

Probe set	C3/C1	C1vC3 p-value	C3/C2	C2vC3 p-value	Gene Symbol	Description
202917_s_at	19,93	3,50E-06	6,67	0,01	S100A8	S100 calcium binding protein A8 (calgranulin A)
205916_at	15,71	0,0001588	15,00	7,25E-03	S100A7	S100 calcium binding protein A7 (psoriasin 1)
217528_at	9,81	1,00E-07	11,42	6,05E-04	CLCA2	chloride channel, calcium activated, family member 2
203535_at	9,45	0,0001029	4,60	0,03	S100A9	S100 calcium binding protein A9 (calgranulin B)
206165_s_at	9,34	p < 1e-07	9,47	5,90E-04	CLCA2	chloride channel, calcium activated, family member 2
207802_at	7,82	0,0002749	11,28	4,62E-03	CRISP3	cysteine-rich secretory protein 3
206166_s_at	7,63	p < 1e-07	7,51	7,74E-04	CLCA2	chloride channel, calcium activated, family member 2
216836_s_at	7,23	1,40E-06	11,71	8,27E-05	ERBB2	v-erb-b2 erythroblastic leukemia viral oncogene homolog 2
214777_at	6,64	0,0001627	5,51	4,94E-03		immunoglobulin kappa light chain VKJ region mRNA, partial
215214_at	6,18	2,06E-05	3,64	0,02	IGLC2	immunoglobulin lambda constant 2 (Kern-Oz- marker)
206793_at	6,15	3,02E-05	8,02	5,40E-03	PNMT	phenylethanolamine N-methyltransferase
217388_s_at	5,98	2,10E-06	3,65	4,59E-03	KYNU	kynureninase (L-kynurenine hydrolase)
210930_s_at	5,82	4,34E-05	7,58	1,15E-03	ERBB2	v-erb-b2 erythroblastic leukemia viral oncogene homolog 2
219612_s_at	5,59	0,0017943	7,88	0,03	FGG	fibrinogen, gamma polypeptide
210761_s_at	5,42	1,00E-06	4,98	1,09E-03	GRB7	growth factor receptor-bound protein 7
208083_s_at	5,07	1,36E-05	4,30	9,31E-03	ITGB6	integrin, beta 6
211138_s_at	4,07	1,90E-05	4,44	5,15E-05	KMO	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
204818_at	3,98	1,10E-05	3,53	0,01	HSD17B2	hydroxysteroid (17-beta) dehydrogenase 2
205267_at	3,85	7,05E-05	2,97	8,54E-03	POU2AF1	POU domain, class 2, associating factor 1
216829_at	3,75	3,01E-05	3,39	7,23E-03	IGKC	immunoglobulin kappa constant
206164_at	3,63	8,00E-07	4,28	1,15E-03	CLCA2	chloride channel, calcium activated, family member 2
204941_s_at	3,21	0,0016419	4,41	2,20E-03	ALDH3B2	aldehyde dehydrogenase 3 family, member B2
212311_at	2,98	1,03E-05	1,88	0,02	KIAA0746	KIAA0746 protein
203767_s_at	2,89	3,34E-05	2,95	5,14E-03	STS	steroid sulfatase (microsomal), arylsulfatase C, isozyme S
212325_at	2,84	6,70E-06	2,67	6,40E-03	KIAA1102	KIAA1102 protein
210663_s_at	2,83	4,50E-06	2,84	1,18E-03	KYNU	kynureninase (L-kynurenine hydrolase)
55616_at	2,83	0,0002087	3,80	1,50E-03	PERLD1	per1-like domain containing 1
221811_at	2,79	0,0003469	4,19	1,20E-03	PERLD1	per1-like domain containing 1
200730_s_at	2,79	0,0006252	1,83	0,04	PTP4A1	protein tyrosine phosphatase type IVA, member 1
208840_s_at	2,73	0,0002022	2,07	0,04	G3BP2	Ras-GTPase activating protein SH3 domain-binding protein
201670_s_at	2,69	0,0003542	2,18	4,04E-04	MARCKS	myristoylated alanine-rich protein kinase C substrate
208084_at	2,68	5,69E-05	2,46	0,01	ITGB6	integrin, beta 6
200727_s_at	2,67	0,0008502	1,96	6,28E-03	ACTR2	ARP2 actin-related protein 2 homolog (yeast)
204942_s_at	2,56	0,0018467	5,94	7,39E-05	ALDH3B2	aldehyde dehydrogenase 3 family, member B2
203740_at	2,54	4,00E-07	2,05	3,77E-03	MPHOSPH6	M-phase phosphoprotein 6
202991_at	2,36	1,69E-05	3,10	1,22E-03	STARD3	START domain containing 3
219551_at	2,33	2,55E-05	2,39	4,95E-03	EAF2	ELL associated factor 2
201669_s_at	2,32	2,80E-06	1,36	0,02	MARCKS	myristoylated alanine-rich protein kinase C substrate
214446_at	2,31	0,0006656	2,93	9,18E-04	ELL2	elongation factor, RNA polymerase II, 2
215501_s_at	2,24	0,0015533	2,60	5,48E-05	DUSP10	dual specificity phosphatase 10
204132_s_at	2,22	4,53E-05	1,54	0,01	FOXO3A	forkhead box O3A
207734_at	2,18	9,88E-05	2,08	9,84E-03	LAX	hypothetical protein FLJ20340
201862_s_at	2,17	0,0006559	2,02	3,84E-03	LRRFIP1	leucine rich repeat (in FLII) interacting protein 1
204385_at	2,14	1,60E-06	1,98	4,81E-03	KYNU	kynureninase (L-kynurenine hydrolase)
203722_at	2,06	0,0004169	2,44	3,09E-03	ALDH4A1	aldehyde dehydrogenase 4 family, member A1
212611_at	2,05	0,0006428	2,06	0,01	DTX4	deltex 4 homolog (Drosophila)
211612_s_at	2,03	0,0007023	2,50	2,45E-04	IL13RA1	interleukin 13 receptor, alpha 1 /// interleukin 13 receptor, alpha 1
216607_s_at	2,02	0,0011431	1,79	0,01	CYP51A1	cytochrome P450, family 51, subfamily A, polypeptide 1
205248_at	2,01	0,0007228	2,27	2,18E-03	C21orf5	chromosome 21 open reading frame 5
218228_s_at	2,01	0,0001923	2,02	9,01E-04	TNKS2	tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2
208997_s_at	2,00	0,0004579	2,97	3,84E-03	UCP2	uncoupling protein 2 (mitochondrial, proton carrier)
205150_s_at	2,00	0,0015589	2,70	1,12E-03	KIAA0644	KIAA0644 gene product
202478_at	1,95	3,90E-05	1,60	0,02	TRIB2	tribbles homolog 2 (Drosophila)
211317_s_at	1,94	0,000489	1,91	0,01	CFLAR	CASP8 and FADD-like apoptosis regulator
200890_s_at	1,94	0,0004676	1,42	6,88E-03	SSR1	signal sequence receptor, alpha (translocon-associated protein)
212902_at	1,93	5,49E-05	1,71	0,01	SEC24A	SEC24 related gene family, member A (S. cerevisiae)
204562_at	1,93	0,0003998	1,81	0,02	IRF4	interferon regulatory factor 4
201204_s_at	1,93	0,0004024	1,73	4,43E-03	RRBP1	ribosome binding protein 1 homolog 180kDa (dog)
200825_s_at	1,92	0,0001201	1,44	0,01	HYOU1	hypoxia up-regulated 1
221563_at	1,92	0,0008559	1,87	2,37E-03	DUSP10	dual specificity phosphatase 10
202195_s_at	1,92	4,80E-05	1,63	3,61E-03	CGI-100	CGI-100 protein
211742_s_at	1,90	0,0013388	1,62	0,04	EVI2B	ecotropic viral integration site 2B /// ecotropic viral integrase
205351_at	1,82	8,95E-05	1,58	0,02	GGCX	gamma-glutamyl carboxylase
209186_at	1,80	0,0007813	1,51	0,01	ATP2A2	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
202067_s_at	1,79	0,0001686	1,63	7,37E-03	LDLR	low density lipoprotein receptor (familial hypercholesterolemia)
219155_at	1,75	0,0018147	1,92	0,01	PITPNC1	phosphatidylinositol transfer protein, cytoplasmic 1
209091_s_at	1,74	0,001123	1,55	4,83E-03	SH3GLB1	SH3-domain GRB2-like endophilin B1
208119_s_at	1,73	2,31E-05	1,60	0,02	ZNF505	zinc finger protein 505 /// zinc finger protein 505
200953_s_at	1,73	0,0003781	2,66	6,98E-04	CCND2	cyclin D2
218113_at	1,71	0,0004672	1,51	0,01	TMEM2	transmembrane protein 2
210555_s_at	1,69	1,54E-05	1,32	0,03	NFATC3	nuclear factor of activated T-cells, cytoplasmic, calcineurin-1
219033_at	1,66	0,0003211	1,89	6,96E-05	PARP8	poly (ADP-ribose) polymerase family, member 8
217168_s_at	1,65	0,0002648	1,58	2,41E-03	HERPUD1	homocysteine-inducible, endoplasmic reticulum stress-inducible
212415_at	1,63	0,0006001	1,65	2,80E-03		6-sep septin 6
204362_at	1,62	0,000738	1,51	0,02	SCAP2	src family associated phosphoprotein 2

218960_at	1,61	0,0017381	1,69	0,02	TMPRSS4	transmembrane protease, serine 4
208033_s_at	1,55	0,0014384	1,69	1,71E-03	ATBF1	AT-binding transcription factor 1
221090_s_at	1,54	0,0001552	1,33	0,02	FLJ10826	hypothetical protein FLJ10826
208743_s_at	1,53	0,0007387	1,29	5,32E-03	YWHAB	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase 4
202702_at	1,52	0,0002541	1,24	0,04	TRIM26	tripartite motif-containing 26
219628_at	1,52	0,0003456	1,45	0,01	WIG1	p53 target zinc finger protein
212293_at	1,51	0,0004013	1,36	0,01	HIPK1	homeodomain interacting protein kinase 1
219865_at	1,51	0,0006032	1,52	0,02	HSPC157	HSPC157 protein
218324_s_at	1,49	0,0010441	1,60	0,01	SPATS2	spermatogenesis associated, serine-rich 2
219397_at	1,48	0,0008397	1,70	1,10E-03	FLJ13448	hypothetical protein FLJ13448
218562_s_at	1,48	0,0009339	1,75	7,76E-03	FLJ10747	hypothetical protein FLJ10747
215044_s_at	1,47	9,17E-05	1,38	9,31E-03	STAM2	signal transducing adaptor molecule (SH3 domain and ITAM)
221937_at	1,47	0,0014225	1,66	0,01	AP1GBP1	AP1 gamma subunit binding protein 1
37831_at	1,46	0,0007042	1,52	8,55E-03	SIPA1L3	signal-induced proliferation-associated 1 like 3
218810_at	1,45	0,0007758	1,43	0,02	FLJ23231	hypothetical protein FLJ23231
217893_s_at	1,42	0,0006359	1,58	5,63E-03	FLJ12666	hypothetical protein FLJ12666
212072_s_at	1,40	0,0010705	1,29	3,45E-03	CSNK2A1	casein kinase 2, alpha 1 polypeptide
218076_s_at	1,39	9,70E-06	1,41	3,96E-03	ARHGAP17	Rho GTPase activating protein 17
219784_at	0,84	0,0005537	0,77	7,49E-03	FBXO31	F-box protein 31
208359_s_at	0,82	0,002498	0,72	6,01E-04	KCNJ4	potassium inwardly-rectifying channel, subfamily J, member 4
205305_at	0,81	0,0014735	0,83	9,28E-03	FGL1	fibrinogen-like 1
213066_at	0,81	0,0008979	0,76	3,96E-04	RUSC2	RUN and SH3 domain containing 2
217205_at	0,80	0,0002861	0,85	0,03		
207690_at	0,79	0,0004309	0,81	6,00E-04	ALX3	aristaless-like homeobox 3
52741_at	0,78	5,61E-05	0,83	0,02	FLJ40452	hypothetical protein FLJ40452
217381_s_at	0,78	0,0005856	0,83	0,02		Full-length cDNA clone CSODJ012YJ02 of T cells (Jurkat cell line)
218372_at	0,77	9,10E-06	0,78	0,01	MED25	mediator subunit 25
211500_at	0,77	0,0003602	0,78	1,59E-03	MAPK11	mitogen-activated protein kinase 11
215367_at	0,77	0,0005108	0,86	5,17E-03	KIAA1614	KIAA1614 protein
202182_at	0,76	0,0010997	0,71	4,65E-03	GCN5L2	GCN5 general control of amino-acid synthesis 5-like 2 (yeast)
211028_s_at	0,75	0,0001049	0,71	1,23E-04	KHK	ketoheksokinase (fructokinase) /// ketoheksokinase (fructokinase)
221109_at	0,75	3,30E-06	0,81	3,65E-03		hypothetical protein DKFZp434C0923
202248_at	0,74	3,60E-06	0,84	0,04	E2F4	E2F transcription factor 4, p107/p130-binding
219364_at	0,74	0,0001176	0,75	9,86E-03	LGP2	likely ortholog of mouse D11lgp2
204706_at	0,72	0,0008932	0,65	1,68E-04	INPP5E	inositol polyphosphate-5-phosphatase, 72 kDa
203171_s_at	0,72	0,000137	0,85	0,01	KIAA0409	KIAA0409 protein
210373_at	0,71	1,08E-05	0,84	0,03	TUSC4	tumor suppressor candidate 4
206298_at	0,71	0,0001183	0,49	4,71E-04	ARHGAP22	Rho GTPase activating protein 22
206814_at	0,71	0,0001523	0,78	0,01	NGFB	nerve growth factor, beta polypeptide
209940_at	0,71	0,0002447	0,74	0,01	PARP3	poly (ADP-ribose) polymerase family, member 3
201470_at	0,70	0,0006444	0,72	3,50E-03	GSTO1	glutathione S-transferase omega 1
214674_at	0,69	9,73E-05	0,83	0,02	USP19	ubiquitin specific protease 19
52159_at	0,69	0,0001779	0,81	2,86E-03	HEMK1	HemK methyltransferase family member 1
212803_at	0,69	0,0001472	0,72	3,55E-03	NAB2	NGFI-A binding protein 2 (EGR1 binding protein 2)
202243_s_at	0,68	0,0013631	0,58	3,11E-04	PSMB4	proteasome (prosome, macropain) subunit, beta type, 4
206441_s_at	0,68	5,38E-05	0,71	0,01	COMMD4	COMMD domain containing 4
214338_at	0,67	8,05E-05	0,79	0,01	DNAJB12	DnaJ (Hsp40) homolog, subfamily B, member 12
221096_s_at	0,67	6,26E-05	0,59	6,29E-04	PRO1580	hypothetical protein PRO1580
205905_s_at	0,67	2,50E-06	0,76	0,04	MICA /// MICB	MHC class I polypeptide-related sequence A /// MHC class I
205086_s_at	0,66	0,0003889	0,63	3,10E-03	384D8-2	hypothetical protein 384D8_6
218664_at	0,65	0,0002926	0,70	0,02	CGI-63	nuclear receptor binding factor 1
203246_s_at	0,65	0,0016892	0,81	0,03	TUSC4	tumor suppressor candidate 4
218590_at	0,64	0,0009672	0,58	8,34E-04	PEO1	progressive external ophthalmoplegia 1
213256_at	0,64	0,0020944	0,33	3,60E-04	MARCH-III	hypothetical protein MGC48332
204121_at	0,64	0,0010808	0,81	9,86E-03	GADD45G	growth arrest and DNA-damage-inducible, gamma
213468_at	0,64	1,60E-05	0,62	6,39E-03	ERCC2	excision repair cross-complementing rodent repair deficient
205224_at	0,64	0,0004458	0,77	0,02	SURF2	surfeit 2
209718_at	0,63	1,30E-06	0,72	2,25E-03	384D8-2	hypothetical protein 384D8_6
218270_at	0,62	0,0012833	0,74	0,02	MRPL24	mitochondrial ribosomal protein L24
213053_at	0,61	1,03E-05	0,66	6,59E-03	KIAA0841	KIAA0841
218722_s_at	0,61	0,0001615	0,72	9,64E-03	FLJ12436	hypothetical protein FLJ12436
203488_at	0,60	2,36E-05	0,79	0,04	LPHN1	latrophilin 1
221649_s_at	0,60	6,00E-07	0,72	4,74E-03	PPAN	peter pan homolog (Drosophila)
203045_at	0,59	0,0004161	0,76	0,01	NINJ1	ninjurin 1
213520_at	0,57	2,80E-05	0,59	3,18E-03		
214919_s_at	0,57	0,0004196	0,70	0,01	EIF4EBP3 /// I	EIF4EBP3 /// I MASK-4E-BP3 alternate reading frame gene /// eukaryotic translation
203655_at	0,55	1,61E-05	0,71	0,03	XRCC1	X-ray repair complementing defective repair in Chinese hamster
202387_at	0,51	0,0003334	0,73	0,01	BAG1	BCL2-associated athanogene
204343_at	0,48	9,74E-05	0,63	0,01	ABCA3	ATP-binding cassette, sub-family A (ABC1), member 3
218862_at	0,48	0,0002035	0,45	7,63E-05	ASB13	ankyrin repeat and SOCS box-containing 13
204497_at	0,47	0,0004747	0,63	5,02E-03	ADCY9	adenylate cyclase 9
212551_at	0,45	3,41E-05	0,50	7,94E-03	CAP2	CAP, adenylate cyclase-associated protein, 2 (yeast)
209339_at	0,45	0,0009976	0,63	2,52E-03	SIAH2	seven in absentia homolog 2 (Drosophila)
203071_at	0,44	0,0001614	0,66	0,03	SEMA3B	sema domain, immunoglobulin domain (Ig), short basic domain
218094_s_at	0,38	4,00E-05	0,63	6,74E-03	C20orf35	chromosome 20 open reading frame 35
207414_s_at	0,34	0,0001331	0,45	3,53E-04	PCSK6	proprotein convertase subtilisin/kexin type 6

203439_s_at	0,24	1,64E-05	0,68	4,97E-03	STC2	stanniocalcin 2
203438_at	0,19	0,0002526	0,43	3,29E-04	STC2	stanniocalcin 2
206754_s_at	0,14	0,0002216	0,81	0,02	CYP2B6	cytochrome P450, family 2, subfamily B, polypeptide 6
202376_at	0,13	5,50E-06	0,27	7,03E-03	SERPINA3	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 ar
205440_s_at	0,13	0,0009498	0,51	0,01		neuropeptide Y receptor Y1