

Table S1

A

Probe set	Gene name	Ratio	P value
202718_at	insulin-like growth factor binding protein 2, 36kDa	59,68	6,07E-10
201427_s_at	selenoprotein P, plasma, 1	14.21	2,15E-07
201069_at	matrix metalloproteinase 2 (gelatinase A, 72kDa gelatinase, 72kDa type IV collagenase)	8.94	8,23E-08
236738_at	Similar to LOC166075	8.82	7,92E-07
228011_at	family with sequence similarity 92, member A2; family with sequence similarity 92, member A1	8,08	1,18E-06
204030_s_at	schwannomin interacting protein 1	6,79	1,18E-06
217975_at	WW domain binding protein 5	5,49	2,15E-07
229620_at	selenoprotein P, plasma, 1	5,42	1,00E-05
209813_x_at	TCR gamma alternate reading frame protein; T cell receptor gamma variable 9; T cell receptor gamma constant 1	5,03	2,54E-06
227235_at	guanylate cyclase 1, soluble, alpha 3	4,99	3,36E-05
204485_s_at	target of myb1 (chicken)-like 1	4,86	7,51E-04
215288_at	transient receptor potential cation channel, subfamily C, member 2 (pseudogene)	4,81	1,64E-08
216920_s_at	TCR gamma alternate reading frame protein; T cell receptor gamma variable 9; T cell receptor gamma constant 1	4,71	2,54E-06
215806_x_at	TCR gamma alternate reading frame protein; T cell receptor gamma variable 9; T cell receptor gamma constant 1	4,66	2,54E-06
211144_x_at	TCR gamma alternate reading frame protein; T cell receptor gamma variable 9; T cell receptor gamma constant 1	4,65	5,16E-06
202890_at	microtubule-associated protein 7	4,56	1,38E-05
215016_x_at	dystonin	4,49	9,85E-05
213110_s_at	collagen, type IV, alpha 5	4,48	1,87E-05
212775_at	obscurin-like 1	4,44	1,64E-08
235391_at	family with sequence similarity 92, member A2; family with sequence similarity 92, member A1	4,23	1,34E-07
206067_s_at	Wilms tumor 1	4,12	5,84E-05
227522_at	carboxymethylenebutenolidase homolog (Pseudomonas)	4,10	1,18E-06
217963_s_at	nerve growth factor receptor (TNFRSF16) associated protein 1	4,09	9,85E-05
212254_s_at	dystonin	4,08	9,15E-04
228885_at	MAM domain containing 2	4,07	2,05E-04
204044_at	quinolinate phosphoribosyltransferase	4,05	5,16E-06
223599_at	TRIM6-TRIM34 readthrough transcript; tripartite motif-containing 6; tripartite motif-containing 34	4,02	1,26E-04
1557383_a_at	ATPase inhibitory factor 1	3,86	5,22E-07
213395_at	megalencephalic leukoencephalopathy with subcortical cysts 1	3,85	1,74E-06
228280_at	zinc finger CCCH-type, antiviral 1-like	3,84	1,34E-07
214321_at	nephroblastoma overexpressed gene	3,79	9,85E-05
240065_at	family with sequence similarity 81, member B	3,79	3,23E-04
227461_at	stonin 2	3,63	5,16E-06
225285_at	branched chain aminotransferase 1, cytosolic	3,60	3,36E-05
223475_at	cysteine-rich secretory protein LCCL domain containing 1	3,53	8,23E-08
205229_s_at	coagulation factor C homolog, cochin (Limulus polyphemus)	3,51	4,45E-05
219304_s_at	platelet derived growth factor D	3,51	1,38E-05
1553808_a_at	NK2 transcription factor related, locus 3 (Drosophila)	3,38	1,87E-05
236892_s_at	similar to hCG2042068	3,38	7,92E-07
243010_at	musashi homolog 2 (Drosophila)	3,17	2,05E-04
235900_at	spinster homolog 3 (Drosophila)	3,16	2,58E-04
204069_at	Meis homeobox 1	3,16	1,74E-06
227860_at	carboxypeptidase X (M14 family), member 1	3,15	7,61E-05
209160_at	aldo-keto reductase family 1, member C3 (3-alpha hydroxysteroid dehydrogenase, type II)	3,12	7,51E-04
235052_at	zinc finger protein 792	3,11	4,02E-04
209392_at	ectonucleotide pyrophosphatase/phosphodiesterase 2	3,11	4,45E-05
228708_at	RAB27B, member RAS oncogene family	3,11	6,13E-04
225112_at	abl interactor 2	3,07	3,38E-07
229530_at	guanylate cyclase 1, soluble, alpha 3	3,04	2,58E-04
203608_at	aldehyde dehydrogenase 5 family, member A1	3,03	7,61E-05
242414_at	quinolinate phosphoribosyltransferase	3,02	1,61E-04
239237_at	amphiphysin	3,00	2,05E-04
241903_at	potassium channel tetramerisation domain containing 3	2,97	1,87E-05
226517_at	branched chain aminotransferase 1, cytosolic	2,95	6,13E-04
221606_s_at	nucleosomal binding protein 1	2,94	1,87E-05
221942_s_at	guanylate cyclase 1, soluble, alpha 3	2,94	3,23E-04
209662_at	centrin, EF-hand protein, 3 (CDC31 homolog, yeast)	2,91	7,23E-06
226190_at	mitogen-activated protein kinase kinase kinase 13	2,90	4,02E-04
230666_at	HOXA11 antisense RNA (non-protein coding)	2,88	7,61E-05
232424_at	PR domain containing 16	2,87	7,51E-04
232979_at		2,87	5,84E-05
215599_at	glucuronidase, beta pseudogene	2,86	4,97E-04
220377_at	family with sequence similarity 30, member A	2,82	2,05E-04
213258_at	tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)	2,81	3,23E-04
203762_s_at	dynein, cytoplasmic 2, light intermediate chain 1	2,77	1,87E-05
226556_at	ribosomal protein L4; ribosomal protein L4 pseudogene 5; ribosomal protein L4 pseudogene 4	2,76	3,23E-04
227509_x_at		2,73	1,38E-05
226134_s_at	musashi homolog 2 (Drosophila)	2,73	1,61E-04
236918_s_at	leucine rich repeat containing 34	2,70	3,36E-05
243128_at	zinc finger protein 175	2,69	1,00E-05
203151_at	microtubule-associated protein 1A	2,69	8,23E-08
225098_at	abl interactor 2	2,66	2,52E-05
230174_at	lysophospholipase-like 1	2,65	4,02E-04
222603_at	endoplasmic reticulum metalloproteinase 1	2,65	4,02E-04
238122_at	RNA binding motif protein 12B	2,64	7,23E-06
228245_s_at	similar to hCG38149	2,64	1,18E-06
239791_at	similar to hCG2042068	2,63	7,61E-05
1561737_at		2,59	2,52E-05
230743_at	hypothetical LOC404266	2,58	5,16E-06
225240_s_at	musashi homolog 2 (Drosophila)	2,56	2,58E-04
243996_at	natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic peptide receptor B)	2,53	5,16E-06
224880_at	v-ral simian leukemia viral oncogene homolog A (ras related)	2,52	7,61E-05
204798_at	v-myb myeloblastosis viral oncogene homolog (avian)	2,51	9,85E-05
1568964_x_at	sialoporphin	2,50	9,85E-05
219372_at	intraflagellar transport 81 homolog (Chlamydomonas)	2,49	5,84E-05
205677_s_at	deleted in lymphocytic leukemia 1 (non-protein coding)	2,47	3,38E-07
218792_s_at	B-box and SPRY domain containing	2,47	1,61E-04

239963_at		2,47	1,38E-05
217892_s_at	LIM domain and actin binding 1	2,46	5,16E-06
201397_at	phosphoglycerate dehydrogenase	2,45	8,23E-08
231852_at	exoribonuclease 1	2,45	3,23E-04
231319_x_at	kinesin family member 9	2,45	4,45E-05
212981_s_at	family with sequence similarity 115, member A; family with sequence similarity 115, member B (pseudogene)	2,43	4,97E-04
226043_at	G-protein signaling modulator 1 (AGS3-like, <i>C. elegans</i>)	2,42	7,61E-05
201664_at	structural maintenance of chromosomes 4	2,42	1,26E-04
227329_at	zinc finger and BTB domain containing 46	2,42	5,84E-05
1566482_at		2,42	6,13E-04
243579_at	musashi homolog 2 (<i>Drosophila</i>)	2,40	9,15E-04
228904_at	homeobox B3	2,40	4,45E-05
209537_at	exostosins (multiple)-like 2	2,40	5,84E-05
208798_x_at	golgi autoantigen, golgin subfamily a, 8B; golgi autoantigen, golgin subfamily a, 8A	2,38	6,13E-04
32723_at	cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kDa	2,38	7,92E-07
206148_at	interleukin 3 receptor, alpha (low affinity)	2,37	9,15E-04
225439_at	NudC domain containing 1	2,36	5,22E-07
200050_at	zinc finger protein 146	2,36	3,64E-06
213891_s_at	transcription factor 4	2,36	2,58E-04
236921_at	embigin homolog (mouse)	2,36	5,84E-05
223000_s_at	F11 receptor	2,35	1,64E-08
203362_s_at	MAD2 mitotic arrest deficient-like 1 (yeast)	2,34	2,52E-05
226339_at	TruB pseudouridine (psi) synthase homolog 1 (<i>E. coli</i>)	2,34	3,64E-06
225256_at		2,34	4,97E-04
220668_s_at	DNA (cytosine-5)-methyltransferase 3 beta	2,33	9,85E-05
244663_at		2,33	9,85E-05
215111_s_at	TSC22 domain family, member 1	2,32	7,51E-04
1554489_a_at	centrosomal protein 70kDa	2,30	4,97E-04
230670_at	immunoglobulin superfamily, member 10	2,30	9,15E-04
220798_x_at	plasticity-related gene 2	2,30	1,26E-04
226157_at	transcription factor Dp-2 (E2F dimerization partner 2)	2,30	2,58E-04
219703_at	meiosis-specific nuclear structural 1	2,29	7,61E-05
224959_at	solute carrier family 26 (sulfate transporter), member 2	2,29	2,58E-04
201294_s_at	WD repeat and SOCS box-containing 1	2,29	1,26E-04
225734_at	FBXO22 opposite strand (non-protein coding); F-box protein 22	2,29	3,36E-05
237388_at	glomulin, FKBP associated protein	2,29	4,45E-05
243707_at		2,28	7,61E-05
1569503_at	HEAT repeat containing 5B	2,28	3,23E-04
208407_s_at	catenin (cadherin-associated protein), delta 1	2,26	3,23E-04
206478_at	KIAA0125	2,26	4,02E-04
202946_s_at	BTB (POZ) domain containing 3	2,26	6,13E-04
229431_at	regulatory factor X-associated protein	2,25	1,87E-05
218643_s_at	cysteine-rich PDZ-binding protein	2,25	5,84E-05
201242_s_at	ATPase, Na ⁺ /K ⁺ transporting, beta 1 polypeptide	2,24	7,51E-04
235779_at	hypothetical protein LOC284408	2,24	3,36E-05
1558700_s_at	zinc finger protein 260	2,24	9,85E-05
231897_at	prostaglandin reductase 1	2,24	9,15E-04
212463_at	CD59 molecule, complement regulatory protein	2,23	4,97E-04
210999_s_at	growth factor receptor-bound protein 10	2,23	9,15E-04
225671_at	spinster homolog 2 (<i>Drosophila</i>)	2,23	1,26E-04
230298_at	metallo-beta-lactamase domain containing 2	2,22	9,85E-05
236953_s_at	NHL repeat containing 3	2,22	3,64E-06
1554973_a_at	zinc finger and BTB domain containing 26	2,22	9,15E-04
213541_s_at	v-ets erythroblastosis virus E26 oncogene homolog (avian)	2,22	9,15E-04
228974_at	zinc finger protein 677	2,21	1,61E-04
226120_at	tetratricopeptide repeat domain 8	2,21	7,92E-07
201111_at	CSE1 chromosome segregation 1-like (yeast)	2,21	9,15E-04
205063_at	survival of motor neuron protein interacting protein 1	2,21	7,92E-07
230281_at	chromosome 16 open reading frame 46	2,21	7,92E-07
212855_at	DCN1, defective in cullin neddylation 1, domain containing 4 (<i>S. cerevisiae</i>)	2,21	7,61E-05
230032_at	O-sialoglycoprotein endopeptidase-like 1	2,20	1,87E-05
242691_at		2,20	3,23E-04
225752_at	non imprinted in Prader-Willi/Angelman syndrome 1	2,20	1,26E-04
213056_at	FERM domain containing 4B	2,19	4,97E-04
225974_at	transmembrane protein 64	2,19	1,87E-05
1559477_s_at	Meis homeobox 1	2,19	6,13E-04
1554769_at	zinc finger protein 688; zinc finger protein 785	2,18	2,05E-04
241815_at		2,18	6,13E-04
211423_s_at	sterol-C5-desaturase (ERG3 delta-5-desaturase homolog, <i>S. cerevisiae</i>)-like	2,18	1,38E-05
219174_at	intraflagellar transport 74 homolog (<i>Chlamydomonas</i>)	2,18	6,13E-04
212905_at	cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kDa, tau variant	2,17	9,85E-05
234423_x_at		2,17	5,16E-06
224968_at	coiled-coil domain containing 104	2,17	2,54E-06
41329_at	SCY1-like 3 (<i>S. cerevisiae</i>)	2,17	1,61E-04
243462_s_at	BTB (POZ) domain containing 3	2,16	1,61E-04
226800_at	EF-hand calcium binding domain 7	2,16	3,23E-04
208002_s_at	acyl-CoA thioesterase 7	2,16	7,61E-05
209695_at	protein tyrosine phosphatase type IVA, member 3	2,16	1,38E-05
231995_at	chromosome 9 open reading frame 82	2,15	1,34E-07
229589_x_at	basic, immunoglobulin-like variable motif containing	2,15	7,51E-04
204142_at	enolase superfamily member 1	2,14	2,05E-04
204045_at	transcription elongation factor A (SII)-like 1	2,14	9,85E-05
224581_s_at	nuclear casein kinase and cyclin-dependent kinase substrate 1	2,14	1,00E-05
235346_at	FUN14 domain containing 1	2,14	3,38E-07
212604_at	mitochondrial ribosomal protein S31	2,13	3,36E-05
238712_at		2,13	1,61E-04
224819_at	transcription elongation factor A (SII)-like 8	2,13	6,13E-04
219169_s_at	transcription factor B1, mitochondrial	2,13	1,00E-05
212386_at	transcription factor 4	2,12	3,23E-04
219583_s_at	spermatogenesis associated 7	2,12	1,38E-05
239091_at	coiled-coil domain containing 52	2,12	1,26E-04

213645_at	enolase superfamily member 1	2,12	1,38E-05
218847_at	insulin-like growth factor 2 mRNA binding protein 2	2,12	7,61E-05
225086_at	family with sequence similarity 98, member B	2,12	2,05E-04
222266_at	chromosome 19 open reading frame 2	2,12	1,26E-04
228597_at	chromosome 21 open reading frame 45	2,12	1,74E-06
209146_at	sterol-C4-methyl oxidase-like	2,12	9,85E-05
231817_at	ubiquitin specific peptidase 53	2,11	4,02E-04
236917_at	leucine rich repeat containing 34	2,11	4,45E-05
227569_at	ligand of numb-protein X 2	2,11	9,15E-04
218889_at	nucleolar complex associated 3 homolog (S. cerevisiae)	2,11	3,23E-04
221652_s_at	chromosome 12 open reading frame 11	2,10	4,02E-04
225028_at	hypothetical LOC550643	2,10	9,15E-04
227301_at	chaperonin containing TCP1, subunit 6 (zeta) pseudogene 1	2,10	3,38E-07
1569540_at	nemo-like kinase	2,10	1,61E-04
202371_at	transcription elongation factor A (SII)-like 4	2,10	4,45E-05
208963_x_at	fatty acid desaturase 1	2,09	7,51E-04
218718_at	platelet derived growth factor C	2,09	3,23E-04
209143_s_at	chloride channel, nucleotide-sensitive, 1A	2,08	1,00E-05
232338_at	zinc finger protein 431	2,08	1,74E-06
209905_at	homeobox A9	2,08	4,02E-04
225537_at	trafficking protein particle complex 6B	2,08	3,36E-05
231970_at	chromosome 14 open reading frame 118	2,08	4,45E-05
225554_s_at	anaphase promoting complex subunit 7	2,08	3,36E-05
226583_at	chromosome 12 open reading frame 76	2,08	4,93E-08
202252_at	RAB13, member RAS oncogene family; similar to hCG24991	2,07	2,58E-04
217988_at	cyclin B1 interacting protein 1	2,07	4,02E-04
203763_at	dynein, cytoplasmic 2, light intermediate chain 1	2,07	2,54E-06
225835_at	solute carrier family 12 (sodium/potassium/chloride transporters), member 2	2,07	2,54E-06
218480_at	ATP/GTP binding protein-like 5	2,06	2,58E-04
219007_at	nucleoporin 43kDa	2,06	3,36E-05
231857_s_at	ATP/GTP binding protein-like 5	2,06	4,97E-04
229700_at	zinc finger protein 738	2,06	3,23E-04
222754_at	tRNA nucleotidyl transferase, CCA-adding, 1	2,06	4,02E-04
229533_x_at	zinc finger protein 680	2,06	7,61E-05
227798_at	SMAD family member 1	2,06	7,51E-04
209272_at	NGFI-A binding protein 1 (EGR1 binding protein 1)	2,06	3,36E-05
235132_at	hypothetical protein LOC254128	2,06	5,84E-05
211064_at	zinc finger protein 493	2,06	3,36E-05
229090_at	hypothetical LOC220930	2,05	9,85E-05
1560133_at	GRB10 interacting GYF protein 2	2,05	2,52E-05
222604_at	general transcription factor IIIc, polypeptide 3, 102kDa	2,04	7,92E-07
225581_s_at	mitochondrial ribosomal protein L50	2,04	1,61E-04
201263_at	threonyl-tRNA synthetase	2,04	7,23E-06
202862_at	fumarylacetoacetate hydrolase (fumarylacetoacetase)	2,04	3,23E-04
211464_x_at	caspase 6, apoptosis-related cysteine peptidase	2,04	4,02E-04
236924_at	glomulin, FKBP associated protein	2,04	2,54E-06
222762_x_at	LIM domains containing 1	2,03	2,52E-05
210145_at	phospholipase A2, group IVA (cytosolic, calcium-dependent)	2,03	6,13E-04
213089_at	ENSG00000219982	2,03	6,13E-04
210705_s_at	tripartite motif-containing 5	2,02	6,13E-04
218395_at	ARP6 actin-related protein 6 homolog (yeast)	2,02	2,15E-07
204162_at	NDC80 homolog, kinetochore complex component (S. cerevisiae)	2,02	7,51E-04
202020_s_at	LanC lantibiotic synthetase component C-like 1 (bacterial)	2,02	1,87E-05
201930_at	minichromosome maintenance complex component 6	2,02	5,16E-06
236704_at	hypothetical protein LOC100134230; similar to KIAA0454 protein; similar to phosphodiesterase 4D interacting protein isoform 2; phosphodiesterase 4D interacting protein	2,02	5,16E-06
227792_at	inositol 1,4,5-triphosphate receptor interacting protein-like 2	2,02	7,51E-04
201892_s_at	IMP (inosine monophosphate) dehydrogenase 2	2,01	6,13E-04
225421_at	peptidase M20 domain containing 2	2,00	4,02E-04

B

Probe set	Gene name	Ratio	P value
213006_at	CCAAT/enhancer binding protein (C/EBP), delta	10,22	2,52E-05
211571_s_at	versican	8,26	1,26E-04
203922_s_at	cytochrome b-245, beta polypeptide	8,23	7,23E-06
209728_at	major histocompatibility complex, class II, DR beta 4; major histocompatibility complex, class II, DR beta 1	7,98	5,84E-05
222496_s_at	RNA binding motif protein 47	7,61	5,84E-05
215646_s_at	versican	7,10	1,26E-04
228325_at	KIAA0146	6,86	1,61E-04
201670_s_at	myristoylated alanine-rich protein kinase C substrate	6,79	2,58E-04
1554406_a_at	C-type lectin domain family 7, member A	6,63	7,61E-05
219593_at	solute carrier family 15, member 3	6,63	2,05E-04
220088_at	complement component 5a receptor 1	6,61	3,23E-04
203923_s_at	cytochrome b-245, beta polypeptide	6,28	2,05E-04
223454_at	chemokine (C-X-C motif) ligand 16	6,22	5,84E-05
212657_s_at	interleukin 1 receptor antagonist	6,15	9,85E-05
212671_s_at	similar to HLA class II histocompatibility antigen, DQ(1) alpha chain precursor (DC-4 alpha chain); major histocompatibility complex, class II, DQ alpha 1	6,03	4,45E-05
213524_s_at	G0/G1switch 2	6,00	2,58E-04
205237_at	ficolin (collagen/fibrinogen domain containing) 1	5,98	9,85E-05
218831_s_at	Fc fragment of IgG, receptor, transporter, alpha	5,97	5,16E-06
1555756_a_at	C-type lectin domain family 7, member A	5,95	9,85E-05
220266_s_at	Kruppel-like factor 4 (gut)	5,88	3,36E-05
205789_at	CD1d molecule	5,87	2,52E-05
202878_s_at	CD93 molecule	5,66	9,85E-05
226818_at	macrophage expressed 1	5,61	4,45E-05
201743_at	CD14 molecule	5,57	4,97E-04
205098_at	chemokine (C-C motif) receptor 1	5,53	2,05E-04
203066_at	carbohydrate (N-acetylgalactosamine 4-sulfate 6-O) sulfotransferase 15	5,53	4,97E-04

221541_at	cysteine-rich secretory protein LCCL domain containing 2	5,41	4,97E-04
201739_at	serum/glucocorticoid regulated kinase 1	5,41	5,22E-07
218559_s_at	v-maf musculoaponeurotic fibrosarcoma oncogene homolog B (avian)	5,40	2,58E-04
221698_s_at	C-type lectin domain family 7, member A	5,37	9,85E-05
225897_at	myristoylated alanine-rich protein kinase C substrate	5,27	9,15E-04
222670_s_at	v-maf musculoaponeurotic fibrosarcoma oncogene homolog B (avian)	5,27	2,58E-04
204588_s_at	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7	5,25	1,61E-04
207697_x_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2	5,22	6,13E-04
238032_at	dehydrogenase/reductase (SDR family) member 3	5,04	5,22E-07
211101_x_at	leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	5,03	1,38E-05
217552_x_at	complement component (3b/4b) receptor 1 (Knops blood group)	4,97	1,26E-04
219505_at	cat eye syndrome chromosome region, candidate 1	4,96	2,52E-05
206631_at	prostaglandin E receptor 2 (subtype EP2), 53kDa	4,96	2,54E-06
214084_x_at	neutrophil cytosolic factor 1; neutrophil cytosolic factor 1C pseudogene	4,92	9,15E-04
203973_s_at	CCAAT/enhancer binding protein (C/EBP), delta	4,92	3,23E-04
221210_s_at	N-acetylneuraminate pyruvate lyase (dihydrodipicolinate synthase)	4,84	2,52E-05
205863_at	S100 calcium binding protein A12	4,84	7,51E-04
218035_s_at	RNA binding motif protein 47	4,82	7,61E-05
202833_s_at	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1	4,81	1,61E-04
212188_at	potassium channel tetramerisation domain containing 12	4,76	1,61E-04
204735_at	phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila)	4,73	5,22E-07
201669_s_at	myristoylated alanine-rich protein kinase C substrate	4,72	2,05E-04
231078_at	solute carrier family 25, member 37	4,71	1,87E-05
206584_at	lymphocyte antigen 96	4,70	1,26E-04
205067_at	interleukin 1, beta	4,69	1,26E-04
211962_s_at	zinc finger protein 36, C3H type-like 1	4,67	8,23E-08
211965_at	zinc finger protein 36, C3H type-like 1	4,57	3,38E-07
202637_s_at	intercellular adhesion molecule 1	4,54	1,87E-05
229937_x_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1	4,54	3,23E-04
1554624_a_at	signal-regulatory protein beta 1	4,53	1,61E-04
1556750_at	hypothetical protein LOC153577	4,47	3,23E-04
208092_s_at	family with sequence similarity 49, member A	4,45	1,74E-06
233555_s_at	sulfatase 2	4,45	3,36E-05
204912_at	interleukin 10 receptor, alpha	4,34	3,36E-05
202838_at	fucosidase, alpha-L- 1, tissue	4,25	2,05E-04
200866_s_at	prosaposin	4,22	1,26E-04
209823_x_at	major histocompatibility complex, class II, DQ beta 1; similar to major histocompatibility complex, class II, DQ beta 1	4,21	4,02E-04
202902_s_at	cathepsin S	4,14	7,61E-05
226841_at	macrophage expressed 1	4,11	6,13E-04
209304_x_at	growth arrest and DNA-damage-inducible, beta	4,10	1,38E-05
221841_s_at	Kruppel-like factor 4 (gut)	4,08	1,87E-05
205992_s_at	interleukin 15	4,07	1,74E-06
209305_s_at	growth arrest and DNA-damage-inducible, beta	4,07	2,58E-04
204103_at	chemokine (C-C motif) ligand 4	4,03	4,02E-04
201422_at	interferon, gamma-inducible protein 30	4,01	1,26E-04
210895_s_at	CD86 molecule	4,00	2,58E-04
202638_s_at	intercellular adhesion molecule 1	4,00	2,58E-04
201566_x_at	inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	3,99	1,34E-07
210904_s_at	interleukin 13 receptor, alpha 1	3,95	9,85E-05
229383_at	membrane-associated ring finger (C3HC4) 1	3,92	1,61E-04
209949_at	neutrophil cytosolic factor 2	3,92	9,15E-04
204620_s_at	versican	3,89	4,02E-04
208949_s_at	lectin, galactoside-binding, soluble, 3	3,88	9,85E-05
219947_at	C-type lectin domain family 4, member A	3,88	9,85E-05
212192_at	potassium channel tetramerisation domain containing 12	3,86	4,02E-04
203761_at	Src-like-adapter	3,85	4,97E-04
202295_s_at	cathepsin H	3,85	4,02E-04
205027_s_at	mitogen-activated protein kinase kinase kinase 8	3,85	5,16E-06
209286_at	CDC42 effector protein (Rho GTPase binding) 3	3,84	5,22E-07
205859_at	lymphocyte antigen 86	3,83	3,36E-05
224790_at	ArfGAP with SH3 domain, ankyrin repeat and PH domain 1	3,82	1,87E-05
218217_at	serine carboxypeptidase 1	3,81	6,13E-04
211100_x_at	leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	3,78	2,52E-05
213537_at	major histocompatibility complex, class II, DP alpha 1	3,77	1,61E-04
206488_s_at	CD36 molecule (thrombospondin receptor)	3,75	9,85E-05
235593_at	zinc finger E-box binding homeobox 2	3,73	1,87E-05
38037_at	heparin-binding EGF-like growth factor	3,71	2,58E-04
205099_s_at	chemokine (C-C motif) receptor 1	3,71	2,58E-04
209555_s_at	CD36 molecule (thrombospondin receptor)	3,70	2,05E-04
224724_at	sulfatase 2	3,70	5,84E-05
221731_x_at	versican	3,68	6,13E-04
225800_at	JAZF zinc finger 1	3,67	7,23E-06
217371_s_at	interleukin 15	3,67	7,92E-07
227645_at	phosphoinositide-3-kinase, regulatory subunit 5	3,67	7,23E-06
214366_s_at	arachidonate 5-lipoxygenase	3,66	4,97E-04
207857_at	leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	3,64	3,23E-04
211656_x_at	major histocompatibility complex, class II, DQ beta 1; similar to major histocompatibility complex, class II, DQ beta 1	3,63	1,26E-04
214590_s_at	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	3,63	7,51E-04
208450_at	lectin, galactoside-binding, soluble, 2	3,63	2,58E-04
202988_s_at	regulator of G-protein signaling 1	3,61	1,61E-04
228918_at	solute carrier family 43, member 2	3,61	6,13E-04
211135_x_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	3,61	6,13E-04
1564077_at	GPRIN family member 3	3,61	3,23E-04

203645_s_at	CD163 molecule	3,60	9,15E-04
202388_at	regulator of G-protein signaling 2, 24kDa	3,57	1,74E-06
209287_s_at	CDC42 effector protein (Rho GTPase binding) 3	3,55	1,74E-06
217388_s_at	kynureninase (L-kynurenine hydrolase)	3,55	9,85E-05
200838_at	cathepsin B	3,54	7,51E-04
223343_at	membrane-spanning 4-domains, subfamily A, member 7	3,52	7,51E-04
209683_at	family with sequence similarity 49, member A	3,52	4,97E-04
207574_s_at	growth arrest and DNA-damage-inducible, beta	3,51	4,45E-05
204440_at	CD83 molecule	3,50	3,64E-06
200785_s_at	low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)	3,49	4,45E-05
205419_at	G protein-coupled receptor 183	3,48	5,84E-05
209288_s_at	CDC42 effector protein (Rho GTPase binding) 3	3,47	1,18E-06
211429_s_at	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1	3,47	2,58E-04
235670_at	syntaxin 11	3,46	7,23E-06
210146_x_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2	3,46	9,15E-04
205639_at	acyloxyacyl hydrolase (neutrophil)	3,46	1,87E-05
225685_at	CDC42 effector protein (Rho GTPase binding) 3	3,42	3,64E-06
211612_s_at	interleukin 13 receptor, alpha 1	3,42	9,85E-05
210423_s_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	3,41	9,15E-04
203773_x_at	biliverdin reductase A	3,37	4,02E-04
242230_at	ataxin 1	3,36	1,61E-04
1555745_a_at	lysozyme (renal amyloidosis)	3,34	1,61E-04
204057_at	interferon regulatory factor 8	3,33	3,36E-05
218486_at	Kruppel-like factor 11	3,32	2,58E-04
1554899_s_at	Fc fragment of IgE, high affinity I, receptor for: gamma polypeptide	3,31	9,15E-04
222218_s_at	paired immunoglobulin-like type 2 receptor alpha	3,30	4,97E-04
230276_at	family with sequence similarity 49, member A	3,29	2,58E-04
209696_at	fructose-1,6-bisphosphatase 1	3,28	9,15E-04
203907_s_at	IQ motif and Sec7 domain 1	3,24	3,36E-05
203932_at	major histocompatibility complex, class II, DM beta	3,24	1,61E-04
242218_at	peroxisome proliferator-activated receptor delta	3,24	4,45E-05
204790_at	SMAD family member 7	3,23	5,16E-06
226140_s_at	OTU domain containing 1	3,21	9,85E-05
201531_at	zinc finger protein 36, C3H type, homolog (mouse)	3,19	3,64E-06
210663_s_at	kynureninase (L-kynurenine hydrolase)	3,17	7,51E-04
225798_at	JAZF zinc finger 1	3,15	7,61E-05
204834_at	fibrinogen-like 2	3,13	9,15E-04
214467_at	G protein-coupled receptor 65	3,12	2,54E-06
229521_at	hypothetical protein FLJ36031	3,11	7,92E-07
205686_s_at	CD86 molecule	3,09	7,51E-04
209037_s_at	EH-domain containing 1	3,09	1,26E-04
203708_at	phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	3,09	1,61E-04
210042_s_at	cathepsin Z	3,08	2,05E-04
221765_at	UDP-glucose ceramide glucosyltransferase	3,08	1,87E-05
224967_at	UDP-glucose ceramide glucosyltransferase	3,08	5,22E-07
223405_at	N-acetylneuraminate pyruvate lyase (dihydrodipicolinate synthase)	3,06	1,61E-04
213274_s_at	cathepsin B	3,00	4,45E-05
208540_x_at	S100 calcium binding protein A11; S100 calcium binding protein A11 pseudogene	3,00	7,51E-04
225142_at	jumonji C domain containing histone demethylase 1 homolog D (S. cerevisiae)	2,98	5,16E-06
1568955_at	SLIT-ROBO Rho GTPase activating protein 2 pseudogene 1	2,97	9,15E-04
203535_at	S100 calcium binding protein A9	2,96	7,51E-04
211991_s_at	major histocompatibility complex, class II, DP alpha 1	2,94	3,23E-04
202896_s_at	signal-regulatory protein alpha	2,92	9,15E-04
203388_at	arrestin, beta 2	2,92	7,61E-05
205685_at	CD86 molecule	2,91	2,58E-04
205786_s_at	integrin, alpha M (complement component 3 receptor 3 subunit)	2,89	2,05E-04
206420_at	immunoglobulin superfamily, member 6	2,89	6,13E-04
224583_at	coactosin-like 1 (Dictyostelium)	2,88	3,36E-05
211810_s_at	galactosylceramidase	2,88	2,58E-04
211133_x_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	2,86	7,51E-04
231035_s_at	OTU domain containing 1	2,85	4,02E-04
232617_at	cathepsin S	2,85	9,85E-05
223922_x_at	membrane-spanning 4-domains, subfamily A, member 6A	2,84	9,15E-04
229067_at	SLIT-ROBO Rho GTPase activating protein 2 pseudogene 1	2,83	2,05E-04
210360_s_at	metastasis suppressor 1	2,82	6,13E-04
215193_x_at	major histocompatibility complex, class II, DR beta 3	2,82	1,61E-04
211764_s_at	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	2,78	6,13E-04
211302_s_at	phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	2,78	7,51E-04
213002_at	myristoylated alanine-rich protein kinase C substrate	2,77	7,51E-04
212830_at	multiple EGF-like-domains 9	2,76	9,15E-04
214696_at	chromosome 17 open reading frame 91	2,74	1,38E-05
204867_at	GTP cyclohydrolase I feedback regulator	2,74	7,92E-07
226756_at	hypothetical protein FLJ36031	2,74	1,87E-05
208891_at	dual specificity phosphatase 6	2,74	6,13E-04
1555419_a_at	N-acylsphingosine amidohydrolase (acid ceramidase) 1	2,73	2,58E-04
221059_s_at	coactosin-like 1 (Dictyostelium)	2,73	2,05E-04
221920_s_at	solute carrier family 25, member 37	2,73	1,38E-05
211458_s_at	GABA(A) receptors associated protein like 3 (pseudogene); GABA(A) receptor-associated protein like 1	2,72	2,52E-05
208306_x_at	major histocompatibility complex, class II, DR beta 4; major histocompatibility complex, class II, DR beta 1	2,71	1,38E-05
226464_at	chromosome 3 open reading frame 58	2,71	1,18E-06
215435_at	ArfGAP with SH3 domain, ankyrin repeat and PH domain 1	2,71	4,97E-04
202201_at	biliverdin reductase B (flavin reductase (NADPH))	2,70	6,13E-04
240118_at	thioredoxin domain containing 11	2,68	1,26E-04
200871_s_at	prosaposin	2,67	2,58E-04
226952_at	ELL associated factor 1	2,66	2,05E-04

216248_s_at	nuclear receptor subfamily 4, group A, member 2	2,66	2,58E-04
209312_x_at	major histocompatibility complex, class II, DR beta 4; major histocompatibility complex, class II, DR beta 1	2,66	9,85E-05
219358_s_at	ArfGAP with dual PH domains 2	2,64	6,13E-04
201360_at	cystatin C	2,64	3,23E-04
226179_at	solute carrier family 25, member 37	2,64	9,85E-05
219622_at	RAB20, member RAS oncogene family	2,64	1,26E-04
201565_s_at	inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	2,63	1,00E-05
1555638_a_at	SAM domain, SH3 domain and nuclear localization signals 1	2,63	4,97E-04
1553137_s_at	Kruppel-like factor 11	2,62	7,92E-07
204621_s_at	nuclear receptor subfamily 4, group A, member 2	2,62	9,15E-04
227961_at	cathepsin B	2,60	4,97E-04
1557797_a_at	zinc finger E-box binding homeobox 2	2,59	4,02E-04
209535_s_at	A kinase (PRKA) anchor protein 13	2,59	1,34E-07
211729_x_at	biliverdin reductase A	2,58	9,15E-04
208892_s_at	dual specificity phosphatase 6	2,58	4,97E-04
224964_s_at	guanine nucleotide binding protein (G protein), gamma 2	2,58	3,23E-04
222221_x_at	EH-domain containing 1	2,57	1,61E-04
228685_at	hypothetical protein LOC150166	2,55	3,23E-04
204670_x_at	major histocompatibility complex, class II, DR beta 4; major histocompatibility complex, class II, DR beta 1	2,55	1,26E-04
204417_at	galactosylceramidase	2,55	1,74E-06
225955_at	meteorin, glial cell differentiation regulator-like; similar to meteorin, glial cell differentiation regulator-like	2,55	4,02E-04
200839_s_at	cathepsin B	2,55	4,97E-04
217362_x_at	major histocompatibility complex, class II, DR beta 6 (pseudogene)	2,54	1,00E-05
228555_at	calcium/calmodulin-dependent protein kinase II delta	2,54	3,23E-04
204118_at	CD48 molecule	2,54	9,85E-05
204122_at	TYRO protein tyrosine kinase binding protein	2,53	7,61E-05
213733_at	myosin IF	2,52	1,26E-04
1553134_s_at	chromosome 9 open reading frame 72	2,52	6,13E-04
1565598_at	RNA binding motif protein 47	2,51	1,61E-04
212737_at	GM2 ganglioside activator	2,51	2,58E-04
239331_at		2,51	9,15E-04
222326_at	phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	2,51	2,58E-04
207224_s_at	sialic acid binding Ig-like lectin 7	2,50	4,97E-04
236248_x_at	transcriptional adaptor 2 (ADA2 homolog, yeast)-beta	2,49	4,02E-04
225252_at	sulfiredoxin 1 homolog (S. cerevisiae)	2,49	1,61E-04
239105_at		2,48	1,00E-05
219582_at	opioid growth factor receptor-like 1	2,48	3,38E-07
204622_x_at	nuclear receptor subfamily 4, group A, member 2	2,47	2,58E-04
206028_s_at	c-mer proto-oncogene tyrosine kinase	2,47	9,15E-04
220987_s_at	chromosome 11 open reading frame 17	2,45	7,23E-06
203507_at	similar to eukaryotic translation initiation factor 4A; small nucleolar RNA, H/ACA box 67; eukaryotic translation initiation factor 4A, isoform 1	2,44	4,97E-04
215671_at	phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	2,44	4,02E-04
216942_s_at	CD58 molecule	2,44	2,58E-04
226810_at	opioid growth factor receptor-like 1	2,43	7,61E-05
226240_at	transcriptional adaptor 2 (ADA2 homolog, yeast)-beta	2,43	2,52E-05
200787_s_at	phosphoprotein enriched in astrocytes 15	2,42	3,36E-05
218611_at	immediate early response 5	2,42	2,54E-06
203217_s_at	ST3 beta-galactoside alpha-2,3-sialyltransferase 5	2,42	5,16E-06
212658_at	lipoma HMGIC fusion partner-like 2	2,41	2,58E-04
210176_at	toll-like receptor 1	2,41	1,61E-04
227897_at	RAP2B, member of RAS oncogene family	2,40	5,84E-05
216336_x_at	metallothionein 1L (gene/pseudogene); metallothionein 1E; metallothionein 1 pseudogene 3; metallothionein 1J (pseudogene)	2,40	3,23E-04
238469_at	opioid growth factor receptor-like 1	2,40	3,64E-06
242335_at	solute carrier family 25, member 37	2,40	2,52E-05
211990_at	major histocompatibility complex, class II, DP alpha 1	2,40	1,26E-04
225604_s_at	GLI pathogenesis-related 2	2,38	6,13E-04
220566_at	phosphoinositide-3-kinase, regulatory subunit 5	2,38	1,87E-05
235479_at	cytoplasmic polyadenylation element binding protein 2	2,37	1,61E-04
201926_s_at	CD55 molecule, decay accelerating factor for complement (Cromer blood group)	2,37	7,23E-06
208894_at	major histocompatibility complex, class II, DR alpha	2,37	2,58E-04
1559249_at	ataxin 1	2,36	4,45E-05
1552807_a_at	sialic acid binding Ig-like lectin 10	2,35	1,61E-04
221718_s_at	A kinase (PRKA) anchor protein 13	2,35	5,84E-05
202205_at	vasodilator-stimulated phosphoprotein	2,35	1,61E-04
208869_s_at	GABA(A) receptors associated protein like 3 (pseudogene); GABA(A) receptor-associated protein like 1	2,35	2,05E-04
223553_s_at	docking protein 3	2,33	2,05E-04
221840_at	protein tyrosine phosphatase, receptor type, E	2,32	1,61E-04
230492_s_at	hypothetical protein KIAA1434	2,32	6,13E-04
201543_s_at	SAR1 homolog A (S. cerevisiae)	2,32	7,51E-04
202901_x_at	cathepsin S	2,32	4,97E-04
1556385_at	cardiotrophin-like cytokine factor 1	2,32	7,61E-05
243819_at	guanine nucleotide binding protein (G protein), gamma 2	2,31	4,45E-05
207072_at	interleukin 18 receptor accessory protein	2,31	4,02E-04
1555950_a_at	CD55 molecule, decay accelerating factor for complement (Cromer blood group)	2,30	2,54E-06
216218_s_at	phospholipase C-like 2	2,29	9,15E-04
213988_s_at	spermidine/spermine N1-acetyltransferase 1	2,28	9,15E-04
210982_s_at	major histocompatibility complex, class II, DR alpha	2,28	2,05E-04
227935_s_at	polycomb group ring finger 5	2,28	7,61E-05
202192_s_at	growth arrest-specific 7	2,27	7,61E-05
212432_at	GrpE-like 1, mitochondrial (E. coli)	2,27	2,52E-05
208868_s_at	GABA(A) receptors associated protein like 3 (pseudogene); GABA(A) receptor-associated protein like 1	2,27	3,23E-04
214211_at	ferritin, heavy polypeptide 1; ferritin, heavy polypeptide-like 16; similar to ferritin, heavy polypeptide 1; ferritin, heavy polypeptide-like 3 pseudogene	2,27	7,51E-04

229756_at		2,27	4,45E-05
211744_s_at	CD58 molecule	2,27	1,61E-04
234574_at	myosin IF	2,27	1,61E-04
217168_s_at	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	2,26	5,84E-05
208872_s_at	receptor accessory protein 5	2,26	9,85E-05
212907_at	solute carrier family 30 (zinc transporter), member 1	2,26	9,15E-04
229733_s_at	chromobox homolog 6	2,25	6,13E-04
220330_s_at	SAM domain, SH3 domain and nuclear localization signals 1	2,24	4,97E-04
244492_at	Src-like-adaptor	2,24	2,52E-05
242738_s_at	zinc finger homeobox 3; hypothetical LOC100132068	2,24	9,15E-04
206571_s_at	mitogen-activated protein kinase kinase kinase kinase 4	2,23	2,54E-06
1559018_at	protein tyrosine phosphatase, receptor type, E	2,22	1,00E-05
244235_at	influenza virus NS1A binding protein	2,22	4,45E-05
217436_x_at	major histocompatibility complex, class I, A	2,22	2,05E-04
243931_at	CD58 molecule	2,22	1,61E-04
204638_at	acid phosphatase 5, tartrate resistant	2,21	6,07E-10
211404_s_at	amyloid beta (A4) precursor-like protein 2	2,21	9,15E-04
242263_at	transmembrane emp24 protein transport domain containing 5	2,20	1,74E-06
201850_at	capping protein (actin filament), gelsolin-like	2,20	5,84E-05
242320_at	forkhead box O3; forkhead box O3B pseudogene	2,20	1,26E-04
243395_at	CD55 molecule, decay accelerating factor for complement (Cromer blood group)	2,19	1,87E-05
226657_at	chromosome 17 open reading frame 103	2,17	9,15E-04
1556321_a_at		2,16	4,45E-05
217741_s_at	similar to zinc finger, AN1-type domain 5; zinc finger, AN1-type domain 5	2,16	5,22E-07
226968_at	kinesin family member 1B	2,16	1,61E-04
222507_s_at	TMEM9 domain family, member B	2,16	4,97E-04
208683_at	calpain 2, (m/II) large subunit	2,15	7,51E-04
217739_s_at	nicotinamide phosphoribosyltransferase	2,14	9,15E-04
224796_at	ArfGAP with SH3 domain, ankyrin repeat and PH domain 1	2,14	2,05E-04
202445_s_at	Notch homolog 2 (Drosophila)	2,14	2,05E-04
217868_s_at	methyltransferase like 9	2,13	2,52E-05
204183_s_at	adrenergic, beta, receptor kinase 2	2,13	1,38E-05
206302_s_at	nudix (nucleoside diphosphate linked moiety X)-type motif 4; nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1	2,13	3,36E-05
204881_s_at	UDP-glucose ceramide glucosyltransferase	2,12	1,26E-04
236297_at	chromosome 10 open reading frame 112	2,12	7,61E-05
225878_at	kinesin family member 1B	2,11	2,58E-04
225582_at	inositol 1,4,5-triphosphate receptor interacting protein	2,11	5,22E-07
221654_s_at	ubiquitin specific peptidase 3	2,10	1,87E-05
226843_s_at	PAP associated domain containing 5	2,10	3,38E-07
204194_at	BTB and CNC homology 1, basic leucine zipper transcription factor 1	2,09	1,26E-04
204088_at	purinergic receptor P2X, ligand-gated ion channel, 4	2,09	3,23E-04
219155_at	phosphatidylinositol transfer protein, cytoplasmic 1	2,09	2,54E-06
224826_at	hypothetical protein KIAA1434	2,09	4,97E-04
244485_at	major histocompatibility complex, class II, DP beta 1	2,09	9,15E-04
229043_at	PAP associated domain containing 5	2,08	5,16E-06
204500_s_at	ATP/GTP binding protein 1	2,08	2,58E-04
228284_at	similar to transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila); transducin-like enhancer of split 1 (E(s	2,08	4,97E-04
201599_at	ornithine aminotransferase (gyrate atrophy)	2,07	4,45E-05
223482_at	transmembrane protein 120A	2,07	7,51E-04
238063_at	transmembrane protein 154	2,07	7,51E-04
238800_s_at	zinc finger, CCHC domain containing 6	2,06	1,38E-05
226554_at	zinc finger and BTB domain containing 7A	2,06	2,05E-04
225251_at	RAB24, member RAS oncogene family	2,06	6,13E-04
208914_at	golgi associated, gamma adaptin ear containing, ARF binding protein 2	2,06	2,58E-04
36564_at	ring finger protein 19B	2,06	9,85E-05
229054_at	chromosome 14 open reading frame 181	2,05	3,23E-04
239404_at	topoisomerase (DNA) I	2,04	4,45E-05
221778_at	jumonji C domain containing histone demethylase 1 homolog D (S. cerevisiae)	2,04	4,97E-04
209124_at	myeloid differentiation primary response gene (88)	2,04	1,00E-05
242814_at	serpin peptidase inhibitor, clade B (ovalbumin), member 9	2,03	7,51E-04
225738_at	Rap guanine nucleotide exchange factor (GEF) 1	2,03	3,36E-05
205547_s_at	transgelin	2,03	6,13E-04
214574_x_at	leukocyte specific transcript 1	2,02	9,15E-04
223679_at	catenin (cadherin-associated protein), beta 1, 88kDa	2,02	9,15E-04

Supplementary table legend

Table S1

Differentially expressed probe sets between high and low *IGFBP2* expression groups (q4 vs. q1)

(A) Probe sets overexpressed in the high *IGFBP2* group. (B) Probe sets underexpressed in the high *IGFBP2* group.