

Probe set	Description	Gene symbol	Fold-change: TKD/WT
218478_s_at	zinc finger, CCHC domain containing 8	<i>ZCCHC8</i>	0.68
204847_at	zinc finger and BTB domain containing 11	<i>ZBTB11</i>	0.70
212665_at	TCDD-inducible poly(ADP-ribose) polymerase	<i>TIPARP</i>	0.54
203948_s_at	myeloperoxidase	<i>MPO</i>	0.18
203949_at	myeloperoxidase	<i>MPO</i>	0.20
207332_s_at	transferrin receptor (p90, CD71)	<i>TFRC</i>	0.58
237215_s_at	transferrin receptor (p90, CD71)	<i>TFRC</i>	0.51
1569652_at	Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3	<i>MLLT3</i>	0.26
204917_s_at	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3	<i>MLLT3</i>	0.37
237299_at	Transcribed locus		0.62
211699_x_at	hemoglobin, alpha 1 /// hemoglobin, alpha 2	<i>HBA1</i> /// <i>HBA2</i>	0.25
217414_x_at	hemoglobin, alpha 1 /// hemoglobin, alpha 2	<i>HBA1</i> /// <i>HBA2</i>	0.30
204018_x_at	hemoglobin, alpha 1 /// hemoglobin, alpha 2	<i>HBA1</i> /// <i>HBA2</i>	0.30
209458_x_at	hemoglobin, alpha 1 /// hemoglobin, alpha 2	<i>HBA1</i> /// <i>HBA2</i>	0.29
211745_x_at	hemoglobin, alpha 1	<i>HBA1</i>	0.30
203502_at	2,3-bisphosphoglycerate mutase	<i>BPGM</i>	0.37
211560_s_at	aminolevulinate, delta-, synthase 2 (sideroblastic/hypochromic anemia)	<i>ALAS2</i>	0.20
205592_at	interleukin 8	<i>IL8</i>	0.19
219672_at	erythroid associated factor	<i>ERAF</i>	0.18
210746_s_at	erythrocyte membrane protein band 4.2	<i>EPB42</i>	0.23
211820_x_at	glycophorin A (MNS blood group)	<i>GYP A</i>	0.17
223670_s_at	hemogen	<i>HEMGN</i>	0.19
211658_at	peroxiredoxin 2	<i>PRDX2</i>	0.41
39729_at	peroxiredoxin 2	<i>PRDX2</i>	0.15
205268_s_at	adducin 2 (beta)	<i>ADD2</i>	0.30
203116_s_at	ferrochelatase (protoporphyrin)	<i>FECH</i>	0.41
210504_at	Kruppel-like factor 1 (erythroid)	<i>KLF1</i>	0.32
239142_at	LOC317671	<i>LOC317671</i>	0.32
206698_at	X-linked Kx blood group (McLeod syndrome)	<i>XK</i>	0.19
221932_s_at	glutaredoxin 5 homolog (<i>S. cerevisiae</i>)	<i>GLRX5</i>	0.53
218883_s_at	MLF1 interacting protein	<i>MLF1IP</i>	0.46
202078_at	COP9 constitutive photomorphogenic homolog subunit 3 (<i>Arabidopsis</i>)	<i>COPS3</i>	0.68
212973_at	ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase)	<i>RPIA</i>	0.66
225051_at	erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)	<i>EPB41</i>	0.55
209273_s_at	HESB like domain containing 2	<i>HBLD2</i>	0.58
209274_s_at	HESB like domain containing 2	<i>HBLD2</i>	0.53
222666_s_at	RNA terminal phosphate cyclase-like 1	<i>RCLI</i>	0.52
205632_s_at	phosphatidylinositol-4-phosphate 5-kinase, type I, beta	<i>PIP5K1B</i>	0.30
218544_s_at	RNA terminal phosphate cyclase-like 1	<i>RCLI</i>	0.62
202947_s_at	glycophorin C (Gerbich blood group)	<i>GYP C</i>	0.30
237194_at	Tetraspanin 5	<i>TSPAN5</i>	0.48
212739_s_at	non-metastatic cells 4, protein expressed in	<i>NME4</i>	0.59
228378_at	chromosome 12 open reading frame 29	<i>C12orf29</i>	0.57
207112_s_at	GRB2-associated binding protein 1	<i>GAB1</i>	0.69

225998_at	GRB2-associated binding protein 1	<i>GABI</i>	0.62
236379_at	Erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)	<i>EPB41</i>	0.60
1554423_a_at	F-box protein 7	<i>FBXO7</i>	0.65
201178_at	F-box protein 7	<i>FBXO7</i>	0.67
206077_at	Kell blood group, metalloendopeptidase	<i>KEL</i>	0.36
208501_at	growth factor independent 1B (potential regulator of CDKN1A, translocated in CML)	<i>GFI1B</i>	0.36
1558942_at	zinc finger protein 765	<i>ZNF765</i>	0.71
208115_x_at	chromosome 10 open reading frame 137	<i>C10orf137</i>	0.64
202968_s_at	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	<i>DYRK2</i>	0.68
202969_at	MRNA; cDNA DKFZp667B0924 (from clone DKFZp667B0924)		0.63
218236_s_at	protein kinase D3	<i>PRKD3</i>	0.39
222565_s_at	protein kinase D3	<i>PRKD3</i>	0.52
226990_at	GPI-anchored membrane protein 1	<i>GPIAP1</i>	0.65
211081_s_at	mitogen-activated protein kinase kinase kinase 5	<i>MAP4K5</i>	0.64
208810_at	DnaJ (Hsp40) homolog, subfamily B, member 6	<i>DNAJB6</i>	0.64
208811_s_at	DnaJ (Hsp40) homolog, subfamily B, member 6	<i>DNAJB6</i>	0.65
220306_at	family with sequence similarity 46, member C	<i>FAM46C</i>	0.43
223708_at	C1q and tumor necrosis factor related protein 4	<i>C1QTNF4</i>	0.27
235061_at	protein phosphatase 1K (PP2C domain containing)	<i>PPM1K</i>	0.53
205284_at	KIAA0133	<i>KIAA0133</i>	0.64
222703_s_at	yrdC domain containing (E. coli)	<i>YRDC</i>	0.68
203795_s_at	B-cell CLL/lymphoma 7A	<i>BCL7A</i>	0.51
204244_s_at	DBF4 homolog (S. cerevisiae)	<i>DBF4</i>	0.61
235170_at	zinc finger protein 92	<i>ZNF92</i>	0.55
202919_at	preimplantation protein 3	<i>PREI3</i>	0.72
219198_at	general transcription factor IIIC, polypeptide 4, 90kDa	<i>GTF3C4</i>	0.67
218954_s_at	BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like	<i>BRF2</i>	0.72
221940_at	RNA pseudouridylylase domain containing 2	<i>RPUSD2</i>	0.72
230274_s_at	Nucleoporin 88kDa	<i>NUP88</i>	0.63
1557954_at	chromosome X open reading frame 15	<i>CXorf15</i>	0.64
227520_at	chromosome X open reading frame 15	<i>CXorf15</i>	0.62
212060_at	U2-associated SR140 protein	<i>SR140</i>	0.69
243176_at	CDNA FLJ30090 fis, clone BNGH41000015		0.68
221749_at	YTH domain family, member 3	<i>YTHDF3</i>	0.71
207941_s_at	RNA binding motif protein 39	<i>RBM39</i>	0.69
229428_at	Poly (ADP-ribose) glycohydrolase	<i>PARG</i>	0.66
202396_at	transcription elongation regulator 1	<i>TCERG1</i>	0.70
206095_s_at	FUS interacting protein (serine/arginine-rich) 1 /// similar to FUS interacting protein (serine-arginine rich) 1	<i>FUSIP1</i> /// <i>LOC642558</i>	0.69
221809_at	RAN binding protein 10	<i>RANBP10</i>	0.57
53987_at	RAN binding protein 10	<i>RANBP10</i>	0.58
202883_s_at	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	<i>PPP2R1B</i>	0.63
213899_at	methionyl aminopeptidase 2	<i>METAP2</i>	0.56
227993_at	methionyl aminopeptidase 2	<i>METAP2</i>	0.59
230129_at	chromosome 10 open reading frame 89	<i>C10orf89</i>	0.53
202353_s_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	<i>PSMD12</i>	0.57
212061_at	U2-associated SR140 protein	<i>SR140</i>	0.56
203358_s_at	enhancer of zeste homolog 2 (Drosophila)	<i>EZH2</i>	0.60
1554251_at	heterochromatin protein 1, binding protein 3	<i>HP1BP3</i>	0.62
221821_s_at	chromosome 12 open reading frame 41	<i>C12orf41</i>	0.57
226370_at	kelch-like 15 (Drosophila)	<i>KLHL15</i>	0.57
204568_at	KIAA0831	<i>KIAA0831</i>	0.63

228380_at	Transcribed locus, moderately similar to NP_689672.2 hypothetical protein MGC45438 [Homo sapiens]		0.67
210054_at	chromosome 4 open reading frame 15	<i>C4orf15</i>	0.57
209056_s_at	CDC5 cell division cycle 5-like (<i>S. pombe</i>)	<i>CDC5L</i>	0.72
213794_s_at	neuroguidin, EIF4E binding protein	<i>NGDN</i>	0.71
219212_at	heat shock 70kDa protein 14	<i>HSPA14</i>	0.62
235037_at	transmembrane protein 41A	<i>TMEM41A</i>	0.65
225455_at	transcriptional adaptor 1 (HFI1 homolog, yeast)-like	<i>TADA1L</i>	0.62
218127_at	nuclear transcription factor Y, beta	<i>NFYB</i>	0.62
218128_at	nuclear transcription factor Y, beta	<i>NFYB</i>	0.63
225906_at	CDNA FLJ38264 fis, clone FCBBF3001657		0.70
209139_s_at	protein kinase, interferon-inducible double stranded RNA dependent activator	<i>PRKRA</i>	0.72
228605_at	CDNA FLJ30010 fis, clone 3NB692000154		0.52
221735_at	WD repeat domain 48	<i>WDR48</i>	0.57
222157_s_at	WD repeat domain 48	<i>WDR48</i>	0.56
65591_at	WD repeat domain 48	<i>WDR48</i>	0.66
226687_at	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	<i>PRPF40A</i>	0.60
236477_at	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	<i>PRPF40A</i>	0.65
202268_s_at	amyloid beta precursor protein binding protein 1	<i>APPBP1</i>	0.72
244103_at	chromosome 1 open reading frame 55	<i>C1orf55</i>	0.59
208051_s_at	poly(A) binding protein interacting protein 1	<i>PAIP1</i>	0.44
209064_x_at	poly(A) binding protein interacting protein 1	<i>PAIP1</i>	0.45
222623_s_at	zinc finger protein 639	<i>ZNF639</i>	0.58
222624_s_at	zinc finger protein 639	<i>ZNF639</i>	0.54
209711_at	solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1	<i>SLC35D1</i>	0.56
212030_at	RNA binding motif protein 25	<i>RBM25</i>	0.65
202560_s_at	chromosome 1 open reading frame 77	<i>C1orf77</i>	0.70
221158_at	chromosome 21 open reading frame 66	<i>C21orf66</i>	0.69
204951_at	ras homolog gene family, member H	<i>RHOH</i>	0.36
204752_x_at	poly (ADP-ribose) polymerase family, member 2	<i>PARP2</i>	0.68
215773_x_at	poly (ADP-ribose) polymerase family, member 2	<i>PARP2</i>	0.69
221517_s_at	cofactor required for Sp1 transcriptional activation, subunit 6, 77kDa	<i>CRSP6</i>	0.55
223297_at	hypothetical protein MGC4268	<i>MGC4268</i>	0.62
1556060_a_at	KIAA1702 protein	<i>KIAA1702</i>	0.57
222430_s_at	YTH domain family, member 2	<i>YTHDF2</i>	0.68
230588_s_at	hypothetical protein LOC285074 /// similar to anaphase promoting complex subunit 1	<i>LOC285074</i> /// <i>LOC653663</i>	0.70
230479_at	Transcribed locus		0.57
240222_at	COP9 constitutive photomorphogenic homolog subunit 8 (<i>Arabidopsis</i>)	<i>COPS8</i>	0.68
202230_s_at	calcium homeostasis endoplasmic reticulum protein	<i>CHERP</i>	0.71
229594_at	SPT2, Suppressor of Ty, domain containing 1 (<i>S. cerevisiae</i>)	<i>SPTY2D1</i>	0.61
217804_s_at	interleukin enhancer binding factor 3, 90kDa	<i>ILF3</i>	0.73
226572_at	Suppressor of cytokine signaling 7	<i>SOCS7</i>	0.72
225157_at	MLX interacting protein	<i>MLXIP</i>	0.68
244304_at	hypothetical protein MGC42174	<i>MGC42174</i>	0.58
211375_s_at	interleukin enhancer binding factor 3, 90kDa	<i>ILF3</i>	0.70
244092_at	zinc finger, RAN-binding domain containing 3	<i>ZRANB3</i>	0.68
223005_s_at	chromosome 9 open reading frame 5	<i>C9orf5</i>	0.69
217830_s_at	NSFL1 (p97) cofactor (p47)	<i>NSFL1C</i>	0.69
216226_at	TAF4b RNA polymerase II, TATA box binding protein (TBP)-associated factor, 105kDa	<i>TAF4B</i>	0.58

235020_at	TAF4b RNA polymerase II, TATA box binding protein (TBP)-associated factor, 105kDa	<i>TAF4B</i>	0.45
219037_at	CGI-115 protein	<i>CGI-115</i>	0.56
205135_s_at	nuclear fragile X mental retardation protein interacting protein 1	<i>NUFIP1</i>	0.58
217961_at	hypothetical protein FLJ20551	<i>FLJ20551</i>	0.56
200993_at	importin 7	<i>IPO7</i>	0.65
200995_at	Importin 7	<i>IPO7</i>	0.54
222404_x_at	protein tyrosine phosphatase-like A domain containing 1	<i>PTPLAD1</i>	0.54
217777_s_at	protein tyrosine phosphatase-like A domain containing 1	<i>PTPLAD1</i>	0.52
222405_at	similar to butyrate-induced transcript 1	<i>LOC652500</i>	0.60
200052_s_at	interleukin enhancer binding factor 2, 45kDa	<i>ILF2</i>	0.69
201946_s_at	chaperonin containing TCP1, subunit 2 (beta)	<i>CCT2</i>	0.59
201947_s_at	chaperonin containing TCP1, subunit 2 (beta)	<i>CCT2</i>	0.61
229256_at	phosphoglucomutase 2-like 1	<i>PGM2L1</i>	0.48
218756_s_at	short-chain dehydrogenase/reductase	<i>MGC4172</i>	0.48
209826_at	EGF-like-domain, multiple 8 /// similar to palmitoyl-protein thioesterase 2	<i>EGFL8 ///</i> <i>LOC653870</i>	0.48
210473_s_at	G protein-coupled receptor 125	<i>GPR125</i>	0.33
213730_x_at	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	<i>TCF3</i>	0.66
210776_x_at	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	<i>TCF3</i>	0.61
213811_x_at	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	<i>TCF3</i>	0.64
239794_at	Isoleucine-tRNA synthetase	<i>IARS</i>	0.59
220607_x_at	TH1-like (Drosophila)	<i>TH1L</i>	0.70
225261_x_at	TH1-like (Drosophila)	<i>TH1L</i>	0.70
225006_x_at	TH1-like (Drosophila)	<i>TH1L</i>	0.70
225865_x_at	TH1-like (Drosophila)	<i>TH1L</i>	0.69
201919_at	Solute carrier family 25, member 36	<i>SLC25A36</i>	0.73
212348_s_at	amine oxidase (flavin containing) domain 2	<i>AOF2</i>	0.69
218192_at	inositol hexaphosphate kinase 2	<i>IHPK2</i>	0.58
47083_at	chromosome 7 open reading frame 26	<i>C7orf26</i>	0.74
223742_at	mitochondrial ribosomal protein L4	<i>MRPL4</i>	0.72
207515_s_at	polymerase (RNA) I polypeptide C, 30kDa	<i>POLR1C</i>	0.63
209317_at	polymerase (RNA) I polypeptide C, 30kDa	<i>POLR1C</i>	0.59
227617_at	hypothetical protein LOC199953	<i>RP13-15M17.2</i>	0.64
225857_s_at	hypothetical LOC388796	<i>LOC388796</i>	0.61
65588_at	hypothetical LOC388796	<i>LOC388796</i>	0.61
206341_at	interleukin 2 receptor, alpha	<i>IL2RA</i>	1.39
224065_at	homeodomain interacting protein kinase 2	<i>HIPK2</i>	1.40
242956_at	Isocitrate dehydrogenase 1 (NADP+), soluble	<i>IDH1</i>	1.49
1562321_at	pyruvate dehydrogenase kinase, isozyme 4	<i>PDK4</i>	1.55
210128_s_at	leukotriene B4 receptor	<i>LTB4R</i>	1.39
216176_at	hepatocellular carcinoma-related HCRP1	<i>HCRP1</i>	1.90
240265_at	TRAF3 interacting protein 3	<i>TRAF3IP3</i>	2.00
228968_at	zinc finger protein 449	<i>ZNF449</i>	1.43
1555317_at	polymerase (DNA directed) kappa	<i>POLK</i>	1.87
233122_at	Keratinocyte associated protein 2 /// MRNA; cDNA DKFZp686L09144 (from clone DKFZp686L09144)	<i>KRTCAP2</i>	1.79
205745_x_at	ADAM metallopeptidase domain 17 (tumor necrosis factor, alpha, converting enzyme)	<i>ADAM17</i>	1.58
205746_s_at	ADAM metallopeptidase domain 17 (tumor necrosis factor, alpha, converting enzyme)	<i>ADAM17</i>	1.90
217416_x_at			1.50

212587_s_at	protein tyrosine phosphatase, receptor type, C	<i>PTPRC</i>	1.79
218501_at	Rho guanine nucleotide exchange factor (GEF) 3	<i>ARHGEF3</i>	2.32
222799_at	HSPC049 protein	<i>HSPC049</i>	1.43
1552690_a_at	calcium channel, voltage-dependent, alpha 2/delta subunit 4	<i>CACNA2D4</i>	3.58
228083_at	calcium channel, voltage-dependent, alpha 2/delta subunit 4	<i>CACNA2D4</i>	4.64
224825_at	deoxynucleotidyltransferase, terminal, interacting protein 1	<i>DNTTIP1</i>	1.41
234942_s_at	deoxynucleotidyltransferase, terminal, interacting protein 1	<i>DNTTIP1</i>	1.64
217852_s_at	ADP-ribosylation factor-like 8B	<i>ARL8B</i>	1.76
222442_s_at	ADP-ribosylation factor-like 8B	<i>ARL8B</i>	1.80
207643_s_at	tumor necrosis factor receptor superfamily, member 1A	<i>TNFRSF1A</i>	2.05
221041_s_at	solute carrier family 17 (anion/sugar transporter), member 5	<i>SLC17A5</i>	1.76
227954_at	hypothetical protein LOC162073	<i>LOC162073</i>	1.49
201389_at	integrin, alpha 5 (fibronectin receptor, alpha polypeptide)	<i>ITGA5</i>	1.90
241464_s_at	FLJ27365 protein	<i>FLJ27365</i>	2.32
205366_s_at	homeobox B6	<i>HOXB6</i>	4.34
205601_s_at	homeobox B5	<i>HOXB5</i>	1.83
205179_s_at	ADAM metallopeptidase domain 8	<i>ADAM8</i>	2.32
214121_x_at	PDZ and LIM domain 7 (enigma)	<i>PDLIM7</i>	1.43
205774_at	coagulation factor XII (Hageman factor)	<i>F12</i>	2.03
226306_at	chromosome 6 open reading frame 1	<i>C6orf1</i>	1.46
1562608_at	Transmembrane protein 144	<i>TMEM144</i>	1.62
213375_s_at	hypothetical gene CG018	<i>CG018</i>	1.67
214906_x_at	hypothetical gene CG018	<i>CG018</i>	1.45
217197_x_at	hypothetical gene CG018	<i>CG018</i>	1.60
1567559_s_at	triggering receptor expressed on myeloid cells-like 4 /// triggering receptor expressed on myeloid cells-like 3	<i>TREML4</i> /// <i>TREML3</i>	1.57
242871_at	progesterone and adiponectin receptor family member V	<i>PAQR5</i>	1.38
214681_at	glycerol kinase	<i>GK</i>	1.57
237264_at	Ring finger protein 13	<i>RNF13</i>	1.47
202357_s_at	complement factor B	<i>CFB</i>	1.38
228923_at	S100 calcium binding protein A6 (calcyclin)	<i>S100A6</i>	1.88
1560011_at	prostate stem cell antigen	<i>PSCA</i>	2.01
214195_at	tripeptidyl peptidase I	<i>TPPI</i>	1.57
215151_at	dedicator of cytokinesis 10	<i>DOCK10</i>	1.68
240392_at	Caspase recruitment domain family, member 14	<i>CARD14</i>	2.23
202625_at	v-src tyrosine kinase viral related oncogene homolog	<i>LYN</i>	1.82
202626_s_at	v-src tyrosine kinase viral related oncogene homolog	<i>LYN</i>	1.69
212119_at	ras homolog gene family, member Q	<i>RHOQ</i>	1.67
212120_at	hypothetical LOC284988	<i>LOC284988</i>	1.84
212117_at	ras homolog gene family, member Q	<i>RHOQ</i>	1.58
214449_s_at	ras homolog gene family, member Q	<i>RHOQ</i>	1.64
1554503_a_at	osteoclast-associated receptor	<i>OSCAR</i>	2.92
203912_s_at	deoxyribonuclease I-like 1	<i>DNASE1L1</i>	1.43
219104_at	ring finger protein 141	<i>RNF141</i>	1.56
230361_at	hypothetical protein KIAA1833	<i>KIAA1833</i>	1.72
211416_x_at	gamma-glutamyltransferase-like activity 4	<i>GGTLA4</i>	1.49
215603_x_at	gamma-glutamyltransferase 1 /// gamma-glutamyltransferase 2 /// gamma-glutamyltransferase-like 4 /// similar to gamma-glutamyltransferase 2 /// similar to Gamma-glutamyltranspeptidase 1 precursor (Gamma-glutamyltransferase 1) (CD224 antigen)	<i>GGT1</i> /// <i>GGT2</i> /// <i>GGTLA4</i> /// <i>LOC643009</i> /// <i>LOC653257</i>	1.56
208284_x_at	gamma-glutamyltransferase 1	<i>GGT1</i>	1.75

211417_x_at	gamma-glutamyltransferase 1	<i>GGT1</i>	1.76
207131_x_at	gamma-glutamyltransferase 1	<i>GGT1</i>	1.87
209919_x_at	gamma-glutamyltransferase 1	<i>GGT1</i>	1.90
218700_s_at	RAB7, member RAS oncogene family-like 1	<i>RAB7L1</i>	1.91
226074_at	protein phosphatase 1M (PP2C domain containing)	<i>PPM1M</i>	1.67
206482_at	PTK6 protein tyrosine kinase 6	<i>PTK6</i>	1.63
205180_s_at	ADAM metalloproteinase domain 8	<i>ADAM8</i>	3.13
209906_at	complement component 3a receptor 1	<i>C3AR1</i>	3.62
219434_at	triggering receptor expressed on myeloid cells 1	<i>TREM1</i>	4.87
204575_s_at	matrix metalloproteinase 19 /// similar to Matrix metalloproteinase-19 precursor (MMP-19) (Matrix metalloproteinase RASI) (MMP-18)	<i>MMP19</i> /// <i>LOC652543</i>	2.29
204043_at	transcobalamin II; macrocytic anemia	<i>TCN2</i>	1.64
204158_s_at	T-cell, immune regulator 1, ATPase, H ⁺ transporting, lysosomal V0 subunit A3	<i>TCIRG1</i>	2.22
222173_s_at	TBC1 domain family, member 2	<i>TBC1D2</i>	1.96
1554285_at	hepatitis A virus cellular receptor 2	<i>HAVCR2</i>	2.40
221641_s_at	acyl-CoA thioesterase 9	<i>ACOT9</i>	1.65
212334_at	glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	<i>GNS</i>	2.68
212335_at	glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	<i>GNS</i>	2.97
200766_at	cathepsin D (lysosomal aspartyl peptidase)	<i>CTSD</i>	2.09
225372_at	chromosome 10 open reading frame 54	<i>C10orf54</i>	3.05
225373_at	chromosome 10 open reading frame 54	<i>C10orf54</i>	2.55
209370_s_at	SH3-domain binding protein 2	<i>SH3BP2</i>	1.68
244654_at	myosin IG	<i>MYO1G</i>	2.43
224846_at	SH3KBP1 binding protein 1	<i>SHKBP1</i>	1.55
211250_s_at	SH3-domain binding protein 2	<i>SH3BP2</i>	1.66
212516_at	centaurin, delta 2	<i>CENTD2</i>	1.51
34206_at	centaurin, delta 2	<i>CENTD2</i>	1.77
201126_s_at	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	<i>MGAT1</i>	1.66
210184_at	integrin, alpha X (complement component 3 receptor 4 subunit)	<i>ITGAX</i>	2.91
211286_x_at	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)	<i>CSF2RA</i>	2.08
222483_at	EF-hand domain family, member D2	<i>EFHD2</i>	1.84
205922_at	vanin 2	<i>VNN2</i>	3.67
217507_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	<i>SLC11A1</i>	2.32
1555116_s_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	<i>SLC11A1</i>	2.02
210423_s_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	<i>SLC11A1</i>	2.71
210422_x_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	<i>SLC11A1</i>	1.87
217473_x_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	<i>SLC11A1</i>	1.75
1554175_at	CD300 molecule-like family member b	<i>CD300LB</i>	2.45
228762_at	lunatic fringe homolog (Drosophila)	<i>LFNG</i>	2.09
1556747_a_at	CDNA FLJ39784 fis, clone SPLEN2002314		1.69
207270_x_at	CD300c molecule	<i>CD300C</i>	2.12
212948_at	calmodulin binding transcription activator 2	<i>CAMTA2</i>	1.57
231099_at	Solute carrier family 31 (copper transporters), member 2	<i>SLC31A2</i>	1.68
216322_at	CD58 molecule	<i>CD58</i>	1.73
211776_s_at	erythrocyte membrane protein band 4.1-like 3	<i>EPB41L3</i>	3.76
212681_at	erythrocyte membrane protein band 4.1-like 3	<i>EPB41L3</i>	3.11
205781_at	chromosome 16 open reading frame 7	<i>C16orf7</i>	1.50
206868_at	START domain containing 8	<i>STARD8</i>	1.59
218821_at	aminopeptidase-like 1	<i>NPEPL1</i>	1.58
220507_s_at	ureidopropionase, beta	<i>UPB1</i>	2.81
209446_s_at	chromosome 7 open reading frame 44	<i>C7orf44</i>	3.44
213700_s_at	Pyruvate kinase, muscle	<i>PKM2</i>	2.95

1568623_a_at	solute carrier family 35, member E4	<i>SLC35E4</i>	1.61
202877_s_at	CD93 molecule	<i>CD93</i>	2.87
240481_at	TRAF3 interacting protein 3	<i>TRAF3IP3</i>	1.75
206600_s_at	solute carrier family 16, member 5 (monocarboxylic acid transporter 6)	<i>SLC16A5</i>	1.72
213590_at	solute carrier family 16, member 5 (monocarboxylic acid transporter 6)	<i>SLC16A5</i>	1.62
206687_s_at	protein tyrosine phosphatase, non-receptor type 6	<i>PTPN6</i>	2.26
223591_at	ring finger protein 135	<i>RNF135</i>	1.84
223592_s_at	ring finger protein 135	<i>RNF135</i>	1.91
1558549_s_at	vanin 1	<i>VNN1</i>	3.64
205844_at	vanin 1	<i>VNN1</i>	8.73
204774_at	ecotropic viral integration site 2A	<i>EVI2A</i>	3.02
223686_at	thiamin pyrophosphokinase 1	<i>TPK1</i>	1.85
228341_at	CDNA FLJ34034 fis, clone FCBBF2004671		1.63
200625_s_at	CAP, adenylate cyclase-associated protein 1 (yeast)	<i>CAP1</i>	1.50
213798_s_at	CAP, adenylate cyclase-associated protein 1 (yeast)	<i>CAP1</i>	1.51
238063_at	transmembrane protein 154	<i>TMEM154</i>	1.92
209933_s_at	CD300a molecule	<i>CD300A</i>	2.31
217078_s_at	CD300a molecule	<i>CD300A</i>	2.18
217909_s_at	MAX-like protein X	<i>MLX</i>	1.50
217910_x_at	MAX-like protein X	<i>MLX</i>	1.40
210205_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4	<i>B3GALT4</i>	1.39
202855_s_at	solute carrier family 16, member 3 (monocarboxylic acid transporter 4)	<i>SLC16A3</i>	2.33
202856_s_at	solute carrier family 16, member 3 (monocarboxylic acid transporter 4)	<i>SLC16A3</i>	2.68
206235_at	ligase IV, DNA, ATP-dependent	<i>LIG4</i>	1.58
227766_at	ligase IV, DNA, ATP-dependent	<i>LIG4</i>	1.86
212685_s_at	transducin (beta)-like 2	<i>TBL2</i>	1.68
222127_s_at	exocyst complex component 1	<i>EXOC1</i>	1.46
219620_x_at	hypothetical protein FLJ20245	<i>FLJ20245</i>	1.51
223261_at	polymerase (DNA directed) kappa	<i>POLK</i>	1.45
203460_s_at	presenilin 1 (Alzheimer disease 3)	<i>PSEN1</i>	1.86
226577_at	CDNA FLJ36867 fis, clone ASTRO2016491		1.55
238816_at	CDNA FLJ36867 fis, clone ASTRO2016491		1.59
202422_s_at	acyl-CoA synthetase long-chain family member 4	<i>ACSL4</i>	2.11
207545_s_at	numb homolog (Drosophila)	<i>NUMB</i>	1.72
230462_at	numb homolog (Drosophila)	<i>NUMB</i>	2.04
203278_s_at	PHD finger protein 21A	<i>PHF21A</i>	1.43
230261_at	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 4	<i>ST8SIA4</i>	2.05
227098_at	dual specificity phosphatase 18	<i>DUSP18</i>	1.56
243969_at	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4	<i>SLC24A4</i>	1.70
205277_at	PR domain containing 2, with ZNF domain	<i>PRDM2</i>	1.66
207122_x_at	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2	<i>SULT1A2</i>	1.37
227352_at	chromosome 19 open reading frame 39	<i>C19orf39</i>	1.49
200761_s_at	ADP-ribosylation-like factor 6 interacting protein 5	<i>ARL6IP5</i>	1.66
1569004_at	Homo sapiens, Similar to neuronal thread protein, clone IMAGE:4106635, mRNA		1.68
242521_at	Homo sapiens, Similar to neuronal thread protein, clone IMAGE:4106635, mRNA		1.81