

Accession	Gene Name	MW(kDa)	Ccr7-/-				Sib3ialV-/-				Nrp-2-/-			
			Coverage	# Peptides	# PSMs	Area	Coverage	# Peptides	# PSMs	Area	Coverage	# Peptides	# PSMs	Area
gil5915682 splP07724.3 ALBU_MOUSE	Alb	68.6	7.07	3	70	6.04E+10	8.06	4	65	6.63E+10	8.06	4	57	4.15E+10
gil122016 splP10853.2 H2B1F_MOUSE	Hist1h2bn	13.9	46.83	7	42	2.56E+09	46.83	7	40	2.25E+09	53.97	9	67	8.33E+09
gil18171239 splQ8CGP6.3 H2A1H_MOUSE	Hist1h2ah	13.9	35.94	4	18	2.36E+09	27.34	3	14	1.85E+09	35.94	4	18	4.75E+09
gil121905 splP15864.2 H12_MOUSE	Hist1h1c	21.3	33.02	9	31	3.14E+09	36.32	10	30	3.99E+09	36.32	10	33	3.31E+09
gil21903465 splP47963.3 RL13_MOUSE	Rpl13	24.3	49.76	11	55	3.38E+09	50.71	12	87	6.31E+09	50.24	15	72	3.25E+09
gil1170151 splP43274.2 H14_MOUSE	Hist1h1e	22.0	19.63	7	32	2.60E+09	30.14	9	28	3.08E+09	19.63	7	32	2.96E+09
gil126302559 splP04104.4 K2C1_MOUSE	Krt1	65.6	6.44	6	24	1.90E+09	6.44	6	22	3.22E+09	6.44	7	28	2.50E+09
gil22001904 splQ9CPR4.3 RL17_MOUSE	Rpl17	21.4	50.54	10	57	2.12E+09	50.54	10	66	3.44E+09	50.54	10	58	2.18E+09
gil66773987 splQ6IFZ6.1 K2C1B_MOUSE	Krt77	61.3	7.52	7	32	1.61E+09	7.52	7	27	2.79E+09	7.52	7	31	2.09E+09
gil306526288 splP14115.5 RL27A_MOUSE	Rpl27a	16.6	37.84	7	32	2.26E+09	48.65	9	36	3.81E+09	31.76	6	27	1.97E+09
gil21264500 splP35980.3 RL18_MOUSE	Rpl18	21.6	44.15	12	62	3.75E+09	45.21	10	56	6.63E+09	45.21	10	45	1.85E+09
gil1892069 splQ6NXH9.1 K2C73_MOUSE	Krt73	58.9	10.58	8	21	1.48E+09	10.58	8	19	2.54E+09	10.58	8	19	1.79E+09
gil1170668 splQ922U2.1 K2C5_MOUSE	Krt5	61.7	25.86	23	57	1.05E+09	31.21	26	62	1.72E+09	27.76	25	68	1.61E+09
gil60416436 splP50446.3 K2C6A_MOUSE	Krt6a	59.3	0	0	0	0.00E+00		0	0	0.00E+00	30.38	26	55	1.59E+09
gil135070 splP47915.2 RL29_MOUSE	Rpl29	17.6	25.00	5	27	1.73E+09	27.50	6	27	3.16E+09	25.00	5	21	1.58E+09
gil116242600 splP02535.3 K1C10_MOUSE	Krt10	57.7	13.51	10	60	1.46E+09	13.51	10	65	3.31E+09	13.51	10	64	1.57E+09
gil123791931 splQ3UV17.1 K22O_MOUSE	Krt76	62.8	9.26	8	19	9.21E+08	9.26	8	21	1.45E+09	10.77	9	23	1.43E+09
gil133015 splP12970.2 RL7A_MOUSE	Rpl7a	30.0	46.62	18	74	1.10E+09	46.24	17	67	1.27E+09	50.00	19	73	1.88E+09
gil23396626 splQ9QWL7.3 K1C17_MOUSE	Krt17	48.1	19.63	11	24	8.24E+08		0	0	0.00E+00	44.34	21	47	1.37E+09
gil205371802 splQ8VDY5.4 MYH9_MOUSE	Mvh9	226.2	10.00	17	20	7.23E+07	67.30	171	1426	1.35E+10	60.97	137	494	1.32E+09
gil302393751 splP62983.2 R27A_MOUSE	Rps27a	17.9	48.72	7	44	1.54E+09	54.49	9	52	1.84E+09	41.03	6	37	1.13E+09
gil13878273 splQ9D1R9.2 RL34_MOUSE	Rpl34	13.3	36.75	6	28	1.67E+09	37.61	7	32	2.15E+09	37.61	7	24	1.10E+09
gil1170153 splP43275.2 H11_MOUSE	Hist1h1a	21.8	0	0	0	0.00E+00	22.54	5	11	1.25E+09	23.00	5	11	1.05E+09
gil1170154 splP43276.2 H15_MOUSE	Hist1h1b	22.6	26.46	8	18	6.74E+08	29.60	9	20	8.00E+08	27.35	9	24	1.05E+09
gil1891691 splQ6IFX2.1 K1C42_MOUSE	Krt42	50.1	18.81	9	20	3.34E+08	18.14	9	17	8.54E+08	28.98	14	42	1.03E+09
gil33302628 splQ61781.2 K1C14_MOUSE	Krt14	52.8	19.42	10	47	6.28E+08	20.25	10	45	1.40E+09	38.43	19	84	9.88E+08
gil62296680 splP47911.3 RL6_MOUSE	Rpl6	33.5	51.01	18	66	1.34E+09	45.27	17	54	1.15E+09	51.01	18	59	9.21E+08
gil23396632 splQ9Z2K1.3 K1C16_MOUSE	Krt16	51.6	13.43	7	40	6.05E+08		0	0	0.00E+00	21.54	12	77	9.13E+08
gil138536 splP20152.3 VIME_MOUSE	Vim	53.7	61.37	29	94	4.10E+08	77.04	36	150	1.60E+09	68.67	33	137	8.99E+08
gil1879424 splQ8VEK3.1 HNRPU_MOUSE	Hnrpu	87.9	28.13	26	92	8.69E+08	31.62	26	83	1.27E+09	30.88	25	83	8.34E+08
gil2500367 splQ09167.3 RL21_MOUSE	Rpl21	18.6	45.00	8	29	4.79E+08	50.63	10	44	1.34E+09	46.25	10	33	7.84E+08
gil730549 splP41105.2 RL28_MOUSE	Rpl28	15.7	59.85	14	55	1.17E+09	59.12	13	66	2.86E+09	54.74	10	38	7.71E+08
gil22001911 splQ9D8E6.3 RL4_MOUSE	Rpl4	47.1	44.39	23	84	7.58E+08	44.15	23	86	1.30E+09	43.20	20	63	7.66E+08
gil1350769 splP14148.2 RL7_MOUSE	Rpl7	31.4	48.15	24	80	9.88E+08	55.19	27	80	1.33E+09	47.78	24	76	7.26E+08
gil123796763 splQ3TTY5.1 K22E_MOUSE	Krt2	70.9	10.04	11	36	7.32E+08	10.04	11	39	1.45E+09	10.18	12	39	7.24E+08
gil1350733 splP47964.2 RL36_MOUSE	Rpl36	12.2	31.43	5	23	6.96E+08	30.48	4	23	1.37E+09	23.81	3	17	7.18E+08
gil136478 splP19253.4 RL13A_MOUSE	Rpl13a	23.4	47.29	14	45	8.48E+08	43.35	13	49	1.25E+09	41.38	12	40	7.09E+08
gil52783239 splQ9CR57.3 RL14_MOUSE	Rpl14	23.5	31.80	8	30	6.57E+08	37.33	10	38	1.28E+09	34.56	9	30	7.04E+08
gil48428680 splQ9D823.3 RL37_MOUSE	Rpl37	11.1	15.46	4	5	6.04E+08	21.65	5	17	2.34E+09	19.59	4	6	6.32E+08
gil81916088 splQ91VM5.1 IRMXL1_MOUSE	Rbmxl1	42.1	38.14	15	49	3.09E+08	44.07	18	74	8.82E+08	38.14	15	50	3.38E+08
gil81917893 splQ9VW02.1 IRBMX_MOUSE	RbmX	42.3	34.78	15	44	3.09E+08	38.62	17	61	8.82E+08	37.60	15	47	3.38E+08
gil391358189 splP02301.3 H3C_MOUSE	Gm7426	15.3	25.74	5	15	2.64E+08	30.15	6	18	3.00E+08	25.74	5	13	3.27E+08
gil3122836 splP97351.3 IRS3A_MOUSE	Rps3a1	29.9	53.79	15	55	2.22E+08	53.79	16	70	3.30E+08	61.36	18	69	3.24E+08
gil341942150 splP27659.3 RL3_MOUSE	Rpl3	46.1	48.88	24	73	3.66E+08	49.88	22	80	5.53E+08	50.12	22	66	3.11E+08
gil519668679 splP21107.3 TPM3_MOUSE	Tpm3	33.0	11.58	3	4	1.71E+07	37.54	15	33	8.40E+08	34.74	15	29	2.82E+08
gil54036077 splQ9CY57.2 CHTOP_MOUSE	Chtop	26.6	32.53	9	51	4.32E+08	34.54	9	52	5.20E+08	32.13	8	46	2.10E+08
gil120702 splP16858.2 IG3P_MOUSE	Gapdh	35.8	42.94	10	32	2.46E+08	45.65	10	30	3.66E+08	42.94	10	29	1.87E+08
gil22002065 splP47962.3 RL5_MOUSE	Rpl5	34.4	31.99	11	32	1.06E+08	35.35	13	34	1.40E+08	31.99	12	35	1.72E+08
gil1880390 splQ9CQN7.1 IRM41_MOUSE	Mrp41	15.3	57.04	7	18	1.76E+08	34.07	4	5	3.83E+07	62.22	8	18	1.50E+08
gil3122833 splP97461.3 IRS5_MOUSE	Rps5	22.9	27.45	5	10	6.32E+07	27.45	6	29	2.47E+08	35.29	9	22	1.50E+08
gil1703188 splP51881.3 ADT2_MOUSE	Sic25a5	32.9	16.11	5	11	4.36E+07	28.19	9	15	2.08E+08	23.49	8	14	1.42E+08
gil341940836 splQ9WTL4.2 INSRR_MOUSE	Insr	144.8	0	0	0	0.00E+00		0	0	0.00E+00	0.62	2	2	1.31E+08
gil341940334 splP47774.2 CCR7_MOUSE	Ccr7	42.8	0	0	0	0.00E+00		0	0	0.00E+00	8.47	3	6	1.16E+08
gil45477334 splQ9ERU3.2 ZNF22_MOUSE	Zfp422	27.3	33.76	8	20	6.12E+07	45.99	13	29	1.47E+08	37.13	9	23	1.11E+08
gil22001919 splQ9JH8.3 RL38_MOUSE	Rpl38	8.2	32.86	4	5	5.71E+07	32.86	4	9	1.40E+08	32.86	4	6	1.07E+08
gil2506545 splP20029.3 GRP78_MOUSE	Hspa5	72.4	41.53	24	44	1.02E+08	43.51	27	45	1.95E+08	42.75	24	38	9.73E+07
gil21903459 splP51410.2 RL9_MOUSE	Rpl9	21.9	22.92	4	7	2.22E+07	22.92	4	12	8.79E+07	29.17	5	14	9.34E+07
gil547923 splP35700.1 PRDX1_MOUSE	Prdx1	22.2	21.11	4	9	9.85E+07	45.73	10	16	1.29E+08	41.21	9	14	8.62E+07
gil54036480 splQ6IRU2.3 TPM4_MOUSE	Tpm4	28.5	0	0	0	0.00E+00	49.60	17	36	2.10E+08	52.02	18	32	8.43E+07
gil116849 splP18760.3 COF1_MOUSE	Cof1	18.5	52.41	7	12	1.42E+08	42.17	7	11	1.86E+08	27.11	3	7	8.37E+07
gil20139648 splQ9N9N1.1 IRM34_MOUSE	Mrp34	10.5	0	0	0	0.00E+00		0	0	0.00E+00	23.91	2	4	6.70E+07
gil341940436 splQ61656.2 DDX5_MOUSE	Ddx5	69.2	9.93	7	9	4.06E+07	25.57	16	22	1.13E+08	19.38	13	17	6.69E+07
gil416677 splQ03265.1 ATPA_MOUSE	Atpa5a1	59.7	13.56	6	8	2.52E+07	24.59	11	18	9.35E+07	26.22	12	18	6.66E+07
gil23396881 splQ9JHJ0.1 TMOD3_MOUSE	Tmod3	39.5	0	0	0	0.00E+00	48.01	20	53	9.61E+08	34.66	12	16	6.54E+07

gil146345448 splP52480.4IKPYM_MOUSE	Pkm	57.8	35.78	16	27	1,17E+08	34.09	16	24	1,70E+08	40.11	17	22	6.25E+07
gil83305636 splQ8K2Y7.2IRM47_MOUSE	Mrp147	29.7	28.97	8	19	3.93E+07	11.51	3	3	6.91E+06	25.00	7	17	6.23E+07
gil47605546 splQ9DB20.1IATPO_MOUSE	Atp5a	23.3	23.00	4	9	4.56E+07	15.02	3	8	9.56E+07	31.92	6	10	6.20E+07
gil158518540 splQ04750.2ITOP1_MOUSE	Top1	90.8	4.95	3	4	4.95E+06	5.74	4	5	8.01E+06	28.03	20	32	6.15E+07
gil1363776 splP17182.3IENOA_MOUSE	Eno1	47.1	45.39	15	29	9.51E+07	47.24	16	28	1.58E+08	41.24	14	24	6.12E+07
gil28381362 splP13020.3IGELS_MOUSE	Gsn	85.9	3.08	2	2	3.29E+06	47.56	31	127	2.07E+09	19.87	9	12	5.68E+07
gil145566796 splQ3THE2.2IML12B_MOUSE	Myl12b	19.8		0	0	0.00E+00	62.21	8	31	5.47E+08	40.12	6	9	5.66E+07
gil32130449 splP35979.2IRL12_MOUSE	Rpl12	17.8	63.64	8	11	4.74E+07	53.94	7	15	9.01E+07	53.94	7	10	5.36E+07
gil399173 splP31725.3IS10A9_MOUSE	S100a9	13.0	38.94	5	10	8.82E+07	31.86	3	7	6.30E+07	28.32	4	7	5.11E+07
gil24212382 splQ9D2R8.1IRT33_MOUSE	Mmps33	12.5	33.96	4	9	4.23E+07	33.96	4	9	6.10E+07	33.96	4	8	4.83E+07
gil73920210 splP35550.2IFBRL_MOUSE	Fbl	34.3	25.99	9	12	3.15E+07	26.91	8	13	5.49E+07	22.63	8	13	4.79E+07
gil18202241 splQ88593.1IPGRP1_MOUSE	Pglyrp1	20.5		0	0	0.00E+00		0	0	0.00E+00	17.58	2	2	4.73E+07
gil126048 splP06151.3ILDHA_MOUSE	Ldha	36.5	24.40	7	11	6.58E+07	12.35	4	5	7.04E+07	21.39	7	9	4.33E+07
gil38258939 splQ64339.4ISG15_MOUSE	Isg15	17.9	16.77	2	3	2.19E+07	40.37	4	8	2.20E+08	16.77	2	4	4.30E+07
gil18203578 splQ9VVK4.1EHD1_MOUSE	Ehd1	60.6		0	0	0.00E+00		0	0	0.00E+00	32.96	13	14	4.25E+07
gil1346247 splP10922.4IH10_MOUSE	H110	20.8	15.98	3	8	8.79E+07	20.62	4	6	1.20E+08	15.98	3	5	4.16E+07
gil17376706 splQ61543.1IGSLG1_MOUSE	Glg1	133.6	14.89	18	28	3.25E+07		0	0	0.00E+00	16.17	20	31	4.09E+07
gil1709972 splP53026.3IRL10A_MOUSE	Rpl10a	24.9	19.82	5	8	2.52E+07	9.68	3	5	4.25E+07	19.82	6	8	4.03E+07
gil729000 splQ99020.1IROAA_MOUSE	Hnmpab	30.8	15.44	4	8	3.94E+07	18.95	5	9	5.77E+07	18.95	5	8	3.74E+07
gil81904744 splQ9D116.1IRM14_MOUSE	Mrp14	15.9	24.14	3	6	1.77E+07	38.62	5	8	4.05E+07	42.07	5	7	3.70E+07
gil1346470 splP19973.2ILSP1_MOUSE	Lsp1	36.7		0	0	0.00E+00	59.09	16	51	9.20E+08	28.48	7	11	3.43E+07
gil126599 splP17897.1LYZ1_MOUSE	Lyz1	16.8	12.16	2	4	5.70E+07	12.16	2	3	3.01E+07	12.16	2	3	3.41E+07
gil81871888 splQ5SUAS.1IMYO1G_MOUSE	Myo1g	117.2		0	0	0.00E+00	55.18	50	175	1.21E+09	17.97	15	22	3.39E+07
gil29840809 splQ61233.4IPLSL_MOUSE	Lcp1	70.1	16.11	8	9	2.43E+07	36.04	18	25	1.06E+08	18.02	10	11	3.19E+07
gil123783231 splQ3TEA8.1IHP1B3_MOUSE	Hp1bp3	60.8	5.23	3	5	7.25E+06	18.05	10	16	3.54E+07	15.16	8	14	3.17E+07
gil30316201 splQ8BG05.1IROA3_MOUSE	Hnmpa3	39.6	16.89	5	7	1.97E+07	20.84	6	10	5.72E+07	24.27	7	11	3.09E+07
gil81916834 splQ9CX86.1IROA0_MOUSE	Hnmpa0	30.5	8.85	2	3	1.28E+07	18.69	5	9	6.32E+07	21.31	5	8	3.04E+07
gil81917122 splQ9D945.1ILLPH_MOUSE	Llph	15.4	29.23	3	5	1.96E+07	49.23	6	14	8.23E+07	43.85	5	10	2.99E+07
gil60390645 splQ9DCN2.3INB5B3_MOUSE	Cyb5r3	34.1	21.59	7	11	2.84E+07	18.60	6	12	6.97E+07	27.91	8	11	2.84E+07
gil32363402 splQ9DCJ5.3INDUA8_MOUSE	Ndufa8	20.0	35.47	5	11	3.10E+07	36.63	5	12	5.63E+07	43.02	6	9	2.58E+07
gil81869559 splQ9CQZ8.3IH2AY_MOUSE	H2afy	39.7		0	0	0.00E+00		0	0	0.00E+00	7.80	2	3	2.57E+07
gil1168737 splQ09200.1IB4GN1_MOUSE	B4gaint1	59.2		0	0	0.00E+00		0	0	0.00E+00	16.14	9	10	2.50E+07
gil76364159 splQ78IK4.1IAP00L_MOUSE	Apool	29.2		0	0	0.00E+00	9.06	2	3	2.54E+07	24.53	5	5	2.48E+07
gil1345696 splP47754.3ICAZA2_MOUSE	Capza2	32.9		0	0	0.00E+00	65.38	14	26	5.30E+08	15.38	3	4	2.46E+07
gil34925411 splQ9CR47.1INSA2_MOUSE	Nsa2	30.0		0	0	0.00E+00	12.31	3	4	2.00E+07	17.31	5	6	2.40E+07
gil118105 splP17742.2PIPIA_MOUSE	Ppia	18.0	10.98	2	3	3.51E+07	21.95	4	4	5.30E+07	10.98	2	3	2.22E+07
gil81914411 splQ8CH25.1ISLTM_MOUSE	Sltm	116.8	8.92	8	13	1.65E+07	9.70	9	12	2.42E+07	11.35	10	16	2.19E+07
gil115602 splP07091.1IS10A4_MOUSE	S100a4	11.7		0	0	0.00E+00	16.83	2	2	2.70E+07	16.83	2	3	2.14E+07
gil1705635 splP28293.2ICATG_MOUSE	Ctsq	29.1	9.58	2	2	2.81E+06		0	0	0.00E+00	16.86	4	7	2.08E+07
gil417489 splP23506.3PIIMT_MOUSE	Pcmt1	24.6	28.63	5	9	2.84E+07	38.77	5	11	6.62E+07	23.79	4	7	2.07E+07
gil11705560 splP54751.1ISIA4_MOUSE	St3gal1	39.0		0	0	0.00E+00		0	0	0.00E+00	16.32	5	5	2.05E+07
gil341941148 splP08752.5IGNAI2_MOUSE	Gnai2	40.5		0	0	0.00E+00		0	0	0.00E+00	4.79	2	3	2.03E+07
gil81916767 splQ9CQP0.1IRM33_MOUSE	Mrp133	7.4		0	0	0.00E+00		0	0	0.00E+00	41.54	3	4	2.01E+07
gil341940350 splQ9LL3.2ICHSTC_MOUSE	Chst12	49.4		0	0	0.00E+00		0	0	0.00E+00	7.40	2	2	1.97E+07
gil118573686 splQ8VDT9.2IRM50_MOUSE	Mrp150	18.2	49.69	6	8	8.85E+07		0	0	0.00E+00	25.79	3	4	1.95E+07
gil338818072 splE90557.1IDESP_MOUSE	Dsp	332.7	2.57	7	7	5.72E+06	2.53	7	7	4.67E+06	6.00	17	17	1.89E+07
gil41688745 splQ08784.1ITCOF_MOUSE	Tcof1	134.9	6.14	6	16	7.02E+07	3.86	4	6	1.74E+07	4.70	5	10	1.87E+07
gil408360268 splP38647.3IGRP75_MOUSE	Hspa9	73.4	10.31	5	7	1.13E+07	21.21	12	17	2.16E+07	20.77	11	13	1.86E+07
gil113951 splP07356.2ANXA2_MOUSE	Anxa2	38.7	10.32	3	4	4.05E+07	21.53	7	8	5.99E+07	10.32	3	4	1.81E+07
gil83305343 splQ02257.3IPLAK_MOUSE	Jup	81.7	5.77	4	4	5.00E+06	11.95	8	8	1.17E+07	13.42	10	11	1.78E+07
gil128843 splP09405.2INUCL_MOUSE	Ncl	76.7	3.68	3	4	1.01E+07	12.16	8	11	3.49E+07	15.42	10	12	1.71E+07
gil550544292 splP47878.2IBP3_MOUSE	Igf1bp3	31.7	6.16	2	6	9.99E+06	8.90	3	8	1.73E+07	8.90	3	8	1.63E+07
gil81880358 splQ9CQ06.1IRM24_MOUSE	Mrp124	24.9	12.50	3	3	1.44E+07		0	0	0.00E+00	25.00	5	5	1.62E+07
gil613504949 splQ9QX31.3IPLC_MOUSE	Plec	533.9		0	0	0.00E+00	2.64	11	14	1.18E+07	0.66	3	6	1.62E+07
gil21263432 splQ91VR2.1IATPG_MOUSE	Atp5c1	32.9		0	0	0.00E+00	14.77	4	6	2.52E+07	11.07	3	5	1.61E+07
gil2500582 splQ61937.1INPM_MOUSE	Npm1	32.5	8.90	3	4	7.99E+06	14.04	5	8	2.06E+07	11.30	4	6	1.60E+07
gil408360135 splQ00P19.2IHNR12_MOUSE	Hnmpu12	84.9	1.74	2	4	8.48E+06		0	0	0.00E+00	8.32	6	7	1.56E+07
gil45477105 splQ81Q05.1INMES1_MOUSE	AA467197	9.6	46.99	4	7	2.65E+07	46.99	4	5	2.46E+07	28.92	3	3	1.52E+07
gil341940433 splQ91VN6.2IDDX41_MOUSE	Ddx41	69.8		0	0	0.00E+00	27.49	15	18	5.79E+07	9.32	6	7	1.49E+07
gil2829839 splP97430.1ISLPI_MOUSE	Slpi	14.3	12.98	2	2	9.21E+06		0	0	0.00E+00	12.98	2	2	1.48E+07
gil21759257 splQ8VEM8.1IMPCC_MOUSE	Slc25a3	39.6		0	0	0.00E+00		0	0	0.00E+00	5.32	2	3	1.47E+07
gil81880345 splQ9CPT5.1INOP16_MOUSE	Nop16	21.1		0	0	0.00E+00	8.43	2	3	1.26E+07	24.16	4	6	1.43E+07
gil46397808 splQ89053.5ICOR1A_MOUSE	Coro1a	51.0	6.94	3	5	1.48E+07	6.94	3	5	1.76E+07	10.85	5	7	1.41E+07
gil114290 splP16460.1IASSY_MOUSE	Ass1	46.6	9.71	4	5	1.39E+07	14.81	9	11	4.39E+07	13.11	6	7	1.40E+07
gil215273926 splP24369.2IPPIB_MOUSE	Ppiib	23.7		0	0	0.00E+00	9.26	2	2	1.51E+07	16.67	4	4	1.37E+07
gil417240 splP32067.1ILA_MOUSE	Ssb	47.7	7.71	3	3	3.02E+06	7.47	3	3	3.26E+06	9.64	4	6	1.34E+07
gil24212379 splQ9N87.1IRT05_MOUSE	Mmps5	48.2	4.86	2	4	1.25E+07		0	0	0.00E+00	8.56	4	6	1.34E+07
gil85695427 splQ9D1B9.3IRM28_MOUSE	Mrp128	30.2	8.56	2	6	1.71E+07		0	0	0.00E+00	12.06	3	5	1.31E+07

gii20139649 spl Q99N92.1 RM27_MOUSE	Mrpl27	15.9		0	0	0,00E+00		0	0	0,00E+00	22,97	3	4	1,25E+07
gii71153525 spl Q08288.2 LYAR_MOUSE	Lyar	43.7		0	0	0,00E+00		0	0	0,00E+00	18,56	6	7	1,25E+07
gii50401030 spl Q8BG81.1 PDI3_MOUSE	Poldip3	46.1	10,00	4	6	6,00E+06	18,57	8	15	2,02E+07	18,57	9	12	1,23E+07
gii124028629 spl Q89569.2 ROA2_MOUSE	Hnrpa2b1	37.4	9,07	3	5	1,19E+07	16,43	5	7	2,69E+07	11,61	4	5	1,21E+07
gii48474242 spl Q08583.3 THOC4_MOUSE	Alvret	26.9	8,24	2	3	1,18E+07	8,24	2	3	1,51E+07	19,22	4	4	1,18E+07
gii218512147 spl Q9WVAA.4 TAGL2_MOUSE	Tagln2	22.4	21,11	4	5	1,41E+07	9,55	2	3	2,56E+07	15,58	3	4	1,17E+07
gii66773932 spl Q91XV3.3 BASP1_MOUSE	Basp1	22.1	21,68	3	3	8,27E+06	27,88	4	4	1,44E+07	27,88	4	5	1,15E+07
gii127294 spl P24668.1 MPRD_MOUSE	M6pr	31.2	11,51	3	3	1,26E+07		0	0	0,00E+00	11,51	3	3	1,14E+07
gii462608 spl P26041.3 MOES_MOUSE	Msn	67.7	14,56	10	11	2,17E+07	10,92	7	7	2,65E+07	9,71	7	7	1,14E+07
gii20455479 spl P56480.2 ATPB_MOUSE	Atp5b	56.3	8,88	3	3	7,46E+06	16,07	6	7	2,83E+08	11,15	4	4	1,13E+07
gii33516945 spl Q60668.2 HNRPD_MOUSE	Hnrmpd	38.3		0	0	0,00E+00	5,07	2	2	1,55E+07	5,07	2	2	1,12E+07
gii32363398 spl Q9CQZ5.1 INDUA6_MOUSE	Ndufa6	15.3	25,19	3	4	2,11E+07		0	0	0,00E+00	19,08	2	2	1,12E+07
gii156633648 spl Q8R2N0.2 TAP2_MOUSE	Ccdc59	28.0		0	0	0,00E+00	9,17	2	2	1,02E+07	9,17	2	2	1,11E+07
gii302425115 spl A2AJ15.1 MA1B1_MOUSE	Man1b1	75.1		0	0	0,00E+00		0	0	0,00E+00	7,14	5	5	1,09E+07
gii134881 spl P16254.1 SRP14_MOUSE	Srp14	12.5		0	0	0,00E+00	23,64	2	2	1,13E+07	31,82	3	3	1,04E+07
gii18187888 spl Q8R1F0.1 L10K_MOUSE	D8Erd738e	10.2		0	0	0,00E+00		0	0	0,00E+00	25,53	3	3	1,01E+07
gii21362934 spl Q9D903.1 EBP2_MOUSE	Ebna1bp2	34.7		0	0	0,00E+00		0	0	0,00E+00	9,15	3	4	1,01E+07
gii408360150 spl Q9JKF1.2 IQGA1_MOUSE	Iqgap1	188.6		0	0	0,00E+00	38,93	54	92	3,20E+08	2,72	3	3	9,71E+06
gii20140237 spl Q8D8Y0.1 EFHD2_MOUSE	Efh2d	26.8		0	0	0,00E+00	67,50	16	58	1,51E+09	10,83	2	2	9,65E+06
gii62510933 spl Q922R8.3 PDIA6_MOUSE	Pdia6	48.1		0	0	0,00E+00		0	0	0,00E+00	6,36	2	2	9,31E+06
gii1881612 spl Q9D8M4.1 R1L7L_MOUSE	Rpl7l1	28.5		0	0	0,00E+00		0	0	0,00E+00	6,91	2	2	9,22E+06
gii113607 spl P05064.2 ALDOA_MOUSE	Aldoa	39.3	32,14	10	13	5,03E+07	23,08	7	9	3,11E+07	16,48	6	7	9,17E+06
gii20140777 spl Q9CQ69.3 QCR8_MOUSE	Uccrq	9.8	29,27	4	5	1,15E+07	29,27	4	10	3,02E+07	45,12	5	7	9,13E+06
gii729023 spl P24452.2 CAPG_MOUSE	Capg	39.2	15,63	5	7	2,25E+07	9,09	3	5	6,34E+07	9,09	3	4	9,02E+06
gii24212196 spl Q99N85.1 IRT18A_MOUSE	Mrps18a	22.3		0	0	0,00E+00		0	0	0,00E+00	12,76	3	3	8,72E+06
gii462064 spl Q05816.3 FABP5_MOUSE	Fabp5	15.1	13,33	2	2	1,56E+07	13,33	2	2	1,82E+07	13,33	2	2	8,54E+06
gii3913473 spl O70503.1 DHB12_MOUSE	Hsd17b12	34.7	4,81	2	3	3,73E+06	9,94	3	6	1,22E+07	9,94	3	6	8,33E+06
gii30913111 spl Q99M87.1 DNJA3_MOUSE	Dnaja3	52.4		0	0	0,00E+00		0	0	0,00E+00	5,00	2	2	8,30E+06
gii1350822 spl P49312.2 ROA1_MOUSE	Hnrpa1	34.2		0	0	0,00E+00	16,88	4	6	1,11E+07	12,81	3	4	8,30E+06
gii113945 spl P10107.2 IANXA1_MOUSE	Anxa1	38.7	5,49	2	2	1,41E+07	5,49	2	3	2,23E+07	5,49	2	2	7,89E+06
gii24636295 spl Q9D8B3.2 CHM4B_MOUSE	Chmp4b	24.9	10,71	2	3	7,49E+06	10,71	2	3	6,10E+06	10,71	2	3	7,86E+06
gii60416372 spl Q61495.2 DSG1A_MOUSE	Dsg1a	114.5	1,99	2	2	9,72E+06		0	0	0,00E+00	1,51	2	2	7,79E+06
gii1917732 spl Q9QYL7.1 ABT1_MOUSE	Abt1	30.6		0	0	0,00E+00		0	0	0,00E+00	7,43	2	2	7,50E+06
gii353678156 spl D3YXK2.2 SAFB1_MOUSE	Safb	105.0	2,35	3	3	4,50E+06	2,35	3	4	1,34E+07	3,09	3	4	7,11E+06
gii60943276 spl P70279.1 SURF6_MOUSE	Surf6	41.2		0	0	0,00E+00		0	0	0,00E+00	10,14	4	4	6,71E+06
gii122239 spl P14483.1 HB2A_MOUSE	H2-Ab1	30.1		0	0	0,00E+00	18,49	4	4	1,80E+07	9,43	2	2	6,67E+06
gii90185245 spl Q69ZN2.1 GNPTA_MOUSE	Gnptab	140.9		0	0	0,00E+00		0	0	0,00E+00	1,38	2	3	6,64E+06
gii110825747 spl P52875.2 TM165_MOUSE	Tmem165	34.8		0	0	0,00E+00		0	0	0,00E+00	8,36	2	2	6,54E+06
gii50401658 spl Q99N89.1 RM43_MOUSE	Mrpl43	20.2	14,21	3	4	6,70E+06		0	0	0,00E+00	10,38	2	3	6,52E+06
gii341942168 spl Q64012.3 RALY_MOUSE	Raly	33.2		0	0	0,00E+00		0	0	0,00E+00	6,73	2	2	6,38E+06
gii416621 spl P32261.1 ANT3_MOUSE	Serpinc1	52.0	9,46	5	5	2,94E+07	4,30	2	2	1,51E+07	3,44	2	2	6,16E+06
gii2494835 spl Q64692.1 SIA8D_MOUSE	St8sia4	41.2		0	0	0,00E+00		0	0	0,00E+00	11,98	4	5	6,10E+06
gii1345668 spl P47757.3 CAPZB_MOUSE	Capzb	31.3		0	0	0,00E+00	43,32	14	43	4,28E+08	10,47	3	3	6,08E+06
gii71151920 spl Q91VE6.1 MK671_MOUSE	Nrkf	36.2		0	0	0,00E+00		0	0	0,00E+00	8,20	2	2	6,04E+06
gii52782750 spl Q9CPO8.1 ATP5L_MOUSE	Atp5l	11.4		0	0	0,00E+00		0	0	0,00E+00	25,24	2	2	5,90E+06
gii12230747 spl Q88342.3 WDR1_MOUSE	Wdr1	66.4		0	0	0,00E+00		0	0	0,00E+00	6,93	4	4	5,89E+06
gii54041950 spl Q99N94.2 RM09_MOUSE	Mrpl9	30.2	10,94	3	3	5,54E+06		0	0	0,00E+00	7,92	2	2	5,77E+06
gii3913652 spl Q09046.1 IOXLA_MOUSE	Ii4i1	70.1		0	0	0,00E+00		0	0	0,00E+00	3,81	2	2	5,59E+06
gii3419411149 spl Q8B8T8.5 FLNA_MOUSE	Flna	281.0	1,40	3	3	2,60E+06	6,16	13	13	1,46E+07	2,46	5	5	5,10E+06
gii73620958 spl Q62095.2 DDX3Y_MOUSE	Ddx3y	73.4		0	0	0,00E+00	6,08	4	5	1,66E+07	4,86	3	3	5,02E+06
gii13633865 spl Q9QU10.1 RHOA_MOUSE	Rhoa	21.8		0	0	0,00E+00	8,29	2	2	5,27E+06	22,28	5	8	4,88E+06
gii158705865 spl A2AJ10.1 MA7D1_MOUSE	Map7d1	93.2	9,34	7	9	8,78E+06	6,38	5	8	9,65E+06	2,36	2	3	4,84E+06
gii1880318 spl Q99N93.1 RM16_MOUSE	Mrpl16	28.8		0	0	0,00E+00		0	0	0,00E+00	10,76	2	2	4,84E+06
gii3799719 spl Q8R0X7.1 ISGPL1_MOUSE	Sgpl1	63.6		0	0	0,00E+00		0	0	0,00E+00	5,81	3	3	4,82E+06
gii122134 spl P14430.1 HA18_MOUSE	H2-Q8	37.4	8,90	3	3	8,62E+06		0	0	0,00E+00	5,83	2	2	4,79E+06
gii18202285 spl P58252.2 EF2_MOUSE	Eef2	95.3	4,66	4	4	5,38E+06	2,10	2	2	1,05E+07	2,91	3	3	4,51E+06
gii548926 spl P30882.2 CCCL5_MOUSE	Ccl5	10.1		0	0	0,00E+00		0	0	0,00E+00	20,88	2	2	4,50E+06
gii119364598 spl P54987.2 IRG1_MOUSE	Irg1	53.7	3,28	2	2	2,64E+06		0	0	0,00E+00	7,38	4	4	4,48E+06
gii226693542 spl Q9WIT7.2 MYO1C_MOUSE	Myo1c	121.9		0	0	0,00E+00	20,13	17	21	6,89E+07	2,54	2	2	4,06E+06
gii3024089 spl Q61792.1 LASP1_MOUSE	Lasp1	30.0	5,32	2	2	1,72E+06		0	0	0,00E+00	10,27	3	3	3,65E+06
gii20455216 spl Q9Z204.1 HNRPC_MOUSE	Hnrmpc	34.4		0	0	0,00E+00	18,53	7	8	2,41E+07	7,03	2	2	3,61E+06
gii62510510 spl Q9CRB9.1 CHCH3_MOUSE	Chchd3	26.3	15,42	3	4	3,11E+06	8,81	2	3	4,13E+06	8,81	2	3	3,61E+06
gii46395721 spl Q7TPR4.1 ACTN1_MOUSE	Actn1	103.0		0	0	0,00E+00	40,13	32	52	1,22E+08	1,91	2	2	3,59E+06
gii62510642 spl Q8C111.2 GNL3_MOUSE	Gnl3	60.7		0	0	0,00E+00	4,65	2	3	3,46E+06	7,62	4	4	3,58E+06
gii85701216 spl Q9CW46.2 RAVR1_MOUSE	Raver1	79.3		0	0	0,00E+00		0	0	0,00E+00	2,67	2	2	3,50E+06
gii52783420 spl Q9CQ17.1 RU2B_MOUSE	Snrbp2	25.3		0	0	0,00E+00		0	0	0,00E+00	11,56	2	2	3,36E+06
gii18804111 spl Q9CR59.1 G45IP_MOUSE	Gadd45qjp1	25.8	8,56	2	3	1,57E+07		0	0	0,00E+00	8,56	2	2	3,30E+06
gii25089776 spl Q8DQC2.3 ATP5H_MOUSE	Atp5h	18.7		0	0	0,00E+00		0	0	0,00E+00	13,04	2	2	3,24E+06

gii711538261spiQ7TPV4.2IMBB1A_MOUSE	Mybbp1a	151.9		0	0	0.00E+00		0	0	0.00E+00	1.93	2	2	3.14E+06
gii341942181spiP14685.3IPSM3_MOUSE	Psm3d	60.7		0	0	0.00E+00		0	0	0.00E+00	6.79	3	3	2.79E+06
gii83305133spiP52503.2INDUS6_MOUSE	Ndufs6	13.0		0	0	0.00E+00	22.41	2	2	5.70E+06	22.41	2	2	2.52E+06
gii1351218spiP47226.1ITES_MOUSE	Tes	48.0		0	0	0.00E+00		0	0	0.00E+00	5.44	2	2	2.33E+06
gii52783213spiQ9CY58.2IPAIRB_MOUSE	Serbp1	44.7		0	0	0.00E+00		0	0	0.00E+00	6.63	2	2	2.29E+06
gii3419409261spiP27046.2IMAZA1_MOUSE	Man2a1	131.5		0	0	0.00E+00		0	0	0.00E+00	1.48	2	2	1.96E+06
gii1358351spiP26350.2IPTMA_MOUSE	Ptma	12.2	13.51	3	3	4.12E+06	13.51	3	4	9.37E+06	13.51	3	3	1.76E+06
gii612119301spiQ9EP78.1ICHST7_MOUSE	Chst7	54.7		0	0	0.00E+00		0	0	0.00E+00	2.48	2	2	1.71E+06
gii81896062spiQ8BGZ7.1IK2C75_MOUSE	Krt75	59.7	15.97	12	26	8.80E+08		0	0	0.00E+00		0	0	0.00E+00
gii123797712spiQ8VED5.2IK2C79_MOUSE	Krt79	57.5	9.42	9	30	7.32E+08		0	0	0.00E+00		0	0	0.00E+00
gii5490781spiP10639.3ITHIO_MOUSE	Txn1	11.7	20.95	2	3	4.47E+07		0	0	0.00E+00		0	0	0.00E+00
gii484281481spiQ9CQP1.3ICOX6C_MOUSE	Cox6c	8.5	11.84	2	3	3.38E+07	21.05	3	4	5.09E+07		0	0	0.00E+00
gii146345481spiP09411.4IPGK1_MOUSE	Pgk1	44.5	16.79	5	6	3.04E+07	16.79	5	5	3.56E+07		0	0	0.00E+00
gii3419412441spiP49290.2IPERE_MOUSE	Epx	81.3	9.36	7	10	2.60E+07	3.07	2	2	6.51E+06		0	0	0.00E+00
gii201417891spiP99029.2IPRD5_MOUSE	Prdx5	21.9	23.81	4	4	1.85E+07	15.71	3	4	1.51E+07		0	0	0.00E+00
gii1193621spiP08113.2ENPL_MOUSE	Hsp90b1	92.4	5.24	4	4	1.66E+07	7.48	6	7	2.20E+07		0	0	0.00E+00
gii32363971spiQ9CCQ7.3INDUB4_MOUSE	Ndufb4	15.1	35.66	3	3	1.65E+07		0	0	0.00E+00		0	0	0.00E+00
gii213629041spiQ9CYH6.1IRRS1_MOUSE	Rrs1	41.5	5.48	2	2	1.38E+07		0	0	0.00E+00		0	0	0.00E+00
gii7309561spiP40142.1ITKT_MOUSE	Tkt	67.6	9.47	6	6	1.31E+07	6.58	4	6	1.85E+07		0	0	0.00E+00
gii21759071spiQ9CQI6.3ICOTL1_MOUSE	Cott1	15.9	11.27	2	3	1.18E+07	11.27	2	3	2.25E+07		0	0	0.00E+00
gii3419406301spiQ9VW32.4IARC1B_MOUSE	Arpc1b	41.0	8.33	2	3	1.17E+07	42.74	16	33	4.96E+08		0	0	0.00E+00
gii1175021spiP14211.1ICALR_MOUSE	Calr	48.0	5.29	3	3	1.12E+07	6.49	3	3	2.07E+07		0	0	0.00E+00
gii321304231spiP51174.2IACADL_MOUSE	Acadl	47.9	5.58	2	2	1.05E+07	13.49	5	5	3.71E+07		0	0	0.00E+00
gii3419411971spiQ35375.2INRP2_MOUSE	Nrp2	104.6	2.79	2	4	9.26E+06		0	0	0.00E+00		0	0	0.00E+00
gii410177931spiQ9CQL5.1IRM18_MOUSE	Mrp18	20.7	11.67	2	3	7.88E+06		0	0	0.00E+00		0	0	0.00E+00
gii25005661spiP97425.1IECP2_MOUSE	Ear2	17.6	12.18	2	2	7.31E+06		0	0	0.00E+00		0	0	0.00E+00
gii3535263541spiP17751.4ITPIS_MOUSE	Tpi1	32.2	13.38	3	5	7.13E+06	16.72	4	6	1.47E+07		0	0	0.00E+00
gii1149301spiP28667.2IMRP_MOUSE	Marcks1	20.2	14.00	2	2	7.11E+06	14.00	2	2	1.67E+07		0	0	0.00E+00
gii13519421spiP48036.1IANXA5_MOUSE	Anxa5	35.7	12.54	5	5	6.99E+06	11.29	4	4	1.39E+07		0	0	0.00E+00
gii3419412391spiP09103.2IPDIA1_MOUSE	P4hb	57.0	6.29	3	4	6.65E+06	4.72	2	2	1.09E+07		0	0	0.00E+00
gii46396651spiQ9EQU5.1ISET_MOUSE	Set	33.4	8.30	2	2	5.96E+06		0	0	0.00E+00		0	0	0.00E+00
gii3419408301spiQ64282.2IFIT1_MOUSE	Ifit1	53.7	6.05	2	2	5.03E+06		0	0	0.00E+00		0	0	0.00E+00
gii885660571spiQ8VJL6.1ISFPQ_MOUSE	Sfpq	75.4	4.15	2	2	5.01E+06		0	0	0.00E+00		0	0	0.00E+00
gii819035021spiQ9CPY1.1IRM51_MOUSE	Mrp51	15.1	17.97	3	3	4.08E+06		0	0	0.00E+00		0	0	0.00E+00
gii17082591spiP30681.3IHMG82_MOUSE	Hmg82	24.1	13.81	2	2	3.96E+06		0	0	0.00E+00		0	0	0.00E+00
gii28515961spiQ93092.2ITALDO_MOUSE	Takdo1	37.4	5.04	2	2	3.91E+06	5.04	2	2	3.15E+06		0	0	0.00E+00
gii24947031spiQ61599.3IGDIR2_MOUSE	Arhgdib	22.8	10.50	2	2	3.80E+06		0	0	0.00E+00		0	0	0.00E+00
gii173686191spiQ9ME9.3INOG1_MOUSE	Gtpbp4	74.1	2.84	2	2	2.47E+06		0	0	0.00E+00		0	0	0.00E+00
gii711519831spiQ6URW6.1IMYH14_MOUSE	Mvh14	228.4		0	0	0.00E+00	4.95	12	53	7.23E+09		0	0	0.00E+00
gii471165731spiQ9JY9.3IARP3_MOUSE	Actr3	47.3		0	0	0.00E+00	45.69	17	43	5.70E+08		0	0	0.00E+00
gii1108257061spiQ9CVB6.3IARPC2_MOUSE	Arpc2	34.3		0	0	0.00E+00	64.00	20	57	5.70E+08		0	0	0.00E+00
gii342237301spiP47753.4ICAZA1_MOUSE	Capza1	32.9		0	0	0.00E+00	65.03	14	24	5.01E+08		0	0	0.00E+00
gii597995241spiQ9J28.1IFLI1_MOUSE	Flii	144.7		0	0	0.00E+00	2.52	3	4	4.26E+08		0	0	0.00E+00
gii625104601spiQ9CPW4.3IARPC5_MOUSE	Arpc5	16.3		0	0	0.00E+00	53.64	8	15	3.91E+08		0	0	0.00E+00
gii628998931spiQ9JM76.3IARPC3_MOUSE	Arpc3	20.5		0	0	0.00E+00	30.90	6	23	2.84E+08		0	0	0.00E+00
gii38372251spiQ99MU3.2IDSRAD_MOUSE	Adar	130.4		0	0	0.00E+00	2.89	3	3	2.42E+08		0	0	0.00E+00
gii3419409831spiQ99104.2IMYO5A_MOUSE	Myo5a	215.4		0	0	0.00E+00	28.01	52	82	1.48E+08		0	0	0.00E+00
gii219033821spiP48962.4IADT1_MOUSE	Slc25a4	32.9		0	0	0.00E+00	16.78	5	12	1.35E+08		0	0	0.00E+00
gii131239461spiP57780.1IACNA1_MOUSE	Actn4	104.9		0	0	0.00E+00	34.54	27	37	1.06E+08		0	0	0.00E+00
gii818795961spiQ91WK0.1ILRRF2_MOUSE	Lrrfp2	47.1		0	0	0.00E+00	22.17	9	13	6.11E+07		0	0	0.00E+00
gii628996731spiQ70200.1IAIF1_MOUSE	Aif1	16.9		0	0	0.00E+00	26.53	4	5	5.93E+07		0	0	0.00E+00
gii246382181spiQ91YW3.1ISH3L3_MOUSE	Sh3bgrl3	10.5		0	0	0.00E+00	26.88	2	3	5.32E+07		0	0	0.00E+00
gii242115581spiQ9QYB1.3ICLIC4_MOUSE	Clic4	28.7		0	0	0.00E+00	5.53	2	3	3.55E+07		0	0	0.00E+00
gii39135811spiQ61878.1IPRG2_MOUSE	Prg2	24.2		0	0	0.00E+00	13.45	3	5	3.17E+07		0	0	0.00E+00
gii3419406101spiQ60604.3IADSV_MOUSE	Scn	80.2		0	0	0.00E+00	14.13	9	10	3.09E+07		0	0	0.00E+00
gii1463454801spiP27773.2IPDIA3_MOUSE	Pdia3	56.6		0	0	0.00E+00	19.01	7	7	2.86E+07		0	0	0.00E+00
gii818824071spiQ569Z6.1ITR150_MOUSE	Thrap3	108.1		0	0	0.00E+00	5.89	5	8	2.23E+07		0	0	0.00E+00
gii173803331spiQ9D855.3IQCR7_MOUSE	Uacr7	13.5		0	0	0.00E+00	21.62	2	5	1.75E+07		0	0	0.00E+00
gii136293691spiQ9WU43.1IK6PP_MOUSE	Pfkp	85.4		0	0	0.00E+00	5.23	4	4	1.69E+07		0	0	0.00E+00
gii146345421spiQ61553.4IFSCN1_MOUSE	Fscn1	54.5		0	0	0.00E+00	4.26	2	3	1.63E+07		0	0	0.00E+00
gii24940221spiQ62426.1ICYTB_MOUSE	Cstb	11.0		0	0	0.00E+00	32.65	2	2	1.58E+07		0	0	0.00E+00
gii818815921spiQ9D898.1IARPSL_MOUSE	Arpc5l	17.0		0	0	0.00E+00	46.41	4	4	1.42E+07		0	0	0.00E+00
gii1164871spiP08207.2IS10A_MOUSE	S100a10	11.2		0	0	0.00E+00	27.84	2	5	1.37E+07		0	0	0.00E+00
gii1267521spiP26645.2IMARCS_MOUSE	Marcks	29.6		0	0	0.00E+00	11.65	2	3	1.23E+07		0	0	0.00E+00
gii465771161spiQ9D1G1.1IRAB1B_MOUSE	Rab1b	22.2		0	0	0.00E+00	19.40	4	4	1.15E+07		0	0	0.00E+00
gii819024781spiQ91Z49.1IUIF_MOUSE	Fytd1	35.9		0	0	0.00E+00	6.94	2	2	9.93E+06		0	0	0.00E+00
gii341940791spiP49710.2IHCLS1_MOUSE	Hcls1	54.2		0	0	0.00E+00	4.53	2	2	6.35E+06		0	0	0.00E+00
gii122305891spiQ9JU8.1ISH3L1_MOUSE	Sh3bgrl	12.8		0	0	0.00E+00	20.18	2	2	6.12E+06		0	0	0.00E+00
gii559762011spiQ9D0E1.3IHNRPM_MOUSE	Hnrpm	77.6		0	0	0.00E+00	3.29	2	2	4.91E+06		0	0	0.00E+00
gii347595711spiQ52K18.2ISRRM1_MOUSE	Srrm1	106.8		0	0	0.00E+00	2.22	2	2	4.34E+06		0	0	0.00E+00
gii540410711spiQ9WUM4.2ICOR1C_MOUSE	Coro1c	53.1		0	0	0.00E+00	7.81	3	3	3.84E+06		0	0	0.00E+00
gii212633741spiQ91V92.1IACLY_MOUSE	Acly	119.7		0	0	0.00E+00	1.56	2	2	2.66E+06		0	0	0.00E+00
gii1108164181spiQ9D877.2ISLIRP_MOUSE	Sllrp	12.6		0	0	0.00E+00	16.96	2	2	1.67E+06		0	0	0.00E+00