

Suppl Mat 3. 158 SG-enriched (red) and 226 IG-enriched transcripts (green)

Probe	Gene Name	Symbol	Unigene	Ttest Pair	Ttest Grp	ALR SG-IG	Fold
213920_at	cut-like 2 (Drosophila)	CUTL2	Hs.124953	0.00000	0.00000	4.80	27.94
206339_at	cocaine- and amphetamine-regulated transcript	CART	Hs.1707	0.00000	0.00000	3.80	13.96
205625_s_at	calbindin 1, 28kDa	CALB1	Hs.65425	0.00000	0.00000	3.34	10.16
205626_s_at	calbindin 1, 28kDa	CALB1	Hs.65425	0.00000	0.00000	3.23	9.36
206935_at	protocadherin 8	PCDH8	Hs.19492	0.00000	0.00000	3.17	9.00
231391_at	CDNA clone IMAGE:4611512, partial cds		Hs.66194	0.00000	0.00000	2.80	6.98
219463_at	chromosome 20 open reading frame 103	C20orf103	Hs.22920	0.00000	0.00000	2.77	6.84
1557475_at	CDNA clone IMAGE:5274197, partial cds		Hs.385496	0.00000	0.00000	2.62	6.13
243337_at	FRAS1 related extracellular matrix 3	FREM3	Hs.252714	0.00000	0.00000	2.60	6.06
243582_at	SH3 domain containing ring finger 2	SH3RF2	Hs.443728	0.00000	0.00000	2.34	5.06
230237_at	Adenylate cyclase activating polypeptide 1 (pituitary)	ADCYAP1	Hs.531719	0.00000	0.00000	2.27	4.84
221729_at	collagen, type V, alpha 2	COL5A2	Hs.445827	0.00000	0.00000	2.27	4.81
205923_at	reelin	RELN	Hs.132439	0.00000	0.00000	2.24	4.73
1552519_at	activin A receptor, type IC	ACVR1C	Hs.352338	0.00000	0.00000	2.24	4.72
215695_s_at	glycogenin 2	GYG2	Hs.541209	0.00000	0.00001	2.22	4.65
229247_at	hypothetical protein FLJ37440	FLJ37440	Hs.437696	0.00000	0.00007	2.09	4.27
243969_at	solute carrier family 24 (sodium/potassium/calcium exchanger)	SLC24A4	Hs.510281	0.00000	0.00000	2.09	4.25
238309_x_at	CDNA FLJ44654 fis, clone BRACE2047232		Hs.128823	0.00000	0.00003	2.07	4.21
1568870_at	Homo sapiens, clone IMAGE:4818211, mRNA		Hs.385530	0.00000	0.00000	2.06	4.17
221730_at	collagen, type V, alpha 2	COL5A2	Hs.445827	0.00004	0.00000	2.04	4.10
205428_s_at	calbindin 2, 29kDa (calretinin)	CALB2	Hs.106857	0.00000	0.00000	2.03	4.10
1556128_a_at	Ras protein-specific guanine nucleotide-releasing factor 2	RASGRF2	Hs.162129	0.00000	0.00000	2.00	4.01
222484_s_at	chemokine (C-X-C motif) ligand 14	CXCL14	Hs.483444	0.00000	0.00000	2.00	3.99
1560692_at	hypothetical protein LOC285878	LOC285878	Hs.548149	0.00000	0.00000	1.99	3.97
203510_at	met proto-oncogene (hepatocyte growth factor receptor)	MET	Hs.132966	0.00009	0.00000	1.99	3.96
213894_at	KIAA0960 protein	KIAA0960	Hs.120855	0.00000	0.00000	1.94	3.84
228109_at	Ras protein-specific guanine nucleotide-releasing factor 2	RASGRF2	Hs.162129	0.00000	0.00000	1.90	3.73
214920_at	hypothetical protein LOC221981	LOC221981		0.00000	0.00000	1.89	3.71
201063_at	reticulocalbin 1, EF-hand calcium binding domain	RCN1	Hs.97887	0.00000	0.00000	1.88	3.68
1554530_at	hypothetical protein MGC33530	MGC33530	Hs.335933	0.00000	0.00000	1.83	3.57
1563182_at	CDNA clone IMAGE:4796641, partial cds		Hs.541710	0.00000	0.00000	1.82	3.53
229159_at	KIAA0960 protein	KIAA0960	Hs.120855	0.00000	0.00000	1.80	3.47
230008_at	hypothetical protein LOC221981	LOC221981		0.00000	0.00000	1.78	3.43
222953_at	G protein-coupled receptor 83	GPR83	Hs.272385	0.00000	0.00000	1.77	3.40
230109_at	phosphodiesterase 7B	PDE7B	Hs.546364	0.00000	0.00000	1.73	3.33
233095_at	CDNA FLJ10095 fis, clone HEMBA1002430		Hs.272208	0.00000	0.00000	1.73	3.32
233135_at	Homo sapiens, clone IMAGE:4820713, mRNA /// Homo sapiens, clone IM		Hs.535863	0.00000	0.00000	1.71	3.27
230117_at	hypothetical protein LOC285878	LOC285878	Hs.548149	0.00000	0.00000	1.69	3.22
1553864_at	hypothetical protein DKFZp761H2121	DKFZp761	Hs.134065	0.00000	0.00000	1.67	3.19
236902_at	CDNA FLJ37214 fis, clone BRALZ2008484		Hs.445241	0.00000	0.00000	1.66	3.16
230238_at	hypothetical protein LOC134548	LOC134548	Hs.13308	0.00000	0.00000	1.66	3.16
206204_at	growth factor receptor-bound protein 14	GRB14	Hs.411881	0.00000	0.00000	1.65	3.14
232275_s_at	heparan sulfate 6-O-sulfotransferase 3	HS6ST3	Hs.171001	0.00000	0.00000	1.64	3.12
218002_s_at	chemokine (C-X-C motif) ligand 14	CXCL14	Hs.483444	0.00000	0.00007	1.64	3.12
236374_at	CDNA clone IMAGE:4611512, partial cds		Hs.66194	0.00000	0.00000	1.64	3.11
236308_at	hypothetical protein LOC285878	LOC285878	Hs.548149	0.00000	0.00000	1.63	3.09
232276_at	heparan sulfate 6-O-sulfotransferase 3	HS6ST3	Hs.171001	0.00000	0.00000	1.59	3.01
203416_at	CD53 antigen	CD53	Hs.443057	0.00000	0.00000	1.58	2.98
212768_s_at	olfactomedin 4	OLFM4	Hs.508113	0.00000	0.00000	1.57	2.98
1553865_a_at	hypothetical protein DKFZp761H2121	DKFZp761	Hs.134065	0.00000	0.00000	1.57	2.98
242524_at	cerebellin 4 precursor	CBLN4	Hs.126141	0.00000	0.00000	1.57	2.97
202291_s_at	matrix Gla protein	MGP	Hs.365706	0.00000	0.00000	1.57	2.96
206577_at	vasoactive intestinal peptide	VIP	Hs.53973	0.00000	0.00000	1.57	2.96
228913_at	LOC440135 /// Homo sapiens, clone IMAGE:2900578, mRNA, partial cds		Hs.375766	0.00000	0.00000	1.55	2.92
230882_at	hypothetical protein FLJ34048	FLJ34048	Hs.34969	0.00000	0.00003	1.51	2.86
211597_s_at	homeodomain-only protein /// homeodomain-only protein	HOP	Hs.121443	0.00000	0.00000	1.49	2.81
215073_s_at	nuclear receptor subfamily 2, group F, member 2	NR2F2	Hs.347991	0.00000	0.00000	1.48	2.79
235085_at	hypothetical protein DKFZp761P0423	DKFZp761	Hs.29068	0.00000	0.00000	1.48	2.79
220359_s_at	cyclic AMP-regulated phosphoprotein, 21 kD	ARPP-21	Hs.475902	0.00000	0.00000	1.47	2.78
230440_at	zinc finger protein 469	ZNF469	Hs.54925	0.00000	0.00000	1.47	2.78
209120_at	nuclear receptor subfamily 2, group F, member 2	NR2F2	Hs.347991	0.00000	0.00000	1.47	2.76
1553822_at	receptor transporting protein 1	RTP1	Hs.518480	0.00000	0.00000	1.47	2.76
205632_s_at	phosphatidylinositol-4-phosphate 5-kinase, type I, beta	PIP5K1B	Hs.534371	0.00000	0.00000	1.46	2.76
229714_at	heparan sulfate 6-O-sulfotransferase 3	HS6ST3	Hs.171001	0.00000	0.00000	1.44	2.72
201505_at	laminin, beta 1	LAMB1	Hs.489646	0.00000	0.00001	1.44	2.71
234331_s_at	NSE1	NSE1	Hs.260855	0.00000	0.00000	1.44	2.71
237556_at	Similar to BTG3 associated nuclear protein isoform b; BANP homolog; SM		Hs.460515	0.00000	0.00000	1.43	2.70

201141_at	glycoprotein (transmembrane) nmb	GNPMB	Hs.190495	0.00000	0.00270	1.42	2.67
1553823_a_at	receptor transporting protein 1	RTP1	Hs.518480	0.00000	0.00001	1.41	2.66
213791_at	proenkephalin	PENK	Hs.339831	0.00002	0.00000	1.41	2.65
213924_at	Metallophosphoesterase 1	MPPE1	Hs.514713	0.00000	0.00000	1.40	2.64
236677_at	neuroglobin	NGB	Hs.274363	0.00004	0.00004	1.39	2.63
228329_at	CDNA FLJ30779 fis, clone FEBRA2000815		Hs.4204	0.00000	0.00000	1.38	2.59
212552_at	hippocalcin-like 1	HPCAL1	Hs.467696	0.00000	0.00000	1.35	2.55
210021_s_at	uracil-DNA glycosylase 2	UNG2	Hs.3041	0.00013	0.00013	1.35	2.55
225975_at	protocadherin 18	PCDH18	Hs.518989	0.00000	0.00000	1.34	2.54
236714_at	Similar to Dedicator of cytokinesis protein 9 (Cdc42 guanine nucleotide ex		Hs.56400	0.00000	0.00000	1.34	2.54
237449_at	Sp8 transcription factor	SP8	Hs.195922	0.00000	0.00000	1.34	2.52
209763_at	chordin-like 1	CHRDL1	Hs.496587	0.00000	0.00008	1.33	2.51
214180_at	MRNA; cDNA DKFZp564H203 (from clone DKFZp564H203)		Hs.546727	0.00000	0.00000	1.33	2.51
243438_at	phosphodiesterase 7B	PDE7B	Hs.546364	0.00000	0.00000	1.31	2.49
240867_at	Regulatory factor X, 3 (influences HLA class II expression)	RFX3	Hs.136829	0.00000	0.00005	1.31	2.48
228742_at	Full length insert cDNA clone YQ54B06		Hs.133261	0.00000	0.00000	1.30	2.47
214091_s_at	glutathione peroxidase 3 (plasma)	GPX3	Hs.386793	0.00000	0.00003	1.30	2.47
233002_at	KIAA1622	KIAA1622	Hs.259599	0.00000	0.00000	1.30	2.46
227140_at	Inhibin, beta A (activin A, activin AB alpha polypeptide)	INHBA	Hs.28792	0.00000	0.00000	1.29	2.45
203661_s_at	tropomodulin 1	TMOD1	Hs.404289	0.00000	0.00000	1.29	2.45
223550_s_at	carbonic anhydrase X	CA10	Hs.463466	0.00000	0.00000	1.29	2.45
243139_at	CDNA FLJ37676 fis, clone BRHIP2012627		Hs.552096	0.00003	0.00000	1.29	2.45
1561691_at	hypothetical protein LOC285735	LOC285735	Hs.407549	0.00000	0.00004	1.28	2.43
236088_at	netrin G1	NTNG1	Hs.211236	0.00004	0.00000	1.27	2.41
234335_s_at	NSE1	NSE1	Hs.260855	0.00000	0.00000	1.27	2.40
219949_at	leucine rich repeat containing 2	LRRC2	Hs.517936	0.00000	0.00000	1.26	2.39
241320_at	R3H domain (binds single-stranded nucleic acids) containing	R3HDM	Hs.412462	0.00000	0.00208	1.24	2.36
230271_at	Homo sapiens, clone IMAGE:4512785, mRNA		Hs.391856	0.00000	0.00000	1.24	2.36
216086_at	synaptic vesicle glycoprotein 2C	SV2C	Hs.225974	0.00001	0.00003	1.24	2.36
235420_at	hyaluronan and proteoglycan link protein 4	HAPLN4	Hs.367829	0.00000	0.00000	1.22	2.33
238663_x_at	glutamate receptor, ionotropic, AMPA 4	GRIA4	Hs.503743	0.00000	0.00000	1.22	2.32
222664_at	potassium channel tetramerisation domain containing 15	KCTD15	Hs.221873	0.00000	0.00000	1.22	2.32
220313_at	G-protein coupled receptor 88	GPR88	Hs.170053	0.00000	0.00000	1.22	2.32
1560554_a_at				0.00001	0.00001	1.21	2.32
209119_x_at	nuclear receptor subfamily 2, group F, member 2	NR2F2	Hs.347991	0.00000	0.00000	1.20	2.30
240690_at	Hypothetical protein DKFZp761P0423	DKFZp761P0423	Hs.29068	0.00000	0.00000	1.19	2.29
236385_at	Cholinergic receptor, nicotinic, alpha polypeptide 7	CHRNA7	Hs.511772	0.00000	0.00000	1.18	2.26
233882_s_at	sema domain, transmembrane domain (TM), and cytoplasmic	SEMA6D	Hs.511265	0.00000	0.00009	1.18	2.26
212364_at	myosin IB	MYO1B	Hs.439620	0.00000	0.00000	1.17	2.26
1568803_at	Homo sapiens, clone IMAGE:5263088, mRNA		Hs.347841	0.00003	0.00010	1.17	2.25
212558_at	sprouty homolog 1, antagonist of FGF signaling (Drosophila)	SPRY1	Hs.436944	0.00000	0.00000	1.16	2.24
211651_s_at	laminin, beta 1 /// laminin, beta 1	LAMB1	Hs.489646	0.00000	0.00005	1.15	2.22
230403_at	CDNA FLJ45905 fis, clone OCBBF3026576		Hs.438858	0.00000	0.00000	1.15	2.21
1557667_at	CDNA FLJ36588 fis, clone TRACH2013991		Hs.49053	0.00000	0.00000	1.14	2.21
232111_at	Homo sapiens, clone IMAGE:4214962, mRNA		Hs.455955	0.00000	0.00000	1.14	2.21
201348_at	glutathione peroxidase 3 (plasma)	GPX3	Hs.386793	0.00000	0.00000	1.14	2.21
244617_at	G protein-coupled receptor 26	GPR26	Hs.12751	0.00000	0.00000	1.14	2.20
222809_x_at	chromosome 14 open reading frame 65	C14orf65	Hs.497591	0.00000	0.00000	1.14	2.20
203476_at	trophoblast glycoprotein	TPBG	Hs.82128	0.00004	0.00001	1.14	2.20
231859_at	chromosome 14 open reading frame 132	C14orf132	Hs.6434	0.00000	0.00001	1.13	2.18
242093_at				0.00000	0.00000	1.13	2.18
213325_at	poliovirus receptor-related 3	PVRL3	Hs.293917	0.00000	0.00000	1.12	2.18
215005_at	EF hand calcium binding protein 2	EFCBP2	Hs.140950	0.00000	0.00001	1.12	2.18
237094_at	family with sequence similarity 19 (chemokine (C-C motif)-like)	FAM19A5	Hs.436854	0.00000	0.00000	1.12	2.17
219140_s_at	retinol binding protein 4, plasma /// KIAA1922 protein	RBP4 /// KIAA1922	Hs.546471	0.00000	0.00001	1.11	2.16
241970_at	Poliovirus receptor-related 3	PVRL3	Hs.293917	0.00000	0.00001	1.11	2.16
238834_at	Transcribed locus		Hs.159282	0.00002	0.00000	1.11	2.16
205259_at	nuclear receptor subfamily 3, group C, member 2	NR3C2	Hs.163924	0.00000	0.00000	1.11	2.16
216268_s_at	jagged 1 (Alagille syndrome)	JAG1	Hs.224012	0.00000	0.00000	1.10	2.15
209493_at	PDZ domain containing 3	PDZK3	Hs.481819	0.00000	0.00000	1.10	2.15
212667_at	secreted protein, acidic, cysteine-rich (osteonectin)	SPARC	Hs.111779	0.00000	0.00001	1.10	2.14
209099_x_at	jagged 1 (Alagille syndrome)	JAG1	Hs.224012	0.00000	0.00000	1.10	2.14
231183_s_at	Jagged 1 (Alagille syndrome)	JAG1	Hs.224012	0.00000	0.00000	1.09	2.13
1566528_at	Cut-like 2 (Drosophila)	CUTL2	Hs.124953	0.00001	0.00007	1.09	2.13
203662_s_at	tropomodulin 1	TMOD1	Hs.404289	0.00000	0.00000	1.09	2.12
206282_at	neurogenic differentiation 1	NEUROD1	Hs.72981	0.00000	0.00000	1.08	2.12
206281_at	adenylate cyclase activating polypeptide 1 (pituitary)	ADCYAP1	Hs.531719	0.00000	0.00001	1.07	2.10
220325_at	TAF7-like RNA polymerase II, TATA box binding protein (TBP	TAF7L	Hs.223806	0.00000	0.00000	1.07	2.10
202421_at	immunoglobulin superfamily, member 3	IGSF3	Hs.171057	0.00000	0.00000	1.07	2.10

1566772_at	Schwannomin interacting protein 1	SCHIP1	Hs.134665	0.00000	0.00021	1.07	2.10
239673_at	Nuclear receptor subfamily 3, group C, member 2	NR3C2	Hs.163924	0.00000	0.00000	1.07	2.10
243526_at	hypothetical protein LOC349136	LOC349136	Hs.174373	0.00000	0.00044	1.07	2.10
202436_s_at	cytochrome P450, family 1, subfamily B, polypeptide 1	CYP1B1	Hs.154654	0.00000	0.00000	1.07	2.10
209822_s_at	very low density lipoprotein receptor	VLDLR	Hs.370422	0.00000	0.00000	1.06	2.09
220345_at	leucine rich repeat transmembrane neuronal 4	LRRTM4	Hs.285782	0.00000	0.00000	1.06	2.09
230788_at	glucosaminyl (N-acetyl) transferase 2, l-branching enzyme	GCNT2	Hs.519884	0.00016	0.00003	1.06	2.08
213452_at	zinc finger protein 184 (Kruppel-like)	ZNF184	Hs.158174	0.00000	0.00008	1.06	2.08
222668_at	potassium channel tetramerisation domain containing 15	KCTD15	Hs.221873	0.00001	0.00002	1.05	2.07
217671_at	Regulatory factor X, 3 (influences HLA class II expression)	RFX3	Hs.136829	0.00001	0.00026	1.05	2.07
205249_at	early growth response 2 (Krox-20 homolog, Drosophila)	EGR2	Hs.1395	0.00000	0.00161	1.05	2.06
220994_s_at	syntaxin binding protein 6 (amisyn)	STXBP6	Hs.508958	0.00000	0.00000	1.04	2.05
220889_s_at	carbonic anhydrase X	CA10	Hs.463466	0.00000	0.00000	1.03	2.05
226905_at	hypothetical protein MGC45871	MGC45871	Hs.345588	0.00000	0.00001	1.03	2.04
231439_at	NSE1	NSE1	Hs.260855	0.00000	0.00000	1.02	2.03
219461_at	p21(CDKN1A)-activated kinase 6	PAK6	Hs.513645	0.00000	0.00009	1.02	2.03
232720_at	Hypothetical protein FLJ31810	FLJ31810	Hs.145642	0.00000	0.00000	1.02	2.03
1555310_a_at	p21(CDKN1A)-activated kinase 6	PAK6	Hs.513645	0.00000	0.00007	1.02	2.03
219564_at	potassium inwardly-rectifying channel, subfamily J, member 16	KCNJ16	Hs.463985	0.00000	0.00018	1.02	2.02
203789_s_at	sema domain, immunoglobulin domain (Ig), short basic domain	SEMA3C	Hs.269109	0.00000	0.00000	1.01	2.02
237472_at	SRY (sex determining region Y)-box 1	SOX1	Hs.202526	0.00000	0.00000	1.00	2.00
219304_s_at	platelet derived growth factor l	PDGFD	Hs.352298	0.00001	0.00000	1.00	2.00
236870_at				0.00007	0.00004	-1.00	2.00
232856_at	FLJ45686 protein	FLJ45686	Hs.199853	0.00001	0.00001	-1.00	2.00
228776_at	gap junction protein, alpha 7, 45kDa (connexin 45)	GJA7	Hs.532593	0.00000	0.00000	-1.01	2.01
239468_at	chromosome 10 open reading frame 48	C10orf48	Hs.128193	0.00000	0.00000	-1.01	2.01
213880_at	leucine-rich repeat-containing G protein-coupled receptor 5	LGR5	Hs.172176	0.00001	0.00003	-1.01	2.01
239202_at	RAB3B, member RAS oncogene family	RAB3B	Hs.123072	0.00000	0.00000	-1.02	2.02
228060_at	solute carrier family 35, member F1	SLC35F1	Hs.549284	0.00000	0.00000	-1.02	2.02
1556325_at	filamin A interacting protein 1	FILIP1	Hs.526972	0.00000	0.00131	-1.02	2.03
233085_s_at	hypothetical protein FLJ22833	FLJ22833	Hs.118183	0.00000	0.00000	-1.02	2.03
204378_at	breast carcinoma amplified sequence 1	BCAS1	Hs.400556	0.00000	0.00000	-1.02	2.03
205011_at	loss of heterozygosity, 11, chromosomal region 2, gene A	LOH11CR1	Hs.152944	0.00008	0.00990	-1.02	2.03
202589_at	thymidylate synthetase	TYMS	Hs.369762	0.00011	0.00980	-1.03	2.04
219750_at	hypothetical protein FLJ11155	FLJ11155	Hs.176227	0.00006	0.00351	-1.03	2.04
218170_at	isochorismatase domain containing 1	ISOC1	Hs.483296	0.00000	0.00000	-1.03	2.04
223707_at	hypothetical protein MGC10850	MGC10850		0.00000	0.00022	-1.03	2.05
205590_at	RAS guanyl releasing protein 1 (calcium and DAG-regulated)	RASGRP1	Hs.511010	0.00000	0.00000	-1.03	2.05
210102_at	loss of heterozygosity, 11, chromosomal region 2, gene A	LOH11CR1	Hs.152944	0.00000	0.00000	-1.03	2.05
239144_at				0.00000	0.00000	-1.04	2.05
216617_s_at	myelin associated glycoprotein	MAG	Hs.515354	0.00022	0.00617	-1.04	2.05
221150_at	matrix, extracellular phosphoglycoprotein with ASARM motif (d	MEPE	Hs.189587	0.00002	0.00000	-1.04	2.06
238066_at	retinol binding protein 7, cellular	RBP7	Hs.422688	0.00000	0.00029	-1.05	2.07
213103_at	START domain containing 13	STARD13	Hs.507704	0.00000	0.00001	-1.06	2.08
204872_at	transducin-like enhancer of split 4 (E(sp1) homolog, Drosophila)	TLE4	Hs.444213	0.00000	0.00000	-1.06	2.09
229288_at	EPH receptor A7	EPHA7	Hs.73962	0.00000	0.00000	-1.06	2.09
235616_at	Chromosome 20 open reading frame 17	C20orf17	Hs.473117	0.00000	0.00001	-1.07	2.09
241772_at	Protocadherin 11 Y-linked	PCDH11Y	Hs.546448	0.00022	0.00018	-1.07	2.10
201012_at	annexin A1	ANXA1	Hs.494173	0.00008	0.00011	-1.07	2.10
225666_at	hypothetical protein FLJ14624	FLJ14624	Hs.190983	0.00000	0.00094	-1.07	2.10
204400_at	embryonal Fyn-associated substrate	EFS	Hs.24587	0.00000	0.00956	-1.07	2.10
229944_at	CDNA FLJ13136 fis, clone NT2RP3003139		Hs.106795	0.00001	0.00002	-1.08	2.11
230206_at	Dedicator of cytokinesis 5	DOCK5	Hs.195403	0.00000	0.00063	-1.08	2.11
1555958_at	cartilage acidic protein 1	CRTAC1	Hs.500741	0.00000	0.00115	-1.08	2.11
1568812_at	Homo sapiens, clone IMAGE:5294477, mRNA		Hs.382116	0.00000	0.00001	-1.08	2.12
226342_at	Spectrin, beta, non-erythrocytic 1	SPTBN1	Hs.503178	0.00000	0.00044	-1.09	2.12
226302_at	ATPase, Class I, type 8B, member 1	ATP8B1	Hs.216623	0.00000	0.00000	-1.09	2.12
236234_at	phosphodiesterase 1A, calmodulin-dependent	PDE1A	Hs.416061	0.00000	0.00000	-1.09	2.12
209771_x_at	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	CD24	Hs.375108	0.00004	0.00119	-1.10	2.14
1556200_a_at	chromosome 10 open reading frame 90	C10orf90	Hs.44939	0.00001	0.00092	-1.10	2.14
223557_s_at	transmembrane protein with EGF-like and two follistatin-like do	TMEFF2	Hs.144513	0.00000	0.00000	-1.10	2.15
228891_at	hypothetical protein LOC349236	LOC349236	Hs.494406	0.00000	0.00006	-1.11	2.15
216379_x_at	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	CD24	Hs.375108	0.00006	0.00140	-1.11	2.15
225727_at	pleckstrin homology domain containing, family H (with MyTH4	PLEKHH1	Hs.546414	0.00000	0.00957	-1.11	2.15
206456_at	gamma-aminobutyric acid (GABA) A receptor, alpha 5	GABRA5	Hs.24969	0.00000	0.00164	-1.11	2.16
235765_at	transducin-like enhancer of split 4 (E(sp1) homolog, Drosophila)	TLE4	Hs.444213	0.00000	0.00003	-1.12	2.17
226834_at	Adipocyte-specific adhesion molecule	ASAM	Hs.504187	0.00001	0.00000	-1.12	2.18
210657_s_at	septin 4	4-Sep	Hs.287518	0.00000	0.00401	-1.13	2.18
227856_at	hypothetical protein FLJ39370	FLJ39370	Hs.23439	0.00058	0.00062	-1.13	2.18

228438_at	CDNA FLJ41321 fis, clone BRAMY2045299		Hs.535929	0.00000	0.00004	-1.13	2.20
209842_at	SRY (sex determining region Y)-box 10	SOX10	Hs.376984	0.00000	0.00005	-1.14	2.20
228485_s_at	CDW92 antigen	CDW92	Hs.494700	0.00000	0.00520	-1.14	2.20
210247_at	synapsin II	SYN2	Hs.445503	0.00000	0.00001	-1.14	2.21
207659_s_at	myelin-associated oligodendrocyte basic protein	MOBP	Hs.121333	0.00000	0.00000	-1.14	2.21
223775_at	hedgehog interacting protein	HHIP	Hs.507991	0.00003	0.00103	-1.14	2.21
230207_s_at	Dedicator of cytokinesis 5	DOCK5	Hs.195403	0.00000	0.00088	-1.14	2.21
1561368_at	Homo sapiens, clone IMAGE:5194369, mRNA		Hs.427900	0.00008	0.00010	-1.15	2.21
233910_at	transmembrane protein with EGF-like and two follistatin-like do	TMEFF2	Hs.144513	0.00000	0.00039	-1.15	2.22
229228_at	cAMP responsive element binding protein 5	CREB5	Hs.437075	0.00000	0.00056	-1.15	2.22
230263_s_at	dedicator of cytokinesis 5	DOCK5	Hs.195403	0.00000	0.00144	-1.15	2.22
212922_s_at	SET and MYND domain containing 2	SMYD2	Hs.66170	0.00000	0.00000	-1.16	2.23
221868_at	KIAA1155 protein	KIAA1155	Hs.416735	0.00000	0.00000	-1.16	2.23
203042_at	lysosomal-associated membrane protein 2	LAMP2	Hs.496684	0.00000	0.00035	-1.16	2.24
205656_at	protocadherin 17	PCDH17	Hs.106511	0.00008	0.00007	-1.17	2.25
228653_at	Similar to RIKEN cDNA E130306M17 gene		Hs.269122	0.00000	0.00000	-1.17	2.25
213601_at				0.00000	0.00000	-1.17	2.25
227995_at	MRNA; cDNA DKFZp564O0862 (from clone DKFZp564O0862)		Hs.99472	0.00000	0.00000	-1.17	2.25
226136_at				0.00000	0.00001	-1.17	2.25
225822_at	hypothetical protein MGC17299	MGC17299	Hs.104476	0.00004	0.00125	-1.17	2.25
213273_at	odz, odd Oz/ten-m homolog 4 (Drosophila)	ODZ4	Hs.213087	0.00000	0.00000	-1.18	2.26
209072_at	myelin basic protein	MBP	Hs.501262	0.00000	0.00036	-1.18	2.27
208396_s_at	phosphodiesterase 1A, calmodulin-dependent	PDE1A	Hs.416061	0.00000	0.00000	-1.19	2.27
214064_at	transferrin	TF	Hs.518267	0.00000	0.00017	-1.19	2.28
1558541_at	hypothetical protein FLJ36980	FLJ36980	Hs.202521	0.00001	0.00364	-1.19	2.28
223349_s_at	BCL2-related ovarian killer	BOK	Hs.293753	0.00007	0.00028	-1.19	2.29
204529_s_at	thymus high mobility group box protein TOX	TOX	Hs.491805	0.00000	0.00003	-1.19	2.29
219429_at	fatty acid 2-hydroxylase	FA2H	Hs.461329	0.00000	0.00019	-1.20	2.29
205404_at	hydroxysteroid (11-beta) dehydrogenase 1	HSD11B1	Hs.195040	0.00004	0.00284	-1.20	2.30
204073_s_at	chromosome 11 open reading frame 9	C11orf9	Hs.473109	0.00000	0.00002	-1.21	2.31
207191_s_at	immunoglobulin superfamily containing leucine-rich repeat	ISLR	Hs.513022	0.00001	0.00000	-1.22	2.33
1552742_at	potassium voltage-gated channel, subfamily H (eag-related), n	KCNH8	Hs.475656	0.00000	0.00424	-1.22	2.34
233220_at	glutamate receptor, ionotropic, N-methyl-D-aspartate 3A	GRIN3A	Hs.283852	0.00000	0.00000	-1.23	2.34
231213_at	phosphodiesterase 1A, calmodulin-dependent	PDE1A	Hs.416061	0.00000	0.00001	-1.23	2.35
1557943_at	2',3'-cyclic nucleotide 3' phosphodiesterase	CNP	Hs.273621	0.00000	0.00007	-1.23	2.35
228807_at	Transcribed locus		Hs.80714	0.00000	0.00227	-1.24	2.36
213110_s_at	collagen, type IV, alpha 5 (Alport syndrome)	COL4A5	Hs.369089	0.00000	0.00421	-1.24	2.36
204519_s_at	plasma membrane proteolipid (plasmolipin)	PLLP	Hs.200821	0.00000	0.00600	-1.24	2.37
230537_at				0.00015	0.00093	-1.26	2.39
206498_at	oculocutaneous albinism II (pink-eye dilution homolog, mouse)	OCA2	Hs.130937	0.00000	0.00000	-1.26	2.39
228863_at	Hypothetical protein LOC144997	PCDH17	Hs.106511	0.00000	0.00000	-1.26	2.40
266_s_at	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	CD24	Hs.375108	0.00006	0.00177	-1.26	2.40
214475_x_at	calpain 3, (p94)	CAPN3	Hs.143261	0.00006	0.00817	-1.26	2.40
230863_at	Low density lipoprotein-related protein 2	LRP2	Hs.470538	0.00000	0.00000	-1.27	2.41
35974_at	lymphoid-restricted membrane protein	LRMP	Hs.124922	0.00000	0.00000	-1.27	2.41
238246_at	Metallothionein-like 5, testis-specific (tesmin)	MTL5	Hs.145932	0.00000	0.00005	-1.27	2.42
228984_at	KIAA1394 protein	KIAA1394	Hs.502982	0.00000	0.00631	-1.28	2.43
206135_at	suppression of tumorigenicity 18 (breast carcinoma) (zinc fing	ST18	Hs.549092	0.00000	0.00055	-1.28	2.43
206805_at	sema domain, immunoglobulin domain (Ig), short basic domain	SEMA3A	Hs.252451	0.00000	0.00009	-1.29	2.44
213059_at	cAMP responsive element binding protein 3-like 1	CREB3L1	Hs.405961	0.00000	0.00000	-1.29	2.45
226117_at	TRAF-interacting protein with a forkhead-associated domain	TIFA	Hs.310640	0.00000	0.00000	-1.29	2.45
204733_at	kallikrein 6 (neurosin, zyme)	KLK6	Hs.79361	0.00004	0.00016	-1.29	2.45
224794_s_at	cerebral endothelial cell adhesion molecule 1	CEECAM1	Hs.495230	0.00000	0.00009	-1.30	2.46
228601_at	hypothetical LOC401022	LOC40102	Hs.98661	0.00000	0.00006	-1.30	2.47
226213_at	v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (av	ERBB3	Hs.118681	0.00000	0.00007	-1.31	2.48
228090_at	nicotinamide nucleotide adenyltransferase 3	NMNAT3	Hs.208673	0.00000	0.00030	-1.32	2.49
209955_s_at	fibroblast activation protein, alpha	FAP	Hs.516493	0.00000	0.00001	-1.32	2.50
204621_s_at	nuclear receptor subfamily 4, group A, member 2	NR4A2	Hs.165258	0.00001	0.00001	-1.32	2.50
209631_s_at	G protein-coupled receptor 37 (endothelin receptor type B-like)	GPR37	Hs.406094	0.00000	0.00085	-1.33	2.51
232165_at	epiplakin 1	EPPK1	Hs.200412	0.00002	0.00001	-1.33	2.51
205989_s_at	myelin oligodendrocyte glycoprotein	MOG	Hs.141308	0.00003	0.00004	-1.33	2.51
226636_at	phospholipase D1, phosphatidylcholine-specific	PLD1	Hs.478230	0.00001	0.00051	-1.33	2.52
230135_at	CDNA FLJ42405 fis, clone ASTRO3000474		Hs.548089	0.00000	0.00055	-1.34	2.53
205801_s_at	RAS guanyl releasing protein 3 (calcium and DAG-regulated)	RASGRP3	Hs.143674	0.00000	0.00005	-1.34	2.53
211303_x_at	prostate-specific membrane antigen-like	PSMAL	Hs.511922	0.00000	0.00008	-1.34	2.54
227289_at	hypothetical protein LOC144997	LOC144997		0.00000	0.00055	-1.35	2.55
214803_at	CDNA clone IMAGE:4152983, partial cds		Hs.124776	0.00000	0.00000	-1.36	2.57
204037_at				0.00000	0.00002	-1.36	2.57
209869_at	adrenergic, alpha-2A-, receptor /// adrenergic, alpha-2A-, rece	ADRA2A	Hs.249159	0.00000	0.00001	-1.37	2.59

219414_at	calsyntenin 2	CLSTN2	Hs.158529	0.00000	0.00000	-1.38	2.60
223058_at	chromosome 10 open reading frame 45	C10orf45	Hs.446315	0.00000	0.00000	-1.38	2.60
204036_at	endothelial differentiation, lysophosphatidic acid G-protein-cou	EDG2	Hs.126667	0.00000	0.00030	-1.38	2.61
229781_at	LOC442530		Hs.304253	0.00009	0.00310	-1.39	2.63
214841_at	cornichon homolog 3 (Drosophila)	CNIH3	Hs.97338	0.00000	0.00000	-1.39	2.63
210839_s_at	ectonucleotide pyrophosphatase/phosphodiesterase 2 (autota	ENPP2	Hs.190977	0.00000	0.00352	-1.39	2.63
208358_s_at	UDP glycosyltransferase 8 (UDP-galactose ceramide galactos	UGT8	Hs.144197	0.00000	0.00000	-1.40	2.64
223741_s_at	tweety homolog 2 (Drosophila)	TTYH2	Hs.27935	0.00000	0.00314	-1.40	2.64
210944_s_at	calpain 3, (p94)	CAPN3	Hs.143261	0.00000	0.00009	-1.40	2.65
222608_s_at	anillin, actin binding protein (scraps homolog, Drosophila)	ANLN	Hs.62180	0.00000	0.00009	-1.40	2.65
228956_at	UDP glycosyltransferase 8 (UDP-galactose ceramide galactos	UGT8	Hs.144197	0.00001	0.00837	-1.41	2.65
242765_at	myelin-associated oligodendrocyte basic protein	MOBP	Hs.121333	0.00000	0.00599	-1.41	2.66
204204_at	solute carrier family 31 (copper transporters), member 2	SLC31A2	Hs.24030	0.00001	0.00915	-1.41	2.66
238763_at	RNA binding motif protein 20	RBM20	Hs.116630	0.00000	0.00000	-1.41	2.67
227550_at	hypothetical protein LOC143381	LOC143381	Hs.388347	0.00000	0.00133	-1.42	2.67
240841_at	insulinoma-associated 2	INSM2	Hs.62813	0.00000	0.00000	-1.42	2.67
204530_s_at	thymus high mobility group box protein TOX	TOX	Hs.491805	0.00000	0.00000	-1.42	2.68
241717_at	hypothetical protein LOC285281	LOC285281		0.00000	0.00000	-1.42	2.68
204777_s_at	mal, T-cell differentiation protein	MAL	Hs.80395	0.00000	0.00220	-1.43	2.69
34210_at	CD52 antigen (CAMPATH-1 antigen)	CD52	Hs.276770	0.00002	0.00000	-1.43	2.70
237187_at	Transcribed locus, strongly similar to XP_509406.1 similar to hypothetical		Hs.153034	0.00000	0.00000	-1.44	2.71
230087_at	Proline rich membrane anchor 1	PRIMA1	Hs.432401	0.00000	0.00015	-1.44	2.71
207323_s_at	myelin basic protein	MBP	Hs.501262	0.00002	0.00617	-1.44	2.72
213183_s_at	Cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	Hs.106070	0.00000	0.00000	-1.45	2.72
241898_at	Transcribed locus, moderately similar to XP_517655.1 similar to KIAA082		Hs.351126	0.00000	0.00000	-1.45	2.74
214586_at	G protein-coupled receptor 37 (endothelin receptor type B-like)	GPR37	Hs.406094	0.00001	0.00779	-1.46	2.75
238756_at	Growth arrest-specific 2 like 3	GAS2L3	Hs.20575	0.00000	0.00000	-1.46	2.75
205943_at	tryptophan 2,3-dioxygenase	TDO2	Hs.183671	0.00000	0.00000	-1.47	2.77
227123_at	RAB3B, member RAS oncogene family	RAB3B	Hs.123072	0.00000	0.00001	-1.49	2.82
206030_at	aspartoacylase (aminoacylase 2, Canavan disease)	ASPA	Hs.171142	0.00000	0.00051	-1.50	2.82
243154_at	Transcribed locus		Hs.86650	0.00000	0.00000	-1.51	2.84
214597_at	somatostatin receptor 2	SSTR2	Hs.514451	0.00000	0.00000	-1.51	2.84
217992_s_at	EF hand domain family, member D2	EFHD2	Hs.465374	0.00000	0.00000	-1.51	2.84
231911_at	KIAA1189	KIAA1189	Hs.443894	0.00000	0.00085	-1.51	2.85
230360_at	collomin	COLM	Hs.526441	0.00000	0.00059	-1.51	2.85
1570515_a_at	filamin A interacting protein 1	FILIP1	Hs.526972	0.00000	0.00000	-1.51	2.86
205303_at	potassium inwardly-rectifying channel, subfamily J, member 8	KCNJ8	Hs.102308	0.00001	0.00000	-1.52	2.87
212236_x_at	keratin 17	KRT17	Hs.2785	0.00000	0.00000	-1.53	2.88
205157_s_at	keratin 17	KRT17	Hs.2785	0.00011	0.00001	-1.53	2.89
211675_s_at	MyoD family inhibitor domain containing /// MyoD family inhibi	MDFIC	Hs.546385	0.00000	0.00000	-1.53	2.90
230593_at	Transcribed locus		Hs.128848	0.00000	0.00000	-1.54	2.90
212354_at	sulfatase 1	SULF1	Hs.409602	0.00000	0.00000	-1.55	2.92
239031_at	somatostatin receptor 2	SSTR2	Hs.514451	0.00000	0.00000	-1.56	2.95
228186_s_at	thrombospondin, type I, domain containing 2	THSD2	Hs.135254	0.00000	0.00000	-1.57	2.97
206343_s_at	neuregulin 1	NRG1	Hs.453951	0.00000	0.00000	-1.58	2.98
220120_s_at	erythrocyte membrane protein band 4.1 like 4A	EPB41L4A	Hs.551557	0.00000	0.00000	-1.58	2.99
38149_at	Rho GTPase activating protein 25	ARHGAP2	Hs.531807	0.00000	0.00000	-1.58	2.99
204774_at	ecotropic viral integration site 2A	EVI2A	Hs.113874	0.00000	0.00059	-1.58	2.99
205860_x_at	folate hydrolase (prostate-specific membrane antigen) 1	FOLH1	Hs.380325	0.00000	0.00097	-1.59	3.01
219643_at	low density lipoprotein-related protein 1B (deleted in tumors)	LRP1B	Hs.470117	0.00000	0.00000	-1.59	3.01
205549_at	Purkinje cell protein 4	PCP4	Hs.80296	0.00000	0.00000	-1.59	3.02
204674_at	lymphoid-restricted membrane protein	LRMP	Hs.124922	0.00000	0.00000	-1.60	3.02
236576_at	Transcribed locus		Hs.21375	0.00000	0.00000	-1.60	3.03
214063_s_at	transferrin	TF	Hs.518267	0.00000	0.00009	-1.60	3.04
232001_at	hypothetical gene supported by AY007155	LOC43994	Hs.13262	0.00000	0.00000	-1.63	3.09
215363_x_at	folate hydrolase (prostate-specific membrane antigen) 1	FOLH1	Hs.380325	0.00000	0.00022	-1.64	3.11
204882_at	Rho GTPase activating protein 25	ARHGAP2	Hs.531807	0.00000	0.00000	-1.64	3.11
225710_at	CDNA FLJ34013 fis, clone FCBBF2002111		Hs.173030	0.00000	0.00000	-1.65	3.15
210445_at	fatty acid binding protein 6, ileal (gastrotropin)	FABP6	Hs.519719	0.00000	0.00000	-1.68	3.21
244521_at	Transcribed locus		Hs.529382	0.00000	0.00000	-1.68	3.21
229552_at	hypothetical protein LOC283454	LOC283454	Hs.26605	0.00000	0.00000	-1.69	3.22
233072_at	netrin G2	NTNG2	Hs.163642	0.00000	0.00010	-1.71	3.28
226272_at	Full length insert cDNA clone ZD79H10		Hs.22380	0.00000	0.00000	-1.73	3.32
223699_at	carnosine dipeptidase 1 (metallopeptidase M20 family)	CNDP1	Hs.400613	0.00000	0.00019	-1.73	3.33
220595_at	PDZ domain containing RING finger 4	PDZRN4	Hs.380044	0.00000	0.00000	-1.74	3.33
1556378_a_at	hypothetical gene supported by BC032955 /// hypothetical gen	LOC40153	Hs.521524	0.00000	0.00001	-1.74	3.34
205304_s_at	potassium inwardly-rectifying channel, subfamily J, member 8	KCNJ8	Hs.102308	0.00000	0.00000	-1.76	3.40
228406_at	SET and MYND domain containing 1	SMYD1	Hs.516176	0.00000	0.00000	-1.77	3.40
235794_at	myelin-associated oligodendrocyte basic protein	MOBP	Hs.121333	0.00000	0.00000	-1.80	3.48

212353_at	sulfatase 1	SULF1	Hs.409602	0.00000	0.00000	-1.80	3.48
238577_s_at	Transcribed locus		Hs.32135	0.00000	0.00000	-1.83	3.57
227867_at	hypothetical protein LOC129293	LOC129293	Hs.469134	0.00000	0.00000	-1.86	3.63
221911_at	ets variant gene 1 /// hypothetical protein LOC221810	ETV1 /// LOC221810	Hs.22634	0.00000	0.00001	-1.87	3.65
210038_at	protein kinase C, theta	PRKCQ	Hs.498570	0.00000	0.00000	-1.87	3.66
240418_at	Transcribed locus		Hs.32135	0.00000	0.00000	-1.92	3.78
207307_at	5-hydroxytryptamine (serotonin) receptor 2C	HTR2C	Hs.149037	0.00000	0.00000	-1.98	3.93
213348_at	Cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	Hs.106070	0.00000	0.00000	-2.00	4.01
210121_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2	B3GALT2	Hs.518834	0.00000	0.00000	-2.05	4.13
210729_at	neuropeptide Y receptor Y2	NPY2R	Hs.37125	0.00000	0.00000	-2.07	4.19
214414_x_at	hemoglobin, alpha 2 /// hemoglobin, alpha 2	HBA2	Hs.398636	0.00005	0.00331	-2.07	4.21
217452_s_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2	B3GALT2	Hs.518834	0.00000	0.00000	-2.09	4.26
231945_at	filamin A interacting protein 1	FILIP1	Hs.526972	0.00000	0.00000	-2.16	4.47
226415_at	KIAA1576 protein	KIAA1576	Hs.461405	0.00000	0.00000	-2.26	4.78
235183_at	Homo sapiens, clone IMAGE:5312689, mRNA		Hs.548069	0.00000	0.00000	-2.26	4.80
206454_s_at	rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal recessive)	RHO	Hs.247565	0.00000	0.00000	-2.27	4.82
229580_at	Transcribed locus		Hs.7413	0.00000	0.00000	-2.29	4.90
213385_at	Chimerin (chimaerin) 2	CHN2	Hs.203663	0.00000	0.00000	-2.31	4.96
1552715_a_at	leucine-rich repeat-containing G protein-coupled receptor 7	LGR7	Hs.196119	0.00000	0.00000	-2.31	4.96
228915_at	dachshund homolog 1 (Drosophila)	DACH1	Hs.129452	0.00000	0.00000	-2.34	5.07
233972_s_at	zinc finger protein 312	ZNF312	Hs.241523	0.00000	0.00000	-2.35	5.08
229538_s_at	IQ motif containing GTPase activating protein 3	IQGAP3	Hs.133294	0.00000	0.00000	-2.41	5.30
229490_s_at	IQ motif containing GTPase activating protein 3	IQGAP3	Hs.133294	0.00000	0.00000	-2.42	5.35
204622_x_at	nuclear receptor subfamily 4, group A, member 2	NR4A2	Hs.165258	0.00000	0.00000	-2.45	5.46
205374_at	sarcolipin	SLN	Hs.334629	0.00000	0.00000	-2.47	5.55
218469_at	gremlin 1 homolog, cysteine knot superfamily (Xenopus laevis)	GREM1	Hs.40098	0.00000	0.00000	-2.53	5.76
229670_at	5.5 kb mRNA upregulated in retinoic acid treated HL-60 neutrophilic cells.		Hs.180284	0.00000	0.00000	-2.56	5.89
231804_at	Transcribed locus			0.00000	0.00000	-2.57	5.94
219370_at	reprimin 1, TP53 dependant G2 arrest mediator candidate	RPRM	Hs.100890	0.00000	0.00000	-2.62	6.16
216248_s_at	nuclear receptor subfamily 4, group A, member 2	NR4A2	Hs.165258	0.00000	0.00000	-2.65	6.30
235201_at	Transcribed locus			0.00000	0.00000	-2.79	6.93
210809_s_at	periostin, osteoblast specific factor	POSTN	Hs.136348	0.00000	0.00000	-2.82	7.05
243278_at	Transcribed locus		Hs.547776	0.00000	0.00000	-2.95	7.74
229070_at	chromosome 6 open reading frame 105	C6orf105	Hs.126409	0.00000	0.00000	-3.05	8.28
238206_at	Transcribed locus			0.00000	0.00000	-3.07	8.41
228206_at	heparan sulfate (glucosamine) 3-O-sulfotransferase 4	HS3ST4	Hs.549096	0.00000	0.00000	-3.09	8.52
1558971_at	chromosome 6 open reading frame 190	C6orf190	Hs.380210	0.00000	0.00000	-3.41	10.66
206941_x_at	sema domain, immunoglobulin domain (Ig), short basic domain	SEMA3E	Hs.528721	0.00000	0.00000	-3.61	12.20
219697_at	heparan sulfate (glucosamine) 3-O-sulfotransferase 2	HS3ST2	Hs.115830	0.00000	0.00000	-4.18	18.11
221086_s_at	zinc finger protein 312	ZNF312	Hs.241523	0.00000	0.00000	-4.50	22.56
1558972_s_at	chromosome 6 open reading frame 19	C6orf190	Hs.380210	0.00000	0.00000	-5.30	39.41