA.
Trabd	Trabd.dSep08	300142	4185	1045	7	153	putative protein of ancient origin (Trabd) alternative variant dSep08, mRNA.
Tradd	Tradd.aSep08	246756	2780	2556		208	TNFRSF1A-associated via death domain (23.2 kD) (Tradd) mRNA.
Traf2	Traf2.bSep08	311786	4387	766	5	255	tnf receptor-associated factor 2 (Traf2) alternative variant bSep08, mRNA.
Traf2	Traf2.cSep08	311786	5701	770	7	163	tnf receptor-associated factor 2 (Traf2) alternative variant cSep08, mRNA.
Traf3	Traf3.bSep08	362788	20393	617	3	107	tnf receptor-associated factor 3 (10.0 kD) (Traf3) alternative variant bSep08, mRNA.
Traf3	Traf3.cSep08	362788	995	295	2	98	tnf receptor-associated factor 3 (Traf3) alternative variant cSep08, mRNA.
Traf3	Traf3.dSep08	362788	3071	3012	2	108	tnf receptor-associated factor 3 (11.7 kD) (Traf3) alternative variant dSep08, mRNA.
Traf3ip1	Traf3ip1.bSep08	363286	2040	715	1	63	TNF receptor-associated factor 3 interacting protein 1 (7.1 kD) (Traf3ip1) alternative variant bSep08, mRNA.

Traf3ip2	Traf3ip2.bSep08	361857	31839	1550	1	157	traf3 interacting protein 2 (18.4 kD) (Traf3ip2) alternative variant bSep08, mRNA.
Traf3ip3	Traf3ip3.bSep08	360900	12962	724	7	241	TRAF3 interacting protein 3 (Traf3ip3) alternative variant bSep08, mRNA.
Traf3ip3	Traf3ip3.cSep08	360900	3193	694	3	230	TRAF3 interacting protein 3 (Traf3ip3) alternative variant cSep08, mRNA.
Traf3ip3	Traf3ip3.dSep08	360900	6142	1338	4	211	TRAF3 interacting protein 3 (24.0 kD) (Traf3ip3) alternative variant dSep08, mRNA.
Traf3ip3	Traf3ip3.eSep08	360900	5647	814	4	157	TRAF3 interacting protein 3 (17.8 kD) (Traf3ip3) alternative variant eSep08, mRNA.
Traf4	Traf4.bSep08	303285	4479	773	4	125	tnf receptor associated factor 4 (Traf4) alternative variant bSep08, mRNA.
Traf4af1	Traf4af1.bSep08	311325	16327	684	1	169	TRAF4 associated factor 1 (Traf4af1) alternative variant bSep08, mRNA.
Traf7	Traf7.bSep08	360491	14514	897	9	264	tnf receptor-associated factor 7 (Traf7) alternative variant bSep08, mRNA.
Traf7	Traf7.cSep08	360491	933	585	4	194	TNF receptor-associated factor 7 (Traf7) alternative variant cSep08, mRNA.
Traf7	Traf7.dSep08	360491	14207	696	8	177	tnf receptor-associated factor 7 (19.1 kD) (Traf7) alternative variant dSep08, mRNA.
Traf7	Traf7.eSep08	360491	2085	817	5	156	tnf receptor-associated factor 7 (Traf7) alternative variant eSep08, mRNA.
Traf7	Traf7.fSep08	360491	2038	1103	5	96	tnf receptor-associated factor 7 (10.5 kD) (Traf7) alternative variant fSep08, mRNA.
Trafd1	Trafd1.aSep08	114635	13424	1972	12	609	putative protein of eukaryotic origin (Trafd1) alternative variant aSep08, mRNA.
Trafd1	Trafd1.bSep08	114635	5906	2644	6	189	zinc finger (20.7 kD) (Trafd1) alternative variant bSep08, mRNA.
Trafd1	Trafd1.cSep08	114635	6974	491	5	113	putative protein of eukaryotic origin (Trafd1) alternative variant cSep08, mRNA.
Trafd1	Trafd1.dSep08	114635	3720	382	2	100	putative protein (Trafd1) alternative variant dSep08, mRNA.
Traip	Traip.bSep08	367167	2791	873	1	159	TRAF-interacting protein (Traip) alternative variant bSep08, mRNA.
Trak2	Trak2.bSep08	171086	9513	873	4	177	trafficking protein, kinesin binding 2 (Trak2) alternative variant bSep08, mRNA.
Trak2	Trak2.cSep08	171086	1790	799	2	148	trafficking protein, kinesin binding 2 (Trak2) alternative variant cSep08, mRNA.
TRAM.0	TRAM.0.aSep08		157536	1223		184	CDK5 regulatory associated protein 1-like 1 (TRAM.0) mRNA.
Tram1	Tram1.bSep08	312903	20168	1126		242	translocating chain-associating membrane protein 1 (Tram1) alternative variant bSep08, mRNA.
TRAM_LAG1_CLN8.0	TRAM_LAG1_CLN8.0.aSep08		4993	2622	3	168	CRA b (TRAM_LAG1_CLN8.0) mRNA.
TRAM_LAG1_CLN8.1	TRAM_LAG1_CLN8.1.aSep08		11694	988		260	membrane protein 2 (TRAM_LAG1_CLN8.1) mRNA.
Trappc1	Trappc1.aSep08	287427	1685	740	4	145	trafficking protein particle complex 1 (16.9 kD) (Trappc1) alternative variant aSep08, complete mRNA.

Trappc1	Trappc1.dSep08	287427	1603	448	2	75	trafficking protein particle complex 1 (8.8 kD) (Trappc1) alternative variant dSep08, complete mRNA.
Trappc1	Trappc1.eSep08	287427	1713	1153	3	62	trafficking protein particle complex 1 (7.0 kD) (Trappc1) alternative variant eSep08, complete mRNA.
Trappc2	Trappc2.aSep08	501550	6462	809	6	146	trafficking protein particle complex 2 (17.4 kD) (Trappc2) alternative variant aSep08, mRNA.
Trappc2	Trappc2.bSep08	501550	11273	1548	6	140	trafficking protein particle complex 2 (16.4 kD) (Trappc2) alternative variant bSep08, mRNA.
Trappc2l	Trappc2l.bSep08	292074	2429	621	4	109	trafficking protein particle complex 2-like (12.5 kD) (Trappc2l) alternative variant bSep08, mRNA.
Trappc2l	Trappc2l.cSep08	292074	1325	521	1	54	trafficking protein particle complex 2-like (6.4 kD) (Trappc2l) alternative variant cSep08, mRNA.
Trappc4	Trappc4.bSep08	367073	3635	682	3	196	trafficking protein particle complex 4 (Trappc4) alternative variant bSep08, mRNA.
Trappc4	Trappc4.cSep08	367073	3341	557	2	154	trafficking protein particle complex 4 (Trappc4) alternative variant cSep08, mRNA.
Trappc4	Trappc4.dSep08	367073	2720	614	2	151	trafficking protein particle complex 4 (16.8 kD) (Trappc4) alternative variant dSep08, mRNA.
Trappc5	Trappc5.bSep08	363858	2191	532	2	113	trafficking protein particle complex 5 (12.5 kD) (Trappc5) alternative variant bSep08, mRNA.
Trappc6a	Trappc6a.bSep08	680465	6890	625	5	126	trafficking protein particle complex 6A (Trappc6a) alternative variant bSep08, mRNA.
Trdmt1	Trdmt1.bSep08	291324	4568	633	4	210	tRNA aspartic acid methyltransferase 1 (Trdmt1) alternative variant bSep08, mRNA.
Trdmt1	Trdmt1.cSep08	291324	15438	758	10	149	tRNA aspartic acid methyltransferase 1 (Trdmt1) alternative variant cSep08, mRNA.
Trdn	Trdn.bSep08	59299	128953	1225	10	229	triadin (26.3 kD) (Trdn) alternative variant bSep08, mRNA.
Trdn	Trdn.cSep08	59299	46085	916	4	58	triadin (Trdn) alternative variant cSep08, mRNA.
Trdn	Trdn.dSep08	59299	34348	883	3	47	triadin (Trdn) alternative variant dSep08, mRNA.
Treh	Treh.aSep08	60576	1355	652		139	trehalase (brush-border membrane glycoprotein) (Treh) mRNA.
Trem2	Trem2.aSep08	301227	6464	881	3	253	triggering receptor expressed on myeloid cells 2 (Trem2) alternative variant aSep08, mRNA.
Trem2	Trem2.bSep08	301227	4307	1012	2	207	triggering receptor expressed on myeloid cells 2 (22.5 kD) (Trem2) alternative variant bSep08, mRNA.
Treml1	Treml1.aSep08	501096	34585	1220		318	triggering receptor expressed on myeloid cells-like 1 (Treml1) mRNA.
Treml2	Treml2.aSep08	680844	1783	732		87	triggering receptor expressed on myeloid cells-like 2 (Treml2) alternative variant aSep08, mRNA.
Trerf1	Trerf1.bSep08	316219	33199	1909	9	435	transcriptional regulating factor 1 (48.1 kD) (Trerf1) alternative variant bSep08, mRNA.
Trerf1	Trerf1.cSep08	316219	6572	654	4	218	transcriptional regulating factor 1 (Trerf1) alternative variant cSep08, mRNA.
Trf	Trf.bSep08	24825	11255	1083	6	235	transferrin (Trf) alternative variant bSep08, mRNA.
Trf	Trf.cSep08	24825	8301	759	6	232	transferrin precursor (24.7 kD) (Trf) alternative variant cSep08, mRNA.

Trf	Trf.dSep08	24825	2309	410	3	128	transferrin (Trf) alternative variant dSep08, mRNA.
Trfr2	Trfr2.bSep08	288562	16893	3015	19	798	transferrin receptor 2 (88.3 kD) (Trfr2) alternative variant bSep08, complete mRNA.
Trfr2	Trfr2.cSep08	288562	3690	719	2	101	transferrin receptor 2 (11.3 kD) (Trfr2) alternative variant cSep08, mRNA.
Trfr2	Trfr2.dSep08	288562	966	497	6	41	transferrin receptor 2 (Trfr2) alternative variant dSep08, mRNA.
Trib2	Trib2.cSep08	313974	47696	502	2	70	tribbles homolog 2 (Drosophila) (Trib2) alternative variant cSep08, mRNA.
Trim2	Trim2.bSep08	361970	9494	546	4	138	tripartite motif protein 2 (Trim2) alternative variant bSep08, mRNA.
Trim2	Trim2.cSep08	361970	66168	336	3	95	tripartite motif protein 2 (Trim2) alternative variant cSep08, mRNA.
Trim2	Trim2.dSep08	361970	48013	277	2	91	tripartite motif protein 2 (Trim2) alternative variant dSep08, mRNA.
Trim2	Trim2.eSep08	361970	117171	376	2	80	tripartite motif protein 2 (Trim2) alternative variant eSep08, mRNA.
Trim3	Trim3.bSep08	83616	16564	631	5	180	tripartite motif-containing 3 (Trim3) alternative variant bSep08, mRNA.
Trim3	Trim3.cSep08	83616	16115	524	3	157	putative protein (Trim3) alternative variant cSep08, mRNA.
Trim3	Trim3.dSep08	83616	1326	922	3	100	tripartite motif-containing 3 CRA e (10.9 kD) (Trim3) alternative variant dSep08, mRNA.
Trim6	Trim6.aSep08	293294	4499	1738	4	212	tripartite motif-containing 6 (24.4 kD) (Trim6) alternative variant aSep08, mRNA.
Trim8	Trim8.bSep08	688785	2982	1715	4	129	tripartite motif-containing 8 and hypothetical protein LOC689889 (14.8 kD) (Trim8) alternative variant bSep08, mRNA.
Trim8	Trim8.bSep08	689889	2982	1715	4	129	tripartite motif-containing 8 and hypothetical protein LOC689889 (14.8 kD) (Trim8) alternative variant bSep08, mRNA.
Trim8	Trim8.dSep08	688785	383	250	2	44	tripartite motif-containing 8 and hypothetical protein LOC689889 (4.9 kD) (Trim8) alternative variant dSep08, mRNA.
Trim8	Trim8.dSep08	689889	383	250	2	44	tripartite motif-containing 8 and hypothetical protein LOC689889 (4.9 kD) (Trim8) alternative variant dSep08, mRNA.
Trim9	Trim9.cSep08	155812	1989	463	2	43	tripartite motif-containing 9 (Trim9) alternative variant cSep08, mRNA.
Trim10	Trim10.bSep08	294210	6033	702	1	234	tripartite motif protein 10 (Trim10) alternative variant bSep08, mRNA.
Trim11	Trim11.bSep08	360534	13035	3472	5	152	tripartite motif-containing 11 (17.2 kD) (Trim11) alternative variant bSep08, mRNA.
Trim11	Trim11.cSep08	360534	648	560	2	91	tripartite motif-containing 11 (Trim11) alternative variant cSep08, mRNA.
Trim13andKcnrg	Trim13andKcnrg.bSep08	305947	15418	667	1	115	potassium channel regulator (Trim13andKcnrg) alternative variant bSep08, mRNA.

Trim13andKcnrg	Trim13andKcnrg.bSep08	364398	15418	667	1	115	potassium channel regulator (Trim13andKcnrg) alternative variant bSep08, mRNA.
Trim14	Trim14.aSep08	313236	25337	3087	3	440	tripartite motif protein 14 (49.8 kD) (Trim14) alternative variant aSep08, mRNA.
Trim16	Trim16.aSep08	303214	9074	641		213	tripartite motif protein 16 (Trim16) mRNA.
Trim17	Trim17.aSep08	64702	4931	760		234	tripartite motif-containing 17 (Trim17) mRNA.
Trim23	Trim23.aSep08	81002	13644	863	3	258	tripartite motif-containing 23 (Trim23) alternative variant aSep08, mRNA.
Trim23	Trim23.bSep08	81002	10598	604	1	155	tripartite motif-containing 23 (16.8 kD) (Trim23) alternative variant bSep08, mRNA.
Trim24	Trim24.bSep08	500084	24843	2431	12	612	tripartite motif-containing 24 (Trim24) alternative variant bSep08, mRNA.
Trim24	Trim24.cSep08	500084	7227	725	5	109	tripartite motif-containing 24 (Trim24) alternative variant cSep08, mRNA.
Trim24	Trim24.dSep08	500084	3040	801	2	52	tripartite motif-containing 24 (6.0 kD) (Trim24) alternative variant dSep08, mRNA.
Trim25	Trim25.aSep08	494338	28533	1863		492	tripartite motif-containing 25 (Trim25) mRNA.
Trim26	Trim26.bSep08	309586	3319	2316	4	284	tripartite motif-containing 26 (Trim26) alternative variant bSep08, mRNA.
Trim27	Trim27.aSep08	291171	12355	2081	6	611	ret finger protein (Trim27) alternative variant aSep08, mRNA.
Trim27	Trim27.cSep08	291171	3382	1383	2	216	ret finger protein (24.1 kD) (Trim27) alternative variant cSep08, mRNA.
Trim28	Trim28.aSep08	116698	6691	2989	17	835	tripartite motif-containing 28 precursor (89.0 kD) (Trim28) alternative variant aSep08, mRNA.
Trim28	Trim28.bSep08	116698	1835	1159	8	259	CRA a (Trim28) alternative variant bSep08, mRNA.
Trim28	Trim28.cSep08	116698	534	429	2	142	CRA a (Trim28) alternative variant cSep08, mRNA.
Trim28	Trim28.dSep08	116698	1430	1261	3	130	tripartite motif-containing 28 (14.0 kD) (Trim28) alternative variant dSep08, mRNA.
Trim28	Trim28.eSep08	116698	872	764	2	98	tripartite motif-containing 28 (Trim28) alternative variant eSep08, mRNA.
Trim29	Trim29.bSep08	300656	8839	1785	1	503	tripartite motif protein 29 (Trim29) alternative variant bSep08, mRNA.
Trim32	Trim32.aSep08	313264	10903	3212	2	655	tripartite motif protein 32 (72.2 kD) (Trim32) alternative variant aSep08, mRNA.
Trim32	Trim32.cSep08	313264	14018	719	2	166	tripartite motif protein 32 (Trim32) alternative variant cSep08, mRNA.
Trim33	Trim33.aSep08	365894	26069	3396	10	582	tripartite motif-containing 33 (Trim33) alternative variant aSep08, mRNA.
Trim33	Trim33.bSep08	365894	2100	577		121	tripartite motif-containing 33 (Trim33) alternative variant bSep08, mRNA.
Trim35	Trim35.bSep08	498538	1129	421	3	100	tripartite motif-containing 35 (Trim35) alternative variant bSep08, mRNA.
Trim36	Trim36.bSep08	291597	3962	2567	2	240	tripartite motif-containing 36 (Trim36) alternative variant bSep08, mRNA.

Trim37	Trim37.bSep08	360592	132133	3380	24	967	tripartite motif-containing 37 (108.4 kD) (Trim37) alternative variant bSep08, mRNA.
Trim37	Trim37.cSep08	360592	20932	1591	4	141	tripartite motif-containing 37 (Trim37) alternative variant cSep08, mRNA.
Trim37	Trim37.dSep08	360592	31508	373	2	78	tripartite motif-containing 37 (Trim37) alternative variant dSep08, mRNA.
Trim37	Trim37.fSep08	360592	15081	270	2	14	tripartite motif-containing 37 (Trim37) alternative variant fSep08, mRNA.
Trim39	Trim39.bSep08	309591	5300	893	1	120	tripartite motif-containing 39 (Trim39) alternative variant bSep08, mRNA.
Trim44	Trim44.bSep08	362172	55649	1186	4	250	tripartite motif-containing 44 (Trim44) alternative variant bSep08, mRNA.
Trim46	Trim46.aSep08	310641	10455	1907	5	626	tripartite motif-containing 46 (Trim46) alternative variant aSep08, mRNA.
Trim46	Trim46.bSep08	310641	5620	647	5	207	tripartite motif-containing 46 (Trim46) alternative variant bSep08, mRNA.
Trim46	Trim46.cSep08	310641	1970	729	1	173	tripartite motif-containing 46 (Trim46) alternative variant cSep08, mRNA.
Trim47l	Trim47l.aSep08	287824	8022	1947		623	tripartite motif protein 47-like (Trim47l) complete mRNA.
Trim50	Trim50.aSep08	288596	12321	780		207	tripartite motif-containing 50 (Trim50) mRNA.
Trim54	Trim54.aSep08	362708	5803	1443	6	436	tripartite motif-containing 54 (Trim54) alternative variant aSep08, mRNA.
Trim54	Trim54.cSep08	362708	13879	697	3	179	tripartite motif-containing 54 (Trim54) alternative variant cSep08, mRNA.
Trim55	Trim55.bSep08	365751	19360	704	3	131	tripartite motif-containing 55 (Trim55) alternative variant bSep08, mRNA.
Trim62	Trim62.aSep08	313045	6767	859		218	tripartite motif-containing 62 (Trim62) mRNA.
Trim66	Trim66.aSep08	361623	56784	2413		804	tripartite motif-containing 66 (Trim66) alternative variant aSep08, mRNA.
Trim66	Trim66.bSep08	361623	10629	1191		350	tripartite motif-containing 66 (Trim66) alternative variant bSep08, mRNA.
Trim66	Trim66.cSep08	361623	2525	520		173	tripartite motif-containing 66 (Trim66) alternative variant cSep08, mRNA.
Trim69	Trim69.aSep08	311373	14621	820	2	215	tripartite motif-containing 69 (Trim69) alternative variant aSep08, mRNA.
Trim69	Trim69.bSep08	311373	11738	802	2	163	tripartite motif-containing 69 (18.8 kD) (Trim69) alternative variant bSep08, mRNA.
Trio	Trio.bSep08	310192	14198	1839	10	612	triple functional domain (PTPRF interacting) (Trio) alternative variant bSep08, mRNA.
Trio	Trio.cSep08	310192	18213	1783	3	545	triple functional domain (PTPRF interacting) (Trio) alternative variant cSep08, mRNA.
Trio	Trio.dSep08	310192	5705	431	3	125	triple functional domain (PTPRF interacting) (Trio) alternative variant dSep08, mRNA.
Triobp	Triobp.aSep08	362956	24291	2540		621	TRIO and F-actin binding protein (Triobp) alternative variant aSep08, mRNA.
Triobp	Triobp.bSep08	362956	8743	1776		33	TRIO and F-actin binding protein (Triobp) alternative variant bSep08, mRNA.



Trip10	Trip10.bSep08	116717	5256	783	1	243	thyroid hormone receptor interactor 10 (Trip10) alternative variant bSep08, mRNA.
Trip10	Trip10.cSep08	116717	5995	928	5	220	thyroid hormone receptor interactor 10 (Trip10) alternative variant cSep08, mRNA.
Trip11	Trip11.aSep08	314393	11111	722		137	thyroid hormone receptor interactor 11 (Trip11) mRNA.
Trip12	Trip12.aSep08	316575	38152	6905	24	1173	thyroid hormone receptor interactor 12 (Trip12) alternative variant aSep08, mRNA.
Trip12	Trip12.bSep08	316575	18535	1557	10	474	thyroid hormone receptor interactor 12 (Trip12) alternative variant bSep08, mRNA.
Trip12	Trip12.cSep08	316575	10733	1226	10	408	thyroid hormone receptor interactor 12 (Trip12) alternative variant cSep08, mRNA.
Trip13	Trip13.bSep08	292206	5502	1315	2	94	thyroid hormone receptor interactor 13 (10.7 kD) (Trip13) alternative variant bSep08, mRNA.
Trit1	Trit1.bSep08	362586	9581	408	5	135	tRNA isopentenyltransferase 1 (Trit1) alternative variant bSep08, mRNA.
Trmt1	Trmt1.aSep08	288914	10180	1941	16	470	trm1 tRNA methyltransferase 1 homolog CRA b (51.6 kD) (Trmt1) alternative variant aSep08, complete mRNA.
Trmt1	Trmt1.cSep08	288914	5388	762	6	244	trm1 tRNA methyltransferase 1 homolog CRA b (Trmt1) alternative variant cSep08, mRNA.
Trmt1	Trmt1.dSep08	288914	1824	710	6	204	trm1 tRNA methyltransferase 1 homolog (22.4 kD) (Trmt1) alternative variant dSep08, mRNA.
Trmt1	Trmt1.eSep08	288914	3729	493	2	135	trm1 tRNA methyltransferase 1 homolog CRA c (Trmt1) alternative variant eSep08, mRNA.
Trmt1	Trmt1.fSep08	288914	449	377	2	125	trm1 tRNA methyltransferase 1 homolog CRA b (Trmt1) alternative variant fSep08, mRNA.
Trmt1	Trmt1.gSep08	288914	543	427	2	110	trm1 tRNA methyltransferase 1 homolog (11.9 kD) (Trmt1) alternative variant gSep08, mRNA.
Trmt5	Trmt5.bSep08	362754	6037	1436	3	478	TRM5 tRNA methyltransferase 5 homolog (S. cerevisiae) (Trmt5) alternative variant bSep08, mRNA.
Trmt5	Trmt5.cSep08	362754	7034	1307	2	258	TRM5 tRNA methyltransferase 5 homolog (S. cerevisiae) (Trmt5) alternative variant cSep08, mRNA.
Trmt5	Trmt5.dSep08	362754	4401	882	3	189	TRM5 tRNA methyltransferase 5 homolog (S. cerevisiae) (Trmt5) alternative variant dSep08, mRNA.
Trmt11	Trmt11.bSep08	378794	50805	1779	3	157	tRNA methyltransferase 11 homolog (S. cerevisiae) (Trmt11) alternative variant bSep08, mRNA.
Trmu	Trmu.aSep08	362976	16526	1623	10	459	trna methyltransferase CRA a (Trmu) alternative variant aSep08, mRNA.
Trmu	Trmu.bSep08	362976	12174	768	8	255	tRNA -methyltransferase (Trmu) alternative variant bSep08, mRNA.
Trmu	Trmu.cSep08	362976	12884	659	6	170	tRNA -methyltransferase (Trmu) alternative variant cSep08, mRNA.
Trmu	Trmu.dSep08	362976	1670	706	2	90	trna methyltransferase CRA a (Trmu) alternative variant dSep08, mRNA.
Trnt1	Trnt1.bSep08	312616	7247	612	5	179	tRNA nucleotidyl transferase, CCA-adding, 1 (Trnt1) alternative variant bSep08, mRNA.
Troap	Troap.aSep08	300219	4842	743	6	247	trophinin associated protein (Troap) alternative variant aSep08, mRNA.

Troap	Troap.bSep08	300219	2018	766	2	159	trophinin associated protein (Troap) alternative variant bSep08, mRNA.
Trp53	Trp53.bSep08	24842	10162	2363	8	259	transformation related protein 53 (28.6 kD) (Trp53) alternative variant bSep08, mRNA.
Trp53	Trp53.cSep08	24842	1756	711	4	199	transformation related protein 53 (Trp53) alternative variant cSep08, mRNA.
Trp53	Trp53.dSep08	24842	1918	1154	3	177	transformation related protein 53 (Trp53) alternative variant dSep08, mRNA.
Trp53	Trp53.eSep08	24842	8224	823	3	89	transformation related protein 53 (Trp53) alternative variant eSep08, mRNA.
Trp53	Trp53.fSep08	24842	8045	1233	3	91	transformation related protein 53 (9.5 kD) (Trp53) alternative variant fSep08, mRNA.
Trp53	Trp53.gSep08	24842	1329	793	2	70	transformation related protein 53 (7.4 kD) (Trp53) alternative variant gSep08, complete mRNA.
Trp53bp1	Trp53bp1.bSep08	296099	3573	825	5	217	transformation related protein 53 binding protein 1 (Trp53bp1) alternative variant bSep08, mRNA.
Trp53bp1	Trp53bp1.cSep08	296099	6660	618	5	171	transformation related protein 53 binding protein 1 (Trp53bp1) alternative variant cSep08, mRNA.
Trp53bp1	Trp53bp1.dSep08	296099	1201	370	3	93	transformation related protein 53 binding protein 1 (Trp53bp1) alternative variant dSep08, mRNA.
Trp53bp2	Trp53bp2.aSep08	305025	16757	2116		465	transformation related protein 53 binding protein 2 (Trp53bp2) mRNA.
Trp53i11	Trp53i11.aSep08	311209	15159	2805	5	204	transformation related protein 53 inducible protein 11 (22.5 kD) (Trp53i11) alternative variant aSep08, complete mRNA.
Trp53i11	Trp53i11.bSep08	311209	11108	367	2	75	transformation related protein 53 inducible protein 11 (Trp53i11) alternative variant bSep08, mRNA.
Trp53i11	Trp53i11.cSep08	311209	10957	421	1	61	transformation related protein 53 inducible protein 11 (Trp53i11) alternative variant cSep08, mRNA.
Trp53i13	Trp53i13.bSep08	287550	4835	713	7	215	tumor protein p53 inducible 13 CRA a like (Trp53i13) alternative variant bSep08, mRNA.
Trp53i13	Trp53i13.cSep08	287550	3938	747	5	203	tumor protein p53 inducible 13 CRA a like (Trp53i13) alternative variant cSep08, mRNA.
Trp53i13	Trp53i13.dSep08	287550	4782	683	5	84	tumor protein p53 inducible 13 CRA a like (Trp53i13) alternative variant dSep08, mRNA.
Trp53i13	Trp53i13.eSep08	287550	3458	600	4	72	tumor protein p53 inducible 13 CRA a like (Trp53i13) alternative variant eSep08, mRNA.
Trp53i13	Trp53i13.fSep08	287550	1001	808	2	51	transformation related protein 53 inducible 13 like (5.3 kD) (Trp53i13) alternative variant fSep08, mRNA.
Trp53inp1	Trp53inp1.bSep08	297822	3069	485	2	123	transformation related protein 53 inducible nuclear protein 1 (Trp53inp1) alternative variant bSep08, mRNA.
Trp53inp2	Trp53inp2.aSep08	362246	8658	4559		221	tumor protein p53 inducible nuclear protein 2 (24.4 kD) (Trp53inp2) mRNA.
Trpc1	Trpc1.bSep08	89821	16441	1798	6	360	transient receptor potential cation channel, subfamily C, member 1 (42.1 kD) (Trpc1) alternative variant bSep08, mRNA.

Trpc2	Trpc2.aSep08	64573	21413	3087	10	407	transient receptor potential cation channel, subfamily C, member 2 (44.4 kD) (Trpc2) alternative variant aSep08, mRNA.
Trpc2	Trpc2.bSep08	64573	5309	470	2	20	transient receptor potential cation channel, subfamily C, member 2 (2.2 kD) (Trpc2) alternative variant bSep08, mRNA.
Trpc3	Trpc3.aSep08	60395	19255	1264	4	170	transient receptor potential cation channel, subfamily C, member 3 (Trpc3) alternative variant aSep08, mRNA.
Trpc3	Trpc3.bSep08	60395	15254	790	3	70	transient receptor potential cation channel, subfamily C, member 3 (8.2 kD) (Trpc3) alternative variant bSep08, mRNA.
Trpc4ap	Trpc4ap.bSep08	362247	25537	855	6	285	transient receptor potential cation channel, subfamily C, member 4 associated protein (Trpc4ap) alternative variant bSep08, mRNA.
Trpc7	Trpc7.aSep08	282822	69644	702		233	transient receptor potential cation channel, subfamily C, member 7 (Trpc7) mRNA.
Trpd52l3	Trpd52l3.bSep08	293894	1700	297	2	71	tumor protein D52-like 3 (Trpd52l3) alternative variant bSep08, mRNA.
Trpm1	Trpm1.cSep08	361586	21549	942	5	233	transient receptor potential cation channel subfamily M member 1 (Trpm1) alternative variant cSep08, mRNA.
Trpm1	Trpm1.dSep08	361586	7560	626	3	161	transient receptor potential cation channel subfamily M member 1 (Trpm1) alternative variant dSep08, mRNA.
Trpm2	Trpm2.bSep08	294329	4447	740	6	246	transient receptor potential cation channel, subfamily M, member 2 (Trpm2) alternative variant bSep08, mRNA.
Trpm2	Trpm2.cSep08	294329	7061	746	7	115	transient receptor potential cation channel, subfamily M, member 2 (Trpm2) alternative variant cSep08, mRNA.
Trpm2	Trpm2.dSep08	294329	2953	869	3	113	transient receptor potential cation channel, subfamily M, member 2 (Trpm2) alternative variant dSep08, mRNA.
Trpm3	Trpm3.aSep08	309407	72931	1182	2	394	transient receptor potential cation channel, subfamily M, member 3 (Trpm3) alternative variant aSep08, mRNA.
Trpm3	Trpm3.bSep08	309407	18375	664	1	221	transient receptor potential cation channel, subfamily M, member 3 (Trpm3) alternative variant bSep08, mRNA.
Trpm4	Trpm4.aSep08	171143	8588	994		331	transient receptor potential cation channel, subfamily M, member 4 (Trpm4) mRNA.
Trpm5	Trpm5.aSep08	365391	5900	1755	4	219	transient receptor potential cation channel, subfamily M, member 5 (Trpm5) alternative variant aSep08, mRNA.
Trpm5	Trpm5.bSep08	365391	3326	330	2	110	transient receptor potential cation channel, subfamily M, member 5 (Trpm5) alternative variant bSep08, mRNA.
Trpm6	Trpm6.aSep08	293874	25514	1043	10	347	transient receptor potential cation channel, subfamily M, member 6 (Trpm6) alternative variant aSep08, mRNA.
Trpm6	Trpm6.bSep08	293874	4079	878	1	76	transient receptor potential cation channel, subfamily M, member 6 (8.7 kD) (Trpm6) alternative variant bSep08, mRNA.
Trpm7	Trpm7.aSep08	679906	14866	2439	12	386	transient receptor potential cation channel, subfamily M, member 7 (Trpm7) alternative variant aSep08, mRNA.
Trpm7	Trpm7.bSep08	679906	8494	385	8	71	transient receptor potential cation channel, subfamily M, member 7 (Trpm7) alternative variant bSep08, mRNA.

Trps1	Trps1.aSep08	299897	10428	1903		420	trichorhinophalangeal syndrome I (human) (Trps1) mRNA.
Trpv1	Trpv1.bSep08	83810	5126	461	1	36	transient receptor potential cation channel, subfamily V, member 1 (Trpv1) alternative variant bSep08, mRNA.
Trpv2	Trpv2.cSep08	29465	2798	398	2	84	putative protein (8.9 kD) (Trpv2) alternative variant cSep08, mRNA.
Trpv2	Trpv2.dSep08	29465	2855	394	2	40	putative protein (Trpv2) alternative variant dSep08, mRNA.
Trpv6	Trpv6.bSep08	114246	1477	411	2	83	transient receptor potential cation channel subfamily V member 6 CRA c (Trpv6) alternative variant bSep08, mRNA.
Trpv6	Trpv6.cSep08	114246	2037	397	4	72	putative protein (Trpv6) alternative variant cSep08, mRNA.
TRP_2.0	TRP_2.0.aSep08		20098	806		268	transient receptor potential cation channel subfamily C member 3 (TRP_2.0) mRNA.
Trrap	Trrap.bSep08	288471	5289	729	5	242	transformation transcription domain-associated protein CRA b (Trrap) alternative variant bSep08, mRNA.
Trrap	Trrap.cSep08	288471	3655	2240	3	238	transformation transcription domain-associated protein CRA b (26.7 kD) (Trrap) alternative variant cSep08, mRNA.
Trrap	Trrap.dSep08	288471	2200	396	3	131	transformation transcription domain-associated protein CRA b (Trrap) alternative variant dSep08, mRNA.
Trrap	Trrap.eSep08	288471	781	529	2	99	transformation transcription domain-associated protein CRA e (Trrap) alternative variant eSep08, mRNA.
Trrap	Trrap.fSep08	288471	2282	402	3	93	transformation transcription domain-associated protein CRA d (Trrap) alternative variant fSep08, mRNA.
Trub1	Trub1.aSep08	361775	46447	1796	7	545	TruB pseudouridine (psi) synthase homolog 1 (E. coli) (Trub1) alternative variant aSep08, complete mRNA.
Trub1	Trub1.cSep08	361775	43101	732	6	239	TruB pseudouridine (psi) synthase homolog 1 (E. coli) (25.0 kD) (Trub1) alternative variant cSep08, mRNA.
Trub1	Trub1.dSep08	361775	46378	1054	7	223	TruB pseudouridine (psi) synthase homolog 1 (E. coli) (25.3 kD) (Trub1) alternative variant dSep08, mRNA.
Trub1	Trub1.eSep08	361775	3416	1782	3	119	TruB pseudouridine (psi) synthase homolog 1 (E. coli) (Trub1) alternative variant eSep08, mRNA.
Trypsin.0	Trypsin.0.aSep08		1298	705		163	mast cell protease (Trypsin.0) mRNA.
Trypsin.1	Trypsin.1.aSep08		1397	655		180	granzyme B (Trypsin.1) mRNA.
Trypsin.3	Trypsin.3.aSep08		8030	1814		426	regeneration associated muscle protease (Trypsin.3) mRNA.
Trypsin.4	Trypsin.4.aSep08		1119	660		200	kallikrein (Trypsin.4) mRNA.
Tsc2	Tsc2.bSep08	24855	4450	724	7	240	tuberous sclerosis 2 (Tsc2) alternative variant bSep08, mRNA.
Tsc2	Tsc2.cSep08	24855	4656	2134	10	179	tuberous sclerosis 2 (20.2 kD) (Tsc2) alternative variant cSep08, mRNA.
Tsc2	Tsc2.dSep08	24855	924	730	2	89	tuberous sclerosis 2 (10.3 kD) (Tsc2) alternative variant dSep08, mRNA.
Tsc22d1	Tsc22d1.cSep08	498545	915	806	2	96	TSC22 domain family, member 1 (10.5 kD) (Tsc22d1) alternative variant cSep08, mRNA.
Tsc22d1	Tsc22d1.dSep08	498545	29016	877	3	86	TSC22 domain family, member 1 (9.4 kD) (Tsc22d1) alternative variant dSep08, mRNA.

Tsc22d1	Tsc22d1.eSep08	498545	445	246	2	82	TSC22 domain family, member 1 (Tsc22d1) alternative variant eSep08, mRNA.
Tsc22d2	Tsc22d2.aSep08	499624	22509	759		145	TSC22 domain family, member 2 (16.2 kD) (Tsc22d2) mRNA.
Tsc22d3	Tsc22d3.aSep08	83514	58159	1139	4	254	TSC22 domain family 3 (Tsc22d3) alternative variant aSep08, mRNA.
Tsc22d3	Tsc22d3.bSep08	83514	58736	762	4	194	TSC22 domain family 3 (Tsc22d3) alternative variant bSep08, mRNA.
Tsc22d3	Tsc22d3.dSep08	83514	2092	1105	2	108	TSC22 domain family 3 (12.0 kD) (Tsc22d3) alternative variant dSep08, mRNA.
Tsc22d3	Tsc22d3.eSep08	83514	2247	898	3	77	TSC22 domain family 3 (8.7 kD) (Tsc22d3) alternative variant eSep08, mRNA.
Tsc22d4	Tsc22d4.bSep08	684980	14643	829	3	260	tsc22 domain family 4 (Tsc22d4) alternative variant bSep08, mRNA.
Tsc22d4	Tsc22d4.cSep08	684980	1570	1109	3	206	CRA c (Tsc22d4) alternative variant cSep08, mRNA.
Tsc22d4	Tsc22d4.dSep08	684980	6181	1055	3	200	uncharacterized protein homolog (23.5 kD) (Tsc22d4) alternative variant dSep08, mRNA.
Tsc22d4	Tsc22d4.hSep08	684980	756	633	2	67	CRA c like (Tsc22d4) alternative variant hSep08, mRNA.
Tsen2	Tsen2.bSep08	312649	8994	709	2	107	tRNA splicing endonuclease 2 homolog (SEN2, <i>S. cerevisiae</i> ) (Tsen2) alternative variant bSep08, mRNA.
Tsen34	Tsen34.bSep08	292534	5991	1797	4	270	tRNA splicing endonuclease 34 homolog (SEN34, <i>S. cerevisiae</i> ) (Tsen34) alternative variant bSep08, mRNA.
Tsen34	Tsen34.cSep08	292534	1814	797	4	243	tRNA splicing endonuclease 34 homolog (SEN34, <i>S. cerevisiae</i> ) (Tsen34) alternative variant cSep08, mRNA.
Tsen34	Tsen34.dSep08	292534	3816	673	2	108	tRNA splicing endonuclease 34 homolog (SEN34, <i>S. cerevisiae</i> ) (12.0 kD) (Tsen34) alternative variant dSep08, mRNA.
Tsen54	Tsen54.bSep08	690308	7510	2068	12	159	tRNA splicing endonuclease 54 homolog ( <i>S. cerevisiae</i> ) (18.1 kD) (Tsen54) alternative variant bSep08, complete mRNA.
Tsen54	Tsen54.cSep08	690308	3909	1783	7	167	tRNA splicing endonuclease 54 homolog ( <i>S. cerevisiae</i> ) (Tsen54) alternative variant cSep08, mRNA.
Tsg101	Tsg101.aSep08	292925	15646	1610	3	276	tumor susceptibility gene 101 (Tsg101) alternative variant aSep08, mRNA.
Tsg101	Tsg101.bSep08	292925	17845	1040	4	236	tumor susceptibility gene 101 (Tsg101) alternative variant bSep08, mRNA.
Tsga10	Tsga10.aSep08	252923	79457	2902	16	794	testis specific 10 CRA d (Tsga10) alternative variant aSep08, mRNA.
Tsga10	Tsga10.bSep08	252923	23818	776	3	212	testis specific 10 CRA c (Tsga10) alternative variant bSep08, mRNA.
Tsga10	Tsga10.cSep08	252923	800	714	2	80	testis specific 10 CRA d (Tsga10) alternative variant cSep08, mRNA.
Tsga10ip	Tsga10ip.bSep08	361707	2397	751	3	250	testis specific 10 interacting protein (Tsga10ip) alternative variant bSep08, mRNA.
Tsga10ip	Tsga10ip.cSep08	361707	11940	289	2	96	testis specific 10 interacting protein (Tsga10ip) alternative variant cSep08, mRNA.

Tsga14	Tsga14.bSep08	500069	32199	389	6	129	testis specific gene A14 (Tsga14) alternative variant bSep08, mRNA.
Tsga14	Tsga14.cSep08	500069	13251	431	3	73	testis specific gene A14 (Tsga14) alternative variant cSep08, mRNA.
Tshr	Tshr.bSep08	25360	111073	788	2	226	thyroid stimulating hormone receptor (Tshr) alternative variant bSep08, mRNA.
Tsku	Tsku.bSep08	308843	9444	625	2	185	tsukushin (Tsku) alternative variant bSep08, mRNA.
Tslp	Tslp.aSep08	688621	2467	612	1	85	thymic stromal lymphopoietin (Tslp) alternative variant aSep08, mRNA.
Tslp	Tslp.bSep08	688621	3009	990	2	52	thymic stromal lymphopoietin (Tslp) alternative variant bSep08, mRNA.
Tsnax	Tsnax.aSep08	64028	13639	2171	5	284	translin-associated factor X (Tsnax) alternative variant aSep08, mRNA.
Tsnaxip1	Tsnaxip1.bSep08	498944	1106	767	4	166	translin-associated factor X interacting protein 1 (Tsnaxip1) alternative variant bSep08, mRNA.
Tsnaxip1	Tsnaxip1.cSep08	498944	19190	561	3	101	translin-associated factor X interacting protein 1 (Tsnaxip1) alternative variant cSep08, mRNA.
Tsnaxip1	Tsnaxip1.dSep08	498944	5803	533	2	89	translin-associated factor X interacting protein 1 like (Tsnaxip1) alternative variant dSep08, mRNA.
Tsnaxip1	Tsnaxip1.eSep08	498944	1764	705	2	47	translin-associated factor X interacting protein 1 (Tsnaxip1) alternative variant eSep08, mRNA.
Tsnaxip1	Tsnaxip1.fSep08	498944	13585	687	7	89	translin-associated factor X interacting protein 1 (Tsnaxip1) alternative variant fSep08, mRNA.
Tsnaxip1	Tsnaxip1.gSep08	498944	1572	554	4	65	translin-associated factor X interacting protein 1 (Tsnaxip1) alternative variant gSep08, mRNA.
Tspan1	Tspan1.aSep08	298436	9065	1783	6	409	tetraspanin 1 (Tspan1) alternative variant aSep08, mRNA.
Tspan1	Tspan1.cSep08	298436	4144	828	6	225	tetraspanin 1 (24.9 kD) (Tspan1) alternative variant cSep08, mRNA.
Tspan2	Tspan2.bSep08	64521	38800	772	2	239	tetraspanin 2 (Tspan2) alternative variant bSep08, mRNA.
Tspan3	Tspan3.bSep08	300733	8078	783	1	131	tetraspanin 3 (14.7 kD) (Tspan3) alternative variant bSep08, mRNA.
Tspan4	Tspan4.bSep08	293627	17373	707	3	216	tetraspanin 4 (Tspan4) alternative variant bSep08, mRNA.
Tspan4	Tspan4.cSep08	293627	2288	393	1	124	tetraspanin 4 (Tspan4) alternative variant cSep08, mRNA.
Tspan5	Tspan5.bSep08	362048	2533	676	3	114	tetraspanin 5 (Tspan5) alternative variant bSep08, mRNA.
Tspan5	Tspan5.cSep08	362048	135860	515	2	76	tetraspanin 5 (Tspan5) alternative variant cSep08, mRNA.
Tspan6	Tspan6.bSep08	302313	4077	799	5	141	tetraspanin 6 (Tspan6) alternative variant bSep08, mRNA.
Tspan6	Tspan6.cSep08	302313	5112	582	6	119	tetraspanin 6 (Tspan6) alternative variant cSep08, mRNA.
Tspan7	Tspan7.aSep08	363447	16991	1644	6	224	tetraspanin 7 (Tspan7) alternative variant aSep08, mRNA.
Tspan7	Tspan7.cSep08	363447	955	684	1	45	tetraspanin 7 (Tspan7) alternative variant cSep08, mRNA.
Tspan8	Tspan8.bSep08	171048	16443	703	5	156	tetraspanin 8 (Tspan8) alternative variant bSep08, mRNA.
Tspan9	Tspan9.bSep08	312728	160580	728	6	169	tetraspanin 9 (Tspan9) alternative variant bSep08, mRNA.
Tspan9	Tspan9.cSep08	312728	141055	551	5	126	tetraspanin 9 (Tspan9) alternative variant cSep08, mRNA.
Tspan9	Tspan9.dSep08	312728	159585	720	5	110	tetraspanin 9 (Tspan9) alternative variant dSep08, mRNA.
Tspan9	Tspan9.eSep08	312728	170755	453	3	75	tetraspanin 9 (Tspan9) alternative variant eSep08, mRNA.

Tspan11	Tspan11.bSep08	312727	29068	458	1	99	tetraspanin 11 (Tspan11) alternative variant bSep08, mRNA.
Tspan12	Tspan12.bSep08	362326	50726	724	5	233	tetraspanin 12 (Tspan12) alternative variant bSep08, mRNA.
Tspan12	Tspan12.cSep08	362326	78771	970	6	232	tetraspanin 12 (27.2 kD) (Tspan12) alternative variant cSep08, mRNA.
Tspan12	Tspan12.dSep08	362326	62599	1174	9	206	tetraspanin 12 (24.0 kD) (Tspan12) alternative variant dSep08, mRNA.
Tspan12	Tspan12.eSep08	362326	48833	607	6	154	tetraspanin 12 (Tspan12) alternative variant eSep08, mRNA.
Tspan12	Tspan12.gSep08	362326	3390	383	3	93	tetraspanin 12 (Tspan12) alternative variant gSep08, mRNA.
Tspan13	Tspan13.aSep08	366602	27110	1795	6	409	tetraspanin 13 (Tspan13) alternative variant aSep08, complete mRNA.
Tspan13	Tspan13.cSep08	366602	20953	377	4	68	tetraspanin 13 (7.1 kD) (Tspan13) alternative variant cSep08, mRNA.
Tspan14	Tspan14.aSep08	306324	55515	1292	8	325	tetraspanin 14 (Tspan14) alternative variant aSep08, mRNA.
Tspan14	Tspan14.bSep08	306324	52003	2509	9	290	tetraspanin 14 (Tspan14) alternative variant bSep08, mRNA.
Tspan14	Tspan14.dSep08	306324	19533	360	4	119	tetraspanin 14 (Tspan14) alternative variant dSep08, mRNA.
Tspan14	Tspan14.eSep08	306324	5877	965	4	40	tetraspanin 14 (Tspan14) alternative variant eSep08, mRNA.
Tspan18	Tspan18.aSep08	311210	137387	1777	2	453	tetraspanin 18 (Tspan18) alternative variant aSep08, mRNA.
Tspan31	Tspan31.aSep08	362890	3757	2342	6	212	tetraspanin 31 (22.9 kD) (Tspan31) alternative variant aSep08, mRNA.
Tspan32	Tspan32.aSep08	685765	13614	1365	7	210	tetraspanin 32 (Tspan32) alternative variant aSep08, mRNA.
Tspan32	Tspan32.bSep08	685765	13028	756	7	174	tetraspanin 32 (19.7 kD) (Tspan32) alternative variant bSep08, mRNA.
Tspan32	Tspan32.cSep08	685765	13793	760	6	156	tetraspanin 32 (Tspan32) alternative variant cSep08, mRNA.
Tspan32	Tspan32.dSep08	685765	13852	718	5	143	tetraspanin 32 (Tspan32) alternative variant dSep08, mRNA.
Tspan32	Tspan32.fSep08	685765	6675	430	5	79	tetraspanin 32 (Tspan32) alternative variant fSep08, mRNA.
Tspan33	Tspan33.bSep08	500065	15532	282	3	80	tetraspanin 33 (Tspan33) alternative variant bSep08, mRNA.
Tspan33	Tspan33.cSep08	500065	2891	397	2	78	tetraspanin 33 (Tspan33) alternative variant cSep08, mRNA.
Tspsy2	Tspsy2.aSep08	302612	4909	2007	7	476	TSPY-like 2 (Tspsy2) alternative variant aSep08, mRNA.
Tspsy2	Tspsy2.bSep08	302612	1241	792	4	200	TSPY-like 2 (Tspsy2) alternative variant bSep08, mRNA.
TSP_1.0	TSP_1.0.aSep08		16556	1098	1	289	with thrombospondin type 1 motif 2 (TSP_1.0) alternative variant aSep08, mRNA.

TSP_1.0	TSP_1.0.bSep08		16276	818	1	264	with thrombospondin type 1 motif 2 (TSP_1.0) alternative variant bSep08, mRNA.
TSP_1.2	TSP_1.2.aSep08		24826	726		241	thrombospondin type-1 domain-containing protein 7A (TSP_1.2) mRNA.
TSP_3.0	TSP_3.0.aSep08		5828	917		305	thrombospondin (TSP_3.0) mRNA.
TSP_C.0	TSP_C.0.aSep08		3023	721	4	185	thrombospondin 4 (TSP_C.0) alternative variant aSep08, mRNA.
Tssc1	Tssc1.bSep08	362721	92124	673	5	173	tumor suppressing subtransferable candidate 1 (Tssc1) alternative variant bSep08, mRNA.
Tssc1	Tssc1.cSep08	362721	4544	809	3	138	tumor suppressing subtransferable candidate 1 (Tssc1) alternative variant cSep08, mRNA.
Tssc1	Tssc1.dSep08	362721	5064	518	3	88	tumor suppressing subtransferable candidate 1 (Tssc1) alternative variant dSep08, mRNA.
Tssc1	Tssc1.eSep08	362721	1444	658	2	48	tumor suppressing subtransferable candidate 1 (5.3 kD) (Tssc1) alternative variant eSep08, mRNA.
Tssc4	Tssc4.bSep08	361682	1708	1379	3	321	tumor-suppressing subchromosomal transferable fragment 4 (34.3 kD) (Tssc4) alternative variant bSep08, mRNA.
Tssc4	Tssc4.dSep08	361682	1109	750	3	42	tumor-suppressing subchromosomal transferable fragment 4 (4.5 kD) (Tssc4) alternative variant dSep08, mRNA.
Tssk3	Tssk3.bSep08	297891	994	762	2	49	testis-specific serine kinase 3 (5.5 kD) (Tssk3) alternative variant bSep08, mRNA.
Tssk4	Tssk4.aSep08	290229	1477	789	1	223	testis-specific serine kinase 4 (Tssk4) alternative variant aSep08, mRNA.
Tssk5	Tssk5.aSep08	315095	1352	717	2	171	testis-specific serine kinase 5 (Tssk5) alternative variant aSep08, mRNA.
Tssk5	Tssk5.bSep08	315095	3025	2049	5	137	testis-specific serine kinase 5 (15.7 kD) (Tssk5) alternative variant bSep08, mRNA.
Tsta3	Tsta3.bSep08	300036	4839	1195	10	278	tissue specific transplantation antigen P35B like (30.9 kD) (Tsta3) alternative variant bSep08, complete mRNA.
Tsta3	Tsta3.cSep08	300036	3413	741	7	221	tissue specific transplantation antigen P35B like (Tsta3) alternative variant cSep08, mRNA.
Tsta3	Tsta3.dSep08	300036	2831	1633	8	172	tissue specific transplantation antigen P35B like (19.0 kD) (Tsta3) alternative variant dSep08, mRNA.
Tsta3	Tsta3.eSep08	300036	1557	604	3	105	tissue specific transplantation antigen P35B like (11.2 kD) (Tsta3) alternative variant eSep08, mRNA.
Tsta3	Tsta3.fSep08	300036	1833	814	3	105	tissue specific transplantation antigen P35B like (11.2 kD) (Tsta3) alternative variant fSep08, mRNA.
Ttc1	Ttc1.bSep08	287208	10184	511	3	145	putative nuclear protein of vertebrate origin (16.5 kD) (Ttc1) alternative variant bSep08, complete mRNA.
Ttc1	Ttc1.cSep08	287208	2090	474	2	39	putative protein (Ttc1) alternative variant cSep08, mRNA.
Ttc3	Ttc3.aSep08	360702	6811	618	7	178	putative protein of vertebrate origin (Ttc3) alternative variant aSep08, mRNA.
Ttc3	Ttc3.bSep08	360702	18172	575	6	170	putative protein of vertebrate origin (Ttc3) alternative variant bSep08, mRNA.
Ttc3	Ttc3.cSep08	360702	3854	366	4	121	putative protein of vertebrate origin (Ttc3) alternative variant cSep08, mRNA.



Ttc3	Ttc3.dSep08	360702	3594	318	4	106	putative protein of mammalian origin (Ttc3) alternative variant dSep08, mRNA.
Ttc3	Ttc3.eSep08	360702	41997	425	5	51	putative protein (6.2 kD) (Ttc3) alternative variant eSep08, mRNA.
Ttc3	Ttc3.fSep08	360702	40749	388	4	56	putative protein (Ttc3) alternative variant fSep08, mRNA.
Ttc4	Ttc4.aSep08	362556	17405	3205	5	386	tetratricopeptide TPR 1 (44.2 kD) (Ttc4) alternative variant aSep08, complete mRNA.
Ttc4	Ttc4.cSep08	362556	6418	682	1	216	tetratricopeptide TPR 1 (Ttc4) alternative variant cSep08, mRNA.
Ttc5	Ttc5.bSep08	305837	12273	630	6	182	putative protein, with a coiled coil domain, of eukaryotic origin (20.6 kD) (Ttc5) alternative variant bSep08, complete mRNA.
Ttc5	Ttc5.cSep08	305837	1420	652	3	164	putative protein of eukaryotic origin (Ttc5) alternative variant cSep08, mRNA.
Ttc5	Ttc5.dSep08	305837	2375	784	2	56	putative protein of metazoan origin (Ttc5) alternative variant dSep08, mRNA.
Ttc6	Ttc6.aSep08	299067	21341	809	5	269	tetratricopeptide TPR 1 (Ttc6) alternative variant aSep08, mRNA.
Ttc6	Ttc6.bSep08	299067	34571	712	5	237	tetratricopeptide TPR 1 (Ttc6) alternative variant bSep08, mRNA.
Ttc6	Ttc6.cSep08	299067	42021	673	6	168	tetratricopeptide TPR 1 (Ttc6) alternative variant cSep08, mRNA.
Ttc7	Ttc7.aSep08	362696	107497	4481	18	941	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc7) alternative variant aSep08, mRNA.
Ttc7	Ttc7.bSep08	362696	4137	620	1	157	putative protein (Ttc7) alternative variant bSep08, mRNA.
Ttc7b	Ttc7b.bSep08	362768	70974	1712	6	291	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 and tetratricopeptide TPR 4 (Ttc7b) alternative variant bSep08, mRNA.
Ttc7b	Ttc7b.cSep08	362768	30381	619	6	206	putative protein of metazoan origin (Ttc7b) alternative variant cSep08, mRNA.
Ttc7b	Ttc7b.dSep08	362768	46971	768	5	140	putative protein of bilateral origin (Ttc7b) alternative variant dSep08, mRNA.
Ttc8	Ttc8.aSep08	299246	54287	2300	14	505	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (57.4 kD) (Ttc8) alternative variant aSep08, mRNA.
Ttc8	Ttc8.cSep08	299246	17222	390	5	115	putative protein of eukaryotic origin (Ttc8) alternative variant cSep08, mRNA.
Ttc8	Ttc8.dSep08	299246	16693	422	5	102	putative protein of eukaryotic origin (Ttc8) alternative variant dSep08, mRNA.
Ttc8	Ttc8.fSep08	299246	513	421	2	26	putative protein (3.0 kD) (Ttc8) alternative variant fSep08, mRNA.
Ttc9	Ttc9.aSep08	500689	34990	2234	4	250	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc9) alternative variant aSep08, mRNA.
Ttc9	Ttc9.cSep08	500689	82852	407	4	132	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc9) alternative variant cSep08, mRNA.
Ttc9	Ttc9.dSep08	500689	701	603	2	63	putative protein of vertebrate origin (Ttc9) alternative variant dSep08, mRNA.

Ttc9c	Ttc9c.aSep08	309196	9705	1473	3	171	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (20.1 kD) (Ttc9c) alternative variant aSep08, mRNA.
Ttc9c	Ttc9c.bSep08	309196	9595	1104	5	171	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (20.1 kD) (Ttc9c) alternative variant bSep08, mRNA.
Ttc9c	Ttc9c.dSep08	309196	4709	512	3	28	putative protein (3.0 kD) (Ttc9c) alternative variant dSep08, complete mRNA.
Ttc12	Ttc12.aSep08	300696	38323	1453		478	tetratricopeptide repeat 12 (Ttc12) mRNA.
Ttc13	Ttc13.aSep08	292095	56057	3283	20	791	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc13) alternative variant aSep08, mRNA.
Ttc13	Ttc13.bSep08	292095	8522	712	6	222	putative protein of eukaryotic origin (Ttc13) alternative variant bSep08, mRNA.
Ttc13	Ttc13.cSep08	292095	2846	534	3	118	putative protein of eukaryotic origin (Ttc13) alternative variant cSep08, mRNA.
Ttc13	Ttc13.dSep08	292095	2130	579	1	39	putative protein (Ttc13) alternative variant dSep08, mRNA.
Ttc15	Ttc15.aSep08	314013	65179	3190	12	797	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (87.5 kD) (Ttc15) alternative variant aSep08, mRNA.
Ttc15	Ttc15.cSep08	314013	6851	606	3	116	putative protein of metazoan origin (Ttc15) alternative variant cSep08, mRNA.
Ttc17	Ttc17.aSep08	311224	109195	3487	19	1162	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc17) alternative variant aSep08, mRNA.
Ttc17	Ttc17.cSep08	311224	34859	1706	2	352	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc17) alternative variant cSep08, mRNA.
Ttc18	Ttc18.bSep08	361006	10901	418	4	139	repeat 18 (Ttc18) alternative variant bSep08, mRNA.
Ttc18	Ttc18.cSep08	361006	3547	403	2	33	putative protein (Ttc18) alternative variant cSep08, mRNA.
Ttc19	Ttc19.bSep08	691506	2040	255	2	62	putative protein of vertebrate origin (Ttc19) alternative variant bSep08, mRNA.
ttc21a	ttc21a.aSep08	301065	4220	697	5	232	tetratricopeptide repeat 21A (ttc21a) alternative variant aSep08, mRNA.
Ttc21b	Ttc21b.aSep08	295654	3866	433		143	tetratricopeptide TPR 1 and tetratricopeptide TPR 2 (Ttc21b) mRNA.
Ttc23	Ttc23.aSep08	308708	51026	1391	7	463	putative protein (Ttc23) alternative variant aSep08, mRNA.
Ttc23	Ttc23.bSep08	308708	50807	1130	6	227	putative protein of metazoan origin (Ttc23) alternative variant bSep08, mRNA.
Ttc23	Ttc23.cSep08	308708	14190	650	5	183	putative protein of metazoan origin (20.8 kD) (Ttc23) alternative variant cSep08, mRNA.
Ttc23	Ttc23.dSep08	308708	62513	754	4	153	putative protein of mammalian origin (Ttc23) alternative variant dSep08, mRNA.
Ttc23	Ttc23.eSep08	308708	21851	728	4	142	putative protein of metazoan origin (Ttc23) alternative variant eSep08, mRNA.
Ttc25	Ttc25.bSep08	303534	9875	648	3	111	putative protein of ancient origin (Ttc25) alternative variant bSep08, mRNA.
Ttc25	Ttc25.cSep08	303534	1262	356	3	84	putative protein of eukaryotic origin (Ttc25) alternative variant cSep08, mRNA.
Ttc26	Ttc26.bSep08	500086	18049	2684		176	tetratricopeptide repeat 26 (Ttc26) alternative variant bSep08, mRNA.

Ttc27	Ttc27.bSep08	298782	7348	634	4	176	putative protein, with a coiled coil domain, of eukaryotic origin (Ttc27) alternative variant bSep08, mRNA.
Ttc27	Ttc27.cSep08	298782	30573	690	5	126	putative protein, with a coiled coil domain, of eukaryotic origin (Ttc27) alternative variant cSep08, mRNA.
Ttf1	Ttf1.aSep08	499766	6421	727		242	transcription termination factor, RNA polymerase I (Ttf1) mRNA.
Ttf2	Ttf2.aSep08	295324	15843	2658	13	484	transcription termination factor, RNA polymerase II (Ttf2) alternative variant aSep08, mRNA.
Ttf2	Ttf2.bSep08	295324	3097	1480	2	71	transcription termination factor, RNA polymerase II (8.3 kD) (Ttf2) alternative variant bSep08, mRNA.
Ttk	Ttk.bSep08	315852	15496	846	7	281	tkk protein kinase (Ttk) alternative variant bSep08, mRNA.
Ttk	Ttk.cSep08	315852	2038	453	2	84	tkk protein kinase (Ttk) alternative variant cSep08, mRNA.
Ttl1	Ttl1.bSep08	362969	5554	415		71	tubulin tyrosine ligase-like 1 (Ttl1) alternative variant bSep08, mRNA.
Ttl3	Ttl3.aSep08	362415	22199	2772	11	662	tubulin tyrosine ligase-like family member 3 (Ttl3) alternative variant aSep08, mRNA.
Ttl3	Ttl3.cSep08	362415	3803	365	2	121	tubulin tyrosine ligase-like family member 3 (Ttl3) alternative variant cSep08, mRNA.
Ttl3	Ttl3.eSep08	362415	3932	763	2	112	tubulin tyrosine ligase-like family member 3 CRA b (Ttl3) alternative variant eSep08, mRNA.
Ttl3	Ttl3.fSep08	362415	10051	783	5	101	ligase-like protein 3 (Ttl3) alternative variant fSep08, mRNA.
Ttl4	Ttl4.aSep08	690512	2736	718		239	tubulin tyrosine ligase-like family, member 4 (Ttl4) mRNA.
Ttl7	Ttl7.aSep08	310982	44655	4038		219	tubulin tyrosine ligase-like family, member 7 (Ttl7) mRNA.
Ttl9	Ttl9.bSep08	311548	6130	527	4	117	tubulin tyrosine ligase-like family, member 9 (13.8 kD) (Ttl9) alternative variant bSep08, mRNA.
Ttl9	Ttl9.cSep08	311548	5320	288	4	88	tubulin tyrosine ligase-like family, member 9 (Ttl9) alternative variant cSep08, mRNA.
Ttl10	Ttl10.bSep08	298692	3120	1044	5	301	tubulin tyrosine ligase-like family, member 10 (Ttl10) alternative variant bSep08, mRNA.
Ttl10	Ttl10.cSep08	298692	11694	1842	12	270	tubulin tyrosine ligase-like family, member 10 (Ttl10) alternative variant cSep08, mRNA.
Ttl10	Ttl10.dSep08	298692	4325	810	8	219	tubulin tyrosine ligase-like family, member 10 (25.2 kD) (Ttl10) alternative variant dSep08, mRNA.
Ttl11	Ttl11.aSep08	689746	197141	2529	7	513	tubulin tyrosine ligase-like family, member 11 (Ttl11) alternative variant aSep08, mRNA.
Ttl11	Ttl11.bSep08	689746	123313	644	6	214	tubulin tyrosine ligase-like family, member 11 (Ttl11) alternative variant bSep08, mRNA.
Ttl12	Ttl12.aSep08	300105	20140	3106		637	tubulin tyrosine ligase-like family, member 12 (Ttl12) mRNA.
Ttn	Ttn.aSep08	84015	6079	4229	6	923	titin (Ttn) alternative variant aSep08, mRNA.
Ttpa	Ttpa.bSep08	25571	21634	2925	2	209	tocopherol (alpha) transfer protein (24.0 kD) (Ttpa) alternative variant bSep08, mRNA.
Ttr	Ttr.bSep08	24856	4097	510	2	170	transthyretin (Ttr) alternative variant bSep08, mRNA.
Ttr	Ttr.cSep08	24856	3875	300	2	93	transthyretin (Ttr) alternative variant cSep08, mRNA.

Ttyh1	Ttyh1.aSep08	292597	18529	3091	14	481	tweety homolog 1 CRA e (Ttyh1) alternative variant aSep08, mRNA.
Ttyh1	Ttyh1.bSep08	292597	11365	2106	11	319	tweety homolog 1 CRA a (35.4 kD) (Ttyh1) alternative variant bSep08, mRNA.
Ttyh1	Ttyh1.cSep08	292597	6191	944	6	184	tweety homolog 1 CRA d (20.1 kD) (Ttyh1) alternative variant cSep08, mRNA.
Ttyh1	Ttyh1.dSep08	292597	3419	399	2	125	tweety 1 (Ttyh1) alternative variant dSep08, mRNA.
Ttyh1	Ttyh1.eSep08	292597	5334	372	3	124	tweety 1 (Ttyh1) alternative variant eSep08, mRNA.
Ttyh1	Ttyh1.gSep08	292597	3939	812	3	107	tweety homolog 1 CRA c (12.0 kD) (Ttyh1) alternative variant gSep08, mRNA.
Ttyh2	Ttyh2.aSep08	287803	11269	2290		160	tweety homolog 2 (Drosophila) (Ttyh2) mRNA.
Ttyh3	Ttyh3.cSep08	304315	2682	495	2	57	tweety homolog 3 (Drosophila) (Ttyh3) alternative variant cSep08, mRNA.
Tub.0	Tub.0.aSep08		5933	2443	5	210	tubby (Tub.0) alternative variant aSep08, mRNA.
Tub.0	Tub.0.bSep08		1658	791	2	88	tubby (10.0 kD) (Tub.0) alternative variant bSep08, mRNA.
Tuba1b	Tuba1b.bSep08	500929	664	415	1	128	tubulin, alpha 1B (Tuba1b) alternative variant bSep08, mRNA.
Tuba1b	Tuba1b.cSep08	500929	1384	318		79	tubulin, alpha 1B (Tuba1b) alternative variant cSep08, mRNA.
Tuba1c	Tuba1c.bSep08	300218	6761	823	1	131	tubulin, alpha 1C (Tuba1c) alternative variant bSep08, mRNA.
Tuba4a	Tuba4a.aSep08	316531	2502	2024	2	452	tubulin, alpha 4A (Tuba4a) alternative variant aSep08, mRNA.
Tuba4a	Tuba4a.cSep08	316531	3024	781	2	248	tubulin, alpha 4A (Tuba4a) alternative variant cSep08, mRNA.
Tubb2a	Tubb2a.cSep08	498736	1608	339	2	90	tubulin, beta 2a (Tubb2a) alternative variant cSep08, mRNA.
Tubb2a	Tubb2a.eSep08	498736	21869	286	4	54	tubulin, beta 2a (Tubb2a) alternative variant eSep08, mRNA.
Tubb2b	Tubb2b.bSep08	291081	1053	733	2	244	tubulin, beta 2b (Tubb2b) alternative variant bSep08, mRNA.
Tube1	Tube1.bSep08	361856	14996	1129	10	346	epsilon-tubulin 1 (Tube1) alternative variant bSep08, mRNA.
Tube1	Tube1.cSep08	361856	17988	2369	12	151	epsilon-tubulin 1 (16.5 kD) (Tube1) alternative variant cSep08, mRNA.
Tube1	Tube1.dSep08	361856	2449	417	2	113	epsilon-tubulin 1 (Tube1) alternative variant dSep08, mRNA.
Tube1	Tube1.eSep08	361856	11732	797	8	76	epsilon-tubulin 1 (8.4 kD) (Tube1) alternative variant eSep08, mRNA.
Tubg1	Tubg1.bSep08	252921	4290	738	6	214	tubulin, gamma 1 (Tubg1) alternative variant bSep08, mRNA.
Tubg1	Tubg1.cSep08	252921	802	288	3	96	tubulin, gamma 1 (Tubg1) alternative variant cSep08, mRNA.
Tubg1	Tubg1.dSep08	252921	1001	471	4	93	tubulin, gamma 1 (Tubg1) alternative variant dSep08, mRNA.

Tubg1	Tubg1.eSep08	252921	1427	783	2	88	tubulin, gamma 1 (Tubg1) alternative variant eSep08, mRNA.
Tubg1	Tubg1.fSep08	252921	6540	368	3	58	tubulin, gamma 1 (6.5 kD) (Tubg1) alternative variant fSep08, complete mRNA.
Tubg2	Tubg2.aSep08	680991	1715	868	5	276	tubulin, gamma 2 (Tubg2) alternative variant aSep08, mRNA.
Tubg2	Tubg2.bSep08	680991	680	472	2	95	tubulin, gamma 2 (Tubg2) alternative variant bSep08, mRNA.
Tubgcp2	Tubgcp2.bSep08	309098	4215	787	5	262	tubulin, gamma complex associated protein 2 (Tubgcp2) alternative variant bSep08, mRNA.
Tubgcp2	Tubgcp2.cSep08	309098	2093	525	2	137	tubulin, gamma complex associated protein 2 (15.1 kD) (Tubgcp2) alternative variant cSep08, mRNA.
Tubgcp3	Tubgcp3.bSep08	306599	22043	1012	3	337	tubulin, gamma complex associated protein 3 (Tubgcp3) alternative variant bSep08, mRNA.
Tubgcp3	Tubgcp3.cSep08	306599	16348	679	3	127	tubulin, gamma complex associated protein 3 (Tubgcp3) alternative variant cSep08, mRNA.
Tubgcp6	Tubgcp6.bSep08	362980	1737	1570	3	149	tubulin, gamma complex associated protein 6 (Tubgcp6) alternative variant bSep08, mRNA.
Tubgcp6	Tubgcp6.cSep08	362980	1766	1685	2	125	tubulin, gamma complex associated protein 6 (Tubgcp6) alternative variant cSep08, mRNA.
tuby	tuby.aSep08		1326	980		142	putative secreted or extracellular protein precursor of mammalian origin (15.5 kD) (tuby) mRNA.
tuchy	tuchy.aSep08		1545	310		41	putative protein (tuchy) mRNA.
tudar	tudar.aSep08		1471	428		142	hydrocephalus inducing (tudar) mRNA.
tuflo	tuflo.aSep08		2111	276		56	putative protein (6.5 kD) (tuflo) mRNA.
tuflu	tuflu.cSep08		587	394	2	71	putative protein (tuflu) alternative variant cSep08, mRNA.
Tufm	Tufm.bSep08	293481	3069	1387	7	229	tu translation elongation factor, mitochondrial (25.0 kD) (Tufm) alternative variant bSep08, mRNA.
Tufm	Tufm.cSep08	293481	1451	807	4	189	tu translation elongation factor, mitochondrial (21.1 kD) (Tufm) alternative variant cSep08, mRNA.
Tuft1	Tuft1.aSep08	365864	46099	2361	10	365	tuftelin 1 (41.4 kD) (Tuft1) alternative variant aSep08, complete mRNA.
Tuft1	Tuft1.cSep08	365864	32157	579	5	193	tuftelin 1 (Tuft1) alternative variant cSep08, mRNA.
tugar	tugar.aSep08		2475	742	4	163	putative protein (17.9 kD) (tugar) alternative variant aSep08, mRNA.
tugar	tugar.bSep08		831	379	3	69	unc-13 homolog B CRA b (tugar) alternative variant bSep08, mRNA.
tuja	tuja.aSep08		3349	1001		260	telomerase protein 1 (tuja) mRNA.
tujey	tujey.bSep08		3226	763	2	59	putative protein (tujey) alternative variant bSep08, mRNA.
tukee	tukee.aSep08		6087	675		66	putative protein (7.3 kD) (tukee) mRNA.
tukler	tukler.aSep08		692	569		42	putative protein (4.8 kD) (tukler) mRNA.
tulo	tulo.aSep08		2442	550	3	111	putative protein (12.2 kD) (tulo) alternative variant aSep08, mRNA.
Tulp1	Tulp1.bSep08	309900	2699	855	8	271	tubby like protein 1 (Tulp1) alternative variant bSep08, mRNA.

Tulp2	Tulp2.aSep08	361576	9103	1931	13	555	tubby-like protein 2 (62.3 kD) (Tulp2) alternative variant aSep08, complete mRNA.
Tulp2	Tulp2.cSep08	361576	2966	682	5	217	tubby-like protein 2 (Tulp2) alternative variant cSep08, mRNA.
Tulp2	Tulp2.dSep08	361576	4040	719	7	162	tubby-like protein 2 (Tulp2) alternative variant dSep08, mRNA.
Tulp2	Tulp2.eSep08	361576	666	396	3	82	tubby-like protein 2 (Tulp2) alternative variant eSep08, mRNA.
tumee	tumee.aSep08		7353	457		40	putative protein (4.7 kD) (tumee) mRNA.
tunoy	tunoy.aSep08		58254	236		62	putative protein (tunoy) mRNA.
tupor	tupor.aSep08		20271	398		42	putative protein (4.9 kD) (tupor) mRNA.
tusa	tusa.aSep08		2372	494	1	96	putative mitochondrial protein (10.8 kD) (tusa) alternative variant aSep08, mRNA.
tusa	tusa.bSep08		12249	685	4	69	CRA b like (8.1 kD) (tusa) alternative variant bSep08, mRNA.
Tusc3	Tusc3.aSep08	290783	99844	1783	6	448	tumor suppressor candidate 3 (Tusc3) alternative variant aSep08, mRNA.
Tusc4	Tusc4.bSep08	363138	1603	735	4	244	tumor suppressor candidate 4 (Tusc4) alternative variant bSep08, mRNA.
Tusc4	Tusc4.cSep08	363138	1825	708	6	171	tumor suppressor candidate 4 (Tusc4) alternative variant cSep08, mRNA.
Tusc4	Tusc4.dSep08	363138	1769	1185	4	163	tumor suppressor candidate 4 (18.1 kD) (Tusc4) alternative variant dSep08, mRNA.
Tusc5	Tusc5.aSep08	360576	26528	1249		173	tumor suppressor candidate 5 (18.7 kD) (Tusc5) mRNA.
tushee	tushee.aSep08		6346	438		20	putative protein (tushee) mRNA.
Tut1	Tut1.bSep08	499314	4944	2725	1	205	terminal uridylyl transferase 1, U6 snRNA-specific (Tut1) alternative variant bSep08, mRNA.
tutu	tutu.bSep08		624	344	3	45	putative protein (tutu) alternative variant bSep08, mRNA.
tuvar	tuvar.aSep08		983	501		166	putative protein of vertebrate origin (tuvar) mRNA.
tuwey	tuwey.aSep08		1092	667		40	putative protein of mammalian origin (4.7 kD) (tuwey) mRNA.
Twf1	Twf1.bSep08	315265	5273	1541	4	147	twinfilin, actin-binding protein, homolog 1 (Drosophila) (17.3 kD) (Twf1) alternative variant bSep08, mRNA.
Twsg1	Twsg1.bSep08	363294	7404	641	3	54	twisted gastrulation homolog 1 (Drosophila) (5.7 kD) (Twsg1) alternative variant bSep08, mRNA.
Txk	Txk.bSep08	305311	3759	742	1	70	TXK tyrosine kinase (Txk) alternative variant bSep08, mRNA.
Txlnb	Txlnb.aSep08	308622	10818	294		97	taxilin beta (Txlnb) mRNA.
Txndc1	Txndc1.bSep08	362751	6191	766	7	229	thioredoxin domain containing protein (Txndc1) alternative variant bSep08, mRNA.
Txndc1	Txndc1.cSep08	362751	3327	420	3	111	thioredoxin domain containing protein (Txndc1) alternative variant cSep08, mRNA.
Txndc2	Txndc2.aSep08	316777	2326	1783	1	550	thioredoxin domain containing protein (61.1 kD) (Txndc2) alternative variant aSep08, mRNA.
Txndc2	Txndc2.cSep08	316777	3874	970	1	323	putative protein of eukaryotic origin (Txndc2) alternative variant cSep08, mRNA.

Txndc3	Txndc3.aSep08	364729	23597	713	4	237	putative protein of ancient origin (Txndc3) alternative variant aSep08, mRNA.
Txndc3	Txndc3.bSep08	364729	19551	797	9	226	thioredoxin domain containing protein (Txndc3) alternative variant bSep08, mRNA.
Txndc4	Txndc4.bSep08	298066	94252	2556	10	369	thioredoxin domain containing protein (42.8 kD) (Txndc4) alternative variant bSep08, complete mRNA.
Txndc4	Txndc4.cSep08	298066	93169	1288	9	238	putative protein of eukaryotic origin (27.9 kD) (Txndc4) alternative variant cSep08, mRNA.
Txndc4	Txndc4.dSep08	298066	77372	610	4	153	putative protein of eukaryotic origin (Txndc4) alternative variant dSep08, mRNA.
Txndc4	Txndc4.eSep08	298066	60246	415	2	137	thioredoxin domain containing protein (Txndc4) alternative variant eSep08, mRNA.
Txndc4	Txndc4.fSep08	298066	81527	774	7	125	putative protein of ancient origin (Txndc4) alternative variant fSep08, mRNA.
Txndc4	Txndc4.gSep08	298066	60276	449	2	82	thioredoxin domain containing protein (Txndc4) alternative variant gSep08, mRNA.
Txndc9	Txndc9.bSep08	280671	6143	549	4	183	ATP binding protein associated with cell differentiation like (Txndc9) alternative variant bSep08, mRNA.
Txndc9	Txndc9.cSep08	280671	3619	742	2	101	ATP binding protein like (Txndc9) alternative variant cSep08, mRNA.
Txndc11	Txndc11.bSep08	302899	32944	933	3	275	putative protein, with a coiled coil domain, of eukaryotic origin (Txndc11) alternative variant bSep08, mRNA.
Txndc11	Txndc11.dSep08	302899	6008	947	2	131	putative protein of vertebrate origin (Txndc11) alternative variant dSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.aSep08	298370	6264	1472	3	350	KTI12 homolog chromatin associated (38.4 kD) (Txndc12andKti12) alternative variant aSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.aSep08	685656	6264	1472	3	350	KTI12 homolog chromatin associated (38.4 kD) (Txndc12andKti12) alternative variant aSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.cSep08	298370	25030	1315	7	217	putative protein of ancient origin (Txndc12andKti12) alternative variant cSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.cSep08	685656	25030	1315	7	217	putative protein of ancient origin (Txndc12andKti12) alternative variant cSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.dSep08	298370	22852	922	6	191	endoplasmic reticulum protein ERp19 (Txndc12andKti12) alternative variant dSep08, mRNA.
Txndc12andKti12	Txndc12andKti12.dSep08	685656	22852	922	6	191	endoplasmic reticulum protein ERp19 (Txndc12andKti12) alternative variant dSep08, mRNA.
Txndc13	Txndc13.aSep08	296182	43393	4026		336	thioredoxin domain containing protein (37.5 kD) (Txndc13) alternative variant aSep08, mRNA.
Txndc15	Txndc15.bSep08	307180	10748	816	3	247	putative protein of metazoan origin (Txndc15) alternative variant bSep08, mRNA.
Txndc15	Txndc15.cSep08	307180	6578	760	2	135	putative protein of vertebrate origin (Txndc15) alternative variant cSep08, mRNA.
Txndc16	Txndc16.aSep08	361025	10832	1661	3	269	putative protein of vertebrate origin (Txndc16) alternative variant aSep08, mRNA.
Txndc16	Txndc16.bSep08	361025	4610	426	1	21	putative protein (Txndc16) alternative variant bSep08, mRNA.

Txnip	Txnip.bSep08	117514	1182	967	3	111	thioredoxin interacting protein (11.8 kD) (Txnip) alternative variant bSep08, mRNA.
Txnip	Txnip.dSep08	117514	915	794	2	51	thioredoxin interacting protein (Txnip) alternative variant dSep08, mRNA.
Txnip	Txnip.eSep08	117514	516	408	2	34	thioredoxin interacting protein (Txnip) alternative variant eSep08, mRNA.
Txn1	Txn1.bSep08	140922	13399	3613	3	86	thioredoxin-like 1 (9.8 kD) (Txn1) alternative variant bSep08, mRNA.
Txn1	Txn1.dSep08	140922	2044	511	2	29	thioredoxin-like 1 (3.3 kD) (Txn1) alternative variant dSep08, mRNA.
Txn14b	Txn14b.bSep08	292008	3298	313	1	66	thioredoxin-like 4B (Txn14b) alternative variant bSep08, mRNA.
Txn14b	Txn14b.cSep08	292008	4238	403	1	54	thioredoxin-like 4B (Txn14b) alternative variant cSep08, mRNA.
Txnrd1	Txnrd1.bSep08	58819	60961	702	7	204	thioredoxin reductase 1 (Txnrd1) alternative variant bSep08, mRNA.
Txnrd1	Txnrd1.cSep08	58819	13735	289	4	95	thioredoxin reductase 1 (Txnrd1) alternative variant cSep08, mRNA.
Txnrd2	Txnrd2.bSep08	50551	21493	1755	7	275	thioredoxin reductase 2 (Txnrd2) alternative variant bSep08, mRNA.
Txnrd2	Txnrd2.cSep08	50551	25141	717	9	237	thioredoxin reductase 2 (Txnrd2) alternative variant cSep08, mRNA.
Txnrd3	Txnrd3.aSep08	297437	20799	1990	10	434	thioredoxin reductase 3 (Txnrd3) alternative variant aSep08, mRNA.
Txnrd3	Txnrd3.bSep08	297437	17782	336	2	112	thioredoxin reductase 3 (Txnrd3) alternative variant bSep08, mRNA.
tyby	tyby.aSep08		3880	1176		28	putative protein (tyby) mRNA.
tychy	tychy.aSep08		5387	816		6	putative protein (0.7 kD) (tychy) mRNA.
tydar	tydar.aSep08		1678	725		69	hydrocephalus inducing like (tydar) mRNA.
tyflo	tyflo.aSep08		1341	606	3	52	putative protein (6.2 kD) (tyflo) alternative variant aSep08, mRNA.
tyflu	tyflu.aSep08		1524	414		137	semaphorin 4B (tyflu) mRNA.
tygar	tygar.aSep08		4984	595	2	198	B-cell novel protein 1 (tygar) alternative variant aSep08, mRNA.
tygar	tygar.bSep08		2909	383	1	127	B-cell novel protein 1 (tygar) alternative variant bSep08, mRNA.
tyja	tyja.aSep08		13130	679		67	putative protein (tyja) mRNA.
tyjey	tyjey.aSep08		1726	384		35	putative protein (3.6 kD) (tyjey) mRNA.
tykee	tykee.aSep08		1175	267		28	putative protein (3.2 kD) (tykee) mRNA.
tykler	tykler.aSep08		13149	344		42	putative protein (4.7 kD) (tykler) mRNA.
tylo	tylo.aSep08		1776	325		55	putative protein (tylo) mRNA.
tymee	tymee.aSep08		448	330		72	putative cytoplasmic protein (8.2 kD) (tymee) mRNA.
Tymp	Tymp.bSep08	315219	2111	1266	2	258	cytochrome oxidase deficient homolog 2 (29.3 kD) (Tymp) alternative variant bSep08, mRNA.
Tymp	Tymp.cSep08	315219	1832	698	5	205	thymidine phosphorylase (Tymp) alternative variant cSep08, mRNA.



Tymp	Tymp.eSep08	315219	512	376	2	125	thymidine phosphorylase (Tymp) alternative variant eSep08, mRNA.
Tymp	Tymp.fSep08	315219	1584	767	2	117	thymidine phosphorylase (Tymp) alternative variant fSep08, mRNA.
Tymp	Tymp.gSep08	315219	1467	571	3	94	thymidine phosphorylase (Tymp) alternative variant gSep08, mRNA.
tynoy	tynoy.aSep08		13817	710	5	236	dedicator of cytokinesis 10 (tynoy) alternative variant aSep08, mRNA.
tynoy	tynoy.bSep08		1926	507	3	151	dedicator of cytokinesis 10 (tynoy) alternative variant bSep08, mRNA.
typor	typor.bSep08		456	361	2	55	putative protein (typor) alternative variant bSep08, mRNA.
Tyro3	Tyro3.bSep08	25232	8564	934	7	311	TYRO3 protein tyrosine kinase 3 (Tyro3) alternative variant bSep08, mRNA.
Tyro3	Tyro3.cSep08	25232	768	406	2	92	TYRO3 protein tyrosine kinase 3 (Tyro3) alternative variant cSep08, mRNA.
Tyro3	Tyro3.dSep08	25232	1638	734	2	68	TYRO3 protein tyrosine kinase 3 (Tyro3) alternative variant dSep08, mRNA.
Tyrobp	Tyrobp.bSep08	361537	3835	497	5	106	tyro protein tyrosine kinase binding protein (Tyrobp) alternative variant bSep08, mRNA.
Tyrobp	Tyrobp.dSep08	361537	3859	669	4	35	tyro protein tyrosine kinase binding protein (Tyrobp) alternative variant dSep08, mRNA.
tysa	tysa.bSep08		1026	350		29	putative protein (3.3 kD) (tysa) alternative variant bSep08, mRNA.
tyshee	tyshee.aSep08		1699	722		117	putative protein (tyshee) mRNA.
Tysnd1	Tysnd1.bSep08	365571	3067	1063	1	192	putative peroxisomal protein of ancient origin (20.4 kD) (Tysnd1) alternative variant bSep08, mRNA.
tytu	tytu.aSep08		15129	3562		81	CRA a like (tytu) mRNA.
tyvar	tyvar.aSep08		1421	347		8	putative protein (0.6 kD) (tyvar) mRNA.
Tyw1	Tyw1.bSep08	304423	54300	1777	1	506	tRNA-yW synthesizing protein 1 homolog (S. cerevisiae) (Tyw1) alternative variant bSep08, mRNA.
Tyw3	Tyw3.aSep08	499731	22332	1109	5	173	tRNA-yW synthesizing protein 3 homolog (S. cerevisiae) (19.1 kD) (Tyw3) alternative variant aSep08, mRNA.
Tyw3	Tyw3.bSep08	499731	10478	807	1	81	tRNA-yW synthesizing protein 3 homolog (S. cerevisiae) (9.1 kD) (Tyw3) alternative variant bSep08, mRNA.
Tyw3	Tyw3.cSep08	499731	19657	652	2	62	tRNA-yW synthesizing protein 3 homolog (S. cerevisiae) (Tyw3) alternative variant cSep08, mRNA.
tywey	tywey.aSep08		10673	625		78	putative protein (tywey) mRNA.
U1snrnpbp	U1snrnpbp.aSep08	360803	7496	925	1	270	u11/U12 snRNP 35K (U1snrnpbp) alternative variant aSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.bSep08	361542	1437	621	6	174	small nuclear RNA factor 1 (U2af114andRGD1563574) alternative variant bSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.bSep08	499126	1437	621	6	174	small nuclear RNA factor 1 (U2af114andRGD1563574) alternative variant bSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.cSep08	361542	1170	401	6	133	small nuclear RNA factor 1 (U2af114andRGD1563574) alternative variant cSep08, mRNA.

U2af114andRGD1563574	U2af114andRGD1563574.cSep08	499126	1170	401	6	133	small nuclear RNA factor 1 (U2af114andRGD1563574) alternative variant cSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.dSep08	361542	2110	860	8	122	small nuclear RNA factor 1 (14.5 kD) (U2af114andRGD1563574) alternative variant dSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.dSep08	499126	2110	860	8	122	small nuclear RNA factor 1 (14.5 kD) (U2af114andRGD1563574) alternative variant dSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.eSep08	361542	2137	1379	3	100	u2 small nuclear RNA auxiliary factor 1-like 4 (10.5 kD) (U2af114andRGD1563574) alternative variant eSep08, mRNA.
U2af114andRGD1563574	U2af114andRGD1563574.eSep08	499126	2137	1379	3	100	u2 small nuclear RNA auxiliary factor 1-like 4 (10.5 kD) (U2af114andRGD1563574) alternative variant eSep08, mRNA.
Uap1	Uap1.aSep08	498272	34138	2324	10	522	UDP-N-acetylglucosamine pyrophosphorylase 1 (58.5 kD) (Uap1) alternative variant aSep08, mRNA.
Uap1	Uap1.bSep08	498272	14944	1253	5	260	UDP-N-acetylglucosamine pyrophosphorylase 1 (Uap1) alternative variant bSep08, mRNA.
Uap1	Uap1.cSep08	498272	1715	908	1	73	UDP-N-acetylglucosamine pyrophosphorylase 1 (8.0 kD) (Uap1) alternative variant cSep08, mRNA.
UBA.0	UBA.0.aSep08		21556	328		109	ubiquitin-associated protein 2 (UBA.0) mRNA.
Uba1	Uba1.bSep08	314432	3739	1124	7	240	ubiquitin-like modifier activating enzyme 1 (Uba1) alternative variant bSep08, mRNA.
Uba1	Uba1.cSep08	314432	2711	696	5	231	ubiquitin-like modifier activating enzyme 1 (Uba1) alternative variant cSep08, mRNA.
Uba1	Uba1.dSep08	314432	8469	349	4	116	ubiquitin-like modifier activating enzyme 1 (Uba1) alternative variant dSep08, mRNA.
Uba1	Uba1.eSep08	314432	745	253	2	84	ubiquitin-like modifier activating enzyme 1 (Uba1) alternative variant eSep08, mRNA.
Uba2	Uba2.aSep08	308508	22434	2245	14	538	ubiquitin-like modifier activating enzyme 2 (Uba2) alternative variant aSep08, mRNA.
Uba3	Uba3.aSep08	117553	22164	2161	18	460	ubiquitin-like modifier activating enzyme 3 (Uba3) alternative variant aSep08, mRNA.
Uba3	Uba3.bSep08	117553	16723	702	8	187	ubiquitin-like modifier activating enzyme 3 (Uba3) alternative variant bSep08, mRNA.
Uba3	Uba3.cSep08	117553	16747	807	9	126	ubiquitin-like modifier activating enzyme 3 (Uba3) alternative variant cSep08, mRNA.
Uba5	Uba5.bSep08	300968	3493	1506	4	118	ubiquitin-like modifier activating enzyme 5 (13.5 kD) (Uba5) alternative variant bSep08, mRNA.
Uba5	Uba5.cSep08	300968	789	323	3	107	ubiquitin-like modifier activating enzyme 5 (Uba5) alternative variant cSep08, mRNA.
Uba52	Uba52.cSep08	64156	1944	446	4	148	ubiquitin A-52 residue ribosomal protein fusion product 1 (Uba52) alternative variant cSep08, mRNA.
Uba52	Uba52.dSep08	64156	2133	477	4	128	ubiquitin A-52 residue ribosomal protein fusion product 1 (14.7 kD) (Uba52) alternative variant dSep08, complete mRNA.

Uba52	Uba52.eSep08	64156	2133	700	4	128	ubiquitin A-52 residue ribosomal protein fusion product 1 (14.7 kD) (Uba52) alternative variant eSep08, complete mRNA.
Ubac1	Ubac1.bSep08	362087	2507	768	3	145	ubiquitin-associated (Ubac1) alternative variant bSep08, mRNA.
Ubac1	Ubac1.cSep08	362087	8809	646	3	56	putative protein of vertebrate origin (Ubac1) alternative variant cSep08, mRNA.
Ubap1	Ubap1.aSep08	362502	40690	2586	2	527	ubiquitin-associated protein 1 (Ubap1) alternative variant aSep08, complete mRNA.
Ubap2	Ubap2.aSep08	313169	41010	3869	24	1009	ubiquitin-associated protein 2 (Ubap2) alternative variant aSep08, mRNA.
Ubap2	Ubap2.bSep08	313169	1285	769	5	256	ubiquitin-associated protein 2 (Ubap2) alternative variant bSep08, mRNA.
Ubap2	Ubap2.dSep08	313169	1715	422	2	71	ubiquitin-associated protein 2 (Ubap2) alternative variant dSep08, mRNA.
Ubash3b	Ubash3b.aSep08	315579	143447	1801	2	463	ubiquitin-associated (Ubash3b) alternative variant aSep08, mRNA.
Ubash3b	Ubash3b.bSep08	315579	112594	781	2	245	ubiquitin-associated (Ubash3b) alternative variant bSep08, mRNA.
Ubash3b	Ubash3b.cSep08	315579	4277	1576	3	107	putative protein of bilateral origin (Ubash3b) alternative variant cSep08, mRNA.
Ubc	Ubc.aSep08	50522	4958	3225	2	962	ubiquitin C (108.2 kD) (Ubc) alternative variant aSep08, mRNA.
Ubc	Ubc.bSep08	50522	4737	724	3	202	ubiquitin C (22.7 kD) (Ubc) alternative variant bSep08, complete mRNA.
Ubc	Ubc.cSep08	50522	1762	505	2	140	ubiquitin C (Ubc) alternative variant cSep08, mRNA.
Ube1l	Ube1l.bSep08	301000	1255	716	6	238	ubiquitin-activating enzyme E1-like (Ube1l) alternative variant bSep08, mRNA.
Ube1l	Ube1l.cSep08	301000	1183	640	3	153	ubiquitin-activating enzyme E1-like (Ube1l) alternative variant cSep08, mRNA.
Ube1l	Ube1l.dSep08	301000	885	812	2	42	ubiquitin-activating enzyme E1-like (Ube1l) alternative variant dSep08, mRNA.
Ube2a	Ube2a.aSep08	298317	10360	1350		214	ubiquitin-conjugating enzyme E2A, RAD6 homolog (S. cerevisiae) (Ube2a) alternative variant aSep08, mRNA.
Ube2b	Ube2b.aSep08	81816	13388	588	7	186	enzyme ubiquitin-conjugating E2 (Ube2b) alternative variant aSep08, mRNA.
Ube2b	Ube2b.cSep08	81816	3894	2539	2	93	ubiquitin-conjugating enzyme E2B (10.5 kD) (Ube2b) alternative variant cSep08, mRNA.
Ube2b	Ube2b.dSep08	81816	5390	796	3	83	enzyme ubiquitin-conjugating E2 (Ube2b) alternative variant dSep08, mRNA.
Ube2b	Ube2b.eSep08	81816	4515	289	3	48	putative protein (Ube2b) alternative variant eSep08, mRNA.
Ube2cbp	Ube2cbp.bSep08	315863	166668	1212		217	ubiquitin-conjugating enzyme E2C binding protein (Ube2cbp) alternative variant bSep08, mRNA.
Ube2d2	Ube2d2.bSep08	641452	9020	1965	3	117	ubiquitin-conjugating enzyme E2D 2 (Ube2d2) alternative variant bSep08, mRNA.

Ube2d2	Ube2d2.cSep08	641452	4978	371	1	65	ubiquitin-conjugating enzyme E2D 2 (Ube2d2) alternative variant cSep08, mRNA.
Ube2d3	Ube2d3.bSep08	81920	28674	1135	8	147	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (16.7 kD) (Ube2d3) alternative variant bSep08, mRNA.
Ube2d3	Ube2d3.cSep08	81920	29156	2136	8	147	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (16.7 kD) (Ube2d3) alternative variant cSep08, mRNA.
Ube2d3	Ube2d3.dSep08	81920	28346	1152	8	147	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (16.7 kD) (Ube2d3) alternative variant dSep08, mRNA.
Ube2d3	Ube2d3.eSep08	81920	27896	753	9	118	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (13.6 kD) (Ube2d3) alternative variant eSep08, mRNA.
Ube2d3	Ube2d3.fSep08	81920	27836	784	7	109	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (12.6 kD) (Ube2d3) alternative variant fSep08, mRNA.
Ube2d3	Ube2d3.gSep08	81920	6853	902	5	87	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (Ube2d3) alternative variant gSep08, mRNA.
Ube2d3	Ube2d3.hSep08	81920	11730	592	4	45	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (Ube2d3) alternative variant hSep08, mRNA.
Ube2d3	Ube2d3.iSep08	81920	2479	391	2	22	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) (Ube2d3) alternative variant iSep08, mRNA.
Ube2e2	Ube2e2.aSep08	361013	279501	1429	3	201	ubiquitin-conjugating enzyme E2E 2 (UBC4/5 homolog, yeast) (22.3 kD) (Ube2e2) alternative variant aSep08, mRNA.
Ube2e3	Ube2e3.bSep08	295686	54296	807	1	268	ubiquitin-conjugating enzyme E2E 3, UBC4/5 homolog (yeast) (Ube2e3) alternative variant bSep08, mRNA.
Ube2e3	Ube2e3.cSep08	295686	53537	730	2	130	ubiquitin-conjugating enzyme E2E 3, UBC4/5 homolog (yeast) (Ube2e3) alternative variant cSep08, mRNA.
Ube2f	Ube2f.aSep08	363284	35654	1304	5	185	ubiquitin-conjugating enzyme E2F (putative) (21.1 kD) (Ube2f) alternative variant aSep08, complete mRNA.
Ube2f	Ube2f.bSep08	363284	35495	1101	5	185	ubiquitin-conjugating enzyme E2F (putative) (21.1 kD) (Ube2f) alternative variant bSep08, complete mRNA.
Ube2f	Ube2f.cSep08	363284	4856	692	1	50	ubiquitin-conjugating enzyme E2F (putative) (Ube2f) alternative variant cSep08, mRNA.
Ube2g1	Ube2g1.bSep08	64631	18967	583	5	128	ubiquitin-conjugating enzyme E2G 1 CRA a (Ube2g1) alternative variant bSep08, mRNA.
Ube2g1	Ube2g1.cSep08	64631	19360	3390	6	124	ubiquitin-conjugating enzyme E2G 1 CRA a (Ube2g1) alternative variant cSep08, mRNA.
Ube2g1	Ube2g1.dSep08	64631	75910	1110	7	99	ubiquitin-conjugating enzyme E2G 1 CRA b (11.4 kD) (Ube2g1) alternative variant dSep08, mRNA.
Ube2g1	Ube2g1.eSep08	64631	41733	632	5	45	putative protein (5.2 kD) (Ube2g1) alternative variant eSep08, mRNA.
Ube2i	Ube2i.aSep08	25573	12745	529	6	176	ubiquitin-conjugating enzyme E2I (Ube2i) alternative variant aSep08, mRNA.

Ube2i	Ube2i.cSep08	25573	10434	390	5	129	ubiquitin-conjugating enzyme E2I (Ube2i) alternative variant cSep08, mRNA.
Ube2i	Ube2i.dSep08	25573	6555	374	4	124	ubiquitin-conjugating enzyme E2I (Ube2i) alternative variant dSep08, mRNA.
Ube2i	Ube2i.eSep08	25573	11656	1079	6	118	ubiquitin-conjugating enzyme E2I (Ube2i) alternative variant eSep08, mRNA.
Ube2i	Ube2i.fSep08	25573	4923	599	3	101	ubiquitin-conjugating enzyme E2I (11.3 kD) (Ube2i) alternative variant fSep08, mRNA.
Ube2i	Ube2i.gSep08	25573	3196	1034	2	88	ubiquitin-conjugating enzyme E2I (9.6 kD) (Ube2i) alternative variant gSep08, mRNA.
Ube2i	Ube2i.iSep08	25573	4722	333	2	36	ubiquitin-conjugating enzyme E2I (Ube2i) alternative variant iSep08, mRNA.
Ube2j1	Ube2j1.aSep08	297961	21186	2105	2	323	ubiquitin-conjugating enzyme E2, J1 (Ube2j1) alternative variant aSep08, mRNA.
Ube2j1	Ube2j1.bSep08	297961	19558	1367	1	296	ubiquitin-conjugating enzyme E2, J1 (Ube2j1) alternative variant bSep08, mRNA.
Ube2j2	Ube2j2.bSep08	298689	14390	2615	7	172	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (19.4 kD) (Ube2j2) alternative variant bSep08, complete mRNA.
Ube2j2	Ube2j2.cSep08	298689	13096	638	5	162	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (Ube2j2) alternative variant cSep08, mRNA.
Ube2j2	Ube2j2.dSep08	298689	13345	1607	7	160	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (18.4 kD) (Ube2j2) alternative variant dSep08, mRNA.
Ube2j2	Ube2j2.eSep08	298689	13077	787	6	160	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (18.4 kD) (Ube2j2) alternative variant eSep08, complete mRNA.
Ube2j2	Ube2j2.fSep08	298689	13046	751	6	156	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (Ube2j2) alternative variant fSep08, mRNA.
Ube2j2	Ube2j2.gSep08	298689	13045	745	5	143	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (Ube2j2) alternative variant gSep08, mRNA.
Ube2j2	Ube2j2.hSep08	298689	1256	417	3	98	ubiquitin-conjugating enzyme E2, J2 homolog (yeast) (Ube2j2) alternative variant hSep08, mRNA.
Ube2k	Ube2k.bSep08	289623	23726	854	5	97	ubiquitin-conjugating enzyme E2-25K precursor (10.7 kD) (Ube2k) alternative variant bSep08, mRNA.
Ube2k	Ube2k.cSep08	289623	20613	336	3	96	enzyme Ubiquitin-Conjugating E2 (Ube2k) alternative variant cSep08, mRNA.
Ube2k	Ube2k.dSep08	289623	55505	381	3	76	ubiquitin-conjugating enzyme E2K (Ube2k) alternative variant dSep08, mRNA.
Ube2k	Ube2k.eSep08	289623	1458	1233	2	61	ubiquitin-conjugating enzyme E2-25K (6.8 kD) (Ube2k) alternative variant eSep08, mRNA.
Ube2l3	Ube2l3.bSep08	363836	39906	403	1	41	ubiquitin-conjugating enzyme E2L 3 (4.8 kD) (Ube2l3) alternative variant bSep08, mRNA.
Ube2m	Ube2m.bSep08	361509	2413	1230	1	87	ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast) (Ube2m) alternative variant bSep08, mRNA.
Ube2n	Ube2n.bSep08	116725	896	326	1	62	ubiquitin-conjugating enzyme E2N (Ube2n) alternative variant bSep08, mRNA.
Ube2o	Ube2o.aSep08	303689	3797	2633		339	ubiquitin-conjugating enzyme E2O (Ube2o) mRNA.
Ube2q1	Ube2q1.bSep08	295252	5745	795	10	264	ubiquitin-conjugating enzyme E2Q (putative) 1 (Ube2q1) alternative variant bSep08, mRNA.

Ube2q1	Ube2q1.cSep08	295252	4861	2745	9	217	ubiquitin-conjugating enzyme E2Q (putative) 1 (24.3 kD) (Ube2q1) alternative variant cSep08, mRNA.
Ube2q1	Ube2q1.eSep08	295252	1361	420	4	71	ubiquitin-conjugating enzyme E2Q (putative) 1 (8.0 kD) (Ube2q1) alternative variant eSep08, mRNA.
Ube2q1	Ube2q1.fSep08	295252	2026	1416	2	97	ubiquitin-conjugating enzyme E2Q (putative) 1 (10.7 kD) (Ube2q1) alternative variant fSep08, mRNA.
Ube2q2	Ube2q2.aSep08	363065	61136	1285	2	356	ubiquitin-conjugating enzyme E2Q (putative) 2 (Ube2q2) alternative variant aSep08, mRNA.
Ube2q2	Ube2q2.bSep08	363065	44402	947	2	315	ubiquitin-conjugating enzyme E2Q (putative) 2 (Ube2q2) alternative variant bSep08, mRNA.
Ube2q2	Ube2q2.cSep08	363065	36150	706	1	235	ubiquitin-conjugating enzyme E2Q (putative) 2 (Ube2q2) alternative variant cSep08, mRNA.
Ube2q2	Ube2q2.dSep08	363065	3919	1635		32	ubiquitin-conjugating enzyme E2Q (putative) 2 (Ube2q2) alternative variant dSep08, mRNA.
Ube2t	Ube2t.bSep08	360847	5457	570	4	75	ubiquitin-conjugating enzyme E2T (putative) (Ube2t) alternative variant bSep08, mRNA.
Ube2v1	Ube2v1.bSep08	296390	7774	449	1	53	ubiquitin-conjugating enzyme E2 variant 1 (Ube2v1) alternative variant bSep08, mRNA.
Ube2v2	Ube2v2.aSep08	287927	31010	757	4	240	ubiquitin-conjugating enzyme E2 variant 2 (Ube2v2) alternative variant aSep08, mRNA.
Ube2z	Ube2z.aSep08	303478	19140	2875	7	374	ubiquitin-conjugating enzyme E2Z (putative) (Ube2z) alternative variant aSep08, mRNA.
Ube3a	Ube3a.aSep08	361585	27493	1145		241	ubiquitin protein ligase E3A (27.9 kD) (Ube3a) mRNA.
Ube3c	Ube3c.aSep08	362294	102007	2024	6	557	ubiquitin protein ligase E3C (Ube3c) alternative variant aSep08, mRNA.
Ube3c	Ube3c.bSep08	362294	56898	3347	11	535	ubiquitin protein ligase E3C (Ube3c) alternative variant bSep08, mRNA.
Ube3c	Ube3c.cSep08	362294	16603	474	3	158	ubiquitin protein ligase E3C (Ube3c) alternative variant cSep08, mRNA.
Ube4a	Ube4a.bSep08	315608	4115	685	4	190	ubiquitination factor E4A, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4a) alternative variant bSep08, mRNA.
Ube4a	Ube4a.cSep08	315608	12268	543	4	135	ubiquitination factor E4A, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4a) alternative variant cSep08, mRNA.
Ube4a	Ube4a.eSep08	315608	7422	909	3	34	ubiquitination factor E4A, UFD2 homolog ( <i>S. cerevisiae</i> ) (3.7 kD) (Ube4a) alternative variant eSep08, mRNA.
Ube4a	Ube4a.hSep08	315608	2112	352	2	28	ubiquitination factor E4A, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4a) alternative variant hSep08, mRNA.
Ube4b	Ube4b.aSep08	298652	45523	3834	20	855	ubiquitination factor E4B, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4b) alternative variant aSep08, mRNA.
Ube4b	Ube4b.bSep08	298652	7931	528	3	176	ubiquitination factor E4B, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4b) alternative variant bSep08, mRNA.
Ube4b	Ube4b.cSep08	298652	10127	534	5	171	ubiquitination factor E4B, UFD2 homolog ( <i>S. cerevisiae</i> ) (Ube4b) alternative variant cSep08, mRNA.
Ubfd1	Ubfd1.aSep08	293454	10263	2423	6	313	ubiquitin (Ubfd1) alternative variant aSep08, mRNA.
Ubfd1	Ubfd1.cSep08	293454	1506	709	2	236	ubiquitin-binding protein homolog like (Ubfd1) alternative variant cSep08, mRNA.

Ubiad1	Ubiad1.bSep08	313706	23430	929	1	191	putative endoplasmic reticulum protein, with a transmembrane domain, of ancient origin (21.0 kD) (Ubiad1) alternative variant bSep08, mRNA.
ubiquitin.3	ubiquitin.3.aSep08		78155	602		68	parkin (ubiquitin.3) mRNA.
Ubl5	Ubl5.aSep08	500954	1902	568	5	73	ubiquitin-like 5 (8.5 kD) (Ubl5) alternative variant aSep08, complete mRNA.
Ubl5	Ubl5.cSep08	500954	1689	1543	2	47	ubiquitin-like 5 (5.4 kD) (Ubl5) alternative variant cSep08, complete mRNA.
Ubl5	Ubl5.dSep08	500954	1048	604	3	47	ubiquitin-like 5 (5.4 kD) (Ubl5) alternative variant dSep08, complete mRNA.
Ubl7	Ubl7.bSep08	300744	10207	950	9	316	ubiquitin-like 7 (bone marrow stromal cell-derived) (Ubl7) alternative variant bSep08, mRNA.
Ubl7	Ubl7.cSep08	300744	9600	782	7	233	ubiquitin-like 7 (bone marrow stromal cell-derived) (Ubl7) alternative variant cSep08, mRNA.
Ubl7	Ubl7.dSep08	300744	9827	741	8	232	ubiquitin-like 7 (bone marrow stromal cell-derived) (Ubl7) alternative variant dSep08, mRNA.
Ubl7	Ubl7.eSep08	300744	8261	563	6	187	ubiquitin-like 7 (bone marrow stromal cell-derived) (Ubl7) alternative variant eSep08, mRNA.
Ublcp1	Ublcp1.bSep08	360514	8299	750	8	216	ubiquitin (Ublcp1) alternative variant bSep08, mRNA.
Ubn1	Ubn1.bSep08	302935	6829	896	3	248	ubiquitin 1 (Ubn1) alternative variant bSep08, mRNA.
Ubn1	Ubn1.cSep08	302935	6315	329	3	109	ubiquitin 1 (Ubn1) alternative variant cSep08, mRNA.
Ubp1	Ubp1.aSep08	301038	46332	3956	17	619	upstream binding protein 1 (Ubp1) alternative variant aSep08, mRNA.
Ubp1	Ubp1.bSep08	301038	27111	1742	4	580	upstream binding protein 1 (Ubp1) alternative variant bSep08, mRNA.
Ubp1	Ubp1.cSep08	301038	16085	1024	9	321	upstream binding protein 1 (Ubp1) alternative variant cSep08, mRNA.
Ubp1	Ubp1.dSep08	301038	24420	549	4	182	upstream binding protein 1 (Ubp1) alternative variant dSep08, mRNA.
Ubp1	Ubp1.eSep08	301038	14047	330	3	109	upstream binding protein 1 (Ubp1) alternative variant eSep08, mRNA.
Ubp1	Ubp1.fSep08	301038	4515	2711	2	87	upstream binding protein 1 (9.3 kD) (Ubp1) alternative variant fSep08, mRNA.
Ubqln1	Ubqln1.bSep08	114590	3156	842	3	159	ubiquilin 1 (Ubqln1) alternative variant bSep08, mRNA.
Ubqln1	Ubqln1.cSep08	114590	3447	2190	2	144	ubiquilin 1 CRA b (14.8 kD) (Ubqln1) alternative variant cSep08, mRNA.
Ubtd2	Ubtd2.aSep08	287178	16627	1799		211	dendritic cell-derived ubiquitin-like protein (Ubtd2) mRNA.
Ubtf	Ubtf.cSep08	25574	3882	854	5	186	upstream binding transcription factor RNA polymerase I like (21.7 kD) (Ubtf) alternative variant cSep08, mRNA.
Ubtf	Ubtf.dSep08	25574	1272	1178	2	150	putative mitochondrial protein (16.8 kD) (Ubtf) alternative variant dSep08, mRNA.
Ubtf	Ubtf.eSep08	25574	1989	706	5	137	upstream binding transcription factor RNA polymerase I like (Ubtf) alternative variant eSep08, mRNA.
Ubtf	Ubtf.hSep08	25574	2839	352	4	83	upstream binding transcription factor RNA polymerase I CRA b like (Ubtf) alternative variant hSep08, mRNA.

Ubx1	Ubx1.bSep08	363332	4769	1519	8	383	UBX domain protein 6 (Ubx1) alternative variant bSep08, mRNA.
Ubx1	Ubx1.cSep08	363332	4755	1974	8	208	UBX domain (23.0 kD) (Ubx1) alternative variant cSep08, complete mRNA.
Ubx1	Ubx1.dSep08	363332	3176	747	4	137	UBX domain (15.8 kD) (Ubx1) alternative variant dSep08, mRNA.
Ubx1	Ubx1.eSep08	363332	1253	398	1	132	UBX domain (Ubx1) alternative variant eSep08, mRNA.
Ubx2	Ubx2.bSep08	304766	32047	1997	12	351	UBX domain (39.5 kD) (Ubx2) alternative variant bSep08, mRNA.
Ubx2	Ubx2.cSep08	304766	15513	974	8	324	UBX domain (Ubx2) alternative variant cSep08, mRNA.
Ubx2	Ubx2.dSep08	304766	11748	741	5	219	UBX domain (Ubx2) alternative variant dSep08, mRNA.
Ubx2	Ubx2.eSep08	304766	2049	659	2	37	putative protein (4.3 kD) (Ubx2) alternative variant eSep08, mRNA.
Ubx4andRGD1561716	Ubx4andRGD1561716.bSep08	500624	25536	1486	6	196	UBX domain (22.4 kD) (Ubx4andRGD1561716) alternative variant bSep08, mRNA.
Ubx4andRGD1561716	Ubx4andRGD1561716.bSep08	685859	25536	1486	6	196	UBX domain (22.4 kD) (Ubx4andRGD1561716) alternative variant bSep08, mRNA.
Ubx5	Ubx5.bSep08	192207	14721	807	8	268	UBX domain (Ubx5) alternative variant bSep08, mRNA.
Ubx5	Ubx5.cSep08	192207	12590	819	7	164	UBX domain (18.1 kD) (Ubx5) alternative variant cSep08, mRNA.
Ubx5	Ubx5.dSep08	192207	835	674	3	149	UBX domain (16.3 kD) (Ubx5) alternative variant dSep08, mRNA.
Ubx5	Ubx5.eSep08	192207	3730	589	4	126	putative protein, with a coiled coil domain, of mammalian origin (14.0 kD) (Ubx5) alternative variant eSep08, mRNA.
Ubx5	Ubx5.fSep08	192207	20298	433	6	116	UBX domain (13.0 kD) (Ubx5) alternative variant fSep08, mRNA.
Ubx6	Ubx6.bSep08	290802	12136	708	1	146	reproduction 8 like (Ubx6) alternative variant bSep08, mRNA.
Ubx7	Ubx7.aSep08	303878	58006	1070	7	356	CRA a (Ubx7) alternative variant aSep08, mRNA.
UCH.0	UCH.0.aSep08		4217	2008		122	ubiquitin specific peptidase 12 (UCH.0) mRNA.
Uchl1	Uchl1.bSep08	29545	3046	505	3	123	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (Uchl1) alternative variant bSep08, mRNA.
Uchl1	Uchl1.dSep08	29545	1465	345	2	90	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (Uchl1) alternative variant dSep08, mRNA.
Uchl1	Uchl1.eSep08	29545	585	473	2	65	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (7.0 kD) (Uchl1) alternative variant eSep08, mRNA.
Uchl5	Uchl5.aSep08	360853	36791	1294	6	363	ubiquitin carboxyl-terminal hydrolase L5 (Uchl5) alternative variant aSep08, mRNA.
Uchl5	Uchl5.cSep08	360853	36767	1159	5	291	ubiquitin carboxyl-terminal hydrolase L5 (33.3 kD) (Uchl5) alternative variant cSep08, complete mRNA.
Uchl5	Uchl5.dSep08	360853	11951	815	2	204	ubiquitin carboxyl-terminal hydrolase L5 (Uchl5) alternative variant dSep08, mRNA.
Uchl5ip	Uchl5ip.aSep08	293844	19083	1366	8	368	UCHL5 interacting protein (40.6 kD) (Uchl5ip) alternative variant aSep08, complete mRNA.



Uchl5ip	Uchl5ip.bSep08	293844	8981	769	4	183	UCHL5 interacting protein (20.6 kD) (Uchl5ip) alternative variant bSep08, mRNA.
Uchl5ip	Uchl5ip.cSep08	293844	5165	911	2	143	UCHL5 interacting protein (Uchl5ip) alternative variant cSep08, mRNA.
Uck1	Uck1.aSep08	311864	2369	1275	4	360	uridine-cytidine kinase 1 (Uck1) alternative variant aSep08, complete mRNA.
Uck1	Uck1.bSep08	311864	5840	1889	7	283	uridine-cytidine kinase 1 (31.7 kD) (Uck1) alternative variant bSep08, complete mRNA.
Uck1	Uck1.cSep08	311864	3390	679	4	194	uridine-cytidine kinase 1 (Uck1) alternative variant cSep08, mRNA.
Uck1	Uck1.dSep08	311864	526	362	2	107	uridine-cytidine kinase 1 (11.5 kD) (Uck1) alternative variant dSep08, mRNA.
Uck1	Uck1.eSep08	311864	3461	989	4	87	uridine-cytidine kinase 1 (Uck1) alternative variant eSep08, mRNA.
Uck2	Uck2.bSep08	304944	52785	497	1	144	uridine-cytidine kinase 2 (Uck2) alternative variant bSep08, mRNA.
Uckl1	Uckl1.bSep08	499956	1067	850	3	94	uridine-cytidine kinase 1-like 1 (10.7 kD) (Uckl1) alternative variant bSep08, mRNA.
Ucn3	Ucn3.aSep08	498791	6256	1164	2	245	urocortin 3 (stresscopin) (Ucn3) alternative variant aSep08, mRNA.
Ucp1	Ucp1.bSep08	24860	1272	609	1	126	uncoupling protein 1 (mitochondrial, proton carrier) (13.4 kD) (Ucp1) alternative variant bSep08, mRNA.
Ucp3	Ucp3.bSep08	25708	7269	734	1	196	uncoupling protein 3 (mitochondrial, proton carrier) (Ucp3) alternative variant bSep08, mRNA.
uDENN.0	uDENN.0.aSep08		6976	434		93	CRA b (uDENN.0) mRNA.
uDENN.1	uDENN.1.aSep08		4304	1139	4	242	containing DENN MADD domain 4 (26.4 kD) (uDENN.1) alternative variant aSep08, mRNA.
uDENN.1	uDENN.1.bSep08		2789	919	1	122	putative cytoplasmic protein of metazoan origin (12.9 kD) (uDENN.1) alternative variant bSep08, mRNA.
uDENN.2	uDENN.2.aSep08		1486	372		124	protein CRA e (uDENN.2) mRNA.
UEV.0	UEV.0.aSep08		7288	353		117	ubiquitin-conjugating enzyme E2-like (UEV.0) mRNA.
Uevld	Uevld.aSep08	691172	5311	408		135	UEV and lactate/malate dehydrogenase domains (Uevld) mRNA.
Ufc1	Ufc1.bSep08	445268	5457	391	2	93	ubiquitin-fold modifier conjugating enzyme 1 (11.4 kD) (Ufc1) alternative variant bSep08, mRNA.
Ufc1	Ufc1.cSep08	445268	5819	889	1	93	ubiquitin-fold modifier conjugating enzyme 1 (11.4 kD) (Ufc1) alternative variant cSep08, mRNA.
Ufc1	Ufc1.dSep08	445268	6630	985	4	87	ubiquitin-fold modifier conjugating enzyme 1 (9.6 kD) (Ufc1) alternative variant dSep08, mRNA.
Ufd1l	Ufd1l.bSep08	84478	16127	731	9	177	ubiquitin fusion degradation 1 like (yeast) (Ufd1l) alternative variant bSep08, mRNA.
Ufm1	Ufm1.bSep08	365797	6695	690	6	93	ubiquitin-fold modifier 1 (10.3 kD) (Ufm1) alternative variant bSep08, mRNA.
Ufsp2	Ufsp2.bSep08	361151	10476	869	7	284	peptidase 2 (Ufsp2) alternative variant bSep08, mRNA.
Ufsp2	Ufsp2.cSep08	361151	14082	2219	7	183	peptidase 2 (21.2 kD) (Ufsp2) alternative variant cSep08, complete mRNA.

Ufsp2	Ufsp2.dSep08	361151	9305	909	6	168	peptidase 2 (Ufsp2) alternative variant dSep08, mRNA.
Ufsp2	Ufsp2.eSep08	361151	9303	738	7	164	peptidase 2 (Ufsp2) alternative variant eSep08, mRNA.
Ufsp2	Ufsp2.fSep08	361151	8001	715	6	128	peptidase 2 (Ufsp2) alternative variant fSep08, mRNA.
Ufsp2	Ufsp2.gSep08	361151	7986	646	6	128	peptidase 2 (Ufsp2) alternative variant gSep08, mRNA.
Ufsp2	Ufsp2.hSep08	361151	2969	569	3	91	peptidase 2 (10.1 kD) (Ufsp2) alternative variant hSep08, mRNA.
Ugcg	Ugcg.bSep08	83626	3206	262	1	63	UDP-glucose ceramide glucosyltransferase (Ugcg) alternative variant bSep08, mRNA.
Ugcgl1	Ugcgl1.bSep08	171129	8738	487	3	148	UDP-glucose ceramide glucosyltransferase-like 1 (Ugcgl1) alternative variant bSep08, mRNA.
Ugcgl1	Ugcgl1.cSep08	171129	4154	338	4	112	UDP-glucose ceramide glucosyltransferase-like 1 (Ugcgl1) alternative variant cSep08, mRNA.
Ugdh	Ugdh.bSep08	83472	1374	520	2	100	UDP-glucose dehydrogenase (11.4 kD) (Ugdh) alternative variant bSep08, mRNA.
Ugp2	Ugp2.bSep08	289827	32626	1081	6	360	UDP-glucose pyrophosphorylase 2 (Ugp2) alternative variant bSep08, mRNA.
Ugp2	Ugp2.cSep08	289827	30333	742	5	243	UDP-glucose pyrophosphorylase 2 (Ugp2) alternative variant cSep08, mRNA.
Ugp2	Ugp2.dSep08	289827	2032	832	2	129	udp-glucose pyrophosphorylase 2 (14.4 kD) (Ugp2) alternative variant dSep08, mRNA.
Ugt2a3	Ugt2a3.aSep08	289533	6037	641		213	UDP glucuronosyltransferase 2 family, polypeptide A3 (Ugt2a3) mRNA.
Ugt2b5	Ugt2b5.aSep08	29623	16269	1532		510	UDP-glucuronosyltransferase 2 family, member 5 (Ugt2b5) mRNA.
Ugt2b36	Ugt2b36.cSep08	83808	14887	1076	3	78	UDP glucuronosyltransferase 2 family, polypeptide B36 (Ugt2b36) alternative variant cSep08, mRNA.
Uhmk1	Uhmk1.bSep08	246332	5280	740	4	192	U2AF homology motif (UHM) kinase 1 (Uhmk1) alternative variant bSep08, mRNA.
Uhmk1	Uhmk1.cSep08	246332	1813	202	2	36	U2AF homology motif (UHM) kinase 1 (Uhmk1) alternative variant cSep08, mRNA.
Uhrf1	Uhrf1.bSep08	316129	18532	3557	13	782	ubiquitin-like with PHD and ring finger domains 1 (88.2 kD) (Uhrf1) alternative variant bSep08, mRNA.
Uhrf1	Uhrf1.cSep08	316129	7584	791	2	185	ubiquitin-like with PHD and ring finger domains 1 (Uhrf1) alternative variant cSep08, mRNA.
Uhrf1	Uhrf1.dSep08	316129	2318	431	2	143	ubiquitin-like with PHD and ring finger domains 1 (Uhrf1) alternative variant dSep08, mRNA.
Uhrf1bp1l	Uhrf1bp1l.bSep08	363009	1591	1067	2	355	UHRF1 binding protein 1-like (Uhrf1bp1l) alternative variant bSep08, mRNA.
Uhrf1bp1l	Uhrf1bp1l.cSep08	363009	9809	2765	6	325	UHRF1 binding protein 1-like (Uhrf1bp1l) alternative variant cSep08, mRNA.
Uhrf1bp1l	Uhrf1bp1l.dSep08	363009	9011	642	4	213	UHRF1 binding protein 1-like (Uhrf1bp1l) alternative variant dSep08, mRNA.
Uhrf1bp1l	Uhrf1bp1l.eSep08	363009	1557	845	2	123	UHRF1 binding protein 1-like (13.7 kD) (Uhrf1bp1l) alternative variant eSep08, mRNA.
Uhrf2	Uhrf2.bSep08	309331	9177	2564	8	265	ubiquitin-like, containing PHD and RING finger domains 2 (30.4 kD) (Uhrf2) alternative variant bSep08, mRNA.

Uhrf2	Uhrf2.cSep08	309331	7386	685	6	217	ubiquitin-like, containing PHD and RING finger domains 2 (Uhrf2) alternative variant cSep08, mRNA.
Uhrf2	Uhrf2.dSep08	309331	2599	734	3	129	ubiquitin-like, containing PHD and RING finger domains 2 (Uhrf2) alternative variant dSep08, mRNA.
Uhrf2	Uhrf2.eSep08	309331	4695	753	4	71	ubiquitin-like, containing PHD and RING finger domains 2 (8.2 kD) (Uhrf2) alternative variant eSep08, mRNA.
Uimc1	Uimc1.bSep08	290997	21546	853	7	284	retinoid x receptor interacting protein (Uimc1) alternative variant bSep08, mRNA.
Uimc1	Uimc1.cSep08	290997	24307	643	5	166	retinoid x receptor interacting protein (18.7 kD) (Uimc1) alternative variant cSep08, complete mRNA.
Uimc1	Uimc1.dSep08	290997	23038	561	4	144	retinoid x receptor interacting protein (16.6 kD) (Uimc1) alternative variant dSep08, complete mRNA.
Uimc1	Uimc1.fSep08	290997	7674	1464	4	58	retinoid x receptor interacting protein (6.6 kD) (Uimc1) alternative variant fSep08, mRNA.
Ulk1	Ulk1.bSep08	360827	1529	891	2	65	kinase 1 (Ulk1) alternative variant bSep08, mRNA.
Ulk2	Ulk2.aSep08	303206	22662	3427	1	382	kinase 2 (Ulk2) alternative variant aSep08, mRNA.
Ulk3	Ulk3.aSep08	691171	4010	1585		384	kinase 3 (43.7 kD) (Ulk3) mRNA.
Umod	Umod.bSep08	25128	7231	1798	2	376	uromodulin (Umod) alternative variant bSep08, mRNA.
Umodl1	Umodl1.aSep08	365544	9468	1316		133	uromodulin-like 1 (Umodl1) mRNA.
Unc5a	Unc5a.bSep08	60629	5992	929	4	192	unc-5 homolog A CRA a (Unc5a) alternative variant bSep08, mRNA.
Unc5a	Unc5a.cSep08	60629	804	720	2	123	unc-5 homolog A CRA a (Unc5a) alternative variant cSep08, mRNA.
Unc5c	Unc5c.bSep08	362049	2995	359	2	85	unc-5 homolog c (Unc5c) alternative variant bSep08, mRNA.
Unc13a	Unc13a.aSep08	64829	6969	2845		217	unc-13 homolog (Unc13a) mRNA.
Unc13b	Unc13b.bSep08	64830	4963	1411	8	343	unc-13 homolog B (Unc13b) alternative variant bSep08, mRNA.
Unc13b	Unc13b.cSep08	64830	51295	403	5	134	unc-13 homolog B CRA a (Unc13b) alternative variant cSep08, mRNA.
Unc45a	Unc45a.bSep08	308759	5449	761	7	234	smooth muscle cell associated (Unc45a) alternative variant bSep08, mRNA.
Unc45a	Unc45a.cSep08	308759	4890	915	6	218	smooth muscle cell associated (Unc45a) alternative variant cSep08, mRNA.
Unc45a	Unc45a.dSep08	308759	736	376	3	67	putative protein of mammalian origin (Unc45a) alternative variant dSep08, mRNA.
Unc45b	Unc45b.bSep08	303373	9019	917	2	305	unc-45 homolog b (Unc45b) alternative variant bSep08, mRNA.
Unc50	Unc50.bSep08	192356	7884	2131	3	251	unc-50 homolog CRA b (29.5 kD) (Unc50) alternative variant bSep08, complete mRNA.
Unc50	Unc50.cSep08	192356	7093	1320	4	226	unc-50 homolog CRA b (26.9 kD) (Unc50) alternative variant cSep08, mRNA.
Unc50	Unc50.dSep08	192356	6298	686	3	67	putative protein (7.5 kD) (Unc50) alternative variant dSep08, mRNA.
Unc84a	Unc84a.bSep08	360773	30368	1322	12	440	unc-84 homolog A CRA a (Unc84a) alternative variant bSep08, mRNA.

Unc84a	Unc84a.cSep08	360773	34947	1059	11	307	unc-84 homolog A CRA a (Unc84a) alternative variant cSep08, mRNA.
Unc84a	Unc84a.dSep08	360773	11933	702	6	220	unc-84 homolog A CRA a (Unc84a) alternative variant dSep08, mRNA.
Unc84a	Unc84a.eSep08	360773	22827	790	7	220	unc-84 homolog A CRA a (Unc84a) alternative variant eSep08, mRNA.
Unc84a	Unc84a.fSep08	360773	22798	789	7	215	unc-84 homolog A CRA a (Unc84a) alternative variant fSep08, mRNA.
Unc84a	Unc84a.gSep08	360773	22823	604	5	201	unc-84 homolog A CRA a (Unc84a) alternative variant gSep08, mRNA.
Unc84a	Unc84a.hSep08	360773	2383	937	5	153	unc-84 homolog A CRA a (Unc84a) alternative variant hSep08, mRNA.
Unc84a	Unc84a.iSep08	360773	4887	412	5	137	unc-84 homolog A CRA d (Unc84a) alternative variant iSep08, mRNA.
Unc84a	Unc84a.jSep08	360773	21394	489	4	124	unc-84 homolog A CRA a (Unc84a) alternative variant jSep08, mRNA.
Unc84a	Unc84a.kSep08	360773	19002	353	3	117	unc-84 homolog A CRA a (Unc84a) alternative variant kSep08, mRNA.
Unc84b	Unc84b.aSep08	315135	17158	4034	17	625	unc-84 homolog B (Unc84b) alternative variant aSep08, mRNA.
Unc84b	Unc84b.bSep08	315135	4546	1317	8	313	unc-84 homolog B (Unc84b) alternative variant bSep08, mRNA.
Unc93b1	Unc93b1.bSep08	361689	4846	1066	8	354	unc-93 homolog B1 CRA b (Unc93b1) alternative variant bSep08, mRNA.
Unc93b1	Unc93b1.cSep08	361689	27493	749	6	249	unc-93 homolog B1 CRA b (Unc93b1) alternative variant cSep08, mRNA.
Unc93b1	Unc93b1.dSep08	361689	1439	716	3	153	unc-93 homolog B1 (Unc93b1) alternative variant dSep08, mRNA.
Unc93b1	Unc93b1.fSep08	361689	3829	639	5	104	unc-93 homolog B1 CRA b (Unc93b1) alternative variant fSep08, mRNA.
Unc119	Unc119.cSep08	29402	1159	898	2	80	unc-119 homolog CRA a (9.6 kD) (Unc119) alternative variant cSep08, mRNA.
Unc119	Unc119.dSep08	29402	897	701	2	65	unc-119 homolog CRA e (Unc119) alternative variant dSep08, mRNA.
Ung	Ung.bSep08	304577	6770	748	6	211	uracil-DNA glycosylase (Ung) alternative variant bSep08, mRNA.
Ung	Ung.cSep08	304577	1556	669	3	140	uracil-DNA glycosylase (Ung) alternative variant cSep08, mRNA.
Ung	Ung.dSep08	304577	5980	738	6	120	uracil-DNA glycosylase (Ung) alternative variant dSep08, mRNA.
Unk	Unk.bSep08	360663	3371	725	4	241	zinc finger ccch type containing 5 CRA a (Unk) alternative variant bSep08, mRNA.
Unk	Unk.cSep08	360663	2458	1830	2	164	zinc finger ccch type containing 5 CRA a (18.1 kD) (Unk) alternative variant cSep08, mRNA.
Unk	Unk.dSep08	360663	2359	375	4	124	zinc finger ccch type containing 5 CRA a (Unk) alternative variant dSep08, mRNA.

Unk	Unk.eSep08	360663	955	390	2	91	zinc finger ccch type containing 5 CRA a (Unk) alternative variant eSep08, mRNA.
Upb1	Upb1.bSep08	116593	9744	773	5	186	ureidopropionase, beta (Upb1) alternative variant bSep08, mRNA.
Upb1	Upb1.cSep08	116593	2208	1393	2	105	ureidopropionase, beta (Upb1) alternative variant cSep08, mRNA.
Upb1	Upb1.dSep08	116593	7596	1777	3		
Upb1	Upb1.eSep08	116593	8475	554	4	47	ureidopropionase, beta (5.5 kD) (Upb1) alternative variant eSep08, mRNA.
Upf3b	Upf3b.aSep08	313449	9906	1771	6	316	UPF3 regulator of nonsense transcripts homolog B (yeast) (Upf3b) alternative variant aSep08, mRNA.
Upf3b	Upf3b.cSep08	313449	4791	347	5	29	UPF3 regulator of nonsense transcripts homolog B (yeast) (3.6 kD) (Upf3b) alternative variant cSep08, mRNA.
UPF0560.1	UPF0560.1.aSep08		8694	1658	8	552	putative protein (UPF0560.1) alternative variant aSep08, mRNA.
UPF0560.1	UPF0560.1.bSep08		2842	1042	4	181	putative protein of vertebrate origin (UPF0560.1) alternative variant bSep08, mRNA.
UPF0560.1	UPF0560.1.eSep08		6817	371	4	101	putative protein (UPF0560.1) alternative variant eSep08, mRNA.
UPF0560.1	UPF0560.1.fSep08		3018	305	4	80	family with sequence similarity 171 member A1 (UPF0560.1) alternative variant fSep08, mRNA.
Upk2	Upk2.bSep08	689093	1414	605		86	uroplakin 2 (Upk2) alternative variant bSep08, mRNA.
Upk3b	Upk3b.aSep08	360790	5993	1783		535	uroplakin 3B (Upk3b) mRNA.
Uprt	Uprt.aSep08	317237	20053	701		221	uracil phosphoribosyltransferase (FUR1) homolog (S. cerevisiae) (Uprt) mRNA.
Uqcr	Uqcr.aSep08	690848	4641	494	3	89	ubiquinol-cytochrome c reductase, 6.4kDa subunit (9.8 kD) (Uqcr) alternative variant aSep08, complete mRNA.
Uqcr	Uqcr.dSep08	690848	534	419	2	39	ubiquinol-cytochrome c reductase, 6.4kDa subunit (Uqcr) alternative variant dSep08, mRNA.
Uqcrbl	Uqcrbl.bSep08	362897	4801	1058	1	90	ubiquinol-cytochrome c reductase binding protein-like (10.8 kD) (Uqcrbl) alternative variant bSep08, complete mRNA.
Uqcrc1	Uqcrc1.bSep08	301011	8988	1042	8	339	ubiquinol-cytochrome c reductase core protein I (Uqcrc1) alternative variant bSep08, mRNA.
Uqcrc1	Uqcrc1.cSep08	301011	855	435	2	68	putative protein (7.4 kD) (Uqcrc1) alternative variant cSep08, mRNA.
Uqcrc2	Uqcrc2.bSep08	293448	14196	644	7	158	cytochrome Bc1 complex (Uqcrc2) alternative variant bSep08, mRNA.
Uqcrc2	Uqcrc2.cSep08	293448	6279	618	5	116	core protein II (12.6 kD) (Uqcrc2) alternative variant cSep08, complete mRNA.
Uqcrc2	Uqcrc2.dSep08	293448	5446	793	4	113	core protein II (12.2 kD) (Uqcrc2) alternative variant dSep08, complete mRNA.
Uqcrc2	Uqcrc2.eSep08	293448	3189	700	3	109	ubiquinol cytochrome c reductase core protein 2 CRA a (Uqcrc2) alternative variant eSep08, mRNA.
Uqcrq	Uqcrq.bSep08	497902	2297	415	3	82	ubiquinol-cytochrome c reductase, complex III subunit VII (9.8 kD) (Uqcrq) alternative variant bSep08, mRNA.
UQ_con.1	UQ_con.1.aSep08		95405	4250	6	192	ubiquitin-conjugating enzyme (UQ_con.1) alternative variant aSep08, mRNA.

UQ_con.1	UQ_con.1.bSep08		1530	430	1	78	ubiquitin-conjugating enzyme E2 H like (UQ_con.1) alternative variant bSep08, mRNA.
Urg4	Urg4.bSep08	305493	21958	381	6	127	up-regulated gene 4 (Urg4) alternative variant bSep08, mRNA.
Urg4	Urg4.cSep08	305493	53278	595	6	82	up-regulated gene 4 (Urg4) alternative variant cSep08, mRNA.
Urg4	Urg4.dSep08	305493	790	455	2	44	up-regulated gene 4 (Urg4) alternative variant dSep08, mRNA.
Urg4	Urg4.eSep08	305493	46149	315	3	82	up-regulated gene 4 (Urg4) alternative variant eSep08, mRNA.
Urm1	Urm1.aSep08	311840	17284	1235	3	101	ubiquitin related modifier 1 homolog (S. cerevisiae) (11.3 kD) (Urm1) alternative variant aSep08, complete mRNA.
Urm1	Urm1.bSep08	311840	14472	611	1	63	ubiquitin related modifier 1 homolog (S. cerevisiae) (7.1 kD) (Urm1) alternative variant bSep08, mRNA.
Uroc1	Uroc1.aSep08	685999	7297	1732		155	putative protein (17.5 kD) (Uroc1) alternative variant aSep08, mRNA.
Urod	Urod.bSep08	29421	1995	837	6	218	uroporphyrinogen decarboxylase (24.2 kD) (Urod) alternative variant bSep08, complete mRNA.
Urod	Urod.cSep08	29421	1402	1305	2	138	uroporphyrinogen decarboxylase (15.3 kD) (Urod) alternative variant cSep08, mRNA.
Urod	Urod.dSep08	29421	2194	1241	4	110	uroporphyrinogen decarboxylase (12.3 kD) (Urod) alternative variant dSep08, mRNA.
Uros	Uros.bSep08	309070	8392	403	4	81	uroporphyrinogen III synthase (Uros) alternative variant bSep08, mRNA.
Use1	Use1.aSep08	290627	2324	749	6	242	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (Use1) alternative variant aSep08, mRNA.
Use1	Use1.bSep08	290627	2695	829	7	241	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (27.3 kD) (Use1) alternative variant bSep08, complete mRNA.
Use1	Use1.cSep08	290627	2693	824	7	240	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (27.2 kD) (Use1) alternative variant cSep08, complete mRNA.
Use1	Use1.dSep08	290627	2673	1388	5	148	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (16.4 kD) (Use1) alternative variant dSep08, complete mRNA.
Use1	Use1.eSep08	290627	2620	687	7	139	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (Use1) alternative variant eSep08, mRNA.
Use1	Use1.fSep08	290627	2209	818	6	123	uncharacterized Hematopoietic stem progenitor cells protein MDS032 (13.8 kD) (Use1) alternative variant fSep08, mRNA.
Use1	Use1.gSep08	290627	2685	1791	2	84	CRA c (9.7 kD) (Use1) alternative variant gSep08, complete mRNA.
Usf1	Usf1.bSep08	83586	6895	964	1	222	upstream transcription factor 1 (Usf1) alternative variant bSep08, mRNA.
Usf2	Usf2.bSep08	81817	9788	756	9	251	upstream transcription factor 2 (Usf2) alternative variant bSep08, mRNA.

Usf2	Usf2.cSep08	81817	2042	1962	2	143	upstream transcription factor 2 (Usf2) alternative variant cSep08, mRNA.
Ush1c	Ush1c.bSep08	308596	7917	1793	1	254	usher syndrome 1C homolog (human) (Ush1c) alternative variant bSep08, mRNA.
Ushbp1	Ushbp1.bSep08	290629	3598	695	2	231	usher syndrome 1C binding protein 1 (Ushbp1) alternative variant bSep08, mRNA.
Ushbp1	Ushbp1.cSep08	290629	4146	666	2	160	usher syndrome 1C binding protein 1 (Ushbp1) alternative variant cSep08, mRNA.
Usmg5.1	Usmg5.1.aSep08	171069	5689	663	5	99	upregulated during skeletal muscle growth 5 (11.3 kD) (Usmg5.1) alternative variant aSep08, mRNA.
Usmg5.1	Usmg5.1.bSep08	171069	781	694	2	61	upregulated during skeletal muscle growth 5 (Usmg5.1) alternative variant bSep08, mRNA.
Usmg5.1	Usmg5.1.dSep08	171069	6890	410	4	58	upregulated during skeletal muscle growth 5 (6.4 kD) (Usmg5.1) alternative variant dSep08, mRNA.
Usmg5.1	Usmg5.1.eSep08	171069	6810	369	5	58	upregulated during skeletal muscle growth 5 (6.4 kD) (Usmg5.1) alternative variant eSep08, complete mRNA.
Uso1	Uso1.bSep08	56042	11656	735	1	244	USO1 homolog, vesicle docking protein (yeast) (Uso1) alternative variant bSep08, mRNA.
Usp1	Usp1.bSep08	313387	1843	693		230	ubiquitin specific peptidase 1 (Usp1) alternative variant bSep08, mRNA.
Usp2	Usp2.bSep08	115771	24934	705	7	234	ubiquitin specific peptidase 2 (Usp2) alternative variant bSep08, mRNA.
Usp2	Usp2.cSep08	115771	6930	575	7	191	ubiquitin specific peptidase 2 (Usp2) alternative variant cSep08, mRNA.
Usp2	Usp2.dSep08	115771	8947	790	2	190	ubiquitin specific peptidase 2 (Usp2) alternative variant dSep08, mRNA.
Usp2	Usp2.eSep08	115771	1307	465	3	88	ubiquitin specific peptidase 2 (Usp2) alternative variant eSep08, mRNA.
Usp3	Usp3.aSep08	363084	43814	1294	8	283	ubiquitin specific peptidase 3 (Usp3) alternative variant aSep08, mRNA.
Usp3	Usp3.bSep08	363084	14870	1226	6	219	ubiquitin specific peptidase 3 (Usp3) alternative variant bSep08, mRNA.
Usp3	Usp3.cSep08	363084	12082	436	2	79	ubiquitin specific peptidase 3 (Usp3) alternative variant cSep08, mRNA.
Usp3	Usp3.dSep08	363084	30261	2367	1	41	ubiquitin specific peptidase 3 (Usp3) alternative variant dSep08, mRNA.
Usp4	Usp4.aSep08	290864	43298	3358	22	961	ubiquitin specific peptidase 4 (proto-oncogene) (108.4 kD) (Usp4) alternative variant aSep08, complete mRNA.
Usp4	Usp4.cSep08	290864	3669	511	3	121	ubiquitin specific peptidase 4 (proto-oncogene) (Usp4) alternative variant cSep08, mRNA.
Usp5	Usp5.bSep08	297593	6223	1661	9	371	ubiquitin specific peptidase 5 (isopeptidase T) (Usp5) alternative variant bSep08, mRNA.
Usp5	Usp5.cSep08	297593	762	684	2	116	ubiquitin specific peptidase 5 (isopeptidase T) (Usp5) alternative variant cSep08, mRNA.
Usp7	Usp7.bSep08	360471	77306	1798	10	563	ubiquitin 7 (Usp7) alternative variant bSep08, mRNA.
Usp7	Usp7.cSep08	360471	3044	1789	4	469	ubiquitin 7 (Usp7) alternative variant cSep08, mRNA.
Usp7	Usp7.dSep08	360471	8145	1863	8	212	ubiquitin 7 (Usp7) alternative variant dSep08, mRNA.

Usp7	Usp7.eSep08	360471	4353	736	8	141	ubiquitin 7 (Usp7) alternative variant eSep08, mRNA.
Usp8	Usp8.bSep08	296121	20690	683	5	166	ubiquitin specific peptidase 8 (Usp8) alternative variant bSep08, mRNA.
Usp9x	Usp9x.aSep08	363445	9168	2975	7	325	ubiquitin specific 9 X-linked (Usp9x) alternative variant aSep08, mRNA.
Usp9x	Usp9x.bSep08	363445	16849	3667	3	279	ubiquitin specific peptidase 9 X-linked CRA a (Usp9x) alternative variant bSep08, mRNA.
Usp9x	Usp9x.cSep08	363445	3435	1036	4	194	ubiquitin specific peptidase 9 X-linked CRA b (Usp9x) alternative variant cSep08, mRNA.
Usp10	Usp10.cSep08	307905	1681	480	2	66	ubiquitin specific peptidase 10 (Usp10) alternative variant cSep08, mRNA.
Usp11	Usp11.bSep08	408217	1136	862	2	202	ubiquitin specific peptidase 11 (22.5 kD) (Usp11) alternative variant bSep08, mRNA.
Usp12	Usp12.aSep08	360763	43645	780		259	ubiquitin specific peptidase 12 (Usp12) mRNA.
Usp13	Usp13.bSep08	310306	1945	354	2	52	ubiquitin specific (Usp13) alternative variant bSep08, mRNA.
Usp14	Usp14.bSep08	291796	35453	1319	14	267	ubiquitin specific peptidase 14 (30.0 kD) (Usp14) alternative variant bSep08, mRNA.
Usp14	Usp14.cSep08	291796	1122	676	2	36	ubiquitin specific peptidase 14 (4.2 kD) (Usp14) alternative variant cSep08, mRNA.
Usp15	Usp15.bSep08	171329	14858	1800	5	341	ubiquitin specific peptidase 15 (Usp15) alternative variant bSep08, mRNA.
Usp15	Usp15.cSep08	171329	23303	711	5	237	ubiquitin specific peptidase 15 CRA c (Usp15) alternative variant cSep08, mRNA.
Usp15	Usp15.dSep08	171329	9292	2174	3	127	ubiquitin specific peptidase 15 CRA b (Usp15) alternative variant dSep08, mRNA.
Usp15	Usp15.eSep08	171329	23058	394	4	124	ubiquitin specific peptidase 15 CRA c (Usp15) alternative variant eSep08, mRNA.
Usp16	Usp16.aSep08	288306	29074	2845	18	825	ubiquitin specific peptidase 16 (93.7 kD) (Usp16) alternative variant aSep08, mRNA.
Usp16	Usp16.bSep08	288306	18914	1245	11	287	ubiquitin specific peptidase 16 (32.6 kD) (Usp16) alternative variant bSep08, mRNA.
Usp16	Usp16.cSep08	288306	1989	427	2	24	ubiquitin specific peptidase 16 (2.5 kD) (Usp16) alternative variant cSep08, mRNA.
Usp18	Usp18.bSep08	312688	5071	726	4	149	ubiquitin specific peptidase 18 (17.1 kD) (Usp18) alternative variant bSep08, mRNA.
Usp18	Usp18.cSep08	312688	5971	595	5	113	ubiquitin specific peptidase 18 (Usp18) alternative variant cSep08, mRNA.
Usp19	Usp19.bSep08	361190	2523	860	8	286	ubiquitin specific peptidase 19 CRA d (Usp19) alternative variant bSep08, mRNA.
Usp19	Usp19.cSep08	361190	1390	855	6	233	ubiquitin specific peptidase 19 CRA c (Usp19) alternative variant cSep08, mRNA.
Usp19	Usp19.dSep08	361190	3027	2844	3	191	ubiquitin specific peptidase 19 (Usp19) alternative variant dSep08, mRNA.
Usp19	Usp19.fSep08	361190	3241	1558	4	121	ubiquitin specific peptidase 19 CRA a (13.3 kD) (Usp19) alternative variant fSep08, mRNA.



Usp19	Usp19.hSep08	361190	518	409	2	97	ubiquitin specific peptidase 19 (Usp19) alternative variant hSep08, mRNA.
Usp19	Usp19.iSep08	361190	701	376	3	94	ubiquitin specific peptidase 19 CRA d (Usp19) alternative variant iSep08, mRNA.
Usp19	Usp19.jSep08	361190	2304	437	5	92	ubiquitin specific peptidase 19 CRA d (Usp19) alternative variant jSep08, mRNA.
Usp19	Usp19.lSep08	361190	603	365	2	30	putative protein (Usp19) alternative variant lSep08, mRNA.
Usp20	Usp20.bSep08	311856	15022	517	7	171	ubiquitin specific peptidase 20 (Usp20) alternative variant bSep08, mRNA.
Usp20	Usp20.cSep08	311856	7833	939	6	168	ubiquitin specific peptidase 20 (Usp20) alternative variant cSep08, mRNA.
Usp20	Usp20.eSep08	311856	2382	1913	2	82	ubiquitin specific peptidase 20 CRA a (8.9 kD) (Usp20) alternative variant eSep08, mRNA.
Usp22	Usp22.aSep08	303201	6304	428		142	ubiquitin specific protease 22 (Usp22) mRNA.
Usp24	Usp24.aSep08	313427	3361	644		214	ubiquitin specific protease 24 (Usp24) mRNA.
Usp25	Usp25.bSep08	304150	19432	563	3	146	ubiquitin specific peptidase 25 (Usp25) alternative variant bSep08, mRNA.
Usp25	Usp25.cSep08	304150	1600	348	3	110	ubiquitin specific peptidase 25 (Usp25) alternative variant cSep08, mRNA.
Usp28	Usp28.bSep08	315639	15465	1779	11	518	ubiquitin specific peptidase 28 (Usp28) alternative variant bSep08, mRNA.
Usp28	Usp28.cSep08	315639	2896	371	4	123	ubiquitin specific peptidase 28 (Usp28) alternative variant cSep08, mRNA.
Usp30	Usp30.bSep08	304579	15660	627	7	209	ubiquitin specific peptidase 30 (Usp30) alternative variant bSep08, mRNA.
Usp30	Usp30.cSep08	304579	37607	738	4	144	ubiquitin specific peptidase 30 (Usp30) alternative variant cSep08, mRNA.
Usp30	Usp30.dSep08	304579	1226	805	2	115	ubiquitin specific peptidase 30 (13.0 kD) (Usp30) alternative variant dSep08, mRNA.
Usp32	Usp32.cSep08	303394	24754	363	3	95	ubiquitin specific protease 32 (Usp32) alternative variant cSep08, mRNA.
Usp33	Usp33.aSep08	310960	17009	2311	12	453	ubiquitin specific peptidase 33 CRA a (Usp33) alternative variant aSep08, mRNA.
Usp33	Usp33.bSep08	310960	5894	469	5	144	ubiquitin specific peptidase 33 (Usp33) alternative variant bSep08, mRNA.
Usp33	Usp33.cSep08	310960	4883	411	5	137	ubiquitin specific 33 (Usp33) alternative variant cSep08, mRNA.
Usp33	Usp33.dSep08	310960	1169	1071	2	110	ubiquitin specific 33 (12.5 kD) (Usp33) alternative variant dSep08, mRNA.
Usp33	Usp33.eSep08	310960	3421	1242	3	91	ubiquitin specific peptidase 33 CRA d (10.1 kD) (Usp33) alternative variant eSep08, mRNA.
Usp33	Usp33.fSep08	310960	1624	707	2	47	ubiquitin specific 33 (Usp33) alternative variant fSep08, mRNA.
Usp35	Usp35.aSep08	308834	8190	754	1	250	ubiquitin specific peptidase 35 (Usp35) alternative variant aSep08, mRNA.
Usp35	Usp35.bSep08	308834	5051	816	1	225	ubiquitin specific peptidase 35 (Usp35) alternative variant bSep08, mRNA.

Usp35	Usp35.cSep08	308834	4787	707	3	137	ubiquitin specific peptidase 35 (Usp35) alternative variant cSep08, mRNA.
Usp36	Usp36.bSep08	303700	8760	3189	7	404	ubiquitin specific peptidase 36 (Usp36) alternative variant bSep08, mRNA.
Usp38	Usp38.aSep08	307764	10810	1700	5	566	ubiquitin specific peptidase 38 (Usp38) alternative variant aSep08, mRNA.
Usp39	Usp39.bSep08	297336	32411	2603	15	564	ubiquitin specific peptidase 39 (65.2 kD) (Usp39) alternative variant bSep08, mRNA.
Usp39	Usp39.cSep08	297336	19242	1790	7	377	ubiquitin specific peptidase 39 (Usp39) alternative variant cSep08, mRNA.
Usp39	Usp39.dSep08	297336	10232	994	5	179	ubiquitin specific peptidase 39 (Usp39) alternative variant dSep08, mRNA.
Usp39	Usp39.eSep08	297336	5320	513	4	116	ubiquitin specific peptidase 39 (Usp39) alternative variant eSep08, mRNA.
Usp39	Usp39.fSep08	297336	3196	693	4	82	ubiquitin specific peptidase 39 (9.5 kD) (Usp39) alternative variant fSep08, mRNA.
Usp40	Usp40.bSep08	316599	5920	661	6	219	ubiquitin specific peptidase 40 (Usp40) alternative variant bSep08, mRNA.
Usp40	Usp40.cSep08	316599	15939	930	11	209	ubiquitin specific peptidase 40 (Usp40) alternative variant cSep08, mRNA.
Usp40	Usp40.dSep08	316599	2542	519	4	135	ubiquitin specific peptidase 40 (Usp40) alternative variant dSep08, mRNA.
Usp42	Usp42.bSep08	288482	1288	515	2	69	ubiquitin specific peptidase 42 (Usp42) alternative variant bSep08, mRNA.
Usp45	Usp45.bSep08	313098	5938	674	2	224	ubiquitin specific peptidase 45 (Usp45) alternative variant bSep08, mRNA.
Usp45	Usp45.cSep08	313098	3785	2143	2	73	ubiquitin specific peptidase 45 (8.2 kD) (Usp45) alternative variant cSep08, mRNA.
Usp46	Usp46.aSep08	289584	16485	728		162	ubiquitin specific peptidase 46 (Usp46) mRNA.
Usp47	Usp47.bSep08	308896	31913	1787	6	472	ubiquitin specific peptidase 47 (Usp47) alternative variant bSep08, mRNA.
Usp47	Usp47.cSep08	308896	17823	971	9	323	ubiquitin specific peptidase 47 (Usp47) alternative variant cSep08, mRNA.
Usp48	Usp48.bSep08	362636	24145	1413	12	278	ubiquitin specific 48 (31.1 kD) (Usp48) alternative variant bSep08, mRNA.
Usp48	Usp48.dSep08	362636	662	561	2	57	ubiquitin specific 48 (Usp48) alternative variant dSep08, mRNA.
Usp50	Usp50.aSep08	311399	5008	720	3	193	ubiquitin specific peptidase 50 (Usp50) alternative variant aSep08, mRNA.
Usp50	Usp50.bSep08	311399	5072	858	3	139	ubiquitin specific peptidase 50 (Usp50) alternative variant bSep08, mRNA.
Usp50	Usp50.cSep08	311399	3382	684	2	93	ubiquitin specific peptidase 50 (Usp50) alternative variant cSep08, mRNA.
Usp50	Usp50.dSep08	311399	3373	749	2	84	ubiquitin specific peptidase 50 (Usp50) alternative variant dSep08, mRNA.
Usp52	Usp52.bSep08	408200	3212	819	8	272	ubiquitin specific peptidase 52 CRA e (Usp52) alternative variant bSep08, mRNA.

Usp52	Usp52.dSep08	408200	6729	1780	6	132	ubiquitin specific peptidase 52 CRA b (Usp52) alternative variant dSep08, mRNA.
Usp52	Usp52.eSep08	408200	3480	1883	4	121	specific 2 (13.9 kD) (Usp52) alternative variant eSep08, mRNA.
Usp52	Usp52.gSep08	408200	2309	755	3	57	ubiquitin specific peptidase 52 CRA b (Usp52) alternative variant gSep08, mRNA.
Usp53	Usp53.bSep08	295425	2854	906	2	279	ubiquitin specific peptidase 53 (Usp53) alternative variant bSep08, mRNA.
Usp53	Usp53.cSep08	295425	1783	403	2	19	ubiquitin specific peptidase 53 (Usp53) alternative variant cSep08, mRNA.
Usp54	Usp54.bSep08	408223	6155	1692	2	438	ubiquitin specific peptidase 54 (47.8 kD) (Usp54) alternative variant bSep08, mRNA.
Usp54	Usp54.cSep08	408223	2150	759	2	252	ubiquitin specific peptidase 54 (Usp54) alternative variant cSep08, mRNA.
Usp54	Usp54.dSep08	408223	6256	448	2	22	ubiquitin specific peptidase 54 (2.7 kD) (Usp54) alternative variant dSep08, mRNA.
Uspl1	Uspl1.aSep08	288447	11126	2395	4	735	ubiquitin specific peptidase like 1 (Uspl1) alternative variant aSep08, mRNA.
Uspl1	Uspl1.dSep08	288447	7935	768	3	235	ubiquitin specific peptidase like 1 (Uspl1) alternative variant dSep08, mRNA.
Uteroglobin.0	Uteroglobin.0.aSep08		4240	513		117	PBPC1BS like (Uteroglobin.0) alternative variant aSep08, mRNA.
Uteroglobin.0	Uteroglobin.0.bSep08		4143	562	1	37	CRA a like (4.4 kD) (Uteroglobin.0) alternative variant bSep08, mRNA.
Utp6	Utp6.aSep08	360574	6257	1796		197	UTP6, small subunit (SSU) processome component, homolog (yeast) (Utp6) mRNA.
Utp14a	Utp14a.bSep08	317579	5157	1184	4	348	UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast) (Utp14a) alternative variant bSep08, mRNA.
Utp14a	Utp14a.cSep08	317579	4239	930	3	292	UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast) (Utp14a) alternative variant cSep08, mRNA.
Utp14a	Utp14a.dSep08	317579	2590	1262	4	206	UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast) (Utp14a) alternative variant dSep08, mRNA.
Utp14a	Utp14a.eSep08	317579	2644	556	2	132	UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast) (Utp14a) alternative variant eSep08, mRNA.
Utp15	Utp15.bSep08	310019	17730	1788	5	445	UTP15, U3 small nucleolar ribonucleoprotein, homolog (yeast) (Utp15) alternative variant bSep08, mRNA.
Utp15	Utp15.cSep08	310019	8979	1453	7	171	UTP15, U3 small nucleolar ribonucleoprotein, homolog (yeast) (Utp15) alternative variant cSep08, mRNA.
Utp18	Utp18.aSep08	303456	26976	2506		560	UTP18, small subunit (SSU) processome component, homolog (yeast) (Utp18) mRNA.
Utp20	Utp20.aSep08	314713	10798	1981		543	UTP20, small subunit (SSU) processome component, homolog (yeast) (Utp20) mRNA.
Utrn	Utrn.aSep08	25600	148596	1202	5	321	utrophin (36.9 kD) (Utrn) alternative variant aSep08, mRNA.
Utrn	Utrn.bSep08	25600	8769	405	3	135	utrophin (Utrn) alternative variant bSep08, mRNA.
Utx	Utx.aSep08	317178	14015	390		130	ubiquitously transcribed tetratricopeptide repeat (Utx) mRNA.

Uvrag	Uvrag.bSep08	308846	29975	2292	3	306	UV radiation resistance associated (33.2 kD) (Uvrag) alternative variant bSep08, mRNA.
Uvrag	Uvrag.cSep08	308846	107006	899	5	299	UV radiation resistance associated (Uvrag) alternative variant cSep08, mRNA.
Uvrag	Uvrag.dSep08	308846	13808	904	2	123	UV radiation resistance associated (Uvrag) alternative variant dSep08, mRNA.
Uxs1	Uxs1.aSep08	246232	52493	1820	3	370	UDP-glucuronate decarboxylase 1 (Uxs1) alternative variant aSep08, mRNA.
Uxs1	Uxs1.bSep08	246232	38835	701	1	233	UDP-glucuronate decarboxylase 1 (Uxs1) alternative variant bSep08, mRNA.
Uxt	Uxt.bSep08	299313	12493	842	5	175	ubiquitously expressed transcript (Uxt) alternative variant bSep08, mRNA.
Uxt	Uxt.cSep08	299313	10717	1082	3	107	ubiquitously expressed transcript (12.1 kD) (Uxt) alternative variant cSep08, mRNA.
Uxt	Uxt.dSep08	299313	12042	571	4	103	ubiquitously expressed transcript (11.7 kD) (Uxt) alternative variant dSep08, complete mRNA.
Uxt	Uxt.eSep08	299313	12538	780	4	89	ubiquitously expressed transcript (10.1 kD) (Uxt) alternative variant eSep08, mRNA.
V-set.0	V-set.0.aSep08		1012	921		119	t-cell receptor like (13.3 kD) (V-set.0) mRNA.
V-set.1	V-set.1.aSep08		956	734		129	T cell receptor (V-set.1) mRNA.
V-set.2	V-set.2.aSep08		708	590		148	T-cell receptor region like (V-set.2) mRNA.
V-set.3	V-set.3.aSep08		528	348		109	T cell receptor (V-set.3) mRNA.
V-set.4	V-set.4.aSep08		570	399		109	T-cell receptor like (V-set.4) mRNA.
V-set.5	V-set.5.aSep08		488	340		113	T-cell receptor like (V-set.5) mRNA.
V-set.6	V-set.6.aSep08		587	493		113	t-cell receptor like (V-set.6) mRNA.
V-set.7	V-set.7.aSep08		520	340		108	T cell receptor (V-set.7) mRNA.
V-set.8	V-set.8.aSep08		826	735		119	t-cell receptor like (13.3 kD) (V-set.8) mRNA.
V-set.9	V-set.9.aSep08		519	339		112	T cell receptor AV10S10 like (V-set.9) mRNA.
V-set.10	V-set.10.aSep08		568	473		113	t-cell receptor like (V-set.10) mRNA.
V-set.11	V-set.11.aSep08		629	534		177	T-cell receptor like (V-set.11) mRNA.
V-set.12	V-set.12.aSep08		528	395		111	T cell receptor (V-set.12) mRNA.
V-set.13	V-set.13.aSep08		7671	704		128	T cell receptor (V-set.13) mRNA.
V-set.14	V-set.14.aSep08		956	734		129	T cell receptor (V-set.14) mRNA.
V-set.15	V-set.15.aSep08		588	395		131	T cell receptor V delta 5 (V-set.15) mRNA.
V-set.16	V-set.16.aSep08		629	534		177	T-cell receptor like (V-set.16) mRNA.
V-set.17	V-set.17.aSep08		483	365		121	T-cell receptor like (V-set.17) mRNA.
V-set.18	V-set.18.aSep08		484	389		115	immunoglobulin lambda chain (V-set.18) mRNA.
V-set.19	V-set.19.aSep08		514	393		131	this CDS feature is included show the translation corresponding V region. Presently qualifiers on V region features are illegal (V-set.19) mRNA.
V-set.20	V-set.20.aSep08		459	375		116	immunoglobulin heavy chain variable region (V-set.20) mRNA.
V-set.21	V-set.21.aSep08		525	415		138	heavy chain (V-set.21) mRNA.
V-set.22	V-set.22.aSep08		455	371		123	H prechain like (V-set.22) mRNA.

V-set.23	V-set.23.aSep08		434	350		116	immunoglobulin heavy chain variable region (V-set.23) mRNA.
V-set.24	V-set.24.aSep08		434	350		116	immunoglobulin heavy chain variable region (V-set.24) mRNA.
V-set.25	V-set.25.aSep08		486	378		126	immunoglobulin heavy chain (V-set.25) mRNA.
V-set.26	V-set.26.aSep08		535	421		139	heavy chain (V-set.26) mRNA.
V-set.27	V-set.27.aSep08		438	354		117	heavy chain variable region (V-set.27) mRNA.
V-set.28	V-set.28.aSep08		443	359		119	immunoglobulin heavy chain variable region (V-set.28) mRNA.
V-set.29	V-set.29.aSep08		518	404		134	immunoglobulin heavy chain (V-set.29) mRNA.
V-set.30	V-set.30.aSep08		535	420		117	immunoglobulin heavy chain (V-set.30) mRNA.
V-set.31	V-set.31.aSep08		461	375		115	heavy chain (V-set.31) mRNA.
V-set.32	V-set.32.aSep08		494	398		132	immunoglobulin heavy chain (V-set.32) mRNA.
V-set.33	V-set.33.aSep08		480	401		119	heavy chain (V-set.33) mRNA.
V-set.34	V-set.34.aSep08		480	394		117	immunoglobulin (V-set.34) mRNA.
V-set.35	V-set.35.aSep08		468	393		118	heavy chain (V-set.35) mRNA.
V-set.36	V-set.36.aSep08		526	426		142	immunoglobulin heavy chain (V-set.36) mRNA.
V-set.37	V-set.37.aSep08		459	377		125	immunoglobulin (V-set.37) mRNA.
V-set.38	V-set.38.aSep08		521	411		119	immunoglobulin heavy chain (V-set.38) mRNA.
V-set.39	V-set.39.aSep08		453	372		123	heavy chain (V-set.39) mRNA.
V-set.40	V-set.40.aSep08		454	371		123	heavy chain (V-set.40) mRNA.
V-set.41	V-set.41.aSep08		469	386		120	heavy chain (V-set.41) mRNA.
V-set.42	V-set.42.aSep08		447	363		120	immunoglobulin heavy chain variable region (V-set.42) mRNA.
V-set.43	V-set.43.aSep08		435	352		116	heavy chain (V-set.43) mRNA.
V-set.44	V-set.44.aSep08		439	355		118	immunoglobulin heavy chain variable region (V-set.44) mRNA.
V-set.60	V-set.60.aSep08		738	498		118	T-cell receptor like (V-set.60) mRNA.
V-set.61	V-set.61.aSep08		575	491		116	T cell receptor (V-set.61) mRNA.
V-set.62	V-set.62.aSep08		2898	399	1	133	T cell receptor (V-set.62) alternative variant aSep08, mRNA.
V-set.62	V-set.62.bSep08		894	774	1	116	T cell receptor (V-set.62) alternative variant bSep08, mRNA.
V-set.63	V-set.63.aSep08		781	400		121	immunoglobulin light chain (V-set.63) mRNA.
V-set.64	V-set.64.aSep08		459	348		115	kappa chain (V-set.64) mRNA.
V-set.65	V-set.65.aSep08		489	374		124	immunoglobulin 4G6 light chain variable region (V-set.65) mRNA.
V-set.66	V-set.66.aSep08		460	340		112	immunoglobulin light chain (V-set.66) mRNA.
V-set.67	V-set.67.aSep08		480	357		118	immunoglobulin light chain variable region (V-set.67) mRNA.
V-set.68	V-set.68.aSep08		484	356		118	immunoglobulin chain (V-set.68) mRNA.
V-set.69	V-set.69.aSep08		521	322		107	chain kappa (V-set.69) mRNA.
V-set.70	V-set.70.aSep08		484	360		119	immunoglobulin chain (V-set.70) mRNA.

V-set.71	V-set.71.aSep08		491	366		121	immunoglobulin chain (V-set.71) mRNA.
V-set.72	V-set.72.aSep08		694	329		109	light chain (V-set.72) mRNA.
V-set.73	V-set.73.aSep08		544	360		120	immunoglobulin chain (V-set.73) mRNA.
V-set.76	V-set.76.aSep08		4264	1271		226	15 -related cell adhesion molecule (V-set.76) mRNA.
V-set.77	V-set.77.aSep08		10349	2207	5	475	pregnancy specific glycoprotein (53.1 kD) (V-set.77) alternative variant aSep08, complete mRNA.
V-set.78	V-set.78.aSep08		524	414		137	heavy chain (V-set.78) mRNA.
V-set.79	V-set.79.aSep08		483	365		121	T-cell receptor like (V-set.79) mRNA.
vaby	vaby.aSep08		2966	347		29	putative protein (3.2 kD) (vaby) mRNA.
Vac14	Vac14.bSep08	307842	8483	809	3	123	vac14 homolog (S. cerevisiae) (Vac14) alternative variant bSep08, mRNA.
Vac14	Vac14.cSep08	307842	5770	397	2	106	vac14 homolog (S. cerevisiae) (11.2 kD) (Vac14) alternative variant cSep08, mRNA.
vachy	vachy.aSep08		6710	464		154	gene associated (vachy) mRNA.
vadar	vadar.aSep08		14876	544		17	putative protein (2.2 kD) (vadar) mRNA.
vadoy	vadoy.aSep08		1356	287		53	putative protein (vadoy) mRNA.
vaflo	vaflo.aSep08		4649	325		100	putative protein (vaflo) mRNA.
vaflu	vaflu.aSep08		7815	512		54	putative protein (vaflu) mRNA.
vagar	vagar.aSep08		5553	826		47	putative protein (vagar) mRNA.
vaja	vaja.aSep08		561	402		32	T cell receptor (vaja) mRNA.
vajey	vajey.aSep08		8593	645		214	CRA a (vajey) mRNA.
vakee	vakee.aSep08		7979	688		33	putative protein (4.1 kD) (vakee) mRNA.
vakler	vakler.aSep08		2459	687		115	putative protein (vakler) mRNA.
valo	valo.aSep08		11513	771		109	tyrosine kinase 3 (valo) mRNA.
vamee	vamee.aSep08		44889	510		131	folliculin interacting protein 1 (14.5 kD) (vamee) mRNA.
Vamp1	Vamp1.aSep08	25624	6937	1214	2	170	vesicle-associated membrane protein 1 (Vamp1) alternative variant aSep08, mRNA.
Vamp1	Vamp1.cSep08	25624	6683	724	3	149	vesicle-associated membrane protein 1 (Vamp1) alternative variant cSep08, mRNA.
Vamp1	Vamp1.dSep08	25624	4697	960	1	141	vesicle-associated membrane protein 1 (Vamp1) alternative variant dSep08, mRNA.
Vamp2	Vamp2.bSep08	24803	1458	359	4	119	vesicle-associated membrane protein 2 (Vamp2) alternative variant bSep08, mRNA.
Vamp2	Vamp2.cSep08	24803	1392	380	3	103	vesicle-associated membrane protein 2 (Vamp2) alternative variant cSep08, mRNA.
Vamp4	Vamp4.aSep08	364033	19483	1413	8	141	vesicle-associated membrane protein 4 (16.4 kD) (Vamp4) alternative variant aSep08, mRNA.
Vamp4	Vamp4.cSep08	364033	6574	668	4	52	vesicle-associated membrane protein 4 (6.0 kD) (Vamp4) alternative variant cSep08, mRNA.
Vamp5	Vamp5.bSep08	89818	1700	702	3	91	vesicle-associated membrane protein 5 (9.7 kD) (Vamp5) alternative variant bSep08, mRNA.
Vamp7	Vamp7.aSep08	85491	38808	3305	8	220	vesicle-associated membrane protein 7 (24.8 kD) (Vamp7) alternative variant aSep08, mRNA.

Vangl1	Vangl1.cSep08	690366	1686	393	2	88	vang-like 1 (van gogh, Drosophila) (Vangl1) alternative variant cSep08, mRNA.
vanoy	vanoy.aSep08		8049	705		47	putative protein (5.7 kD) (vanoy) mRNA.
Vapa	Vapa.cSep08	58857	23921	721	5	176	vesicle-associated membrane protein, associated protein a (Vapa) alternative variant cSep08, mRNA.
Vapa	Vapa.eSep08	58857	10580	602	3	128	vesicle-associated membrane protein, associated protein a (14.4 kD) (Vapa) alternative variant eSep08, mRNA.
vapor	vapor.aSep08		5808	1012		65	putative protein (vapor) mRNA.
varby	varby.aSep08		5401	2125	2	106	serine threonine protein kinase MST4 (varby) alternative variant aSep08, mRNA.
varby	varby.bSep08		2049	367	1	49	putative protein (5.4 kD) (varby) alternative variant bSep08, mRNA.
varchy	varchy.aSep08		3900	3539		69	zinc finger protein 106 (varchy) mRNA.
vardar	vardar.aSep08		26093	348		62	putative protein (7.0 kD) (vardar) mRNA.
vardoy	vardoy.aSep08		25456	731	4	211	niemann-pick disease type c1 like (vardoy) alternative variant aSep08, mRNA.
vardoy	vardoy.bSep08		3374	308	2	102	niemann-pick C1 (vardoy) alternative variant bSep08, mRNA.
varflo	varflo.aSep08		6585	574		102	putative nuclear protein (11.1 kD) (varflo) mRNA.
varflu	varflu.aSep08		1818	601		49	putative protein (varflu) mRNA.
vargar	vargar.aSep08		6547	484		102	putative protein (11.3 kD) (vargar) mRNA.
varja	varja.aSep08		11724	1285		9	putative protein (0.9 kD) (varja) mRNA.
varjey	varjey.aSep08		7149	552		183	FRYL protein (varjey) mRNA.
varkee	varkee.aSep08		10192	738		51	putative protein (5.5 kD) (varkee) mRNA.
varkler	varkler.aSep08		21749	1203		400	intersectin 2 CRA a (varkler) mRNA.
varlo	varlo.aSep08		7684	763		41	putative protein (4.5 kD) (varlo) mRNA.
varmee	varmee.aSep08		30923	207		52	putative protein (varmee) mRNA.
varnoy	varnoy.aSep08		2371	674		108	dis3 mitotic control homolog -like 2 (varnoy) mRNA.
varpor	varpor.aSep08		2158	739		44	putative protein (varpor) mRNA.
Vars2	Vars2.bSep08	25009	1727	835	8	278	valyl-tRNA synthetase 2 (Vars2) alternative variant bSep08, mRNA.
Vars2	Vars2.cSep08	25009	752	570	2	189	valyl-tRNA synthetase 2 (Vars2) alternative variant cSep08, mRNA.
Vars2	Vars2.dSep08	25009	2996	738	7	184	valyl-tRNA synthetase 2 (Vars2) alternative variant dSep08, mRNA.
Vars2	Vars2.eSep08	25009	1217	579	3	170	valyl-tRNA synthetase 2 (Vars2) alternative variant eSep08, mRNA.
Vars2	Vars2.fSep08	25009	814	422	4	140	valyl-tRNA synthetase 2 (Vars2) alternative variant fSep08, mRNA.
Vars2	Vars2.hSep08	25009	830	393	3	98	valyl-tRNA synthetase 2 (Vars2) alternative variant hSep08, mRNA.
Vars2l	Vars2l.bSep08	309596	3583	1821	8	348	valyl-tRNA synthetase 2-like (Vars2l) alternative variant bSep08, mRNA.
Vars2l	Vars2l.cSep08	309596	1292	777	7	244	valyl-tRNA synthetase 2-like (Vars2l) alternative variant cSep08, mRNA.

Vars2l	Vars2l.dSep08	309596	914	691	3	145	valyl-tRNA synthetase 2-like (Vars2l) alternative variant dSep08, mRNA.
Vars2l	Vars2l.eSep08	309596	2187	781	8	109	valyl-tRNA synthetase 2-like (Vars2l) alternative variant eSep08, mRNA.
varsa	varsa.aSep08		1902	1200		244	ligase-like 8 (varsa) mRNA.
varshee	varshee.aSep08		21424	587		85	putative protein (varshee) mRNA.
vartu	vartu.aSep08		2491	293		97	gene regulated by estrogen in breast cancer protein like (vartu) mRNA.
varvar	varvar.aSep08		4963	785		261	zinc finger castor 1 (varvar) mRNA.
varwey	varwey.aSep08		33004	1777	1	64	putative protein (7.2 kD) (varwey) alternative variant aSep08, mRNA.
varwey	varwey.bSep08		11301	729	2	63	putative protein (7.3 kD) (varwey) alternative variant bSep08, mRNA.
vasa	vasa.aSep08		67432	269		89	tbc1 domain family member 22a (vasa) mRNA.
Vash1	Vash1.aSep08	503052	3019	770		156	vasohibin 1 (Vash1) mRNA.
Vash2	Vash2.aSep08	498309	17442	1434	7	277	vasohibin 2 (Vash2) alternative variant aSep08, mRNA.
Vash2	Vash2.cSep08	498309	7618	904	4	36	vasohibin 2 (4.3 kD) (Vash2) alternative variant cSep08, mRNA.
vashee	vashee.aSep08		2853	1764		457	insulin receptor-related receptor CRA a (vashee) mRNA.
Vasp	Vasp.bSep08	361517	7854	1677	3	361	vasodilator-stimulated phosphoprotein (Vasp) alternative variant bSep08, mRNA.
Vasp	Vasp.cSep08	361517	3730	661	2	220	vasodilator-stimulated phosphoprotein (Vasp) alternative variant cSep08, mRNA.
Vat1	Vat1.dSep08	287721	650	225	3	60	vesicle amine transport protein 1 homolog (T californica) (Vat1) alternative variant dSep08, mRNA.
vatu	vatu.aSep08		7450	301		100	neuroblastoma-amplified protein (vatu) mRNA.
Vav1	Vav1.bSep08	25156	23242	728	6	242	vav 1 oncogene (Vav1) alternative variant bSep08, mRNA.
Vav1	Vav1.cSep08	25156	2649	616	6	175	vav 1 oncogene (Vav1) alternative variant cSep08, mRNA.
Vav1	Vav1.dSep08	25156	12048	1733	5	158	vav 1 oncogene (Vav1) alternative variant dSep08, mRNA.
Vav2	Vav2.cSep08	296603	4422	753	3	79	vav2 oncogene (Vav2) alternative variant cSep08, mRNA.
Vav3	Vav3.aSep08	295378	102000	2389		311	vav 3 oncogene (Vav3) mRNA.
vavar	vavar.aSep08		1709	778		259	putative protein of mammalian origin (vavar) mRNA.
vawby	vawby.aSep08		620	385	2	75	putative protein (vawby) alternative variant aSep08, mRNA.
vawby	vawby.bSep08		2090	770	3	71	putative protein (vawby) alternative variant bSep08, mRNA.
vawby	vawby.cSep08		2426	1075	3	62	putative protein (vawby) alternative variant cSep08, mRNA.
vawby	vawby.dSep08		2523	1226	4	49	CRA d like (6.1 kD) (vawby) alternative variant dSep08, mRNA.
vawchy	vawchy.aSep08		706	490		40	putative protein (4.3 kD) (vawchy) mRNA.
vawdar	vawdar.aSep08		3955	768		102	putative protein (11.0 kD) (vawdar) mRNA.
vawdoy	vawdoy.aSep08		1098	393		45	putative protein (vawdoy) mRNA.
vawey	vawey.aSep08		5120	531		96	putative protein (10.7 kD) (vawey) mRNA.
vawflo	vawflo.aSep08		575	224		42	putative protein (4.7 kD) (vawflo) mRNA.
vawflu	vawflu.aSep08		14026	674		50	putative protein (vawflu) mRNA.



vawgar	vawgar.aSep08		14238	704		61	putative protein (7.0 kD) (vawgar) mRNA.
vawja	vawja.aSep08		667	581		35	putative protein (3.7 kD) (vawja) mRNA.
vawjey	vawjey.aSep08		7953	726	6	233	FRY protein (vawjey) alternative variant aSep08, mRNA.
vawjey	vawjey.bSep08		3591	342	4	113	FRY protein (vawjey) alternative variant bSep08, mRNA.
vawkee	vawkee.aSep08		1459	626		75	putative protein (vawkee) mRNA.
vawkler	vawkler.aSep08		2626	249		43	putative protein (4.9 kD) (vawkler) mRNA.
vawlo	vawlo.aSep08		2986	575		39	putative protein (4.4 kD) (vawlo) mRNA.
vawmee	vawmee.aSep08		2981	448		53	putative protein (6.2 kD) (vawmee) mRNA.
vawnoy	vawnoy.aSep08		627	271		89	putative protein (vawnoy) mRNA.
vawpor	vawpor.aSep08		9019	389		25	putative protein (vawpor) mRNA.
vawsa	vawsa.aSep08		22412	738		176	ligase-like 8 (vawsa) mRNA.
vawshee	vawshee.aSep08		7481	430		70	putative protein (vawshee) mRNA.
vawtu	vawtu.aSep08		1361	380		101	putative protein (10.7 kD) (vawtu) mRNA.
vawvar	vawvar.aSep08		6800	1800	2	599	zinc finger castor homolog 1 (vawvar) alternative variant aSep08, mRNA.
vawvar	vawvar.bSep08		4847	528	3	175	zinc finger castor homolog 1 (vawvar) alternative variant bSep08, mRNA.
vawvar	vawvar.dSep08		4054	1783	2	33	putative protein (3.6 kD) (vawvar) alternative variant dSep08, mRNA.
vawwey	vawwey.aSep08		894	779		96	putative protein (vawwey) mRNA.
VBS.0	VBS.0.aSep08		21958	663		221	talin 2 (VBS.0) mRNA.
Vcan	Vcan.aSep08	114122	36808	3013	8	364	versican (Vcan) alternative variant aSep08, mRNA.
Vcp	Vcp.bSep08	116643	2854	863	6	282	valosin-containing protein (Vcp) alternative variant bSep08, mRNA.
Vcp	Vcp.dSep08	116643	4866	760	3	64	valosin-containing protein (Vcp) alternative variant dSep08, mRNA.
Vcsa1	Vcsa1.bSep08	24867	56043	630	2	162	variable coding sequence A1 (Vcsa1) alternative variant bSep08, mRNA.
Vdac2	Vdac2.bSep08	83531	9057	1202	8	274	voltage-dependent anion channel 2 (29.6 kD) (Vdac2) alternative variant bSep08, mRNA.
Vdac2	Vdac2.cSep08	83531	6681	895	5	120	voltage-dependent anion channel 2 (13.1 kD) (Vdac2) alternative variant cSep08, mRNA.
Vdac3	Vdac3.bSep08	83532	14962	1177	7	245	voltage-dependent anion channel 3 (26.9 kD) (Vdac3) alternative variant bSep08, mRNA.
veeby	veeby.aSep08		85284	705		209	putative protein of bilateral origin (veeby) mRNA.
veechy	veechy.aSep08		1057	668	2	83	putative protein of metazoan origin (veechy) alternative variant aSep08, mRNA.
veedar	veedar.aSep08		8463	281		47	putative protein (veedar) mRNA.
veedoy	veedoy.aSep08		24440	268		71	putative nuclear protein (8.0 kD) (veedoy) mRNA.
veeflo	veeflo.aSep08		2023	352		60	sideroflexin 4 (veeflo) mRNA.
veeflu	veeflu.aSep08		1190	348		53	putative protein (veeflu) mRNA.
veefly	veefly.aSep08		91088	733		97	putative protein of mammalian origin (veefly) mRNA.
veegar	veegar.aSep08		64740	307		73	CRA b like (veegar) mRNA.

veeja	veeja.aSep08		954	385		128	apoptotic chromatin condensation inducer 1 (veeja) mRNA.
veejey	veejey.aSep08		31436	1790		548	FRY protein (veejey) alternative variant aSep08, mRNA.
veejey	veejey.bSep08		4556	2589		191	fryl protein (21.3 kD) (veejey) alternative variant bSep08, mRNA.
veekee	veekee.aSep08		1590	739		99	putative protein (veekee) mRNA.
veekler	veekler.aSep08		9250	332		69	putative protein of eukaryotic origin (veekler) mRNA.
veelo	veelo.aSep08		1750	237		29	putative protein (veelo) mRNA.
veemee	veemee.aSep08		2723	298		53	putative protein (veemee) mRNA.
veenoy	veenoy.aSep08		68659	2324	9	185	grb10 interacting GYF protein 2 (20.7 kD) (veenoy) alternative variant aSep08, mRNA.
veenoy	veenoy.bSep08		1305	803	2	30	putative protein (3.4 kD) (veenoy) alternative variant bSep08, mRNA.
veepor	veepor.aSep08		1938	410		136	inhibitor of Bruton's tyrosine kinase (veepor) mRNA.
veesa	veesa.aSep08		2350	530		111	putative protein (veesa) alternative variant aSep08, mRNA.
veeshee	veeshee.aSep08		744	284		47	putative protein (veeshee) mRNA.
veetu	veetu.bSep08		23704	467	4	85	putative cytoplasmic protein (9.9 kD) (veetu) alternative variant bSep08, mRNA.
veetu	veetu.cSep08		4443	393	2	32	putative protein (3.5 kD) (veetu) alternative variant cSep08, mRNA.
veevar	veevar.bSep08		5220	421	4	59	putative protein (veevar) alternative variant bSep08, mRNA.
veewey	veewey.aSep08		635	541		21	putative protein (veewey) mRNA.
Vegfa	Vegfa.dSep08	83785	7879	1074	5	248	vascular endothelial growth factor A (Vegfa) alternative variant dSep08, mRNA.
Vegfa	Vegfa.eSep08	83785	12441	513	7	170	vascular endothelial growth factor A (20.0 kD) (Vegfa) alternative variant eSep08, mRNA.
Vegfa	Vegfa.fSep08	83785	12458	541	6	148	vascular endothelial growth factor A (17.5 kD) (Vegfa) alternative variant fSep08, mRNA.
Vegfa	Vegfa.hSep08	83785	12458	409	5	104	vascular endothelial growth factor A (12.3 kD) (Vegfa) alternative variant hSep08, mRNA.
Vegfa	Vegfa.iSep08	83785	5675	292	5	97	vascular endothelial growth factor A (Vegfa) alternative variant iSep08, mRNA.
Vegfa	Vegfa.jSep08	83785	3091	281	2	93	vascular endothelial growth factor A (Vegfa) alternative variant jSep08, mRNA.
Vegfa	Vegfa.lSep08	83785	11748	1401	6	82	vascular endothelial growth factor A (Vegfa) alternative variant lSep08, mRNA.
Vegfa	Vegfa.mSep08	83785	4615	1333	3	78	vascular endothelial growth factor A (Vegfa) alternative variant mSep08, mRNA.
Vegfa	Vegfa.oSep08	83785	6314	793	3	41	vascular endothelial growth factor A (5.0 kD) (Vegfa) alternative variant oSep08, mRNA.
Vegfa	Vegfa.pSep08	83785	12458	105	2	23	vascular endothelial growth factor A (2.8 kD) (Vegfa) alternative variant pSep08, mRNA.
Vegfb	Vegfb.bSep08	89811	4396	563	6	168	vascular endothelial growth factor B (Vegfb) alternative variant bSep08, mRNA.
Vegfb	Vegfb.cSep08	89811	1791	923	4	101	vascular endothelial growth factor B (Vegfb) alternative variant cSep08, mRNA.

Vegfb	Vegfb.dSep08	89811	997	711	2	89	vascular endothelial growth factor B (Vegfb) alternative variant dSep08, mRNA.
Vegfb	Vegfb.eSep08	89811	3328	649	2	81	vascular endothelial growth factor B (8.8 kD) (Vegfb) alternative variant eSep08, complete mRNA.
Vegfc	Vegfc.bSep08	114111	115257	1174	1	269	vascular endothelial growth factor C (Vegfc) alternative variant bSep08, mRNA.
Vegp2	Vegp2.bSep08	94106	4036	489		107	von Ebners gland protein 2 (Vegp2) alternative variant bSep08, mRNA.
Veph1	Veph1.bSep08	361954	92826	813	4	165	ventricular zone expressed PH domain homolog 1 (zebrafish) (Veph1) alternative variant bSep08, mRNA.
Veph1	Veph1.cSep08	361954	9529	433	2	126	ventricular zone expressed PH domain homolog 1 (zebrafish) (Veph1) alternative variant cSep08, mRNA.
verby	verby.aSep08		5412	745		113	putative nuclear protein (13.0 kD) (verby) mRNA.
verchy	verchy.aSep08		6846	375		56	putative protein (6.1 kD) (verchy) mRNA.
verdar	verdar.aSep08		823	350		62	putative protein (verdar) mRNA.
verdoy	verdoy.aSep08		1339	638		46	putative protein (5.0 kD) (verdoy) mRNA.
verflu	verflu.aSep08		9461	505		41	putative protein (verflu) mRNA.
verfly	verfly.aSep08		443	250		19	putative protein (verfly) mRNA.
vergar	vergar.aSep08		55658	619	1	87	SEC7-like (vergar) alternative variant aSep08, mRNA.
vergar	vergar.bSep08		35258	324	1	75	SEC7-like (vergar) alternative variant bSep08, mRNA.
verja	verja.aSep08		2470	614		204	putative protein of eukaryotic origin (verja) mRNA.
verjey	verjey.aSep08		1586	530		39	putative protein (verjey) mRNA.
verkee	verkee.aSep08		10883	1337		44	putative protein (5.1 kD) (verkee) mRNA.
verkler	verkler.aSep08		8561	754		118	putative protein (12.6 kD) (verkler) mRNA.
verlo	verlo.aSep08		7044	450		68	putative protein (verlo) mRNA.
vermee	vermee.aSep08		6760	294		92	putative protein (vermee) mRNA.
vernoy	vernoy.aSep08		575	404		134	trinucleotide repeat containing 15 CRA c (vernoy) mRNA.
verpor	verpor.aSep08		2984	397		113	inhibitor of Bruton tyrosine kinase (verpor) mRNA.
versa	versa.aSep08		1271	733	4	164	histone deacetylase 10 (versa) alternative variant aSep08, mRNA.
versa	versa.bSep08		869	799	1	77	histone deacetylase 10 (8.8 kD) (versa) alternative variant bSep08, mRNA.
vershee	vershee.aSep08		8876	647		80	CRA b like (8.7 kD) (vershee) mRNA.
vertu	vertu.aSep08		2623	286		65	putative protein (7.4 kD) (vertu) mRNA.
vervar	vervar.aSep08		6081	1670	6	516	finger castor 1 (vervar) alternative variant aSep08, mRNA.
verwey	verwey.aSep08		46678	512		40	putative protein (4.3 kD) (verwey) mRNA.
veybor	veybor.aSep08		2543	564		48	putative protein (veybor) mRNA.
veyby	veyby.aSep08		889	395		28	putative protein (veyby) mRNA.
veychy	veychy.aSep08		13670	724	7	241	ubiquitin protein ligase E3 component n-recognin 1 (veychy) alternative variant aSep08, mRNA.
veychy	veychy.bSep08		8617	375	4	97	ubiquitin protein ligase E3 component n-recognin 1 (veychy) alternative variant bSep08, mRNA.
veydar	veydar.aSep08		652	377		36	putative protein (veydar) mRNA.

veydoy	veydoy.aSep08		39517	704		83	putative protein (veydoy) mRNA.
veyflo	veyflo.aSep08		3535	451		34	putative protein (3.9 kD) (veyflo) mRNA.
veyflu	veyflu.aSep08		79095	302		93	putative protein (veyflu) mRNA.
veyfly	veyfly.aSep08		1487	335		55	putative protein (veyfly) mRNA.
veygar	veygar.aSep08		85806	440		146	putative protein of metazoan origin (veygar) mRNA.
veyja	veyja.aSep08		1309	842	3	109	putative endoplasmic reticulum protein (12.0 kD) (veyja) alternative variant aSep08, mRNA.
veyjey	veyjey.aSep08		8297	650		216	nuclear transcription factor X-box binding-like 1 (veyjey) mRNA.
veykee	veykee.aSep08		545	359		55	putative protein (veykee) mRNA.
veykler	veykler.aSep08		12788	754		118	putative nuclear protein (12.6 kD) (veykler) mRNA.
veylo	veylo.aSep08		7461	739		219	CRA a (veylo) mRNA.
veymee	veymee.aSep08		397	246	1	67	putative protein (veymee) alternative variant aSep08, mRNA.
veymee	veymee.bSep08		87164	933	6	107	putative protein (12.2 kD) (veymee) alternative variant bSep08, mRNA.
veynoy	veynoy.aSep08		8478	875		291	GRB10 interacting GYF protein 2 (veynoy) mRNA.
veypor	veypor.aSep08		774	683		161	putative protein (18.0 kD) (veypor) mRNA.
veysa	veysa.aSep08		1143	229		61	mitogen-activated protein kinase 11 (veysa) mRNA.
veyshee	veyshee.bSep08		1786	1460	1	80	putative protein (8.9 kD) (veyshee) alternative variant bSep08, mRNA.
veytu	veytu.aSep08		1240	450		105	putative protein (veytu) mRNA.
veyvar	veyvar.aSep08		1927	597	1	84	CRA a like (veyvar) alternative variant aSep08, mRNA.
veyvar	veyvar.bSep08		4219	588	2	79	apoptosis-inducing domain 1 (veyvar) alternative variant bSep08, mRNA.
veywey	veywey.aSep08		1933	369		61	putative protein (veywey) mRNA.
VeZF1andCuedc1	VeZF1andCuedc1.bSep08	287615	6379	3358	2	184	zinc finger 1 (VeZF1andCuedc1) alternative variant bSep08, mRNA.
VeZF1andCuedc1	VeZF1andCuedc1.bSep08	303419	6379	3358	2	184	zinc finger 1 (VeZF1andCuedc1) alternative variant bSep08, mRNA.
VeZF1andCuedc1	VeZF1andCuedc1.cSep08	287615	4778	748	4	55	CUE domain-containing 1 like (6.3 kD) (VeZF1andCuedc1) alternative variant cSep08, mRNA.
VeZF1andCuedc1	VeZF1andCuedc1.cSep08	303419	4778	748	4	55	CUE domain-containing 1 like (6.3 kD) (VeZF1andCuedc1) alternative variant cSep08, mRNA.
VeZt	VeZt.bSep08	299738	50435	1335	8	411	vezatin, adherens junctions transmembrane protein (VeZt) alternative variant bSep08, mRNA.
VeZt	VeZt.cSep08	299738	16955	1176	5	363	vezatin, adherens junctions transmembrane protein (VeZt) alternative variant cSep08, mRNA.
VeZt	VeZt.eSep08	299738	4023	617	2	41	vezatin, adherens junctions transmembrane protein (VeZt) alternative variant eSep08, mRNA.
Vgf	Vgf.bSep08	29461	3302	2798	1	617	VGF nerve growth factor inducible (68.2 kD) (Vgf) alternative variant bSep08, mRNA.
Vgll1	Vgll1.aSep08	363508	3328	468		112	vestigial like 1 homolog (Drosophila) (Vgll1) mRNA.
Vgll3	Vgll3.aSep08	498038	21974	297		98	vestigial like 3 (Drosophila) (Vgll3) mRNA.

Vgll4	Vgll4.bSep08	297523	12698	841	4	212	vestigial like 4 (Drosophila) (Vgll4) alternative variant bSep08, mRNA.
Vgll4	Vgll4.cSep08	297523	110302	793	5	171	vestigial like 4 (Drosophila) (Vgll4) alternative variant cSep08, mRNA.
Vgll4	Vgll4.dSep08	297523	109284	613	5	155	vestigial like 4 (Drosophila) (Vgll4) alternative variant dSep08, mRNA.
Vgll4	Vgll4.eSep08	297523	854	536	2	139	vestigial like 4 (Drosophila) (Vgll4) alternative variant eSep08, mRNA.
Vhlh	Vhlh.cSep08	24874	4479	329	3	35	von Hippel-Lindau syndrome homolog (Vhlh) alternative variant cSep08, mRNA.
VHP.0	VHP.0.aSep08		7166	1232	7	335	supervillin (VHP.0) alternative variant aSep08, mRNA.
Vil1	Vil1.bSep08	316521	3509	762	6	253	villin 1 (Vil1) alternative variant bSep08, mRNA.
Vill	Vill.aSep08	301057	4153	1001	5	268	villin-like (Vill) alternative variant aSep08, mRNA.
Vill	Vill.bSep08	301057	3530	763	5	254	villin-like (Vill) alternative variant bSep08, mRNA.
Vill	Vill.cSep08	301057	2542	636	3	207	villin-like (Vill) alternative variant cSep08, mRNA.
Vill	Vill.dSep08	301057	1330	609	4	172	villin-like (Vill) alternative variant dSep08, mRNA.
Vim	Vim.bSep08	81818	4395	906	4	301	vimentin (Vim) alternative variant bSep08, mRNA.
Vim	Vim.cSep08	81818	3627	838	5	172	vimentin (Vim) alternative variant cSep08, mRNA.
Vim	Vim.dSep08	81818	5817	479	5	159	vimentin (Vim) alternative variant dSep08, mRNA.
Vip	Vip.aSep08	117064	8083	1502	1	219	vasoactive intestinal polypeptide (Vip) alternative variant aSep08, mRNA.
Vip	Vip.bSep08	117064	8079	1495	1	216	vasoactive intestinal polypeptide (Vip) alternative variant bSep08, mRNA.
Visa	Visa.bSep08	311430	12039	888	5	227	virus-induced signaling adapter (Visa) alternative variant bSep08, mRNA.
Visa	Visa.cSep08	311430	7384	487	4	132	virus-induced signaling adapter (Visa) alternative variant cSep08, mRNA.
Visa	Visa.eSep08	311430	6235	390	4	43	virus-induced signaling adapter (Visa) alternative variant eSep08, mRNA.
Visa	Visa.fSep08	311430	676	472	2	38	virus-induced signaling adapter (Visa) alternative variant fSep08, mRNA.
Vit	Vit.aSep08	313831	2514	974		159	vitrin (Vit) mRNA.
Vkorc111	Vkorc111.bSep08	399684	49057	1628	3	130	vitamin K epoxide reductase complex, subunit 1-like 1 (Vkorc111) alternative variant bSep08, mRNA.
Vmo1	Vmo1.aSep08	360553	6713	455	2	151	vitelline membrane outer layer 1 homolog (chicken) (Vmo1) alternative variant aSep08, mRNA.
Vmo1	Vmo1.bSep08	360553	1250	937	1	105	vitelline membrane outer layer 1 homolog (chicken) (11.6 kD) (Vmo1) alternative variant bSep08, mRNA.
Vnn1	Vnn1.bSep08	29142	4470	755	1	172	vanin 1 (Vnn1) alternative variant bSep08, mRNA.
voby	voby.aSep08		999	367		63	protein CRA a (voby) mRNA.
vochy	vochy.aSep08		6601	463	2	118	glucosidase alpha (vochy) alternative variant aSep08, mRNA.
vochy	vochy.bSep08		5264	1411	2	79	glucosidase alpha (8.9 kD) (vochy) alternative variant bSep08, mRNA.

vochy	vochy.cSep08		4878	379	2	73	glucosidase alpha (vochy) alternative variant cSep08, mRNA.
vodar	vodar.aSep08		13063	497		52	putative protein (5.9 kD) (vodar) mRNA.
vodoy	vodoy.aSep08		9104	727		242	niemann-pick C1 (vodoy) mRNA.
voflo	voflo.aSep08		659	248		36	putative protein (4.3 kD) (voflo) mRNA.
voflu	voflu.aSep08		29468	730		76	putative mitochondrial protein (8.4 kD) (voflu) mRNA.
vogar	vogar.aSep08		1174	330		24	putative protein (vogar) mRNA.
voja	voja.aSep08		1790	447		41	putative protein (4.7 kD) (voja) mRNA.
vojey	vojey.aSep08		4368	871		289	furry homolog-like (vojey) mRNA.
vokee	vokee.aSep08		35878	402		27	putative protein (2.9 kD) (vokee) mRNA.
vokler	vokler.aSep08		27848	734	3	169	putative protein of metazoan origin (vokler) alternative variant aSep08, mRNA.
vokler	vokler.bSep08		14932	912	1	155	putative protein of metazoan origin (vokler) alternative variant bSep08, mRNA.
volo	volo.aSep08		2781	358		91	putative protein (volo) mRNA.
Vom2r3	Vom2r3.bSep08	502213	12363	552	2	48	vomeronasal 2 receptor, 3 and similar to vomeronasal 2, receptor, 10 (Vom2r3) alternative variant bSep08, mRNA.
Vom2r3	Vom2r3.bSep08	679643	12363	552	2	48	vomeronasal 2 receptor, 3 and similar to vomeronasal 2, receptor, 10 (Vom2r3) alternative variant bSep08, mRNA.
Vom2r57	Vom2r57.bSep08	299683	12239	738	2	113	vomeronasal 2 receptor, 57 (Vom2r57) alternative variant bSep08, mRNA.
Vom2r57	Vom2r57.cSep08	299683	570	294	2	89	vomeronasal 2 receptor, 57 (Vom2r57) alternative variant cSep08, mRNA.
vomee	vomee.aSep08		568	358		19	putative protein (vomee) mRNA.
vonoy	vonoy.aSep08		1237	1030			
vopor	vopor.aSep08		1753	417		41	putative protein (vopor) mRNA.
vorbor	vorbor.aSep08		1181	888	2	112	CRA c (12.7 kD) (vorbor) alternative variant aSep08, mRNA.
vorbor	vorbor.cSep08		11193	697	4	49	putative protein of mammalian origin (vorbor) alternative variant cSep08, mRNA.
vorbor	vorbor.dSep08		11054	609	5	33	putative protein of mammalian origin (3.9 kD) (vorbor) alternative variant dSep08, mRNA.
vorbor	vorbor.eSep08		10673	504	4	49	CRA b like (5.4 kD) (vorbor) alternative variant eSep08, mRNA.
vorby	vorby.aSep08		12378	588		196	solute carrier family 9 (vorby) mRNA.
vorchy	vorchy.aSep08		10424	857		285	ubiquitin protein ligase E3 component n-recognin 1 (vorchy) mRNA.
vordar	vordar.aSep08		8407	380		93	putative protein (vordar) mRNA.
vordoy	vordoy.aSep08		2132	260		40	putative protein (4.5 kD) (vordoy) mRNA.
vorflo	vorflo.aSep08		836	596		155	hepatocellular carcinoma-associated antigen MHCA108 like (vorflo) mRNA.
vorflu	vorflu.aSep08		641	534	2	65	putative protein (vorflu) alternative variant aSep08, mRNA.
vorfly	vorfly.aSep08		152142	1973		588	CRA d (vorfly) alternative variant aSep08, mRNA.
vorfly	vorfly.bSep08		7466	459		81	putative protein, with a coiled coil domain, of mammalian origin (vorfly) alternative variant bSep08, mRNA.

vorgar	vorgar.aSep08		3301	785		76	putative protein (8.4 kD) (vorgar) mRNA.
vorja	vorja.bSep08		20677	549	3	91	putative protein (9.2 kD) (vorja) alternative variant bSep08, mRNA.
vorjey	vorjey.aSep08		2103	402		94	nuclear transcription factor X-box binding-like 1 CRA d (vorjey) mRNA.
vorkee	vorkee.aSep08		1385	362		119	putative protein (vorkee) mRNA.
vorkler	vorkler.aSep08		8576	754		118	putative protein (12.6 kD) (vorkler) mRNA.
vorlo	vorlo.aSep08		8762	886		190	lemur tyrosine kinase 2 CRA a (vorlo) alternative variant aSep08, mRNA.
vormee	vormee.aSep08		7703	521		82	putative protein (vormee) mRNA.
vornoy	vornoy.aSep08		5728	2140		150	grb10 interacting GYF protein 2 (vornoy) mRNA.
vorpor	vorpor.aSep08		1030	338		58	putative protein (vorpor) mRNA.
vorsa	vorsa.aSep08		1451	807		37	putative protein (4.0 kD) (vorsa) mRNA.
vorshee	vorshee.bSep08		1428	529	2	37	putative protein (4.2 kD) (vorshee) alternative variant bSep08, mRNA.
vortu	vortu.aSep08		2789	555		43	putative protein (4.8 kD) (vortu) mRNA.
vorvar	vorvar.aSep08		564	522		58	putative protein (6.0 kD) (vorvar) mRNA.
vorwey	vorwey.aSep08		9214	374		28	putative protein (vorwey) mRNA.
voshee	voshee.aSep08		1493	426		142	IQ motif containing GTPase activating protein 3 (voshee) mRNA.
votu	votu.aSep08		508	333	2	111	putative protein of mammalian origin (votu) alternative variant aSep08, mRNA.
vovar	vovar.aSep08		811	477		69	putative protein (vovar) mRNA.
vowey	vowey.aSep08		11543	1657	3	376	putative protein (vowey) alternative variant aSep08, mRNA.
vowey	vowey.dSep08		12767	266	2	27	putative protein (3.0 kD) (vowey) alternative variant dSep08, mRNA.
voybor	voybor.aSep08		25075	698		76	putative protein (8.5 kD) (voybor) mRNA.
voyby	voyby.aSep08		1961	1099		138	putative mitochondrial protein of mammalian origin (15.2 kD) (voyby) mRNA.
voychy	voychy.aSep08		9483	551	3	183	transmembrane protein 62 CRA b (voychy) alternative variant aSep08, mRNA.
voychy	voychy.bSep08		2052	434	1	138	transmembrane protein 62 (voychy) alternative variant bSep08, mRNA.
voydar	voydar.aSep08		9099	474		66	calcium binding protein like (voydar) mRNA.
voydoy	voydoy.aSep08		17627	274		91	putative protein of mammalian origin (voydoy) mRNA.
voyflo	voyflo.aSep08		21423	376		84	putative protein (voyflo) mRNA.
voyflu	voyflu.aSep08		5708	2124	2	127	putative endoplasmic reticulum protein of eukaryotic origin (13.8 kD) (voyflu) alternative variant aSep08, mRNA.
voyfly	voyfly.aSep08		4651	536		178	CRA c (voyfly) mRNA.
voygar	voygar.aSep08		7903	1536		125	putative mitochondrial protein of eukaryotic origin (14.4 kD) (voygar) mRNA.
voyja	voyja.aSep08		10406	716		125	putative protein (13.7 kD) (voyja) mRNA.
voyjey	voyjey.aSep08		60485	551	5	122	ATPase class V type 10D (voyjey) alternative variant aSep08, mRNA.

voykee	voykee.aSep08		2777	1210		100	putative protein (voykee) mRNA.
voykler	voykler.aSep08		18770	736		36	putative protein (4.0 kD) (voykler) mRNA.
voylo	voylo.aSep08		8440	539		53	lemur tyrosine kinase 2 CRA a (voylo) mRNA.
voymee	voymee.aSep08		2710	350		85	putative protein (voymee) mRNA.
voynoy	voynoy.aSep08		6558	499		166	neuronal guanine nucleotide exchange factor (voynoy) mRNA.
voypor	voypor.aSep08		10762	610		202	dopey family member 1 (voypor) mRNA.
voysa	voysa.aSep08		3285	954		317	C-jun-amino-terminal kinase-interacting protein 2 (voysa) mRNA.
voyshee	voyshee.aSep08		15110	402		18	putative protein (voyshee) mRNA.
voytu	voytu.aSep08		19763	1885		44	putative protein (4.7 kD) (voytu) mRNA.
voyvar	voyvar.aSep08		3834	413		131	ubiquitination factor E4B (voyvar) mRNA.
voywey	voywey.aSep08		9467	708		83	putative protein (voywey) mRNA.
Vprbp	Vprbp.aSep08	315987	44058	394		96	vpr (HIV-1) binding protein (Vprbp) mRNA.
Vpreb3	Vpreb3.aSep08	365550	1492	742	2	122	pre-B lymphocyte gene 3 (13.5 kD) (Vpreb3) alternative variant aSep08, mRNA.
Vps4a	Vps4a.cSep08	246772	5372	380	3	32	vacuolar protein sorting 4a (yeast) (Vps4a) alternative variant cSep08, mRNA.
Vps4b	Vps4b.bSep08	360834	2046	402	1	72	vacuolar protein sorting 4b (yeast) (Vps4b) alternative variant bSep08, mRNA.
Vps8	Vps8.aSep08	287990	111144	1651		430	vacuolar protein sorting 8 homolog ( <i>S. cerevisiae</i> ) (Vps8) mRNA.
Vps11	Vps11.aSep08	315600	14417	3162	13	941	vacuolar protein sorting 11 (yeast) (107.8 kD) (Vps11) alternative variant aSep08, complete mRNA.
Vps11	Vps11.bSep08	315600	1758	442	1	147	vacuolar protein sorting 11 (yeast) (Vps11) alternative variant bSep08, mRNA.
Vps13a	Vps13a.bSep08	309243	12783	677	4	225	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant bSep08, mRNA.
Vps13a	Vps13a.cSep08	309243	28340	612	6	165	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant cSep08, mRNA.
Vps13a	Vps13a.dSep08	309243	7589	444	6	148	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant dSep08, mRNA.
Vps13a	Vps13a.eSep08	309243	4323	400	4	133	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant eSep08, mRNA.
Vps13a	Vps13a.fSep08	309243	20598	405	4	112	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant fSep08, mRNA.
Vps13a	Vps13a.gSep08	309243	7551	298	3	98	vacuolar protein sorting 13A (yeast) (Vps13a) alternative variant gSep08, mRNA.
Vps13d	Vps13d.bSep08	313825	87000	1961	11	539	vacuolar protein sorting 13D (yeast) (Vps13d) alternative variant bSep08, mRNA.
Vps13d	Vps13d.cSep08	313825	8714	523	5	173	vacuolar protein sorting 13D (yeast) (Vps13d) alternative variant cSep08, mRNA.
Vps13d	Vps13d.dSep08	313825	10336	719	3	126	vacuolar protein sorting 13D (yeast) (14.9 kD) (Vps13d) alternative variant dSep08, mRNA.
Vps13d	Vps13d.gSep08	313825	10039	575	4	74	vacuolar protein sorting 13D (yeast) (8.5 kD) (Vps13d) alternative variant gSep08, mRNA.



Vps16	Vps16.bSep08	296159	1797	840	7	280	vacuolar protein sorting 16 (yeast) (Vps16) alternative variant bSep08, mRNA.
Vps16	Vps16.cSep08	296159	1908	805	6	213	vacuolar protein sorting 16 (yeast) (Vps16) alternative variant cSep08, mRNA.
Vps16	Vps16.dSep08	296159	1926	1034	5	166	vacuolar protein sorting 16 (yeast) (18.9 kD) (Vps16) alternative variant dSep08, mRNA.
Vps16	Vps16.eSep08	296159	1248	554	4	85	vacuolar protein sorting 16 (yeast) (Vps16) alternative variant eSep08, mRNA.
Vps16	Vps16.fSep08	296159	749	667	2	82	vacuolar protein sorting 16 (yeast) (Vps16) alternative variant fSep08, mRNA.
Vps25	Vps25.aSep08	681059	5489	1094	6	176	vacuolar protein sorting 25 homolog (S. cerevisiae) (20.8 kD) (Vps25) alternative variant aSep08, complete mRNA.
Vps25	Vps25.bSep08	681059	4406	1762	5	161	vacuolar protein sorting 25 homolog (S. cerevisiae) (Vps25) alternative variant bSep08, mRNA.
Vps25	Vps25.cSep08	681059	2263	404	4	119	vacuolar protein sorting 25 homolog (S. cerevisiae) (Vps25) alternative variant cSep08, complete mRNA.
Vps25	Vps25.dSep08	681059	3951	973	6	100	vacuolar protein sorting 25 homolog (S. cerevisiae) (11.7 kD) (Vps25) alternative variant dSep08, mRNA.
Vps25	Vps25.eSep08	681059	3596	429	5	99	vacuolar protein sorting 25 homolog (S. cerevisiae) (Vps25) alternative variant eSep08, mRNA.
Vps25	Vps25.fSep08	681059	2984	286	4	95	vacuolar protein sorting 25 homolog (S. cerevisiae) (Vps25) alternative variant fSep08, mRNA.
Vps26a	Vps26a.bSep08	361846	17074	730	5	212	vacuolar protein sorting 26 homolog A (yeast) (Vps26a) alternative variant bSep08, mRNA.
Vps26b	Vps26b.bSep08	300472	16403	1329	1	364	vacuolar protein sorting 26 homolog B (S. pombe) (Vps26b) alternative variant bSep08, mRNA.
Vps28	Vps28.aSep08	300052	3774	923	10	228	vacuolar protein sorting 28 (26.1 kD) (Vps28) alternative variant aSep08, complete mRNA.
Vps28	Vps28.cSep08	300052	2047	679	6	120	vacuolar protein sorting 28 (Vps28) alternative variant cSep08, mRNA.
Vps28	Vps28.dSep08	300052	2238	721	5	65	vacuolar protein sorting 28 (7.3 kD) (Vps28) alternative variant dSep08, mRNA.
Vps28	Vps28.eSep08	300052	776	693	2	61	vacuolar protein sorting 28 (7.1 kD) (Vps28) alternative variant eSep08, mRNA.
Vps29	Vps29.bSep08	288666	3677	918	3	181	vacuolar protein sorting 29 (S. pombe) (20.4 kD) (Vps29) alternative variant bSep08, mRNA.
Vps33a	Vps33a.aSep08	65081	19388	3625		446	vacuolar protein sorting 33A (yeast) (Vps33a) mRNA.
Vps33b	Vps33b.bSep08	64060	1217	420	5	124	vacuolar protein sorting 33B (yeast) (Vps33b) alternative variant bSep08, mRNA.
Vps33b	Vps33b.cSep08	64060	4568	497	7	119	vacuolar protein sorting 33B (yeast) (Vps33b) alternative variant cSep08, mRNA.
Vps33b	Vps33b.dSep08	64060	1318	451	2	100	vacuolar protein sorting 33B (yeast) (Vps33b) alternative variant dSep08, mRNA.
Vps36	Vps36.bSep08	290851	14116	749	5	150	vacuolar protein sorting 36 (yeast) (17.3 kD) (Vps36) alternative variant bSep08, mRNA.
Vps36	Vps36.cSep08	290851	14840	545	6	71	vacuolar protein sorting 36 (yeast) (8.2 kD) (Vps36) alternative variant cSep08, complete mRNA.

Vps37b	Vps37b.bSep08	288659	20252	726	2	241	vacuolar protein sorting 37B (yeast) (Vps37b) alternative variant bSep08, mRNA.
Vps37c	Vps37c.aSep08	308178	25523	2637	4	353	vacuolar protein sorting 37C (yeast) (38.4 kD) (Vps37c) alternative variant aSep08, mRNA.
Vps37c	Vps37c.bSep08	308178	1857	755	1	180	vacuolar protein sorting 37C (yeast) (Vps37c) alternative variant bSep08, mRNA.
Vps39	Vps39.aSep08	362199	39856	4972	25	875	vacuolar protein sorting 39 (yeast) (100.6 kD) (Vps39) alternative variant aSep08, complete mRNA.
Vps39	Vps39.bSep08	362199	8961	610	5	169	vacuolar protein sorting 39 (yeast) (Vps39) alternative variant bSep08, mRNA.
Vps39	Vps39.cSep08	362199	9870	563	7	140	vacuolar protein sorting 39 (yeast) (Vps39) alternative variant cSep08, mRNA.
Vps45	Vps45.bSep08	64516	3521	1446	4	129	vacuolar protein sorting (Vps45) alternative variant bSep08, mRNA.
Vps45	Vps45.cSep08	64516	883	332	2	95	vacuolar protein sorting (Vps45) alternative variant cSep08, mRNA.
Vps52	Vps52.bSep08	25218	2099	765	6	254	vacuolar protein sorting 52 CRA a (Vps52) alternative variant bSep08, mRNA.
Vps52	Vps52.cSep08	25218	1870	561	5	169	vacuolar protein sorting 52 CRA d (Vps52) alternative variant cSep08, mRNA.
Vps52	Vps52.dSep08	25218	1665	1083	4	131	vacuolar protein sorting 52 CRA a (14.7 kD) (Vps52) alternative variant dSep08, mRNA.
Vps52	Vps52.fSep08	25218	1033	775	2	63	vacuolar protein sorting 52 (7.4 kD) (Vps52) alternative variant fSep08, mRNA.
Vps53	Vps53.bSep08	287535	124750	2671	21	803	vacuolar protein sorting 53 (yeast) (91.1 kD) (Vps53) alternative variant bSep08, complete mRNA.
Vps53	Vps53.cSep08	287535	31107	739	6	199	vacuolar protein sorting 53 (yeast) (Vps53) alternative variant cSep08, mRNA.
Vps53	Vps53.dSep08	287535	57187	762	8	129	vacuolar protein sorting 53 (yeast) (Vps53) alternative variant dSep08, mRNA.
Vps54	Vps54.bSep08	286932	8912	856	6	161	vacuolar protein sorting 54 (yeast) (Vps54) alternative variant bSep08, mRNA.
Vps54	Vps54.cSep08	286932	3452	545	2	49	vacuolar protein sorting 54 (yeast) (Vps54) alternative variant cSep08, mRNA.
Vps72	Vps72.bSep08	310661	11555	1182	5	197	vacuolar protein sorting 72 (yeast) (22.5 kD) (Vps72) alternative variant bSep08, complete mRNA.
Vrk1	Vrk1.bSep08	362779	51166	898	9	271	vaccinia related kinase 1 (Vrk1) alternative variant bSep08, mRNA.
Vrk1	Vrk1.cSep08	362779	51073	1204	10	261	vaccinia related kinase 1 (Vrk1) alternative variant cSep08, mRNA.
Vrk1	Vrk1.dSep08	362779	50421	734	8	214	vaccinia related kinase 1 (Vrk1) alternative variant dSep08, mRNA.
Vrk1	Vrk1.eSep08	362779	44368	793	6	124	vaccinia related kinase 1 (Vrk1) alternative variant eSep08, mRNA.
Vrk1	Vrk1.gSep08	362779	31337	417	4	91	vaccinia related kinase 1 (Vrk1) alternative variant gSep08, mRNA.

Vrk2	Vrk2.aSep08	360991	171197	1559	12	504	vaccinia related kinase 2 (Vrk2) alternative variant aSep08, mRNA.
Vrk3	Vrk3.bSep08	361565	13065	706	6	178	vaccinia related kinase 3 (Vrk3) alternative variant bSep08, mRNA.
Vrk3	Vrk3.cSep08	361565	11798	720	5	147	vaccinia related kinase 3 (Vrk3) alternative variant cSep08, mRNA.
Vrk3	Vrk3.dSep08	361565	9333	385	4	112	vaccinia related kinase 3 (Vrk3) alternative variant dSep08, mRNA.
Vrk3	Vrk3.eSep08	361565	1808	477	2	66	vaccinia related kinase 3 (Vrk3) alternative variant eSep08, mRNA.
Vrk3	Vrk3.fSep08	361565	2886	318	2	60	vaccinia related kinase 3 (Vrk3) alternative variant fSep08, mRNA.
Vsig4	Vsig4.aSep08	312102	23344	705		220	immunoglobulin V-set (25.0 kD) (Vsig4) mRNA.
Vstm2a	Vstm2a.aSep08	689106	25901	1079	3	252	CRA b (Vstm2a) alternative variant aSep08, mRNA.
Vstm2a	Vstm2a.bSep08	689106	22000	740	2	179	CRA b (19.9 kD) (Vstm2a) alternative variant bSep08, mRNA.
Vstm2a	Vstm2a.cSep08	689106	2135	693	1	164	putative protein of vertebrate origin (Vstm2a) alternative variant cSep08, mRNA.
Vta1	Vta1.aSep08	292640	47366	907	2	295	vps20-associated 1 homolog (S. cerevisiae) (Vta1) alternative variant aSep08, mRNA.
Vta1	Vta1.cSep08	292640	28801	1216	1	213	vps20-associated 1 homolog (S. cerevisiae) (23.1 kD) (Vta1) alternative variant cSep08, mRNA.
Vti1a	Vti1a.bSep08	65277	175651	578	7	123	vesicle transport through interaction with t-SNAREs homolog 1A (yeast) (Vti1a) alternative variant bSep08, mRNA.
Vti1a	Vti1a.cSep08	65277	139885	511	5	120	vesicle transport through interaction with t-SNAREs homolog 1A (yeast) (Vti1a) alternative variant cSep08, mRNA.
Vti1a	Vti1a.dSep08	65277	12313	444	3	63	vesicle transport through interaction with t-SNAREs homolog 1A (yeast) (Vti1a) alternative variant dSep08, mRNA.
Vtn	Vtn.bSep08	29169	1331	1166	1	201	vitronectin (22.5 kD) (Vtn) alternative variant bSep08, mRNA.
vuby	vuby.aSep08		5168	367	1	86	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 1 (vuby) alternative variant aSep08, mRNA.
vuby	vuby.bSep08		4891	419	2	70	putative protein (7.6 kD) (vuby) alternative variant bSep08, mRNA.
vuchy	vuchy.aSep08		420	335		111	spectrin beta non-erythrocytic 5 orthologue like (vuchy) mRNA.
vudar	vudar.aSep08		1789	409		48	putative protein (5.3 kD) (vudar) mRNA.
vudoy	vudoy.aSep08		1482	976		48	putative protein (4.9 kD) (vudoy) mRNA.
vuflo	vuflo.aSep08		2596	631		107	solute carrier family 18 member 2 (vuflo) mRNA.
vufly	vufly.aSep08		1278	615		49	polyprotein (5.5 kD) (vufly) mRNA.
vugar	vugar.aSep08		3961	402		90	putative protein (10.2 kD) (vugar) mRNA.
vuja	vuja.aSep08		2758	1525		31	putative protein (3.6 kD) (vuja) mRNA.

vujey	vujey.aSep08		3638	394		89	furry homolog-like (vujey) mRNA.
vukee	vukee.aSep08		586	382		127	putative protein (vukee) mRNA.
vukler	vukler.aSep08		11778	420		84	putative protein (vukler) mRNA.
vulo	vulo.aSep08		2567	307		95	putative protein (vulo) mRNA.
vumee	vumee.aSep08		12733	1373		432	folliculin interacting protein 1 (vumee) mRNA.
vunoy	vunoy.aSep08		4919	429		125	putative protein (vunoy) mRNA.
vupor	vupor.aSep08		4357	630		43	putative protein (5.0 kD) (vupor) mRNA.
vusa	vusa.aSep08		2026	582		70	putative protein (vusa) mRNA.
vushee	vushee.aSep08		625	367		78	putative protein (vushee) mRNA.
vutu	vutu.aSep08		3953	713		37	putative protein (4.4 kD) (vutu) mRNA.
vuvar	vuvar.aSep08		593	387		29	putative protein (3.3 kD) (vuvar) mRNA.
vuwey	vuwey.aSep08		1614	478	1	60	CRA a like (6.5 kD) (vuwey) alternative variant aSep08, mRNA.
vuwey	vuwey.bSep08		1543	374	1	60	CRA a like (6.5 kD) (vuwey) alternative variant bSep08, mRNA.
VWA.0	VWA.0.aSep08		4229	673		198	type VI alpha 1 (VWA.0) mRNA.
VWA.1	VWA.1.aSep08		19117	3611	4	927	collagen type VI alpha 3 (VWA.1) alternative variant aSep08, mRNA.
VWA.1	VWA.1.bSep08		13113	1881	2	436	type VI alpha 3 (VWA.1) alternative variant bSep08, mRNA.
VWA.2	VWA.2.aSep08		18557	1332		444	procollagen type XII alpha 1 CRA a (VWA.2) mRNA.
VWA.3	VWA.3.aSep08		67477	394		112	putative protein (VWA.3) alternative variant aSep08, mRNA.
VWA.3	VWA.3.bSep08		67661	740	1	108	putative mitochondrial protein (10.9 kD) (VWA.3) alternative variant bSep08, mRNA.
VWA.4	VWA.4.aSep08		8566	980		326	von Willebrand factor (VWA.4) mRNA.
Vwa1	Vwa1.aSep08	298683	3940	1089	3	344	von Willebrand factor, type A and fibronectin, type III (Vwa1) alternative variant aSep08, mRNA.
Vwa1	Vwa1.bSep08	298683	3633	711	3	159	von Willebrand factor, type A precursor (17.1 kD) (Vwa1) alternative variant bSep08, mRNA.
Vwa3a	Vwa3a.bSep08	293449	9246	556	4	155	CRA c (Vwa3a) alternative variant bSep08, mRNA.
Vwa3a	Vwa3a.cSep08	293449	6182	307	2	101	CRA c (Vwa3a) alternative variant cSep08, mRNA.
VWC.0	VWC.0.aSep08		84913	426		84	procollagen type V alpha 2 (VWC.0) mRNA.
VWC.1	VWC.1.aSep08		1872	750		249	cysteine-rich motor neuron 2 protein (VWC.1) mRNA.
VWC.2	VWC.2.aSep08		2972	473		157	cysteine rich bmp regulator 2 (VWC.2) mRNA.
VWC.3	VWC.3.aSep08		3120	791		197	thrombospondin 2 (VWC.3) mRNA.
Vwc2	Vwc2.bSep08	501231	130912	776	2	75	putative protein (8.5 kD) (Vwc2) alternative variant bSep08, mRNA.
Vwce	Vwce.aSep08	309209	3259	690		86	von Willebrand factor C and EGF domains (Vwce) mRNA.
VWD.0	VWD.0.aSep08		7964	623		207	CRA a (VWD.0) mRNA.
VWD.1	VWD.1.aSep08		4719	1214		404	CRA a (VWD.1) mRNA.
VWD.2	VWD.2.aSep08		2427	1660		553	IgG binding protein like (VWD.2) mRNA.
VWD.3	VWD.3.aSep08		1439	1103		367	fbp protein (VWD.3) mRNA.

VWD.4	VWD.4.aSep08		3279	773		208	fbp protein (VWD.4) mRNA.
VWD.5	VWD.5.aSep08		11989	1016	4	334	CRA a (VWD.5) alternative variant aSep08, mRNA.
Vwf	Vwf.aSep08	116669	20368	1712	3	570	von Willebrand factor homolog (Vwf) alternative variant aSep08, mRNA.
Vwf	Vwf.bSep08	116669	8149	798	1	228	von Willebrand factor homolog (Vwf) alternative variant bSep08, mRNA.
Vwf	Vwf.cSep08	116669	8164	735	2	225	von Willebrand factor homolog (Vwf) alternative variant cSep08, mRNA.
vyby	vyby.aSep08		5656	507		168	SWI SNF related matrix associated actin dependent regulator of chromatin subfamily a member 1 (vyby) mRNA.
vychy	vychy.aSep08		4768	1138		112	putative protein, with a coiled coil domain, of mammalian origin (vychy) mRNA.
vydar	vydar.aSep08		905	320		26	putative protein (vydar) mRNA.
vydoy	vydoy.aSep08		997	571	2	76	CRA a like (vydoy) alternative variant aSep08, mRNA.
vydoy	vydoy.bSep08		3774	1047	3	74	putative protein (vydoy) alternative variant bSep08, mRNA.
vyflo	vyflo.aSep08		3099	325		42	CRA b like (vyflo) mRNA.
vyflu	vyflu.aSep08		1585	410		55	putative protein (vyflu) mRNA.
vygar	vygar.aSep08		1219	542	3	140	putative protein (vygar) alternative variant aSep08, mRNA.
vygar	vygar.bSep08		1078	503	2	127	putative protein (vygar) alternative variant bSep08, mRNA.
vyja	vyja.aSep08		3187	199		63	gag protein like (vyja) mRNA.
vyjey	vyjey.aSep08		112861	621		120	putative protein (vyjey) mRNA.
vykee	vykee.aSep08		1956	241		79	putative protein (vykee) mRNA.
vykler	vykler.aSep08		833	717		53	putative protein (5.7 kD) (vykler) mRNA.
vylo	vylo.aSep08		7350	818		189	tyrosine kinase 3 (vylo) mRNA.
vymee	vymee.aSep08		11235	945		184	folliculin interacting protein 1 (vymee) mRNA.
vynoy	vynoy.aSep08		2382	718		61	putative protein (vynoy) mRNA.
vypor	vypor.aSep08		2646	373		50	putative protein (vypor) mRNA.
vysa	vysa.aSep08		21379	682		44	putative protein (vysa) mRNA.
vyshee	vyshee.aSep08		2118	607		167	insulin receptor-related receptor CRA a (vyshee) mRNA.
vytu	vytu.aSep08		44238	265		25	putative protein (3.0 kD) (vytu) mRNA.
vyvar	vyvar.aSep08		2359	720		145	putative protein human specific (vyvar) mRNA.
vywey	vywey.aSep08		1174	805		138	putative protein (vywey) mRNA.
wabor	wabor.aSep08		1343	539	2	80	putative protein (wabor) alternative variant aSep08, mRNA.
wabor	wabor.bSep08		29168	995	2	40	putative protein (wabor) alternative variant bSep08, mRNA.
wabor	wabor.cSep08		11829	627	3	63	putative protein (7.3 kD) (wabor) alternative variant cSep08, mRNA.
waby	waby.aSep08		1751	759		218	putative protein of mammalian origin (waby) mRNA.
wachy	wachy.aSep08		7007	687		163	tubulin gamma complex associated protein 4 (wachy) mRNA.
WAC_Acf1_DNA_bd.0	WAC_Acf1_DNA_bd.0.aSep08		22711	1398		210	bromodomain adjacent zinc finger domain 1B (WAC_Acf1_DNA_bd.0) mRNA.
wadar	wadar.aSep08		1414	386		122	putative protein (wadar) mRNA.

wadoy	wadoy.aSep08		19790	310		29	putative protein (wadoy) mRNA.
waflo	waflo.aSep08		3034	1091	3	143	dynamamin binding protein like (16.7 kD) (waflo) alternative variant aSep08, mRNA.
wafllu	wafllu.aSep08		5565	840		49	putative protein (wafllu) mRNA.
wafly	wafly.aSep08		5319	298		21	putative protein (2.4 kD) (wafly) mRNA.
wagar	wagar.aSep08		15964	1776		37	putative protein (4.0 kD) (wagar) mRNA.
waja	waja.aSep08		4567	403		78	putative protein (waja) mRNA.
wajey	wajey.aSep08		22115	1790		518	ATPase class V type 10D (wajey) mRNA.
wakee	wakee.aSep08		3367	354		64	polyprotein (wakee) mRNA.
wakler	wakler.aSep08		2509	732		44	putative protein (4.9 kD) (wakler) mRNA.
walo	walo.aSep08		1335	494		26	putative protein (walo) mRNA.
wamee	wamee.aSep08		5015	265		58	putative protein (wamee) mRNA.
wanoy	wanoy.aSep08		6019	442		49	putative protein (wanoy) mRNA.
wapor	wapor.aSep08		3674	356		118	dopey family member 1 (wapor) mRNA.
warbor	warbor.aSep08		664	381		94	putative protein (warbor) mRNA.
warby	warby.aSep08		11120	442		146	family member 2 (warby) mRNA.
warchy	warchy.aSep08		4370	1420		156	spastic paraplegia 11 (warchy) mRNA.
wardar	wardar.aSep08		1765	487		162	genetic suppressor element 1 (wardar) mRNA.
wardoy	wardoy.aSep08		983	272		51	putative protein (wardoy) mRNA.
warflu	warflu.aSep08		8512	429		110	odd Oz ten-m homolog 4 (warflu) mRNA.
warfly	warfly.aSep08		8233	1365	2	80	ac1576 like (9.1 kD) (warfly) mRNA.
wargar	wargar.aSep08		5651	430		143	palladin (wargar) mRNA.
warja	warja.aSep08		723	622		35	putative protein (warja) mRNA.
warjey	warjey.aSep08		8457	673		102	putative protein (11.4 kD) (warjey) mRNA.
warkee	warkee.aSep08		16389	423		85	putative nuclear protein (9.1 kD) (warkee) mRNA.
warkler	warkler.aSep08		803	478		91	putative protein (warkler) mRNA.
warlo	warlo.aSep08		930	482		84	CRA a like (warlo) mRNA.
warmee	warmee.aSep08		4223	398		15	putative protein (warmee) mRNA.
warnoy	warnoy.aSep08		663	275		22	putative protein (warnoy) mRNA.
warpor	warpor.aSep08		2418	388		129	CRA a (warpor) mRNA.
Wars	Wars.aSep08	314442	31195	1857	2	481	tryptophanyl-tRNA synthetase (54.1 kD) (Wars) alternative variant aSep08, complete mRNA.
Wars	Wars.bSep08	314442	6022	1665	1	111	tryptophanyl-tRNA synthetase (Wars) alternative variant bSep08, mRNA.
Wars2	Wars2.aSep08	690654	79582	542	4	180	tryptophanyl tRNA synthetase 2 (mitochondrial) (Wars2) alternative variant aSep08, mRNA.
Wars2	Wars2.bSep08	690654	3700	576	1	33	tryptophanyl tRNA synthetase 2 (mitochondrial) (Wars2) alternative variant bSep08, mRNA.
warsa	warsa.aSep08		18515	735		245	leucine-rich repeat kinase 2 (warsa) mRNA.
warshee	warshee.aSep08		1305	371		66	putative protein (warshee) mRNA.
wartu	wartu.aSep08		7609	826		80	putative protein (9.5 kD) (wartu) mRNA.
warvar	warvar.aSep08		2277	335		91	putative protein (9.9 kD) (warvar) mRNA.

warwey	warwey.aSep08		1627	417		139	putative protein of vertebrate origin (warwey) mRNA.
Was	Was.bSep08	317371	8728	1741	12	333	wiskott-Aldrich syndrome homolog (human) (36.2 kD) (Was) alternative variant bSep08, complete mRNA.
Was	Was.cSep08	317371	2817	763	7	240	wiskott-Aldrich syndrome homolog (human) (Was) alternative variant cSep08, mRNA.
wasa	wasa.aSep08		1316	458		49	putative protein (wasa) mRNA.
Wasf1	Wasf1.bSep08	294568	49882	722	5	180	WASP family 1 (Wasf1) alternative variant bSep08, mRNA.
washee	washee.aSep08		1271	568		67	putative protein (washee) mRNA.
watu	watu.aSep08		8735	730	3	175	tata box binding protein -associated factor RNA polymerase I b like (watu) alternative variant aSep08, mRNA.
watu	watu.bSep08		1177	665	1	35	putative protein (4.2 kD) (watu) alternative variant bSep08, mRNA.
wavar	wavar.aSep08		1725	1223	1	187	putative protein (21.1 kD) (wavar) alternative variant aSep08, mRNA.
wavar	wavar.bSep08		1814	1095	1	159	putative protein (18.0 kD) (wavar) alternative variant bSep08, mRNA.
wawbor	wawbor.aSep08		1301	732		52	putative protein (5.2 kD) (wawbor) mRNA.
wawby	wawby.aSep08		12755	705		69	putative protein (wawby) mRNA.
wawchy	wawchy.aSep08		25031	1767		537	spastic paraplegia 11 (wawchy) mRNA.
wawdar	wawdar.aSep08		1993	371		123	genetic suppressor element 1 (wawdar) mRNA.
wawdoy	wawdoy.aSep08		70720	714		60	putative protein (7.0 kD) (wawdoy) mRNA.
wawey	wawey.aSep08		10124	744		107	putative protein of eukaryotic origin (wawey) mRNA.
wawflu	wawflu.aSep08		1931	246		36	putative protein (4.1 kD) (wawflu) mRNA.
wawfly	wawfly.aSep08		11908	301		99	G protein-coupled receptor 126 (wawfly) mRNA.
wawgar	wawgar.aSep08		15908	864	1	68	putative protein (7.3 kD) (wawgar) alternative variant aSep08, mRNA.
wawgar	wawgar.bSep08		15920	659		68	putative protein (7.3 kD) (wawgar) alternative variant bSep08, mRNA.
wawja	wawja.aSep08		3198	781		100	putative protein (11.3 kD) (wawja) mRNA.
wawjey	wawjey.aSep08		40245	366		31	putative protein (wawjey) mRNA.
wawkee	wawkee.aSep08		1571	506	1	109	putative protein (wawkee) alternative variant aSep08, mRNA.
wawkee	wawkee.bSep08		1138	243	1	43	putative protein (wawkee) alternative variant bSep08, mRNA.
wawkler	wawkler.aSep08		2060	318		106	cytochrome p450 (wawkler) mRNA.
wawlo	wawlo.aSep08		3519	923	5	252	putative protein of eukaryotic origin (wawlo) alternative variant aSep08, mRNA.
wawmee	wawmee.aSep08		1271	650		151	obscurin (wawmee) mRNA.
wawnoy	wawnoy.aSep08		2642	251		51	putative protein (wawnoy) mRNA.
wawpor	wawpor.aSep08		1964	946	5	110	putative protein (12.1 kD) (wawpor) alternative variant aSep08, mRNA.
wawpor	wawpor.bSep08		1682	567	4	49	putative protein (wawpor) alternative variant bSep08, mRNA.
wawpor	wawpor.cSep08		789	353	3	47	putative protein (wawpor) alternative variant cSep08, mRNA.

wawpor	wawpor.dSep08		1793	1694	2	37	putative protein (4.1 kD) (wawpor) alternative variant dSep08, mRNA.
wawsa	wawsa.aSep08		5765	1625		120	ab2-143 like (12.8 kD) (wawsa) mRNA.
wawshee	wawshee.aSep08		1287	668		222	putative protein, with a transmembrane domain, of bilateral origin (wawshee) mRNA.
wawtu	wawtu.aSep08		409	264		34	putative protein (wawtu) mRNA.
wawvar	wawvar.bSep08		455	208	2	51	putative protein (wawvar) alternative variant bSep08, mRNA.
wawwey	wawwey.aSep08		17908	1090	2	78	putative protein (9.7 kD) (wawwey) alternative variant aSep08, mRNA.
wawwey	wawwey.bSep08		798	429	1	69	putative protein (wawwey) alternative variant bSep08, mRNA.
Wbp2	Wbp2.bSep08	192645	4082	675	7	224	WW domain binding protein 2 (Wbp2) alternative variant bSep08, mRNA.
Wbp2	Wbp2.cSep08	192645	7100	618	6	196	WW domain binding protein 2 (Wbp2) alternative variant cSep08, mRNA.
Wbp2	Wbp2.dSep08	192645	1471	1360	2	109	WW domain binding protein 2 (11.2 kD) (Wbp2) alternative variant dSep08, mRNA.
Wbp4	Wbp4.bSep08	114765	18087	2741	5	214	formin binding protein 21 like (23.9 kD) (Wbp4) alternative variant bSep08, mRNA.
Wbp4	Wbp4.cSep08	114765	15526	760	4	170	formin binding protein 21 like (18.9 kD) (Wbp4) alternative variant cSep08, mRNA.
Wbp4	Wbp4.dSep08	114765	7917	786	5	155	formin binding protein 21 like (17.6 kD) (Wbp4) alternative variant dSep08, mRNA.
Wbp5	Wbp5.aSep08	294067	2235	1012	1	182	WW domain binding protein 5 like (Wbp5) alternative variant aSep08, mRNA.
Wbp5	Wbp5.aSep08	680354	2235	1012	1	182	WW domain binding protein 5 like (Wbp5) alternative variant aSep08, mRNA.
Wbp7	Wbp7.aSep08	361543	5995	2515	10	734	WW domain binding protein 7 (Wbp7) alternative variant aSep08, mRNA.
Wbp7	Wbp7.bSep08	361543	3742	794	6	264	WW domain binding protein 7 (Wbp7) alternative variant bSep08, mRNA.
Wbp7	Wbp7.dSep08	361543	854	683	3	112	WW domain binding protein 7 (Wbp7) alternative variant dSep08, mRNA.
Wbp11	Wbp11.bSep08	297695	2715	721	3	144	WW domain binding protein 11 like (Wbp11) alternative variant bSep08, mRNA.
Wbp11	Wbp11.cSep08	297695	3414	467	2	86	WW domain binding protein 11 like (Wbp11) alternative variant cSep08, mRNA.
Wbp11	Wbp11.dSep08	297695	460	281	2	36	WW domain binding protein 11 like (Wbp11) alternative variant dSep08, mRNA.
Wbscr16	Wbscr16.bSep08	360796	7491	590	4	170	putative protein of bilateral origin (Wbscr16) alternative variant bSep08, mRNA.
Wbscr16	Wbscr16.cSep08	360796	17540	782	8	146	putative protein of ancient origin (Wbscr16) alternative variant cSep08, mRNA.
Wbscr16	Wbscr16.dSep08	360796	7528	668	3	117	regulator of chromosome condensation, RCC1 (Wbscr16) alternative variant dSep08, mRNA.



Wbscr16	Wbscr16.eSep08	360796	2433	364	3	37	putative protein (4.3 kD) (Wbscr16) alternative variant eSep08, mRNA.
Wbscr22	Wbscr22.aSep08	360830	8729	1172	10	254	methyltransferase type 11 and methyltransferase type 12 (Wbscr22) alternative variant aSep08, mRNA.
Wbscr22	Wbscr22.bSep08	360830	925	840	2	88	putative nuclear protein of mammalian origin (10.5 kD) (Wbscr22) alternative variant bSep08, mRNA.
Wbscr22	Wbscr22.cSep08	360830	2987	1555	4	77	putative protein of eukaryotic origin (Wbscr22) alternative variant cSep08, mRNA.
Wbscr27	Wbscr27.bSep08	688407	9414	467	4	104	williams-Beuren syndrome region 27 like (Wbscr27) alternative variant bSep08, mRNA.
Wbscr28	Wbscr28.aSep08	288604	1140	424		134	putative protein of mammalian origin (Wbscr28) mRNA.
WD40.0	WD40.0.aSep08		12649	515		171	WD repeat domain 44 (WD40.0) mRNA.
WD40.1	WD40.1.aSep08		1843	419		139	pwp2 periodic tryptophan protein homolog (WD40.1) mRNA.
WD40.2	WD40.2.aSep08		22850	3394	9	187	WD-40 repeat (WD40.2) alternative variant aSep08, mRNA.
WD40.2	WD40.2.bSep08		11382	2858	5	182	da1-6 (WD40.2) alternative variant bSep08, mRNA.
WD40.3	WD40.3.aSep08		11036	841		279	lysosomal trafficking regulator (WD40.3) mRNA.
WD40.5	WD40.5.aSep08		7918	408	4	135	hira (WD40.5) alternative variant aSep08, mRNA.
WD40.6	WD40.6.aSep08		14881	385	1	127	hira (WD40.6) alternative variant aSep08, mRNA.
WD40.6	WD40.6.bSep08		21890	789	2	88	histone cell cycle regulation defective homolog A (WD40.6) alternative variant bSep08, mRNA.
WD40.7	WD40.7.aSep08		2278	1165		388	WD repeat domain 24 (WD40.7) mRNA.
WD40.8	WD40.8.bSep08		5926	798	1	123	WD repeat domain 16 (13.6 kD) (WD40.8) alternative variant bSep08, mRNA.
WD40.10	WD40.10.aSep08		2779	1654	2	126	dmx-like 2 (WD40.10) alternative variant aSep08, mRNA.
WD40.11	WD40.11.aSep08		2437	391		130	guanine nucleotide exchange factor p532 (WD40.11) mRNA.
WD40.12	WD40.12.aSep08		3874	3518	5	958	WD repeat domain 6 (WD40.12) alternative variant aSep08, mRNA.
WD40.13	WD40.13.aSep08		2087	890		296	CRA c (WD40.13) mRNA.
WD40.14	WD40.14.aSep08		16603	790	7	263	WD repeat domain 48 (WD40.14) alternative variant aSep08, mRNA.
WD40.14	WD40.14.bSep08		6942	260	1	46	WD repeat domain 48 like (WD40.14) alternative variant bSep08, mRNA.
WD40.15	WD40.15.aSep08		128577	2093		583	WD repeat domain 51B (WD40.15) alternative variant aSep08, mRNA.
WD40.16	WD40.16.bSep08	685491	10793	581	5	66	retinoblastoma binding protein 4 like (8.1 kD) (WD40.16) alternative variant bSep08, complete mRNA.
WD40.17	WD40.17.bSep08		3417	541		113	CRA b (WD40.17) alternative variant bSep08, mRNA.
WD40.18	WD40.18.aSep08		5577	840	5	155	damage specific DNA binding protein 2 like (WD40.18) alternative variant aSep08, mRNA.
WD40.19	WD40.19.aSep08		1049	617		205	zinc finger protein 106 (WD40.19) mRNA.
WD40.20	WD40.20.aSep08		15438	737		245	CRA b (WD40.20) mRNA.

WD40.21	WD40.21.aSep08		74545	1579	5	526	WD-40 repeat (WD40.21) alternative variant aSep08, mRNA.
WD40.21	WD40.21.bSep08		3578	654	3	218	WD-40 repeat (WD40.21) alternative variant bSep08, mRNA.
WD40.22	WD40.22.aSep08		10180	955		306	WD repeat domain 62 (WD40.22) mRNA.
WD40.23	WD40.23.aSep08		9883	670		222	WD-40 repeat (WD40.23) mRNA.
WD40.24	WD40.24.aSep08		6833	801	5	266	CRA b (WD40.24) alternative variant aSep08, mRNA.
WD40.24	WD40.24.bSep08		936	812	1	83	putative protein of eukaryotic origin (WD40.24) alternative variant bSep08, mRNA.
WD40.25	WD40.25.aSep08		7249	580		165	WD-40 repeat (WD40.25) mRNA.
WD40.26	WD40.26.bSep08		703	278	2	86	guanine nucleotide-binding protein like (WD40.26) alternative variant bSep08, mRNA.
Wdfy1	Wdfy1.aSep08	301549	40364	707	7	220	WD-40 repeat (Wdfy1) alternative variant aSep08, mRNA.
Wdfy1	Wdfy1.cSep08	301549	10674	3536	5	157	zinc finger, FYVE-type and WD-40 repeat (Wdfy1) alternative variant cSep08, mRNA.
Wdfy1	Wdfy1.eSep08	301549	36455	647	6	102	WD-40 repeat (Wdfy1) alternative variant eSep08, mRNA.
Wdfy1	Wdfy1.fSep08	301549	85256	658	7	76	WD-40 repeat (8.5 kD) (Wdfy1) alternative variant fSep08, mRNA.
Wdfy3	Wdfy3.aSep08	305164	6533	3476	2	188	putative protein (Wdfy3) alternative variant aSep08, mRNA.
Wdfy3	Wdfy3.bSep08	305164	7326	787	4	151	alfy (Wdfy3) alternative variant bSep08, mRNA.
Wdfy3	Wdfy3.cSep08	305164	869	638	2	136	alfy (Wdfy3) alternative variant cSep08, mRNA.
Wdhd1	Wdhd1.bSep08	305827	6053	717	1	238	WD repeat and HMG-box DNA binding protein 1 (Wdhd1) alternative variant bSep08, mRNA.
Wdr1	Wdr1.bSep08	360950	21725	773	7	237	WD repeat domain 1 (Wdr1) alternative variant bSep08, mRNA.
Wdr1	Wdr1.cSep08	360950	11313	721	6	196	WD repeat domain 1 (Wdr1) alternative variant cSep08, mRNA.
Wdr1	Wdr1.dSep08	360950	1034	404	4	114	WD repeat domain 1 (Wdr1) alternative variant dSep08, mRNA.
Wdr1	Wdr1.eSep08	360950	1146	402	2	46	WD repeat domain 1 (Wdr1) alternative variant eSep08, mRNA.
Wdr3	Wdr3.bSep08	310720	4679	876	5	161	WD repeat domain 3 (Wdr3) alternative variant bSep08, mRNA.
Wdr3	Wdr3.cSep08	310720	3691	971	4	57	WD repeat domain 3 (Wdr3) alternative variant cSep08, mRNA.
Wdr7	Wdr7.bSep08	66031	19514	995	4	331	WD repeat domain 7 (Wdr7) alternative variant bSep08, mRNA.
Wdr12	Wdr12.bSep08	363237	7538	851	5	231	WD repeat domain 12 (Wdr12) alternative variant bSep08, mRNA.
Wdr12	Wdr12.cSep08	363237	19107	803	5	203	WD repeat domain 12 (Wdr12) alternative variant cSep08, mRNA.
Wdr12	Wdr12.dSep08	363237	19226	941	5	118	WD repeat domain 12 (13.7 kD) (Wdr12) alternative variant dSep08, mRNA.
Wdr12	Wdr12.eSep08	363237	49601	422	3	51	WD repeat domain 12 (Wdr12) alternative variant eSep08, mRNA.

Wdr13	Wdr13.bSep08	317370	2351	792	4	166	WD repeat domain 13 (Wdr13) alternative variant bSep08, mRNA.
Wdr18	Wdr18.bSep08	314617	5655	847	2	280	WD repeat domain 18 (Wdr18) alternative variant bSep08, mRNA.
Wdr18	Wdr18.cSep08	314617	5920	756	3	242	WD repeat domain 18 (26.0 kD) (Wdr18) alternative variant cSep08, mRNA.
Wdr18	Wdr18.dSep08	314617	6327	690	3	227	WD repeat domain 18 (Wdr18) alternative variant dSep08, mRNA.
Wdr19	Wdr19.aSep08	305349	12935	1784	9	391	WD repeat domain 19 (Wdr19) alternative variant aSep08, mRNA.
Wdr19	Wdr19.bSep08	305349	5708	768	5	178	WD repeat domain 19 (Wdr19) alternative variant bSep08, mRNA.
Wdr20a	Wdr20a.aSep08	314453	67026	2091	5	579	WD repeat domain 20a (Wdr20a) alternative variant aSep08, mRNA.
Wdr20a	Wdr20a.bSep08	314453	67292	2109	3	448	WD repeat domain 20a (Wdr20a) alternative variant bSep08, mRNA.
Wdr20a	Wdr20a.dSep08	314453	4877	503	3	88	WD repeat domain 20a (Wdr20a) alternative variant dSep08, mRNA.
Wdr22	Wdr22.aSep08	314273	88877	5318		864	WD repeat domain 22 (94.7 kD) (Wdr22) mRNA.
Wdr23	Wdr23.bSep08	305895	5081	1442	9	283	WD repeat domain 23 (32.0 kD) (Wdr23) alternative variant bSep08, mRNA.
Wdr23	Wdr23.cSep08	305895	5756	1451	11	258	WD repeat domain 23 (Wdr23) alternative variant cSep08, mRNA.
Wdr23	Wdr23.dSep08	305895	2064	978	4	127	WD repeat domain 23 (Wdr23) alternative variant dSep08, mRNA.
Wdr23	Wdr23.eSep08	305895	3639	768	5	118	WD repeat domain 23 (Wdr23) alternative variant eSep08, mRNA.
Wdr23	Wdr23.fSep08	305895	3040	1004	3	117	WD repeat domain 23 (Wdr23) alternative variant fSep08, mRNA.
Wdr23	Wdr23.hSep08	305895	1340	672	3	77	WD repeat domain 23 (Wdr23) alternative variant hSep08, mRNA.
Wdr24	Wdr24.aSep08	360497	1698	1301	1	365	WD repeat domain 24 (40.2 kD) (Wdr24) alternative variant aSep08, mRNA.
Wdr24	Wdr24.bSep08	360497	813	617	2	156	WD repeat domain 24 (Wdr24) alternative variant bSep08, mRNA.
Wdr25	Wdr25.aSep08	299317	46858	989		272	WD repeat domain 25 (Wdr25) mRNA.
Wdr26	Wdr26.bSep08	498301	24167	1002	9	334	WD repeat domain 26 (Wdr26) alternative variant bSep08, mRNA.
Wdr26	Wdr26.cSep08	498301	30897	1277	11	309	WD repeat domain 26 (35.5 kD) (Wdr26) alternative variant cSep08, mRNA.
Wdr26	Wdr26.dSep08	498301	12845	391	3	130	WD repeat domain 26 (Wdr26) alternative variant dSep08, mRNA.
Wdr27	Wdr27.aSep08	308222	11882	928		308	WD repeat domain 27 (Wdr27) mRNA.
Wdr33	Wdr33.bSep08	307524	24393	2664	7	609	WD repeat domain 33 (Wdr33) alternative variant bSep08, mRNA.
Wdr33	Wdr33.cSep08	307524	19606	971	5	323	WD repeat domain 33 (Wdr33) alternative variant cSep08, mRNA.

Wdr33	Wdr33.dSep08	307524	35108	1803	6	319	WD repeat domain 33 (37.4 kD) (Wdr33) alternative variant dSep08, complete mRNA.
Wdr33	Wdr33.eSep08	307524	7765	648	7	215	WD repeat domain 33 (Wdr33) alternative variant eSep08, mRNA.
Wdr34	Wdr34.bSep08	296618	4114	1783	5	365	WD repeat domain 34 (Wdr34) alternative variant bSep08, mRNA.
Wdr34	Wdr34.cSep08	296618	13675	754	5	222	WD repeat domain 34 (Wdr34) alternative variant cSep08, mRNA.
Wdr34	Wdr34.dSep08	296618	1615	673	3	198	WD repeat domain 34 (Wdr34) alternative variant dSep08, mRNA.
Wdr35	Wdr35.aSep08	298876	11574	1080		360	WD repeat domain 35 (Wdr35) mRNA.
Wdr37	Wdr37.bSep08	307075	16874	1586	6	412	WD repeat domain 37 (Wdr37) alternative variant bSep08, mRNA.
Wdr37	Wdr37.cSep08	307075	28931	3519	7	253	WD repeat domain 37 (Wdr37) alternative variant cSep08, mRNA.
Wdr37	Wdr37.dSep08	307075	19622	794	8	223	WD repeat domain 37 (Wdr37) alternative variant dSep08, mRNA.
Wdr37	Wdr37.fSep08	307075	9620	404	2	40	WD repeat domain 37 (Wdr37) alternative variant fSep08, mRNA.
Wdr38	Wdr38.aSep08	366035	1651	828		208	WD repeat domain 38 (Wdr38) mRNA.
Wdr41	Wdr41.aSep08	361879	50133	3089	13	460	WD repeat domain 41 (51.4 kD) (Wdr41) alternative variant aSep08, mRNA.
Wdr41	Wdr41.cSep08	361879	7933	513	4	137	WD repeat domain 41 (Wdr41) alternative variant cSep08, mRNA.
Wdr41	Wdr41.dSep08	361879	4149	732	4	113	WD repeat domain 41 CRA a (12.9 kD) (Wdr41) alternative variant dSep08, mRNA.
Wdr41	Wdr41.eSep08	361879	4357	1152	3	113	WD repeat domain 41 CRA a (12.9 kD) (Wdr41) alternative variant eSep08, complete mRNA.
Wdr41	Wdr41.fSep08	361879	43921	1064	10	110	WD repeat domain 41 (11.8 kD) (Wdr41) alternative variant fSep08, mRNA.
Wdr42a	Wdr42a.bSep08	364050	21100	1763	5	141	WD repeat domain 42A (Wdr42a) alternative variant bSep08, mRNA.
Wdr42a	Wdr42a.cSep08	364050	4932	872	3	92	WD repeat domain 42A (Wdr42a) alternative variant cSep08, mRNA.
Wdr43	Wdr43.aSep08	362703	29820	1801	16	511	CRA b (57.6 kD) (Wdr43) alternative variant aSep08, mRNA.
Wdr43	Wdr43.bSep08	362703	16637	837	6	278	WD repeat domain 43 like (Wdr43) alternative variant bSep08, mRNA.
Wdr43	Wdr43.cSep08	362703	12950	871	8	198	WD repeat domain 43 like (Wdr43) alternative variant cSep08, mRNA.
Wdr43	Wdr43.dSep08	362703	12519	848	10	178	WD repeat domain 43 like (Wdr43) alternative variant dSep08, mRNA.
Wdr43	Wdr43.eSep08	362703	6701	2097	4	121	WD repeat domain 43 like (13.8 kD) (Wdr43) alternative variant eSep08, mRNA.
Wdr44	Wdr44.aSep08	246152	50710	840	5	186	WD repeat domain 44 (Wdr44) alternative variant aSep08, mRNA.

Wdr45	Wdr45.aSep08	302559	5480	1385	11	360	WD repeat domain 45 (39.8 kD) (Wdr45) alternative variant aSep08, complete mRNA.
Wdr45	Wdr45.cSep08	302559	4207	624	7	157	WD repeat domain 45 (Wdr45) alternative variant cSep08, mRNA.
Wdr45l	Wdr45l.bSep08	360682	3134	622	4	138	wdr45 like (Wdr45l) alternative variant bSep08, mRNA.
Wdr45l	Wdr45l.cSep08	360682	3096	1539	3	122	wdr45 like (Wdr45l) alternative variant cSep08, mRNA.
Wdr47	Wdr47.aSep08	310785	77620	4106	14	529	repeat 47 (58.0 kD) (Wdr47) alternative variant aSep08, complete mRNA.
Wdr47	Wdr47.bSep08	310785	12191	1518	6	277	wd repeat domain 47 CRA b (30.1 kD) (Wdr47) alternative variant bSep08, mRNA.
Wdr47	Wdr47.dSep08	310785	7110	537	3	142	repeat 47 (Wdr47) alternative variant dSep08, mRNA.
Wdr47	Wdr47.fSep08	310785	14891	303	1	36	wd repeat domain 47 CRA b (Wdr47) alternative variant fSep08, mRNA.
Wdr48	Wdr48.aSep08	363164	14667	1573	1	429	WD repeat domain 48 (Wdr48) alternative variant aSep08, mRNA.
Wdr48	Wdr48.bSep08	363164	8076	2356	2	202	WD repeat domain 48 (Wdr48) alternative variant bSep08, mRNA.
Wdr55	Wdr55.bSep08	307494	2323	741	3	183	WD repeat domain 55 (Wdr55) alternative variant bSep08, mRNA.
Wdr55	Wdr55.cSep08	307494	1883	486	1	93	WD repeat domain 55 (Wdr55) alternative variant cSep08, mRNA.
Wdr59	Wdr59.aSep08	307855	51551	2478	7	389	WD repeat domain 59 (Wdr59) alternative variant aSep08, mRNA.
Wdr59	Wdr59.bSep08	307855	40693	617	3	141	WD repeat domain 59 (Wdr59) alternative variant bSep08, mRNA.
Wdr60	Wdr60.aSep08	314523	17873	1208	5	341	WD repeat domain 60 (Wdr60) alternative variant aSep08, mRNA.
Wdr60	Wdr60.bSep08	314523	6183	1585	1	53	WD repeat domain 60 (Wdr60) alternative variant bSep08, mRNA.
Wdr61	Wdr61.bSep08	363064	17733	1158	10	305	WD repeat domain 61 (Wdr61) alternative variant bSep08, complete mRNA.
Wdr61	Wdr61.cSep08	363064	11061	820	6	219	WD repeat domain 61 (Wdr61) alternative variant cSep08, mRNA.
Wdr61	Wdr61.dSep08	363064	5613	549	4	118	WD repeat domain 61 (Wdr61) alternative variant dSep08, mRNA.
Wdr61	Wdr61.eSep08	363064	6356	373	5	61	WD repeat domain 61 (Wdr61) alternative variant eSep08, mRNA.
Wdr62	Wdr62.aSep08	308492	11467	2454		789	WD repeat domain 62 (Wdr62) mRNA.
Wdr66	Wdr66.aSep08	304498	29181	722		240	WD repeat domain 66 (Wdr66) mRNA.
Wdr67	Wdr67.bSep08	299949	11011	668	5	222	WD repeat domain 67 (Wdr67) alternative variant bSep08, mRNA.
Wdr67	Wdr67.cSep08	299949	10123	581	3	128	WD repeat domain 67 (Wdr67) alternative variant cSep08, mRNA.
Wdr70	Wdr70.bSep08	294783	52062	739	6	212	WD repeat domain 70 (Wdr70) alternative variant bSep08, mRNA.

Wdr70	Wdr70.cSep08	294783	62225	647	8	102	WD repeat domain 70 (11.5 kD) (Wdr70) alternative variant cSep08, mRNA.
Wdr70	Wdr70.dSep08	294783	5878	368	4	87	WD repeat domain 70 (9.6 kD) (Wdr70) alternative variant dSep08, mRNA.
Wdr73	Wdr73.bSep08	308751	6080	745	6	180	WD repeat domain 73 (Wdr73) alternative variant bSep08, mRNA.
Wdr73	Wdr73.dSep08	308751	1154	687	2	109	WD repeat domain 73 (Wdr73) alternative variant dSep08, mRNA.
Wdr73	Wdr73.eSep08	308751	6468	775	7	102	WD repeat domain 73 (Wdr73) alternative variant eSep08, mRNA.
Wdr73	Wdr73.fSep08	308751	1741	654	5	28	WD repeat domain 73 (Wdr73) alternative variant fSep08, mRNA.
Wdr73	Wdr73.gSep08	308751	1793	420	5	52	WD repeat domain 73 (Wdr73) alternative variant gSep08, mRNA.
Wdr74	Wdr74.aSep08	690229	5288	1791	11	385	WD repeat domain 74 and hypothetical protein LOC690257 (42.7 kD) (Wdr74) alternative variant aSep08, complete mRNA.
Wdr74	Wdr74.aSep08	690257	5288	1791	11	385	WD repeat domain 74 and hypothetical protein LOC690257 (42.7 kD) (Wdr74) alternative variant aSep08, complete mRNA.
Wdr74	Wdr74.cSep08	690229	874	711	3	80	WD repeat domain 74 and hypothetical protein LOC690257 (Wdr74) alternative variant cSep08, mRNA.
Wdr74	Wdr74.cSep08	690257	874	711	3	80	WD repeat domain 74 and hypothetical protein LOC690257 (Wdr74) alternative variant cSep08, mRNA.
Wdr75	Wdr75.bSep08	314545	21764	1456	13	485	WD repeat domain 75 (Wdr75) alternative variant bSep08, mRNA.
Wdr75	Wdr75.cSep08	314545	6546	590	4	124	WD repeat domain 75 (Wdr75) alternative variant cSep08, mRNA.
Wdr75	Wdr75.dSep08	314545	3707	459	3	111	WD repeat domain 75 (Wdr75) alternative variant dSep08, mRNA.
Wdr75	Wdr75.fSep08	314545	771	473	2	58	WD repeat domain 75 (6.8 kD) (Wdr75) alternative variant fSep08, mRNA.
Wdr77	Wdr77.bSep08	310769	8152	1197	8	287	WD repeat domain 77 (30.8 kD) (Wdr77) alternative variant bSep08, mRNA.
Wdr77	Wdr77.cSep08	310769	2537	1347	3	119	WD repeat domain 77 (Wdr77) alternative variant cSep08, mRNA.
Wdr78	Wdr78.bSep08	313417	22651	1703	1	136	WD repeat domain 78 (14.7 kD) (Wdr78) alternative variant bSep08, mRNA.
Wdr79	Wdr79.bSep08	287432	16100	1181	5	279	WD repeat domain 79 (30.7 kD) (Wdr79) alternative variant bSep08, mRNA.
Wdr81	Wdr81.bSep08	303312	1875	736	1	245	WD repeat domain 81 (Wdr81) alternative variant bSep08, mRNA.
Wdr88	Wdr88.aSep08	292809	17522	710		236	WD repeat domain 88 (Wdr88) mRNA.
Wdr91	Wdr91.aSep08	312225	37324	2584	15	753	WD repeat domain 91 (Wdr91) alternative variant aSep08, mRNA.
Wdr91	Wdr91.cSep08	312225	2507	547	3	179	WD repeat domain 91 (Wdr91) alternative variant cSep08, mRNA.

Wdr91	Wdr91.dSep08	312225	10648	2341	5	151	WD repeat domain 91 (16.1 kD) (Wdr91) alternative variant dSep08, mRNA.
Wdr91	Wdr91.eSep08	312225	2935	794	3	138	WD repeat domain 91 (Wdr91) alternative variant eSep08, mRNA.
Wdsof1	Wdsof1.aSep08	362902	35349	1501	11	467	WD-40 repeat and sof1-like protein (Wdsof1) alternative variant aSep08, mRNA.
Wdsof1	Wdsof1.bSep08	362902	2469	664	2	36	putative protein of metazoan origin (4.2 kD) (Wdsof1) alternative variant bSep08, mRNA.
Wdsof1	Wdsof1.cSep08	362902	2226	674	2	31	putative protein of vertebrate origin (Wdsof1) alternative variant cSep08, mRNA.
Wdsub1	Wdsub1.bSep08	362137	5800	723	1	80	U box (9.2 kD) (Wdsub1) alternative variant bSep08, mRNA.
Wdte1	Wdte1.cSep08	313020	2252	1342	2	56	WD and tetratricopeptide repeats 1 (6.2 kD) (Wdte1) alternative variant cSep08, mRNA.
Wee1	Wee1.bSep08	308937	5946	574	4	162	wee 1 homolog (S. pombe) (Wee1) alternative variant bSep08, mRNA.
Wee1	Wee1.cSep08	308937	791	584	2	60	wee 1 homolog (S. pombe) (Wee1) alternative variant cSep08, mRNA.
weebor	weebor.aSep08		2343	825	3	84	putative protein (weebor) alternative variant aSep08, mRNA.
weeby	weeby.aSep08		510	399		48	CRA b like (weeby) mRNA.
weechy	weechy.aSep08		4152	282		93	tripartite motif-containing 69 (weechy) mRNA.
weedar	weedar.aSep08		1366	391		129	genetic suppressor element 1 (weedar) mRNA.
weedoy	weedoy.aSep08		22039	1169		93	putative protein (10.5 kD) (weedoy) mRNA.
weeflu	weeflu.aSep08		2527	1657		248	ubiquitin 35 (weeflu) mRNA.
weefly	weefly.aSep08		65254	687		57	putative protein (6.7 kD) (weefly) mRNA.
weegar	weegar.aSep08		4846	935		83	putative nuclear protein (9.4 kD) (weegar) mRNA.
weeja	weeja.aSep08		1465	385		77	CRA a like (weeja) mRNA.
weejey	weejey.aSep08		2341	265	2	43	putative protein (weejey) alternative variant aSep08, mRNA.
weekee	weekee.aSep08		4098	1121		38	putative protein (weekee) mRNA.
weekler	weekler.aSep08		25767	330		43	putative protein (4.9 kD) (weekler) mRNA.
weelo	weelo.aSep08		3054	764		254	putative protein of eukaryotic origin (weelo) mRNA.
weemee	weemee.aSep08		1550	1136		160	obscurin (weemee) mRNA.
weenoy	weenoy.aSep08		39260	414		53	putative protein (weenoy) mRNA.
weepor	weepor.aSep08		2373	385		77	putative cytoplasmic protein (8.7 kD) (weepor) mRNA.
weesa	weesa.aSep08		4831	692	2	111	ab2-143 like (11.9 kD) (weesa) alternative variant aSep08, mRNA.
weeshee	weeshee.aSep08		6177	299		52	putative protein (weeshee) mRNA.
weetu	weetu.aSep08		771	567		63	putative protein (weetu) mRNA.
weevar	weevar.aSep08		7564	309		42	putative protein (weevar) mRNA.
weewey	weewey.aSep08		29080	1088	2	97	putative mitochondrial protein (11.3 kD) (weewey) alternative variant aSep08, mRNA.
weewey	weewey.bSep08		21660	784	1	55	putative protein (6.0 kD) (weewey) alternative variant bSep08, mRNA.

werbor	werbor.aSep08		4681	2471		823	hla-b associated transcript 2 (werbor) mRNA.
werby	werby.aSep08		1666	830	2	126	zinc finger protein 185 (14.0 kD) (werby) alternative variant aSep08, mRNA.
werchy	werchy.aSep08		52765	387		43	putative protein (werchy) mRNA.
werdar	werdar.aSep08		981	775		158	genetic suppressor element 1 (18.2 kD) (werdar) mRNA.
werdoy	werdoy.aSep08		29483	769		16	putative protein (1.8 kD) (werdoy) mRNA.
werflu	werflu.aSep08		3376	412		137	ubiquitin 35 (werflu) mRNA.
werfly	werfly.aSep08		41821	449		50	putative protein (werfly) mRNA.
wergar	wergar.aSep08		17210	407		64	putative protein (wergar) mRNA.
werja	werja.aSep08		1337	305		101	mast cell protease (werja) mRNA.
werjey	werjey.aSep08		4475	833	2	64	putative protein (werjey) alternative variant aSep08, mRNA.
werkee	werkee.aSep08		1922	961		53	putative protein (6.0 kD) (werkee) mRNA.
werkler	werkler.aSep08		3537	737	2	86	putative cytoplasmic protein (9.8 kD) (werkler) alternative variant aSep08, mRNA.
werkler	werkler.bSep08		1028	244	1	25	putative protein (werkler) alternative variant bSep08, mRNA.
werlo	werlo.aSep08		2706	393		130	putative protein of metazoan origin (werlo) mRNA.
wermee	wermee.aSep08		2846	722		173	obscurin (wermee) mRNA.
wernoy	wernoy.aSep08		1422	300		42	putative protein (5.1 kD) (wernoy) mRNA.
werpor	werpor.aSep08		3839	755		122	putative protein of eukaryotic origin (werpor) mRNA.
wersa	wersa.aSep08		4208	2115		511	putative nuclear protein, with a coiled coil domain, of bilateral origin (58.6 kD) (wersa) mRNA.
wershee	wershee.aSep08		3966	760		53	putative protein (wershee) mRNA.
wertu	wertu.aSep08		6010	5331		194	putative protein (20.8 kD) (wertu) mRNA.
wervar	wervar.aSep08		911	813		123	putative protein (14.1 kD) (wervar) mRNA.
werwey	werwey.aSep08		4306	362		52	putative protein (5.9 kD) (werwey) mRNA.
weybor	weybor.aSep08		7404	271		20	putative protein (weybor) mRNA.
weyby	weyby.aSep08		2295	360		108	putative protein (weyby) mRNA.
weychy	weychy.aSep08		6792	661		112	putative protein, with a transmembrane domain (13.1 kD) (weychy) mRNA.
weydar	weydar.aSep08		3147	720		239	genetic suppressor element 1 (weydar) mRNA.
weydoy	weydoy.aSep08		12570	1778		99	nucleolar protein 4 (weydoy) mRNA.
weyflu	weyflu.aSep08		7244	587		100	ubiquitin 35 (weyflu) mRNA.
weyfly	weyfly.aSep08		25903	396		46	putative protein (5.1 kD) (weyfly) mRNA.
weygar	weygar.aSep08		16114	496		67	putative protein (7.6 kD) (weygar) mRNA.
weyja	weyja.aSep08		2650	484		86	ring finger protein 17 (weyja) mRNA.
weyjey	weyjey.aSep08		12882	922		307	WD repeat (weyjey) mRNA.
weykee	weykee.aSep08		1100	668		18	putative protein (2.3 kD) (weykee) mRNA.
weykler	weykler.aSep08		2575	716		58	putative protein (weykler) mRNA.
weylo	weylo.aSep08		20326	556		141	sidekick 1 (weylo) mRNA.
weymee	weymee.aSep08		7426	1784		452	putative protein (weymee) alternative variant aSep08, mRNA.



weymee	weymee.bSep08		3138	884		142	putative protein (15.1 kD) (weymee) alternative variant bSep08, mRNA.
weynoy	weynoy.aSep08		1814	260		47	putative protein (weynoy) mRNA.
weypor	weypor.aSep08		21650	740		215	putative protein (weypor) mRNA.
weysa	weysa.aSep08		3722	456		43	putative protein (weysa) mRNA.
weyshee	weyshee.aSep08		1625	384	3	128	phospholipid-transporting ATPase ID (weyshee) alternative variant aSep08, mRNA.
weytu	weytu.aSep08		1037	365		56	putative protein (weytu) mRNA.
weyvar	weyvar.aSep08		4243	248		27	putative protein (weyvar) mRNA.
weyvo	weyvo.aSep08		1622	1277		199	gag-pro-pol polyprotein (weyvo) mRNA.
weywey	weywey.aSep08		3814	286		94	putative protein (weywey) mRNA.
Wfdc1	Wfdc1.aSep08	171112	17240	992	1	296	WAP four-disulfide core domain 1 (Wfdc1) alternative variant aSep08, mRNA.
Wfdc2	Wfdc2.bSep08	286888	4658	751	3	119	WAP four-disulfide core domain 2 (12.6 kD) (Wfdc2) alternative variant bSep08, mRNA.
Wfdc2	Wfdc2.cSep08	286888	16190	616	2	44	WAP four-disulfide core domain 2 (Wfdc2) alternative variant cSep08, mRNA.
Wfdc2	Wfdc2.dSep08	286888	25414	871	4	54	WAP four-disulfide core domain 2 (Wfdc2) alternative variant dSep08, mRNA.
Wfdc2	Wfdc2.eSep08	286888	5429	752	4	59	WAP four-disulfide core domain 2 (6.4 kD) (Wfdc2) alternative variant eSep08, mRNA.
Wfdc3	Wfdc3.bSep08	296366	10548	586	2	120	WAP four-disulfide core domain 3 (12.7 kD) (Wfdc3) alternative variant bSep08, mRNA.
Wfdc6a	Wfdc6a.aSep08	685153	2883	982		94	WAP four-disulfide core domain 6A (Wfdc6a) mRNA.
Wfdc15a	Wfdc15a.bSep08	408226	1215	355	3	66	WAP four-disulfide core domain 15A (Wfdc15a) alternative variant bSep08, mRNA.
Wfikkn2	Wfikkn2.aSep08	287631	5992	3521		622	kazal immunoglobulin kunitz ntr domain-containing protein 2 (Wfikkn2) mRNA.
Whsc1	Whsc1.aSep08	680537	26724	3866	14	762	wolf-Hirschhorn syndrome candidate 1 like (Whsc1) alternative variant aSep08, mRNA.
Whsc1	Whsc1.bSep08	680537	18317	1271	7	377	wolf-Hirschhorn syndrome candidate 1 like (Whsc1) alternative variant bSep08, mRNA.
Whsc1	Whsc1.cSep08	680537	4270	2904	3	133	wolf-Hirschhorn syndrome candidate 1 like (Whsc1) alternative variant cSep08, mRNA.
Whsc1	Whsc1.eSep08	680537	1132	332	2	93	wolf-Hirschhorn syndrome candidate 1 like (Whsc1) alternative variant eSep08, mRNA.
Whsc111	Whsc111.bSep08	290831	11634	2415	3	231	wolf-Hirschhorn syndrome candidate 1-like 1 (human) (25.6 kD) (Whsc111) alternative variant bSep08, mRNA.
Whsc111	Whsc111.dSep08	290831	10339	530	4	118	wolf-Hirschhorn syndrome candidate 1-like 1 (human) (Whsc111) alternative variant dSep08, mRNA.
Whsc111	Whsc111.fSep08	290831	105208	1810	3	13	wolf-Hirschhorn syndrome candidate 1-like 1 (human) (Whsc111) alternative variant fSep08, mRNA.
Whsc2	Whsc2.bSep08	305455	3644	1388	6	284	wolf-Hirschhorn syndrome candidate 2 (human) (Whsc2) alternative variant bSep08, mRNA.
Whsc2	Whsc2.cSep08	305455	8149	610	3	167	wolf-Hirschhorn syndrome candidate 2 (human) (Whsc2) alternative variant cSep08, mRNA.

Wif1	Wif1.bSep08	114557	65014	763	7	253	wnt inhibitory factor 1 (Wif1) alternative variant bSep08, mRNA.
Wif1	Wif1.cSep08	114557	48505	802	8	224	wnt inhibitory factor 1 (Wif1) alternative variant cSep08, mRNA.
Wif1	Wif1.dSep08	114557	6106	404	2	34	wnt inhibitory factor 1 (Wif1) alternative variant dSep08, mRNA.
Wif1	Wif1.eSep08	114557	29127	313	3		
Wipf2	Wipf2.aSep08	360620	9761	1782		155	WAS/WASL interacting protein family, member 2 (Wipf2) mRNA.
Wipi1	Wipi1.bSep08	303630	19786	628	4	167	WD repeat domain, phosphoinositide interacting 1 (Wipi1) alternative variant bSep08, mRNA.
Wipi1	Wipi1.cSep08	303630	10781	266	2	61	WD repeat domain, phosphoinositide interacting 1 (Wipi1) alternative variant cSep08, mRNA.
Wisp1	Wisp1.bSep08	65154	21479	713	3	139	WNT1 inducible signaling pathway protein 1 (15.4 kD) (Wisp1) alternative variant bSep08, mRNA.
Wisp1	Wisp1.cSep08	65154	1130	662	2	121	WNT1 inducible signaling pathway protein 1 (Wisp1) alternative variant cSep08, mRNA.
Wisp2	Wisp2.bSep08	29576	10551	720		124	WNT1 inducible signaling pathway protein 2 (Wisp2) alternative variant bSep08, mRNA.
Wnk1	Wnk1.bSep08	116477	14231	4060	6	490	WNK lysine deficient protein kinase 1 (Wnk1) alternative variant bSep08, mRNA.
Wnk1	Wnk1.cSep08	116477	81767	1367	8	372	WNK lysine deficient protein kinase 1 (Wnk1) alternative variant cSep08, mRNA.
Wnk1	Wnk1.dSep08	116477	11249	767	4	197	WNK lysine deficient protein kinase 1 (Wnk1) alternative variant dSep08, mRNA.
Wnk1	Wnk1.eSep08	116477	2557	479	2	132	WNK lysine deficient protein kinase 1 (Wnk1) alternative variant eSep08, mRNA.
Wnk4	Wnk4.bSep08	287715	1448	1361	2	213	WNK lysine deficient protein kinase 4 (Wnk4) alternative variant bSep08, mRNA.
Wnk4	Wnk4.cSep08	287715	834	731	2	116	WNK lysine deficient protein kinase 4 (Wnk4) alternative variant cSep08, mRNA.
Wnt2	Wnt2.aSep08	114487	22643	738		246	wingless-related MMTV integration site 2 (Wnt2) mRNA.
Wnt2b	Wnt2b.aSep08	116466	6435	1245		215	wingless related MMTV integration site 2b (Wnt2b) mRNA.
Wnt4	Wnt4.aSep08	84426	18863	1717	2	508	wingless-related MMTV integration site 4 (Wnt4) alternative variant aSep08, complete mRNA.
Wnt4	Wnt4.cSep08	84426	17605	461	1	142	wingless-related MMTV integration site 4 (Wnt4) alternative variant cSep08, mRNA.
Wnt5b	Wnt5b.bSep08	282582	148258	2265	1	372	wingless-related MMTV integration site 5B (41.9 kD) (Wnt5b) alternative variant bSep08, mRNA.
Wnt7a	Wnt7a.cSep08	114850	2032	1283	2	79	wingless-related MMTV integration site 7A (Wnt7a) alternative variant cSep08, mRNA.
Wnt11	Wnt11.aSep08	140584	9213	551		183	wingless-related MMTV integration site 11 (Wnt11) mRNA.
wobor	wobor.aSep08		2975	385		38	putative protein (4.1 kD) (wobor) mRNA.
woby	woby.aSep08		53021	472	4	56	putative protein (woby) alternative variant aSep08, mRNA.
woby	woby.cSep08		23735	401	6	52	putative protein (woby) alternative variant cSep08, mRNA.
woby	woby.dSep08		6074	312	5	48	putative protein (woby) alternative variant dSep08, mRNA.

woby	woby.eSep08		6096	285	5	55	putative protein (woby) alternative variant eSep08, mRNA.
wochy	wochy.aSep08		16363	447	5	149	putative protein of metazoan origin (wochy) alternative variant aSep08, mRNA.
wodar	wodar.aSep08		3543	288		95	genetic suppressor element 1 (wodar) mRNA.
wodoy	wodoy.aSep08		934	743		69	putative protein (8.4 kD) (wodoy) mRNA.
woflu	woflu.aSep08		5082	445		60	prolylcarboxypeptidase (6.8 kD) (woflu) mRNA.
wofly	wofly.aSep08		1311	682		128	CRA a like (wofly) mRNA.
wogar	wogar.aSep08		7181	356		28	putative protein (wogar) mRNA.
woja	woja.aSep08		938	626		124	copine VI (woja) mRNA.
wojey	wojey.aSep08		6304	549	2	72	lipogenin like (8.3 kD) (wojey) alternative variant aSep08, mRNA.
wojey	wojey.bSep08		29890	386	2	51	putative protein (wojey) alternative variant bSep08, mRNA.
wokee	wokee.aSep08		1104	989		60	f11 receptor like (wokee) mRNA.
wokler	wokler.aSep08		6512	360	3	62	putative protein (7.1 kD) (wokler) alternative variant aSep08, mRNA.
wokler	wokler.bSep08		27948	570	4	85	GTPase activating protein testicular GAP1 (10.1 kD) (wokler) alternative variant bSep08, mRNA.
wolo	wolo.aSep08		6773	806	3	190	CRA b (wolo) alternative variant aSep08, mRNA.
wolo	wolo.bSep08		2908	406	3	135	CRA b (wolo) alternative variant bSep08, mRNA.
womee	womee.aSep08		1126	603		88	gem associated protein 5 (womee) mRNA.
wonoy	wonoy.aSep08		3090	439		59	putative protein (6.9 kD) (wonoy) mRNA.
wopor	wopor.aSep08		6600	585		52	putative protein (wopor) mRNA.
worbor	worbor.aSep08		1904	1071	2	132	putative protein (worbor) mRNA.
worby	worby.aSep08		495	280		41	putative protein (worby) mRNA.
worchy	worchy.aSep08		4107	760		69	putative protein (worchy) mRNA.
wordar	wordar.aSep08		2371	427		102	putative protein of vertebrate origin (wordar) mRNA.
wordoy	wordoy.aSep08		130311	407		135	nucleolar protein 4 (wordoy) mRNA.
worflu	worflu.aSep08		2775	349		69	putative protein (worflu) mRNA.
worfly	worfly.aSep08		21560	358		49	putative protein (worfly) mRNA.
worgar	worgar.aSep08		5947	402		39	putative protein (worgar) mRNA.
worja	worja.aSep08		13345	310		103	poly polymerase 4 (worja) mRNA.
worjey	worjey.aSep08		2396	421	4	140	WD repeat domain 19 (worjey) alternative variant aSep08, mRNA.
workee	workee.aSep08		1015	537		168	cell surface glycoprotein gp42 (workee) mRNA.
workler	workler.aSep08		5389	807		60	putative protein (6.7 kD) (workler) mRNA.
worlo	worlo.bSep08		1775	413	3	24	putative protein (worlo) alternative variant bSep08, mRNA.
worlo	worlo.cSep08		1677	267	2	20	putative protein (worlo) alternative variant cSep08, mRNA.
wormee	wormee.aSep08		1301	265		88	obscurin (wormee) mRNA.
wornoy	wornoy.aSep08		13491	442		24	putative protein (2.9 kD) (wornoy) alternative variant aSep08, mRNA.
worpor	worpor.aSep08		798	722		22	putative protein (worpor) mRNA.
worsa	worsa.aSep08		13818	260		84	putative protein (worsa) mRNA.

worshee	worshee.aSep08		1362	231		45	putative protein (worshee) mRNA.
wortu	wortu.aSep08		2664	456		43	putative protein (wortu) mRNA.
worvar	worvar.aSep08		6013	628		140	putative protein (worvar) mRNA.
worvo	worvo.aSep08		3756	637		50	lymphocyte antigen 96 like (worvo) mRNA.
worwey	worwey.bSep08		2219	1236	2	48	putative protein (5.5 kD) (worwey) alternative variant bSep08, mRNA.
worwey	worwey.cSep08		4120	853	2	35	putative protein (3.9 kD) (worwey) alternative variant cSep08, mRNA.
wosa	wosa.bSep08		3464	776	3	33	putative protein (4.0 kD) (wosa) alternative variant bSep08, mRNA.
woshee	woshee.aSep08		5171	638		53	putative protein (5.9 kD) (woshee) mRNA.
wotu	wotu.aSep08		28018	824	4	42	putative protein (4.7 kD) (wotu) alternative variant aSep08, mRNA.
wotu	wotu.bSep08		96841	423	2	49	putative protein (wotu) alternative variant bSep08, mRNA.
wovar	wovar.aSep08		4438	2203		107	putative protein (wovar) mRNA.
wowey	wowey.aSep08		33545	818		78	putative mitochondrial protein (8.4 kD) (wowey) alternative variant aSep08, mRNA.
woybor	woybor.aSep08		6404	335		49	putative protein (5.2 kD) (woybor) mRNA.
woyby	woyby.aSep08		1388	250		49	plexin B3 like (woyby) mRNA.
woychy	woychy.aSep08		6183	406	2	48	putative protein (woychy) alternative variant aSep08, mRNA.
woydar	woydar.aSep08		690	437	2	86	putative mitochondrial protein (9.6 kD) (woydar) alternative variant aSep08, mRNA.
woydoy	woydoy.aSep08		44788	331		25	putative protein (woydoy) mRNA.
woyflu	woyflu.aSep08		3644	360		48	putative protein (woyflu) mRNA.
woyfly	woyfly.aSep08		9125	769		66	putative protein (7.0 kD) (woyfly) mRNA.
woygar	woygar.aSep08		4040	702		41	putative protein (4.9 kD) (woygar) mRNA.
woyja	woyja.aSep08		4667	398		132	poly polymerase 4 (woyja) mRNA.
woyjey	woyjey.aSep08		1752	296		91	WD repeat domain 19 (woyjey) mRNA.
woykee	woykee.aSep08		27109	1801		585	putative protein (woykee) mRNA.
woykler	woykler.aSep08		10065	352		55	putative protein (woykler) mRNA.
woylo	woylo.aSep08		8482	1800			
woymee	woymee.aSep08		6779	519		50	putative protein (5.4 kD) (woymee) mRNA.
woynoy	woynoy.aSep08		39084	422	4	140	IQ containing AAA domain (woynoy) alternative variant aSep08, mRNA.
woypor	woypor.aSep08		2220	278		92	putative protein (woypor) mRNA.
woysa	woysa.aSep08		7032	292		97	putative protein (woysa) mRNA.
woyshee	woyshee.aSep08		24645	402		29	putative protein (3.3 kD) (woyshee) mRNA.
woytu	woytu.aSep08		24331	560	4	60	putative protein (woytu) alternative variant aSep08, mRNA.
woyvar	woyvar.aSep08		5931	592		196	calmodulin binding transcription activator 1 like (woyvar) mRNA.
woyvo	woyvo.aSep08		10449	606		97	putative protein (woyvo) mRNA.
woywey	woywey.aSep08		12046	682		48	putative protein (woywey) mRNA.

Wrb	Wrb.bSep08	288233	7701	729	6	126	tryptophan rich basic protein (14.4 kD) (Wrb) alternative variant bSep08, mRNA.
Wrb	Wrb.cSep08	288233	12649	2032	6	88	tryptophan rich basic protein (10.3 kD) (Wrb) alternative variant cSep08, mRNA.
Wrnip1	Wrnip1.bSep08	282835	19385	1724	1	474	werner helicase interacting protein 1 (Wrnip1) alternative variant bSep08, mRNA.
Wsb1	Wsb1.cSep08	303336	5647	691	5	121	WD repeat SOCS box-containing 1 (Wsb1) alternative variant cSep08, mRNA.
Wsb1	Wsb1.dSep08	303336	1317	717	3	75	WD repeat SOCS box-containing 1 (Wsb1) alternative variant dSep08, mRNA.
Wsb2	Wsb2.bSep08	288692	19791	1197	7	344	WD repeat and SOCS box-containing 2 (38.8 kD) (Wsb2) alternative variant bSep08, mRNA.
Wsb2	Wsb2.cSep08	288692	2706	1533	2	163	WD repeat and SOCS box-containing 2 (Wsb2) alternative variant cSep08, mRNA.
Wscd1	Wscd1.bSep08	287466	12490	370	2	119	putative protein (Wscd1) alternative variant bSep08, mRNA.
Wscd1	Wscd1.cSep08	287466	10371	510	2	74	putative protein (Wscd1) alternative variant cSep08, mRNA.
Wscd1	Wscd1.dSep08	287466	9169	536	2	60	putative protein (Wscd1) alternative variant dSep08, mRNA.
Wt1	Wt1.bSep08	24883	43651	2356	10	305	wilms tumor 1 (34.8 kD) (Wt1) alternative variant bSep08, mRNA.
Wt1	Wt1.cSep08	24883	2578	744	2	75	wilms tumor 1 (Wt1) alternative variant cSep08, mRNA.
Wtap	Wtap.bSep08	499020	2677	1047	2	258	wilms' tumour 1-associating protein (Wtap) alternative variant bSep08, mRNA.
Wtip	Wtip.aSep08	361552	33511	1273	6	245	WT1-interacting protein (Wtip) alternative variant aSep08, mRNA.
Wtip	Wtip.bSep08	361552	27163	744	5	212	WT1-interacting protein (Wtip) alternative variant bSep08, mRNA.
Wtip	Wtip.cSep08	361552	8404	1020	3	133	WT1-interacting protein (15.0 kD) (Wtip) alternative variant cSep08, mRNA.
wubor	wubor.aSep08		573	418		86	putative protein (wubor) mRNA.
wuby	wuby.aSep08		11356	573		190	ATPase class VI type 11 (wuby) mRNA.
wuchy	wuchy.aSep08		3452	1464		256	putative protein of vertebrate origin (28.5 kD) (wuchy) mRNA.
wudar	wudar.aSep08		575	333		110	putative protein (wudar) mRNA.
wudoy	wudoy.aSep08		2593	1955		124	desmocollin 2 CRA c (wudoy) mRNA.
wuflo	wuflo.aSep08		13910	1179		193	carboxypeptidase N polypeptide 1 (21.5 kD) (wuflo) mRNA.
wuflu	wuflu.aSep08		1947	1301	3	26	putative protein (wuflu) alternative variant aSep08, mRNA.
wuflly	wuflly.aSep08		56581	377		56	utrophin (wuflly) mRNA.
wugar	wugar.aSep08		1133	230		54	putative protein (wugar) mRNA.
wuja	wuja.aSep08		3304	921		199	leucine rich repeat containing 16b (wuja) alternative variant aSep08, mRNA.
wuja	wuja.bSep08		17244	1922		123	leucine rich repeat containing 16b (wuja) alternative variant bSep08, mRNA.

wujey	wujey.aSep08		42701	574		191	ATPase aminophospholipid transporter class I type 8A member 1 (wujey) mRNA.
wukee	wukee.aSep08		1081	373	2	86	CRA b (wukee) alternative variant aSep08, mRNA.
wukee	wukee.bSep08		875	657	1	66	CRA b (7.4 kD) (wukee) alternative variant bSep08, mRNA.
wukler	wukler.aSep08		1704	594		84	putative protein (9.1 kD) (wukler) mRNA.
wulo	wulo.aSep08		1237	549	3	60	putative protein (wulo) alternative variant aSep08, mRNA.
wumee	wumee.aSep08		31116	498		143	gem associated protein 5 CRA a (wumee) mRNA.
wunoy	wunoy.aSep08		1184	465		36	putative protein (wunoy) mRNA.
wupor	wupor.aSep08		10138	742		246	dopey family member 1 CRA b (wupor) mRNA.
wusa	wusa.aSep08		34608	851		51	putative protein (wusa) mRNA.
wushee	wushee.aSep08		3835	560		123	thrombospondin 3 (wushee) mRNA.
wutu	wutu.aSep08		1859	664	1	62	putative protein (6.9 kD) (wutu) alternative variant aSep08, mRNA.
wutu	wutu.bSep08		1815	615	1	62	putative protein (6.9 kD) (wutu) alternative variant bSep08, mRNA.
wuvar	wuvar.aSep08		5595	2697	8	448	calsyntenin 1 (wuvar) alternative variant aSep08, mRNA.
wuvar	wuvar.bSep08		1041	305	2	66	putative protein (wuvar) alternative variant bSep08, mRNA.
wuwey	wuwey.aSep08		30962	371		123	TEA domain family member 4 (wuwey) mRNA.
WW.0	WW.0.aSep08		39815	901		191	WW/Rsp5/WWP (WW.0) mRNA.
WW.1	WW.1.aSep08		59354	783	4	261	amyloid beta protein-binding family B member 2 like (WW.1) alternative variant aSep08, mRNA.
WW.1	WW.1.bSep08		38398	630	5	150	amyloid beta protein-binding family B member 2 like (WW.1) alternative variant bSep08, mRNA.
Wwc1	Wwc1.aSep08	303039	12866	1150	4	269	KIBRA (Wwc1) alternative variant aSep08, mRNA.
Wwc1	Wwc1.bSep08	303039	3563	1522	1	171	KIBRA (19.9 kD) (Wwc1) alternative variant bSep08, mRNA.
Wwox	Wwox.bSep08	292041	13303	717	5	138	WW domain-containing oxidoreductase (15.7 kD) (Wwox) alternative variant bSep08, mRNA.
Wwox	Wwox.cSep08	292041	36611	564	2	64	WW domain-containing oxidoreductase (Wwox) alternative variant cSep08, mRNA.
Wwp1	Wwp1.bSep08	297930	12530	730	6	208	CRA a (Wwp1) alternative variant bSep08, mRNA.
Wwp2	Wwp2.bSep08	291999	95605	1584	10	394	WW/Rsp5/WWP (42.7 kD) (Wwp2) alternative variant bSep08, mRNA.
Wwp2	Wwp2.cSep08	291999	1205	691	2	229	WW/Rsp5/WWP (Wwp2) alternative variant cSep08, mRNA.
Wwp2	Wwp2.dSep08	291999	8664	766	10	168	WW/Rsp5/WWP (19.8 kD) (Wwp2) alternative variant dSep08, mRNA.
Wwp2	Wwp2.eSep08	291999	1996	473	3	157	HECT (Wwp2) alternative variant eSep08, mRNA.
Wwp2	Wwp2.fSep08	291999	13649	393	3	130	putative protein of vertebrate origin (Wwp2) alternative variant fSep08, mRNA.
Wwp2	Wwp2.gSep08	291999	35786	325	3	108	putative protein (Wwp2) alternative variant gSep08, mRNA.
Wwp2	Wwp2.hSep08	291999	1325	1049	2	72	putative protein of eukaryotic origin (Wwp2) alternative variant hSep08, mRNA.
wybor	wybor.aSep08		583	277		92	class I (wybor) mRNA.

wyby	wyby.aSep08		4232	343		114	atpase class VI type 11C (wyby) mRNA.
wychy	wychy.aSep08		13992	2501	10	371	tubulin gamma complex associated protein 4 (wychy) alternative variant aSep08, mRNA.
wychy	wychy.bSep08		8006	715	7	238	tubulin gamma complex associated protein 4 (wychy) alternative variant bSep08, mRNA.
wydar	wydar.aSep08		9460	753		77	putative protein (8.6 kD) (wydar) mRNA.
wydoy	wydoy.aSep08		34726	364		32	putative protein (wydoy) mRNA.
wyflo	wyflo.aSep08		975	729		59	putative protein (wyflo) mRNA.
wyflu	wyflu.aSep08		375	277		36	putative protein (wyflu) mRNA.
wyfly	wyfly.aSep08		11921	588		196	utrophin (wyfly) mRNA.
wygar	wygar.aSep08		2200	316		36	putative protein (wygar) mRNA.
wyja	wyja.aSep08		2283	226		40	putative protein (wyja) mRNA.
wyjey	wyjey.aSep08		68757	832		175	ATPase aminophospholipid transporter class I type 8A member 1 (20.1 kD) (wyjey) mRNA.
wykee	wykee.aSep08		2707	377		49	putative protein (5.7 kD) (wykee) mRNA.
wykler	wykler.aSep08		4666	651		61	putative protein (wykler) mRNA.
wylo	wylo.aSep08		6837	2002		69	putative protein (7.6 kD) (wylo) mRNA.
wymee	wymee.aSep08		2306	565		187	gem associated protein 5 CRA a (wymee) mRNA.
wynoy	wynoy.aSep08		3138	267		72	putative protein (wynoy) mRNA.
wypor	wypor.aSep08		1006	363		120	dopey family member 1 (wypor) mRNA.
wysa	wysa.aSep08		3105	241		48	putative protein (5.0 kD) (wysa) mRNA.
wyshee	wyshee.aSep08		1093	705		160	putative protein (wyshee) mRNA.
wytu	wytu.aSep08		6083	411		71	putative protein (wytu) mRNA.
wyvar	wyvar.aSep08		6060	394		130	calsyntenin 1 (wyvar) mRNA.
wywey	wywey.aSep08		28435	418	3	138	TEA domain family member 4 (wywey) alternative variant aSep08, mRNA.
wywey	wywey.bSep08		4630	354	1	117	TEA domain family member 4 (wywey) alternative variant bSep08, mRNA.
Xab2	Xab2.bSep08	245976	1417	790	1	226	XPA binding protein 2 (Xab2) alternative variant bSep08, mRNA.
XAF1	XAF1.aSep08	679600	5846	470		156	XIAP associated factor-1 (XAF1) mRNA.
Xbp1	Xbp1.bSep08	289754	3563	1065	4	219	X-box binding protein 1 (24.9 kD) (Xbp1) alternative variant bSep08, mRNA.
Xbp1	Xbp1.cSep08	289754	2284	831	2	120	X-box binding protein 1 (13.6 kD) (Xbp1) alternative variant cSep08, mRNA.
Xbp1	Xbp1.dSep08	289754	3172	607	4	100	X-box binding protein 1 (Xbp1) alternative variant dSep08, mRNA.
Xirp2	Xirp2.aSep08	311098	15505	2046		415	xin actin-binding repeat containing 2 (Xirp2) mRNA.
Xlink.0	Xlink.0.aSep08		4768	2498	16	792	stabilin 1 (Xlink.0) alternative variant aSep08, mRNA.
Xlink.0	Xlink.0.bSep08		1795	730	7	243	stabilin 1 (Xlink.0) alternative variant bSep08, mRNA.
Xlink.0	Xlink.0.cSep08		947	625	3	184	stabilin 1 (Xlink.0) alternative variant cSep08, mRNA.
Xlink.0	Xlink.0.dSep08		477	386	1	93	stabilin 1 CRA c (Xlink.0) alternative variant dSep08, mRNA.

Xlink.0	Xlink.0.eSep08		1198	692	3	94	stabilin 1 (Xlink.0) alternative variant eSep08, mRNA.
Xlr4a	Xlr4a.aSep08	293841	3889	662		117	X-linked lymphocyte-regulated 4A (Xlr4a) mRNA.
Xpa	Xpa.bSep08	298074	24971	1314	6	235	xeroderma pigmentosum, complementation group A (27.4 kD) (Xpa) alternative variant bSep08, complete mRNA.
Xpc	Xpc.dSep08	312560	1802	816	2	116	xeroderma pigmentosum, complementation group C (13.1 kD) (Xpc) alternative variant dSep08, mRNA.
Xpnpep1	Xpnpep1.bSep08	170751	42748	1820	12	470	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (Xpnpep1) alternative variant bSep08, mRNA.
Xpnpep1	Xpnpep1.cSep08	170751	34007	1458	9	366	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (Xpnpep1) alternative variant cSep08, mRNA.
Xpnpep1	Xpnpep1.dSep08	170751	13722	1566	11	314	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (34.7 kD) (Xpnpep1) alternative variant dSep08, mRNA.
Xpnpep1	Xpnpep1.eSep08	170751	65048	668	6	211	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (Xpnpep1) alternative variant eSep08, mRNA.
Xpnpep1	Xpnpep1.gSep08	170751	839	718	2	75	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble (Xpnpep1) alternative variant gSep08, mRNA.
Xpnpep2	Xpnpep2.bSep08	117522	12125	764	8	254	X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound (Xpnpep2) alternative variant bSep08, mRNA.
Xpnpep2	Xpnpep2.cSep08	117522	9211	1130	8	228	X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound (Xpnpep2) alternative variant cSep08, mRNA.
Xpo1	Xpo1.bSep08	85252	3849	1438	2	178	exportin 1, CRM1 homolog (yeast) (Xpo1) alternative variant bSep08, mRNA.
Xpo6	Xpo6.bSep08	293476	19603	765	4	254	exportin 6 (Xpo6) alternative variant bSep08, mRNA.
Xpo6	Xpo6.cSep08	293476	7070	750	3	228	exportin 6 (Xpo6) alternative variant cSep08, mRNA.
Xpo6	Xpo6.dSep08	293476	16428	754	7	91	exportin 6 (Xpo6) alternative variant dSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.aSep08	290359	18005	1358	11	452	exportin 7 (Xpo7andNpm2) alternative variant aSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.aSep08	361070	18005	1358	11	452	exportin 7 (Xpo7andNpm2) alternative variant aSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.bSep08	290359	17338	2702	11	412	exportin 7 (Xpo7andNpm2) alternative variant bSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.bSep08	361070	17338	2702	11	412	exportin 7 (Xpo7andNpm2) alternative variant bSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.fSep08	290359	16316	397	3	58	exportin 7 CRA c (Xpo7andNpm2) alternative variant fSep08, mRNA.
Xpo7andNpm2	Xpo7andNpm2.fSep08	361070	16316	397	3	58	exportin 7 CRA c (Xpo7andNpm2) alternative variant fSep08, mRNA.
Xpot	Xpot.bSep08	314879	18912	1786	10	415	exportin, tRNA (nuclear export receptor for tRNAs) (Xpot) alternative variant bSep08, mRNA.
Xpot	Xpot.cSep08	314879	14157	1411	8	290	exportin, tRNA (nuclear export receptor for tRNAs) (Xpot) alternative variant cSep08, mRNA.
Xpot	Xpot.dSep08	314879	8993	748	7	249	exportin, tRNA (nuclear export receptor for tRNAs) (Xpot) alternative variant dSep08, mRNA.
Xpot	Xpot.eSep08	314879	11999	3127	4	90	exportin, tRNA (nuclear export receptor for tRNAs) (10.5 kD) (Xpot) alternative variant eSep08, mRNA.



Xpot	Xpot.fSep08	314879	1705	667	2	45	exportin, tRNA (nuclear export receptor for tRNAs) (Xpot) alternative variant fSep08, mRNA.
Xpr1	Xpr1.bSep08	289424	108496	2831	9	392	xenotropic and polytropic retrovirus receptor 1 (45.9 kD) (Xpr1) alternative variant bSep08, mRNA.
Xrcc1	Xrcc1.bSep08	84495	26843	2256	14	369	X-ray repair complementing 1 (Xrcc1) alternative variant bSep08, mRNA.
Xrcc1	Xrcc1.cSep08	84495	8717	589	5	196	repair protein (Xrcc1) alternative variant cSep08, mRNA.
Xrcc1	Xrcc1.dSep08	84495	6323	888	7	135	X-ray repair complementing 1 (15.3 kD) (Xrcc1) alternative variant dSep08, mRNA.
Xrcc1	Xrcc1.eSep08	84495	1852	1476	3	115	putative nuclear protein (12.4 kD) (Xrcc1) alternative variant eSep08, mRNA.
Xrcc1	Xrcc1.fSep08	84495	770	404	3	114	putative protein (Xrcc1) alternative variant fSep08, mRNA.
Xrcc4	Xrcc4.bSep08	309995	111530	619	4	142	X-ray repair complementing defective repair in Chinese hamster cells 4 (Xrcc4) alternative variant bSep08, mRNA.
Xrcc4	Xrcc4.cSep08	309995	26091	539	4	105	X-ray repair complementing defective repair in Chinese hamster cells 4 (Xrcc4) alternative variant cSep08, mRNA.
Xrcc4	Xrcc4.dSep08	309995	22761	690	2	68	X-ray repair complementing defective repair in Chinese hamster cells 4 (Xrcc4) alternative variant dSep08, mRNA.
Xrcc5	Xrcc5.bSep08	363247	44611	946	8	315	X-ray repair complementing defective in Chinese hamster cells 5 (Xrcc5) alternative variant bSep08, mRNA.
Xrcc5	Xrcc5.cSep08	363247	8754	776	5	205	X-ray repair complementing defective in Chinese hamster cells 5 (Xrcc5) alternative variant cSep08, mRNA.
Xrcc5	Xrcc5.dSep08	363247	10242	789	5	185	X-ray repair complementing defective in Chinese hamster cells 5 (Xrcc5) alternative variant dSep08, mRNA.
Xrcc5	Xrcc5.eSep08	363247	3749	461	2	102	X-ray repair complementing defective in Chinese hamster cells 5 (Xrcc5) alternative variant eSep08, mRNA.
Xrcc6	Xrcc6.bSep08	25019	2601	735	2	96	X-ray repair complementing defective repair in Chinese hamster cells 6 (10.6 kD) (Xrcc6) alternative variant bSep08, mRNA.
Xrn1	Xrn1.aSep08	300944	55096	2566	18	696	5'-3' exoribonuclease 1 (Xrn1) alternative variant aSep08, mRNA.
Xrn1	Xrn1.bSep08	300944	26042	711	8	236	5'-3' exoribonuclease 1 (Xrn1) alternative variant bSep08, mRNA.
Xrn1	Xrn1.cSep08	300944	1225	676	1	146	5'-3' exoribonuclease 1 (Xrn1) alternative variant cSep08, mRNA.
Xrn2	Xrn2.bSep08	362229	15244	586	6	195	5'-3' exoribonuclease 2 (Xrn2) alternative variant bSep08, mRNA.
Xrn2	Xrn2.cSep08	362229	6443	514	5	46	5'-3' exoribonuclease 2 (5.3 kD) (Xrn2) alternative variant cSep08, mRNA.
XRN_N.0	XRN_N.0.aSep08		8997	521		173	5'-3' exoribonuclease 1 (XRN_N.0) mRNA.
Xylb	Xylb.bSep08	316067	7090	747	5	151	xylokinase homolog (H. influenzae) (Xylb) alternative variant bSep08, mRNA.
Xylb	Xylb.cSep08	316067	4110	328	3	73	xylokinase homolog (H. influenzae) (Xylb) alternative variant cSep08, mRNA.
Xylt2	Xylt2.bSep08	64134	2928	758	3	252	xylosyltransferase II (Xylt2) alternative variant bSep08, mRNA.

Xylt2	Xylt2.cSep08	64134	2090	749	1	135	xylosyltransferase II (Xylt2) alternative variant cSep08, mRNA.
Yaf2	Yaf2.bSep08	690262	55074	1755	2	145	YY1 associated factor 2 (Yaf2) alternative variant bSep08, mRNA.
Yaf2	Yaf2.cSep08	690262	53932	809	4	109	YY1 associated factor 2 (Yaf2) alternative variant cSep08, mRNA.
Yaf2	Yaf2.dSep08	690262	34300	651	2	41	YY1 associated factor 2 (4.7 kD) (Yaf2) alternative variant dSep08, mRNA.
Yap1	Yap1.bSep08	363014	26310	887	1	238	yes-associated protein 1 (Yap1) alternative variant bSep08, mRNA.
Yars	Yars.bSep08	313047	10701	415	3	138	tyrosyl-tRNA synthetase (Yars) alternative variant bSep08, mRNA.
Ybx1	Ybx1.aSep08	500538	16649	1427	7	322	Y box protein 1 (35.7 kD) (Ybx1) alternative variant aSep08, mRNA.
Ybx1	Ybx1.bSep08	500538	12593	750	4	209	Y box protein 1 (Ybx1) alternative variant bSep08, mRNA.
Ybx1	Ybx1.cSep08	500538	723	382	1	95	Y box protein 1 (Ybx1) alternative variant cSep08, mRNA.
Ybx2	Ybx2.aSep08	303250	980	597		59	Y box protein 2 (Ybx2) mRNA.
Yc2	Yc2.bSep08	494500	26208	912	7	221	glutathione S-transferase Yc2 subunit (25.3 kD) (Yc2) alternative variant bSep08, mRNA.
Yc2	Yc2.cSep08	494500	21590	741	5	153	glutathione S-transferase Yc2 subunit (17.5 kD) (Yc2) alternative variant cSep08, mRNA.
YdjC	YdjC.aSep08	287938	1271	695	2	231	YdjC homolog (bacterial) (YdjC) alternative variant aSep08, mRNA.
Yeats2	Yeats2.bSep08	498112	8658	406	2	123	putative protein of vertebrate origin (Yeats2) alternative variant bSep08, mRNA.
Yeats4	Yeats4.bSep08	299810	7072	1784	1	64	putative protein (7.1 kD) (Yeats4) alternative variant bSep08, complete mRNA.
Yes1	Yes1.bSep08	24884	31829	1073	6	286	yamaguchi sarcoma viral (v-yes) oncogene homolog 1 (Yes1) alternative variant bSep08, mRNA.
Yif1b	Yif1b.aSep08	292768	10063	1794	1	598	yip1 interacting factor homolog B (S. cerevisiae) (Yif1b) alternative variant aSep08, mRNA.
Yif1b	Yif1b.dSep08	292768	8414	775	3	250	yip1 interacting factor homolog B (S. cerevisiae) (Yif1b) alternative variant dSep08, mRNA.
Yif1b	Yif1b.eSep08	292768	2019	667	1	82	yip1 interacting factor homolog B (S. cerevisiae) (9.2 kD) (Yif1b) alternative variant eSep08, mRNA.
Yipf1	Yipf1.bSep08	298312	13783	685	5	176	yip1 domain family member 1 like (19.5 kD) (Yipf1) alternative variant bSep08, mRNA.
Yipf1	Yipf1.cSep08	298312	21211	683	5	151	yip1 domain family member 1 like (Yipf1) alternative variant cSep08, mRNA.
Yipf1	Yipf1.dSep08	298312	18430	758	7	146	yip1 domain family member 1 like (16.6 kD) (Yipf1) alternative variant dSep08, mRNA.
Yipf1	Yipf1.eSep08	298312	5989	406	3	134	yip1 domain family member 1 (Yipf1) alternative variant eSep08, mRNA.
Yipf1	Yipf1.fSep08	298312	2702	730	3	127	yip1 domain family member 1 (Yipf1) alternative variant fSep08, mRNA.
Yipf1	Yipf1.gSep08	298312	8104	606	2	49	putative protein (5.4 kD) (Yipf1) alternative variant gSep08, mRNA.

Yipf2	Yipf2.bSep08	363027	2699	830	6	213	yip1 domain family, member 2 (Yipf2) alternative variant bSep08, mRNA.
Yipf2	Yipf2.cSep08	363027	2587	796	5	194	yip1 domain family, member 2 (Yipf2) alternative variant cSep08, mRNA.
Yipf2	Yipf2.dSep08	363027	8798	785	3	64	yip1 domain family, member 2 (Yipf2) alternative variant dSep08, mRNA.
Yipf2	Yipf2.fSep08	363027	512	429	2	37	yip1 domain family, member 2 (Yipf2) alternative variant fSep08, mRNA.
Yipf3	Yipf3.aSep08	301245	4261	1760	5	498	yip1 domain family, member 3 (Yipf3) alternative variant aSep08, mRNA.
Yipf3	Yipf3.cSep08	301245	1279	604	2	129	yip1 domain family, member 3 (14.0 kD) (Yipf3) alternative variant cSep08, mRNA.
Yipf4	Yipf4.bSep08	362699	7332	483	4	24	yip1 domain family, member 4 (Yipf4) alternative variant bSep08, mRNA.
Yipf5	Yipf5.bSep08	361315	11356	1120	5	168	yip1 domain family, member 5 (18.4 kD) (Yipf5) alternative variant bSep08, complete mRNA.
Yipf6	Yipf6.bSep08	363476	4587	518	5	83	yip1 domain family, member 6 (Yipf6) alternative variant bSep08, mRNA.
Yippee.0	Yippee.0.aSep08		12299	900	3	118	yippee-like 3 (13.5 kD) (Yippee.0) alternative variant aSep08, mRNA.
Yippee.0	Yippee.0.bSep08		9144	1808	1	51	putative protein (5.2 kD) (Yippee.0) alternative variant bSep08, mRNA.
Ykt6	Ykt6.bSep08	64351	8197	475	3	115	YKT6 homolog (S. Cerevisiae) (Ykt6) alternative variant bSep08, mRNA.
Ykt6	Ykt6.cSep08	64351	5134	2204	2	60	YKT6 homolog (S. Cerevisiae) (6.8 kD) (Ykt6) alternative variant cSep08, mRNA.
Ylpm1	Ylpm1.aSep08	299199	35179	2377	16	649	YLP motif-containing protein 1 (Ylpm1) alternative variant aSep08, mRNA.
Ylpm1	Ylpm1.bSep08	299199	6064	1405	3	126	YLP motif containing 1 (14.9 kD) (Ylpm1) alternative variant bSep08, mRNA.
Ylpm1	Ylpm1.cSep08	299199	8356	440	4	119	YLP motif containing 1 (Ylpm1) alternative variant cSep08, mRNA.
Ylpm1	Ylpm1.dSep08	299199	10499	903	3	72	ylp motif containing 1 (Ylpm1) alternative variant dSep08, mRNA.
Ylpm1	Ylpm1.eSep08	299199	21351	846	5	60	YLP motif-containing protein 1 like (Ylpm1) alternative variant eSep08, mRNA.
Yme111	Yme111.bSep08	114217	15545	697	6	218	YME1-like 1 (S. cerevisiae) (Yme111) alternative variant bSep08, mRNA.
Ypel4	Ypel4.bSep08	502643	770	612	2	62	yippee-like 4 (Drosophila) (6.9 kD) (Ypel4) alternative variant bSep08, mRNA.
Ypel5	Ypel5.aSep08	298792	9820	618	4	121	yippee-like 5 (Drosophila) (13.8 kD) (Ypel5) alternative variant aSep08, mRNA.
Ypel5	Ypel5.bSep08	298792	9651	768	3	121	yippee-like 5 (Drosophila) (13.8 kD) (Ypel5) alternative variant bSep08, mRNA.
Ypel5	Ypel5.cSep08	298792	13051	757	4	121	yippee-like 5 (Drosophila) (13.8 kD) (Ypel5) alternative variant cSep08, mRNA.

Ypel5	Ypel5.dSep08	298792	13247	784	3	121	yippee-like 5 (Drosophila) (13.8 kD) (Ypel5) alternative variant dSep08, mRNA.
Ypel5	Ypel5.eSep08	298792	4553	990	3	121	yippee-like 5 (Drosophila) (13.8 kD) (Ypel5) alternative variant eSep08, mRNA.
Ypel5	Ypel5.hSep08	298792	4234	333	2	34	yippee-like 5 (Drosophila) (Ypel5) alternative variant hSep08, mRNA.
Yt521	Yt521.bSep08	170956	14491	1412	11	359	splicing factor CRA a (Yt521) alternative variant bSep08, mRNA.
Yt521	Yt521.cSep08	170956	5170	749	3	136	splicing factor CRA c (Yt521) alternative variant cSep08, mRNA.
Yt521	Yt521.dSep08	170956	2622	1263	4	132	splicing factor CRA c (15.5 kD) (Yt521) alternative variant dSep08, mRNA.
Yt521	Yt521.eSep08	170956	4377	725	5	129	splicing factor CRA c (15.9 kD) (Yt521) alternative variant eSep08, mRNA.
Yt521	Yt521.fSep08	170956	11126	650	4	118	splicing factor CRA a (Yt521) alternative variant fSep08, mRNA.
Yt521	Yt521.gSep08	170956	4180	760	3	82	putative protein (Yt521) alternative variant gSep08, mRNA.
Ythdc2	Ythdc2.aSep08	307446	8681	970		267	putative protein of ancient origin (Ythdc2) mRNA.
Ythdf2	Ythdf2.aSep08	313053	25194	2629	2	605	YTH domain family 2 (Ythdf2) alternative variant aSep08, mRNA.
Ythdf3	Ythdf3.aSep08	361920	14338	4344	2	396	YTH domain family 3 (Ythdf3) alternative variant aSep08, mRNA.
Ythdf3	Ythdf3.bSep08	361920	20016	1153	4	182	YTH domain family 3 (Ythdf3) alternative variant bSep08, mRNA.
Ythdf3	Ythdf3.eSep08	361920	31225	769	4	52	YTH domain family 3 (5.8 kD) (Ythdf3) alternative variant eSep08, mRNA.
Ywhab	Ywhab.bSep08	56011	20956	788		127	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide (Ywhab) alternative variant bSep08, mRNA.
Ywhae	Ywhae.aSep08	29753	37995	2016	1	272	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide (Ywhae) alternative variant aSep08, mRNA.
Ywhah	Ywhah.bSep08	25576	8414	1804	1	113	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, eta polypeptide (12.1 kD) (Ywhah) alternative variant bSep08, mRNA.
Ywhaq	Ywhaq.bSep08	25577	26000	866	3	225	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide (Ywhaq) alternative variant bSep08, mRNA.
Ywhaq	Ywhaq.cSep08	25577	25318	355	3	95	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide (Ywhaq) alternative variant cSep08, mRNA.
Ywhaz	Ywhaz.aSep08	25578	18911	789	3	255	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide (Ywhaz) alternative variant aSep08, mRNA.
Yy1	Yy1.bSep08	24919	23648	700	2	160	YY1 transcription factor (18.0 kD) (Yy1) alternative variant bSep08, mRNA.

Y_phosphatase.0	Y_phosphatase.0.aSep08		13841	375		117	protein tyrosine phosphatase (Y_phosphatase.0) mRNA.
Y_phosphatase.1	Y_phosphatase.1.aSep08		10782	1303	3	179	protein tyrosine phosphatase receptor type D (Y_phosphatase.1) alternative variant aSep08, mRNA.
Y_phosphatase.1	Y_phosphatase.1.bSep08		2993	369	1	119	protein tyrosine phosphatase receptor type D (Y_phosphatase.1) alternative variant bSep08, mRNA.
Y_phosphatase.2	Y_phosphatase.2.aSep08		1786	895		133	protein tyrosine phosphatase (Y_phosphatase.2) mRNA.
zabor	zabor.aSep08		6696	881		85	putative protein of vertebrate origin (zabor) mRNA.
zaby	zaby.aSep08		635	404	3	134	plexin B1 CRA c (zaby) alternative variant aSep08, mRNA.
zachy	zachy.aSep08		3777	247		74	putative protein (zachy) mRNA.
zadar	zadar.aSep08		5305	735		66	putative protein (6.9 kD) (zadar) mRNA.
Zadh1	Zadh1.aSep08	299194	30292	1564	10	351	alcohol dehydrogenase, zinc-binding (38.1 kD) (Zadh1) alternative variant aSep08, mRNA.
zadoy	zadoy.aSep08		1485	588		74	putative protein (zadoy) mRNA.
zaflu	zaflu.aSep08		27938	215		59	putative protein (zaflu) mRNA.
zafly	zafly.aSep08		1456	472		95	putative nuclear protein (10.6 kD) (zafly) mRNA.
zagar	zagar.aSep08		2139	540	2	100	putative protein (zagar) alternative variant aSep08, mRNA.
zagar	zagar.bSep08		2074	538	2	51	putative protein (zagar) alternative variant bSep08, mRNA.
zaja	zaja.aSep08		16086	746		96	putative protein of eukaryotic origin (zaja) mRNA.
zajey	zajey.aSep08		25517	1833		113	putative protein of metazoan origin (zajey) mRNA.
zakee	zakee.aSep08		770	545		86	putative protein (9.7 kD) (zakee) mRNA.
zakler	zakler.aSep08		428	265		25	putative protein (zakler) mRNA.
zalo	zalo.aSep08		1275	782		100	putative protein (zalo) mRNA.
zamee	zamee.aSep08		6079	3285		81	target of 2 CRA f (zamee) mRNA.
zanoy	zanoy.aSep08		5536	914		43	putative protein (zanoy) mRNA.
Zap70	Zap70.bSep08	301348	1353	732	4	84	zeta-chain (TCR) associated protein kinase (Zap70) alternative variant bSep08, mRNA.
zapor	zapor.aSep08		7468	813		199	zic family member 4 (zapor) alternative variant aSep08, mRNA.
zarbor	zarbor.aSep08		2680	433		35	putative protein (3.6 kD) (zarbor) mRNA.
zarby	zarby.aSep08		2285	455		151	host cell factor (zarby) mRNA.
zarchy	zarchy.aSep08		26203	1470	10	490	uncharacterized Protein (zarchy) alternative variant aSep08, mRNA.
zardar	zardar.aSep08		1727	606		185	pecanex-like 2 (zardar) mRNA.
zardoy	zardoy.aSep08		27908	395		131	putative protein (zardoy) mRNA.
zarflu	zarflu.aSep08		10331	934	2	284	x-ray radiation resistance associated 1 (zarflu) alternative variant aSep08, mRNA.
zarflu	zarflu.bSep08		5908	758	1	252	x-ray radiation resistance associated 1 protein (zarflu) alternative variant bSep08, mRNA.
zarfly	zarfly.aSep08		1453	923		24	putative protein (zarfly) mRNA.
zargar	zargar.aSep08		6491	397		131	odd Oz ten-m homolog 3 (zargar) mRNA.
zarja	zarja.aSep08		2325	670		95	putative protein (zarja) mRNA.

zarjey	zarjey.aSep08		3498	502		89	putative protein (10.0 kD) (zarjey) mRNA.
zarkee	zarkee.aSep08		19632	562		65	putative protein (zarkee) mRNA.
zarkler	zarkler.aSep08		3903	620		43	putative protein (5.0 kD) (zarkler) mRNA.
zarlo	zarlo.aSep08		28064	1016	2	45	reverse transcriptase (zarlo) alternative variant aSep08, mRNA.
zarmee	zarmee.aSep08		18920	473		157	sperm antigen with calponin homology coiled-coil domains 1 like (zarmee) mRNA.
zarnoy	zarnoy.aSep08		1862	1297		31	putative protein (3.6 kD) (zarnoy) mRNA.
zarpor	zarpor.aSep08		19098	426		29	putative protein (zarpor) mRNA.
zarsa	zarsa.aSep08		989	640		30	putative protein (3.3 kD) (zarsa) complete mRNA.
zarshee	zarshee.aSep08		2489	1045		205	cingulin (zarshee) mRNA.
zartu	zartu.aSep08		1015	361		29	putative protein (zartu) mRNA.
zarvar	zarvar.aSep08		2007	777		259	nucleolar protein 9 (zarvar) mRNA.
zarvo	zarvo.aSep08		3950	298		34	putative protein (zarvo) mRNA.
zarwey	zarwey.aSep08		4564	337		29	putative protein (3.2 kD) (zarwey) mRNA.
zasa	zasa.aSep08		3956	2429		359	prickle homolog 1 (zasa) mRNA.
zashee	zashee.aSep08		1076	549		74	putative protein (zashee) mRNA.
zatu	zatu.aSep08		5899	748		66	putative protein (zatu) mRNA.
zavar	zavar.aSep08		1229	348	3	61	putative protein (zavar) alternative variant aSep08, mRNA.
zavo	zavo.aSep08		1649	569		63	putative protein (zavo) mRNA.
zawbor	zawbor.aSep08		9529	838		120	putative protein (12.6 kD) (zawbor) mRNA.
zawby	zawby.aSep08		990	467	1	45	putative protein (zawby) alternative variant aSep08, mRNA.
zawby	zawby.bSep08		936	451	1	58	CRA b like (6.6 kD) (zawby) alternative variant bSep08, mRNA.
zawchy	zawchy.aSep08		8219	1078		358	transient receptor potential cation channel subfamily M member 7 (zawchy) mRNA.
zawdar	zawdar.aSep08		27145	295		55	putative protein (6.0 kD) (zawdar) mRNA.
zawdoy	zawdoy.aSep08		25112	821		237	putative protein of vertebrate origin (zawdoy) mRNA.
zawey	zawey.bSep08		14497	886	3	108	putative nuclear protein (12.1 kD) (zawey) alternative variant bSep08, mRNA.
zawey	zawey.cSep08		10503	588	2	40	putative protein (4.7 kD) (zawey) alternative variant cSep08, mRNA.
zawflu	zawflu.aSep08		4300	410		136	CRA b (zawflu) mRNA.
zawfly	zawfly.aSep08		11950	748		9	putative cytoplasmic protein (8.6 kD) (zawfly) mRNA.
zawgar	zawgar.aSep08		3259	1256	2	64	putative protein (zawgar) alternative variant aSep08, mRNA.
zawja	zawja.aSep08		15051	346		71	putative protein (zawja) mRNA.
zawjey	zawjey.aSep08		13026	545		181	repeat-containing protein (zawjey) mRNA.
zawkee	zawkee.aSep08		4785	429		30	putative protein (3.4 kD) (zawkee) mRNA.
zawkler	zawkler.aSep08		1223	521		96	CRA a like (zawkler) mRNA.
zawlo	zawlo.aSep08		1087	463	2	52	putative protein (5.4 kD) (zawlo) alternative variant aSep08, mRNA.

zawmee	zawmee.aSep08		18222	2429	11	534	nuclear receptor co-repressor 1 (zawmee) alternative variant aSep08, mRNA.
zawmee	zawmee.bSep08		3934	740	3	246	nuclear receptor co-repressor 1 CRA b (zawmee) alternative variant bSep08, mRNA.
zawmee	zawmee.cSep08		9130	707	5	235	nuclear receptor co-repressor 1 CRA b (zawmee) alternative variant cSep08, mRNA.
zawmee	zawmee.dSep08		2203	318	2	105	nuclear receptor co-repressor 1 CRA a (zawmee) alternative variant dSep08, mRNA.
zawnoy	zawnoy.aSep08		16255	684		60	putative protein (6.7 kD) (zawnoy) mRNA.
zawpor	zawpor.aSep08		3208	1709		111	solute carrier family 9 (zawpor) mRNA.
zawsa	zawsa.aSep08		9844	332		110	putative protein (zawsa) mRNA.
zawshee	zawshee.aSep08		667	350		116	cingulin (zawshee) mRNA.
zawtu	zawtu.aSep08		9998	240		79	oligomeric golgi complex (zawtu) mRNA.
zawvar	zawvar.aSep08		2859	467		155	chromodomain helicase DNA binding protein 5 like (zawvar) mRNA.
zawvo	zawvo.aSep08		1707	698		29	putative protein (zawvo) mRNA.
zawwey	zawwey.aSep08		139952	358		51	putative protein (5.9 kD) (zawwey) mRNA.
Zbbx	Zbbx.aSep08	361964	41631	1785	2	562	putative protein of mammalian origin (Zbbx) alternative variant aSep08, mRNA.
Zbbx	Zbbx.cSep08	361964	2095	290	2	77	putative protein of mammalian origin (Zbbx) alternative variant cSep08, mRNA.
Zbp1	Zbp1.aSep08	171091	2550	681		103	Z-DNA binding protein 1 (Zbp1) mRNA.
Zbtb5	Zbtb5.aSep08	298084	20531	2945	3	692	BTB/POZ and zinc finger, C2H2-type (Zbtb5) alternative variant aSep08, mRNA.
Zbtb7a	Zbtb7a.bSep08	117107	8026	620	2	206	BTB/POZ (Zbtb7a) alternative variant bSep08, mRNA.
Zbtb8a	Zbtb8a.aSep08	313049	27973	1912	1	441	BTB/POZ and zinc finger, C2H2-type (49.9 kD) (Zbtb8a) alternative variant aSep08, mRNA.
Zbtb8os	Zbtb8os.aSep08	297885	11464	1623	7	167	putative protein of ancient origin (19.6 kD) (Zbtb8os) alternative variant aSep08, mRNA.
Zbtb8os	Zbtb8os.bSep08	297885	5332	567	4	74	putative protein of ancient origin (Zbtb8os) alternative variant bSep08, mRNA.
Zbtb10	Zbtb10.bSep08	80338	13458	611	2	203	putative protein of vertebrate origin (Zbtb10) alternative variant bSep08, mRNA.
Zbtb16	Zbtb16.bSep08	353227	87052	785	1	193	zinc finger (Zbtb16) alternative variant bSep08, mRNA.
Zbtb17	Zbtb17.bSep08	313666	18176	1040	7	288	BTB/POZ (Zbtb17) alternative variant bSep08, mRNA.
Zbtb17	Zbtb17.cSep08	313666	1504	833	6	277	zinc finger, C2H2-type (Zbtb17) alternative variant cSep08, mRNA.
Zbtb17	Zbtb17.dSep08	313666	1021	851	3	207	zinc finger, C2H2-type (Zbtb17) alternative variant dSep08, mRNA.
Zbtb20	Zbtb20.bSep08	288105	40855	1277	3	321	BTB/POZ (Zbtb20) alternative variant bSep08, mRNA.
Zbtb40	Zbtb40.aSep08	362635	5064	692		176	zinc finger, C2H2-type (Zbtb40) mRNA.
Zbtb43	Zbtb43.bSep08	311872	15096	768		197	zinc finger (Zbtb43) alternative variant bSep08, mRNA.
Zbtb45	Zbtb45.bSep08	308366	2702	1378	1	424	BTB/POZ (Zbtb45) alternative variant bSep08, mRNA.
Zc3h3	Zc3h3.bSep08	300032	20841	821	5	134	zinc finger CCCH type containing 3 (Zc3h3) alternative variant bSep08, mRNA.

Zc3h3	Zc3h3.cSep08	300032	17336	368	2	63	zinc finger CCCH type containing 3 (Zc3h3) alternative variant cSep08, mRNA.
Zc3h7a	Zc3h7a.bSep08	360466	6044	655	5	218	zinc finger CCCH type containing 7 A (Zc3h7a) alternative variant bSep08, mRNA.
Zc3h7a	Zc3h7a.cSep08	360466	7650	707	6	218	zinc finger CCCH type containing 7 A (Zc3h7a) alternative variant cSep08, mRNA.
Zc3h7a	Zc3h7a.dSep08	360466	2708	778	3	172	zinc finger CCCH type containing 7 A (Zc3h7a) alternative variant dSep08, mRNA.
Zc3h7b	Zc3h7b.bSep08	315158	28520	799	9	221	zinc finger CCCH-type containing 7B (Zc3h7b) alternative variant bSep08, mRNA.
Zc3h7b	Zc3h7b.cSep08	315158	653	394	3	89	zinc finger CCCH-type containing 7B (Zc3h7b) alternative variant cSep08, mRNA.
Zc3h10	Zc3h10.aSep08	685928	3243	1231		410	zinc finger CCCH type containing 10 (Zc3h10) mRNA.
Zc3h13	Zc3h13.aSep08	305955	36706	1131		275	zinc finger CCCH type containing 13 (Zc3h13) mRNA.
Zc3h14	Zc3h14.cSep08	192359	10746	973	5	184	zinc finger CCCH-type containing 14 (Zc3h14) alternative variant cSep08, mRNA.
Zc3h14	Zc3h14.dSep08	192359	3286	654	2	154	nuclear protein UKp68 (Zc3h14) alternative variant dSep08, mRNA.
Zc3h14	Zc3h14.eSep08	192359	3115	724	4	152	zinc finger CCCH-type containing 14 (Zc3h14) alternative variant eSep08, mRNA.
Zc3h14	Zc3h14.fSep08	192359	22495	706	6	117	zinc finger CCCH-type containing 14 (Zc3h14) alternative variant fSep08, mRNA.
Zc3h14	Zc3h14.gSep08	192359	10228	621	3	84	zinc finger CCCH-type containing 14 (9.2 kD) (Zc3h14) alternative variant gSep08, mRNA.
Zc3h14	Zc3h14.iSep08	192359	3109	237	3	20	putative protein (Zc3h14) alternative variant iSep08, mRNA.
Zc3h15	Zc3h15.bSep08	362154	13442	682	5	227	zinc finger CCCH-type containing 15 (Zc3h15) alternative variant bSep08, mRNA.
Zc3h15	Zc3h15.cSep08	362154	15319	759	6	172	zinc finger CCCH-type containing 15 (19.7 kD) (Zc3h15) alternative variant cSep08, mRNA.
Zc3h15	Zc3h15.dSep08	362154	8258	551	3	103	zinc finger CCCH-type containing 15 (Zc3h15) alternative variant dSep08, mRNA.
Zc3h15	Zc3h15.eSep08	362154	835	789	2	48	zinc finger CCCH-type containing 15 (Zc3h15) alternative variant eSep08, mRNA.
Zc3h15	Zc3h15.fSep08	362154	448	402	2	32	zinc finger CCCH-type containing 15 (3.6 kD) (Zc3h15) alternative variant fSep08, mRNA.
Zc3h18	Zc3h18.bSep08	292067	1630	1109	3	108	zinc finger CCCH-type containing 18 (Zc3h18) alternative variant bSep08, mRNA.
Zc3hav1	Zc3hav1.bSep08	252832	13776	2371	6	246	zinc finger CCCH type, antiviral 1 (Zc3hav1) alternative variant bSep08, mRNA.
Zc3hav1	Zc3hav1.cSep08	252832	780	409	2	87	zinc finger CCCH type, antiviral 1 (Zc3hav1) alternative variant cSep08, mRNA.
Zc3hav1l	Zc3hav1l.aSep08	362341	3662	547		182	zinc finger CCCH-type, antiviral 1-like (Zc3hav1l) mRNA.
Zc3hc1	Zc3hc1.bSep08	296957	13624	763	3	208	zinc finger, C3HC-type 1 (Zc3hc1) alternative variant bSep08, mRNA.
Zc3hc1	Zc3hc1.cSep08	296957	8986	1347	2	175	zinc finger, C3HC-type 1 (Zc3hc1) alternative variant cSep08, mRNA.



Zcchc2	Zcchc2.bSep08	304695	33307	2057	8	266	putative protein of metazoan origin (Zcchc2) alternative variant bSep08, mRNA.
Zcchc2	Zcchc2.cSep08	304695	4975	2334	2	142	zinc finger, CCHC-type (Zcchc2) alternative variant cSep08, mRNA.
Zcchc2	Zcchc2.dSep08	304695	15730	1067	5	111	zinc finger, CCHC-type (Zcchc2) alternative variant dSep08, mRNA.
Zcchc7	Zcchc7.bSep08	298086	6659	1966	3	250	zinc finger, CCHC-type (29.9 kD) (Zcchc7) alternative variant bSep08, mRNA.
Zcchc8	Zcchc8.bSep08	288661	2499	1583	2	251	putative nuclear protein of vertebrate origin (26.6 kD) (Zcchc8) alternative variant bSep08, mRNA.
Zcchc9	Zcchc9.bSep08	309986	2400	945	2	73	zinc finger, CCHC-type (8.1 kD) (Zcchc9) alternative variant bSep08, mRNA.
Zcchc9	Zcchc9.cSep08	309986	3359	1348	4	37	putative protein (4.6 kD) (Zcchc9) alternative variant cSep08, mRNA.
Zcchc10	Zcchc10.aSep08	360524	10471	1240		173	putative protein of eukaryotic origin (Zcchc10) mRNA.
Zcchc11	Zcchc11.bSep08	313481	15019	1257	6	355	zinc finger, CCHC-type (Zcchc11) alternative variant bSep08, mRNA.
Zcchc11	Zcchc11.cSep08	313481	14019	1506	5	290	zinc finger, CCHC-type (Zcchc11) alternative variant cSep08, mRNA.
Zcchc11	Zcchc11.eSep08	313481	9421	369	4	17	putative protein (2.1 kD) (Zcchc11) alternative variant eSep08, mRNA.
Zcchc12	Zcchc12.bSep08	313436	1583	490	5	46	putative protein (Zcchc12) alternative variant bSep08, mRNA.
Zcchc12	Zcchc12.cSep08	313436	1719	761	4	46	putative protein (Zcchc12) alternative variant cSep08, mRNA.
Zcchc12	Zcchc12.dSep08	313436	1449	439	4	44	putative protein (Zcchc12) alternative variant dSep08, mRNA.
Zcchc12	Zcchc12.eSep08	313436	1169	399	4	38	putative protein (Zcchc12) alternative variant eSep08, mRNA.
Zcchc12	Zcchc12.fSep08	313436	1613	531	5	34	putative protein (Zcchc12) alternative variant fSep08, mRNA.
Zcchc12	Zcchc12.gSep08	313436	2218	1108	4	33	putative protein (Zcchc12) alternative variant gSep08, mRNA.
Zcchc12	Zcchc12.hSep08	313436	1056	438	3	51	putative protein (Zcchc12) alternative variant hSep08, mRNA.
Zcchc14	Zcchc14.aSep08	365018	1791	736		115	zinc finger, CCHC-type (Zcchc14) mRNA.
Zcchc17	Zcchc17.bSep08	500555	41226	2070	7	193	putative nuclear protein of ancient origin (22.1 kD) (Zcchc17) alternative variant bSep08, complete mRNA.
Zcchc17	Zcchc17.cSep08	500555	9509	545	2	94	putative nuclear protein of mammalian origin (10.9 kD) (Zcchc17) alternative variant cSep08, mRNA.
Zcrb1	Zcrb1.bSep08	362990	6475	746	2	156	zinc finger CCHC-type and RNA binding motif 1 (17.8 kD) (Zcrb1) alternative variant bSep08, mRNA.
Zcwpw1	Zcwpw1.aSep08	304368	20199	1695		515	zinc finger, CW-type with PWWP domain 1 (Zcwpw1) mRNA.
Zdbf2	Zdbf2.aSep08	501153	20682	1426		449	zinc finger, DBF-type containing 2 (Zdbf2) mRNA.
Zdhhc1	Zdhhc1.bSep08	291967	15644	2213	10	234	putative nuclear protein of mammalian origin (25.5 kD) (Zdhhc1) alternative variant bSep08, mRNA.

Zdhhc1	Zdhhc1.cSep08	291967	4932	1838	6	129	putative protein of vertebrate origin (14.9 kD) (Zdhhc1) alternative variant cSep08, mRNA.
Zdhhc1	Zdhhc1.dSep08	291967	3371	744	3	69	putative protein (Zdhhc1) alternative variant dSep08, mRNA.
Zdhhc1	Zdhhc1.eSep08	291967	19032	470	5	109	putative protein of eukaryotic origin (Zdhhc1) alternative variant eSep08, mRNA.
Zdhhc1	Zdhhc1.fSep08	291967	14140	412	5	41	putative protein (4.7 kD) (Zdhhc1) alternative variant fSep08, mRNA.
Zdhhc2	Zdhhc2.bSep08	246326	35607	1196	4	82	putative protein of metazoan origin (9.6 kD) (Zdhhc2) alternative variant bSep08, mRNA.
Zdhhc2	Zdhhc2.cSep08	246326	20902	2000	3	55	putative protein (6.4 kD) (Zdhhc2) alternative variant cSep08, mRNA.
Zdhhc4	Zdhhc4.bSep08	304291	8341	753	6	148	putative protein (15.9 kD) (Zdhhc4) alternative variant bSep08, mRNA.
Zdhhc4	Zdhhc4.cSep08	304291	6042	721	6	120	putative protein, with a transmembrane domain, of vertebrate origin (Zdhhc4) alternative variant cSep08, mRNA.
Zdhhc5	Zdhhc5.bSep08	362156	22361	392	3	130	zinc finger, DHHC-type (Zdhhc5) alternative variant bSep08, mRNA.
Zdhhc5	Zdhhc5.cSep08	362156	4931	426	2	41	putative protein (Zdhhc5) alternative variant cSep08, mRNA.
Zdhhc5	Zdhhc5.dSep08	362156	4775	400	3	36	putative protein (Zdhhc5) alternative variant dSep08, mRNA.
Zdhhc6	Zdhhc6.cSep08	361771	4002	794	3	42	putative protein (5.0 kD) (Zdhhc6) alternative variant cSep08, mRNA.
Zdhhc8	Zdhhc8.bSep08	303796	544	379	2	81	zinc finger (Zdhhc8) alternative variant bSep08, mRNA.
Zdhhc11	Zdhhc11.bSep08	499000	2140	772	1	98	putative protein (10.9 kD) (Zdhhc11) alternative variant bSep08, mRNA.
Zdhhc12	Zdhhc12.bSep08	366014	2179	458		152	zinc finger, DHHC-type (Zdhhc12) alternative variant bSep08, mRNA.
Zdhhc13	Zdhhc13.bSep08	365252	33957	2137	14	712	zinc finger DHHC-type containing 13 CRA a (Zdhhc13) alternative variant bSep08, mRNA.
Zdhhc13	Zdhhc13.cSep08	365252	39137	2344	16	638	zinc finger DHHC-type containing 13 CRA b (Zdhhc13) alternative variant cSep08, mRNA.
Zdhhc13	Zdhhc13.dSep08	365252	6864	388	4	95	zinc finger DHHC-type containing 13 CRA a (Zdhhc13) alternative variant dSep08, mRNA.
Zdhhc13	Zdhhc13.eSep08	365252	4342	551	1	83	zinc finger DHHC-type containing 13 CRA b (Zdhhc13) alternative variant eSep08, mRNA.
Zdhhc17	Zdhhc17.bSep08	366889	5172	3594	3	103	zinc finger DHHC-type containing 17 CRA d (Zdhhc17) alternative variant bSep08, mRNA.
Zdhhc17	Zdhhc17.cSep08	366889	2571	268	2	31	putative protein (Zdhhc17) alternative variant cSep08, mRNA.
Zdhhc18	Zdhhc18.aSep08	362613	19913	547	3	176	zinc finger DHHC-type containing 18 CRA b (Zdhhc18) alternative variant aSep08, mRNA.
Zdhhc18	Zdhhc18.bSep08	362613	6090	425	1	103	zinc finger DHHC-type containing 18 precursor (Zdhhc18) alternative variant bSep08, mRNA.

Zdhhc19	Zdhhc19.bSep08	288045	1336	576	1	106	CRA b like (11.2 kD) (Zdhhc19) alternative variant bSep08, mRNA.
Zdhhc20	Zdhhc20.aSep08	305923	56408	2551	7	387	zinc finger DHHC-type containing 20 CRA d (Zdhhc20) alternative variant aSep08, complete mRNA.
Zdhhc20	Zdhhc20.bSep08	305923	28238	1177	4	296	zinc finger DHHC-type containing 20 CRA d (Zdhhc20) alternative variant bSep08, mRNA.
Zdhhc20	Zdhhc20.cSep08	305923	55459	1579	7	265	zinc finger DHHC-type containing 20 CRA d (31.0 kD) (Zdhhc20) alternative variant cSep08, mRNA.
Zdhhc20	Zdhhc20.dSep08	305923	20041	582		193	zinc finger DHHC-type containing 20 CRA d (Zdhhc20) alternative variant dSep08, mRNA.
Zeb1	Zeb1.aSep08	25705	59169	5124	7	1092	zinc finger E-box binding homeobox 1 (Zeb1) alternative variant aSep08, mRNA.
zeebor	zeebor.aSep08		2505	396		34	putative protein (3.5 kD) (zeebor) mRNA.
zeeby	zeeby.aSep08		1765	1089	2	126	binding protein 4 CRA a like (zeeby) alternative variant aSep08, mRNA.
zeechy	zeechy.aSep08		1282	314		104	transient receptor potential cation channel subfamily M member 7 (zeechy) mRNA.
zeedar	zeedar.aSep08		3436	767	1	21	putative protein (2.5 kD) (zeedar) alternative variant aSep08, mRNA.
zeedar	zeedar.bSep08		3556	762		21	putative protein (2.5 kD) (zeedar) alternative variant bSep08, mRNA.
zeedoy	zeedoy.aSep08		11629	1079	1	186	putative protein of metazoan origin (zeedoy) alternative variant aSep08, mRNA.
zeedoy	zeedoy.bSep08		8823	473	1	157	putative protein of metazoan origin (zeedoy) alternative variant bSep08, mRNA.
zeeflu	zeeflu.aSep08		6816	523		173	CRA b (zeeflu) mRNA.
zeefly	zeefly.aSep08		1663	838		65	putative protein (7.2 kD) (zeefly) mRNA.
zeegar	zeegar.aSep08		5420	354		64	polyprotein (zeegar) mRNA.
zeeja	zeeja.bSep08		2307	444	2	63	putative protein (zeeja) alternative variant bSep08, mRNA.
zeejey	zeejey.aSep08		12899	315		56	putative protein (zeejey) mRNA.
zeekee	zeekee.aSep08		3400	403		79	putative secreted or extracellular protein precursor (8.8 kD) (zeekee) mRNA.
zeekler	zeekler.aSep08		7928	923		43	putative protein (5.0 kD) (zeekler) mRNA.
zeelo	zeelo.aSep08		2100	588		31	putative protein (3.4 kD) (zeelo) mRNA.
zeemee	zeemee.aSep08		9913	1343		447	nuclear receptor co-repressor (zeemee) mRNA.
zeenoy	zeenoy.aSep08		5423	699		83	putative cytoplasmic protein (9.6 kD) (zeenoy) mRNA.
zeepor	zeepor.aSep08		12118	268		36	putative protein (4.1 kD) (zeepor) mRNA.
zeesa	zeesa.bSep08		2422	373	4	37	putative protein (zeesa) alternative variant bSep08, mRNA.
zeeshee	zeeshee.aSep08		3485	342		113	cingulin (zeeshee) mRNA.
zeetu	zeetu.aSep08		62255	571		190	oligomeric golgi complex (zeetu) mRNA.
zeevar	zeevar.aSep08		1961	783		40	chromodomain helicase DNA binding protein 5 like (zeevar) mRNA.
zeevo	zeevo.aSep08		4371	356		118	nuclear receptor coactivator 2 (zeevo) mRNA.
zeewey	zeewey.aSep08		5415	729		90	putative protein (zeewey) mRNA.

Zer1	Zer1.aSep08	311842	9310	1257	9	359	zyg-11 homolog B -like (Zer1) alternative variant aSep08, mRNA.
Zer1	Zer1.bSep08	311842	3087	1688	2	74	zyg-11 homolog B -like (Zer1) alternative variant bSep08, mRNA.
zerbor	zerbor.aSep08		5607	4276		153	binding protein 1 like (zerbor) mRNA.
zerby	zerby.aSep08		3029	1106		111	putative protein (11.8 kD) (zerby) mRNA.
zerchy	zerchy.aSep08		1296	336		112	transient receptor potential cation channel subfamily M member 7 (zerchy) mRNA.
zerdar	zerdar.aSep08		8751	497		45	putative protein (5.0 kD) (zerdar) mRNA.
zerdoy	zerdoy.aSep08		11982	261		87	putative protein (zerdoy) mRNA.
zerflu	zerflu.aSep08		2832	269		89	CRA a like (zerflu) mRNA.
zerfly	zerfly.aSep08		18393	742		49	putative protein (zerfly) mRNA.
zergar	zergar.aSep08		3942	545		126	CRA a (zergar) mRNA.
zerja	zerja.aSep08		4205	861	3	141	PHD finger protein 11 (zerja) alternative variant aSep08, mRNA.
zerjey	zerjey.aSep08		656	312		28	putative protein (zerjey) mRNA.
zerkee	zerkee.aSep08		12527	711		86	centrosomal protein (zerkee) mRNA.
zerkler	zerkler.aSep08		6491	397		131	odd Oz ten-m homolog 3 (zerkler) mRNA.
zerlo	zerlo.aSep08		6772	361		70	putative protein of mammalian origin (zerlo) mRNA.
zermee	zermee.aSep08		18773	817	4	271	nuclear receptor co-repressor 1 (zermee) alternative variant aSep08, mRNA.
zermee	zermee.bSep08		9098	411	1	136	putative protein of vertebrate origin (zermee) alternative variant bSep08, mRNA.
zernoy	zernoy.aSep08		24302	1165		222	histone deacetylase 4 (zernoy) mRNA.
zerpor	zerpor.aSep08		8353	974		166	ataxia telangiectasia Rad3 related (zerpor) mRNA.
zersa	zersa.aSep08		4819	678		33	putative protein (zersa) mRNA.
zershee	zershee.aSep08		17774	664	2	58	putative protein (zershee) alternative variant aSep08, mRNA.
zershee	zershee.bSep08		856	753	1	44	putative protein (zershee) alternative variant bSep08, mRNA.
zertu	zertu.aSep08		79596	1227	3	319	oligomeric golgi complex (35.4 kD) (zertu) alternative variant aSep08, mRNA.
zertu	zertu.bSep08		27491	514	1	132	oligomeric golgi complex (zertu) alternative variant bSep08, mRNA.
zervar	zervar.aSep08		5185	717		64	putative protein (zervar) mRNA.
zervo	zervo.aSep08		773	417		138	nuclear receptor coactivator 2 (zervo) mRNA.
zerwey	zerwey.aSep08		8902	619	2	43	putative protein (5.0 kD) (zerwey) alternative variant aSep08, mRNA.
zerwey	zerwey.bSep08		1944	430	1	38	putative protein (4.4 kD) (zerwey) alternative variant bSep08, mRNA.
zeybor	zeybor.aSep08		1629	304	2	76	putative protein (zeybor) mRNA.
zeychy	zeychy.aSep08		1677	279		93	transient receptor potential cation channel subfamily m member 7 CRA b (zeychy) mRNA.
zeydar	zeydar.aSep08		3137	366	2	37	putative protein (zeydar) alternative variant aSep08, mRNA.

zeydar	zeydar.bSep08		3100	307	2	34	putative protein (zeydar) alternative variant bSep08, mRNA.
zeydoy	zeydoy.aSep08		2568	1115		52	putative protein (5.7 kD) (zeydoy) mRNA.
zeyflu	zeyflu.aSep08		2754	333		110	CRA a (zeyflu) mRNA.
zeyfly	zeyfly.aSep08		3948	907		72	putative mitochondrial protein (7.7 kD) (zeyfly) mRNA.
zeygar	zeygar.aSep08		13140	1287	2	428	CRA c (zeygar) alternative variant aSep08, mRNA.
zeygar	zeygar.bSep08		3752	764	1	173	CRA a (zeygar) alternative variant bSep08, mRNA.
zeyja	zeyja.aSep08		9073	961	5	220	SET domain bifurcated protein (zeyja) alternative variant aSep08, mRNA.
zeyja	zeyja.bSep08		13220	370	4	123	SET domain bifurcated protein (zeyja) alternative variant bSep08, mRNA.
zeyjey	zeyjey.aSep08		686	360		111	putative protein (zeyjey) mRNA.
zeykee	zeykee.aSep08		2195	692		230	centrosomal protein 170kDa (zeykee) mRNA.
zeykler	zeykler.aSep08		2837	400		71	putative mitochondrial protein (7.8 kD) (zeykler) mRNA.
zeylo	zeylo.aSep08		8659	820	3	54	putative protein (6.3 kD) (zeylo) alternative variant aSep08, mRNA.
zeylo	zeylo.bSep08		8747	185	2	22	putative protein (zeylo) alternative variant bSep08, mRNA.
zeymee	zeymee.aSep08		5290	326		108	nuclear receptor co-repressor 1 (zeymee) mRNA.
zeynoy	zeynoy.aSep08		16715	519		72	histone deacetylase 4 (zeynoy) mRNA.
zeypor	zeypor.aSep08		1773	709		129	5'-3' exoribonuclease 1 (zeypor) mRNA.
zeyroy	zeyroy.aSep08		2953	376		35	putative protein (3.8 kD) (zeyroy) mRNA.
zeysa	zeysa.aSep08		2940	409		135	AT rich interactive domain 2 (zeysa) mRNA.
zeyshee	zeyshee.aSep08		14182	658	3	35	putative protein (4.1 kD) (zeyshee) alternative variant aSep08, mRNA.
zeytu	zeytu.aSep08		3106	375		107	catalytic gamma polypeptide (zeytu) mRNA.
zeyvar	zeyvar.aSep08		2732	731		92	putative protein (10.1 kD) (zeyvar) mRNA.
zeyvo	zeyvo.aSep08		2884	667		44	putative protein (5.2 kD) (zeyvo) mRNA.
zeywey	zeywey.aSep08		22333	554		82	putative protein of mammalian origin (zeywey) mRNA.
zf-AD.0	zf-AD.0.aSep08		4495	1000		333	zinc finger protein 276 (zf-AD.0) mRNA.
zf-B_box.0	zf-B_box.0.aSep08		62824	781		260	tripartite motif-containing (zf-B_box.0) mRNA.
zf-C2H2.0	zf-C2H2.0.aSep08		1314	1183		394	transcription factor yy2 (zf-C2H2.0) mRNA.
zf-C2H2.1	zf-C2H2.1.aSep08		1466	394		131	zinc finger, C2H2-type (zf-C2H2.1) mRNA.
zf-C2H2.3	zf-C2H2.3.aSep08		2887	900		224	zinc finger protein 187 (zf-C2H2.3) mRNA.
zf-C2H2.16	zf-C2H2.16.aSep08		3355	2745		486	CRA a (56.9 kD) (zf-C2H2.16) mRNA.
zf-C2H2.17	zf-C2H2.17.aSep08		11949	907		36	CRA b like (4.1 kD) (zf-C2H2.17) mRNA.
zf-C2H2.38	zf-C2H2.38.aSep08		19669	910	3	201	zinc finger, C2H2-type (zf-C2H2.38) alternative variant aSep08, mRNA.
zf-C2H2.38	zf-C2H2.38.bSep08		18104	453	1	39	putative protein (zf-C2H2.38) alternative variant bSep08, mRNA.
zf-C2H2.45	zf-C2H2.45.aSep08		11829	515	4	123	CRA c like (zf-C2H2.45) alternative variant aSep08, mRNA.
zf-C2H2.52	zf-C2H2.52.aSep08		3997	591	2	197	zinc finger protein 406 CRA a (zf-C2H2.52) alternative variant aSep08, mRNA.
zf-C2H2.52	zf-C2H2.52.bSep08		30067	444	3	106	zinc finger (zf-C2H2.52) alternative variant bSep08, mRNA.

zf-C2H2.52	zf-C2H2.52.cSep08		22217	374	2	103	zinc finger (zf-C2H2.52) alternative variant cSep08, mRNA.
zf-C2H2.53	zf-C2H2.53.aSep08		6552	3715	2	152	zinc finger protein 740 (17.9 kD) (zf-C2H2.53) alternative variant aSep08, mRNA.
zf-C2H2.53	zf-C2H2.53.bSep08		3202	801	1	92	putative protein (zf-C2H2.53) alternative variant bSep08, mRNA.
zf-C2H2.64	zf-C2H2.64.aSep08		2259	558		137	zinc finger, C2H2-type (zf-C2H2.64) mRNA.
zf-C2H2.65	zf-C2H2.65.aSep08		1591	781		260	zinc finger protein 462 CRA c (zf-C2H2.65) mRNA.
zf-C2H2.66	zf-C2H2.66.aSep08		12237	372		123	zinc finger (zf-C2H2.66) mRNA.
zf-C2H2.67	zf-C2H2.67.aSep08		1377	487		118	zinc finger, C2H2-type (zf-C2H2.67) mRNA.
zf-C2H2.74	zf-C2H2.74.aSep08		5530	714	1	115	putative protein (12.1 kD) (zf-C2H2.74) alternative variant aSep08, mRNA.
zf-C2H2.74	zf-C2H2.74.bSep08		6092	1533	1	115	putative protein (12.1 kD) (zf-C2H2.74) alternative variant bSep08, mRNA.
zf-C2H2.77	zf-C2H2.77.aSep08		3376	1299		208	zinc finger protein 341 (zf-C2H2.77) mRNA.
zf-C2H2.78	zf-C2H2.78.aSep08		3241	881	6	293	zinc finger protein 335 (zf-C2H2.78) alternative variant aSep08, mRNA.
zf-C2H2.79	zf-C2H2.79.aSep08		5198	732		244	zinc finger protein 335 (zf-C2H2.79) mRNA.
zf-C2H2.86	zf-C2H2.86.aSep08		4167	3173		497	zinc finger protein 687 (zf-C2H2.86) mRNA.
zf-C2H2.90	zf-C2H2.90.aSep08		5055	3010		574	pleiomorphic adenoma gene-like 1 (zf-C2H2.90) mRNA.
zf-C2H2.91	zf-C2H2.91.aSep08		3151	2255	2	448	KRAB box and zinc finger, C2H2-type (zf-C2H2.91) alternative variant aSep08, mRNA.
zf-C2H2.91	zf-C2H2.91.bSep08		15277	3356	5	211	zinc finger, C2H2-type (24.5 kD) (zf-C2H2.91) alternative variant bSep08, complete mRNA.
zf-C2H2.91	zf-C2H2.91.eSep08		4435	552	4	24	putative protein (2.8 kD) (zf-C2H2.91) alternative variant eSep08, mRNA.
zf-C2H2.92	zf-C2H2.92.aSep08		3926	200	2	66	putative protein (zf-C2H2.92) alternative variant aSep08, mRNA.
zf-C2H2.92	zf-C2H2.92.bSep08		3642	1175	2	72	zinc finger, C2H2-type (zf-C2H2.92) alternative variant bSep08, mRNA.
zf-C2H2.92	zf-C2H2.92.cSep08		15949	673	6	65	putative protein (zf-C2H2.92) alternative variant cSep08, mRNA.
zf-C2H2.93	zf-C2H2.93.aSep08		9230	762		219	zinc finger, C2H2-type (zf-C2H2.93) mRNA.
zf-C2H2.94	zf-C2H2.94.aSep08		9335	575		191	zinc finger protein 780B (zf-C2H2.94) mRNA.
zf-C2H2.95	zf-C2H2.95.aSep08		10828	514		66	CRA a like (zf-C2H2.95) mRNA.
zf-C2H2.96	zf-C2H2.96.aSep08		2834	280	2	68	CRA a like (zf-C2H2.96) alternative variant aSep08, mRNA.
zf-C2H2.97	zf-C2H2.97.aSep08		7640	353	3	55	putative protein (6.1 kD) (zf-C2H2.97) alternative variant aSep08, mRNA.
zf-C2H2.97	zf-C2H2.97.bSep08		11204	825	3	52	putative protein (5.8 kD) (zf-C2H2.97) alternative variant bSep08, mRNA.
zf-C2H2.97	zf-C2H2.97.dSep08		9672	199	2	29	putative protein (zf-C2H2.97) alternative variant dSep08, mRNA.
zf-C2H2.98	zf-C2H2.98.aSep08		11438	888		222	zinc finger, C2H2-type (zf-C2H2.98) mRNA.
zf-C2HC5.0	zf-C2HC5.0.aSep08		90177	3798	15	685	thyroid hormone receptor interactor 4 (77.6 kD) (zf-C2HC5.0) alternative variant aSep08, mRNA.

zf-C2HC5.0	zf-C2HC5.0.bSep08		34478	676	5	225	thyroid hormone receptor interactor 4 (zf-C2HC5.0) alternative variant bSep08, mRNA.
zf-C2HC5.0	zf-C2HC5.0.cSep08		8246	746	4	173	thyroid hormone receptor interactor 4 (zf-C2HC5.0) alternative variant cSep08, mRNA.
zf-C2HC5.0	zf-C2HC5.0.dSep08		6698	689	2	35	thyroid hormone receptor interactor 4 (4.0 kD) (zf-C2HC5.0) alternative variant dSep08, mRNA.
zf-C3HC4.2	zf-C3HC4.2.aSep08		6868	716	2	238	putative protein of mammalian origin (zf-C3HC4.2) alternative variant aSep08, mRNA.
zf-C3HC4.2	zf-C3HC4.2.bSep08		10682	821	2	234	ring finger WD repeat domain 3 (zf-C3HC4.2) alternative variant bSep08, mRNA.
zf-C3HC4.3	zf-C3HC4.3.aSep08		2207	865		175	LON peptidase N-terminal domain ring finger 1 (zf-C3HC4.3) mRNA.
zf-C3HC4.4	zf-C3HC4.4.aSep08		38009	1529	1	326	ring finger WD repeat domain 2 CRA a (zf-C3HC4.4) alternative variant aSep08, mRNA.
zf-C3HC4.4	zf-C3HC4.4.bSep08		37218	726	1	241	ring finger WD repeat domain 2 CRA a (zf-C3HC4.4) alternative variant bSep08, mRNA.
zf-C3HC4.5	zf-C3HC4.5.aSep08		3697	730		242	CRA a (zf-C3HC4.5) mRNA.
zf-C3HC4.6	zf-C3HC4.6.aSep08		1523	365		121	cas-Br-M ecotropic retroviral transforming sequence (zf-C3HC4.6) mRNA.
zf-C4.0	zf-C4.0.aSep08		2220	361		120	steroidogenic factor 1 (zf-C4.0) mRNA.
zf-CCCH.0	zf-CCCH.0.aSep08		11232	636		187	small nuclear auxiliary factor 2 RNA 1-like (zf-CCCH.0) mRNA.
zf-CCCH.1	zf-CCCH.1.aSep08		18704	616	2	204	zinc finger, CCCH-type (zf-CCCH.1) alternative variant aSep08, mRNA.
zf-CCHC.6	zf-CCHC.6.aSep08		5366	947		315	retinoblastoma binding protein 6 like (zf-CCHC.6) mRNA.
zf-CHCC.0	zf-CHCC.0.aSep08		493	235	2	31	nadh dehydrogenase protein 6 (zf-CHCC.0) alternative variant aSep08, mRNA.
zf-CXXC.0	zf-CXXC.0.aSep08		643	389		129	myeloid lymphoid leukemia like (zf-CXXC.0) mRNA.
zf-DHHC.0	zf-DHHC.0.bSep08	308081	31408	1351	1	195	putative protein of metazoan origin (zf-DHHC.0) alternative variant bSep08, mRNA.
zf-HIT.0	zf-HIT.0.aSep08		6275	893	4	153	zinc finger HIT 1 (17.4 kD) (zf-HIT.0) alternative variant aSep08, mRNA.
zf-HIT.0	zf-HIT.0.cSep08		3110	669	2	102	putative secreted or extracellular protein precursor (10.7 kD) (zf-HIT.0) alternative variant cSep08, mRNA.
zf-HIT.0	zf-HIT.0.dSep08		762	478	2	91	zinc finger HIT 1 (zf-HIT.0) alternative variant dSep08, mRNA.
zf-NF-X1.0	zf-NF-X1.0.aSep08		712	403		134	nuclear transcription factor X-box binding-like 1 CRA d (zf-NF-X1.0) mRNA.
zf-TAZ.0	zf-TAZ.0.aSep08		1208	289		96	E1A binding protein p300 like (zf-TAZ.0) mRNA.
zf-UBP.0	zf-UBP.0.aSep08		18068	1531	7	453	ubiquitin specific 33 (zf-UBP.0) alternative variant aSep08, mRNA.
zf-UBP.0	zf-UBP.0.bSep08		4156	739	1	153	ubiquitin specific 33 (zf-UBP.0) alternative variant bSep08, mRNA.
Zfand1	Zfand1.aSep08	361917	4080	367		121	zinc finger, AN1-type domain 1 (Zfand1) mRNA.
Zfand2a	Zfand2a.bSep08	360772	2720	585	3	120	zinc finger, AN1-type domain 2A (Zfand2a) alternative variant bSep08, mRNA.

Zfand2b	Zfand2b.bSep08	363253	1141	409	5	135	zinc finger, AN1 type domain 2B (Zfand2b) alternative variant bSep08, mRNA.
Zfand2b	Zfand2b.cSep08	363253	1115	386	5	128	zinc finger, AN1 type domain 2B (Zfand2b) alternative variant cSep08, mRNA.
Zfand2b	Zfand2b.dSep08	363253	1151	592	4	89	zinc finger, AN1 type domain 2B (Zfand2b) alternative variant dSep08, mRNA.
Zfand2b	Zfand2b.eSep08	363253	609	397	2	76	zinc finger, AN1 type domain 2B (Zfand2b) alternative variant eSep08, mRNA.
Zfand2b	Zfand2b.fSep08	363253	1082	423	4	69	zinc finger, AN1 type domain 2B (Zfand2b) alternative variant fSep08, mRNA.
Zfand3	Zfand3.bSep08	361816	16811	1286	1	114	zinc finger, AN1-type domain 3 (13.2 kD) (Zfand3) alternative variant bSep08, mRNA.
Zfand5	Zfand5.aSep08	293960	9767	2684	6	298	zinc finger, AN1-type domain 5 and hypothetical protein LOC679612 (32.3 kD) (Zfand5) alternative variant aSep08, mRNA.
Zfand5	Zfand5.aSep08	679612	9767	2684	6	298	zinc finger, AN1-type domain 5 and hypothetical protein LOC679612 (32.3 kD) (Zfand5) alternative variant aSep08, mRNA.
Zfand5	Zfand5.bSep08	293960	3774	744	4	131	zinc finger, AN1-type domain 5 and hypothetical protein LOC679612 (Zfand5) alternative variant bSep08, mRNA.
Zfand5	Zfand5.bSep08	679612	3774	744	4	131	zinc finger, AN1-type domain 5 and hypothetical protein LOC679612 (Zfand5) alternative variant bSep08, mRNA.
Zfand6andFah	Zfand6andFah.cSep08	29383	13890	811	8	234	fumarylacetoacetate hydrolase (Zfand6andFah) alternative variant cSep08, mRNA.
Zfand6andFah	Zfand6andFah.cSep08	293067	13890	811	8	234	fumarylacetoacetate hydrolase (Zfand6andFah) alternative variant cSep08, mRNA.
Zfand6andFah	Zfand6andFah.dSep08	29383	63191	1095	7	223	zinc finger domain (24.0 kD) (Zfand6andFah) alternative variant dSep08, mRNA.
Zfand6andFah	Zfand6andFah.dSep08	293067	63191	1095	7	223	zinc finger domain (24.0 kD) (Zfand6andFah) alternative variant dSep08, mRNA.
Zfand6andFah	Zfand6andFah.eSep08	29383	74117	839	6	208	zinc finger domain (22.5 kD) (Zfand6andFah) alternative variant eSep08, mRNA.
Zfand6andFah	Zfand6andFah.eSep08	293067	74117	839	6	208	zinc finger domain (22.5 kD) (Zfand6andFah) alternative variant eSep08, mRNA.
Zfand6andFah	Zfand6andFah.fSep08	29383	60781	567	6	188	zinc finger domain (Zfand6andFah) alternative variant fSep08, mRNA.
Zfand6andFah	Zfand6andFah.fSep08	293067	60781	567	6	188	zinc finger domain (Zfand6andFah) alternative variant fSep08, mRNA.
Zfand6andFah	Zfand6andFah.gSep08	29383	72095	744	6	174	zinc finger domain (Zfand6andFah) alternative variant gSep08, mRNA.
Zfand6andFah	Zfand6andFah.gSep08	293067	72095	744	6	174	zinc finger domain (Zfand6andFah) alternative variant gSep08, mRNA.
Zfand6andFah	Zfand6andFah.hSep08	29383	72004	807	7	172	zinc finger domain 6 CRA b (Zfand6andFah) alternative variant hSep08, mRNA.
Zfand6andFah	Zfand6andFah.hSep08	293067	72004	807	7	172	zinc finger domain 6 CRA b (Zfand6andFah) alternative variant hSep08, mRNA.



Zfand6andFah	Zfand6andFah.iSep08	29383	60799	746	7	171	zinc finger domain (Zfand6andFah) alternative variant iSep08, mRNA.
Zfand6andFah	Zfand6andFah.iSep08	293067	60799	746	7	171	zinc finger domain (Zfand6andFah) alternative variant iSep08, mRNA.
Zfand6andFah	Zfand6andFah.jSep08	29383	71996	761	6	145	zinc finger domain (Zfand6andFah) alternative variant jSep08, mRNA.
Zfand6andFah	Zfand6andFah.jSep08	293067	71996	761	6	145	zinc finger domain (Zfand6andFah) alternative variant jSep08, mRNA.
Zfand6andFah	Zfand6andFah.kSep08	29383	60754	643	6	145	zinc finger domain (Zfand6andFah) alternative variant kSep08, mRNA.
Zfand6andFah	Zfand6andFah.kSep08	293067	60754	643	6	145	zinc finger domain (Zfand6andFah) alternative variant kSep08, mRNA.
Zfand6andFah	Zfand6andFah.lSep08	29383	52099	388	4	128	zinc finger domain (Zfand6andFah) alternative variant lSep08, mRNA.
Zfand6andFah	Zfand6andFah.lSep08	293067	52099	388	4	128	zinc finger domain (Zfand6andFah) alternative variant lSep08, mRNA.
Zfand6andFah	Zfand6andFah.mSep08	29383	26716	780	3	58	zinc finger domain 6 (Zfand6andFah) alternative variant mSep08, mRNA.
Zfand6andFah	Zfand6andFah.mSep08	293067	26716	780	3	58	zinc finger domain 6 (Zfand6andFah) alternative variant mSep08, mRNA.
Zfand6andFah	Zfand6andFah.nSep08	29383	3992	2110	2	58	zinc finger domain (6.6 kD) (Zfand6andFah) alternative variant nSep08, mRNA.
Zfand6andFah	Zfand6andFah.nSep08	293067	3992	2110	2	58	zinc finger domain (6.6 kD) (Zfand6andFah) alternative variant nSep08, mRNA.
Zfand6andFah	Zfand6andFah.oSep08	29383	63285	749	6	98	zinc finger domain (Zfand6andFah) alternative variant oSep08, mRNA.
Zfand6andFah	Zfand6andFah.oSep08	293067	63285	749	6	98	zinc finger domain (Zfand6andFah) alternative variant oSep08, mRNA.
Zfand6andFah	Zfand6andFah.qSep08	29383	4869	625	5	59	putative protein (Zfand6andFah) alternative variant qSep08, mRNA.
Zfand6andFah	Zfand6andFah.qSep08	293067	4869	625	5	59	putative protein (Zfand6andFah) alternative variant qSep08, mRNA.
Zfat	Zfat.aSep08	362925	90637	1786		390	zinc finger protein 406 CRA b (Zfat) alternative variant aSep08, mRNA.
Zfat	Zfat.bSep08	362925	25287	1154		179	zinc finger (Zfat) alternative variant bSep08, mRNA.
Zfhx2	Zfhx2.bSep08	305888	2359	395	1	131	zinc finger homeobox 2 (Zfhx2) alternative variant bSep08, mRNA.
Zfhx3	Zfhx3.aSep08	307829	4882	827		275	zinc finger homeobox 3 (Zfhx3) mRNA.
Zfhx4	Zfhx4.aSep08	310250	8084	612		203	zinc finger homeodomain 4 (Zfhx4) mRNA.
Zfml	Zfml.bSep08	312491	13078	1378	2	459	zinc finger, matrin-like (Zfml) alternative variant bSep08, mRNA.
Zfml	Zfml.cSep08	312491	9165	1834	5	420	zinc finger, matrin-like (47.2 kD) (Zfml) alternative variant cSep08, mRNA.
Zfml	Zfml.dSep08	312491	5395	605	4	119	zinc finger, matrin-like (Zfml) alternative variant dSep08, mRNA.
Zfml	Zfml.eSep08	312491	2983	544	3	99	zinc finger, matrin-like (Zfml) alternative variant eSep08, mRNA.

Zfp12	Zfp12.aSep08	288486	17400	567		125	zinc finger protein 12 (13.6 kD) (Zfp12) mRNA.
Zfp28	Zfp28.aSep08	502294	6245	413		120	zinc finger protein 28 (Zfp28) mRNA.
Zfp36l1	Zfp36l1.bSep08	29344	3328	2837	2	309	zinc finger protein 36, C3H type-like 1 (Zfp36l1) alternative variant bSep08, mRNA.
Zfp37	Zfp37.bSep08	115768	32249	3162	3	559	zinc finger protein 37 (63.4 kD) (Zfp37) alternative variant bSep08, mRNA.
Zfp37	Zfp37.cSep08	115768	32900	3342	4	134	zinc finger protein 37 (Zfp37) alternative variant cSep08, mRNA.
Zfp37	Zfp37.dSep08	115768	4019	472	2	70	zinc finger protein 37 (Zfp37) alternative variant dSep08, mRNA.
Zfp40	Zfp40.aSep08	690661	22454	3192	4	730	zinc finger protein 40 (86.0 kD) (Zfp40) alternative variant aSep08, mRNA.
Zfp40	Zfp40.bSep08	690661	4273	411	3	137	zinc finger protein 40 (Zfp40) alternative variant bSep08, mRNA.
Zfp40	Zfp40.dSep08	690661	33513	264	4	87	zinc finger protein 40 (Zfp40) alternative variant dSep08, mRNA.
Zfp40	Zfp40.eSep08	690661	76785	1783	4	80	zinc finger protein 40 (Zfp40) alternative variant eSep08, mRNA.
Zfp40	Zfp40.fSep08	690661	3041	621	4	42	zinc finger protein 40 (4.7 kD) (Zfp40) alternative variant fSep08, mRNA.
Zfp40	Zfp40.gSep08	690661	15653	426	4	42	zinc finger protein 40 (4.7 kD) (Zfp40) alternative variant gSep08, mRNA.
Zfp41	Zfp41.aSep08	315081	1626	386		42	zinc finger protein 41 (Zfp41) mRNA.
Zfp53	Zfp53.bSep08	308236	26677	1593	4	485	zinc finger protein 53 (Zfp53) alternative variant bSep08, mRNA.
Zfp53	Zfp53.cSep08	308236	18385	417	3	55	zinc finger protein 53 (6.4 kD) (Zfp53) alternative variant cSep08, mRNA.
Zfp59	Zfp59.aSep08	685610	12565	589		195	zinc finger protein 59 (Zfp59) mRNA.
Zfp61	Zfp61.bSep08	499094	3517	1007	5	335	zinc finger protein 61 (Zfp61) alternative variant bSep08, mRNA.
Zfp61	Zfp61.cSep08	499094	2981	906	4	233	zinc finger protein 61 (Zfp61) alternative variant cSep08, mRNA.
Zfp61	Zfp61.dSep08	499094	2100	684	3	82	zinc finger protein 61 (9.7 kD) (Zfp61) alternative variant dSep08, mRNA.
Zfp61	Zfp61.eSep08	499094	1813	686	2	66	zinc finger protein 61 (7.2 kD) (Zfp61) alternative variant eSep08, mRNA.
Zfp64	Zfp64.bSep08	311661	14577	414	1	138	zinc finger protein 64 (Zfp64) alternative variant bSep08, mRNA.
Zfp68	Zfp68.bSep08	304337	13885	3578	1	554	zinc finger protein 68 (64.7 kD) (Zfp68) alternative variant bSep08, mRNA.
Zfp68	Zfp68.cSep08	304337	842	780	2	141	zinc finger protein 68 (Zfp68) alternative variant cSep08, mRNA.
Zfp74	Zfp74.aSep08	365224	1666	769		256	zinc finger protein 74 (Zfp74) mRNA.
Zfp90	Zfp90.bSep08	498945	5613	543	1	165	zinc finger protein 90 (Zfp90) alternative variant bSep08, mRNA.

Zfp90	Zfp90.cSep08	498945	9637	741	5	102	zinc finger protein 90 (11.7 kD) (Zfp90) alternative variant cSep08, mRNA.
Zfp91	Zfp91.aSep08	246282	36872	5211	11	474	zinc finger protein 91 (Zfp91) alternative variant aSep08, mRNA.
Zfp91	Zfp91.bSep08	246282	26519	735	6	162	zinc finger protein 91 (Zfp91) alternative variant bSep08, mRNA.
Zfp91	Zfp91.cSep08	246282	2056	340	4	113	zinc finger protein 91 (Zfp91) alternative variant cSep08, mRNA.
Zfp91	Zfp91.eSep08	246282	32863	476	4	24	zinc finger protein 91 (Zfp91) alternative variant eSep08, mRNA.
Zfp93	Zfp93.bSep08	296399	12908	718	1	213	zinc finger protein 64 (Zfp93) alternative variant bSep08, mRNA.
Zfp110	Zfp110.bSep08	308362	13868	1884	6	292	zinc finger protein 110 (33.6 kD) (Zfp110) alternative variant bSep08, mRNA.
Zfp131	Zfp131.bSep08	310375	27720	3077	7	585	zinc finger protein 131 (66.7 kD) (Zfp131) alternative variant bSep08, complete mRNA.
Zfp131	Zfp131.dSep08	310375	14368	578	4	135	zinc finger protein 131 (Zfp131) alternative variant dSep08, mRNA.
Zfp131	Zfp131.eSep08	310375	2056	397	2	20	zinc finger protein 131 (2.2 kD) (Zfp131) alternative variant eSep08, mRNA.
Zfp143	Zfp143.bSep08	361627	13375	688	1	183	zinc finger protein 143 (Zfp143) alternative variant bSep08, mRNA.
Zfp148	Zfp148.aSep08	58820	111472	4084	9	800	zinc finger protein 148 (89.5 kD) (Zfp148) alternative variant aSep08, mRNA.
Zfp157	Zfp157.aSep08	360775	25989	735		176	zinc finger protein 157 (20.0 kD) (Zfp157) mRNA.
Zfp167	Zfp167.aSep08	363170	7800	2148	5	363	zinc finger protein 167 (Zfp167) alternative variant aSep08, mRNA.
Zfp167	Zfp167.bSep08	363170	3328	761	2	181	zinc finger protein 167 (Zfp167) alternative variant bSep08, mRNA.
Zfp179	Zfp179.bSep08	24916	1792	672	1	224	zinc finger protein 179 (Zfp179) alternative variant bSep08, mRNA.
Zfp180	Zfp180.bSep08	246279	17012	641	5	102	zinc finger protein 180 (Zfp180) alternative variant bSep08, mRNA.
Zfp184	Zfp184.aSep08	306966	35530	1186	7	274	zinc finger protein 184 (Kruppel-like) (Zfp184) alternative variant aSep08, mRNA.
Zfp184	Zfp184.cSep08	306966	5190	638	5	79	zinc finger protein 184 (Kruppel-like) (Zfp184) alternative variant cSep08, mRNA.
Zfp185	Zfp185.aSep08	689949	4574	352		117	zinc finger protein 185 (Zfp185) mRNA.
Zfp187	Zfp187.aSep08	266792	14523	954	1	115	zinc finger protein 187 (Zfp187) alternative variant aSep08, mRNA.
Zfp187	Zfp187.bSep08	266792	14710	389	3	84	zinc finger protein 187 (9.0 kD) (Zfp187) alternative variant bSep08, mRNA.
Zfp187	Zfp187.cSep08	266792	13874	416	1	98	zinc finger protein 187 (10.7 kD) (Zfp187) alternative variant cSep08, mRNA.
Zfp191	Zfp191.bSep08	360204	9092	3223	2	368	zinc finger protein 191 (41.9 kD) (Zfp191) alternative variant bSep08, mRNA.

Zfp191	Zfp191.cSep08	360204	3016	773	1	173	zinc finger protein 191 (Zfp191) alternative variant cSep08, mRNA.
Zfp192	Zfp192.aSep08	306974	6376	652		216	zinc finger protein 192 (Zfp192) mRNA.
Zfp207	Zfp207.bSep08	303763	9054	1263	10	420	zinc finger protein 207 (Zfp207) alternative variant bSep08, mRNA.
Zfp207	Zfp207.cSep08	303763	8397	1181	7	262	zinc finger protein 207 (28.5 kD) (Zfp207) alternative variant cSep08, mRNA.
Zfp207	Zfp207.dSep08	303763	9109	1405	6	82	zinc finger protein 207 (9.1 kD) (Zfp207) alternative variant dSep08, mRNA.
Zfp212	Zfp212.aSep08	297066	12353	2621		514	zinc finger protein 212 (Zfp212) mRNA.
Zfp217	Zfp217.bSep08	311764	30518	641	4	32	zinc finger protein 217 (3.7 kD) (Zfp217) alternative variant bSep08, mRNA.
Zfp219	Zfp219.cSep08	305848	11787	668	3	81	zinc finger protein 219 (8.7 kD) (Zfp219) alternative variant cSep08, mRNA.
Zfp236	Zfp236.aSep08	291409	8344	418		138	zinc finger protein 236 (Zfp236) mRNA.
Zfp251	Zfp251.aSep08	366954	689	359		119	zinc finger protein 251 (Zfp251) mRNA.
Zfp259andBud13	Zfp259andBud13.bSep08	300687	14989	1583	17	460	zinc finger protein 259 (50.5 kD) (Zfp259andBud13) alternative variant bSep08, mRNA.
Zfp259andBud13	Zfp259andBud13.bSep08	500989	14989	1583	17	460	zinc finger protein 259 (50.5 kD) (Zfp259andBud13) alternative variant bSep08, mRNA.
Zfp259andBud13	Zfp259andBud13.cSep08	300687	2612	660	5	155	zinc finger protein 259 (Zfp259andBud13) alternative variant cSep08, mRNA.
Zfp259andBud13	Zfp259andBud13.cSep08	500989	2612	660	5	155	zinc finger protein 259 (Zfp259andBud13) alternative variant cSep08, mRNA.
Zfp259andBud13	Zfp259andBud13.dSep08	300687	3068	765	4	111	zinc finger, ZPR1-type (Zfp259andBud13) alternative variant dSep08, mRNA.
Zfp259andBud13	Zfp259andBud13.dSep08	500989	3068	765	4	111	zinc finger, ZPR1-type (Zfp259andBud13) alternative variant dSep08, mRNA.
Zfp263	Zfp263.bSep08	287076	2509	399	1	132	zinc finger protein 263 (Zfp263) alternative variant bSep08, mRNA.
Zfp276	Zfp276.aSep08	307924	3885	1434	4	234	zinc finger protein (C2H2 type) 276 (26.8 kD) (Zfp276) alternative variant aSep08, mRNA.
Zfp276	Zfp276.bSep08	307924	684	389	1	105	zinc finger protein (C2H2 type) 276 (Zfp276) alternative variant bSep08, mRNA.
Zfp278	Zfp278.bSep08	305471	17873	2607	4	409	zinc finger (Zfp278) alternative variant bSep08, mRNA.
Zfp278	Zfp278.cSep08	305471	3213	753	2	163	POZ AT hook containing zinc finger 1 (Zfp278) alternative variant cSep08, mRNA.
Zfp278	Zfp278.dSep08	305471	15427	1202	3	160	zinc finger (17.3 kD) (Zfp278) alternative variant dSep08, mRNA.
Zfp278	Zfp278.eSep08	305471	2412	1066	2	105	zinc finger (11.5 kD) (Zfp278) alternative variant eSep08, mRNA.
Zfp282	Zfp282.aSep08	297065	23601	1993	3	664	zinc finger protein 282 (Zfp282) alternative variant aSep08, mRNA.
Zfp282	Zfp282.bSep08	297065	10884	732	3	206	zinc finger protein 282 (Zfp282) alternative variant bSep08, mRNA.

Zfp286	Zfp286.aSep08	497923	8629	1179	5	303	zinc finger protein 286 (Zfp286) alternative variant aSep08, mRNA.
Zfp287	Zfp287.bSep08	303212	4000	893	4	171	zinc finger protein 287 (Zfp287) alternative variant bSep08, mRNA.
Zfp294	Zfp294.bSep08	288308	2798	696	4	189	zinc finger protein 294 (Zfp294) alternative variant bSep08, mRNA.
Zfp294	Zfp294.cSep08	288308	2496	289	3	49	zinc finger protein 294 (Zfp294) alternative variant cSep08, mRNA.
Zfp296	Zfp296.aSep08	365511	1979	419		139	zinc finger protein 296 (Zfp296) mRNA.
Zfp313	Zfp313.bSep08	362277	10687	560	5	107	zinc finger protein 313 (11.9 kD) (Zfp313) alternative variant bSep08, complete mRNA.
Zfp316	Zfp316.bSep08	304293	5592	369	2	123	zinc finger protein 316 (Zfp316) alternative variant bSep08, mRNA.
Zfp322a	Zfp322a.aSep08	680201	11487	693	2	81	zinc finger protein 322a (3.9 kD) (Zfp322a) alternative variant aSep08, mRNA.
Zfp322a	Zfp322a.bSep08	680201	6096	396		36	zinc finger protein 322a (Zfp322a) alternative variant bSep08, mRNA.
Zfp329andVom2r-ps60	Zfp329andVom2r-ps60.bSep08	292612	12019	1758	2	551	vomer nasal 2 receptor, pseudogene 60 and zinc finger protein 329 (Zfp329andVom2r-ps60) alternative variant bSep08, mRNA.
Zfp329andVom2r-ps60	Zfp329andVom2r-ps60.bSep08	308361	12019	1758	2	551	vomer nasal 2 receptor, pseudogene 60 and zinc finger protein 329 (Zfp329andVom2r-ps60) alternative variant bSep08, mRNA.
Zfp330	Zfp330.bSep08	361387	6091	784	6	210	zinc finger protein 330 (Zfp330) alternative variant bSep08, mRNA.
Zfp330	Zfp330.cSep08	361387	1834	855	2	88	zinc finger protein 330 (Zfp330) alternative variant cSep08, mRNA.
Zfp335	Zfp335.aSep08	259270	5781	1643	12	547	zinc finger protein 335 (Zfp335) alternative variant aSep08, mRNA.
Zfp335	Zfp335.bSep08	259270	2283	1245	5	278	zinc finger protein 335 (Zfp335) alternative variant bSep08, mRNA.
Zfp335	Zfp335.cSep08	259270	721	632	2	36	zinc finger protein 335 (4.0 kD) (Zfp335) alternative variant cSep08, mRNA.
Zfp341	Zfp341.aSep08	690571	12868	782		260	zinc finger protein 341 (Zfp341) mRNA.
Zfp346	Zfp346.bSep08	306765	63657	1001	8	272	zinc finger protein 346 (30.5 kD) (Zfp346) alternative variant bSep08, mRNA.
Zfp354a	Zfp354a.bSep08	24522	9994	653	5	217	zinc finger protein 354A (Zfp354a) alternative variant bSep08, mRNA.
Zfp354a	Zfp354a.cSep08	24522	9927	589	5	131	zinc finger protein 354A (Zfp354a) alternative variant cSep08, mRNA.
Zfp354b	Zfp354b.aSep08	497898	2922	654		85	zinc finger protein 354B (Zfp354b) mRNA.
Zfp362	Zfp362.aSep08	297879	14281	2147	5	228	zinc finger protein 362 (Zfp362) alternative variant aSep08, mRNA.
Zfp362	Zfp362.cSep08	297879	672	381	2	60	zinc finger protein 362 (Zfp362) alternative variant cSep08, mRNA.
Zfp364	Zfp364.bSep08	362002	31759	2834	8	224	rabring 7 (25.2 kD) (Zfp364) alternative variant bSep08, mRNA.

Zfp364	Zfp364.cSep08	362002	57806	957	5	214	finger protein (Zfp364) alternative variant cSep08, mRNA.
Zfp364	Zfp364.dSep08	362002	1916	720	2	81	rabring 7 (9.4 kD) (Zfp364) alternative variant dSep08, mRNA.
Zfp365	Zfp365.bSep08	499425	14800	395	3	131	zinc finger protein 365 (Zfp365) alternative variant bSep08, mRNA.
Zfp367	Zfp367.bSep08	306695	24449	1067	1	218	zinc finger protein 367 (25.3 kD) (Zfp367) alternative variant bSep08, mRNA.
Zfp382	Zfp382.bSep08	246264	9753	730	2	242	zinc finger protein 382 (Zfp382) alternative variant bSep08, mRNA.
Zfp384	Zfp384.dSep08	171018	28815	1778	7	383	zinc finger protein 384 (Zfp384) alternative variant dSep08, mRNA.
Zfp384	Zfp384.eSep08	171018	15696	633	5	211	zinc finger protein 384 CRA d (Zfp384) alternative variant eSep08, mRNA.
Zfp384	Zfp384.fSep08	171018	8626	703	3	186	zinc finger protein 384 (Zfp384) alternative variant fSep08, mRNA.
Zfp384	Zfp384.gSep08	171018	15914	707	4	166	zinc finger protein 384 CRA d (Zfp384) alternative variant gSep08, mRNA.
Zfp385a	Zfp385a.aSep08	685474	20904	2450	7	415	zinc finger protein 385A (43.6 kD) (Zfp385a) alternative variant aSep08, mRNA.
Zfp385a	Zfp385a.bSep08	685474	8332	622	3	207	zinc finger protein 385A (Zfp385a) alternative variant bSep08, mRNA.
Zfp385a	Zfp385a.cSep08	685474	18850	557	4	185	zinc finger protein 385A (Zfp385a) alternative variant cSep08, mRNA.
Zfp385a	Zfp385a.dSep08	685474	16319	433	2	143	zinc finger protein 385A (Zfp385a) alternative variant dSep08, mRNA.
Zfp385a	Zfp385a.eSep08	685474	19227	508	4	126	zinc finger protein 385A (Zfp385a) alternative variant eSep08, mRNA.
Zfp385c	Zfp385c.aSep08	303537	28072	3150		501	zinc finger protein 385C (Zfp385c) mRNA.
Zfp386	Zfp386.bSep08	25165	12417	414	3	83	zinc finger protein 386 (Kruppel-like) (Zfp386) alternative variant bSep08, mRNA.
Zfp386	Zfp386.cSep08	25165	8822	366	2	83	zinc finger protein 386 (Kruppel-like) (9.7 kD) (Zfp386) alternative variant cSep08, mRNA.
Zfp407	Zfp407.aSep08	307213	257593	861		286	zinc finger protein 407 (Zfp407) mRNA.
Zfp410	Zfp410.bSep08	314310	13172	1135	6	243	zinc finger protein 410 (Zfp410) alternative variant bSep08, mRNA.
Zfp414	Zfp414.aSep08	299647	2220	1528	4	344	zinc finger protein 414 (38.2 kD) (Zfp414) alternative variant aSep08, mRNA.
Zfp414	Zfp414.cSep08	299647	1072	583	3	157	zinc finger protein 414 (Zfp414) alternative variant cSep08, mRNA.
Zfp414	Zfp414.dSep08	299647	2599	1845	5	120	zinc finger protein 414 (13.7 kD) (Zfp414) alternative variant dSep08, complete mRNA.
Zfp414	Zfp414.eSep08	299647	1373	767	2	107	zinc finger protein 414 (11.3 kD) (Zfp414) alternative variant eSep08, mRNA.
Zfp418	Zfp418.aSep08	292548	9267	762		222	zinc finger protein 418 (Zfp418) mRNA.
Zfp422	Zfp422.cSep08	360389	2081	443	2	65	putative protein (Zfp422) alternative variant cSep08, mRNA.

Zfp423	Zfp423.bSep08	94188	7107	543	1	113	zinc finger protein 423 (Zfp423) alternative variant bSep08, mRNA.
Zfp426	Zfp426.aSep08	367034	23240	680	5	113	CRA b like (Zfp426) alternative variant aSep08, mRNA.
Zfp426	Zfp426.bSep08	367034	23531	764	6	107	KRAB box (Zfp426) alternative variant bSep08, mRNA.
Zfp426	Zfp426.cSep08	367034	1179	428	1	55	putative protein (6.2 kD) (Zfp426) alternative variant cSep08, mRNA.
Zfp428	Zfp428.bSep08	361519	5997	512	2	111	zinc finger protein 428 (11.9 kD) (Zfp428) alternative variant bSep08, mRNA.
Zfp428	Zfp428.cSep08	361519	8662	509	2	106	zinc finger protein 428 (Zfp428) alternative variant cSep08, mRNA.
Zfp428	Zfp428.dSep08	361519	8401	662	2	69	zinc finger protein 428 (Zfp428) alternative variant dSep08, mRNA.
Zfp444	Zfp444.aSep08	292569	17389	1507	4	333	zinc finger protein 444 (36.1 kD) (Zfp444) alternative variant aSep08, complete mRNA.
Zfp444	Zfp444.bSep08	292569	11396	780	2	119	zinc finger protein 444 (Zfp444) alternative variant bSep08, mRNA.
Zfp451	Zfp451.bSep08	316312	6963	710	4	226	zinc finger protein 451 (Zfp451) alternative variant bSep08, mRNA.
Zfp451	Zfp451.cSep08	316312	12018	638	4	193	zinc finger protein 451 (Zfp451) alternative variant cSep08, mRNA.
Zfp451	Zfp451.dSep08	316312	30651	662	6	186	zinc finger protein 451 (Zfp451) alternative variant dSep08, mRNA.
Zfp451	Zfp451.eSep08	316312	7447	1126	2	79	zinc finger protein 451 (9.0 kD) (Zfp451) alternative variant eSep08, mRNA.
Zfp451	Zfp451.gSep08	316312	14460	620	2	32	zinc finger protein 451 (3.6 kD) (Zfp451) alternative variant gSep08, mRNA.
Zfp455	Zfp455.bSep08	286979	6620	908	1	115	zinc finger protein 455 (Zfp455) alternative variant bSep08, mRNA.
Zfp462	Zfp462.aSep08	362522	85096	1586		375	zinc finger protein 462 (Zfp462) alternative variant aSep08, mRNA.
Zfp462	Zfp462.bSep08	362522	39579	1273		172	zinc finger protein 462 (Zfp462) alternative variant bSep08, mRNA.
Zfp467	Zfp467.bSep08	500110	3099	837	2	216	zinc finger protein 467 (Zfp467) alternative variant bSep08, mRNA.
Zfp469	Zfp469.bSep08	304302	4253	780	2	259	zinc finger protein 469 (Zfp469) alternative variant bSep08, mRNA.
Zfp469	Zfp469.bSep08	679549	4253	780	2	259	zinc finger protein 469 (Zfp469) alternative variant bSep08, mRNA.
Zfp469	Zfp469.bSep08	679577	4253	780	2	259	zinc finger protein 469 (Zfp469) alternative variant bSep08, mRNA.
Zfp469	Zfp469.dSep08	304302	1234	724	2	84	zinc finger protein 469 (10.3 kD) (Zfp469) alternative variant dSep08, mRNA.
Zfp469	Zfp469.dSep08	679549	1234	724	2	84	zinc finger protein 469 (10.3 kD) (Zfp469) alternative variant dSep08, mRNA.
Zfp469	Zfp469.dSep08	679577	1234	724	2	84	zinc finger protein 469 (10.3 kD) (Zfp469) alternative variant dSep08, mRNA.
Zfp474	Zfp474.aSep08	307310	29629	1309		322	zinc finger protein 474 (Zfp474) mRNA.

zfp507	zfp507.bSep08	292816	17043	1799	1	179	zinc finger protein 507 (zfp507) alternative variant bSep08, mRNA.
Zfp509	Zfp509.bSep08	305428	13581	765	6	212	zinc finger protein 509 (Zfp509) alternative variant bSep08, mRNA.
Zfp509	Zfp509.dSep08	305428	4708	362	2	71	zinc finger protein 509 CRA b (Zfp509) alternative variant dSep08, mRNA.
Zfp511	Zfp511.bSep08	293586	4036	922	5	118	zinc finger protein 511 (13.6 kD) (Zfp511) alternative variant bSep08, mRNA.
Zfp512	Zfp512.aSep08	313906	31759	3376	12	523	zinc finger protein 512 (Zfp512) alternative variant aSep08, mRNA.
Zfp512	Zfp512.bSep08	313906	6398	721	7	240	zinc finger protein 512 (Zfp512) alternative variant bSep08, mRNA.
Zfp512	Zfp512.cSep08	313906	4082	741	1	161	zinc finger protein 512 (18.4 kD) (Zfp512) alternative variant cSep08, mRNA.
Zfp513	Zfp513.aSep08	313913	1581	1494		384	zinc finger protein 513 (Zfp513) mRNA.
Zfp516	Zfp516.aSep08	291406	16482	4778		286	zinc finger protein 516 (Zfp516) mRNA.
Zfp518b	Zfp518b.aSep08	498390	3520	2959	1	202	zinc finger protein 518B (Zfp518b) alternative variant aSep08, mRNA.
Zfp518b	Zfp518b.bSep08	498390	3411	2388	1	75	zinc finger protein 518B (Zfp518b) alternative variant bSep08, mRNA.
Zfp523	Zfp523.bSep08	361809	5702	730	1	243	zinc finger protein 523 (Zfp523) alternative variant bSep08, mRNA.
Zfp536	Zfp536.aSep08	292820	199725	421	1	46	zinc finger protein 536 (Zfp536) alternative variant aSep08, mRNA.
Zfp536	Zfp536.bSep08	292820	144023	346	1	46	zinc finger protein 536 (Zfp536) alternative variant bSep08, mRNA.
Zfp568	Zfp568.aSep08	308488	6073	615		117	zinc finger protein 568 (Zfp568) mRNA.
Zfp575	Zfp575.aSep08	308430	2097	1058	1	239	zinc finger protein 575 (26.4 kD) (Zfp575) alternative variant aSep08, mRNA.
Zfp592	Zfp592.bSep08	293038	996	616	3	204	zinc finger protein 592 (Zfp592) alternative variant bSep08, mRNA.
Zfp597	Zfp597.bSep08	266774	16003	1471	3	401	zinc finger protein 597 (46.0 kD) (Zfp597) alternative variant bSep08, mRNA.
Zfp597	Zfp597.cSep08	266774	1422	699	2	84	zinc finger protein 597 (9.6 kD) (Zfp597) alternative variant cSep08, mRNA.
Zfp598	Zfp598.bSep08	287119	2106	1781	4	343	zinc finger protein 598 (35.3 kD) (Zfp598) alternative variant bSep08, mRNA.
Zfp606	Zfp606.aSep08	292610	10520	449		33	zinc finger protein 606 (Zfp606) mRNA.
Zfp609	Zfp609.aSep08	363412	104849	7329	9	777	zinc finger protein 609 (81.7 kD) (Zfp609) alternative variant aSep08, mRNA.
Zfp609	Zfp609.bSep08	363412	1540	392	2	130	zinc finger protein 609 (Zfp609) alternative variant bSep08, mRNA.
Zfp612	Zfp612.cSep08	307839	5259	649	3	93	zinc finger protein 612 (Zfp612) alternative variant cSep08, mRNA.
Zfp612	Zfp612.dSep08	307839	4356	636	4	74	putative protein (8.0 kD) (Zfp612) alternative variant dSep08, mRNA.



Zfp612	Zfp612.eSep08	307839	4352	582	4	71	putative protein (7.6 kD) (Zfp612) alternative variant eSep08, mRNA.
Zfp622	Zfp622.bSep08	294846	1452	559	1	79	zinc finger protein 622 (Zfp622) alternative variant bSep08, mRNA.
Zfp637	Zfp637.aSep08	362425	4815	1164	3	325	zinc finger protein 637 (Zfp637) alternative variant aSep08, mRNA.
Zfp637	Zfp637.bSep08	362425	2646	672	3	200	zinc finger protein 637 (Zfp637) alternative variant bSep08, mRNA.
Zfp637	Zfp637.cSep08	362425	4134	525	3	174	zinc finger protein 637 (Zfp637) alternative variant cSep08, mRNA.
Zfp644	Zfp644.aSep08	305127	18422	2241		233	zinc finger protein 644 (Zfp644) mRNA.
Zfp646	Zfp646.aSep08	309003	3285	1550		365	zinc finger protein 646 (Zfp646) mRNA.
Zfp651	Zfp651.aSep08	316087	1997	504		168	zinc finger protein 651 (Zfp651) mRNA.
Zfp652	Zfp652.bSep08	497984	3965	283	1	94	zinc finger protein 652 (Zfp652) alternative variant bSep08, mRNA.
Zfp653	Zfp653.aSep08	300446	12586	1874	5	608	zinc finger protein 653 (Zfp653) alternative variant aSep08, mRNA.
Zfp653	Zfp653.cSep08	300446	831	362	1	41	zinc finger protein 653 (Zfp653) alternative variant cSep08, mRNA.
Zfp655	Zfp655.bSep08	360764	14120	453	4	117	zinc finger protein 655 (Zfp655) alternative variant bSep08, mRNA.
Zfp655	Zfp655.cSep08	360764	14125	626	5	116	zinc finger protein 655 (12.9 kD) (Zfp655) alternative variant cSep08, mRNA.
Zfp655	Zfp655.dSep08	360764	6366	1441	3	110	zinc finger protein 655 (12.2 kD) (Zfp655) alternative variant dSep08, complete mRNA.
Zfp655	Zfp655.eSep08	360764	4019	396	4	109	zinc finger protein 655 (Zfp655) alternative variant eSep08, mRNA.
Zfp655	Zfp655.fSep08	360764	14294	1181	5	103	zinc finger protein 655 (11.6 kD) (Zfp655) alternative variant fSep08, mRNA.
Zfp655	Zfp655.gSep08	360764	22710	389	3	64	zinc finger protein 655 (7.0 kD) (Zfp655) alternative variant gSep08, mRNA.
Zfp667	Zfp667.bSep08	308326	5544	1292	2	363	zinc finger protein 667 (Zfp667) alternative variant bSep08, mRNA.
Zfp668	Zfp668.bSep08	309002	7596	1028	2	280	zinc finger protein 668 (Zfp668) alternative variant bSep08, mRNA.
Zfp668	Zfp668.cSep08	309002	7317	464	1	154	zinc finger protein 668 (Zfp668) alternative variant cSep08, mRNA.
Zfp672	Zfp672.bSep08	303165	6482	996	3	207	zinc finger protein 672 (Zfp672) alternative variant bSep08, mRNA.
Zfp672	Zfp672.cSep08	303165	5770	711	3	48	zinc finger protein 672 (5.3 kD) (Zfp672) alternative variant cSep08, mRNA.
Zfp672	Zfp672.dSep08	303165	612	492	2	50	zinc finger protein 672 (Zfp672) alternative variant dSep08, mRNA.
Zfp688	Zfp688.aSep08	293511	3029	870	2	227	zinc finger protein 688 (Zfp688) alternative variant aSep08, mRNA.
Zfp688	Zfp688.bSep08	293511	2390	420	2	140	zinc finger protein 688 (Zfp688) alternative variant bSep08, mRNA.

Zfp689	Zfp689.bSep08	286996	4775	1104	3	277	zinc finger protein 689 (Zfp689) alternative variant bSep08, mRNA.
Zfp689	Zfp689.cSep08	286996	15241	733	4	169	zinc finger protein 689 (Zfp689) alternative variant cSep08, mRNA.
Zfp689	Zfp689.dSep08	286996	20460	671	5	20	zinc finger protein 689 (2.1 kD) (Zfp689) alternative variant dSep08, mRNA.
Zfp689	Zfp689.eSep08	286996	797	588	2	49	zinc finger protein 689 (5.5 kD) (Zfp689) alternative variant eSep08, mRNA.
Zfp692	Zfp692.aSep08	303164	8024	2430	12	533	zinc finger protein 692 (59.2 kD) (Zfp692) alternative variant aSep08, mRNA.
Zfp692	Zfp692.bSep08	303164	7028	1183	7	315	zinc finger protein 692 CRA a (Zfp692) alternative variant bSep08, mRNA.
Zfp692	Zfp692.dSep08	303164	3369	822	3	183	zinc finger protein 692 (20.3 kD) (Zfp692) alternative variant dSep08, mRNA.
Zfp692	Zfp692.eSep08	303164	2467	1261	6	172	zinc finger protein 692 CRA a (Zfp692) alternative variant eSep08, mRNA.
Zfp692	Zfp692.gSep08	303164	3124	686	2	60	putative protein (6.4 kD) (Zfp692) alternative variant gSep08, mRNA.
Zfp704	Zfp704.aSep08	310233	7165	437	2	85	zinc finger protein 704 (Zfp704) alternative variant aSep08, mRNA.
Zfp704	Zfp704.bSep08	310233	78072	549	2	40	zinc finger protein 704 (Zfp704) alternative variant bSep08, mRNA.
Zfp706	Zfp706.aSep08	500855	8153	2345	4	148	zinc finger protein 706 (Zfp706) alternative variant aSep08, mRNA.
Zfp709	Zfp709.bSep08	266773	645	431	2	36	zinc finger protein 709 (4.1 kD) (Zfp709) alternative variant bSep08, mRNA.
Zfp710	Zfp710.bSep08	293044	5869	424	2	84	zinc finger protein 710 (8.9 kD) (Zfp710) alternative variant bSep08, mRNA.
Zfp746	Zfp746.aSep08	312303	4143	2691	1	402	zinc finger protein 746 (Zfp746) alternative variant aSep08, mRNA.
Zfp746	Zfp746.bSep08	312303	1871	416	1	98	zinc finger protein 746 (Zfp746) alternative variant bSep08, mRNA.
Zfp770	Zfp770.aSep08	691610	5167	1669		448	zinc finger protein 770 (Zfp770) alternative variant aSep08, mRNA.
Zfp770	Zfp770.bSep08	691610	3896	398		34	zinc finger protein 770 (Zfp770) alternative variant bSep08, mRNA.
Zfp771	Zfp771.bSep08	308992	9672	552			
Zfp775	Zfp775.bSep08	312309	15314	362	2	120	zinc finger protein 775 (Zfp775) alternative variant bSep08, mRNA.
Zfp777	Zfp777.bSep08	502764	3613	861	2	251	zinc finger protein 777 (Zfp777) alternative variant bSep08, mRNA.
Zfp787	Zfp787.bSep08	365176	12791	1758	2	164	zinc finger protein 787 (Zfp787) alternative variant bSep08, mRNA.
Zfp787	Zfp787.cSep08	365176	775	446	2	33	zinc finger protein 787 (Zfp787) alternative variant cSep08, mRNA.
Zfp819	Zfp819.bSep08	308561	8738	745	2	165	zinc finger protein 819 (Zfp819) alternative variant bSep08, mRNA.

Zfp819	Zfp819.cSep08	308561	35524	754	2	152	zinc finger protein 819 (Zfp819) alternative variant cSep08, mRNA.
Zfp821	Zfp821.aSep08	307834	18218	1758	8	369	zinc finger protein 821 (41.5 kD) (Zfp821) alternative variant aSep08, complete mRNA.
Zfp821	Zfp821.bSep08	307834	12088	986	4	124	putative protein (Zfp821) alternative variant bSep08, mRNA.
Zfp821	Zfp821.cSep08	307834	10168	379	4	89	putative protein (Zfp821) alternative variant cSep08, mRNA.
Zfpm2	Zfpm2.bSep08	314930	121690	346	1	110	zinc finger protein, multitype 2 (Zfpm2) alternative variant bSep08, mRNA.
Zfr	Zfr.aSep08	365703	29830	3312	13	666	zinc finger RNA binding protein (Zfr) alternative variant aSep08, mRNA.
Zfr	Zfr.bSep08	365703	32856	1666	9	457	zinc finger RNA binding protein (Zfr) alternative variant bSep08, mRNA.
Zfr	Zfr.cSep08	365703	33336	1787	6	373	zinc finger RNA binding protein (Zfr) alternative variant cSep08, mRNA.
Zfr	Zfr.dSep08	365703	13926	1733	5	128	zinc finger RNA binding protein (14.4 kD) (Zfr) alternative variant dSep08, mRNA.
Zfr	Zfr.eSep08	365703	5864	797	4	122	zinc finger RNA binding protein (Zfr) alternative variant eSep08, mRNA.
Zfr	Zfr.fSep08	365703	612	501	2	66	zinc finger RNA binding protein (Zfr) alternative variant fSep08, mRNA.
Zfr	Zfr.gSep08	365703	523	234	2	61	zinc finger RNA binding protein (Zfr) alternative variant gSep08, mRNA.
Zfx	Zfx.dSep08	367832	22214	477	3	51	zinc finger protein X-linked (Zfx) alternative variant dSep08, mRNA.
Zfyve1	Zfyve1.bSep08	299188	12268	1800	8	452	putative protein of metazoan origin (Zfyve1) alternative variant bSep08, mRNA.
Zfyve1	Zfyve1.cSep08	299188	11305	779	5	225	putative protein of metazoan origin (Zfyve1) alternative variant cSep08, mRNA.
Zfyve1	Zfyve1.eSep08	299188	6816	765	5	128	putative protein of metazoan origin (Zfyve1) alternative variant eSep08, mRNA.
Zfyve1	Zfyve1.gSep08	299188	2025	404	2	38	putative protein (Zfyve1) alternative variant gSep08, mRNA.
Zfyve1	Zfyve1.hSep08	299188	1914	352	2	41	putative protein (Zfyve1) alternative variant hSep08, mRNA.
Zfyve9	Zfyve9.bSep08	313477	38577	916	5	274	putative cytoplasmic protein of metazoan origin (30.7 kD) (Zfyve9) alternative variant bSep08, mRNA.
Zfyve9	Zfyve9.cSep08	313477	38334	742	2	84	putative protein of metazoan origin (Zfyve9) alternative variant cSep08, mRNA.
Zfyve16	Zfyve16.aSep08	499508	19778	1724	9	574	zinc finger, FYVE-type (Zfyve16) alternative variant aSep08, mRNA.
Zfyve16	Zfyve16.bSep08	499508	5157	608	1	106	putative protein of metazoan origin (Zfyve16) alternative variant bSep08, mRNA.
Zfyve19	Zfyve19.bSep08	499871	6256	1655	10	313	putative protein, with a coiled coil domain, of fungal and metazoan origin (35.0 kD) (Zfyve19) alternative variant bSep08, mRNA.

Zfyve19	Zfyve19.cSep08	499871	6003	830	7	252	zinc finger, FYVE-type (Zfyve19) alternative variant cSep08, mRNA.
Zfyve19	Zfyve19.dSep08	499871	984	409	4	99	putative protein of vertebrate origin (Zfyve19) alternative variant dSep08, mRNA.
Zfyve19	Zfyve19.eSep08	499871	1667	882	3	96	putative protein (Zfyve19) alternative variant eSep08, mRNA.
Zfyve19	Zfyve19.fSep08	499871	2698	1013	3	76	zinc finger, FYVE-type (8.4 kD) (Zfyve19) alternative variant fSep08, mRNA.
Zfyve20	Zfyve20.cSep08	312562	1117	409	2	15	hypothetical protein LOC688545 (1.8 kD) (Zfyve20) alternative variant cSep08, mRNA.
Zfyve20	Zfyve20.cSep08	688545	1117	409	2	15	hypothetical protein LOC688545 (1.8 kD) (Zfyve20) alternative variant cSep08, mRNA.
Zfyve21	Zfyve21.bSep08	362789	15844	738	2	246	zinc finger, FYVE-type (Zfyve21) alternative variant bSep08, mRNA.
Zfyve21	Zfyve21.cSep08	362789	19661	795	2	245	zinc finger, FYVE-type (Zfyve21) alternative variant cSep08, mRNA.
Zfyve21	Zfyve21.dSep08	362789	16627	1337	2	234	zinc finger, FYVE-type (26.0 kD) (Zfyve21) alternative variant dSep08, complete mRNA.
Zfyve21	Zfyve21.eSep08	362789	16625	1332	2	233	zinc finger, FYVE-type (25.9 kD) (Zfyve21) alternative variant eSep08, complete mRNA.
Zfyve27	Zfyve27.bSep08	309376	13003	1058	8	294	putative endoplasmic reticulum protein, with at least 2 transmembrane domains, of vertebrate origin (32.5 kD) (Zfyve27) alternative variant bSep08, mRNA.
Zfyve27	Zfyve27.dSep08	309376	4183	928	3	116	putative protein of metazoan origin (12.8 kD) (Zfyve27) alternative variant dSep08, mRNA.
Zfyve27	Zfyve27.eSep08	309376	4684	522	4	97	zinc finger, FYVE-type (10.7 kD) (Zfyve27) alternative variant eSep08, mRNA.
Zfyve27	Zfyve27.fSep08	309376	5077	771	5	97	zinc finger, FYVE-type (10.7 kD) (Zfyve27) alternative variant fSep08, mRNA.
Zfyve27	Zfyve27.gSep08	309376	1074	537	2	66	putative protein of mammalian origin (Zfyve27) alternative variant gSep08, mRNA.
Zhx1	Zhx1.bSep08	171159	3881	630	2	148	zinc fingers and homeoboxes 1 (Zhx1) alternative variant bSep08, mRNA.
Zhx1	Zhx1.cSep08	171159	21784	1927	2	84	zinc fingers and homeoboxes 1 (10.1 kD) (Zhx1) alternative variant cSep08, mRNA.
Zic1	Zic1.bSep08	64618	3199	2144	2	135	zinc finger protein of the cerebellum 1 (Zic1) alternative variant bSep08, mRNA.
Zic4	Zic4.cSep08	315882	4220	740	2	78	zinc finger protein of the cerebellum 4 (Zic4) alternative variant cSep08, mRNA.
Zkscan2	Zkscan2.aSep08	368120	3319	1332		369	zinc finger with KRAB and SCAN domains 2 (Zkscan2) mRNA.
Zkscan3	Zkscan3.cSep08	306977	6639	772	2	134	zinc finger with KRAB and SCAN domains 3 (Zkscan3) alternative variant cSep08, mRNA.
Zkscan3	Zkscan3.eSep08	306977	14991	1799	3	51	zinc finger with KRAB and SCAN domains 3 (Zkscan3) alternative variant eSep08, mRNA.
Zkscan5	Zkscan5.bSep08	304275	13180	1571	1	260	zinc finger with KRAB and SCAN domains 5 (29.6 kD) (Zkscan5) alternative variant bSep08, mRNA.

Zkscan6	Zkscan6.bSep08	303226	7023	1259	1	326	zinc finger with KRAB and SCAN domains 6 (Zkscan6) alternative variant bSep08, mRNA.
Zkscan17	Zkscan17.aSep08	287361	17044	1992	2	482	zinc finger with KRAB and SCAN domains 17 (Zkscan17) alternative variant aSep08, mRNA.
Zkscan17	Zkscan17.bSep08	287361	2959	758	1	147	zinc finger with KRAB and SCAN domains 17 (Zkscan17) alternative variant bSep08, mRNA.
Zmat2	Zmat2.bSep08	307491	1882	339	2	69	zinc finger, matrin type 2 (7.6 kD) (Zmat2) alternative variant bSep08, mRNA.
Zmat5	Zmat5.aSep08	501926	15602	657		153	zinc finger, matrin type 5 (Zmat5) mRNA.
Zmiz1	Zmiz1.bSep08	361103	5114	1720	3	420	zinc finger, MIZ-type containing 1 (Zmiz1) alternative variant bSep08, mRNA.
Zmiz1	Zmiz1.dSep08	361103	82509	1101	5	121	zinc finger, MIZ-type containing 1 (14.1 kD) (Zmiz1) alternative variant dSep08, mRNA.
Zmiz1	Zmiz1.eSep08	361103	116056	706	6	65	zinc finger, MIZ-type containing 1 (7.6 kD) (Zmiz1) alternative variant eSep08, mRNA.
Zmiz2	Zmiz2.aSep08	289783	8699	3118	13	692	zinc finger, MIZ-type containing 2 (Zmiz2) alternative variant aSep08, mRNA.
Zmiz2	Zmiz2.bSep08	289783	2730	2646	2	246	zinc finger, MIZ-type containing 2 (25.5 kD) (Zmiz2) alternative variant bSep08, mRNA.
Zmiz2	Zmiz2.cSep08	289783	3295	719	6	229	zinc finger, MIZ-type containing 2 (Zmiz2) alternative variant cSep08, mRNA.
Zmiz2	Zmiz2.dSep08	289783	1138	319	3	106	zinc finger, MIZ-type containing 2 (Zmiz2) alternative variant dSep08, mRNA.
Zmpste24	Zmpste24.bSep08	313564	24225	798	5	93	zinc metalloproteinase, STE24 homolog (S. cerevisiae) (Zmpste24) alternative variant bSep08, mRNA.
Zmpste24	Zmpste24.cSep08	313564	12852	300	3	65	zinc metalloproteinase, STE24 homolog (S. cerevisiae) (Zmpste24) alternative variant cSep08, mRNA.
Zmym1	Zmym1.aSep08	313604	17100	3327	4	1086	HAT dimerisation (Zmym1) alternative variant aSep08, mRNA.
Zmym1	Zmym1.cSep08	313604	16221	1796	3	439	zinc finger MYM-type 1 CRA d (Zmym1) alternative variant cSep08, mRNA.
Zmym2	Zmym2.aSep08	305913	37442	1807	5	518	zinc finger, MYM-type 2 (Zmym2) alternative variant aSep08, mRNA.
Zmym2	Zmym2.bSep08	305913	12919	877	5	292	zinc finger, MYM-type 2 (Zmym2) alternative variant bSep08, mRNA.
Zmym2	Zmym2.cSep08	305913	11909	1308	6	261	zinc finger, MYM-type 2 (30.0 kD) (Zmym2) alternative variant cSep08, mRNA.
Zmym2	Zmym2.dSep08	305913	12776	698	5	232	zinc finger, MYM-type 2 (Zmym2) alternative variant dSep08, mRNA.
Zmym2	Zmym2.eSep08	305913	9614	686	5	227	zinc finger, MYM-type 2 (Zmym2) alternative variant eSep08, mRNA.
Zmym3	Zmym3.bSep08	317260	3849	1141	5	365	putative protein of vertebrate origin (Zmym3) alternative variant bSep08, mRNA.
Zmym3	Zmym3.cSep08	317260	3982	1223	6	279	zinc finger MYM-type 3 (Zmym3) alternative variant cSep08, mRNA.
Zmym3	Zmym3.dSep08	317260	1185	739	2	166	putative protein (Zmym3) alternative variant dSep08, mRNA.

Zmym3	Zmym3.eSep08	317260	781	492	2	134	zinc finger MYM-type 3 (Zmym3) alternative variant eSep08, mRNA.
Zmym3	Zmym3.fSep08	317260	1016	393	2	122	putative protein (Zmym3) alternative variant fSep08, mRNA.
Zmym4	Zmym4.bSep08	313598	2795	665	3	205	zinc finger, MYM-type 4 (Zmym4) alternative variant bSep08, mRNA.
Zmym4	Zmym4.cSep08	313598	8271	736	3	185	zinc finger, MYM-type 4 (Zmym4) alternative variant cSep08, mRNA.
Zmym4	Zmym4.dSep08	313598	4871	394	3	130	zinc finger, MYM-type 4 (Zmym4) alternative variant dSep08, mRNA.
Zmym6	Zmym6.bSep08	362602	19427	2018	6	198	zinc finger, MYM-type 6 (Zmym6) alternative variant bSep08, mRNA.
Zmynd10	Zmynd10.bSep08	363139	1675	1477	3	132	zinc finger, MYND domain-containing 10 (Zmynd10) alternative variant bSep08, mRNA.
Zmynd10	Zmynd10.cSep08	363139	1640	469	3	92	zinc finger, MYND domain-containing 10 (Zmynd10) alternative variant cSep08, mRNA.
Zmynd10	Zmynd10.dSep08	363139	863	747	2	80	zinc finger, MYND domain-containing 10 (8.9 kD) (Zmynd10) alternative variant dSep08, mRNA.
Zmynd10	Zmynd10.eSep08	363139	1051	606	4	50	zinc finger, MYND domain-containing 10 (Zmynd10) alternative variant eSep08, mRNA.
Zmynd11	Zmynd11.eSep08	291259	53390	521	4	141	hypothetical protein LOC689685 (16.1 kD) (Zmynd11) alternative variant eSep08, mRNA.
Zmynd11	Zmynd11.eSep08	689685	53390	521	4	141	hypothetical protein LOC689685 (16.1 kD) (Zmynd11) alternative variant eSep08, mRNA.
Zmynd11	Zmynd11.fSep08	291259	77907	661	6	59	hypothetical protein LOC689685 (6.8 kD) (Zmynd11) alternative variant fSep08, mRNA.
Zmynd11	Zmynd11.fSep08	689685	77907	661	6	59	hypothetical protein LOC689685 (6.8 kD) (Zmynd11) alternative variant fSep08, mRNA.
Zmynd12	Zmynd12.bSep08	313552	1316	660	3	113	putative protein (12.2 kD) (Zmynd12) alternative variant bSep08, mRNA.
Zmynd12	Zmynd12.cSep08	313552	5276	402	2	70	putative protein of mammalian origin (Zmynd12) alternative variant cSep08, mRNA.
Zmynd17	Zmynd17.bSep08	289904	2788	1036	3	225	putative protein of vertebrate origin (Zmynd17) alternative variant bSep08, mRNA.
Zmynd17	Zmynd17.dSep08	289904	1321	1006	2	104	putative protein of vertebrate origin (11.8 kD) (Zmynd17) alternative variant dSep08, mRNA.
Zmynd17	Zmynd17.eSep08	289904	1440	635	3	97	zinc finger mynd containing 17 (Zmynd17) alternative variant eSep08, mRNA.
Znf124	Znf124.bSep08	314586	4336	443	2	123	zinc finger protein 124 (HZF-16) (Znf124) alternative variant bSep08, mRNA.
Znf142	Znf142.bSep08	316524	5520	1721	4	509	zinc finger protein 142 (clone pHZ-49) (Znf142) alternative variant bSep08, mRNA.
Znf142	Znf142.cSep08	316524	9454	756	4	175	zinc finger protein 142 (clone pHZ-49) (Znf142) alternative variant cSep08, mRNA.
Znf213	Znf213.aSep08	287094	3028	1712	5	570	zinc finger protein 213 (Znf213) alternative variant aSep08, mRNA.

Znf213	Znf213.cSep08	287094	2804	1436	4	199	zinc finger protein 213 (Znf213) alternative variant cSep08, mRNA.
Znf213	Znf213.dSep08	287094	2270	607	5	135	zinc finger protein 213 (Znf213) alternative variant dSep08, mRNA.
Znf213	Znf213.eSep08	287094	1229	457	4	116	zinc finger protein 213 (Znf213) alternative variant eSep08, mRNA.
Znf291	Znf291.aSep08	117521	64373	944		314	zinc finger protein 291 (Znf291) alternative variant aSep08, mRNA.
Znf291	Znf291.bSep08	117521	38994	2408		214	zinc finger protein 291 (24.0 kD) (Znf291) alternative variant bSep08, mRNA.
Znf291	Znf291.cSep08	117521	268132	1728			
Znf307	Znf307.aSep08	291164	7535	1783		479	zinc finger protein 307 (Znf307) alternative variant aSep08, mRNA.
Znf397	Znf397.aSep08	364827	4719	794		256	zinc finger protein 397 (Znf397) mRNA.
Znf498	Znf498.bSep08	363872	5158	1564	2	402	zinc finger protein 498 (44.2 kD) (Znf498) alternative variant bSep08, mRNA.
Znf512b	Znf512b.bSep08	311721	1606	1024	2	83	zinc finger protein 512B (8.6 kD) (Znf512b) alternative variant bSep08, mRNA.
Znf532	Znf532.bSep08	307362	102078	2641	2	816	zinc finger protein 532 (Znf532) alternative variant bSep08, mRNA.
Znf532	Znf532.cSep08	307362	49753	845	2	197	zinc finger protein 532 (Znf532) alternative variant cSep08, mRNA.
Znf532	Znf532.dSep08	307362	48180	409	2	73	zinc finger protein 532 (Znf532) alternative variant dSep08, mRNA.
Znf618	Znf618.aSep08	313253	12307	1695		548	zinc finger protein 618 (Znf618) mRNA.
Znhit3	Znhit3.aSep08	497975	8174	963	2	146	zinc finger, HIT type 3 (Znhit3) alternative variant aSep08, mRNA.
Znhit3	Znhit3.bSep08	497975	10204	1323	3	111	zinc finger, HIT type 3 (12.4 kD) (Znhit3) alternative variant bSep08, mRNA.
Znhit3	Znhit3.cSep08	497975	28074	640	2	102	zinc finger, HIT type 3 (Znhit3) alternative variant cSep08, mRNA.
Znhit3	Znhit3.dSep08	497975	14543	1017	2	93	zinc finger, HIT type 3 (Znhit3) alternative variant dSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.aSep08	297381	6243	3103	9	375	high mobility group AT-hook 1-like 4 (40.5 kD) (Znhit4andWbp1) alternative variant aSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.aSep08	500225	6243	3103	9	375	high mobility group AT-hook 1-like 4 (40.5 kD) (Znhit4andWbp1) alternative variant aSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.dSep08	297381	1975	913	3	230	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant dSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.dSep08	500225	1975	913	3	230	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant dSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.eSep08	297381	1814	691	5	230	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant eSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.eSep08	500225	1814	691	5	230	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant eSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.fSep08	297381	14455	816	8	207	high mobility group AT-hook 1-like 4 (Znhit4andWbp1) alternative variant fSep08, mRNA.

Znhit4andWbp1	Znhit4andWbp1.fSep08	500225	14455	816	8	207	high mobility group AT-hook 1-like 4 (Znhit4andWbp1) alternative variant fSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.gSep08	297381	1684	868	3	167	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant gSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.gSep08	500225	1684	868	3	167	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant gSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.hSep08	297381	1232	906	2	151	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant hSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.hSep08	500225	1232	906	2	151	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant hSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.iSep08	297381	1362	424	4	117	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant iSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.iSep08	500225	1362	424	4	117	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant iSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.jSep08	297381	1650	1509	2	102	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant jSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.jSep08	500225	1650	1509	2	102	WW domain binding protein 1 like (Znhit4andWbp1) alternative variant jSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.kSep08	297381	2807	488	3	29	putative protein (3.2 kD) (Znhit4andWbp1) alternative variant kSep08, mRNA.
Znhit4andWbp1	Znhit4andWbp1.kSep08	500225	2807	488	3	29	putative protein (3.2 kD) (Znhit4andWbp1) alternative variant kSep08, mRNA.
Znrd1	Znrd1.bSep08	361784	3373	587	2	58	putative protein (6.6 kD) (Znrd1) alternative variant bSep08, mRNA.
Znrf1	Znrf1.bSep08	362367	128776	650	5	152	zinc and ring finger 1 (Znrf1) alternative variant bSep08, mRNA.
zobor	zobor.aSep08		840	589		113	putative nuclear protein (12.2 kD) (zobor) mRNA.
zoby	zoby.aSep08		1177	420		140	host cell factor (zoby) mRNA.
zochy	zochy.aSep08		6916	739		246	centrosomal protein 152kDa (zochy) mRNA.
zodar	zodar.aSep08		1636	618		40	putative protein (zodar) mRNA.
zodoy	zodoy.aSep08		102219	579		130	putative protein of metazoan origin (14.6 kD) (zodoy) mRNA.
zoflu	zoflu.aSep08		8263	745		56	putative protein (6.5 kD) (zoflu) mRNA.
zofly	zofly.aSep08		2134	547		87	putative nuclear protein (9.5 kD) (zofly) mRNA.
zogar	zogar.aSep08		21142	634		43	putative protein (zogar) mRNA.
zoja	zoja.bSep08		32225	1138	3	53	putative protein (zoja) alternative variant bSep08, mRNA.
zojey	zojey.aSep08		3444	610		68	putative protein (zojey) mRNA.
zokee	zokee.aSep08		116924	640		183	formin 2 (zokee) mRNA.
zokler	zokler.aSep08		2794	740		63	putative protein of mammalian origin (7.4 kD) (zokler) mRNA.
zolo	zolo.aSep08		6379	1800	8	546	CRA a (zolo) alternative variant aSep08, mRNA.
zolo	zolo.bSep08		7364	1624	13	541	CRA b (zolo) alternative variant bSep08, mRNA.
zolo	zolo.cSep08		3139	982	4	252	CRA b (zolo) alternative variant cSep08, mRNA.
zomee	zomee.aSep08		4039	621		48	solute carrier family 37 member 3 like (5.2 kD) (zomee) mRNA.



zonoy	zonoy.aSep08		8123	403		25	putative protein (zonoy) mRNA.
zopor	zopor.bSep08		5384	803		57	putative protein (6.4 kD) (zopor) alternative variant bSep08, mRNA.
zorbor	zorbor.aSep08		24345	552	1	41	putative protein (zorbor) alternative variant aSep08, mRNA.
zorbor	zorbor.bSep08		24325	300		41	putative protein (4.8 kD) (zorbor) alternative variant bSep08, mRNA.
zorby	zorby.aSep08		907	614		48	putative protein (5.5 kD) (zorby) mRNA.
zorchy	zorchy.aSep08		1679	689		55	putative protein (zorchy) mRNA.
zordar	zordar.aSep08		13006	444		30	putative protein (3.5 kD) (zordar) mRNA.
zordoy	zordoy.aSep08		2047	334		50	putative protein (zordoy) mRNA.
zorflu	zorflu.aSep08		1323	431		143	putative protein of metazoan origin (zorflu) mRNA.
zorfly	zorfly.aSep08		3271	199		63	gag protein like (zorfly) mRNA.
zorgar	zorgar.aSep08		882	296		51	putative protein (zorgar) mRNA.
zorja	zorja.aSep08		10424	774		67	SET domain bifurcated 2 (zorja) mRNA.
zorjey	zorjey.aSep08		57903	403		97	putative protein (10.0 kD) (zorjey) mRNA.
zorkee	zorkee.aSep08		16052	702		31	putative protein (3.1 kD) (zorkee) mRNA.
zorkler	zorkler.aSep08		1049	266		51	putative protein (5.6 kD) (zorkler) mRNA.
zorlo	zorlo.aSep08		12122	619		58	putative protein (6.6 kD) (zorlo) mRNA.
zormee	zormee.aSep08		1882	545		64	nuclear receptor co-repressor 1 (zormee) mRNA.
zornoy	zornoy.aSep08		121415	609		113	putative protein of vertebrate origin (zornoy) mRNA.
zorpor	zorpor.aSep08		14774	811		270	5'-3' exoribonuclease 1 (zorpor) mRNA.
zorroy	zorroy.aSep08		10361	521		173	2 cell cycle control (zorroy) mRNA.
zorsa	zorsa.aSep08		3369	2431	2	785	splicing factor arginine serine-rich 2 interacting protein CRA b (zorsa) alternative variant aSep08, mRNA.
zorsa	zorsa.bSep08		2483	988	4	258	splicing factor arginine serine-rich 2 interacting protein CRA b (zorsa) alternative variant bSep08, mRNA.
zorsa	zorsa.cSep08		2205	707	4	164	splicing factor arginine serine-rich 2 interacting protein CRA b (zorsa) alternative variant cSep08, mRNA.
zorsa	zorsa.dSep08		1205	1021	2	100	splicing factor arginine serine-rich 2 interacting protein (11.5 kD) (zorsa) alternative variant dSep08, mRNA.
zorshee	zorshee.aSep08		551	383		91	CRA b like (zorshee) mRNA.
zortu	zortu.aSep08		100016	1665		139	ataxin 7-like (zortu) mRNA.
zorvar	zorvar.aSep08		646	535	2	69	putative protein (7.3 kD) (zorvar) alternative variant aSep08, mRNA.
zorvo	zorvo.aSep08		22877	917		305	guanine nucleotide-exchange A-inhibited (zorvo) mRNA.
zorwer	zorwer.aSep08		10194	409		44	putative protein (4.9 kD) (zorwer) mRNA.
zorwey	zorwey.aSep08		18628	500		69	putative protein (7.5 kD) (zorwey) mRNA.
zosa	zosa.aSep08		3385	351		116	putative protein (zosa) mRNA.
zoshee	zoshee.aSep08		2864	325		56	putative protein human specific (zoshee) mRNA.
zotu	zotu.aSep08		3681	441		57	putative protein human specific (zotu) mRNA.
zovar	zovar.aSep08		19275	552		36	putative protein (zovar) mRNA.
zovo	zovo.aSep08		39242	435		57	putative protein (zovo) mRNA.
zowey	zowey.aSep08		2460	695		122	putative protein of mammalian origin (zowey) mRNA.

zoybor	zoybor.aSep08		5730	740		65	putative protein (zoybor) mRNA.
zoychy	zoychy.aSep08		8173	675	1	124	adaptor-related protein complex (zoychy) alternative variant aSep08, mRNA.
zoychy	zoychy.bSep08		12957	372	1	123	adaptor-related protein complex (zoychy) alternative variant bSep08, mRNA.
zoydar	zoydar.aSep08		3698	333		58	putative protein (zoydar) mRNA.
zoydoy	zoydoy.aSep08		3545	480		76	putative protein (zoydoy) mRNA.
zoyflu	zoyflu.aSep08		3653	432	2	143	CRA a (zoyflu) alternative variant aSep08, mRNA.
zoyfly	zoyfly.aSep08		8768	417		75	putative protein (zoyfly) mRNA.
zoygar	zoygar.aSep08		3535	733		60	putative protein (zoygar) mRNA.
zoyja	zoyja.aSep08		5307	498		63	ATPase aminophospholipid transporter-like Class I type 8A member 2 (zoyja) mRNA.
zoyjey	zoyjey.aSep08		973	501		32	putative protein (zoyjey) mRNA.
zoykee	zoykee.aSep08		690	483		62	putative protein (zoykee) mRNA.
zoykler	zoykler.aSep08		127118	1288		91	putative protein (9.9 kD) (zoykler) mRNA.
zoylo	zoylo.aSep08		3774	787		149	putative protein (16.1 kD) (zoylo) mRNA.
zoymee	zoymee.bSep08		1067	474	2	21	putative protein (2.4 kD) (zoymee) alternative variant bSep08, mRNA.
zoymee	zoymee.cSep08		1039	253	3	46	putative protein (5.0 kD) (zoymee) alternative variant cSep08, mRNA.
zoynoy	zoynoy.aSep08		5213	317		39	putative protein (4.7 kD) (zoynoy) mRNA.
zoypor	zoypor.aSep08		4047	549		70	putative protein (zoypor) alternative variant aSep08, mRNA.
zoyroy	zoyroy.aSep08		21499	859	2	285	2 cell cycle control (zoyroy) alternative variant aSep08, mRNA.
zoyroy	zoyroy.bSep08		21287	815	1	271	2 cell cycle control (zoyroy) alternative variant bSep08, mRNA.
zoysa	zoysa.aSep08		1407	281		93	splicing factor arginine serine-rich 2 interacting protein (zoysa) mRNA.
zoyshee	zoyshee.aSep08		1202	417		87	putative protein (zoyshee) mRNA.
zoytu	zoytu.aSep08		7472	680	2	63	putative protein (zoytu) alternative variant aSep08, mRNA.
zoytu	zoytu.bSep08		1078	656	1	83	CRA a like (8.9 kD) (zoytu) alternative variant bSep08, mRNA.
zoyvar	zoyvar.aSep08		916	460		52	putative protein (5.7 kD) (zoyvar) mRNA.
zoyvo	zoyvo.aSep08		4967	504		167	guanine nucleotide-exchange A-inhibited (zoyvo) mRNA.
zoywer	zoywer.aSep08		84860	387		68	putative protein (zoywer) mRNA.
zoywey	zoywey.aSep08		2872	412		52	putative protein (zoywey) mRNA.
Zp3r	Zp3r.bSep08	289010	14724	518	1	145	zona pellucida 3 receptor (Zp3r) alternative variant bSep08, mRNA.
Zranb2	Zranb2.bSep08	58821	2265	1487	4	398	putative protein, with a coiled coil domain (Zranb2) alternative variant bSep08, mRNA.
Zranb2	Zranb2.cSep08	58821	11234	1782	5	366	ZIS1 like (Zranb2) alternative variant cSep08, mRNA.
Zranb2	Zranb2.dSep08	58821	8208	1770	7	225	zinc finger protein 265 (Zranb2) alternative variant dSep08, mRNA.

Zranb2	Zranb2.eSep08	58821	5605	774	6	142	putative protein of mammalian origin (15.5 kD) (Zranb2) alternative variant eSep08, mRNA.
Zranb2	Zranb2.fSep08	58821	3857	634	3	132	putative protein of mammalian origin (Zranb2) alternative variant fSep08, mRNA.
Zranb2	Zranb2.gSep08	58821	2082	837	2	85	putative protein of mammalian origin (Zranb2) alternative variant gSep08, mRNA.
Zranb2	Zranb2.iSep08	58821	1882	518	3	54	zinc finger protein 265 CRA a (Zranb2) alternative variant iSep08, mRNA.
Zrsr2	Zrsr2.aSep08	302670	4044	684		104	zinc finger (CCCH type), RNA binding motif and serine/arginine rich 2 (Zrsr2) mRNA.
Zscan10	Zscan10.bSep08	302962	4478	382	1	73	zinc finger (Zscan10) alternative variant bSep08, mRNA.
Zscan12	Zscan12.aSep08	266716	5848	2574		503	transcriptional regulator SCAN and zinc finger, C2H2-type (58.4 kD) (Zscan12) mRNA.
Zscan21	Zscan21.bSep08	304342	15760	1473	3	324	zinc finger (36.7 kD) (Zscan21) alternative variant bSep08, mRNA.
Zscan21	Zscan21.cSep08	304342	5711	698	2	232	zinc finger (Zscan21) alternative variant cSep08, mRNA.
Zswim3	Zswim3.bSep08	311630	15262	2813	1	632	zinc finger SWIM-type containing 3 (71.5 kD) (Zswim3) alternative variant bSep08, mRNA.
Zswim4	Zswim4.bSep08	304655	1496	275	2	32	zinc finger SWIM-type containing 4 (Zswim4) alternative variant bSep08, mRNA.
Zswim5	Zswim5.bSep08	313524	8338	1803	1	462	putative protein of metazoan origin (Zswim5) alternative variant bSep08, mRNA.
ZU5.0	ZU5.0.aSep08		16435	453	3	150	ankyrin 2 (ZU5.0) alternative variant aSep08, mRNA.
ZU5.0	ZU5.0.bSep08		5720	319	1	31	ankyrin 2 like (ZU5.0) alternative variant bSep08, mRNA.
zubor	zubor.aSep08		1841	1717	3	303	putative protein (zubor) alternative variant aSep08, mRNA.
zubor	zubor.bSep08		1055	930	2	56	putative protein (6.1 kD) (zubor) alternative variant bSep08, mRNA.
zuby	zuby.aSep08		963	502		167	host cell factor (zuby) mRNA.
zuchy	zuchy.aSep08		2447	644		40	putative protein (4.3 kD) (zuchy) mRNA.
zudar	zudar.aSep08		3844	477		44	putative protein (4.8 kD) (zudar) mRNA.
zudoy	zudoy.aSep08		6760	303		43	putative protein (zudoy) mRNA.
zuflu	zuflu.aSep08		3932	334	1	111	putative protein (zuflu) alternative variant aSep08, mRNA.
zuflu	zuflu.bSep08		3082	311	1	54	putative protein (zuflu) alternative variant bSep08, mRNA.
zufly	zufly.aSep08		5759	282		76	putative protein (zufly) mRNA.
Zufsp	Zufsp.bSep08	294390	8548	1105	6	249	zinc finger with peptidase domain (28.2 kD) (Zufsp) alternative variant bSep08, complete mRNA.
Zufsp	Zufsp.cSep08	294390	10497	707	4	190	zinc finger with peptidase domain (Zufsp) alternative variant cSep08, mRNA.
Zufsp	Zufsp.dSep08	294390	5635	679	3	185	zinc finger with peptidase domain (Zufsp) alternative variant dSep08, mRNA.
Zufsp	Zufsp.eSep08	294390	10346	765	5	123	zinc finger with peptidase domain (Zufsp) alternative variant eSep08, mRNA.
Zufsp	Zufsp.fSep08	294390	6480	756	3	62	zinc finger with peptidase domain (Zufsp) alternative variant fSep08, mRNA.
zugar	zugar.aSep08		853	493		31	putative protein (zugar) mRNA.

zuja	zuja.aSep08		9918	1491	5	276	zinc finger (zuja) alternative variant aSep08, mRNA.
zujey	zujey.aSep08		15935	2441		180	tbc1 domain family member 1 (20.5 kD) (zujey) mRNA.
zukee	zukee.aSep08		16046	506		30	putative protein (3.6 kD) (zukee) mRNA.
zukler	zukler.aSep08		17806	1638	3	211	putative mitochondrial protein, with a coiled coil domain, of mammalian origin (23.2 kD) (zukler) alternative variant aSep08, mRNA.
zukler	zukler.bSep08		7890	395	2	128	putative protein (zukler) alternative variant bSep08, mRNA.
zulo	zulo.aSep08		2916	1613		105	IQ motif containing e (10.9 kD) (zulo) mRNA.
zumee	zumee.aSep08		1372	415		78	myosin XVA (zumee) mRNA.
zunoy	zunoy.aSep08		5166	458	1	40	putative protein (4.3 kD) (zunoy) alternative variant aSep08, mRNA.
zunoy	zunoy.bSep08		3514	286	1	45	putative protein (zunoy) alternative variant bSep08, mRNA.
zupor	zupor.aSep08		7016	498		124	putative protein (zupor) mRNA.
zusa	zusa.aSep08		116718	767		35	putative protein (zusa) mRNA.
zushee	zushee.aSep08		7921	411	5	80	domain-containing protein (zushee) alternative variant aSep08, mRNA.
zushee	zushee.bSep08		16362	392	4	73	domain-containing protein like (zushee) alternative variant bSep08, mRNA.
zushee	zushee.cSep08		16321	548	6	64	putative protein (zushee) alternative variant cSep08, mRNA.
zutu	zutu.aSep08		6977	602		169	laminin (zutu) mRNA.
zuvar	zuvar.bSep08		689	482	2	86	putative nuclear protein (10.0 kD) (zuvar) alternative variant bSep08, mRNA.
zuvo	zuvo.aSep08		16203	250		30	putative protein (3.6 kD) (zuvo) mRNA.
zuwey	zuwey.aSep08		111215	709		36	putative protein (4.5 kD) (zuwey) mRNA.
Zw10	Zw10.bSep08	363059	5267	642	5	200	ZW10 homolog (Drosophila), centromere/kinetochore protein (Zw10) alternative variant bSep08, mRNA.
Zwilch	Zwilch.aSep08	691493	15692	1755		266	zwilch, kinetochore associated, homolog (Drosophila) (Zwilch) mRNA.
Zxdc	Zxdc.aSep08	362399	28407	3336	3	828	ZXD family zinc finger C (Zxdc) alternative variant aSep08, mRNA.
Zxdc	Zxdc.bSep08	362399	31289	1782	1	356	ZXD family zinc finger C (Zxdc) alternative variant bSep08, mRNA.
Zxdc	Zxdc.cSep08	362399	18481	1763	3	130	ZXD family zinc finger C (Zxdc) alternative variant cSep08, mRNA.
zybor	zybor.aSep08		1037	463		49	putative protein (5.4 kD) (zybor) mRNA.
zyby	zyby.aSep08		854	347		115	plexin B3 (zyby) mRNA.
zychy	zychy.aSep08		2283	255		84	centrosomal protein 152 (zychy) mRNA.
zydar	zydar.aSep08		4109	419		139	CRA b (zydar) mRNA.
zydoy	zydoy.aSep08		13366	1469	4	165	dystrobrevin alpha (zydoy) alternative variant aSep08, mRNA.
zyflu	zyflu.aSep08		1075	646		41	putative protein (zyflu) mRNA.
zyfly	zyfly.aSep08		22573	817		57	putative protein (6.3 kD) (zyfly) mRNA.
Zyg11a	Zyg11a.aSep08	313482	7625	621		206	protein zyg-11 homolog a (Zyg11a) mRNA.

Zyg11b	Zyg11b.aSep08	362559	35597	1379		412	zyg-11 homolog B (Zyg11b) mRNA.
zygar	zygar.aSep08		3468	698		82	putative protein (zygar) mRNA.
zyja	zyja.aSep08		15468	846	1	281	zinc finger (zyja) alternative variant aSep08, mRNA.
zyja	zyja.bSep08		14882	378	1	125	zinc finger (zyja) alternative variant bSep08, mRNA.
zyjey	zyjey.aSep08		46428	440		41	putative protein (zyjey) mRNA.
zykee	zykee.aSep08		1294	658		53	absent in melanoma 2 like (zykee) mRNA.
zykler	zykler.aSep08		7158	682		82	domain-containing protein 3A like (9.1 kD) (zykler) mRNA.
zylo	zylo.aSep08		4123	983		327	CRA c (zylo) mRNA.
zymee	zymee.aSep08		516	408		34	putative protein (zymee) mRNA.
zynoy	zynoy.aSep08		11943	731		82	putative nuclear protein (9.2 kD) (zynoy) mRNA.
zypor	zypor.aSep08		15871	647		47	putative protein (5.5 kD) (zypor) mRNA.
zysa	zysa.aSep08		1829	383		42	putative protein (zysa) mRNA.
zyshee	zyshee.aSep08		4093	477		54	putative protein (zyshee) mRNA.
zytu	zytu.aSep08		26516	482		84	putative cytoplasmic protein (9.7 kD) (zytu) mRNA.
zyvar	zyvar.aSep08		70801	620		109	calmodulin binding transcription activator 1 like (zyvar) mRNA.
zyvo	zyvo.aSep08		13445	524	2	36	putative protein (4.1 kD) (zyvo) alternative variant aSep08, mRNA.
zyvo	zyvo.bSep08		5075	426	2	36	putative protein (4.1 kD) (zyvo) alternative variant bSep08, mRNA.
zywey	zywey.aSep08		1515	512		52	putative protein of mammalian origin (6.2 kD) (zywey) mRNA.
Zyx	Zyx.aSep08	114636	9018	2721	10	682	zyxin (Zyx) alternative variant aSep08, mRNA.
Zyx	Zyx.bSep08	114636	1933	1077	5	358	zyxin (Zyx) alternative variant bSep08, mRNA.
Zyx	Zyx.cSep08	114636	5203	739	3	246	zyxin (Zyx) alternative variant cSep08, mRNA.
Zyx	Zyx.dSep08	114636	1570	850	4	218	zyxin (Zyx) alternative variant dSep08, mRNA.
Zyx	Zyx.eSep08	114636	1705	601	3	146	zyxin (Zyx) alternative variant eSep08, mRNA.
Zyx	Zyx.fSep08	114636	1586	1363	3	132	zyxin (14.7 kD) (Zyx) alternative variant fSep08, mRNA.
Zzef1	Zzef1.aSep08	287476	35940	654		217	zinc finger, ZZ-type with EF hand domain 1 (Zzef1) mRNA.
_Pcdhg	_Pcdhg.pSep08	116782	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	252895	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	252897	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	291635	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	291637	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	364843	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	364844	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.

_Pcdhg	_Pcdhg.pSep08	364845	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	498846	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	498849	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	498850	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	502156	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	553129	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.pSep08	680470	133290	633	3	210	protocadherin gamma subfamily A (_Pcdhg) alternative variant pSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	116782	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	252895	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	252897	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	291635	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	291637	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	364843	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	364844	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	364845	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	498846	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	498849	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	498850	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	502156	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	553129	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_Pcdhg	_Pcdhg.qSep08	680470	6721	2077	2	105	protocadherin gamma subfamily A 6 (_Pcdhg) alternative variant qSep08, mRNA.
_RT1	_RT1.aSep08	24737	4087	2150	8	374	class I MHC (42.0 kD) (_RT1) alternative variant aSep08, mRNA.
_RT1	_RT1.aSep08	24973	4087	2150	8	374	class I MHC (42.0 kD) (_RT1) alternative variant aSep08, mRNA.

_RT1	_RT1.aSep08	24974	4087	2150	8	374	class I MHC (42.0 kD) (_RT1) alternative variant aSep08, mRNA.
_RT1	_RT1.aSep08	309627	4087	2150	8	374	class I MHC (42.0 kD) (_RT1) alternative variant aSep08, mRNA.
_RT1	_RT1.aSep08	415073	4087	2150	8	374	class I MHC (42.0 kD) (_RT1) alternative variant aSep08, mRNA.
_RT1	_RT1.dSep08	24737	2977	1537	4	318	class I (_RT1) alternative variant dSep08, mRNA.
_RT1	_RT1.dSep08	24973	2977	1537	4	318	class I (_RT1) alternative variant dSep08, mRNA.
_RT1	_RT1.dSep08	24974	2977	1537	4	318	class I (_RT1) alternative variant dSep08, mRNA.
_RT1	_RT1.dSep08	309627	2977	1537	4	318	class I (_RT1) alternative variant dSep08, mRNA.
_RT1	_RT1.dSep08	415073	2977	1537	4	318	class I (_RT1) alternative variant dSep08, mRNA.
_RT1	_RT1.eSep08	24737	3019	1450	6	248	mature of major histocompatibility complex class I antigen like (27.5 kD) (_RT1) alternative variant eSep08, mRNA.
_RT1	_RT1.eSep08	24973	3019	1450	6	248	mature of major histocompatibility complex class I antigen like (27.5 kD) (_RT1) alternative variant eSep08, mRNA.
_RT1	_RT1.eSep08	24974	3019	1450	6	248	mature of major histocompatibility complex class I antigen like (27.5 kD) (_RT1) alternative variant eSep08, mRNA.
_RT1	_RT1.eSep08	309627	3019	1450	6	248	mature of major histocompatibility complex class I antigen like (27.5 kD) (_RT1) alternative variant eSep08, mRNA.
_RT1	_RT1.eSep08	415073	3019	1450	6	248	mature of major histocompatibility complex class I antigen like (27.5 kD) (_RT1) alternative variant eSep08, mRNA.
_RT1	_RT1.fSep08	24737	38473	1749	8	247	MHC class I (27.5 kD) (_RT1) alternative variant fSep08, complete mRNA.
_RT1	_RT1.fSep08	24973	38473	1749	8	247	MHC class I (27.5 kD) (_RT1) alternative variant fSep08, complete mRNA.
_RT1	_RT1.fSep08	24974	38473	1749	8	247	MHC class I (27.5 kD) (_RT1) alternative variant fSep08, complete mRNA.
_RT1	_RT1.fSep08	309627	38473	1749	8	247	MHC class I (27.5 kD) (_RT1) alternative variant fSep08, complete mRNA.
_RT1	_RT1.fSep08	415073	38473	1749	8	247	MHC class I (27.5 kD) (_RT1) alternative variant fSep08, complete mRNA.
_RT1	_RT1.gSep08	24737	2795	1196	5	244	class I (_RT1) alternative variant gSep08, mRNA.
_RT1	_RT1.gSep08	24973	2795	1196	5	244	class I (_RT1) alternative variant gSep08, mRNA.
_RT1	_RT1.gSep08	24974	2795	1196	5	244	class I (_RT1) alternative variant gSep08, mRNA.
_RT1	_RT1.gSep08	309627	2795	1196	5	244	class I (_RT1) alternative variant gSep08, mRNA.
_RT1	_RT1.gSep08	415073	2795	1196	5	244	class I (_RT1) alternative variant gSep08, mRNA.
_RT1	_RT1.hSep08	24737	1942	879	3	240	class I antigen like (_RT1) alternative variant hSep08, mRNA.
_RT1	_RT1.hSep08	24973	1942	879	3	240	class I antigen like (_RT1) alternative variant hSep08, mRNA.
_RT1	_RT1.hSep08	24974	1942	879	3	240	class I antigen like (_RT1) alternative variant hSep08, mRNA.
_RT1	_RT1.hSep08	309627	1942	879	3	240	class I antigen like (_RT1) alternative variant hSep08, mRNA.
_RT1	_RT1.hSep08	415073	1942	879	3	240	class I antigen like (_RT1) alternative variant hSep08, mRNA.

_RT1	_RT1.jSep08	24737	1374	754	3	163	class I (_RT1) alternative variant jSep08, mRNA.
_RT1	_RT1.jSep08	24973	1374	754	3	163	class I (_RT1) alternative variant jSep08, mRNA.
_RT1	_RT1.jSep08	24974	1374	754	3	163	class I (_RT1) alternative variant jSep08, mRNA.
_RT1	_RT1.jSep08	309627	1374	754	3	163	class I (_RT1) alternative variant jSep08, mRNA.
_RT1	_RT1.jSep08	415073	1374	754	3	163	class I (_RT1) alternative variant jSep08, mRNA.
_RT1	_RT1.mSep08	24737	616	522	2	146	MHC class I (_RT1) alternative variant mSep08, mRNA.
_RT1	_RT1.mSep08	24973	616	522	2	146	MHC class I (_RT1) alternative variant mSep08, mRNA.
_RT1	_RT1.mSep08	24974	616	522	2	146	MHC class I (_RT1) alternative variant mSep08, mRNA.
_RT1	_RT1.mSep08	309627	616	522	2	146	MHC class I (_RT1) alternative variant mSep08, mRNA.
_RT1	_RT1.mSep08	415073	616	522	2	146	MHC class I (_RT1) alternative variant mSep08, mRNA.
_RT1	_RT1.nSep08	24737	23896	338	3	112	class I (_RT1) alternative variant nSep08, mRNA.
_RT1	_RT1.nSep08	24973	23896	338	3	112	class I (_RT1) alternative variant nSep08, mRNA.
_RT1	_RT1.nSep08	24974	23896	338	3	112	class I (_RT1) alternative variant nSep08, mRNA.
_RT1	_RT1.nSep08	309627	23896	338	3	112	class I (_RT1) alternative variant nSep08, mRNA.
_RT1	_RT1.nSep08	415073	23896	338	3	112	class I (_RT1) alternative variant nSep08, mRNA.
_RT1	_RT1.oSep08	24737	1436	788	4	108	class I MHC heavy chain antigen like (_RT1) alternative variant oSep08, mRNA.
_RT1	_RT1.oSep08	24973	1436	788	4	108	class I MHC heavy chain antigen like (_RT1) alternative variant oSep08, mRNA.
_RT1	_RT1.oSep08	24974	1436	788	4	108	class I MHC heavy chain antigen like (_RT1) alternative variant oSep08, mRNA.
_RT1	_RT1.oSep08	309627	1436	788	4	108	class I MHC heavy chain antigen like (_RT1) alternative variant oSep08, mRNA.
_RT1	_RT1.oSep08	415073	1436	788	4	108	class I MHC heavy chain antigen like (_RT1) alternative variant oSep08, mRNA.
_RT1	_RT1.pSep08	24737	36320	781	5	106	class I MHC heavy chain antigen like (_RT1) alternative variant pSep08, mRNA.
_RT1	_RT1.pSep08	24973	36320	781	5	106	class I MHC heavy chain antigen like (_RT1) alternative variant pSep08, mRNA.
_RT1	_RT1.pSep08	24974	36320	781	5	106	class I MHC heavy chain antigen like (_RT1) alternative variant pSep08, mRNA.
_RT1	_RT1.pSep08	309627	36320	781	5	106	class I MHC heavy chain antigen like (_RT1) alternative variant pSep08, mRNA.
_RT1	_RT1.pSep08	415073	36320	781	5	106	class I MHC heavy chain antigen like (_RT1) alternative variant pSep08, mRNA.
_RT1	_RT1.qSep08	24737	644	512	2	26	putative protein (_RT1) alternative variant qSep08, mRNA.
_RT1	_RT1.qSep08	24973	644	512	2	26	putative protein (_RT1) alternative variant qSep08, mRNA.
_RT1	_RT1.qSep08	24974	644	512	2	26	putative protein (_RT1) alternative variant qSep08, mRNA.
_RT1	_RT1.qSep08	309627	644	512	2	26	putative protein (_RT1) alternative variant qSep08, mRNA.
_RT1	_RT1.qSep08	415073	644	512	2	26	putative protein (_RT1) alternative variant qSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	24861	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	113992	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.



_Ugt1a	_Ugt1a.jSep08	154516	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	301595	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	396527	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	396551	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	396552	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.jSep08	574523	18021	622	4	150	UDP glycosyltransferase 1 family polypeptide (_Ugt1a) alternative variant jSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	24861	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	113992	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	154516	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	301595	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	396527	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	396551	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	396552	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.kSep08	574523	1701	505	2	137	UDP-glucuronosyltransferase (_Ugt1a) alternative variant kSep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	24861	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	113992	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	154516	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	301595	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	396527	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	396551	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	396552	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.
_Ugt1a	_Ugt1a.ISep08	574523	125464	293	3	97	UGT1A7 (_Ugt1a) alternative variant ISep08, mRNA.