

Table 2. Probes with greater than 1.5 fold change following Snf5 inactivation.

Probe set	Gene	Accession no.	LocusLink	Fold change
1437591_a_at	WD repeat domain 1	AW542545	22388	42.51
1421893_a_at	tripeptidyl peptidase II	BG066807	22019	38.07
1416617_at	acetyl-Coenzyme A synthetase 2 (AMP forming)-lik	NM_080575	68738	38
1431213_a_at	gb:BG297038 /DB_XREF=gi:13060290 /DB_XREF	BG297038		15.36
1424459_at	cDNA sequence BC005662	BG068664	210992	14.21
1453724_a_at	serine (or cysteine) proteinase inhibitor, clade F, mε	AK012411	20317	10.94
1425725_s_at	Protein phosphatase 2, regulatory subunit B (B56),	BF136532	26931	10.47
1421972_s_at	host cell factor C1	NM_008224	15161	10.1
1422606_at	C1q and tumor necrosis factor related protein 3	NM_030888	81799	7.17
1430533_a_at	catenin beta	BI134907	12387	6.92
1449939_s_at	delta-like 1 homolog (Drosophila)	NM_010052	13386	6.66
1420953_at	adducin 1 (alpha)	BF140063	11518	6.59
1416371_at	apolipoprotein D	NM_007470	11815	6.34
1456543_at	G protein-coupled receptor 73	BB037474	58182	6.25
1450988_at	G protein-coupled receptor 49	BB751088	14160	6.22
1455892_x_at	gb:BB794742 /DB_XREF=gi:16964361 /DB_XREF	BB794742		5.79
1448201_at	secreted frizzled-related sequence protein 2	NM_009144	20319	5.68
1420549_at	guanylate nucleotide binding protein 1	NM_010259	14468	5.51
1416573_at	protein O-fucosyltransferase 2	BC018194	80294	5.2
1427520_a_at	myosin, heavy polypeptide 1, skeletal muscle, adult	AJ293626	17879	5.13
1427038_at	preproenkephalin 1	M13227	18619	5.11
1432143_a_at	high mobility group box transcription factor 1	AK006835	73389	5.01
1431645_a_at	guanosine diphosphate (GDP) dissociation inhibitor	AK013104	14569	5
1422587_at	transmembrane protein 45a	NM_019631	56277	4.92
1433491_at	erythrocyte protein band 4.1-like 2	BE951907	13822	4.87
1456080_a_at	Tumor differentially expressed 1	BM239368	26943	4.85
1420924_at	tissue inhibitor of metalloproteinase 2	M93954	21858	4.73
1452899_at	RNA imprinted and accumulated in nucleus	AK017440	75745	4.62
1420855_at	elastin	BB229377	13717	4.52
1426027_a_at	Rho GTPase activating protein 10	AF297029	78514	4.51
1451777_at	cDNA sequence BC013672	BC013672	234311	4.45
1427868_x_at	myosin, heavy polypeptide 1, skeletal muscle, adult	AJ002522	17879	4.41
1455869_at	gb:BG862223 /DB_XREF=gi:14212761 /DB_XREF	BG862223		4.39
1427565_a_at	ATP-binding cassette, sub-family C (CFTR/MRP), r	AF213387	27416	4.35
1425326_at	ATP citrate lyase	BI456232	104112	4.28
1421604_a_at	Kruppel-like factor 3 (basic)	NM_008453	16599	4.26
1452907_at	gb:AK010101.1 /DB_XREF=gi:12845310 /FEA=mF	AK010101		4.26
1453196_a_at	2'-5' oligoadenylate synthetase-like 2	BQ033138	23962	4.2
1430980_a_at	eukaryotic translation initiation factor 4A1	BI656407	13681	4.18
1460187_at	secreted frizzled-related sequence protein 1 /// RIK	BI658627	20377 /// 72	4.17
1424041_s_at	complement component 1, s subcomponent	BC022123	50908	4
1437685_x_at	fibromodulin	BB235530	14264	3.97
1451726_at	myotubularin related protein 6	BC020019	219135	3.95
1438966_x_at	gb:AV290700 /DB_XREF=gi:6304731 /DB_XREF=,	AV290700		3.91
1449551_at	myosin IC	NM_008659	17913	3.89
1460250_at	sclerostin domain containing 1	BC021458	66042	3.85
1449340_at	sclerostin domain containing 1	BC021458	66042	3.84
1415823_at	gb:BB459479 /DB_XREF=gi:16426408 /DB_XREF	BG060909		3.83
1416652_at	asporin	NM_025711	66695	3.71
1437324_x_at	fibromodulin	BB504826	14264	3.65
1416515_at	fascin homolog 1, actin bundling protein (Strongylo	NM_007984	14086	3.61

1423348_at	frizzled homolog 8 (Drosophila)	AV345166	14370	3.61
1415832_at	angiotensin II receptor, type 2	NM_007429	11609	3.51
1420899_at	RAB18, member RAS oncogene family	AW542340	19330	3.47
1450033_a_at	signal transducer and activator of transcription 1	AW214029	20846	3.45
1427054_s_at	RIKEN cDNA D930038M13 gene	BC026627	320712	3.35
1456494_a_at	tripartite motif protein 30 /// tripartite motif protein 30	BG068242	20128 /// 20128	3.34
1426439_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linker	AA210261	26900	3.33
1437718_x_at	fibromodulin	BB483571	14264	3.31
1421816_at	glutathione reductase 1	AK019177	14782	3.3
1456084_x_at	fibromodulin	BB532202	14264	3.3
1417009_at	complement component 1, r subcomponent	NM_023143	50909	3.27
1416572_at	matrix metalloproteinase 14 (membrane-inserted)	NM_008608	17387	3.26
1423294_at	mesoderm specific transcript	AW555393	17294	3.26
1416318_at	gb:AF426024.1 /DB_XREF=gi:18034363 /GEN=Seq	AF426024		3.25
1451406_a_at	proprotein convertase subtilisin/kexin type 5	BC013068	18552	3.23
1426278_at	interferon, alpha-inducible protein 27	AY090098	76933	3.22
1436713_s_at	GTL2, imprinted maternally expressed untranslated control	BM119226	17263	3.21
1427053_at	RIKEN cDNA D930038M13 gene	BC026627	320712	3.18
1448380_at	lectin, galactoside-binding, soluble, 3 binding protein	NM_011150	19039	3.18
1426340_at	solute carrier family 1 (glial high affinity glutamate transporter)	BB357585	20512	3.17
1451943_a_at	protein phosphatase 1A, magnesium dependent, alpha	AF259673	19042	3.16
1434357_a_at	karyopherin (importin) beta 1	AW544889	16211	3.13
1416302_at	early B-cell factor 1	BB125261	13591	3.12
1426324_at	MHC (A.CA/J(H-2K-f) class I antigen	M33151	56628	3.1
1450655_at	hect (homologous to the E6-AP (UBE3A) carboxyl terminal domain)	AA214868	235439	3.1
1426852_x_at	nephroblastoma overexpressed gene	X96585	18133	3.08
1425929_a_at	ring finger protein 14	AF249668	56736	3.03
1460248_at	carboxypeptidase X 2 (M14 family)	AF017639	55987	3.03
1425329_a_at	diaphorase 1 (NADH)	AF332060	109754	3.02
1430514_a_at	RIKEN cDNA 2410026K10 gene	AK002762	66478	3.02
1448127_at	ribonucleotide reductase M1	BB758819	20133	3.01
1450779_at	fatty acid binding protein 7, brain	NM_021272	12140	3.01
1418930_at	chemokine (C-X-C motif) ligand 10	NM_021274	15945	2.99
1450403_at	signal transducer and activator of transcription 2	AF088862	20847	2.99
1450089_a_at	signal recognition particle receptor, B subunit	NM_009275	20818	2.96
1432466_a_at	apolipoprotein E	AK019319	11816	2.95
1416321_s_at	proline arginine-rich end leucine-rich repeat	NM_054077	116847	2.94
1452244_at	RIKEN cDNA 6330406I15 gene	AK018128	70717	2.93
1417716_at	glutamate oxaloacetate transaminase 2, mitochondrial	U82470	14719	2.9
1450783_at	interferon-induced protein with tetratricopeptide repeats	NM_008331	15957	2.9
1422450_at	catenin src	NM_007615	12388	2.87
1425527_at	paired related homeobox 1	L06502	18933	2.87
1454159_a_at	insulin-like growth factor binding protein 2	AK011784	16008	2.87
1426937_at	RIKEN cDNA 6330406I15 gene	AK018128	70717	2.86
1427048_at	smoothed homolog (Drosophila)	AW555326	319757	2.84
1439260_a_at	ectonucleotide pyrophosphatase/phosphodiesterase 1	BB039510	209558	2.84
1448416_at	matrix gamma-carboxyglutamate (gla) protein	NM_008597	17313	2.84
1432331_a_at	paired related homeobox 2	AK019971	20204	2.81
1432478_a_at	IBR domain containing 3	AK015966	75234	2.81
1421315_s_at	gb:BI688355 /DB_XREF=gi:15650984 /DB_XREF=	BI688355		2.8
1419332_at	EGF-like-domain, multiple 6	NM_019397	54156	2.79
1420901_a_at	hexokinase 1	NM_010438	15275	2.78
1455061_a_at	acetyl-Coenzyme A acyltransferase 2 (mitochondrial)	BB718075	52538	2.78

1419662_at	osteoglycin	BB542051	18295	2.77
1425911_a_at	fibroblast growth factor receptor 1	M65053	14182	2.77
1450105_at	a disintegrin and metalloprotease domain 10	BG921598	11487	2.77
1425545_x_at	histocompatibility 2, D region	M86502	14980	2.76
1422781_at	toll-like receptor 3	NM_126166	142980	2.75
1427580_a_at	RNA imprinted and accumulated in nucleus	BB649603	75745	2.75
1425803_a_at	methyl-CpG binding domain protein 2	AF072245	17191	2.74
1426438_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linker	AA210261	26900	2.74
1448330_at	glutathione S-transferase, mu 1	NM_010358	14862	2.74
1449082_at	microfibrillar associated protein 5	NM_015776	50530	2.72
1424948_x_at	histocompatibility 2, D region locus 1 /// histocompatibility 2, D region locus 1	L23495	14964 /// 14	2.71
1451683_x_at	histocompatibility 2, D region locus 1 /// histocompatibility 2, D region locus 1	M34962	14964 /// 14	2.71
1451784_x_at	histocompatibility 2, D region locus 1 /// histocompatibility 2, D region locus 1	L36068	14964 /// 14	2.71
1424903_at	jumonji, AT rich interactive domain 1D (Rbp2 like)	AF127244	20592	2.7
1417210_at	eukaryotic translation initiation factor 2, subunit 3, small	NM_012011	26908	2.69
1432411_a_at	F-box and WD-40 domain protein 2	AK009893	30050	2.69
1422185_a_at	diaphorase 1 (NADH)	NM_029787	109754	2.68
1418317_at	LIM homeobox protein 2	NM_010710	16870	2.67
1430692_a_at	Sel1 (suppressor of lin-12) 1 homolog (C. elegans)	AK005023	20338	2.67
1460279_a_at	general transcription factor II I	AF043219	14886	2.67
1427416_x_at	dual specificity phosphatase 7	BC025048	235584	2.66
1428492_at	GLI pathogenesis-related 2	BM208214	384009	2.66
1417343_at	FXYD domain-containing ion transport regulator 6	AB032010	59095	2.65
1427746_x_at	histocompatibility 2, K1, K region	S70184	14972	2.65
1454791_a_at	retinoblastoma binding protein 4	AU040918	19646	2.64
1421346_a_at	solute carrier family 6 (neurotransmitter transporter), member 6	NM_009320	21366	2.63
1451931_x_at	histocompatibility 2, D region locus 1 /// histocompatibility 2, D region locus 1	M69068	14964 /// 14	2.63
1451939_a_at	sushi-repeat-containing protein	AB028050	51795	2.61
1431592_a_at	SH3-domain kinase binding protein 1	AK007283	58194	2.6
1418532_at	frizzled homolog 2 (Drosophila)	BB371406	57265	2.59
1419088_at	tissue inhibitor of metalloproteinase 3	BI111620	21859	2.59
1421813_a_at	prosaposin	BM212050	19156	2.59
1425336_x_at	Histocompatibility 2, K1, K region	BC011306	14972	2.59
1426598_at	ubiquitously transcribed tetratricopeptide repeat gene 1	BB742957	22290	2.59
1420920_a_at	ADP-ribosylation factor 1	NM_007476	11840	2.58
1422631_at	aryl-hydrocarbon receptor	BE989096	11622	2.58
1449556_at	histocompatibility 2, T region locus 23 /// RIKEN cDNA 1500015O10	NM_010398	15040 /// 31	2.58
1450672_a_at	three prime repair exonuclease 1	NM_011637	22040	2.58
1423261_at	RIKEN cDNA 1500015O10 gene	BB426248	78896	2.57
1452471_at	interleukin 17 receptor D	AF459444	171463	2.56
1422903_at	lymphocyte antigen 86	NM_010745	17084	2.54
1459546_s_at	ectonucleotide pyrophosphatase/phosphodiesterase 1	C76301	18605	2.54
1419089_at	tissue inhibitor of metalloproteinase 3	BI111620	21859	2.53
1416013_at	phospholipase D3	NM_011116	18807	2.52
1421217_a_at	lectin, galactose binding, soluble 9	NM_010708	16859	2.52
1437932_a_at	claudin 1	AV227581	12737	2.51
1422620_s_at	phosphatidic acid phosphatase 2a	NM_008903	19012	2.5
1424617_at	interferon-induced protein 35	BC008158	70110	2.5
1426607_at	MRNA similar to hypothetical protein FLJ21657 (cDNA)	BG068672		2.5
1422561_at	a disintegrin-like and metalloprotease (reprolysin type) family 1 member 1	BB658835	23794	2.48
1450429_at	calpain 6	AI747133	12338	2.47
1451782_a_at	solute carrier family 29 (nucleoside transporters), member 1	AF305501	63959	2.47
1416157_at	vinculin	NM_009502	22330	2.46

1426936_at	cDNA sequence BC005512	BC002257	192885	2.46
1438676_at	macrophage activation 2 like	BM241485	100702	2.46
1419561_at	chemokine (C-C motif) ligand 3	NM_011337	20302	2.45
1451241_at	laminin B1 subunit 1	BG970109	16777	2.45
1416416_x_at	glutathione S-transferase, mu 1	NM_010358	14862	2.44
1448494_at	growth arrest specific 1	BB550400	14451	2.44
1416066_at	CD9 antigen	NM_007657	12527	2.43
1417876_at	Fc receptor, IgG, high affinity I	AF143181	14129	2.43
1421992_a_at	insulin-like growth factor binding protein 4	NM_010517	16010	2.41
1427683_at	early growth response 2	X06746	13654	2.41
1435176_a_at	inhibitor of DNA binding 2	BF019883	15902	2.41
1448908_at	phosphatidic acid phosphatase type 2B	NM_080555	67916	2.41
1449645_s_at	chaperonin subunit 3 (gamma)	AL024092	12462	2.41
1421058_at	alcohol dehydrogenase 7 (class IV), mu or sigma p	NM_009626	11529	2.4
1448842_at	cysteine dioxygenase 1, cytosolic	NM_033037	12583	2.4
1450377_at	thrombospondin 1	AI385532	21825	2.4
1421313_s_at	cortactin	BI688355	13043	2.39
1424807_at	laminin, alpha 4	BB053010	16775	2.39
1431744_a_at	stromal membrane-associated protein 1	AK014888	98366	2.39
1433685_a_at	RIKEN cDNA 6430706D22 gene	BM248225	381280	2.39
1435637_at	DNA segment, Chr 8, Wayne State University 49, e	AW554709	71927	2.39
1449335_at	tissue inhibitor of metalloproteinase 3	BI111620	21859	2.39
1454011_a_at	replication protein A2	AK011530	19891	2.39
1426851_a_at	nephroblastoma overexpressed gene	X96585	18133	2.38
1431420_s_at	RIKEN cDNA 2610524G07 gene	BI082843	66494	2.38
1416855_at	growth arrest specific 1	BB550400	14451	2.37
1453556_x_at	RIKEN cDNA 2410026K10 gene	AK002762	66478	2.37
1424010_at	microfibrillar-associated protein 4	BC022666	76293	2.35
1436372_a_at	expressed sequence AA415817	BB026304	94184	2.35
1453651_a_at	brain protein 44-like	AV223468	55951	2.35
1423053_at	ADP-ribosylation factor 4	BI653265	11843	2.34
1448601_s_at	homeo box, msh-like 1	BC016426	17701	2.34
1451413_at	calpastatin	AB026997	12380	2.34
1433446_at	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1	BB705380	208715	2.33
1417127_at	homeo box, msh-like 1	BC016426	17701	2.32
1417455_at	transforming growth factor, beta 3	BC014690	21809	2.32
1425942_a_at	glycoprotein m6b	AF254879	14758	2.32
1450008_a_at	catenin beta	NM_007614	12387	2.32
1415982_at	RIKEN cDNA 5031400M07 gene	AV258950	80517	2.3
1420814_at	guanosine diphosphate (GDP) dissociation inhibitor	NM_008112	14569	2.3
1424596_s_at	LIM and cysteine-rich domains 1	BC019124	30937	2.3
1450007_at	RIKEN cDNA 1500003O03 gene	NM_019769	56398	2.3
1426758_s_at	GTL2, imprinted maternally expressed untranslated c	Y13832	17263	2.29
1417063_at	complement component 1, q subcomponent, beta c	NM_009777	12260	2.28
1417381_at	complement component 1, q subcomponent, alpha	NM_007572	12259	2.28
1448228_at	lysyl oxidase	M65143	16948	2.28
1416800_at	transient receptor potential cation channel, subfam	AV320241	58800	2.26
1419636_at	RIKEN cDNA 4833420G17 gene	BG067119	67392	2.26
1421917_at	platelet derived growth factor receptor, alpha polyp	AW537708	18595	2.26
1427773_a_at	Rab acceptor 1 (prenylated)	L40934	14470	2.26
1449207_a_at	kinesin family member 20A	NM_009004	19348	2.26
1418454_at	microfibrillar associated protein 5	NM_015776	50530	2.25
1449164_at	CD68 antigen	BC021637	12514	2.25

1453321_at	fibronectin type III domain containing 1	AK003938	68655	2.25
1453836_a_at	monoglyceride lipase	AK006949	23945	2.25
1419043_a_at	interferon inducible GTPase 1	BM239828	60440	2.24
1421433_at	zinc finger homeodomain 4	NM_030708	80892	2.24
1423045_at	nuclear cap binding protein subunit 2	BE285362	68092	2.24
1448475_at	olfactomedin-like 3	NM_133859	99543	2.24
1448732_at	cathepsin B	M14222	13030	2.24
1450040_at	tissue inhibitor of metalloproteinase 2	M93954	21858	2.24
1419482_at	complement component 3a receptor 1	NM_009779	12267	2.23
1424895_at	G-protein signalling modulator 2 (AGS3-like, C. ele	BC021308	76123	2.23
1427091_at	expressed sequence AI481105	BC025488	98999	2.23
1431081_a_at	phospholipid scramblase 3	AK011716	70310	2.23
1423379_at	nuclear factor of activated T-cells, cytoplasmic, calc	BF227641	73181	2.22
1423890_x_at	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide	BC027319	11931	2.22
1448218_s_at	tyrosine 3-monooxygenase/tryptophan 5-monooxyc	AV027921	22631	2.22
1418796_at	C-type lectin domain family 11, member a	NM_009131	20256	2.21
1448293_at	early B-cell factor 1	BB125261	13591	2.21
1453307_a_at	anaphase-promoting complex subunit 5	AK003821	59008	2.21
1417979_at	tenomodulin	AF291655	64103	2.2
1427301_at	CD48 antigen	BE634960	12506	2.2
1438661_a_at	ADP-ribosylation factor 2	AV023312	11841	2.2
1426869_at	biregional cell adhesion molecule-related/down-reg	BB005556	117606	2.19
1434370_s_at	Fas-associated factor 1	AV016515	14084	2.19
1420612_s_at	protein tyrosine phosphatase 4a2	BE134116	19244	2.18
1426680_at	selenoprotein N, 1	AK003819	74777	2.17
1448958_at	interferon alpha responsive gene	NM_022329	64164	2.17
1451897_a_at	neighbor of Brca1 gene 1	AF227188	17966	2.17
1452430_s_at	splicing factor, arginine/serine-rich 1 (ASF/SF2)	X66091	110809	2.17
1455961_at	Membrane metallo endopeptidase	AV174022	17380	2.17
1417570_at	anaphase promoting complex subunit 1	NM_008569	17222	2.16
1452398_at	phospholipase C, epsilon 1	AV306884	74055	2.16
1416513_at	laminin, beta 2	BC026051	16779	2.15
1418541_at	RIKEN cDNA 2810429O05 gene	NM_134046	52504	2.15
1423549_at	solute carrier family 1 (glutamate/neutral amino aci	BB277461	55963	2.15
1418457_at	chemokine (C-X-C motif) ligand 14	AF252873	57266	2.14
1419066_at	immediate early response 5-like	NM_030244	72500	2.14
1419442_at	matrilin 2	BC005429	17181	2.14
1430388_a_at	sulfatase 2	AK008108	72043	2.13
1450932_s_at	dedicator of cytokinesis 9	BB795072	105445	2.13
1416072_at	CD34 antigen	NM_133654	12490	2.12
1418126_at	gb:NM_013653.1 /DB_XREF=gi:7305460 /GEN=Sc	NM_013653		2.12
1423062_at	gb:AV175389 /DB_XREF=gi:16384293 /DB_XREF:	AV175389		2.12
1423978_at	SH3-binding kinase 1	BC025837	104175	2.12
1451152_a_at	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide	BC027319	11931	2.12
1451289_at	Double cortin and calcium/calmodulin-dependent pi	AW105916	13175	2.12
1423607_at	lumican	AK014312	17022	2.11
1423627_at	NAD(P)H dehydrogenase, quinone 1	AV158882	18104	2.11
1427692_a_at	calcium/calmodulin-dependent serine protein kinas	Y17137	12361	2.11
1437626_at	zinc finger protein 36, C3H type-like 2	BB031791	12193	2.11
1419015_at	WNT1 inducible signaling pathway protein 2	NM_016873	22403	2.1
1427314_at	transmembrane emp24 protein transport domain cc	AV225029	66676	2.1
1434748_at	cytoskeleton associated protein 2	BM208103	80986	2.1
1448754_at	retinol binding protein 1, cellular	NM_011254	19659	2.1

1450170_x_at	histocompatibility 2, D region locus 1 /// histocompat	NM_010380	14964 /// 14	2.1
1418020_s_at	carboxypeptidase D	NM_007754	12874	2.09
1429514_at	phosphatidic acid phosphatase type 2B	AW111876	67916	2.09
1451884_a_at	LSM2 homolog, U6 small nuclear RNA associated (	AF204156	27756	2.09
1452997_at	RIKEN cDNA 2610005L07 gene	BE692399	381598	2.09
1427567_a_at	tropomyosin 3, gamma	BE380713	59069	2.08
1427831_s_at	zinc finger protein 260	L36316	26466	2.08
1456386_at	RIKEN cDNA B330012G18 gene	BB203348	170791	2.08
1417595_at	mesenchyme homeobox 1	NM_010791	17285	2.07
1424271_at	double cortin and calcium/calmodulin-dependent pr	AW105916	13175	2.07
1424339_at	2'-5' oligoadenylate synthetase-like 1	AB067533	231655	2.07
1448613_at	extracellular matrix protein 1	NM_007899	13601	2.07
1449928_at	t-complex-associated-testis-expressed 1-like	NM_025975	67117	2.07
1454953_at	Ring finger protein 157	BB246182	217340	2.07
1449401_at	complement component 1, q subcomponent, gamr	NM_007574	12262	2.06
1451335_at	placenta-specific 8	AF263458	231507	2.06
1417462_at	CAP, adenylate cyclase-associated protein 1 (yeast)	NM_007598	12331	2.05
1421871_at	SH3-binding domain glutamic acid-rich protein like	NM_019989	56726	2.05
1451912_a_at	fibroblast growth factor receptor-like 1	AF321301	116701	2.05
1452432_at	tissue factor pathway inhibitor	BF451808	21788	2.05
1455868_a_at	tubulin, gamma complex associated protein 2	BB387906	74237	2.05
1415945_at	minichromosome maintenance deficient 5, cell divis	NM_008566	17218	2.04
1418204_s_at	allograft inflammatory factor 1	NM_019467	11629	2.04
1435129_at	gb:AW495875 /DB_XREF=gi:7066156 /DB_XREF= AW495875			2.04
1450291_s_at	gb:NM_022429.1 /DB_XREF=gi:11967954 /GEN=N	NM_022429		2.04
1460549_a_at	CDC23 (cell division cycle 23, yeast, homolog)	BB492440	52563	2.04
1416239_at	argininosuccinate synthetase 1	NM_007494	11898	2.03
1418078_at	proteasome (prosome, macropain) 28 subunit, 3	U60330	19192	2.03
1419042_at	interferon inducible GTPase 1	BM239828	60440	2.03
1426734_at	cDNA sequence BC022623	BB008324	224093	2.03
1416459_at	ADP-ribosylation factor 2	NM_007477	11841	2.02
1421812_at	gb:AK014566.1 /DB_XREF=gi:12852498 /GEN=Ta	AF043943		2.02
1422962_a_at	proteasome (prosome, macropain) subunit, beta ty	NM_010724	16913	2.02
1425373_a_at	tumor necrosis factor superfamily, member 5-induc	BC016606	107047	2.02
1427126_at	heat shock protein 1B	M12573	15511	2.02
1428380_at	RIKEN cDNA 0610007C21 gene	AK002276	381629	2.02
1452925_a_at	membrane-associated ring finger (C3HC4) 5	AK009364	69104	2.02
1417756_a_at	lymphocyte specific 1	NM_019391	16985	2.01
1426998_at	testis expressed gene 27	BG976649	21769	2.01
1451860_a_at	tripartite motif protein 30	AF220015	20128	2.01
1452031_at	solute carrier family 1 (glial high affinity glutamate t	BB357585	20512	2.01
1460295_s_at	interleukin 6 signal transducer	AA717838	16195	2.01
1460331_at	transmembrane 9 superfamily member 2	BB747462	68059	2.01
1416467_at	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3,	L25126	13205	2
1420828_s_at	tyrosine 3-monooxygenase/tryptophan 5-monooxyc	NM_011739	22630	2
1425896_a_at	fibrillin 1	AF007248	14118	2
1429681_a_at	glycoprotein, synaptic 2	AK020708	106529	2
1438887_a_at	germ cell-less homolog (Drosophila)	BM239632	23885	2
1451236_at	RAS-like, estrogen-regulated, growth-inhibitor	BC026463	232441	2
1451832_at	chemokine-like factor	AF401530	75458	2
1428257_s_at	dynein, cytoplasmic, light chain 2A	BG791323	67068	1.99
1417501_at	F-box only protein 6b	NM_015797	50762	1.98
1417793_at	interferon inducible GTPase 2	NM_019440	54396	1.98

1418819_at	ADP-ribosylation factor-like 10C	NM_026011	67166	1.98
1418985_at	cDNA sequence BC003236	NM_030249	80281	1.98
1423263_at	H3 histone, family 3A	BI111967	15078	1.98
1423368_at	lysosomal-associated protein transmembrane 4A	BI695636	17775	1.98
1423771_at	protein kinase C, delta binding protein	BC009660	109042	1.98
1424234_s_at	mesenchyme homeobox 2	BC002076	17286	1.98
1416030_a_at	minichromosome maintenance deficient 7 (S. cerevisiae)	NM_008568	17220	1.97
1456393_at	RIKEN cDNA 2310002J21 gene	AI642124	66360	1.97
1418369_at	DNA primase, p49 subunit	J04620	19075	1.96
1420841_at	protein tyrosine phosphatase, receptor type, F	BF235516	19268	1.96
1422264_s_at	basic transcription element binding protein 1	NM_010638	16601	1.96
1426260_a_at	UDP-glucuronosyltransferase 1 family, member 2 //	D87867	22236 /// 35	1.96
1429534_a_at	inner membrane protein, mitochondrial	BB222675	76614	1.96
1431746_a_at	ubiquitin-activating enzyme E1C	AK014433	22200	1.96
1448620_at	Fc receptor, IgG, low affinity III	NM_010188	14131	1.96
1451987_at	arrestin, beta 2	BC016642	216869	1.96
1454875_a_at	retinoblastoma binding protein 4	BF011461	19646	1.96
1415677_at	dehydrogenase/reductase (SDR family) member 1	NM_026819	52585	1.95
1418698_a_at	ferrochelatase	NM_007998	14151	1.95
1426781_at	radical S-adenosyl methionine and flavodoxin domain	BF021045	100929	1.95
1427552_a_at	glutathione transferase zeta 1 (maleylacetoacetate	AB041613	14874	1.95
1431295_a_at	syntaxin 18	AK016910	71116	1.95
1449060_at	kinesin family member 2C	NM_134471	73804	1.95
1417926_at	RIKEN cDNA 5830426I05 gene	NM_133762	76044	1.94
1423916_s_at	myeloid leukemia factor 2	BC003975	30853	1.94
1429252_at	gb:AK002491.1 /DB_XREF=gi:12832513 /FEA=mF	AK002491		1.94
1431827_a_at	tousled-like kinase 2 (Arabidopsis)	AK014829	24086	1.94
1450112_a_at	growth arrest specific 2	NM_008087	14453	1.94
1452920_a_at	peptidylprolyl isomerase (cyclophilin)-like 2	AK009460	66053	1.94
1422034_a_at	paralemmin	NM_023128	18483	1.93
1422414_a_at	calmodulin 2	NM_007589	12314	1.93
1423675_at	ubiquitin specific protease 1	BC018179	230484	1.93
1425764_a_at	branched chain aminotransferase 2, mitochondrial	AF031467	12036	1.93
1427132_at	myotubularin related protein 13	BG242242	319934	1.93
1428922_at	RIKEN cDNA 1200009O22 gene	AK004681	66873	1.93
1450388_s_at	twisted gastrulation homolog 1 (Drosophila)	BC004850	65960	1.93
1420307_a_at	phosphatidylinositol transfer protein, beta	BG975479	56305	1.92
1433805_at	Janus kinase 1	BQ032637	16451	1.92
1448749_at	pleckstrin	AF181829	56193	1.92
1451285_at	fusion, derived from t(12;16) malignant liposarcoma	AF224264	233908	1.92
1455480_s_at	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homologue)	BG070759	66105	1.92
1460694_s_at	supervillin	BM203457	225115	1.92
1417961_a_at	tripartite motif protein 30	BM240719	20128	1.91
1425351_at	neoplastic progression 3	BC011325	76650	1.91
1450678_at	gb:NM_008404.1 /DB_XREF=gi:6680489 /GEN=Itc	NM_008404		1.91
1454699_at	Sestrin 1	BG076140	140742	1.91
1419554_at	CD47 antigen (Rh-related antigen, integrin-associated	NM_010581	16423	1.9
1438759_x_at	gb:AU046270 /DB_XREF=gi:3982453 /DB_XREF=	AU046270		1.9
1448460_at	activin A receptor, type 1	NM_007394	11477	1.9
1450776_at	expressed sequence AU041707	NM_018743	102247	1.9
1456616_a_at	basigin	AV035166	12215	1.9
1419276_at	ectonucleotide pyrophosphatase/phosphodiesterase	AF339910	18605	1.89
1420915_at	signal transducer and activator of transcription 1	AW214029	20846	1.89

1421491_a_at	RIKEN cDNA 4930579A11 gene	NM_029478	75909	1.89
1428500_at	RIKEN cDNA 2210419D22 gene	AK008985	73661	1.89
1451426_at	DNA segment, Chr 11, Lothar Hennighausen 2, ex	AF316999	80861	1.89
1452429_s_at	ATP-binding cassette, sub-family F (GCN20), mem	BF236176	224742	1.89
1422062_at	macrophage scavenger receptor 1	BC003814	20288	1.88
1434877_at	neuronal pentraxin 1	AI152800	18164	1.88
1448972_at	glutamate receptor, ionotropic, AMPA1 (alpha 1)	NM_008165	14799	1.88
1452077_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-link	AA210261	26900	1.88
1417133_at	peripheral myelin protein	NM_008885	18858	1.87
1423746_at	thioredoxin domain containing 5	BC016252	105245	1.87
1428707_at	RIKEN cDNA 2610009E16 gene	AK011360	69202	1.87
1431686_a_at	glia maturation factor, beta	AK013970	63985	1.87
1448590_at	procollagen, type VI, alpha 1	NM_009933	12833	1.87
1451538_at	gb:BC024958.1 /DB_XREF=gi:19353855 /FEA=FL	BC024958		1.87
1416623_at	thrombospondin 3	NM_013691	21827	1.86
1416625_at	serine (or cysteine) proteinase inhibitor, clade G, m	NM_009776	12258	1.86
1417359_at	microfibrillar-associated protein 2	NM_008546	17150	1.86
1417439_at	CD248 antigen, endosialin	NM_054042	70445	1.86
1419569_a_at	interferon-stimulated protein	BC022751	57444	1.86
1420671_x_at	membrane-spanning 4-domains, subfamily A, mem	NM_029499	64380	1.86
1423077_at	sorting nexin 9	BB703513	66616	1.86
1424956_at	RIKEN cDNA D030015G18 gene	BC019130	230793	1.86
1425039_at	integrin, beta-like 1	BC020152	223272	1.86
1427027_a_at	general transcription factor III A	AV173739	66596	1.86
1431105_a_at	transmembrane protein 33	AK012413	67878	1.86
1434881_s_at	potassium channel tetramerisation domain containi	BM220945	239217	1.86
1436038_a_at	Down syndrome critical region homolog 5 (human)	BF318238	56176	1.86
1439064_at	cDNA sequence BC030046	AV233866	277154	1.86
1451074_at	gb:AF037205.1 /DB_XREF=gi:2746334 /GEN=Rzf	AF037205		1.86
1415756_a_at	SNAP-associated protein	BB667523	20615	1.85
1417298_at	emopamil binding protein-like	BC027422	68177	1.85
1420817_at	3-monooxygenase/tryptophan 5-monooxygenase acti	NM_018871	22628	1.85
1421596_s_at	histocompatibility 28	NM_031367	15061	1.85
1423041_a_at	basic leucine zipper and W2 domains 1	AV144956	66882	1.85
1429144_at	RIKEN cDNA 2310032D16 gene	AV291259	74182	1.85
1438398_at	RIKEN cDNA B330012G18 gene	BB203348	170791	1.85
1452782_a_at	thioredoxin 2	AK002358	56551	1.85
1416942_at	type 1 tumor necrosis factor receptor shedding ami	NM_030711	80898	1.84
1417442_a_at	peroxisomal biogenesis factor 3	AK012827	56535	1.84
1426312_at	brain and reproductive organ-expressed protein	BE984258	107976	1.84
1429246_a_at	annexin A6	AK013026	11749	1.84
1448865_at	hydroxysteroid (17-beta) dehydrogenase 7	NM_010476	15490	1.84
1449299_at	low density lipoprotein receptor-related protein 5	BC011374	16973	1.84
1419586_at	retinitis pigmentosa 2 homolog (human)	BF168611	19889	1.83
1423331_a_at	poliovirus receptor-related 3	BG070704	58998	1.83
1424180_a_at	thyroid hormone receptor associated protein 4	BC005409	23989	1.83
1424775_at	2'-5' oligoadenylate synthetase 1A	BC018470	246730	1.83
1426774_at	gb:BM227980 /DB_XREF=gi:17790458 /DB_XREF	BM227980		1.83
1428708_x_at	RIKEN cDNA 2610009E16 gene	AK011360	69202	1.83
1448332_at	peroxisome biogenesis factor 19	NM_023041	19298	1.83
1448384_at	gb:BC018194.1 /DB_XREF=gi:17390433 /FEA=FL	BC018194		1.83
1415935_at	SPARC related modular calcium binding 2	NM_022315	64074	1.82
1416319_at	adenosine kinase	NM_134079	11534	1.82



1417753_at	polycystic kidney disease 2	AF014010	18764	1.82
1417821_at	DNA segment, Chr 17, human D6S56E 5	NM_033075	110956	1.82
1419483_at	complement component 3a receptor 1	NM_009779	12267	1.82
1421063_s_at	small nuclear ribonucleoprotein N /// SNRPN upstre	NM_033174	20646 /// 84	1.82
1423545_a_at	zinc finger protein 207	BB633247	22680	1.82
1429859_a_at	ADP-ribosylation factor-like 2 binding protein	AK006418	107566	1.82
1448235_s_at	high mobility group box 1 /// RIKEN cDNA 4932431	BF166000	114675 /// 1	1.82
1451789_a_at	receptor-like tyrosine kinase	M98547	20187	1.82
1416213_x_at	surfeit gene 4	BF122782	20932	1.81
1416897_at	poly (ADP-ribose) polymerase family, member 9	NM_030253	80285	1.81
1418536_at	Histocompatibility 2, Q region locus 6	M29881	15018	1.81
1422782_s_at	toll-like receptor 3	NM_126166	142980	1.81
1423840_at	DNA segment, Chr 11, ERATO Doi 99, expressed	BC026206	52469	1.81
1426830_a_at	S-adenosylhomocysteine hydrolase-like 1	BB831090	229709	1.81
1426947_x_at	procollagen, type VI, alpha 2	BI455189	12834	1.81
1427682_a_at	early growth response 2	X06746	13654	1.81
1448591_at	cathepsin S	NM_021281	13040	1.81
1449018_at	profilin 1	NM_011072	18643	1.81
1424007_at	growth differentiation factor 10	L42114	14560	1.8
1424874_a_at	polypyrimidine tract binding protein 1	BC007472	19205	1.8
1426880_at	CDNA sequence BC026657	BM250266	208618	1.8
1450716_at	a disintegrin-like and metalloprotease (reprolysin ty	D67076	11504	1.8
1452037_at	mannoside acetylglucosaminyltransferase 2	AI481328	217664	1.8
1452426_x_at	gb:BC004065.1 /DB_XREF=gi:13278542 /FEA=mF	BC004065		1.8
1416619_at	RIKEN cDNA 4632428N05 gene	BC003967	74048	1.79
1418666_at	pentaxin related gene	NM_008987	19288	1.79
1419037_at	casein kinase II, alpha 1 polypeptide	BB283759	12995	1.79
1423259_at	inhibitor of DNA binding 4	BB121406	15904	1.79
1423653_at	ATPase, Na+/K+ transporting, alpha 1 polypeptide	BC025618	11928	1.79
1424040_at	cDNA sequence BC019977	BC019977	245877	1.79
1424183_at	acetyl-Coenzyme A acetyltransferase 1	BG070487	110446	1.79
1425610_s_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide l	AF348968	108148	1.79
1427762_x_at	histone 1, H2bp	M25487	319188	1.79
1431602_a_at	RIKEN cDNA 1810064L21 gene	AK007746	66435	1.79
1433963_a_at	cDNA sequence BC032204	BG066664	108101	1.79
1434036_at	Metastasis suppressor 1	AV024771	211401	1.79
1451672_at	G protein-coupled receptor kinase 6	AF040748	26385	1.79
1452363_a_at	ATPase, Ca++ transporting, cardiac muscle, slow t	AA245637	11938	1.79
1416369_at	RIKEN cDNA 5730414C17 gene	NM_133680	66631	1.78
1417946_at	abhydrolase domain containing 3	NM_134130	106861	1.78
1419153_at	RIKEN cDNA 2810417H13 gene	AK017673	68026	1.78
1420401_a_at	receptor (calcitonin) activity modifying protein 3	NM_019511	56089	1.78
1421487_a_at	non-catalytic region of tyrosine kinase adaptor prot	NM_010878	17973	1.78
1421546_a_at	Rac GTPase-activating protein 1	NM_012025	26934	1.78
1423493_a_at	nuclear factor I/X	BB315728	18032	1.78
1424669_at	zinc finger, FYVE domain containing 21	BC019521	68520	1.78
1444028_s_at	dedicator of cytokinesis 9	AA410148	105445	1.78
1449175_at	G-protein coupled receptor 65	NM_008152	14744	1.78
1450920_at	cyclin B2	AK013312	12442	1.78
1452490_a_at	adaptor protein complex AP-2, alpha 2 subunit	BC010597	11772	1.78
1453367_a_at	RIKEN cDNA 6330583M11 gene	BB178770	76192	1.78
1454120_a_at	ring finger protein 134	AK016709	71041	1.78
1417423_at	glutamate receptor, ionotropic, N-methyl D-asparat	NM_023168	66168	1.77

1420688_a_at	sarcoglycan, epsilon	NM_011360	20392	1.77
1421424_a_at	alanyl (membrane) aminopeptidase	NM_008486	16790	1.77
1422504_at	glycine receptor, beta subunit	NM_010298	14658	1.77
1423548_s_at	serologically defined breast cancer antigen 84	BB556862	66366	1.77
1423756_s_at	insulin-like growth factor binding protein 4	BC019836	16010	1.77
1423984_a_at	myeloid ecotropic viral integration site-related gene	BC003762	17537	1.77
1426528_at	neuropilin 2	BQ176723	18187	1.77
1435306_a_at	kinesin family member 11	BM234447	16551	1.77
1448196_at	methionine adenosyltransferase II, beta	NM_134017	108645	1.77
1448421_s_at	asporin	NM_025711	66695	1.77
1448847_at	gb:BF139489 /DB_XREF=gi:10978529 /DB_XREF:	NM_012032		1.77
1452268_at	RIKEN cDNA 2810485I05 gene	AV337692	72826	1.77
1452780_at	gb:AK008582.1 /DB_XREF=gi:12842855 /FEA=mF	AV297256		1.77
1455482_at	adaptor protein complex AP-2, alpha 2 subunit	BQ175609	11772	1.77
1416741_at	procollagen, type V, alpha 1	AW744319	12831	1.76
1416985_at	gb:D87968.1 /DB_XREF=gi:1864014 /FEA=FLmR	AB018194		1.76
1423547_at	lysozyme	AW208566	17105	1.76
1426368_at	gb:AK014548.1 /DB_XREF=gi:12852473 /FEA=FL	AK014548		1.76
1449093_at	cardiotrophin 1	NM_007795	13019	1.76
1452014_a_at	insulin-like growth factor 1	AF440694	16000	1.76
1452534_a_at	high mobility group box 2	X67668	97165	1.76
1460692_at	HLA-B associated transcript 8	BI412952	110147	1.76
1417819_at	torsin family 1, member B	BB004887	30934	1.75
1418061_at	latent transforming growth factor beta binding prote	NM_013589	16997	1.75
1419435_at	aldehyde oxidase 1	NM_009676	11761	1.75
1423298_at	adducin 3 (gamma)	BM239842	27360	1.75
1424225_at	ankyrin repeat and SOCS box-containing protein 8	AF398969	78541	1.75
1424849_at	RIKEN cDNA 2310038K02 gene	BC026444	233064	1.75
1425190_a_at	RIKEN cDNA 1700048E23 gene	BC025612	73373	1.75
1425548_a_at	leukocyte specific transcript 1	U72644	16988	1.75
1426576_at	transmembrane protein 23	BE629162	208449	1.75
1426801_at	septin 8	BB756379	20362	1.75
1426906_at	interferon activated gene 203	M74124	15950	1.75
1426948_at	translocated promoter region	BM214109	108989	1.75
1427441_a_at	succinate-Coenzyme A ligase, GDP-forming, beta s	BF608645	20917	1.75
1427705_a_at	nuclear factor of kappa light chain gene enhancer ii	L28118	18033	1.75
1428904_at	RIKEN cDNA E230022H04 gene	AK017359	225339	1.75
1433678_at	expressed sequence AI132321	BB210623	104759	1.75
1435737_a_at	nuclear distribution gene E homolog 1 (A nidulans)	BQ032232	67203	1.75
1436528_at	Kazal-type serine protease inhibitor domain 1	AI842353	107250	1.75
1448491_at	enoyl coenzyme A hydratase 1, peroxisomal	NM_016772	51798	1.75
1448710_at	chemokine (C-X-C motif) receptor 4	D87747	12767	1.75
1449025_at	interferon-induced protein with tetratricopeptide rep	NM_010501	15959	1.75
1453913_a_at	transporter 2, ATP-binding cassette, sub-family B (N	BE691515	21355	1.75
1460205_at	RIKEN cDNA 6720485C15 gene	NM_026551	68087	1.75
1416032_at	RIKEN cDNA 1110006I15 gene	AW494906	68539	1.74
1417424_at	immediate early response 3 interacting protein 1	BE287896	66191	1.74
1418460_at	SH3 domain protein D19	NM_012059	27059	1.74
1419637_s_at	RIKEN cDNA 4833420G17 gene	BG067119	67392	1.74
1425025_at	cDNA sequence BC022145	BC022145	217203	1.74
1425466_at	SUMO/sentrin specific protease 2	AV317107	75826	1.74
1426580_at	polo-like kinase 4 (Drosophila)	BB706079	20873	1.74
1427139_at	a disintegrin-like and metalloprotease (reprolysin ty	BG064835	224697	1.74

1432042_a_at	smu-1 suppressor of mec-8 and unc-52 homolog (C	AK011140	74255	1.74
1433645_at	RIKEN cDNA 2210409B22 gene	BM207133	70174	1.74
1437052_s_at	solute carrier family 2 (facilitated glucose transport	BB414515	20527	1.74
1450059_at	gb:BG072083 /DB_XREF=gi:12554652 /DB_XREF	BG072083		1.74
1452250_a_at	procollagen, type VI, alpha 2	BI455189	12834	1.74
1452770_at	vitamin K epoxide reductase complex, subunit 1	AK003237	27973	1.74
1416629_at	solute carrier family 1 (neutral amino acid transport	NM_009201	20514	1.73
1417183_at	DnaJ (Hsp40) homolog, subfamily A, member 2	C77509	56445	1.73
1418392_a_at	guanylate nucleotide binding protein 4	NM_018734	55932	1.73
1418440_at	procollagen, type VIII, alpha 1	NM_007739	12837	1.73
1419324_at	LIM homeobox protein 9	NM_010714	16876	1.73
1422795_at	cullin 3	AV273804	26554	1.73
1423852_at	RIKEN cDNA 9430059P22 gene	BC024118	219134	1.73
1424131_at	procollagen, type VI, alpha 3	AF064749	12835	1.73
1426025_s_at	lysosomal-associated protein transmembrane 5	U29539	16792	1.73
1427471_at	F-box and leucine-rich repeat protein 3	BF782863	50789	1.73
1427739_a_at	transformation related protein 53	AJ297973	22059	1.73
1429775_a_at	transmembrane 7 superfamily member 1	AK009736	83924	1.73
1434379_at	RIKEN cDNA 2810410A03 gene	BG868949	17122	1.73
1448876_at	Ellis van Creveld gene homolog (human)	NM_021292	59056	1.73
1451080_at	ubiquitin specific protease 1	BC018179	230484	1.73
1452721_a_at	RIKEN cDNA 2900091E11 gene	BM210204	67282	1.73
1416782_s_at	PRA1 domain family 2	NM_138602	54637	1.72
1417326_a_at	anaphase promoting complex subunit 11 homolog (	NM_025389	66156	1.72
1419598_at	membrane-spanning 4-domains, subfamily A, mem	NM_026835	68774	1.72
1419667_at	sarcoglycan, beta (dystrophin-associated glycoprot	AK014381	24051	1.72
1420821_at	sphingosine-1-phosphate phosphatase 1	NM_030750	81535	1.72
1421891_at	ST3 beta-galactoside alpha-2,3-sialyltransferase 2	BB750118	20444	1.72
1423314_s_at	phosphodiesterase 7A	BG070255	18583	1.72
1423478_at	protein kinase C, beta 1	BF660388	18751	1.72
1423743_at	archain 1	BC017124	213827	1.72
1425913_a_at	RIKEN cDNA 2810022L02 gene	BC014764	67198	1.72
1426963_at	phosphofurin acidic cluster sorting protein 1-like	BI412345	217893	1.72
1427318_s_at	dysferlin /// fer-1-like 3, myoferlin (C. elegans)	BI555209	226101 /// 2	1.72
1429947_a_at	Z-DNA binding protein 1	AK008179	58203	1.72
1431030_a_at	ring finger protein 14	AK010162	56736	1.72
1448775_at	interferon activated gene 203	NM_008328	15950	1.72
1449154_at	procollagen, type XI, alpha 1	NM_007729	12814	1.72
1449531_at	leprecan-like 2	BC003726	14789	1.72
1450877_at	cDNA sequence BC005537	BI155782	79555	1.72
1453299_a_at	purine-nucleoside phosphorylase	AK008143	18950	1.72
1417244_a_at	interferon regulatory factor 7	NM_016850	54123	1.71
1418240_at	guanylate nucleotide binding protein 2	NM_010260	14469	1.71
1419225_at	calcium channel, voltage dependent, alpha2/delta s	NM_009785	12294	1.71
1421116_a_at	reticulin 4	NM_024226	68585	1.71
1423499_at	synuclein, alpha interacting protein (synphilin)	AK017012	67847	1.71
1423790_at	death-associated protein	BC024876	223453	1.71
1426766_at	RIKEN cDNA 6330403K07 gene	AK018106	103712	1.71
1427076_at	macrophage expressed gene 1	L20315	17476	1.71
1427884_at	procollagen, type III, alpha 1	AW550625	12825	1.71
1428187_at	CD47 antigen (Rh-related antigen, integrin-associa	AK018679	16423	1.71
1428699_at	spermine synthase	AW744143	20603	1.71
1438716_at	tripartite motif protein 30-like	BG068242	209387	1.71

1448148_at	granulin	M86736	14824	1.71
1448192_s_at	phosphoribosyl pyrophosphate synthetase 1	AK011304	19139	1.71
1449049_at	toll-like receptor 1	AF316985	21897	1.71
1450405_at	mitochondrial ribosomal protein L19	BB041267	56284	1.71
1451091_at	thioredoxin domain containing 5	BC016252	105245	1.71
1451332_at	zinc finger protein 521	BC021376	225207	1.71
1452540_a_at	histone 1, H2bc /// histone 1, H2be /// histone 1, H2M	M25487	319179 /// 3	1.71
1416203_at	aquaporin 1	NM_007472	11826	1.7
1417018_at	epidermal growth factor-containing fibulin-like extra	NM_021474	58859	1.7
1423418_at	farnesyl diphosphate synthetase	BI247584	110196	1.7
1423689_a_at	G-protein signalling modulator 1 (AGS3-like, C. ele	BC026486	67839	1.7
1425120_x_at	RIKEN cDNA 1810023F06 gene	BC021795	217845	1.7
1427548_a_at	chloride channel, nucleotide-sensitive, 1A	U53455	12729	1.7
1428115_a_at	RAB2B, member RAS oncogene family	AK005230	76338	1.7
1432332_a_at	DNA segment, Chr 7, Roswell Park 2 complex, exp	AK008430	110959	1.7
1435504_at	Restin-like 2	BM217861	78785	1.7
1438480_a_at	thymocyte protein thy28	AV301675	77862	1.7
1448119_at	2,3-bisphosphoglycerate mutase	NM_007563	12183	1.7
1449264_at	synaptotagmin 11	BC025207	229521	1.7
1451047_at	integral membrane protein 2A	BI966443	16431	1.7
1452105_a_at	tuberous sclerosis 2	BB759096	22084	1.7
1452384_at	ectonucleotide pyrophosphatase/phosphodiesteras	AV224446	209558	1.7
1452954_at	ubiquitin-conjugating enzyme E2C	AV162459	68612	1.7
1453740_a_at	cyclin L2	BI112766	56036	1.7
1417148_at	platelet derived growth factor receptor, beta polype	NM_008809	18596	1.69
1419873_s_at	colony stimulating factor 1 receptor	AI323359	12978	1.69
1422619_at	gb:NM_008903.1 /DB_XREF=gi:6679430 /GEN=Pf	NM_008903		1.69
1423757_x_at	insulin-like growth factor binding protein 4	BC019836	16010	1.69
1423774_a_at	protein regulator of cytokinesis 1	BC005475	233406	1.69
1424498_at	RIKEN cDNA 5730596K20 gene	BC017138	109168	1.69
1425496_at	ATP-binding cassette, sub-family A (ABC1), membe	AK007703	27410	1.69
1425951_a_at	C-type lectin domain family 4, member n	AF240358	56620	1.69
1426208_x_at	pleiomorphic adenoma gene-like 1	AF147785	22634	1.69
1426652_at	minichromosome maintenance deficient 3 (S. cerev	BI658327	17215	1.69
1434695_at	RIKEN cDNA 2810047L02 gene	AV270035	76843	1.69
1435375_at	cDNA sequence BC052328	BM224662	223433	1.69
1448336_at	developmentally regulated GTP binding protein 1	NM_007879	13494	1.69
1449195_s_at	chemokine (C-X-C motif) ligand 16	BC019961	66102	1.69
1452328_s_at	gb:BF160731 /DB_XREF=gi:11040838 /DB_XREF=	BF160731		1.69
1456741_s_at	glycoprotein m6a	BB348674	234267	1.69
1415907_at	cyclin D3	NM_007632	12445	1.68
1416016_at	gb:BC024897.1 /DB_XREF=gi:19352971 /FEA=FL	AW048052		1.68
1418439_at	DNA segment, Chr 10, ERATO Doi 322, expressed	NM_026065	67270	1.68
1419182_at	polydomain protein	NM_022814	64817	1.68
1422243_at	fibroblast growth factor 7	NM_008008	14178	1.68
1422453_at	pre-mRNA processing factor 8	NM_138659	192159	1.68
1422563_at	complement receptor related protein	NM_013499	12946	1.68
1424046_at	budding uninhibited by benzimidazoles 1 homolog (	AF002823	12235	1.68
1424732_s_at	RIKEN cDNA 3110005G23 gene	BC020157	73067	1.68
1425281_a_at	delta sleep inducing peptide, immunoreactor	AF201289	14605	1.68
1425434_a_at	macrophage scavenger receptor 1	L04274	20288	1.68
1425492_at	bone morphogenetic protein receptor, type 1A	BM939768	12166	1.68
1425654_a_at	DNA segment, Chr 15, ERATO Doi 366, expressed	AF307845	65970	1.68

1426981_at	paired basic amino acid cleaving system 4	BI157485	18553	1.68
1427680_a_at	nuclear factor I/B	Y07687	18028	1.68
1438009_at	histone 1, H2ad	W91024	319165	1.68
1448648_at	RIKEN cDNA 9130005N14 gene	NM_026667	68303	1.68
1450106_a_at	Ena-vasodilator stimulated phosphoprotein	NM_007965	14026	1.68
1450454_at	torsin family 3, member A	NM_023141	30935	1.68
1452154_at	gb:BE915338 /DB_XREF=gi:10414866 /DB_XREF: BE915338			1.68
1460672_at	RIKEN cDNA 2410002F23 gene	BC016099	66976	1.68
1416818_at	parvin, alpha	NM_020606	57342	1.67
1416904_at	muscleblind-like 1 (Drosophila)	NM_020007	56758	1.67
1419313_at	cyclin T1	NM_009833	12455	1.67
1419668_at	sarcoglycan, beta (dystrophin-associated glycoprot	AK014381	24051	1.67
1422601_at	serine (or cysteine) proteinase inhibitor, clade B, m	NM_009256	20723	1.67
1422736_at	RAN binding protein 9	NM_019930	56705	1.67
1423800_at	aspartyl-tRNA synthetase	BC008638	226414	1.67
1428301_at	RIKEN cDNA 2610042L04 gene	BM195235	67055	1.67
1428787_at	hematopoietic protein 1	BM238906	105855	1.67
1434380_at	Diabetic nephropathy-like protein (Dnr12) mRNA, p	BM241271		1.67
1448524_s_at	signal sequence receptor, delta	NM_009279	20832	1.67
1450431_a_at	gb:D85414.1 /DB_XREF=gi:1374781 /GEN=Nedd4	NM_010890		1.67
1450724_at	down-regulated by Ctnnb1, a	NM_053090	84652	1.67
1451161_a_at	EGF-like module containing, mucin-like, hormone r	U66888	13733	1.67
1452173_at	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoac	AW107842	97212	1.67
1416527_at	RAB32, member RAS oncogene family	NM_026405	67844	1.66
1416906_at	anaphase-promoting complex subunit 5	NM_021505	59008	1.66
1417261_at	mbt domain containing 1	NM_134012	103537	1.66
1419821_s_at	gb:A1788952 /DB_XREF=gi:5336668 /DB_XREF=u	A1788952		1.66
1420624_a_at	vesicle-associated membrane protein 8	NM_016794	22320	1.66
1421034_a_at	interleukin 4 receptor, alpha	NM_010557	16190	1.66
1422521_at	dynactin 1	NM_007835	13191	1.66
1422972_s_at	GCN5 general control of amino acid synthesis-like :	NM_020004	14534	1.66
1425391_a_at	oxysterol binding protein-like 5	AB074008	79196	1.66
1425658_at	CD109 antigen	AY083458	235505	1.66
1426160_a_at	serine/threonine kinase 16	AF062076	20872	1.66
1427771_x_at	integrin beta 1 (fibronectin receptor beta)	U37029	16412	1.66
1428179_at	NADH dehydrogenase (ubiquinone) flavoprotein 2	BI692577	72900	1.66
1435477_s_at	Fc receptor, IgG, low affinity IIb	BM224327	14130	1.66
1450065_at	adenylate cyclase 7	NM_007406	11513	1.66
1415972_at	myristoylated alanine rich protein kinase C substrat	AW546141	17118	1.65
1416105_at	nicotinamide nucleotide transhydrogenase	BC008518	18115	1.65
1416301_a_at	early B-cell factor 1	BB125261	13591	1.65
1418629_a_at	gb:BE447456 /DB_XREF=gi:9447033 /DB_XREF=	NM_011317		1.65
1421819_a_at	SET translocation	BF134272	56086	1.65
1421861_at	calsyntenin 1	BG065300	65945	1.65
1421916_at	platelet derived growth factor receptor, alpha polyp	AW537708	18595	1.65
1423382_a_at	heterogeneous nuclear ribonucleoprotein F	AW412441	98758	1.65
1425048_a_at	high mobility group box 1	U00431	15289	1.65
1426640_s_at	tribbles homolog 2 (Drosophila)	BB354684	217410	1.65
1428870_at	gb:BB093407 /DB_XREF=gi:15407802 /DB_XREF: BM213850			1.65
1432029_a_at	stromal membrane-associated protein 1	AK006817	98366	1.65
1448525_a_at	BCL2/adenovirus E1B 19kDa-interacting protein 3-l	AK018668	12177	1.65
1448619_at	7-dehydrocholesterol reductase	NM_007856	13360	1.65
1449209_a_at	retinol dehydrogenase 11	AB030503	17252	1.65

1449334_at	tissue inhibitor of metalloproteinase 3	BI111620	21859	1.65
1450380_at	ependymin related protein 2 (zebrafish)	AF353717	105298	1.65
1450663_at	thrombospondin 2	NM_011581	21826	1.65
1450997_at	serine/threonine kinase 17b (apoptosis-inducing)	AV173139	98267	1.65
1452279_at	properdin factor, complement	BB800282	18636	1.65
1454023_a_at	DNA segment, Chr 1, Brigham & Women's Genetic	AK006466	74241	1.65
1454161_s_at	RIKEN cDNA 0610007P14 gene	AK002308	58520	1.65
1415836_at	pyrroline-5-carboxylate synthetase (glutamate gam	NM_019698	56454	1.64
1416309_at	nucleolar and spindle associated protein 1	BC009096	108907	1.64
1417947_at	proliferating cell nuclear antigen	BC010343	18538	1.64
1418340_at	Fc receptor, IgE, high affinity I, gamma polypeptide	NM_010185	14127	1.64
1420634_a_at	MAD homolog 2 (Drosophila)	NM_010754	17126	1.64
1423111_at	ATP synthase, H+ transporting, mitochondrial F1 cc	BM213627	11946	1.64
1423463_a_at	DNA segment, Chr 2, ERATO Doi 750, expressed	AK012148	51944	1.64
1424321_at	replication factor C (activator 1) 4	BC003335	106344	1.64
1424921_at	bone marrow stromal cell antigen 2	BC008532	69550	1.64
1425483_at	thymocyte selection-associated HMG box gene	BB547854	252838	1.64
1425491_at	bone morphogenetic protein receptor, type 1A	BM939768	12166	1.64
1426088_at	gb:BC004015.1 /DB_XREF=gi:13278405 /FEA=FL	BC004015		1.64
1426349_s_at	thymopoietin	AA153892	21917	1.64
1426831_at	S-adenosylhomocysteine hydrolase-like 1	BB831090	229709	1.64
1433504_at	brain glycogen phosphorylase	AW547988	110078	1.64
1434900_at	MKL (megakaryoblastic leukemia)/myocardin-like 1	BM196656	223701	1.64
1438211_s_at	D site albumin promoter binding protein	BB550183	13170	1.64
1448883_at	legumain	NM_011175	19141	1.64
1451134_a_at	RIKEN cDNA 2410018G23 gene	BC026789	69742	1.64
1453251_at	DEAH (Asp-Glu-Ala-His) box polypeptide 30	AK013413	72831	1.64
1460194_at	phytanoyl-CoA hydroxylase	NM_010726	16922	1.64
1460460_a_at	golgi reassembly stacking protein 2	AK010664	70231	1.64
1415773_at	nucleolin	BF118393	17975	1.63
1415977_at	myo-inositol 1-phosphate synthase A1	NM_023627	71780	1.63
1415983_at	lymphocyte cytosolic protein 1	NM_008879	18826	1.63
1416691_at	GTP binding protein 2	NM_019581	56055	1.63
1416732_at	topoisomerase (DNA) II beta	BB166592	21974	1.63
1417399_at	growth arrest specific 6	NM_019521	14456	1.63
1417534_at	gb:NM_010580.1 /DB_XREF=gi:6754379 /GEN=Itc	NM_010580		1.63
1417870_x_at	cathepsin Z	NM_022325	64138	1.63
1417963_at	phospholipid transfer protein	NM_011125	18830	1.63
1421009_at	radical S-adenosyl methionine domain containing 2	BB741897	58185	1.63
1421033_a_at	transcription elongation regulator 1 (CA150)	AW046403	56070	1.63
1424906_at	RIKEN cDNA E030024M05 gene	BC025220	217430	1.63
1426532_at	zinc finger, MYND domain containing 11	BB832996	66505	1.63
1427161_at	centromere autoantigen F	BE848253	108000	1.63
1427212_at	mitogen-activated protein kinase associated proteir	BC027377	227743	1.63
1430780_a_at	phosphomannomutase 1	BI739353	29858	1.63
1438808_at	gb:BB450072 /DB_XREF=gi:16425416 /DB_XREF:	BB450072		1.63
1450681_at	zinc finger protein 143	NM_009281	20841	1.63
1452046_a_at	protein phosphatase 1, catalytic subunit, gamma is	BG071790	19047	1.63
1452079_s_at	testis derived transcript 3, pseudogene /// testis der	BE851193	114893 /// 5	1.63
1452314_at	kinesin family member 11	BB827235	16551	1.63
1453473_a_at	t-complex testis expressed 1	BG094946	21648	1.63
1454631_at	RIKEN cDNA 6330549H03 gene	BI689897	108667	1.63
1455898_x_at	solute carrier family 2 (facilitated glucose transport	BB414515	20527	1.63

1459909_at	Nuclear factor I/X	BI249259	18032	1.63
1460555_at	RIKEN cDNA 6330500D04 gene	BM242294	193385	1.63
1415980_at	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 c	NM_026468	67942	1.62
1416006_at	midkine	M34328	17242	1.62
1416474_at	neighbor of Punc E11	NM_020043	56741	1.62
1417960_at	cytoplasmic polyadenylation element binding protei	NM_007755	12877	1.62
1418049_at	latent transforming growth factor beta binding prote	NM_008520	16998	1.62
1418456_a_at	chemokine (C-X-C motif) ligand 14	AF252873	57266	1.62
1418529_at	O-sialoglycoprotein endopeptidase	NM_133676	66246	1.62
1423870_at	RIKEN cDNA 2610208M17 gene	BI104583	108909	1.62
1424186_at	RIKEN cDNA 2610001E17 gene	BG074158	67896	1.62
1424410_at	tetratricopeptide repeat domain 8	BC017523	76260	1.62
1424427_at	DNA segment, Chr 1, ERATO Doi 251, expressed	BC027337	27878	1.62
1424708_at	RIKEN cDNA 1110014C03 gene	BI409239	68581	1.62
1425933_a_at	5'-nucleotidase, cytosolic II	BC006028	76952	1.62
1434801_x_at	solute carrier family 25 (mitochondrial carrier, aden	AV026569	11740	1.62
1435986_x_at	succinate dehydrogenase complex, subunit C, inte	AW107712	66052	1.62
1436058_at	radical S-adenosyl methionine domain containing 2	BB132493	58185	1.62
1448632_at	proteasome (prosome, macropain) subunit, beta ty	NM_013640	19171	1.62
1448910_at	peroxisomal trans-2-enoyl-CoA reductase	NM_023523	111175	1.62
1450420_at	stromal antigen 1	BQ174473	20842	1.62
1451025_at	ADP-ribosylation factor-like 1	AW011908	104303	1.62
1451121_a_at	glioma tumor suppressor candidate region gene 2	BC017637	68077	1.62
1451680_at	neoplastic progression 3	BC011325	76650	1.62
1451981_at	gene trap ROSA b-geo 22	AF303106	110012	1.62
1460718_s_at	mitochondrial carrier homolog 1 (C. elegans)	AF192558	56462	1.62
1416121_at	lysyl oxidase	M65143	16948	1.61
1416132_at	RIKEN cDNA C920006C10 gene	BB188557	76740	1.61
1421083_x_at	barrier to autointegration factor 1	NM_011793	23825	1.61
1421462_a_at	leprecan 1	NM_019783	56401	1.61
1422510_at	CTD (carboxy-terminal domain, RNA polymerase II	NM_133710	69274	1.61
1423098_at	calpain 7	BQ257745	12339	1.61
1423316_at	transmembrane protein 39a	AK017817	67846	1.61
1423495_at	2-4-dienoyl-Coenzyme A reductase 2, peroxisomal	BE952632	26378	1.61
1426298_at	Iroquois related homeobox 2 (Drosophila)	AF295369	16372	1.61
1427298_at	neoplastic progression 1	BB542096	18152	1.61
1427516_a_at	biregional cell adhesion molecule-related/down-reg	AF388037	117606	1.61
1433803_at	Janus kinase 1	BQ032637	16451	1.61
1435862_at	Son cell proliferation protein	BG067046	20658	1.61
1438223_at	gb:BG065705 /DB_XREF=gi:12548268 /DB_XREF	BG065705		1.61
1449126_at	zinc finger protein 90	NM_011764	22751	1.61
1450268_at	fidgetin	NM_021716	60344	1.61
1450478_a_at	protein tyrosine phosphatase, non-receptor type 12	X63440	19248	1.61
1450738_at	gb:NM_016705.1 /DB_XREF=gi:7710045 /GEN=Ki	NM_016705		1.61
1451358_a_at	Rac GTPase-activating protein 1	AF212320	26934	1.61
1451564_at	poly (ADP-ribose) polymerase family, member 14	BC021340	432411	1.61
1452829_at	carbamoyl-phosphate synthetase 2, aspartate trans	AK010453	69719	1.61
1456651_a_at	translocated promoter region	BG067858	108989	1.61
1460317_s_at	Guanine nucleotide binding protein, alpha 13	BI662324	14674	1.61
1415792_at	expressed sequence AL033326	NM_019705	24105	1.6
1416695_at	benzodiazepine receptor, peripheral	BC002055	12257	1.6
1417056_at	proteasome (prosome, macropain) 28 subunit, alph	NM_011189	19186	1.6
1423262_a_at	H3 histone, family 3A	BI111967	15078	1.6

1423423_at	glucose regulated protein	BF319868	14827	1.6
1423452_at	serine/threonine kinase 17b (apoptosis-inducing)	AV173139	98267	1.6
1423986_a_at	scotin gene	BC010238	66940	1.6
1425348_a_at	signal recognition particle receptor, B subunit	BC003798	20818	1.6
1426555_at	serine carboxypeptidase 1	AK014680	74617	1.6
1434920_a_at	Ena-vasodilator stimulated phosphoprotein	AW553781	14026	1.6
1435321_at	RIKEN cDNA 3732412D22 gene	BM117827	77569	1.6
1451395_at	DNA segment, Chr 5, Brigham & Women's Genetic	BC021492	215707	1.6
1452428_a_at	beta-2 microglobulin	AI099111	12010	1.6
1454694_a_at	topoisomerase (DNA) II alpha	BM211413	21973	1.6
1415908_at	testis-specific protein, Y-encoded-like 1	AF042180	22110	1.59
1416076_at	cyclin B1	NM_007629	268697	1.59
1416096_at	expressed sequence AI413782	NM_134044	104799	1.59
1416102_at	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase	AV027921	22631	1.59
1416176_at	high mobility group box 1	BF166000	15289	1.59
1416343_a_at	lysosomal membrane glycoprotein 2	NM_010685	16784	1.59
1416539_at	yolk sac gene 2	NM_011734	22619	1.59
1416726_s_at	ubiquitin-conjugating enzyme E2S	NM_133777	77891	1.59
1417730_at	exostoses (multiple) 1	NM_010162	14042	1.59
1418326_at	solute carrier family 7 (cationic amino acid transporter)	BC026131	20539	1.59
1418892_at	ras homolog gene family, member J	AF309564	80837	1.59
1418896_a_at	ribophorin II	NM_019642	20014	1.59
1423600_a_at	Treacher Collins Franceschetti syndrome 1, homolog	AW209012	21453	1.59
1424128_x_at	aurora kinase B	BC003261	20877	1.59
1424648_at	RAB, member of RAS oncogene family-like 4	BC017514	67042	1.59
1425529_s_at	DNA segment, Chr 19, Wayne State University 162	BC026369	226178	1.59
1425702_a_at	ectonucleotide pyrophosphatase/phosphodiesterase	BC011294	83965	1.59
1426434_at	transmembrane protein 43	AI747289	74122	1.59
1427050_at	RIKEN cDNA 5730420B22 gene	BC027108	70561	1.59
1428531_at	RIKEN cDNA 5930412E23 gene	BB457797	77065	1.59
1434054_at	v-maf musculoaponeurotic fibrosarcoma oncogene	BB342453	17134	1.59
1448372_a_at	transmembrane protein 4	NM_019953	56530	1.59
1448777_at	minichromosome maintenance deficient 2 mitotin (S)	NM_008564	17216	1.59
1452393_at	expressed sequence AI597013	BB014626	100182	1.59
1452427_s_at	expressed sequence AW742319	BM950003	57874	1.59
1455364_a_at	ribosomal protein S7	AI414989	20115	1.59
1415759_a_at	hepatitis B virus x interacting protein	BC028547	68576	1.58
1415989_at	vascular cell adhesion molecule 1	BB250384	22329	1.58
1418281_at	RAD51 homolog (S. cerevisiae)	NM_011234	19361	1.58
1419451_at	fizzy/cell division cycle 20 related 1 (Drosophila)	BC006616	56371	1.58
1419703_at	procollagen, type V, alpha 3	NM_016919	53867	1.58
1421014_a_at	citrate lyase beta like	BC023398	69634	1.58
1423092_at	inner centromere protein	BI410774	16319	1.58
1423345_at	degenerative spermatocyte homolog 1 (Drosophila)	AV286991	13244	1.58
1426389_at	calcium/calmodulin-dependent protein kinase ID	BG071931	227541	1.58
1426410_at	pyruvate dehydrogenase kinase, isoenzyme 3	AV086243	236900	1.58
1426612_at	timeless interacting protein	AK011357	66131	1.58
1426897_at	RIKEN cDNA 2610510H01 gene	AV122997	108911	1.58
1427148_at	gb:BF160731 /DB_XREF=gj:11040838 /DB_XREF=	BF160731		1.58
1427296_at	cDNA sequence BC010304	BB553912	218236	1.58
1427447_a_at	TRIO and F-actin binding protein	BC003984	110253	1.58
1430527_a_at	ring finger protein 167	AK017523	70510	1.58
1437406_x_at	insulin-like growth factor binding protein 4	BB787243	16010	1.58



1448933_at	protocadherin beta 17	NM_053142	93888	1.58
1449053_s_at	component of oligomeric golgi complex 1	BB210424	16834	1.58
1450241_a_at	ecotropic viral integration site 2a	NM_010161	14017	1.58
1450757_at	cadherin 11	NM_009866	12552	1.58
1450792_at	TYRO protein tyrosine kinase binding protein	NM_011662	22177	1.58
1450892_a_at	ubiquitin specific protease 4 (proto-oncogene)	BF321773	22258	1.58
1450970_at	glutamate oxaloacetate transaminase 1, soluble	AA792094	14718	1.58
1452712_at	heterogeneous nuclear ribonucleoprotein A3	AK011930	229279	1.58
1453107_s_at	forkhead box M1 /// phosphatidylethanolamine bind	AK008037	14235 /// 23	1.58
1454758_a_at	transforming growth factor beta 1 induced transcrip	AU016382	21807	1.58
1416630_at	inhibitor of DNA binding 3	NM_008321	15903	1.57
1416641_at	ligase I, DNA, ATP-dependent	NM_010715	16881	1.57
1418622_at	RAB2, member RAS oncogene family	NM_021518	59021	1.57
1419041_at	DNA segment, Chr 8, Wayne State University 49, e	BC003977	71927	1.57
1419444_at	Sin3-associated polypeptide 18	NM_009119	20220	1.57
1420981_a_at	LIM domain only 4	NM_010723	16911	1.57
1421846_at	WD repeat and SOCS box-containing 2	BM730566	59043	1.57
1422993_s_at	THO complex 4	NM_019484	21681	1.57
1423049_a_at	tropomyosin 1, alpha	AK002271	22003	1.57
1423894_a_at	DALR anticodon binding domain containing 3	BC013483	67789	1.57
1424727_at	chemokine (C-C motif) receptor 5	D83648	12774	1.57
1424885_at	RIKEN cDNA A630065K24 gene	BC011135	213417	1.57
1425665_a_at	signal recognition particle 54	BC005543	24067	1.57
1426254_at	beta-amyloid binding protein precursor	AF353993	94043	1.57
1426397_at	transforming growth factor, beta receptor II	BG793483	21813	1.57
1426573_at	gb:BM235734 /DB_XREF=gi:17871004 /DB_XREF	BM235734		1.57
1427021_s_at	ferritin heavy chain 1 /// dipeptidase 2	AK002778	14319 /// 31	1.57
1427112_at	RIKEN cDNA 2700049H19 gene	AV127581	109008	1.57
1427511_at	Beta-2 microglobulin	AA170322	12010	1.57
1428050_a_at	RIKEN cDNA 0610007H07 gene	BC027637	68212	1.57
1428850_x_at	RIKEN cDNA 2410026K10 gene	AK004342	66478	1.57
1431339_a_at	EF hand domain containing 2	AK007560	27984	1.57
1431349_at	heterogeneous nuclear ribonucleoprotein A/B	AK013709	15384	1.57
1433496_at	glycosyltransferase 25 domain containing 1	AV122321	234407	1.57
1433892_at	sperm associated antigen 5	BM208112	54141	1.57
1437405_a_at	insulin-like growth factor binding protein 4	BB787243	16010	1.57
1448566_at	solute carrier family 40 (iron-regulated transporter),	AF226613	53945	1.57
1449523_at	B-cell CLL/lymphoma 7C	NM_009746	12055	1.57
1450777_at	5'-3' exoribonuclease 2	NM_011917	24128	1.57
1451604_a_at	activin A receptor, type II-like 1	BC014291	11482	1.57
1451742_a_at	UDP-glucose pyrophosphorylase 2	AF424698	216558	1.57
1452731_x_at	RIKEN cDNA 2610042L04 gene	BM195235	67055	1.57
1453744_a_at	RIKEN cDNA 5530600A18 gene	AK017451	71452	1.57
1455176_a_at	gb:BF457392 /DB_XREF=gi:11523561 /DB_XREF=	BF457392		1.57
1456360_at	gb:BI466259 /DB_XREF=gi:15279137 /DB_XREF=	BI466259		1.57
1417141_at	interferon gamma induced GTPase	NM_018738	16145	1.56
1417823_at	glycine C-acetyltransferase (2-amino-3-ketobutyrate)	AK013138	26912	1.56
1418533_s_at	frizzled homolog 2 (Drosophila)	BB371406	57265	1.56
1418758_a_at	pleckstrin homology, Sec7 and coiled-coil domains	NM_011182	19159	1.56
1419872_at	gb:AI323359 /DB_XREF=gi:4057788 /DB_XREF=n	AI323359		1.56
1423130_a_at	splicing factor, arginine/serine-rich 5 (SRp40, HRS)	AW212917	20384	1.56
1423694_at	potassium channel tetramerisation domain containi	BC006935	330171	1.56
1424105_a_at	pituitary tumor-transforming 1	AF069051	30939	1.56

1424965_at	leupaxin	BC026563	107321	1.56
1426372_a_at	blocked early in transport 1 homolog (S. cerevisiae)	BE686988	54399	1.56
1434045_at	Cyclin-dependent kinase inhibitor 1B (P27)	BB354528	12576	1.56
1435335_a_at	basic domain/leucine zipper transcription factor	BG144467	432486	1.56
1438546_x_at	solute carrier family 25 (mitochondrial carrier, adenine nucleotide)	C81442	11740 /// 43	1.56
1448324_at	ribonucleic acid binding protein S1	NM_009070	19826	1.56
1448395_at	secreted frizzled-related sequence protein 1	BI658627	20377	1.56
1448742_at	snail homolog 1 (Drosophila)	NM_011427	20613	1.56
1449009_at	T-cell specific GTPase	NM_011579	21822	1.56
1449047_at	RIKEN cDNA 1600020H07 gene	NM_019975	56794	1.56
1450344_a_at	prostaglandin E receptor 3 (subtype EP3)	NM_011196	19218	1.56
1450496_a_at	RIKEN cDNA 2810433K01 gene	NM_025581	66468	1.56
1451119_a_at	fibulin 1	BC007140	14114	1.56
1451932_a_at	thrombospondin repeat containing 1	BC016215	229595	1.56
1453175_at	zinc finger protein 50	AK013374	109929	1.56
1460288_a_at	protein phosphatase 4, catalytic subunit	NM_019674	56420	1.56
1415776_at	gb:NM_007437.1 /DB_XREF=gi:6680677 /GEN=AI	NM_007437		1.55
1416251_at	minichromosome maintenance deficient 6 (MIS5 homolog)	NM_008567	17219	1.55
1416389_a_at	chromosome condensation 1-like	NM_134083	105670	1.55
1416415_a_at	H2A histone family, member Z	AY074806	51788	1.55
1417694_at	growth factor receptor bound protein 2-associated protein 1	NM_021356	14388	1.55
1417879_at	RIKEN cDNA 1110060M21 gene	NM_025424	66208	1.55
1418580_at	gb:NM_023386.1 /DB_XREF=gi:12963656 /GEN=EMBL	BC024872		1.55
1418627_at	glutamate-cysteine ligase , modifier subunit	NM_008129	14630	1.55
1421385_a_at	myosin VIIa	NM_008663	17921	1.55
1421911_at	signal transducer and activator of transcription 2	AF088862	20847	1.55
1422520_at	neurofilament 3, medium	NM_008691	18040	1.55
1422559_at	ubiquitin-conjugating enzyme E2N	AY039837	93765	1.55
1423484_at	bicaudal C homolog 1 (Drosophila)	BM217996	83675	1.55
1423566_a_at	heat shock protein 105	BI499717	15505	1.55
1424045_at	RIKEN cDNA 5730437N04 gene	BC019420	70544	1.55
1424659_at	slit homolog 2 (Drosophila)	BG963150	20563	1.55
1425810_a_at	cysteine and glycine-rich protein 1	BF124540	13007	1.55
1433711_s_at	Sestrin 1	BG076140	140742	1.55
1433830_at	heterogeneous nuclear ribonucleoprotein A2/B1	BG064117	53379	1.55
1434846_at	RIKEN cDNA 1700065A05 gene	AW557010	329877	1.55
1436172_at	hypothetical protein 9530028C05	BQ175154	330256	1.55
1436874_x_at	solute carrier family 25 (mitochondrial carrier, adenine nucleotide)	BM210336	11740	1.55
1448557_at	RIKEN cDNA 1200015N20 gene	NM_024244	71721	1.55
1449061_a_at	DNA primase, p49 subunit	J04620	19075	1.55
1450020_at	chemokine (C-X3-C) receptor 1	BC012653	13051	1.55
1450632_at	ras homolog gene family, member A	AF178958	11848	1.55
1451253_at	PX domain containing serine/threonine kinase	BC016131	218699	1.55
1451271_a_at	acetyl-Coenzyme A acetyltransferase 1	BG070487	110446	1.55
1452021_a_at	hairy and enhancer of split 6 (Drosophila)	AF247040	55927	1.55
1452024_a_at	LIM domain binding 1	AF030333	16825	1.55
1452080_a_at	testis derived transcript 3, pseudogene	BE851193	54339	1.55
1415873_a_at	ARP1 actin-related protein 1 homolog A (yeast)	NM_016860	54130	1.54
1416186_at	proline-rich nuclear receptor coactivator 2	NM_026383	52830	1.54
1416746_at	H2A histone family, member X	NM_010436	15270	1.54
1417533_a_at	integrin beta 5	NM_010580	16419	1.54
1417728_at	methyl-CpG binding domain protein 3	NM_013595	17192	1.54
1417868_a_at	cathepsin Z	NM_022325	64138	1.54

1419838_s_at	polo-like kinase 4 (Drosophila)	AI385771	20873	1.54
1420922_at	ubiquitin specific protease 9, X chromosome /// sim	AW107303	22284 /// 43	1.54
1421817_at	glutathione reductase 1	AK019177	14782	1.54
1422763_at	regulator of G-protein signaling 19 interacting prote	NM_018771	67903	1.54
1423140_at	lysosomal acid lipase 1	AI596237	16889	1.54
1423479_at	RIKEN cDNA 150002M01 gene	AK005123	68979	1.54
1423681_at	RIKEN cDNA 1300018I05 gene	AK005044	74157	1.54
1423818_a_at	ADP-ribosylation factor-like 6 interacting protein 1	AF133669	54208	1.54
1423940_at	Yip1 interacting factor homolog (S. cerevisiae)	BC011117	68090	1.54
1424033_at	splicing factor, arginine/serine-rich 7	BC014857	225027	1.54
1424382_at	reticulocalbin 3, EF-hand calcium binding domain	BC025602	52377	1.54
1424570_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 46	BF023426	212880	1.54
1424616_s_at	FGF receptor activating protein 1	BG063931	233575	1.54
1425930_a_at	transcription factor-like 4	AF213670	21428	1.54
1427086_at	gb:AK019479.1 /DB_XREF=gi:12859711 /GEN=Sli	BM570006		1.54
1427504_s_at	splicing factor, arginine/serine-rich 2 (SC-35)	AF250133	20382	1.54
1428868_a_at	ornithine decarboxylase antizyme	BG094492	18245	1.54
1432094_a_at	RIKEN cDNA 1700034M03 gene	AK006602	73288	1.54
1433758_at	nischarin	BB025231	64652	1.54
1436567_a_at	NADH dehydrogenase (ubiquinone) 1 alpha subcor	C88880	66416	1.54
1437432_a_at	tripartite motif protein 12	BM244351	76681	1.54
1448314_at	cell division cycle 2 homolog A (S. pombe)	NM_007659	12534	1.54
1448692_at	ubiquilin 4	BB764994	94232	1.54
1449145_a_at	caveolin, caveolae protein 1	AB029929	12389	1.54
1451180_a_at	RIKEN cDNA C330027I04 gene	BC015307	68106	1.54
1452346_at	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosamin	AV032053	108902	1.54
1452545_a_at	integrin beta 1 (fibronectin receptor beta)	U37029	16412	1.54
1454036_a_at	ubiquitin specific protease 15	AK014891	14479	1.54
1415973_at	Myristoylated alanine rich protein kinase C substrat	AW546141	17118	1.53
1416246_a_at	coronin, actin binding protein 1A	BC002136	12721	1.53
1416308_at	UDP-glucose dehydrogenase	NM_009466	22235	1.53
1416558_at	maternal embryonic leucine zipper kinase	NM_010790	17279	1.53
1416701_at	ras homolog gene family, member E	BC009002	74194	1.53
1417189_at	proteasome (prosome, macropain) 28 subunit, beta	NM_011190	19188	1.53
1418015_at	pumilio 2 (Drosophila)	BI689507	80913	1.53
1419165_at	zinc finger protein 260	D45210	26466	1.53
1419548_at	karyopherin (importin) alpha 1	U20619	16646	1.53
1420752_at	deltex 3 homolog (Drosophila)	NM_030714	80904	1.53
1421731_a_at	flap structure specific endonuclease 1	NM_007999	14156	1.53
1421867_at	nuclear receptor subfamily 3, group C, member 1	NM_008173	14815	1.53
1422016_a_at	centromere autoantigen H	BC025084	26886	1.53
1422440_at	cyclin-dependent kinase 4	NM_009870	12567	1.53
1422516_a_at	fibroblast growth factor (acidic) intracellular binding	NM_021438	58249	1.53
1423972_at	electron transferring flavoprotein, alpha polypeptide	BC003432	110842	1.53
1424156_at	retinoblastoma-like 1 (p107)	U27177	19650	1.53
1424443_at	transmembrane 6 superfamily member 1	AV378394	107769	1.53
1425487_at	DNA segment, Chr 11, ERATO Doi 730, expressed	BC025870	193116	1.53
1425927_a_at	activating transcription factor 5	AF375476	107503	1.53
1426999_at	RIKEN cDNA 2700069A02 gene	BM198642	75553	1.53
1427108_at	RIKEN cDNA 9530068E07 gene	BM233467	213673	1.53
1427181_at	DNA segment, Chr 18, ERATO Doi 653, expressed	BQ032363	52662	1.53
1427468_at	protein phosphatase 3, catalytic subunit, beta isofo	M81483	19056	1.53
1428061_at	histone aminotransferase 1	AK014330	107435	1.53

1430542_a_at	solute carrier family 25 (mitochondrial carrier, adenine nucleotide)	AA823938	11740	1.53
1433429_at	phosphatidylinositol glycan, class S	BB309245	276846	1.53
1434089_at	synaptopodin	BB426294	104027	1.53
1448143_at	aldehyde dehydrogenase 2, mitochondrial	NM_009656	11669	1.53
1448258_a_at	signal peptidase complex subunit 1 homolog (S. cerevisiae)	NM_026911	69019	1.53
1448335_s_at	cyclin I	NM_017367	12453	1.53
1450407_a_at	acidic (leucine-rich) nuclear phosphoprotein 32 family A member 1	AF022957	11737	1.53
1450868_at	DNA segment, Chr 8, ERATO Doi 354, expressed in HeLa cells	BF468249	52120	1.53
1451489_at	RIKEN cDNA 1810012H11 gene	BC019996	71998	1.53
1453014_a_at	SEC31-like 1 (S. cerevisiae)	BM222383	69162	1.53
1453406_a_at	RAB28, member RAS oncogene family	AK012286	100972	1.53
1460180_at	hexosaminidase B	NM_010422	15212	1.53
1415850_at	RAS p21 protein activator 3	NM_009025	19414	1.52
1416052_at	phosphoribosyl pyrophosphate synthetase 1	AK011304	19139	1.52
1416208_at	ubiquitin specific protease 14	BC005571	59025	1.52
1416289_at	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1	NM_011122	18822	1.52
1416315_at	abhydrolase domain containing 4	NM_134076	105501	1.52
1416757_at	RIKEN cDNA 2310031L18 gene	BC027435	68014	1.52
1416871_at	a disintegrin and metalloprotease domain 8	NM_007403	11501	1.52
1418755_at	T-box 15	NM_009323	21384	1.52
1421378_s_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 1	NM_008576	17250	1.52
1422514_at	AE binding protein 1	NM_009636	11568	1.52
1422985_at	frizzled homolog 1 (Drosophila)	NM_021457	14362	1.52
1423078_a_at	sterol-C4-methyl oxidase-like	AK005441	66234	1.52
1423335_at	RIKEN cDNA 1110004F10 gene	AW545354	56372	1.52
1423340_at	transcription factor AP-2 beta	AV334599	21419	1.52
1424857_a_at	tripartite motif protein 34	AF220142	94094	1.52
1428076_s_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 1	BG968046	68194	1.52
1428873_a_at	RIKEN cDNA 4121402D02 gene	AW495537	74026	1.52
1428905_at	Ras-related GTP binding A	AI118026	68441	1.52
1429776_a_at	DnaJ (Hsp40) homolog, subfamily B, member 6	AK005680	23950	1.52
1433490_s_at	Erythrocyte protein band 4.1-like 2	BE951907	13822	1.52
1437434_a_at	RIKEN cDNA 5031439A09 gene	BM241735	68151	1.52
1438058_s_at	prostate tumor over expressed gene 1	BG073526	84113	1.52
1448138_at	protein phosphatase 2A, regulatory subunit B (PRIP)	NM_138748	110854	1.52
1448144_at	heterogeneous nuclear ribonucleoprotein A/B	NM_010448	15384	1.52
1448780_at	solute carrier family 12, member 2	BG069505	20496	1.52
1449528_at	c-fos induced growth factor	NM_010216	14205	1.52
1449936_at	RIKEN cDNA 8430419L09 gene	NM_028982	74525	1.52
1450677_at	checkpoint kinase 1 homolog (S. pombe)	NM_007691	12649	1.52
1450768_at	discs, large homolog 1 (Drosophila)	BQ176806	13383	1.52
1450814_a_at	importin 4	NM_024267	75751	1.52
1451036_at	spastic paraplegia 21 homolog (human)	BB095517	27965	1.52
1451082_at	RIKEN cDNA 1300018I05 gene	AK005044	74157	1.52
1451086_s_at	RAS-related C3 botulinum substrate 1	BC003828	19353	1.52
1451174_at	RIKEN cDNA E430025L02 gene	BC027411	224109	1.52
1452016_at	arachidonate 5-lipoxygenase activating protein	BC026209	11690	1.52
1452026_a_at	phospholipase A2, group XIA	AY007382	66350	1.52
1452174_at	Sterol regulatory element binding factor 2	BM123132	20788	1.52
1452365_at	RIKEN cDNA 4732435N03 gene	AV371987	234356	1.52
1452660_s_at	kelch-like 7 (Drosophila)	AK012326	52323	1.52
1452768_at	testis expressed gene 261	AK013971	21766	1.52
1453849_s_at	heterogeneous nuclear ribonucleoprotein A/B	AK013709	15384	1.52

1416031_s_at	minichromosome maintenance deficient 7 (S. cerevisiae)	NM_008568	17220	1.51
1416452_at	ornithine aminotransferase	BC008119	18242	1.51
1417238_at	Ewing sarcoma breakpoint region 1	NM_007968	14030	1.51
1417601_at	regulator of G-protein signaling 1	NM_015811	50778	1.51
1418194_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide 1	BG965198	171212	1.51
1418279_a_at	A kinase (PRKA) anchor protein 1	BG067335	11640	1.51
1419132_at	toll-like receptor 2	NM_011905	24088	1.51
1419259_at	gb:NM_009105.1 /DB_XREF=gi:6677824 /GEN=R:	NM_009105		1.51
1420886_a_at	X-box binding protein 1	NM_013842	22433	1.51
1421534_at	gb:NM_008016.1 /DB_XREF=gi:6679796 /GEN=Fi	NM_008016		1.51
1422814_at	calmodulin binding protein 1	NM_009791	12316	1.51
1423951_at	RIKEN cDNA 1110025109 gene	BC024620	68634	1.51
1426028_a_at	citron	AF086823	12704	1.51
1426078_a_at	G protein-coupled receptor 108	AF376726	78308	1.51
1426195_a_at	cystatin C	AF483486	13010	1.51
1427056_at	a disintegrin-like and metalloprotease (reprolysin ty	AV228731	235130	1.51
1427253_s_at	suppressor of zeste 12 homolog (Drosophila)	BG066534	52615	1.51
1428235_at	succinate dehydrogenase complex, subunit D, inte	AK013962	66925	1.51
1429239_a_at	StAR-related lipid transfer (START) domain contain	AK014587	170459	1.51
1433665_at	vacuolar protein sorting 41 (yeast)	AW536822	218035	1.51
1436393_a_at	tripartite motif protein 37	BG065227	68729	1.51
1448185_at	homocysteine-inducible, endoplasmic reticulum stre	NM_022331	64209	1.51
1450928_at	inhibitor of DNA binding 4	BB121406	15904	1.51
1460436_at	N-deacetylase/N-sulfotransferase (heparan glucos	BI652065	15531	1.51
1416136_at	matrix metalloproteinase 2	NM_008610	17390	1.5
1417268_at	CD14 antigen	NM_009841	12475	1.5
1419663_at	osteoglycin	BB542051	18295	1.5
1422847_a_at	protein kinase C, delta	AF251036	18753	1.5
1424318_at	RIKEN cDNA 1110067D22 gene	BC019131	216551	1.5
1425526_a_at	paired related homeobox 1	L06502	18933	1.5
1425676_a_at	elongation of very long chain fatty acids (FEN1/Elo	BC006735	54325	1.5
1427058_at	eukaryotic translation initiation factor 4A1	AK010644	13681	1.5
1430811_a_at	cell division cycle associated 1	AK010351	66977	1.5
1431226_a_at	fibronectin type III domain containing 4	AK013203	64339	1.5
1434366_x_at	complement component 1, q subcomponent, beta	AW227993	12260	1.5
1449875_s_at	histocompatibility 2, T region locus 10 /// histocomp	NM_010395	15024 /// 15	1.5
1449930_a_at	signal sequence receptor, beta	NM_025448	66256	1.5
1452296_at	slit homolog 3 (Drosophila)	BM570006	20564	1.5
1456383_at	RIKEN cDNA 2410005K20 gene	BM239987	66409	1.5
1417883_at	glutathione S-transferase, theta 2	BC012707	14872	-1.5
1421050_at	DNA segment, Chr 11, Wayne State University 68,	NM_026776	28084	-1.5
1422592_at	catenin delta 2	NM_008729	18163	-1.5
1427364_a_at	ornithine decarboxylase, structural 1	S64539	18263	-1.5
1437477_at	leucine rich repeat (in FLII) interacting protein 1	AV301584	16978	-1.5
1449957_at	protein tyrosine phosphatase, receptor type, V	NM_007955	13924	-1.5
1451699_at	MRNA similar to RIKEN cDNA 9030624G23 gene (	BC024416		-1.5
1418289_at	nestin	AI413223	18008	-1.51
1418852_at	cholinergic receptor, nicotinic, alpha polypeptide 1 (	NM_007389	11435	-1.51
1421309_at	O-6-methylguanine-DNA methyltransferase	NM_008598	17314	-1.51
1424598_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 6	BC021452	13209	-1.51
1424893_at	nuclear distribution gene E-like homolog 1 (A. nidul	BC021434	83431	-1.51
1424940_s_at	cDNA sequence BC022687	BC022687	217887	-1.51
1427104_at	zinc finger protein 612	BG076162	234725	-1.51

1434537_at	Solute carrier organic anion transporter family, men	BM237089	108116	-1.51
1439295_x_at	hypothetical LOC434179	BB371300	434179	-1.51
1451629_at	limb-bud and heart	BC026827	77889	-1.51
1452320_at	similar to low density lipoprotein receptor-related pr	C80829	381369 /// 4	-1.51
1456011_x_at	acetyl-Coenzyme A acyltransferase 1	BB210491	113868	-1.51
1416298_at	matrix metalloproteinase 9	NM_013599	17395	-1.52
1416811_s_at	cytotoxic T lymphocyte-associated protein 2 alpha /	NM_007796	13024 /// 13	-1.52
1419624_a_at	RIKEN cDNA 1700010I14 gene	NM_025851	66931	-1.52
1420416_at	sema domain, immunoglobulin domain (Ig), short b.	NM_009152	20346	-1.52
1420664_s_at	protein C receptor, endothelial	NM_011171	19124	-1.52
1420989_at	RIKEN cDNA 4933411K20 gene	NM_025747	66756	-1.52
1434987_at	aldehyde dehydrogenase 2, mitochondrial	AI462635	11669	-1.52
1436343_at	chromodomain helicase DNA binding protein 4	BM502696	107932	-1.52
1437100_x_at	proviral integration site 3	BB206220	223775	-1.52
1437848_x_at	ADP-dependent glucokinase	BB490673	72141	-1.52
1449078_at	ST3 beta-galactoside alpha-2,3-sialyltransferase 6	NM_018784	54613	-1.52
1449852_a_at	EH-domain containing 4	NM_133838	98878	-1.52
1419689_at	glypican 6	BC023448	23888	-1.53
1420532_at	activin receptor interacting protein 1	NM_015823	50791	-1.53
1421354_at	protein kinase, cGMP-dependent, type II	NM_008926	19092	-1.53
1422948_s_at	histone 1, H4h /// histone 1, H4c /// histone 1, H4i //	NM_013550	319155 /// 3	-1.53
1423071_x_at	gb:AW549928 /DB_XREF=gi:7195356 /DB_XREF= AW549928			-1.53
1423696_a_at	gb:BC006869.1 /DB_XREF=gi:13905153 /FEA=FLI	BC006869		-1.53
1424699_at	RIKEN cDNA 4921511K06 gene	BC006583	232664	-1.53
1425271_at	proteasome (prosome, macropain) 26S subunit, AT	AB000121	19183	-1.53
1431980_a_at	arsenic (+3 oxidation state) methyltransferase	AK009814	57344	-1.53
1436990_s_at	Nur77 downstream gene 2	AA038464	103172	-1.53
1449357_at	RIKEN cDNA 2310030G06 gene	NM_025865	66952	-1.53
1452590_a_at	hypothetical LOC211623	BB609699	211623	-1.53
1416379_at	pannexin 1	NM_019482	55991	-1.54
1416689_at	tuffelin 1	NM_011656	22156	-1.54
1418285_at	ephrin B1	NM_010110	13641	-1.54
1419712_at	interleukin 3 receptor, alpha chain	NM_008369	16188	-1.54
1423537_at	growth associated protein 43	BB622036	14432	-1.54
1425652_s_at	RNA binding protein gene with multiple splicing	AF148511	19663	-1.54
1425983_x_at	homeodomain interacting protein kinase 2	AF208292	15258	-1.54
1426725_s_at	E26 avian leukemia oncogene 1, 5' domain	BB151715	23871	-1.54
1428935_at	calnexin	AI988026	12330	-1.54
1437218_at	fibronectin 1	BM234360	14268	-1.54
1438415_s_at	RIKEN cDNA 1300010K09 gene	AV166218	74766	-1.54
1448894_at	aldo-keto reductase family 1, member B8	NM_008012	14187	-1.54
1455505_at	Transcribed locus	AI840824		-1.54
1455893_at	RIKEN cDNA 2610028F08 gene	BG067392	239405	-1.54
1416357_a_at	melanoma cell adhesion molecule	NM_023061	84004	-1.55
1417022_at	solute carrier family 7 (cationic amino acid transpor	NM_007515	11989	-1.55
1417853_at	chloride channel calcium activated 1 /// chloride ch	AF047838	12722 /// 80	-1.55
1420991_at	ankyrin repeat domain 1 (cardiac muscle)	AK009959	107765	-1.55
1422573_at	gb:D85596.1 /DB_XREF=gi:2094764 /GEN=AMPD	D85596		-1.55
1422809_at	regulating synaptic membrane exocytosis 2	NM_053271	116838	-1.55
1423506_a_at	neuronatin	AV218841	18111	-1.55
1425268_a_at	RIKEN cDNA 3110045G13 gene	BB752266	73182	-1.55
1425357_a_at	gremlin 1	BC015293	23892	-1.55
1438982_s_at	RIKEN cDNA 2810417J12 gene	BB251185	76917	-1.55

1448154_at	N-myc downstream regulated gene 2	NM_013864	29811	-1.55
1449591_at	caspase 11, apoptosis-related cysteine protease	NM_007609	12363	-1.55
1449982_at	interleukin 11	NM_008350	16156	-1.55
1418835_at	pleckstrin homology-like domain, family A, member	NM_009344	21664	-1.56
1419766_at	SNF1-like kinase	NM_010831	17691	-1.56
1424471_at	Rap guanine nucleotide exchange factor (GEF) 3	BC020532	223864	-1.56
1428882_at	RIKEN cDNA 2510001110 gene	AK010881	74781	-1.56
1431024_a_at	AT rich interactive domain 4B (Rbp1 like)	AK020165	94246	-1.56
1450744_at	elongation factor RNA polymerase II 2	NM_138953	192657	-1.56
1452304_a_at	Rho guanine nucleotide exchange factor (GEF) 5	BC025127	54324	-1.56
1452391_at	coxsackievirus and adenovirus receptor	AK004908	13052	-1.56
1415824_at	stearoyl-Coenzyme A desaturase 2	BG060909	20250	-1.57
1419457_at	Rho-guanine nucleotide exchange factor	BG069493	110596	-1.57
1419806_at	high density lipoprotein (HDL) binding protein	C77256	110611	-1.57
1421201_a_at	trophinin	AF241244	56191	-1.57
1424376_at	CDC42 effector protein (Rho GTPase binding) 1	BC016250	104445	-1.57
1424612_at	RIKEN cDNA 9330161F08 gene	BC016107	223473	-1.57
1425837_a_at	CCR4 carbon catabolite repression 4-like (S. cerev	AF199491	12457	-1.57
1449553_at	RIKEN cDNA 2610200G18 gene	AK009389	67149	-1.57
1451660_a_at	homeo box B6	BC016893	15414	-1.57
1417034_at	RIKEN cDNA 1810073E21 gene	NM_025960	67091	-1.58
1417323_at	RIKEN cDNA 5430413I02 gene	NM_019976	56742	-1.58
1422803_at	follistatin-like 3	NM_031380	83554	-1.58
1423489_at	monocyte to macrophage differentiation-associated	BC021914	67468	-1.58
1424733_at	purinergic receptor P2Y, G-protein coupled, 14	AF177211	140795	-1.58
1424801_at	enabled homolog (Drosophila)	BQ044016	13800	-1.58
1425148_a_at	sorting nexin 6	BC025911	72183	-1.58
1434196_at	DnaJ (Hsp40) homolog, subfamily A, member 4	BQ176119	58233	-1.58
1448857_a_at	zinc finger, DHHC domain containing 12	NM_025428	66220	-1.58
1449283_a_at	mitogen-activated protein kinase 12	BC021640	29857	-1.58
1451245_at	leucine rich repeat containing 3B	BC019794	218763	-1.58
1452180_at	PHD finger protein 17	BG065238	269424	-1.58
1452412_at	homeo box C8	BB283726	15426	-1.58
1454890_at	angiominin	BG067039	27494	-1.58
1425274_at	gb:AF221854.1 /DB_XREF=gi:12655824 /FEA=FLr	AF289490		-1.59
1425964_x_at	heat shock protein 1	U03561	15507	-1.59
1450541_at	plasmacytoma variant translocation 1	NM_011222	19296	-1.59
1451615_at	cDNA sequence BC026374	BC026374	234677	-1.59
1459021_at	gb:BB307071 /DB_XREF=gi:16402263 /DB_XREF: BB307071			-1.59
1417483_at	nuclear factor of kappa light polypeptide gene enha	AB026551	80859	-1.6
1418429_at	kinesin family member 5B	BI328541	16573	-1.6
1419300_at	FMS-like tyrosine kinase 1	NM_010228	14254	-1.6
1420010_at	CDNA, clone:Y0G0116J03, strand:plus, reference:f	C80083		-1.6
1425673_at	LIM domain containing preferred translocation part	BC005613	210126	-1.6
1426165_a_at	caspase 3, apoptosis related cysteine protease	D86352	12367	-1.6
1435866_s_at	histone 3, H2a	AV297651	319162	-1.6
1448728_a_at	nuclear factor of kappa light polypeptide gene enha	AB026551	80859	-1.6
1449352_at	polymerase (DNA directed), lambda	NM_020032	56626	-1.6
1449752_at	gb:AI596360 /DB_XREF=gi:4605408 /DB_XREF=n	AI596360		-1.6
1416779_at	serum deprivation response	BE197945	20324	-1.61
1423310_at	trophoblast glycoprotein	BQ177165	21983	-1.61
1424150_at	cDNA sequence BC024955	BC024955	233552	-1.61
1437835_a_at	RIKEN cDNA 0610011L14 gene	BB357560	68295	-1.61

1449152_at	cyclin-dependent kinase inhibitor 2B (p15, inhibits C	AF059567	12579	-1.61
1450766_at	phosphodiesterase 6H, cGMP-specific, cone, gamr	NM_023898	78600	-1.61
1418509_at	carbonyl reductase 2	BC010758	12409	-1.62
1418937_at	deiodinase, iodothyronine, type II	AF177196	13371	-1.62
1424829_at	RIKEN cDNA A830007P12 gene	BC013092	227612	-1.62
1425558_at	kinesin light chain 3	BC017147	232943	-1.62
1425603_at	RIKEN cDNA 0610011104 gene	BC006049	66058	-1.62
1428794_at	RIKEN cDNA 2810012G08 gene	BQ175052	76364	-1.62
1429787_x_at	ZW10 interactor	AK008144	52696	-1.62
1431170_at	ephrin A3	AA388313	13638	-1.62
1449311_at	BTB and CNC homology 1	NM_007520	12013	-1.62
1451474_a_at	poly (ADP-ribose) polymerase family, member 8	BC022679	52552	-1.62
1420329_at	RIKEN cDNA 4930455C21 gene	NM_024273	76916	-1.63
1420851_at	par-6 partitioning defective 6 homolog gamma (C. ε	NM_053117	93737	-1.63
1422545_at	T-box 2	NM_009324	21385	-1.63
1424854_at	histone 1, H4f /// histone 1, H4i /// histone 1, H4m //	BC019757	319157 /// 3	-1.63
1429692_s_at	GTP cyclohydrolase 1	BB698398	14528	-1.63
1434958_at	Similar to saccin	BG075163	432862	-1.63
1435252_at	UDP-Gal:betaGal beta 1,3-galactosyltransferase, p	AV328064	117592	-1.63
1449563_at	contactin 1	NM_007727	12805	-1.63
1451498_at	cDNA sequence BC004853	BC004853	227618	-1.63
1422168_a_at	brain derived neurotrophic factor	AY057913	12064	-1.64
1435137_s_at	RIKEN cDNA 1200015M12 gene	BB119177	71739	-1.64
1448508_at	expressed sequence AI429613	NM_134000	103213	-1.64
1450781_at	high mobility group AT-hook 2	X58380	15364	-1.64
1451287_s_at	RIKEN cDNA 2810003C17 gene	BC024599	108897	-1.64
1417374_at	tubulin, alpha 4	AW491660	22145	-1.65
1421117_at	dystonin	NM_010081	13518	-1.65
1451751_at	DNA-damage-inducible transcript 4-like	AF335325	73284	-1.65
1419458_at	Rho-guanine nucleotide exchange factor	BG069493	110596	-1.66
1421895_at	eukaryotic translation initiation factor 2, subunit 3, s	NM_012010	26905	-1.66
1422155_at	histone 2, H3c2	BC015270	97114	-1.66
1427321_s_at	coxsackievirus and adenovirus receptor	AK004908	13052	-1.66
1439388_s_at	breast cancer anti-estrogen resistance 1	BB251739	12927	-1.66
1455265_a_at	gb:BB100249 /DB_XREF=gi:8752817 /DB_XREF=	BB100249		-1.66
1416927_at	transformation related protein 53 inducible nuclear	AW495711	60599	-1.67
1417311_at	cysteine rich protein 2	NM_024223	68337	-1.67
1422169_a_at	brain derived neurotrophic factor	AY057913	12064	-1.67
1422945_a_at	kinesin family member 5C	AI844677	16574	-1.67
1424470_a_at	Rap guanine nucleotide exchange factor (GEF) 3	BC020532	223864	-1.67
1426472_at	zinc finger protein 52	BM225280	22710	-1.67
1417373_a_at	tubulin, alpha 4	AW491660	22145	-1.68
1422947_at	histone 1, H4f /// histone 1, H4i /// histone 1, H4m //	NM_013550	319157 /// 3	-1.68
1427482_a_at	carbonic anhydrase 8	X61397	12319	-1.68
1437012_x_at	Rap guanine nucleotide exchange factor (GEF) 3	BB226235	223864	-1.68
1438789_s_at	dihydropyrimidinase-like 3	AV161550	22240	-1.68
1451999_at	LIM domain binding 3	AF114378	24131	-1.68
1452436_at	lysyl oxidase-like 2	AF117951	94352	-1.68
1423277_at	protein tyrosine phosphatase, receptor type, K	AI893646	19272	-1.69
1423488_at	monocyte to macrophage differentiation-associated	BC021914	67468	-1.69
1448024_at	RIKEN cDNA B430320C24 gene	BG080055	319900	-1.69
1449033_at	tumor necrosis factor receptor superfamily, membe	AB013898	18383	-1.69
1421053_at	kinesin family member 1A	NM_008440	16560	-1.7



1418711_at	platelet derived growth factor, alpha	BB371842	18590	-1.71
1424863_a_at	homeodomain interacting protein kinase 2	AF170301	15258	-1.71
1426910_at	PRKC, apoptosis, WT1, regulator	BB398886	114774	-1.71
1427633_a_at	Pregnancy-associated plasma protein A	AF439513	18491	-1.71
1432281_a_at	integrin beta 6	AK019511	16420	-1.71
1438116_x_at	solute carrier family 9 (sodium/hydrogen exchanger	BB805362	26941	-1.71
1449193_at	CD5 antigen-like	NM_009690	11801	-1.71
1422028_a_at	E26 avian leukemia oncogene 1, 5' domain	BC010588	23871	-1.72
1424290_at	cDNA sequence BC010311	BB817847	209212	-1.72
1426037_a_at	gb:U94828.1 /DB_XREF=gi:2605641 /GEN=RGS-r	U94828		-1.72
1426930_at	bruno-like 4, RNA binding protein (Drosophila)	AV327653	108013	-1.72
1449481_at	solute carrier family 25 (mitochondrial carrier, adenine	BC016571	50799	-1.72
1453092_at	RIKEN cDNA 2300002G24 gene	AK009010	74175	-1.72
1460670_at	RIO kinase 3 (yeast)	NM_024182	66878	-1.72
1415932_x_at	ATPase, class II, type 9A	NM_015731	11981	-1.73
1420518_a_at	immunoglobulin superfamily, member 9	AF317839	93842	-1.73
1422742_at	human immunodeficiency virus type I enhancer binding	NM_007772	110521	-1.73
1426010_a_at	erythrocyte protein band 4.1-like 3	AF177146	13823	-1.73
1452104_at	RIKEN cDNA 2600005N12 gene	BI456906	70317	-1.73
1417917_at	calponin 1	NM_009922	12797	-1.74
1422751_at	transducin-like enhancer of split 1, homolog of Drosophila	NM_011599	21885	-1.74
1424800_at	enabled homolog (Drosophila)	BQ044016	13800	-1.74
1427539_a_at	ZW10 interactor	BC013559	52696	-1.74
1450786_x_at	PDZ and LIM domain 5	NM_019808	56376	-1.74
1418872_at	ATP-binding cassette, sub-family B (MDR/TAP), member 1	NM_011075	18669	-1.75
1419613_at	procollagen, type VII, alpha 1	NM_007738	12836	-1.75
1438855_x_at	gb:BB233088 /DB_XREF=gi:8912943 /DB_XREF=	BB233088		-1.75
1439376_x_at	gb:BB357590 /DB_XREF=gi:9069418 /DB_XREF=	BB357590		-1.75
1417703_at	poliovirus receptor-related 2	NM_008990	19294	-1.76
1423988_at	gb:BE373450 /DB_XREF=gi:9318813 /DB_XREF=	BE373450		-1.76
1418076_at	suppression of tumorigenicity 14 (colon carcinoma)	NM_011176	19143	-1.77
1418876_at	forkhead box D1	BB662927	15229	-1.77
1450826_a_at	serum amyloid A 3	NM_011315	20210	-1.78
1454959_s_at	guanine nucleotide binding protein, alpha inhibiting	BQ174580	14677	-1.78
1421425_a_at	Down syndrome critical region gene 1-like 1	NM_030598	53901	-1.79
1422562_at	Ras-related associated with diabetes	NM_019662	56437	-1.79
1424008_a_at	RNA binding protein with multiple splicing 2	BC021788	71973	-1.79
1427485_at	leiomodulin 1 (smooth muscle)	AF237627	93689	-1.79
1418030_at	solute carrier organic anion transporter family, member 1	NM_023908	108116	-1.8
1419719_at	gamma-aminobutyric acid (GABA-A) receptor, subunit 1	NM_008069	14400	-1.8
1432156_a_at	ring finger protein 32	AK012288	56874	-1.8
1460511_at	plakophilin 2	AK004731	67451	-1.8
1416023_at	fatty acid binding protein 3, muscle and heart	NM_010174	14077	-1.81
1417023_a_at	fatty acid binding protein 4, adipocyte	NM_024406	11770	-1.81
1420406_at	paternally expressed 12	NM_013788	27412	-1.81
1435184_at	RIKEN cDNA B430320C24 gene	BG066982	319900	-1.81
1433509_s_at	DNA segment, Chr 6, ERATO Doi 253, expressed	BQ174328	52250	-1.82
1452291_at	centaurin, delta 1	AV375176	212285	-1.82
1423055_at	neuron specific gene family member 1	NM_010942	18196	-1.84
1424966_at	transmembrane protein 40	BC019416	94346	-1.84
1427019_at	protein tyrosine phosphatase, receptor type Z, polypeptide	BC002298	19283	-1.84
1436853_a_at	synuclein, alpha	AI324124	20617	-1.84
1437170_x_at	PDZ and LIM domain 1 (elfin)	AV013697	54132	-1.84

1438115_a_at	solute carrier family 9 (sodium/hydrogen exchanger)	BB805362	26941	-1.84
1448471_a_at	cytotoxic T lymphocyte-associated protein 2 alpha	NM_007796	13024	-1.84
1425521_at	polyadenylate binding protein-interacting protein 1	BC019726	218693	-1.85
1452240_at	bruno-like 4, RNA binding protein (Drosophila)	AV327653	108013	-1.85
1427943_at	acylphosphatase 2, muscle type	BI730288	75572	-1.86
1448468_a_at	potassium voltage-gated channel, shaker-related s	AF033003	16497	-1.86
1422612_at	hexokinase 2	NM_013820	15277	-1.87
1423072_at	RIKEN cDNA 6720475J19 gene	AW549928	68157	-1.87
1423585_at	insulin-like growth factor binding protein 7	AI481026	29817	-1.87
1435695_a_at	RIKEN cDNA A030007L17 gene	AA673177	68252	-1.88
1437548_at	gb:AV367839 /DB_XREF=gi:16397715 /DB_XREF:	AV367839		-1.88
1449491_at	caspase recruitment domain family, member 10	NM_130859	105844	-1.88
1451263_a_at	fatty acid binding protein 4, adipocyte	BC002148	11770	-1.88
1418393_a_at	integrin alpha 7	NM_008398	16404	-1.89
1418778_at	RIKEN cDNA 9030408N13 gene	NM_025779	66815	-1.89
1423500_a_at	SRY-box containing gene 5	AI528773	20678	-1.89
1424967_x_at	troponin T2, cardiac	L47552	21956	-1.89
1448182_a_at	CD24a antigen	NM_009846	12484	-1.89
1418004_a_at	RIKEN cDNA 1810009M01 gene	NM_023056	65963	-1.9
1423471_at	polypyrimidine tract binding protein 2	BB076855	56195	-1.9
1436272_at	RIKEN cDNA 1110059F07 gene	BQ175872	98732	-1.9
1451756_at	FMS-like tyrosine kinase 1	D88690	14254	-1.9
1416007_at	special AT-rich sequence binding protein 1	AV172776	20230	-1.91
1416034_at	CD24a antigen	NM_009846	12484	-1.92
1418726_a_at	troponin T2, cardiac	NM_011619	21956	-1.93
1420411_a_at	phosphatidylinositol 4-kinase type 2 beta	NM_028744	67073	-1.93
1438476_a_at	chromodomain helicase DNA binding protein 4	BB201828	107932	-1.93
1428014_at	histone 1, H4h	BC028550	69386	-1.94
1435162_at	Protein kinase, cGMP-dependent, type II	BB823350	19092	-1.94
1429183_at	plakophilin 2	AK005020	67451	-1.95
1447959_at	zinc finger protein 398	AI506402	272347	-1.95
1448961_at	phospholipid scramblase 2	NM_008880	18828	-1.95
1449187_at	platelet derived growth factor, alpha	BB371842	18590	-1.95
1451612_at	gb:BC027262.1 /DB_XREF=gi:20072672 /FEA=FL	BC027262		-1.96
1420872_at	guanylate cyclase 1, soluble, beta 3	BF472806	54195	-1.98
1421153_at	lysyl oxidase-like 4	NM_053083	67573	-1.99
1423952_a_at	keratin complex 2, basic, gene 7	BC010337	110310	-1.99
1416688_at	synaptosomal-associated protein 91	NM_013669	20616	-2
1417588_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide l	AK019995	14425	-2
1420183_at	loricrin	AI036317	16939	-2
1416008_at	special AT-rich sequence binding protein 1	AV172776	20230	-2.01
1424890_at	basonuclin 1	U88064	12173	-2.01
1418990_at	membrane-spanning 4-domains, subfamily A, mem	NM_025658	66607	-2.02
1419437_at	single-minded homolog 2 (Drosophila)	D63383	20465	-2.02
1421624_a_at	enabled homolog (Drosophila)	NM_010135	13800	-2.02
1426243_at	cystathionase (cystathionine gamma-lyase)	BC019483	107869	-2.02
1448942_at	guanine nucleotide binding protein (G protein), gar	NM_025331	66066	-2.02
1417262_at	prostaglandin-endoperoxide synthase 2	M94967	19225	-2.03
1439368_a_at	solute carrier family 9 (sodium/hydrogen exchanger	AV002797	65962	-2.03
1455149_at	SH3 multiple domains 2	BB072090	59009	-2.03
1418494_at	early B-cell factor 2	U71189	13592	-2.06
1437502_x_at	CD24a antigen	BB560574	12484	-2.06
1448289_at	collapsin response mediator protein 1	AB006714	12933	-2.08

1450329_a_at	arrestin 3, retinal	NM_133205	170735	-2.09
1448260_at	ubiquitin carboxy-terminal hydrolase L1	NM_011670	22223	-2.1
1418933_at	solute carrier family 1 (high affinity aspartate/glutan	NM_009200	20513	-2.11
1448664_a_at	Aortic preferentially expressed gene 1	NM_007463	11790	-2.12
1424568_at	gb:AV234585 /DB_XREF=gi:16386146 /DB_XREF:BC007185			-2.13
1450140_a_at	cyclin-dependent kinase inhibitor 2A	NM_009877	12578	-2.16
1427183_at	epidermal growth factor-containing fibulin-like extra	BC023060	216616	-2.17
1454737_at	dual specificity phosphatase 9	AV295798	75590	-2.17
1422861_s_at	gb:NM_019808.1 /DB_XREF=gi:12083690 /GEN=E	NM_019808		-2.19
1425573_a_at	development and differentiation enhancing	BE943736	13196	-2.19
1435943_at	Dipeptidase 1 (renal)	AI647687	13479	-2.19
1451527_at	procollagen C-endopeptidase enhancer 2	AF352788	76477	-2.2
1417220_at	fumarylacetoacetate hydrolase	NM_010176	14085	-2.21
1418486_at	vanin 1	NM_011704	22361	-2.21
1428955_x_at	solute carrier family 9 (sodium/hydrogen exchanger	AK004710	65962	-2.21
1452124_at	RIKEN cDNA 2900054D09 gene	BB628935	73013	-2.22
1418773_at	fatty acid desaturase 3	BE652876	60527	-2.23
1422316_at	glycoprotein 1b, alpha polypeptide	NM_010326	14723	-2.23
1449823_at	dachshund 2 (Drosophila)	NM_033605	93837	-2.23
1420682_at	cholinergic receptor, nicotinic, beta polypeptide 1 (n	M14537	11443	-2.24
1420992_at	ankyrin repeat domain 1 (cardiac muscle)	AK009959	107765	-2.25
1416045_a_at	SWI/SNF related, matrix associated, actin depende	NM_011418	20587	-2.27
1451415_at	RIKEN cDNA 1810011O10 gene	BC016562	69068	-2.27
1418480_at	chemokine (C-X-C motif) ligand 7	NM_023785	57349	-2.28
1439369_x_at	solute carrier family 9 (sodium/hydrogen exchanger	AV002797	65962	-2.28
1417951_at	enolase 3, beta muscle	NM_007933	13808	-2.29
1421825_at	beta-site APP cleaving enzyme 1	AF200346	23821	-2.29
1417542_at	ribosomal protein S6 kinase, polypeptide 2	BB737182	20112	-2.3
1432517_a_at	nicotinamide N-methyltransferase	AK006371	18113	-2.35
1444052_at	FMS-like tyrosine kinase 4	AI323512	14257	-2.35
1449799_s_at	plakophilin 2	AA516617	67451	-2.35
1449036_at	ring finger protein 128	AK004847	66889	-2.38
1426731_at	desmin	AI156158	13346	-2.39
1426016_a_at	trophinin	AF288606	56191	-2.4
1418710_at	CD59a antigen	NM_007652	12509	-2.41
1418945_at	matrix metalloproteinase 3	NM_010809	17392	-2.41
1434442_at	DNA segment, Chr 5, ERATO Doi 593, expressed	BB667844	52331	-2.41
1422765_at	microtubule-associated protein, RP/EB family, mem	BM228525	13589	-2.46
1416783_at	gb:NM_009311.1 /DB_XREF=gi:6678206 /GEN=Ta	NM_009311		-2.49
1449328_at	lymphocyte antigen 75	NM_013825	17076	-2.49
1438968_x_at	serine protease inhibitor, Kunitz type 2	AV058358	20733	-2.52
1424451_at	3-ketoacyl-CoA thiolase B	BC019882	235674	-2.54
1418318_at	ring finger protein 128	AK004847	66889	-2.55
1419534_at	gb:NM_138648.1 /DB_XREF=gi:20149719 /GEN=C	NM_138648		-2.56
1449356_at	ankyrin repeat and SOCs box-containing protein 5	NM_029569	76294	-2.62
1418497_at	fibroblast growth factor 13	AF020737	14168	-2.65
1419100_at	serine (or cysteine) proteinase inhibitor, clade A, m	NM_009252	20716	-2.67
1448265_x_at	epithelial V-like antigen 1	BC015076	14012	-2.71
1418569_at	RIKEN cDNA 2410043F08 gene	BG070068	74202	-2.72
1425092_at	RIKEN cDNA A830016G23 gene	AF183946	320873	-2.79
1420450_at	matrix metalloproteinase 10	NM_019471	17384	-2.89
1455158_at	integrin alpha 3	BI664675	16400	-2.91
1448595_a_at	reduced expression 3	NM_009052	19716	-2.93

1417109_at	lipocalin 7	BC005738	94242	-2.99
1422245_a_at	MRV integration site 1	U63408	17540	-2.99
1421594_a_at	synaptotagmin-like 2	NM_031394	83671	-3.04
1448871_at	mitogen activated protein kinase 13	NM_011950	26415	-3.05
1421227_at	granzyme E	NM_010373	14942	-3.07
1435493_at	RIKEN cDNA 5730453H04 gene	AV297961	105072	-3.09
1456747_x_at	MIC2 (monoclonal Imperial Cancer Research Fund	BB334959	171486	-3.2
1427735_a_at	actin, alpha 1, skeletal muscle	M12233	11459	-3.22
1456746_a_at	gb:BB334959 /DB_XREF=gi:9043722 /DB_XREF=	BB334959		-3.28
1448926_at	homeo box A5	BC011063	15402	-3.35
1419033_at	RIKEN cDNA 2610018G03 gene	AW556821	70415	-3.38
1448169_at	keratin complex 1, acidic, gene 18	NM_010664	16668	-3.4
1434325_x_at	protein kinase, cAMP dependent regulatory, type I	BB274009	19085	-3.66
1453208_at	RIKEN cDNA 2700089E24 gene	AV030118	381820	-3.67
1417130_s_at	angiopoietin-like 4	NM_020581	57875	-3.68
1418357_at	forkhead box G1	NM_008241	15228	-4.23
1417466_at	gb:NM_133736.1 /DB_XREF=gi:19526933 /GEN=1	NM_133736		-4.28
1435494_s_at	RIKEN cDNA 5730453H04 gene	AV297961	105072	-4.34
1426225_at	retinol binding protein 4, plasma	U63146	19662	-4.36
1416676_at	kininogen 1	NM_023125	16644	-4.41
1433783_at	LIM domain binding 3	BB479063	24131	-4.5
1449965_at	mast cell protease 8	NM_008572	17231	-4.69
1418547_at	tissue factor pathway inhibitor 2	NM_009364	21789	-5.24
1435856_x_at	gb:AV310148 /DB_XREF=gi:6363183 /DB_XREF=,	AV310148		-5.73
1425413_at	regulatory factor X, 3 (influences HLA class II expre	BC017598	19726	-6.29
1417256_at	matrix metalloproteinase 13	NM_008607	17386	-13.26
1436936_s_at	gb:BG806300 /DB_XREF=gi:17953245 /DB_XREF	BG806300		-20.68
1419195_at	RIKEN cDNA 9230106L14 gene	NM_138685	192201	-46.93