

Probe set	C1/C2	C1vC2 p-value	C1/C3	C1vC3 p-value	Gene symbol	Description
205225_at	16,57	p < 1e-07	15,85	p < 1e-07	ESR1	estrogen receptor 1
219197_s_at	11,31	1,04E-05	5,94	0,0002018	SCUBE2	signal peptide, CUB domain, EGF-like 2
209604_s_at	8,61	p < 1e-07	3,27	2,20E-06	GATA3	GATA binding protein 3
203963_at	8,37	2,00E-07	8,30	p < 1e-07	CA12	carbonic anhydrase XII
209603_at	7,64	p < 1e-07	4,24	1,40E-06	GATA3	GATA binding protein 3
209602_s_at	6,96	2,00E-07	3,03	0,0001004	GATA3	GATA binding protein 3
205696_s_at	6,17	1,01E-05	6,17	1,20E-06	GFRA1	GDNF family receptor alpha 1
210735_s_at	6,06	2,58E-05	4,39	7,17E-05	CA12	carbonic anhydrase XII
204508_s_at	5,88	8,00E-07	5,07	4,00E-07	CA12	carbonic anhydrase XII
203571_s_at	5,38	7,60E-06	3,16	0,0001853	C10orf116	chromosome 10 open reading frame 116
203929_s_at	5,09	0,0003715	4,01	0,0005628	MAPT	microtubule-associated protein tau
209460_at	5,00	0,000177	3,60	0,000537	ABAT	4-aminobutyrate aminotransferase
214053_at	4,05	0,0001311	3,01	0,000637		CDNA FLJ44318 fis, clone TRACH300C
211712_s_at	3,92	6,00E-07	2,33	0,00078	ANXA9	annexin A9 /// annexin A9
210652_s_at	3,68	7,18E-05	3,04	0,0002142	C1orf34	chromosome 1 open reading frame 34
204776_at	3,20	0,0019947	2,92	0,001266	THBS4	thrombospondin 4
218692_at	3,10	1,10E-06	3,17	3,20E-06	FLJ20366	hypothetical protein FLJ20366
210085_s_at	3,06	1,20E-06	1,90	0,0013326	ANXA9	annexin A9
212195_at	2,80	0,0011831	2,59	0,000916	IL6ST	interleukin 6 signal transducer (gp130, IL6R)
205116_at	2,68	0,000459	2,31	0,0007733	LAMA2	laminin, alpha 2 (merosin, congenital muscular dystrophy type 1B)
200810_s_at	2,62	3,37E-05	1,77	0,0015435	CIRBP	cold inducible RNA binding protein
206338_at	2,59	5,02E-05	1,87	0,0006175	ELAVL3	ELAV (embryonic lethal, abnormal vision)
217838_s_at	2,50	0,0002818	1,94	0,0012908	EVL	Enah/Vasp-like
213234_at	2,42	6,80E-06	1,82	0,0004548	KIAA1467	KIAA1467 protein
209769_s_at	2,41	0,0003829	2,01	0,0009563	GP1BB	glycoprotein Ib (platelet), beta polypeptide
205050_s_at	2,37	3,98E-05	2,10	9,70E-06	MAPK8IP2	mitogen-activated protein kinase 8 interacting protein 2
221934_s_at	2,31	2,00E-06	1,64	0,0005998	FLJ10496	hypothetical protein FLJ10496
215304_at	2,30	0,0002848	2,24	0,0001244		Human clone 23948 mRNA sequence
206278_at	2,27	8,68E-05	1,93	0,0002488	PTAFR	platelet-activating factor receptor
208682_s_at	2,22	0,0001352	1,74	0,0009878	MAGED2	melanoma antigen, family D, 2
203842_s_at	2,20	0,0002832	1,98	0,0002772	MAPRE3	microtubule-associated protein, RP/EB family class 3 member 3
204378_at	2,19	3,00E-07	1,60	7,73E-05	BCAS1	breast carcinoma amplified sequence 1
203670_at	2,19	0,0001633	1,78	0,000353	TTL3	tubulin tyrosine ligase-like family, member 3
207306_at	2,18	6,89E-05	1,83	0,0002596	TCF15	transcription factor 15 (basic helix-loop-helix)
40016_g_at	2,15	4,30E-06	1,75	7,14E-05	KIAA0303	KIAA0303 protein
217696_at	2,14	0,0003288	1,85	0,0003921	FUT7	fucosyltransferase 7 (alpha (1,3) fucosyltransferase)
205354_at	2,14	0,0001865	2,08	5,83E-05	GAMT	guanidinoacetate N-methyltransferase
205212_s_at	2,13	9,68E-05	1,75	0,0002644	CENTB1	centaurin, beta 1
206416_at	2,12	0,000148	1,85	0,0002187	ZNF205	zinc finger protein 205
222125_s_at	2,11	0,0001055	2,00	0,0005216	PH-4	hypoxia-inducible factor prolyl 4-hydroxylase
210684_s_at	2,10	0,0002629	1,91	0,0001535	DLG4	discs, large homolog 4 (Drosophila)
216289_at	2,10	0,0002215	1,84	0,0002813	GPR144	G protein-coupled receptor 144
215616_s_at	2,09	1,00E-06	1,75	4,70E-06	JMJD2B	jumonji domain containing 2B
209730_at	2,07	5,50E-05	1,92	2,48E-05	SEMA3F	sema domain, immunoglobulin domain
208868_s_at	2,06	4,74E-05	1,55	0,0010661	GABARAPL1	GABA(A) receptor-associated protein like 1
205141_at	2,05	0,0004125	1,91	0,0005375	ANG	angiogenin, ribonuclease, RNase A family
204862_s_at	2,05	0,0004588	1,70	0,0018047	NME3	non-metastatic cells 3, protein expressed in
211692_s_at	2,04	5,20E-05	1,92	1,91E-05	BBC3	BCL2 binding component 3 /// BCL2 binding component 3
217700_at	2,02	0,0002211	1,73	0,0005293		
213550_s_at	2,01	0,0004628	2,22	2,33E-05	PRO1580	hypothetical protein PRO1580
215844_at	2,00	0,0001526	1,77	0,0002054	TNPO2	transportin 2 (importin 3, karyopherin beta 2)
214122_at	2,00	0,000211	1,67	0,0013169		
221946_at	1,99	9,58E-05	1,60	0,0013917	C9orf116	chromosome 9 open reading frame 116
206328_at	1,99	0,0002984	1,88	0,0002345	CDH15	cadherin 15, M-cadherin (myotubule)
201841_s_at	1,96	0,0020072	1,75	0,0013311	HSPB1	heat shock 27kDa protein 1
217058_at	1,93	0,0002015	1,63	0,0016867	GNAS	GNAS complex locus
205451_at	1,91	0,0008512	1,80	0,0002204	MLLT7	myeloid/lymphoid or mixed-lineage leukemia 7
204482_at	1,91	0,0018012	1,96	0,0003043	CLDN5	claudin 5 (transmembrane protein deleted in colorectal cancer)
209697_at	1,90	0,0008172	1,78	0,0004519		
206650_at	1,87	0,0003576	1,75	0,0002134	IQCC	IQ motif containing C
208141_s_at	1,87	0,0004225	1,79	0,0001356	MGC4293	hypothetical protein MGC4293 /// hypothetical protein MGC4293
204878_s_at	1,85	0,0001578	1,65	0,0001033	TAO1	thousand and one amino acid protein kinase 1
204231_s_at	1,84	0,0002371	1,89	2,65E-05	FAAH	fatty acid amide hydrolase
220705_s_at	1,84	0,0003755	1,53	0,0015459	ADAMTS7	a disintegrin-like and metalloprotease (with thrombospondin type 1 motifs) 7
205074_at	1,84	1,80E-05	1,50	0,0009569	SLC22A5	solute carrier family 22 (organic cation/carnitine/cholesterol)
213690_s_at	1,83	0,0005455	1,70	0,000562	PTOV1	prostate tumor overexpressed gene 1
204876_at	1,82	4,75E-05	1,64	6,86E-05	KIAA0296	KIAA0296 gene product

216345_at	1,81	0,0014811	1,70	0,0008961	KIAA0913	KIAA0913
211248_s_at	1,81	5,48E-05	1,63	0,0001223	CHRD	chordin
208102_s_at	1,81	0,0003824	1,68	0,0001942	PSD	pleckstrin and Sec7 domain containing
221013_s_at	1,79	0,0020081	1,87	0,0001043	APOL2	apolipoprotein L, 2 /// apolipoprotein L,
204104_at	1,79	7,33E-05	1,72	1,42E-05	SNAPC2	small nuclear RNA activating complex,
202700_s_at	1,78	0,0005387	1,78	0,0001181	KIAA0792	KIAA0792 gene product
201452_at	1,77	0,0005989	1,57	0,0005311	RHEB	Ras homolog enriched in brain
205730_s_at	1,77	0,0006504	1,67	0,000342	ABLIM3	actin binding LIM protein family, membe
213319_s_at	1,75	0,0005517	1,55	0,0008811		
219849_at	1,75	0,0001031	1,52	0,0005805		
220755_s_at	1,74	0,001477	1,78	1,05E-05	C6orf48	chromosome 6 open reading frame 48
221163_s_at	1,74	0,0006284	1,56	0,0009569	WBSCR14	Williams Beuren syndrome chromosom
213651_at	1,74	0,0001063	1,49	0,0016783	PIB5PA	phosphatidylinositol (4,5) bisphosphate
216821_at	1,74	0,0002177	1,49	0,0024253		
205201_at	1,73	0,0004063	1,57	0,001131		
215712_s_at	1,72	0,0008733	1,76	3,41E-05	IGFALS	insulin-like growth factor binding proteir
201354_s_at	1,72	0,0005304	1,54	0,0008807		
214269_at	1,72	0,0009946	1,67	0,0002871	FLJ22269	hypothetical protein FLJ22269
209224_s_at	1,71	0,001266	1,59	0,0009895	NDUFA2	NADH dehydrogenase (ubiquinone) 1 ε
205441_at	1,70	0,0017104	1,71	7,08E-05	FLJ22709	hypothetical protein FLJ22709
216611_s_at	1,70	0,0007205	1,69	8,53E-05	SLC6A2	solute carrier family 6 (neurotransmitter
202312_s_at	1,69	0,0004789	1,49	0,0012541	COL1A1	collagen, type I, alpha 1
210025_s_at	1,69	0,0005205	1,46	0,0021124	CARD10	caspase recruitment domain family, me
215045_at	1,69	0,0015614	1,51	0,0014933	TNRC4	trinucleotide repeat containing 4
202488_s_at	1,68	0,001561	1,97	1,44E-05	FXYD3	FXYD domain containing ion transport i
212925_at	1,67	8,97E-05	1,48	0,0007695	C19orf21	chromosome 19 open reading frame 21
211512_s_at	1,66	0,0003242	1,58	8,18E-05	OGFR	opioid growth factor receptor
220024_s_at	1,66	0,0004698	1,49	0,0008219	PRX	periaxin
205211_s_at	1,64	0,0001279	1,59	4,96E-05		
216835_s_at	1,64	0,0008615	1,80	5,20E-06	DOK1	docking protein 1, 62kDa (downstream
218262_at	1,64	0,0002735	1,58	0,0002368	FLJ22318	hypothetical protein FLJ22318
217937_s_at	1,62	0,0001506	1,42	0,0016306	HDAC7A	histone deacetylase 7A
209280_at	1,61	0,0005154	1,43	0,0002125	MRC2	mannose receptor, C type 2
200895_s_at	1,61	0,001736	1,62	0,0009367	FKBP4	FK506 binding protein 4, 59kDa
205245_at	1,61	0,0002503	1,43	0,0015565	PARD6A	par-6 partitioning defective 6 homolog ε
201124_at	1,60	0,0009086	1,46	0,0019597	ITGB5	integrin, beta 5
206781_at	1,60	0,0013948	1,47	0,002455	DNAJC4	DnaJ (Hsp40) homolog, subfamily C, m
208104_s_at	1,60	0,0019761	1,53	0,0010521	THG-1	TSC-22-like /// TSC-22-like
214105_at	1,60	0,0005779	1,64	7,82E-05		
213731_s_at	1,59	0,0001993	1,45	0,000535		
89476_r_at	1,58	0,0010408	1,62	0,0002035	NPEPL1	aminopeptidase-like 1
217046_s_at	1,57	0,0003684	1,43	0,0017643	AGER	advanced glycosylation end product-sp
204144_s_at	1,56	0,0012864	1,47	0,0022785	PIGQ	phosphatidylinositol glycan, class Q
218707_at	1,55	0,0007241	1,60	9,36E-05	ZNF444	zinc finger protein 444
221557_s_at	1,55	0,0004885	1,52	0,0001506	LEF1	lymphoid enhancer-binding factor 1
220341_s_at	1,54	6,64E-05	1,56	4,10E-06	LOC51149	truncated calcium binding protein
205561_at	1,53	0,0004947	1,44	0,0005562	FLJ12242	hypothetical protein FLJ12242
209665_at	1,53	0,0013421	1,42	0,0018571	CYB561D2	cytochrome b-561 domain containing 2
217794_at	1,52	2,00E-07	1,28	0,000104	DKFZP564J15	DKFZp564J157 protein
205906_at	1,52	0,0007773	1,45	0,000729	FOXJ1	forkhead box J1
207353_s_at	1,51	0,0018762	1,42	0,0015018	HMX1	homeo box (H6 family) 1
213512_at	1,51	0,00047	1,61	9,70E-06	C14orf79	chromosome 14 open reading frame 79
214125_s_at	1,51	0,0010882	1,39	0,0024183		
202045_s_at	1,51	1,00E-07	1,25	0,0008971	GRLF1	glucocorticoid receptor DNA binding fac
205806_at	1,50	0,000761	1,60	2,10E-05	ROM1	retinal outer segment membrane protei
214002_at	1,50	9,00E-07	1,56	4,00E-07	MYL6	myosin, light polypeptide 6, alkali, smoc
217079_at	1,49	0,0019614	1,49	0,0003226		
214386_at	1,49	0,0006884	1,50	8,76E-05		
219057_at	1,47	0,0010077	1,36	0,0012537	RABEP2	rabaptin, RAB GTPase binding effector
215082_at	1,47	0,0010575	1,38	0,0010943	ELOVL5	ELOVL family member 5, elongation of
208176_at	1,46	0,001821	1,44	0,0007611	DUX1	double homeobox, 1
222228_s_at	1,46	0,0007969	1,46	1,07E-05	FLJ20013	hypothetical protein FLJ20013
219417_s_at	1,44	6,15E-05	1,39	5,10E-06	FLJ20014	hypothetical protein FLJ20014
220193_at	1,44	0,0011884	1,37	0,0009533	FLJ22938	hypothetical protein FLJ22938
203996_s_at	1,44	0,0002625	1,35	0,0006825	C21orf2	chromosome 21 open reading frame 2
210161_at	1,43	0,0008939	1,46	0,000276		
204746_s_at	1,43	0,0006258	1,48	2,06E-05	PRKCABP	protein kinase C, alpha binding protein
204954_s_at	1,43	0,0006438	1,56	9,70E-06	DYRK1B	dual-specificity tyrosine-(Y)-phosphoryl
204648_at	1,42	0,0003444	1,34	0,0004544	NPR1	natriuretic peptide receptor A/guanylate
217054_at	1,42	0,0012435	1,35	0,0024817		CDNA FLJ39484 fis, clone PROST2014

220061_at	1,42	0,0003323	1,30	0,0011668	FLJ20581	hypothetical protein FLJ20581
220430_at	1,40	0,0018631	1,42	0,0001148	FLJ14050	hypothetical protein FLJ14050
209322_s_at	1,39	0,0021262	1,35	0,0011566	SH2B	SH2-B homolog
204660_at	1,38	0,0008522	1,37	0,0002827	GFER	growth factor, augmenter of liver regenera
203682_s_at	1,38	0,0020853	1,55	0,0001552	IVD	isovaleryl Coenzyme A dehydrogenase
207004_at	1,37	0,0019432	1,47	0,0002903	BCL2	B-cell CLL/lymphoma 2
221232_s_at	1,37	0,0005102	1,27	0,001689	ANKRD2	ankyrin repeat domain 2 (stretch respon
204889_s_at	1,36	0,0002208	1,36	4,90E-05	NEURL	neuralized-like (Drosophila)
219766_at	1,36	0,0011097	1,32	0,0008085	MGC4093	hypothetical protein MGC4093
203586_s_at	1,36	0,0016477	1,32	0,0003952	ARF4L	ADP-ribosylation factor 4-like
212047_s_at	1,35	0,0018638	1,31	0,0007459	RNF167	ring finger protein 167
213433_at	1,35	0,0017691	1,33	0,0009181	ARL3	ADP-ribosylation factor-like 3
203273_s_at	1,32	0,0008243	1,22	0,0021979	TUSC2	tumor suppressor candidate 2
219986_s_at	1,32	0,000194	1,35	1,60E-06	ACAD10	acyl-Coenzyme A dehydrogenase famil
206739_at	1,26	2,24E-05	1,26	6,30E-06	HOXC5	homeo box C5
218257_s_at	0,74	0,0005943	0,62	2,00E-07	UGCGL1	UDP-glucose ceramide glucosyltransfe
212752_at	0,67	5,41E-05	0,71	6,65E-05	CLASP1	cytoplasmic linker associated protein 1
218420_s_at	0,67	0,0009414	0,69	5,86E-05	C13orf23	chromosome 13 open reading frame 23
203843_at	0,66	0,0020658	0,71	0,0022472	RPS6KA3	ribosomal protein S6 kinase, 90kDa, pc
221985_at	0,63	0,0002155	0,73	0,0012748	DRE1	DRE1 protein
203746_s_at	0,63	7,96E-05	0,68	0,0003097	HCCS	holocytochrome c synthase (cytochrom
203622_s_at	0,63	0,0008113	0,65	0,0003767	LOC56902	putative 28 kDa protein
220668_s_at	0,62	9,00E-06	0,61	1,83E-05	DNMT3B	DNA (cytosine-5-)-methyltransferase 3
220941_s_at	0,61	0,0001908	0,71	0,0012136	C21orf91	chromosome 21 open reading frame 91
202754_at	0,61	6,57E-05	0,65	1,35E-05	R3HDM	R3H domain (binds single-stranded nuc
219862_s_at	0,60	4,94E-05	0,55	3,88E-05	NARF	nuclear prelamin A recognition factor
208433_s_at	0,60	0,0003524	0,67	0,0002879	LRP8	low density lipoprotein receptor-related
203702_s_at	0,60	0,0014806	0,64	0,0009352	TTLL4	tubulin tyrosine ligase-like family, memt
201393_s_at	0,59	0,002052	0,47	1,11E-05	IGF2R	insulin-like growth factor 2 receptor
210466_s_at	0,58	0,0004903	0,71	0,0010032	PAI-RBP1	PAI-1 mRNA-binding protein
219498_s_at	0,58	6,00E-06	0,75	0,0004189	BCL11A	B-cell CLL/lymphoma 11A (zinc finger p
200693_at	0,57	6,70E-05	0,60	8,25E-05	YWHAQ	tyrosine 3-monooxygenase/tryptophan
212350_at	0,57	8,10E-06	0,69	5,88E-05	TBC1D1	TBC1 (tre-2/USP6, BUB2, cdc16) doma
202370_s_at	0,57	0,0014139	0,63	0,0012305	CBFB	core-binding factor, beta subunit
204094_s_at	0,56	0,0021206	0,57	0,0002786	KIAA0669	KIAA0669 gene product
202511_s_at	0,55	0,0010962	0,56	0,0011034	APG5L	APG5 autophagy 5-like (S. cerevisiae)
218566_s_at	0,55	0,0005554	0,58	0,0001403	CHORDC1	cysteine and histidine-rich domain (CH
217832_at	0,54	0,001183	0,62	0,0014086	SYNCRIP	synaptotagmin binding, cytoplasmic RN
212262_at	0,54	0,0012194	0,59	0,000456	QKI	quaking homolog, KH domain RNA bind
203139_at	0,54	8,20E-06	0,50	2,89E-05	DAPK1	death-associated protein kinase 1
212780_at	0,54	0,0009552	0,67	0,0001938	SOS1	son of sevenless homolog 1 (Drosophil
208398_s_at	0,54	0,0001142	0,55	0,000116	TBPL1	TBP-like 1
202188_at	0,54	1,45E-05	0,68	0,0006159	NUP93	nucleoporin 93kDa
202200_s_at	0,54	0,0005361	0,55	0,0002564	SRPK1	SFRS protein kinase 1
219675_s_at	0,53	0,0001389	0,67	0,0011532	UXS1	UDP-glucuronate decarboxylase 1
211764_s_at	0,53	8,53E-05	0,59	0,0002716	UBE2D1	ubiquitin-conjugating enzyme E2D 1 (U
218497_s_at	0,53	0,0003496	0,67	0,0001946	RNASEH1	ribonuclease H1
219960_s_at	0,52	0,0004888	0,63	0,000463	UCHL5	ubiquitin carboxyl-terminal hydrolase L5
204915_s_at	0,52	0,0015318	0,16	p < 1e-07	SOX11	SRY (sex determining region Y)-box 11
202147_s_at	0,52	0,0017243	0,59	0,0004569	IFRD1	interferon-related developmental regula
218092_s_at	0,52	0,0004283	0,49	8,30E-05	HRB	HIV-1 Rev binding protein
218826_at	0,52	2,49E-05	0,57	0,000117	SLC35F2	solute carrier family 35, member F2
221843_s_at	0,51	4,02E-05	0,66	0,0001168	KIAA1609	KIAA1609 protein
221676_s_at	0,51	0,0006185	0,54	0,0003082	CORO1C	coronin, actin binding protein, 1C
200821_at	0,51	0,0002944	0,59	0,0019771	LAMP2	lysosomal-associated membrane protei
212020_s_at	0,51	7,80E-06	0,67	0,0005729	MKI67	antigen identified by monoclonal antibo
202720_at	0,51	0,0004904	0,55	0,0003327	TES	testis derived transcript (3 LIM domains
218137_s_at	0,50	2,51E-05	0,55	3,56E-05	SMAP1	stromal membrane-associated protein
212263_at	0,50	7,78E-05	0,57	8,65E-05	QKI	quaking homolog, KH domain RNA bind
213102_at	0,50	0,0001722	0,58	0,0004905	ACTR3	ARP3 actin-related protein 3 homolog (
205339_at	0,49	3,50E-06	0,73	0,0006869	SIL	TAL1 (SCL) interrupting locus
203837_at	0,49	0,0004758	0,50	0,0001882	MAP3K5	mitogen-activated protein kinase kinase
212022_s_at	0,49	7,87E-05	0,61	0,0008591	MKI67	antigen identified by monoclonal antibo
203222_s_at	0,48	0,0014137	0,52	0,0002874	TLE1	transducin-like enhancer of split 1 (E(s
201833_at	0,48	0,0002381	0,47	6,95E-05	HDAC2	histone deacetylase 2
204198_s_at	0,47	0,0020493	0,50	0,0014011	RUNX3	runt-related transcription factor 3
205596_s_at	0,47	0,0001256	0,55	0,0007123	SMURF2	SMAD specific E3 ubiquitin protein liga
203315_at	0,47	1,10E-06	0,62	0,0001114	NCK2	NCK adaptor protein 2
218486_at	0,47	3,31E-05	0,67	0,0025557	TIEG2	TGFB inducible early growth response
218726_at	0,46	1,90E-06	0,69	0,00155	DKFZp762E1312	hypothetical protein DKFZp762E1312

220892_s_at	0,46	0,0004598	0,50	4,24E-05	PSAT1	phosphoserine aminotransferase 1
201197_at	0,45	0,0021375	0,50	0,000738	AMD1	adenosylmethionine decarboxylase 1
216640_s_at	0,45	6,38E-05	0,66	0,0005833	TXNDC7	thioredoxin domain containing 7 (protein)
203418_at	0,45	0,0004249	0,52	0,0005556	CCNA2	cyclin A2
222201_s_at	0,45	2,11E-05	0,61	0,0014686	CASP8AP2	CASP8 associated protein 2
219588_s_at	0,45	0,0001128	0,66	0,0012417	MTB	more than blood homolog
221677_s_at	0,44	0,000673	0,48	8,07E-05		
205891_at	0,43	3,10E-06	0,67	0,000115	ADORA2B	adenosine A2b receptor
220094_s_at	0,42	2,90E-05	0,63	0,0012499	C6orf79	chromosome 6 open reading frame 79
211967_at	0,42	0,0001122	0,48	9,06E-05	PORIMIN	pro-oncosis receptor inducing membrar
205668_at	0,41	9,59E-05	0,47	0,0004218	LY75	lymphocyte antigen 75
218319_at	0,40	5,75E-05	0,58	0,0020946	PEL1	pellino homolog 1 (Drosophila)
209205_s_at	0,40	0,000256	0,42	0,000152	LMO4	LIM domain only 4
203405_at	0,40	1,96E-05	0,60	0,0021432	DSCR2	Down syndrome critical region gene 2
217834_s_at	0,40	1,96E-05	0,61	0,0018575	SYNCRIP	synaptotagmin binding, cytoplasmic RN
220651_s_at	0,40	2,68E-05	0,49	0,0012885	MCM10	MCM10 minichromosome maintenance
200602_at	0,40	9,10E-05	0,55	0,0020847	APP	amyloid beta (A4) precursor protein (pr
201589_at	0,39	1,31E-05	0,56	0,0010059	SMC1L1	SMC1 structural maintenance of chrom
214953_s_at	0,39	8,17E-05	0,49	0,0010164	APP	amyloid beta (A4) precursor protein (pr
209773_s_at	0,38	0,0001084	0,35	2,08E-05	RRM2	ribonucleotide reductase M2 polypeptid
210024_s_at	0,38	0,0001766	0,44	0,0004001	UBE2E3	ubiquitin-conjugating enzyme E2E 3 (U
200790_at	0,37	7,50E-05	0,47	0,0002728	ODC1	ornithine decarboxylase 1
204602_at	0,36	0,0003786	0,24	0,0003412	DKK1	dickkopf homolog 1 (Xenopus laevis)
201976_s_at	0,36	0,0001534	0,47	0,0012703	MYO10	myosin X
203510_at	0,33	0,0002367	0,36	2,56E-05	MET	met proto-oncogene (hepatocyte growtl
201088_at	0,33	0,0002656	0,45	0,0025671	KPNA2	karyopherin alpha 2 (RAG cohort 1, im
201890_at	0,33	0,0008105	0,37	0,0005339	RRM2	ribonucleotide reductase M2 polypeptid
203574_at	0,33	7,40E-06	0,54	0,0017526		
219918_s_at	0,33	0,000238	0,49	0,0024498	ASPM	asp (abnormal spindle)-like, microceph
212276_at	0,30	6,00E-07	0,55	3,00E-07	LPIN1	lipin 1
209288_s_at	0,30	8,77E-05	0,50	0,002508	CDC42EP3	CDC42 effector protein (Rho GTPase b
205681_at	0,30	0,0005541	0,53	0,0025246	BCL2A1	BCL2-related protein A1
212274_at	0,30	1,30E-06	0,59	4,12E-05	LPIN1	lipin 1
201579_at	0,30	1,49E-05	0,53	0,0023277	FAT	FAT tumor suppressor homolog 1 (Dros
219497_s_at	0,29	p < 1e-07	0,63	0,0006702	BCL11A	B-cell CLL/lymphoma 11A (zinc finger p
202270_at	0,29	4,96E-05	0,43	0,0014354	GBP1	guanylate binding protein 1, interferon-i
209642_at	0,27	2,00E-07	0,56	0,0002211	BUB1	BUB1 budding uninhibited by benzimid
204962_s_at	0,25	3,80E-06	0,52	0,0021298	CENPA	centromere protein A, 17kDa
204822_at	0,19	p < 1e-07	0,43	4,48E-05		
204304_s_at	0,16	6,60E-05	0,20	1,12E-05	PROM1	prominin 1
205347_s_at	0,14	1,19E-05	0,27	5,75E-05	TMSNB	thymosin, beta, identified in neuroblast
215729_s_at	0,08	2,00E-07	0,39	0,0022683	VGLL1	vestigial like 1 (Drosophila)