

## SUPPORTING DATA SET 1

>gnl|UG|Hs#S3322070 **Homo sapiens matrix metalloproteinase 7** (matrilysin, uterine) (MMP7), mRNA /cds=(48,851) /gb=NM\_002423 /gi=13027804 /ug=Hs.2256 /len=1127

>gnl|UG|Hs#S3219743 **Homo sapiens junctophilin 3** (JPH3), mRNA /cds=(255,2501) /gb=NM\_020655 /gi=21704282 /ug=Hs.4775 /len=3997

>gnl|UG|Hs#S3219164 **Homo sapiens esophageal cancer associated protein** (MGC16824), mRNA /cds=(416,2890) /gb=NM\_020314 /gi=18482378 /ug=Hs.5320 /len=3113

>gnl|UG|Hs#S4807780 **Homo sapiens hypothetical protein FLJ36666** (FLJ36666), mRNA /cds=(35,520) /gb=NM\_152482 /gi=22749014 /ug=Hs.7604 /len=2131

>gnl|UG|Hs#S2294470 **Homo sapiens GK001 protein** (GK001), mRNA /cds=(185,1636) /gb=NM\_020198 /gi=9910241 /ug=Hs.8207 /len=3294

>gnl|UG|Hs#S5978568 **Homo sapiens DKFZP434D1335 protein** (DKFZP434D1335), mRNA /cds=(78,1469) /gb=NM\_015578 /gi=24308092 /ug=Hs.8258 /len=3389

>gnl|UG|Hs#S3219521 **Homo sapiens RNA helicase-related protein** (RNAHP), mRNA /cds=(18,2147) /gb=NM\_007372 /gi=11321631 /ug=Hs.8765 /len=3347

>gnl|UG|Hs#S3219471 **Homo sapiens complement component 1, q subcomponent, beta polypeptide** (C1QB), mRNA /cds=(64,825) /gb=NM\_000491 /gi=11038661 /ug=Hs.8986 /len=1072

>gnl|UG|Hs#S5494967 **Homo sapiens sorting nexin 3** (SNX3), transcript variant 3, mRNA /cds=(326,667) /gb=NM\_152828 /gi=23111042 /ug=Hs.12102 /len=1559

>gnl|UG|Hs#S2140319 **Homo sapiens CGI-147 protein** (CGI-147), mRNA /cds=(128,667) /gb=NM\_016077 /gi=7706350 /ug=Hs.12677 /len=806

>gnl|UG|Hs#S1726209 **Homo sapiens actin, gamma 1** (ACTG1), mRNA /cds=(75,1202) /gb=NM\_001614 /gi=11038618 /ug=Hs.14376 /len=1919

>gnl|UG|Hs#S1731310 **Homo sapiens COP9 subunit 6** (MOV34 homolog, 34 kD) (COPS6), mRNA /cds=(105,998) /gb=NM\_006833 /gi=21359878 /ug=Hs.15591 /len=1148

>gnl|UG|Hs#S3619375 **Homo sapiens hypothetical protein MGC10820** (MGC10820), mRNA /cds=(457,681) /gb=NM\_032648 /gi=14249191 /ug=Hs.18449 /len=959

>gnl|UG|Hs#S2140049 **Homo sapiens comparative gene identification 58** (CGI58), mRNA /cds=(7,1056) /gb=NM\_016006 /gi=7705770 /ug=Hs.19385 /len=1328

>gnl|UG|Hs#S1727443 **Homo sapiens protein phosphatase 1, catalytic subunit, beta isoform** (PPP1CB), mRNA /cds=(259,1242) /gb=NM\_002709 /gi=4506004 /ug=Hs.21537 /len=3590

>gnl|UG|Hs#S2140072 **Homo sapiens HSPC038 protein** (LOC51123), mRNA /cds=(137,367) /gb=NM\_016096 /gi=7705816 /ug=Hs.23528 /len=719

>gnl|UG|Hs#S3547207 **Homo sapiens phospholipase A2, group XII** (PLA2G12), mRNA /cds=(256,825) /gb=NM\_030821 /gi=21361943 /ug=Hs.25206 /len=1708

>gnl|UG|Hs#S2293282 **Homo sapiens hypothetical protein FLJ10300** (FLJ10300), mRNA /cds=(1710,3359) /gb=NM\_018051 /gi=21361686 /ug=Hs.42233 /len=3785

>gnl|UG|Hs#S1728960 **Homo sapiens fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)** (FABP3), mRNA /cds=(46,447) /gb=NM\_004102 /gi=10938020 /ug=Hs.49881 /len=679

>gnl|UG|Hs#S1726296 **Homo sapiens argininosuccinate lyase** (ASL), mRNA /cds=(70,1464) /gb=NM\_000048 /gi=4502256 /ug=Hs.61258 /len=1502

>gnl|UG|Hs#S1727480 **Homo sapiens protein S (alpha) (PROS1)**, mRNA /cds=(147,2177) /gb=NM\_000313 /gi=4506116 /ug=Hs.64016 /len=3309

>gnl|UG|Hs#S3437945 **Homo sapiens mitogen-activated protein kinase kinase 2 (MAP2K2)**, mRNA /cds=(255,1457) /gb=NM\_030662 /gi=21614527 /ug=Hs.72241 /len=1759

>gnl|UG|Hs#S1728144 **Homo sapiens ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast) (UBE2N)**, mRNA /cds=(64,522) /gb=NM\_003348 /gi=4507792 /ug=Hs.75355 /len=1203

>gnl|UG|Hs#S1729349 **Homo sapiens ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (UCHL1)**, mRNA /cds=(75,746) /gb=NM\_004181 /gi=21361090 /ug=Hs.76118 /len=1119

>gnl|UG|Hs#S5518205 **Homo sapiens casein kinase 1, epsilon (CSNK1E)**, transcript variant 1, mRNA /cds=(233,1483) /gb=NM\_152221 /gi=23199990 /ug=Hs.79658 /len=1559

>gnl|UG|Hs#S2139834 **Homo sapiens hypothetical protein BM-009 (BM-009)**, mRNA /cds=(386,1048) /gb=NM\_016623 /gi=7705303 /ug=Hs.92918 /len=1919

>gnl|UG|Hs#S2139766 **Homo sapiens PTD015 protein (PTD015)**, mRNA /cds=(148,504) /gb=NM\_014040 /gi=7662642 /ug=Hs.95870 /len=620

>gnl|UG|Hs#S3603834 **Homo sapiens testis-specific protein TES101RP (MGC4766)**, mRNA /cds=(561,1364) /gb=NM\_031451 /gi=20127665 /ug=Hs.97978 /len=1605

>gnl|UG|Hs#S3219684 **Homo sapiens FK506 binding protein like (FKBPL)**, mRNA /cds=(265,1314) /gb=NM\_022110 /gi=17149852 /ug=Hs.99134 /len=1379

>gnl|UG|Hs#S1729115 **Homo sapiens metastasis associated 1 (MTA1)**, mRNA /cds=(8,2155) /gb=NM\_004689 /gi=14141149 /ug=Hs.101448 /len=2662

>gnl|UG|Hs#S1727214 **Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6, 17kDa (NDUFB6)**, mRNA /cds=(104,490) /gb=NM\_002493 /gi=20149518 /ug=Hs.109646 /len=733

>gnl|UG|Hs#S1824446 **Homo sapiens leucine zipper-EF-hand containing transmembrane protein 1 (LETM1)**, mRNA /cds=(298,2517) /gb=NM\_012318 /gi=6912481 /ug=Hs.120165 /len=3907

>gnl|UG|Hs#S4284762 **Homo sapiens hypothetical protein BC009518 (LOC90799)**, mRNA /cds=(59,2524) /gb=NM\_138363 /gi=19923898 /ug=Hs.135265 /len=2705

>gnl|UG|Hs#S1730282 **Homo sapiens nuclear factor (erythroid-derived 2)-like 2 (NFE2L2)**, mRNA /cds=(114,1931) /gb=NM\_006164 /gi=20149575 /ug=Hs.155396 /len=2439

>gnl|UG|Hs#S1727788 **Homo sapiens chemokine (C-X-C motif) ligand 6 (granulocyte chemotactic protein 2) (CXCL6)**, mRNA /cds=(64,408) /gb=NM\_002993 /gi=4506850 /ug=Hs.164021 /len=1547

>gnl|UG|Hs#S1731058 **Homo sapiens poliovirus receptor (PVR)**, mRNA /cds=(172,1425) /gb=NM\_006505 /gi=19923371 /ug=Hs.171844 /len=3215

>gnl|UG|Hs#S1729935 **Homo sapiens solute carrier family 1 (neutral amino acid transporter), member 5 (SLC1A5)**, mRNA /cds=(591,2216) /gb=NM\_005628 /gi=5032092 /ug=Hs.183556 /len=2856

>gnl|UG|Hs#S2138875 **Homo sapiens neutrophil cytosolic factor 4, 40kDa (NCF4)**, transcript variant 2, mRNA /cds=(157,1203) /gb=NM\_013416 /gi=7382492 /ug=Hs.196352 /len=1421

>gnl|UG|Hs#S2294124 **Homo sapiens MCM10 minichromosome maintenance deficient 10 (S. cerevisiae) (MCM10)**, mRNA /cds=(95,2662) /gb=NM\_018518 /gi=19923533 /ug=Hs.198363 /len=4619

>gnl|UG|Hs#S2138938 **Homo sapiens chromosome 22 open reading frame 2 (C22orf2)**, mRNA /cds=(115,495) /gb=NM\_015373 /gi=21359898 /ug=Hs.227637 /len=1194

>gnl|UG|Hs#S1731210 **Homo sapiens ribosomal protein L14** (RPL14), mRNA /cds=(38,688) /gb=NM\_003973 /gi=16753224 /ug=Hs.235422 /len=843

>gnl|UG|Hs#S1730207 **Homo sapiens ariadne homolog 2 (Drosophila)** (ARIH2), mRNA /cds=(145,1626) /gb=NM\_006321 /gi=5453556 /ug=Hs.241558 /len=3892

>gnl|UG|Hs#S1727552 **Homo sapiens prothymosin, alpha (gene sequence 28)** (PTMA), mRNA /cds=(182,514) /gb=NM\_002823 /gi=21359859 /ug=Hs.250655 /len=1233

>gnl|UG|Hs#S1728284 **Homo sapiens aldolase A, fructose-bisphosphate** (ALDOA), mRNA /cds=(168,1262) /gb=NM\_000034 /gi=4557304 /ug=Hs.273415 /len=1464

>gnl|UG|Hs#S2138743 **Homo sapiens prolactin regulatory element binding** (PREB), mRNA /cds=(132,1385) /gb=NM\_013388 /gi=7019502 /ug=Hs.279784 /len=2059

>gnl|UG|Hs#S2139666 **Homo sapiens KIAA1001 protein** (KIAA1001), mRNA /cds=(459,2036) /gb=NM\_014960 /gi=7662439 /ug=Hs.279937 /len=4304

>gnl|UG|Hs#S3438634 **Homo sapiens RNA processing factor 1** (RPF1), mRNA /cds=(8,1057) /gb=NM\_025065 /gi=18643386 /ug=Hs.287863 /len=1336

>gnl|UG|Hs#S1731856 **Homo sapiens karyopherin alpha 4** (importin alpha 3) (KPNA4), mRNA /cds=(307,1872) /gb=NM\_002268 /gi=27477125 /ug=Hs.288193 /len=3816

>gnl|UG|Hs#S3323004 **Homo sapiens, Similar to neuronal thread protein, clone MGC:2959 IMAGE:3139291**, mRNA, complete cds /cds=(77,262) /gb=BC002534 /gi=12803420 /ug=Hs.306992 /len=1101

>gnl|UG|Hs#S3601614 **Homo sapiens hypothetical protein FLJ23027** (FLJ23027), mRNA /cds=(171,1709) /gb=NM\_032233 /gi=14149937 /ug=Hs.321766 /len=2374

>gnl|UG|Hs#S3619352 **Homo sapiens hypothetical protein MGC10814** (MGC10814), mRNA /cds=(403,519) /gb=NM\_032671 /gi=14249237 /ug=Hs.326741 /len=954

>gnl|UG|Hs#S4631077 **Homo sapiens crystallin, zeta (quinone reductase)-like 1** (CRYZL1), transcript variant 3, mRNA /cds=(86,682) /gb=NM\_145858 /gi=22202615 /ug=Hs.330208 /len=2099

>gnl|UG|Hs#S3940065 **Homo sapiens hypothetical protein MGC19556** (MGC19556), mRNA /cds=(1506,1667) /gb=NM\_033551 /gi=15809039 /ug=Hs.334787 /len=2552

>gnl|UG|Hs#S1731044 **Homo sapiens activated RNA polymerase II transcription cofactor 4** (PC4), mRNA /cds=(57,440) /gb=NM\_006713 /gi=19923783 /ug=Hs.349506 /len=1336

>gnl|UG|Hs#S1727740 **Homo sapiens ribosomal protein S5** (RPS5), mRNA /cds=(54,668) /gb=NM\_001009 /gi=13904869 /ug=Hs.356019 /len=725

>gnl|UG|Hs#S1731106 **Homo sapiens tissue factor pathway inhibitor 2** (TFPI2), mRNA /cds=(57,764) /gb=NM\_006528 /gi=5730090 /ug=Hs.378774 /len=1142

>gnl|UG|Hs#S1726756 **Homo sapiens FK506 binding protein 1A, 12kDa** (FKBP1A), transcript variant 12B, mRNA /cds=(104,430) /gb=NM\_000801 /gi=17149837 /ug=Hs.380080 /len=1578

**LEGEND**

**HuR** Z scores for genes on HuR array  
**IgG1** Z scores for genes on IgG1 array  
**HuR-IgG1** Z differences between genes from HuR minus IgG1 immunoprecipitation

**GB acc#** Accession number  
**UGCluster** Unigene accession number

**Symbol** Gene symbol  
**Name** Gene name

HuR	IgG1	HuR-IgG1	GB acc#	UGCluster	Symbol	Name
9.69	1.28	8.41	BC000305	Hs.3280	LOC90799	hypothetical protein BC009518
8.99	1.47	7.52	BC005323	Hs.182421	ACTB	actin, beta
8.85	1.48	7.37	BC005026	Hs.423756	PPP1CB	protein phosphatase 1, catalytic subunit, beta isoform
8.41	1.12	7.29	BC015937	Hs.109643	ACTB	actin, beta
7.73	1.42	6.31	BC019088	Hs.154332	ACTB	actin, beta
7.24	3.02	4.22	BC012355	Hs.187958	ACTG1	actin, gamma 1
7.04	1.94	5.09	BC007442	Hs.14376	ACTG1	actin, gamma 1
6.93	0.64	6.29	BC018445	Hs.164285	ACTB	actin, beta
6.93	1.76	5.17	BC003550	Hs.107387	ACTG1	actin, gamma 1
6.81	0.12	6.69	BC005105	Hs.336927	ACTB	actin, beta
6.61	1.97	4.64	BC017490	Data not found	ACTG1	actin, gamma 1
6.46	2.26	4.21	BC001341	Hs.6278	ACTG1	actin, gamma 1
5.58	1.42	4.15	BC002953	Hs.279881	GNA-11	likely ortholog of mouse G protein couple-dreceptor alpha 11
5.44	0.35	5.09	BC017117	Hs.155924	NCF4	neutrophil cytosolic factor 4, 40kDa
5.42	2.10	3.32	BC016031	Hs.159161	PTMA	prothymosin, alpha (gene sequence 28)
5.41	2.43	2.98	BC002680	Hs.380230	PTMA	prothymosin, alpha (gene sequence 28)
5.29	4.30	0.98	BC008883	Hs.289072	PC4	activated RNA polymerase II transcription cofactor 4
5.27	0.12	5.14	BC012617	Hs.378774	ACTG2	actin, gamma 2, smooth muscle, enteric
5.16	0.51	4.65	BC008407	In multiple clusters	PCBP2	poly(rC) binding protein 2
5.08	-0.62	5.70	BC011454	Hs.92186	UBE2N	ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
4.99	-0.18	5.17	BC003365	Hs.75355	UBE2N	ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
4.88	4.50	0.38	BC010853	Hs.424551	MGC4766	hypothetical protein MGC4766 similar to testis specific protein TES101RP
4.81	2.31	2.50	BC011683	Hs.109752	DKFZP434D1335	DKFZP434D1335 protein
4.69	-0.32	5.02	BC011642	Hs.137568	MTA1	metastasis associated 1
4.67	3.69	0.98	BC007066	Hs.11810	RPS5	ribosomal protein S5
4.62	1.04	3.57	BC017480	Hs.91684	ACTG1	actin, gamma 1
4.55	0.40	4.15	BC009308	Hs.9003	KIAA1001	KIAA1001 protein
4.43	4.11	0.32	BC009631	Hs.403436	MMP7	matrix metalloproteinase 7 (matrilysin, uterine)
4.42	2.62	1.80	BC012986	Hs.153863	ACTG1	actin, gamma 1
4.37	-0.77	5.14	NM_030662	Hs.72241	MAP2K2	mitogen-activated protein kinase kinase 2
4.35	1.11	3.24	BC017317	Hs.239718	ACTG1	actin, gamma 1
4.34	2.65	1.69	BC003500	Hs.18851	LOC51123	HSPC038 protein
4.30	2.87	1.43	BC002717	Hs.347963	RNAHP	RNA helicase-related protein
4.24	0.34	3.90	BC005223	Hs.99875	CD9	CD9 antigen (p24)
4.17	3.15	1.02	BC000451	Hs.30035	PREB	prolactin regulatory element binding
4.15	2.70	1.45	BC008195	Hs.61258	ASL	argininosuccinate lyase
4.06	2.91	1.15	BC015354	Hs.5565	FABP3	fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)
4.05	2.55	1.50	BC015028	Hs.235935	MGC10814	hypothetical protein MGC10814
4.04	2.66	1.38	BC020590	Hs.108408	C1QB	complement component 1, q subcomponent, beta polypeptide
4.02	2.04	1.98	BC006807	Hs.12677	CGI-147	CGI-147 protein

3.99	2.64	1.35	BC000986	Hs.183556	SLC1A5	solute carrier family 1 (neutral amino acid transporter), member 5
3.97	3.07	0.90	BC013980	Hs.30736	SERF2	small EDRK-rich factor 2
3.94	2.66	1.28	BC000417	Hs.5212	CRYZL1	crystallin, zeta (quinone reductase)-like 1
3.93	2.46	1.48	BC004285	Hs.22222	CGI-58	CGI-58 protein
3.87	1.49	2.37	BC004926	Hs.429142	NA 4-14-03 32	
3.76	1.77	1.99	BC001039	Hs.6844	IFITM3	
3.74	3.04	0.69	BC002756	Hs.12040	PVR	poliovirus receptor
3.68	0.48	3.20	BC001147	Hs.241205	ERH	enhancer of rudimentary homolog (Drosophila)
3.66	0.27	3.39	BC022096	Hs.314295	MGC10820	hypothetical protein MGC10820
3.65	1.41	2.24	BC008216	Data not found	FLJ36666	hypothetical protein FLJ36666
3.53	2.87	0.66	BC009960	Hs.285115	FTH1	ferritin, heavy polypeptide 1
3.50	2.60	0.90	BC005261	Hs.334629	NA 4-14-03 104	
3.46	2.18	1.28	BC008651	Data not found	CXCL6	chemokine (C-X-C motif) ligand 6 (granulocyte chemotactic protein 2)
3.45	2.49	0.96	BC006973	Hs.285847		
3.45	0.52	2.93	BC007489	Hs.12971	CSNK1E	casein kinase 1, epsilon
3.43	0.02	3.41	BC012745	Hs.60103	FLJ10300	hypothetical protein FLJ10300
3.41	0.64	2.78	BC012109	Hs.93564	FKBP1A	FK506 binding protein 1A, 12kDa
3.40	2.05	1.35	BC002707	Hs.288126	ASC	apoptosis-associated speck-like protein containing a CARD
3.38	1.80	1.58	BC002813	Hs.139226	PROS1	protein S (alpha)
3.37	2.84	0.53	BC004348	Hs.210546	COPS6	COP9 subunit 6 (MOV34 homolog, 34 kD)
3.36	3.27	0.09	BC007921	Hs.192861	MGC1223	hypothetical protein MGC1223
3.34	2.51	0.82	BC012489	Hs.111286	FLJ23027	hypothetical protein FLJ23027
3.29	0.16	3.13	BC000675	Hs.6449	BTF3	basic transcription factor 3
3.28	1.21	2.07	BC012938	Hs.52515	JPH3	junctophilin 3
3.28	2.34	0.94	BC008942	Hs.6904	NA 4-14-03 518	
3.27	1.75	1.52	BC011672	Hs.129739	ZFP26	C3HC4-like zinc finger protein
3.26	2.18	1.08	BC002849	Hs.181271	MGC16824	esophageal cancer associated protein
3.24	2.02	1.22	BC003516	Hs.325860	NA 4-14-03 41	
3.22	0.84	2.39	BC008297	Hs.98593	MGC19556	hypothetical protein MGC19556
3.22	0.19	3.03	BC002702	Hs.78457	PFN1	profilin 1
3.22	0.31	2.90	BC016582	Hs.34956	MRPS5	mitochondrial ribosomal protein S5
3.21	1.73	1.48	BC002362	Hs.234489	P17.3	neuronal protein 17.3
3.20	-0.98	4.19	BC007304	Hs.231851	FLJ11730	hypothetical protein FLJ11730
3.16	1.39	1.77	BC018999	Hs.42945	MGC4663	hypothetical protein MGC4663
3.15	0.43	2.73	BC003644	Hs.95577	KIAA0801	RNA helicase
3.15	1.97	1.18	BC012330	Hs.124840	U2AF1	U2(RNU2) small nuclear RNA auxillary factor 1
3.14	-0.76	3.90	BC016754	Hs.288193	KPNA4	karyopherin alpha 4 (importin alpha 3)
3.12	2.17	0.94	BC007067	Hs.24427	MT1L	metallothionein 1L
3.11	2.19	0.92	BC003159	Hs.79402	NA 4-14-03 119	
3.10	2.34	0.76	BC010019	Hs.301756	SNX3	sorting nexin 3
3.07	2.11	0.96	BC021236	Hs.433419	ALDOA	aldolase A, fructose-bisphosphate
3.03	1.39	1.64	BC004312	Hs.433326	C22orf2	chromosome 22 open reading frame 2
3.01	1.91	1.10	BC006106	Hs.103081	NA 4-14-03 429	
2.99	0.73	2.26	BC008870	Hs.32352	SPF30	splicing factor 30, survival of motor neuron-related
2.98	1.42	1.57	BC001304	Hs.12376	PCLO	piccino (presynaptic cytomatrix protein)
2.96	0.03	2.92	BC012611	Hs.79306	EIF4E	eukaryotic translation initiation factor 4E
2.90	0.17	2.73	BC005343	Hs.7086	PTD015	PTD015 protein
2.89	1.50	1.39	BC003682	Hs.146409	CDC42	cell division cycle 42 (GTP binding protein, 25kDa)
2.84	0.68	2.16	BC016280	Hs.5420	GTF2H4	general transcription factor IIH, polypeptide 4, 52kDa
2.83	0.31	2.52	BC011892	Hs.27633	TMSB4X	thymosin, beta 4, X chromosome
2.83	1.62	1.21	BC005097	Hs.77257	NA 4-14-03 431	
2.83	1.89	0.94	BC011806	Hs.29724	ALOX5AP	arachidonate 5-lipoxygenase-activating protein
2.81	2.53	0.28	BC012082	Hs.8358	CPSF6	cleavage and polyadenylation specific factor 6, 68kDa

2.80	1.47	1.33	BC002683	Hs.103291	GOSR2	golgi SNAP receptor complex member 2
2.80	1.64	1.16	BC010432	Data not found	RPF1	RNA processing factor 1
2.79	1.24	1.55	BC009801	Hs.109646	NDUFB6	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6, 17kDa
2.78	1.22	1.57	BC009362	Hs.440942	NA 4-14-03 244	
2.73	1.71	1.02	BC011814	Hs.273357	NUFIP1	nuclear fragile X mental retardation protein interacting protein 1
2.73	0.33	2.39	BC011382	Hs.5056	NCL	nucleolin
2.72	2.15	0.57	BC016300	Hs.81256	S100A4	S100 calcium binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog)
2.71	1.94	0.77	BC011363	Hs.173422	FUS	fusion, derived from t(12;16) malignant liposarcoma
2.70	0.84	1.86	BC010007	Hs.3682	NA 4-14-03 245	
2.69	1.86	0.83	BC010044	Hs.82906	LOC115761	hypothetical protein MGC17429
2.69	0.96	1.73	BC003688	Hs.367675	KIAA0632	KIAA0632 protein
2.68	1.90	0.78	BC019359	Hs.251653	TUBB2	tubulin, beta, 2
2.65	1.82	0.84	BC002928	Hs.43299	MGC2594	hypothetical protein MGC2594
2.64	1.72	0.92	Data not found		LOC115761	hypothetical protein MGC17429
2.63	0.32	2.31	BC010704	Hs.15744	PTDSS1	phosphatidylserine synthase 1
2.63	1.35	1.28	BC015330	Hs.187660	CCL5	chemokine (C-C motif) ligand 5
2.62	2.50	0.12	BC008586	Hs.50848		
2.59	-0.21	2.80	BC013142	Hs.1722	MGC4737	KIAA1882 protein
2.59	0.68	1.91	BC005197	Hs.16755	DKFZP586K0717	FIP1 like 1 (S. cerevisiae)
2.58	1.93	0.65	BC000751	Hs.119140	EIF5A	eukaryotic translation initiation factor 5A
2.57	2.23	0.34	BC008880	Hs.177677	NARF	nuclear prelamin A recognition factor
2.56	0.23	2.33	BC002534	Data not found	NA 4-14-03 78	
2.55	2.14	0.42	BC006311	Hs.421907		Homo sapiens cDNA: FLJ21869 fis, clone HEP02442.
2.54	2.46	0.08	BF695725	Hs.381171		ESTs, Highly similar to YZA1_HUMAN HYPOTHETICAL PROTEIN [H.sapiens]
2.54	2.47	0.08	BC002994	Hs.376253	PCTP	phosphatidylcholine transfer protein
2.54	-0.29	2.83	BC006334	Hs.183255	MGC7036	hypothetical protein MGC7036
2.54	1.06	1.48	BC010124	Hs.325422	TCEB1	transcription elongation factor B (SIII), polypeptide 1 (15kDa, elongin C)
2.54	2.12	0.42	BC004258	Hs.70333	K-ALPHA-1	tubulin, alpha, ubiquitous
2.53	1.67	0.86	BC014547	Hs.118836	MB	myoglobin
2.51	1.34	1.17	BC011369	Hs.27445	GBA	glucosidase, beta; acid (includes glucosylceramidase)
2.50	2.28	0.23	BC006364	Hs.12570	TUBB2	tubulin, beta, 2
2.49	-0.62	3.11	BC005051	Hs.101514	CTPS	CTP synthase
2.49	0.61	1.88	BC009267	Hs.76084	MGC17528	
2.49	0.68	1.81	BC004287	Hs.141693	LOC57146	hypothetical protein from clone 24796
2.48	1.21	1.27	BC005269	Hs.85050	GK001	GK001 protein
2.46	0.38	2.07	BC005871	Hs.334437	MGC4248	hypothetical protein MGC4248
2.45	0.20	2.25	BC010571	Hs.6194	CORO1C	coronin, actin binding protein, 1C
2.43	1.59	0.84	BC005858	Hs.287820	FN1	fibronectin 1
2.41	1.46	0.95	BC002356	Hs.172609	LOC115557	RAC/CDC42 exchange factor
2.41	0.87	1.54	BC001792	Hs.288600	KIAA0101	KIAA0101 gene product
2.41	1.37	1.04	BC007984	Hs.195370	TSSC1	tumor suppressing subtransferable candidate 1
2.40	-0.14	2.54	BC012026	Data not found	NA 4-14-03 200	
2.40	0.37	2.03	BC005949	Hs.333512	ALDH1B1	aldehyde dehydrogenase 1 family, member B1
2.36	0.49	1.87	BC005339	Hs.381154	FLJ38663	hypothetical protein FLJ38663
2.36	1.72	0.64	BC004267	Hs.412173	SSB1	SPRY domain-containing SOCS box protein SSB-1
2.36	1.51	0.85	BC004255	Hs.288539	MSCP	mitochondrial solute carrier protein
2.35	1.51	0.84	BC011770	Hs.75087	KIAA1972	KIAA1972 protein
2.35	0.52	1.83	BC001009	Hs.311609		
2.34	1.10	1.24	BC012732	Hs.146139	NA 4-14-03 38	
2.32	0.63	1.69	BC000141	Hs.79070	MYC	v-myc myelocytomatosis viral oncogene homolog (avian)
2.32	0.44	1.88	BC014898	Hs.3376	NA 4-14-03 312	
2.32	1.41	0.91	BC001809	Hs.279450	MGC3771	hypothetical protein MGC3771
2.31	1.77	0.54	BC001963	Hs.286128	C20orf98	chromosome 20 open reading frame 98

2.30	0.48	1.83	BC005190	Hs.236642	NETO2	neuropilin (NRP) and tolloid (TLL)-like 2
2.30	1.59	0.72	BC004257	Hs.350321	KNSL6	kinesin family member 2C
2.30	0.64	1.66	BC010998	Hs.239069	KIAA0793	FERM, RhoGEF and pleckstrin domain protein 2
2.30	1.86	0.44	BC011680	Hs.67991	SSR4	signal sequence receptor, delta (translocon-associated protein delta)
2.30	0.49	1.81	BC001073	Hs.7987	MIZF	MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein
2.29	-0.58	2.86	BC020116	Hs.263216	CDCA4	cell division cycle associated 4
2.28	-0.38	2.66	BC001311	Hs.25237	LOC51337	mesenchymal stem cell protein DSCD75
2.28	0.15	2.14	BC006981	Hs.278635	CLIC4	chloride intracellular channel 4
2.28	1.18	1.10	BC000790	Hs.77631	GCSH	glycine cleavage system protein H (aminomethyl carrier)
2.27	0.20	2.06	BC006375	Hs.173135	DNASE2	deoxyribonuclease II, lysosomal
2.24	1.92	0.32	BC001253	Hs.80042	ALG6	dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase
2.24	0.11	2.14	BC009390	Hs.144063	UCHL3	ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase)
2.23	-0.99	3.22	BC004954	Hs.431392	RBBP9	retinoblastoma binding protein 9
2.23	0.54	1.69	BC001232	Hs.101359	C6orf32	chromosome 6 open reading frame 32
2.22	0.94	1.28	BC013591	Hs.102541	MGC4771	hypothetical protein MGC4771
2.22	0.78	1.44	BC014840	In multiple clusters	FLJ11029	hypothetical protein FLJ11029
2.22	2.00	0.21	BC011049	Hs.26670	HNRPA1	heterogeneous nuclear ribonucleoprotein A1
2.22	2.11	0.10	BC009290	Hs.389107	ENO3	enolase 3, (beta, muscle)
2.21	2.16	0.05	BC003413	Hs.69851	NOLA1	nucleolar protein family A, member 1 (H/ACA small nucleolar RNPs)
2.20	-0.69	2.89	BC002661	Hs.74070	PTPN9	protein tyrosine phosphatase, non-receptor type 9
2.20	0.89	1.30	BC008585	Hs.198253	FLJ20320	hypothetical protein FLJ20320
2.19	0.99	1.21	BC002369	Hs.433619	ZNF35	zinc finger protein 35 (clone HF.10)
2.17	0.79	1.38	BC011021	Hs.198269	NDUFA3	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3, 9kDa
2.16	0.74	1.42	BC009782	Hs.268115	DJ122O8.2	hypothetical protein DJ122O8.2
2.15	0.32	1.83	BC008899	Hs.5890	NA 4-14-03 223	
2.15	0.77	1.38	BC001894	Hs.91448	FLJ20436	hypothetical protein FLJ20436
2.14	1.24	0.90	BC010160	Hs.38738	SPEC1	small protein effector 1 of Cdc42
2.14	1.12	1.02	BC007389	Hs.389107	MAX	MAX protein
2.14	0.65	1.49	BC018663	Hs.112250	DFFA	DNA fragmentation factor, 45kDa, alpha polypeptide
2.14	0.40	1.74	BC006837	Hs.7179	RAD1	RAD1 homolog (S. pombe)
2.14	0.60	1.54	BC000307	Hs.42826	LOC144845	hypothetical protein BC008631
2.14	0.93	1.21	BC021969	Hs.59584	CSH2	chorionic somatomammotropin hormone 2
2.13	0.39	1.74	BC018788	Hs.301839	MRPL30	mitochondrial ribosomal protein L30
2.11	0.60	1.51	BC007198	Data not found	KIAA0663	KIAA0663 gene product
2.11	1.12	0.99	BC001318	Hs.422336	RGS19	regulator of G-protein signalling 19
2.10	1.51	0.60	BC008254	Data not found	NA 4-14-03 255	
2.10	1.02	1.08	BC007647	Hs.119177	KIAA1921	KIAA1921 protein
2.09	0.06	2.03	BC007345	Hs.431392	DKFZP566I1024	DKFZP566I1024 protein
2.08	0.46	1.62	BC007450	Hs.22937	KBRAS2	I-kappa-B-interacting Ras-like protein 2
2.08	-0.02	2.09	BC009010	Hs.350860	WTAP	Wilms' tumour 1-associating protein
2.08	0.96	1.12	BC016904	Hs.78019	PLEKHB2	pleckstrin homology domain containing, family B (evectins) member 2
2.07	1.17	0.91	BC002945	Hs.334802	DNAJB6	DnaJ (Hsp40) homolog, subfamily B, member 6
2.07	1.09	0.98	BC000405	Hs.173255	LRRFIP1	leucine rich repeat (in FLII) interacting protein 1
2.07	1.47	0.60	BC009200	Hs.83656	LOC163233	similar to Zinc finger protein 93 (Zinc finger protein HTF34)
2.06	0.86	1.20	BC000211	Hs.421608	LOC51759	hepatocellular carcinoma-associated antigen 59
2.06	-0.54	2.60	BC006462	Hs.81424	UBL1	ubiquitin-like 1 (sentrin)
2.06	0.65	1.41	BC014394	Hs.10700	C20orf106	
2.05	0.20	1.86	NM_173562	Hs.188757	MGC14254	hypothetical protein MGC14254
2.05	1.54	0.51	BC011900	Hs.118721	HIG2	hypoxia-inducible protein 2
2.05	0.56	1.49	BC000093	Hs.77602	PSMB10	proteasome (prosome, macropain) subunit, beta type, 10
2.05	0.74	1.31	BC005195	Hs.381047	MRPL24	mitochondrial ribosomal protein L24
2.05	1.51	0.54	BC010570	Hs.831	CALM1	calmodulin 1 (phosphorylase kinase, delta)
2.05	0.18	1.87	BC007768	Data not found	MGC3794	similar to CG9578 gene product

2.04	0.23	1.82	BC003354	Hs.425808	CALM2	calmodulin 2 (phosphorylase kinase, delta)
2.04	0.20	1.83	BC001161	Hs.155204	C1orf8	chromosome 1 open reading frame 8
2.04	-0.13	2.16	BC009970	Hs.89643	TXNRD1	thioredoxin reductase 1
2.03	0.05	1.99	BC007455	Hs.136309	C6orf37	chromosome 6 open reading frame 37
2.03	0.65	1.38	BC004948	Data not found	LOC130589	hypothetical protein BC014916
2.03	0.19	1.85	BC006118	Hs.66774	MMP1	matrix metalloproteinase 1 (interstitial collagenase)
2.03	0.77	1.26	BC004420	Hs.25413	C20orf52	chromosome 20 open reading frame 52
2.02	1.88	0.14	BC009259	Hs.23590	FLJ33069	hypothetical protein FLJ33069
2.01	1.21	0.80	BC000383	Hs.119	MGC18257	hypothetical protein MGC18257
2.00	1.28	0.72	BC013352	Hs.63609	RNPEPL1	arginyl aminopeptidase (aminopeptidase B)-like 1
2.00	0.56	1.44	BC017714	Hs.169853	LIG3	ligase III, DNA, ATP-dependent
2.00	0.38	1.62	BC000365	Hs.89578	GTF2H1	general transcription factor IIH, polypeptide 1, 62kDa
2.00	0.00	2.00	BC002505	Hs.105465	SNRPF	small nuclear ribonucleoprotein polypeptide F
1.99	0.94	1.06	BC008396	Hs.1327	FKBP1A	FK506 binding protein 1A, 12kDa
1.99	-1.12	3.11	BC008455	Hs.16187	LGMN	legumain
1.99	-0.72	2.71	BC002964	Hs.91747	SMS	spermine synthase
1.98	0.97	1.01	BC002826	Hs.8737	BIRC2	baculoviral IAP repeat-containing 2
1.98	0.87	1.11	BC002724	Hs.13794	PADI2	peptidyl arginine deiminase, type II
1.98	0.63	1.35	BC005245	Hs.46967	DPP7	
1.97	-0.25	2.22	BC008832	Hs.57301	SSR3	signal sequence receptor, gamma (translocon-associated protein gamma)
1.96	0.20	1.76	NM_006807	Hs.77254	SP192	hypothetical protein SP192
1.96	1.62	0.34	BC003126	Hs.247741	NDUFS7	NADH dehydrogenase (ubiquinone) Fe-S protein 7, 20kDa (NADH-coenzyme Q reductase)
1.95	1.15	0.80	BC000418	Hs.104925	IHPK1	inositol hexaphosphate kinase 1
1.95	1.46	0.49	BC003625	Hs.737	ETR101	immediate early protein
1.94	1.87	0.07	BC001809	Hs.279450	MGC3771	hypothetical protein MGC3771
1.93	1.12	0.81	BC007047	Hs.107	ZFP36L1	zinc finger protein 36, C3H type-like 1
1.93	1.42	0.51	BC014408	Hs.31431	NONO	
1.93	1.39	0.53	BC008445	Hs.75823	NA 4-14-03 281	
1.92	1.11	0.81	BC009523	Hs.172458	TRIP11	thyroid hormone receptor interactor 11
1.91	-0.35	2.26	NM_004475	Hs.184488	ERP70	protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)
1.90	1.58	0.32	BC008090	Hs.181243	CASP6	caspase 6, apoptosis-related cysteine protease
1.90	-0.25	2.15	BC001883	Hs.75337	FLJ10989	hypothetical protein FLJ10989
1.90	0.42	1.47	BC000610	Hs.273333	NCL	nucleolin
1.89	1.67	0.22	BC003536	Hs.136912	LYAR	hypothetical protein FLJ20425
1.89	1.04	0.85	BC000460	Hs.159425	HSAJ1454	sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 3
1.88	0.27	1.61	BC008244	Hs.82829	TGFB3	transforming growth factor, beta 3
1.87	1.46	0.41	BC006772	Hs.165590	RPS13	ribosomal protein S13
1.86	0.49	1.37	BC000814	Hs.90077	TGIF	TGFB-induced factor (TALE family homeobox)
1.86	1.28	0.58	BC015893	Hs.154230	MGC10871	hypothetical protein similar to RNA-binding protein lark
1.86	-0.05	1.91	BC000988	Hs.306995	MGC5457	hypothetical protein MGC5457
1.86	0.48	1.38	BC001157	Hs.56294	FBL	fibrillarin
1.86	1.67	0.19	BC004979	Hs.73722	LOC65243	hypothetical protein LOC65243
1.86	0.44	1.42	BC000087	Hs.406511	TIP47	cargo selection protein (mannose 6 phosphate receptor binding protein)
1.86	1.84	0.02	BC005216	Hs.425790	ETS2	v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)
1.85	1.71	0.14	BC000583	Hs.78769	RPS17	ribosomal protein S17
1.85	-0.30	2.14	BC020090	Hs.52788	MGC19556	hypothetical protein MGC19556
1.84	0.68	1.16	BC012828	Hs.343824	NM23-H6	non-metastatic cells 6, protein expressed in (nucleoside-diphosphate kinase)
1.84	1.68	0.15	BC018117	Hs.2384	TSNAX	translin-associated factor X
1.83	1.35	0.48	BC011826	Hs.239720	FLJ13614	hypothetical protein FLJ13614
1.83	0.99	0.83	BC011535	Hs.26358	COX15	COX15 homolog, cytochrome c oxidase assembly protein (yeast)
1.82	1.11	0.72	BC008960	Hs.271167	PIPOX	pipecolic acid oxidase
1.82	0.74	1.08	BC011787	Hs.30868	AP2M1	adaptor-related protein complex 2, mu 1 subunit
1.81	0.50	1.31	BC009617	Hs.98855	BCL7A	B-cell CLL/lymphoma 7A



1.81	1.52	0.28	BC006111	Hs.120439	OPTN	optineurin
1.81	0.79	1.02	BC000820	Hs.433410	MNAT1	menage a trois 1 (CAK assembly factor)
1.80	0.27	1.53	BC000171	Hs.262476	AMD1	S-adenosylmethionine decarboxylase 1
1.80	-0.29	2.09	BC016663	Hs.173694	LOC220070	hypothetical protein BC004224
1.80	-0.15	1.95	BC001284	Hs.284741	MGC5149	hypothetical protein MGC5149
1.80	-0.17	1.97	BC007612	Hs.406455	TMSB4X	thymosin, beta 4, X chromosome
1.80	1.22	0.58	BC004349	Hs.176657	RPS9	ribosomal protein S9
1.80	-0.27	2.07	BC006007	Hs.93235	MGC3169	hypothetical protein MGC3169
1.80	0.95	0.85	BC006099	Hs.321231	CCBP2	chemokine binding protein 2
1.80	1.79	0.01	BC009234	Hs.352392	FLJ21156	hypothetical protein FLJ21156
1.78	1.46	0.31	BC004898	Hs.379815	AKR1C1	aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase)
1.78	1.06	0.71	BC012140	Hs.293981	C1orf33	chromosome 1 open reading frame 33
1.77	0.49	1.28	BC000162	Hs.3727	UNRIP	unr-interacting protein
1.77	0.24	1.54	BC003395	Hs.118797	UBE2D3	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)
1.77	0.98	0.80	BC007085	Hs.90458	TUBB	tubulin, beta polypeptide
1.76	0.54	1.22	BC012356	Hs.90011	SIGIRR	single Ig IL-1R-related molecule
1.76	-0.06	1.82	BC006503	Hs.3123	LLGL2	lethal giant larvae homolog 2 (Drosophila)
1.76	1.07	0.69	BC005314	Data not found	LBH	likely ortholog of mouse limb-bud and heart gene
1.76	0.59	1.17	BC000798	Hs.315164	MGC3038	hypothetical protein similar to actin related protein 2/3 complex, subunit 5
1.76	0.61	1.15	BC001079	Hs.240170	MGC2731	hypothetical protein MGC2731
1.75	0.75	1.01	BC000322	Hs.16244	NA 4-14-03 202	
1.75	1.38	0.37	BC016765	Hs.424312	MBLR	MBLR protein
1.75	0.52	1.23	BC000385	Hs.177556	RGR	retinal G protein coupled receptor
1.75	0.32	1.43	BC009182	Hs.285673	LOC51185	protein x 0001
1.75	0.05	1.70	BC015310	Hs.341806	RetSDR2	retinal short-chain dehydrogenase/reductase 2
1.75	-0.23	1.98	BC010123	Hs.109150	DBT	dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)
1.74	0.55	1.19	BC009404	Hs.77039	SLC31A1	solute carrier family 31 (copper transporters), member 1
1.74	-0.39	2.13	BC005976	Hs.77273	GDI2	GDP dissociation inhibitor 2
1.74	0.74	1.00	BC009830	Hs.288880	KPNA3	karyopherin alpha 3 (importin alpha 4)
1.73	0.77	0.97	BC002796	Hs.46446	TALDO1	transaldolase 1
1.73	0.58	1.15	BC017096	Hs.272018	FLJ20013	hypothetical protein FLJ20013
1.73	0.36	1.37	BC008636	Hs.406013	PXF	peroxisomal farnesylated protein
1.73	0.37	1.36	BC018164	Hs.155894	DAP3	death associated protein 3
1.73	1.06	0.66	BC005070	Hs.9015	MGC10485	hypothetical protein MGC10485
1.73	1.08	0.65	BC002669	Hs.239752	CST7	cystatin F (leukocystatin)
1.72	0.27	1.44	BC000918	Hs.288678	MGC5384	hypothetical protein MGC5384
1.71	0.33	1.38	BC019337	Hs.413826	IGHG3	immunoglobulin heavy constant gamma 3 (G3m marker)
1.71	-0.01	1.72	BC008180	Hs.288771	DKFZP586A0522	DKFZP586A0522 protein
1.70	0.52	1.18	BC009866	Hs.293981	CALM2	calmodulin 2 (phosphorylase kinase, delta)
1.70	-1.93	3.63	BC002733	Hs.323817	NA 4-14-03 94	
1.69	0.47	1.22	BC002359	Hs.75066	HZF12	zinc finger protein 12
1.69	-0.09	1.77	BC007457	Hs.36237	TEM7R	tumor endothelial marker 7-related precursor
1.69	0.12	1.56	BC012103	Hs.154729	MRPS25	mitochondrial ribosomal protein S25
1.68	1.62	0.07	BC019314	Hs.26518	PHGDH	phosphoglycerate dehydrogenase
1.68	0.21	1.47	BC008280	Hs.146688	PTGES	prostaglandin E synthase
1.68	0.58	1.10	BC002335	Hs.20597	HCLS1	hematopoietic cell-specific Lyn substrate 1
1.68	1.08	0.60	BC005123	Hs.59403	DKFZP434A0131	DKFZp434A0131 protein
1.67	0.10	1.58	BC004402	Hs.44592	LOC130589	hypothetical protein BC014916
1.67	0.21	1.46	BC010876	Hs.28355	JUN	v-jun sarcoma virus 17 oncogene homolog (avian)
1.67	-0.87	2.54	BC011594	Hs.125133	STOML2	stomatolipin-like 2
1.67	-1.19	2.86	BC009871	Data not found	MGC29761	hypothetical protein MGC29761

1.67	-0.11	1.78	BC000765	Hs.223394	MGC2742	hypothetical protein MGC2742
1.67	0.88	0.79	BC009361	Hs.82280	ESM1	endothelial cell-specific molecule 1
1.67	1.38	0.29	BC007240	Hs.153959	NPM1	nucleophosmin (nucleolar phosphoprotein B23, numatrin)
1.66	-1.24	2.90	BC010119	Hs.5662	MGC21675	hypothetical protein MGC21675
1.66	0.30	1.36	BC017562	Hs.241336	CDX2	caudal type homeo box transcription factor 2
1.65	1.17	0.49	BC004242	Hs.155218	MVK	mevalonate kinase (mevalonic aciduria)
1.65	-0.05	1.69	BC000614	Hs.446552	TUBB2	tubulin, beta, 2
1.65	0.15	1.49	BC001642	Hs.89605	STOML2	stomatin (EPB72)-like 2
1.65	0.53	1.12	BC005371	Hs.7036	KIAA1191	KIAA1191 protein
1.64	0.89	0.76	BC013188	Hs.421194	TM4SF1	transmembrane 4 superfamily member 1
1.64	-0.20	1.84	BC000144	Hs.184270	CAPZA1	capping protein (actin filament) muscle Z-line, alpha 1
1.64	0.36	1.28	BC012550	Hs.74082	KLRC1	killer cell lectin-like receptor subfamily C, member 1
1.63	1.35	0.28	BC009319	Hs.250624	DKFZP564M182	DKFZP564M182 protein
1.63	1.50	0.13	BC002803	Hs.11156	KIAA0020	minor histocompatibility antigen HA-8
1.63	-0.09	1.73	BC004819	Hs.317714	PLDN	pallidin homolog (mouse)
1.63	-1.03	2.66	BC006804	Hs.154718	TPD52L2	tumor protein D52-like 2
1.63	-1.69	3.32	BC010958	Hs.75586	CCND2	cyclin D2
1.63	0.17	1.45	BC008846	Data not found	ELA3B	elastase 3B, pancreatic
1.62	0.18	1.44	BC004881	Hs.25524	SMA3	SMA3
1.62	0.25	1.37	BC005366	Hs.81634	ARHF	ras homolog gene family, member F (in filopodia)
1.62	0.80	0.82	BC000623	Hs.311100	MGC2718	hypothetical protein MGC2718
1.62	0.53	1.08	Data not found		PSMD11	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11
1.61	-0.05	1.66	BC007011	Hs.97411	SLAHBP1	fuse-binding protein-interacting repressor
1.61	0.50	1.11	BC010652	Hs.91668	MAGEA3	melanoma antigen, family A, 3
1.60	-1.01	2.60	BC000416	Hs.150922	SPHK2	sphingosine kinase 2
1.60	0.91	0.69	BC008439	Hs.430150	USP10	ubiquitin specific protease 10
1.59	0.48	1.10	BC000744	Hs.84285	UBE2I	ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)
1.58	1.16	0.43	BC012354	Hs.356360	NA 4-14-03 2	
1.58	0.23	1.36	BC007374	Hs.48650	LOC142937	hypothetical protein BC008131
1.58	-0.74	2.33	BC005231	Data not found	BPGM	2,3-bisphosphoglycerate mutase
1.58	0.96	0.62	BC009790	Hs.1815	MYL3	myosin, light polypeptide 3, alkali; ventricular, skeletal, slow
1.58	0.49	1.09	BC014061	Hs.169138	DKFZP434H132	DKFZP434H132 protein
1.58	0.68	0.89	BC010377	Hs.50984	SAS	sarcoma amplified sequence
1.57	0.03	1.54	BC000744	Hs.84285	UBE2I	ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)
1.56	1.03	0.53	BC014217	Hs.8054	CHEK1	CHK1 checkpoint homolog (S. pombe)
1.56	0.72	0.84	In multiple clusters		MGC20481	hypothetical protein MGC20481
1.56	1.30	0.26	BC000390	Hs.203910	GDI1	GDP dissociation inhibitor 1
1.56	-0.29	1.85	BC008251	Hs.182476	RABIF	RAB interacting factor
1.56	1.14	0.42	BC001821	Hs.323949	KAI1	kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))
1.55	1.09	0.46	BC011583	Hs.159406	EIF3S2	eukaryotic translation initiation factor 3, subunit 2 beta, 36kDa
1.54	0.98	0.56	Data not found		dJ383J4.3	hypothetical gene supported by BC007071
1.54	0.57	0.97	BC003637	Hs.400353	NA 4-14-03 500	
1.54	-0.22	1.76	BC006559	Hs.356360	CHEK2	CHK2 checkpoint homolog (S. pombe)
1.54	0.02	1.51	BC002944	Hs.424099	MAGEA8	melanoma antigen, family A, 8
1.54	0.11	1.43	BC000747	Hs.211973	RRP4	homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease
1.53	0.34	1.20	BC021198	Hs.75244	CGI-128	CGI-128 protein
1.53	0.41	1.12	BC004964	Hs.168669	CALU	calumenin
1.53	0.67	0.86	BC005214	Hs.79381	SLC12A8	solute carrier family 12 (potassium/chloride transporters), member 8
1.53	-0.70	2.23	BC015177	Data not found	STOML2	stomatin (EPB72)-like 2
1.53	-0.40	1.93	BC010621	Hs.158084	NA 4-14-03 1	
1.53	0.22	1.31	BC007516	Hs.326422	MGC4549	hypothetical protein MGC4549
1.52	0.89	0.63	BC002396	Hs.284279	WBP2	WW domain binding protein 2

1.52	0.63	0.88	BC001348	Hs.245710	FLJ11305	hypothetical protein FLJ11305
1.52	0.48	1.03	BC011885	Hs.332404	TPM3	
1.51	0.82	0.69	BC015920	Hs.71642	KCNS3	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3
1.51	-0.05	1.56	BC000993	Hs.334319	LOC130026	hypothetical protein BC000993
1.51	-1.48	2.99	BC007035	Hs.84928	UPK1B	uroplakin 1B
1.51	1.27	0.24	BC011350	Hs.48938	NUP88	nucleoporin 88kDa
1.50	1.32	0.19	BC006439	Hs.284180	VMP1	likely ortholog of rat vacuole membrane protein 1
1.50	0.55	0.96	BC008858	Hs.283976	CCBL1	cysteine conjugate-beta lyase; cytoplasmic (glutamine transaminase K, kyneurenine aminotransferase)
1.50	-0.22	1.72	BC002890	Hs.184519	MRS2L	MRS2-like, magnesium homeostasis factor ( <i>S. cerevisiae</i> )
1.50	0.84	0.66	BC000748	Hs.159154	TUBB4	tubulin, beta, 4
1.49	0.57	0.92	BC000051	Hs.182859	HSPC152	hypothetical protein HSPC152
1.49	1.26	0.22	BC008644	Data not found	SEDL	
1.49	-0.19	1.68	BC007295	Hs.407372	ANXA2	annexin A2
1.49	0.44	1.05	BC017555	Hs.72879	C(27)-3BETA-HSD	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
1.49	0.67	0.82	BC001514	Hs.155584	ICAM4	intercellular adhesion molecule 4, Landsteiner-Wiener blood group
1.49	0.62	0.86	BC012139	Hs.29696	RPS25	ribosomal protein S25
1.48	-0.23	1.72	BC014107	Hs.27262	MGC4266	hypothetical protein MGC4266
1.48	-1.52	3.00	BC007827	Hs.334526		
1.48	1.04	0.43	BC008712	Hs.206501	NAG73	NPC-related protein NAG73
1.47	0.26	1.22	BC005248	Hs.155103	CFL1	cofilin 1 (non-muscle)
1.47	1.07	0.40	BC011556	Hs.65412	NUDC	nuclear distribution gene C homolog ( <i>A. nidulans</i> )
1.47	1.40	0.07	BC007863	Hs.6793		
1.47	0.95	0.51	BC014029	Hs.155287	U2AF1	U2(RNU2) small nuclear RNA auxillary factor 1
1.46	0.23	1.23	BC006133	Hs.65238	EPHX2	epoxide hydrolase 2, cytoplasmic
1.46	0.89	0.57	BC008420	Hs.85119	TUBD1	likely ortholog of mouse tubulin, delta 1
1.46	1.26	0.20	BC006784	Hs.431584	RPS14	ribosomal protein S14
1.46	0.65	0.81	BC001131	Hs.352109	H2BFA	histone 1, H2bg
1.45	-0.57	2.01	BC005023	Hs.9825	BCL2L2	BCL2-like 2
1.44	0.61	0.82	BC000901	Hs.119475	CIRBP	cold inducible RNA binding protein
1.44	0.20	1.24	BC008678	Hs.126256	C21orf97	chromosome 21 open reading frame 97
1.43	0.26	1.18	BC007212	Hs.99256	NPM1	nucleophosmin (nucleolar phosphoprotein B23, numatrin)
1.43	1.22	0.21	BC001674	Hs.27556	FLJ22405	hypothetical protein FLJ22405
1.43	-0.63	2.06	BC016979	Hs.140950	COX7A2L	cytochrome c oxidase subunit VIIa polypeptide 2 like
1.43	-0.52	1.95	BC012469	Data not found	BSG	basigin (OK blood group)
1.43	0.36	1.07	BC008876	Hs.1408	RTN3	reticulon 3
1.43	0.57	0.86	BC007680	Hs.278507	HARSL	histidyl-tRNA synthetase-like
1.43	0.92	0.51	BC013376	Hs.326733	C1orf8	chromosome 1 open reading frame 8
1.43	-0.05	1.48	BC012131	Hs.29706	LAPTM4A	lysosomal-associated protein transmembrane 4 alpha
1.43	1.27	0.16	BC012357	Hs.53447	NDRG1	N-myc downstream regulated gene 1
1.43	0.25	1.18	BC006254	Hs.48348	MANBAL	mannosidase, beta A, lysosomal-like
1.42	0.21	1.21	BC001259	Hs.50628	AP4S1	adaptor-related protein complex 4, sigma 1 subunit
1.42	0.81	0.61	BC021988	Hs.30340	DPP3	dipeptidylpeptidase 3
1.42	1.18	0.24	BC003555	Hs.134200	RPL9	ribosomal protein L9
1.42	0.76	0.66	BC013953	Hs.424931	TPM3	
1.41	1.09	0.32	BC015961	Hs.394	SATB1	special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)
1.41	0.35	1.07	BC000391	Hs.66196	CASP2	caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
1.41	0.70	0.71	BC004987	Hs.47584	GNB3	guanine nucleotide binding protein (G protein), beta polypeptide 3
1.41	0.53	0.88	BC003362	Hs.9030	TONDU	TONDU
1.41	-0.61	2.02	BC009869	Hs.380718	LOC91056	hypothetical protein BC004895
1.41	1.36	0.05	BC014128	Hs.334684		
1.41	0.28	1.13	BC017469	Hs.71040	NA 4-14-03 110	

1.41	1.09	0.31	BC004162	Hs.275711	MGC2452	hypothetical protein MGC2452
1.40	1.13	0.27	BC008438	Hs.109276	PTP4A3	protein tyrosine phosphatase type IVA, member 3
1.40	-1.66	3.06	BC017219	Hs.288995	FLJ20813	hypothetical protein FLJ20813
1.40	0.25	1.15	BC007095	Hs.30888	MGC3234	hypothetical protein MGC3234
1.39	1.08	0.31	BC010080	Hs.296242	CENTB1	centaurin, beta 1
1.39	0.18	1.21	BC012912	Hs.82208	MYCBP	c-myc binding protein
1.39	1.38	0.01	BC012063	Hs.26550	NA 4-14-03 297	
1.39	-0.92	2.31	BC007917	Hs.334527	NA 4-14-03 106	
1.39	-0.05	1.44	BC002812	Hs.154510	KIAA1376	KIAA1376 protein
1.38	0.17	1.22	BC014555	Hs.334		Homo sapiens, clone IMAGE:4850403, mRNA
1.38	0.24	1.14	BC014948	Hs.297405	CYR61	cysteine-rich, angiogenic inducer, 61
1.38	0.09	1.29	BC009881	Hs.178617	STOML2	stomatin (EPB72)-like 2
1.38	0.17	1.21	BC008306	Hs.355906	PBP	prostatic binding protein
1.38	0.69	0.69	BC007402	Hs.75736	MGC3113	hypothetical protein MGC3113
1.38	-0.30	1.68	BC001651	Hs.48855	FLJ10468	hypothetical protein FLJ10468
1.38	1.29	0.08	BC008975	Hs.7049	HSPC052	HSPC052 protein
1.37	0.90	0.48	BC009976	Hs.192285	MGC29762	hypothetical protein MGC29762
1.37	0.80	0.57	BC018652	Hs.3807	SAS	N-acetylneuraminic acid synthase (sialic acid synthase)
1.37	0.92	0.45	BC000043	Hs.183153	LOC56181	hypothetical protein RP1-317E23
1.37	0.95	0.42	BC012178	Hs.5398	IFIT4	interferon-induced protein with tetratricopeptide repeats 4
1.37	-0.90	2.27	BC012595	Hs.23103	BET1	BET1 homolog (S. cerevisiae)
1.37	0.69	0.67	BC013611	Hs.380728	RPS3A	ribosomal protein S3A
1.36	-0.45	1.81	NM_006029	Hs.194709	FLJ11198	hypothetical protein FLJ11198
1.36	0.23	1.13	BC004535	Hs.76662	LOC90141	chromosome 14 open reading frame 143
1.36	0.00	1.36	BC002420	Hs.22412	KIAA1160	KIAA1160 protein
1.35	0.73	0.62	BC001659	Hs.75104	RNPS1	RNA binding protein S1, serine-rich domain
1.35	1.16	0.19	BC000232	Hs.178112	PEN-2	presenilin enhancer 2
1.35	0.72	0.64	BC008664	Hs.2043	LOC113174	hypothetical protein BC012010
1.34	-0.82	2.16	BC010855	Hs.75835	LOC93408	hypothetical protein BC002778
1.34	0.90	0.44	BC017191	Hs.265327	EIF3S6IP	eukaryotic translation initiation factor 3, subunit 6 interacting protein
1.34	0.70	0.63	BC006825	Hs.301404	RBM3	RNA binding motif protein 3
1.33	0.36	0.98	BC007610	Hs.406423	PDCD10	programmed cell death 10
1.33	0.35	0.98	BC010660	Hs.273415	PHGDH	phosphoglycerate dehydrogenase
1.33	0.29	1.04	BC003604	Hs.7579	NFIC	nuclear factor I/C (CCAAT-binding transcription factor)
1.32	0.94	0.38	BC011777	Hs.184276	PPM1F	protein phosphatase 1F (PP2C domain containing)
1.32	0.52	0.81	BC011584	Hs.10487	LR8	LR8 protein
1.32	0.76	0.56	BC014058	Hs.13094	CD84	CD84 antigen (leukocyte antigen)
1.32	0.30	1.02	BC013630	Hs.301613	DKFZP761I2123	hypothetical protein DKFZp761I2123
1.31	-0.06	1.38	BC004957	Hs.54320	HSPC031	hypothetical protein HSPC031
1.31	1.28	0.02	BC001285	Hs.180433	HSRTSBETA	rTS beta protein
1.30	0.59	0.72	NM_032261	Hs.381214	UP	uridine phosphorylase
1.30	0.69	0.62	BC016977	Hs.294101	KIAA1305	KIAA1305 protein
1.30	-0.08	1.38	BC012352	Hs.133331	NINJ1	ninjurin 1
1.30	-1.39	2.70	BC006155	Hs.19121	GM2A	GM2 ganglioside activator protein
1.30	-0.28	1.58	BC020998	Hs.158244	PPARA	peroxisome proliferative activated receptor, alpha
1.30	0.62	0.68	BC002382	Hs.226581	DKFZP566K1924	DKFZP566K1924 protein
1.29	-0.06	1.36	BC010627	Data not found	CIRBP	cold inducible RNA binding protein
1.29	1.23	0.07	BC009461	Hs.231029		Homo sapiens, clone IMAGE:4861097, mRNA
1.29	-0.25	1.54	BC001319	Hs.347950	PIGS	phosphatidylinositol glycan, class S
1.29	1.00	0.29	BC006253	Hs.16390	C20orf97	chromosome 20 open reading frame 97
1.29	-0.23	1.52	BC003137	Hs.414990	DNAJA3	DnaJ (Hsp40) homolog, subfamily A, member 3
1.29	0.55	0.74	BC002833	Hs.87225	ABCE1	ATP-binding cassette, sub-family E (OABP), member 1
1.28	-0.58	1.86	BC012950	Hs.247323	CKB	creatine kinase, brain

1.28	0.50	0.78	BC000147	Hs.75342	ME2	malic enzyme 2, NAD(+)-dependent, mitochondrial
1.28	0.40	0.88	BC000319	Hs.15961	TSPAN-3	tetraspan 3
1.28	0.43	0.85	BC005071	Hs.332382	STK12	serine/threonine kinase 12
1.28	0.51	0.77	BC001052	Hs.235069	RECQL	RecQ protein-like (DNA helicase Q1-like)
1.28	1.11	0.17	BC004233	Hs.27935	SIAHBP1	fuse-binding protein-interacting repressor
1.28	-0.62	1.90	NM_000371	Hs.427202	MGC13007	hypothetical protein MGC13007
1.27	0.21	1.07	BC000732	Hs.152707	GBAS	glioblastoma amplified sequence
1.27	0.98	0.29	BC021239	Hs.153177	PTAFR	platelet-activating factor receptor
1.27	0.24	1.03	BC005246	Hs.84072	STX7	syntaxin 7
1.27	-0.04	1.31	BC013596	Hs.406397	G6PT1	glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1
1.27	1.16	0.11	BC005080	Hs.76095	SF3B4	splicing factor 3b, subunit 4, 49kDa
1.27	-0.54	1.81	BC002553	Hs.83155	ALDH3B1	aldehyde dehydrogenase 3 family, member B1
1.27	1.12	0.15	BC000155	Hs.268530	GPS1	G protein pathway suppressor 1
1.27	1.14	0.12	BC001823	Hs.6061	PDAP1	PDGFA associated protein 1
1.27	0.13	1.14	BC003394	Hs.418901	HNRPC	heterogeneous nuclear ribonucleoprotein C (C1/C2)
1.27	-0.30	1.57	BC003629	Data not found	RNU2	
1.27	0.46	0.81	BC007195	Hs.101695	NCK2	NCK adaptor protein 2
1.27	-0.96	2.22	BC001047	Hs.446459	DBR1	debranching enzyme homolog 1 (S. cerevisiae)
1.26	1.03	0.24	BC012809	Hs.5198	DSCR2	Down syndrome critical region gene 2
1.26	0.21	1.05	BC007063	Hs.180909	My015	chromosome 14 open reading frame 52
1.26	1.02	0.24	BC011936	In multiple clusters	CLN2	ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
1.26	0.70	0.56	BC004523	Hs.13225	NA 4-14-03 501	
1.26	-0.05	1.31	BC009331	Hs.337772	HSPC049	HSPC049 protein
1.25	-0.02	1.28	BC015601	Hs.73722	ALS2CR2	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2
1.25	0.27	0.98	BC021979	Hs.99858	MSN	moesin
1.25	0.51	0.75	BC012113	Hs.424053	TP53	tumor protein p53 (Li-Fraumeni syndrome)
1.25	0.64	0.61	BC011942	Data not found	B4GALT4	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4
1.25	-0.40	1.65	BC009825	Hs.101821	MYOZ2	myozenin 2
1.25	0.42	0.83	BC003360	Hs.100555	DDX18	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)
1.25	0.21	1.04	BC002730	Hs.17144	SPIN	spindlin
1.25	0.63	0.62	BC001952	Hs.7845	MGC2752	hypothetical protein MGC2752
1.24	-0.18	1.42	BC005118	Hs.150580	PCBP3	poly(rC) binding protein 3
1.24	-0.39	1.63	BC012049	Hs.154654	FAF1	Fas (TNFRSF6) associated factor 1
1.24	0.87	0.37	BC007320	Hs.188401	MGC2603	hypothetical protein MGC2603
1.24	-0.10	1.34	BC001037	Hs.288544	MGC10977	tripartite motif-containing 51
1.24	1.09	0.15	BC011349	Hs.1544	MIF	macrophage migration inhibitory factor (glycosylation-inhibiting factor)
1.24	0.57	0.67	BC012167	Hs.284286	FLJ10569	hypothetical protein FLJ10569
1.24	0.71	0.53	BC007560	Hs.334851	SRP9	signal recognition particle 9kDa
1.24	0.26	0.98	BC000084	Hs.22451	FLJ20399	hypothetical protein FLJ20399
1.24	0.15	1.09	BC002805	Hs.333303	COG8	component of oligomeric golgi complex 8
1.24	0.96	0.28	BC000453	Hs.75737	FLJ14775	hypothetical protein FLJ14775
1.24	0.43	0.81	BC005073	Hs.332040	LOC55971	insulin receptor tyrosine kinase substrate
1.24	-0.35	1.59	BC008877	Hs.271815	FLJ12975	hypothetical protein FLJ12975
1.23	0.81	0.42	BC012120	Hs.433504	FLJ10402	importin 9
1.23	0.21	1.02	BC001383	Hs.181874	IFIT4	interferon-induced protein with tetratricopeptide repeats 4
1.23	0.28	0.95	BC002914	Hs.379203	RRP40	exosome component Rrp40
1.22	1.02	0.21	BC009846	Hs.3618	LAMP1	lysosomal-associated membrane protein 1
1.22	-0.82	2.04	BC007242	Hs.77711	LOC116254	hypothetical protein BC014320
1.22	-0.36	1.58	BC015725	Hs.172702	MUF1	MUF1 protein
1.22	-0.84	2.07	BC006553	Hs.70704	LOC54499	putative membrane protein
1.22	-1.03	2.25	BC006098	Hs.421667	ETF1	eukaryotic translation termination factor 1
1.22	-0.27	1.49	BC011839	Hs.334	LSP1	lymphocyte-specific protein 1
1.22	0.64	0.58	BC010903	Hs.14347	FLJ10460	hypothetical protein FLJ10460

1.22	-2.89	4.11	BC006515	Hs.159406	COX6B	cytochrome c oxidase subunit VIb
1.22	-1.15	2.37	BC005004	Hs.86211	HTATSF1	HIV TAT specific factor 1
1.22	0.13	1.09	BC015617	Hs.13526	HMGN2	high-mobility group nucleosomal binding domain 2
1.22	-0.27	1.49	BC006535	Hs.86859	MRP63	mitochondrial ribosomal protein 63
1.21	0.19	1.02	BC000077	Hs.178551	SCMH1	sex comb on midleg homolog 1 (Drosophila)
1.21	0.08	1.13	BC015934	Hs.75207	CGI-135	CGI-135 protein
1.21	0.65	0.56	BC005963	Hs.417816	H2AFL	histone 1, H2ac
1.21	0.02	1.19	BC010947	Hs.2943	SRP19	signal recognition particle 19kDa
1.21	-0.71	1.92	Data not found		MGC4809	serologically defined breast cancer antigen NY-BR-20
1.21	0.71	0.50	BC000910	Hs.36587	PPP1R7	protein phosphatase 1, regulatory subunit 7
1.21	0.06	1.15	BC003400	Hs.60679	TAF9	TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 32kDa
1.21	0.25	0.96	BC000669	Hs.433337	HM13	histocompatibility (minor) 13
1.21	-0.70	1.91	BC001282	Hs.236774	HMGN4	high mobility group nucleosomal binding domain 4
1.21	-0.02	1.23	BC016763	Hs.278255	C20orf100	chromosome 20 open reading frame 100
1.20	-0.49	1.70	BC013999	Hs.158006	NA 4-14-03 252	
1.20	-0.70	1.90	BC010550	Hs.3132	LOC51219	clone FLB5214
1.20	0.40	0.80	BC005289	Hs.42644	SNX12	sorting nexin 12
1.20	0.62	0.58	BC003533	Hs.73452	NA 4-14-03 216	
1.20	0.14	1.06	BC005156	Hs.38044	FLJ23277	FLJ23277 protein
1.20	0.56	0.64	BC002771	Hs.13880	AKR7A2	aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)
1.20	-0.24	1.44	BC008914	Hs.407995	TOLLIP	Toll-interacting protein
1.20	0.31	0.89	BC002842	Hs.180779	KIAA0252	KIAA0252 protein
1.20	0.32	0.87	BC000154	Hs.119178	CIP1	cation-chloride cotransporter-interacting protein
1.19	0.46	0.73	BC011714	Hs.372673	SLC23A1	solute carrier family 23 (nucleobase transporters), member 1
1.18	0.53	0.65	BC022191	Hs.166019	DFFA	DNA fragmentation factor, 45kDa, alpha polypeptide
1.18	0.33	0.85	BC007951	Hs.2316	CALM2	calmodulin 2 (phosphorylase kinase, delta)
1.18	0.98	0.20	BC000116	Hs.75824	KDELR2	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2
1.18	0.26	0.92	BC013748	Hs.279529	DKFZP564K1964	DKFZP564K1964 protein
1.18	0.28	0.90	BC010035	Hs.3712	TFPI	tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)
1.18	-0.19	1.37	BC017068	Hs.105956	QPRT	quinolate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating))
1.17	0.25	0.92	BC008628	Hs.5710	LOC200383	similar to Dynein heavy chain at 16F
1.17	0.59	0.58	BC005830	Hs.279928	RNH	ribonuclease/angiogenin inhibitor
1.16	0.62	0.54	BC015022	Hs.84038	KIAA0111	KIAA0111 gene product
1.16	0.51	0.66	BC010540	Data not found	FLJ23277	FLJ23277 protein
1.16	-0.39	1.55	BC001129	Hs.323634	MGC2463	hypothetical protein MGC2463
1.16	0.36	0.80	BC016707	Hs.10729	SUI1	putative translation initiation factor
1.16	0.23	0.93	BC002357	Hs.100016	CNOT3	CCR4-NOT transcription complex, subunit 3
1.16	1.14	0.02	BC009291	Hs.90998	PPN	likely ortholog of mouse porcupine homolog (Drosophila)
1.16	-0.75	1.91	BC000511	Hs.377973	PTD012	PTD012 protein
1.16	-1.37	2.53	BC011011	Hs.337778	UEV3	ubiquitin-conjugating enzyme E2-like
1.15	0.61	0.54	BC011922	Hs.90998	MGC2663	hypothetical protein MGC2663
1.15	-0.69	1.84	BC007967	Hs.426460	LDHB	lactate dehydrogenase B
1.15	-0.05	1.20	BC012621	Hs.146322	KLRG1	killer cell lectin-like receptor subfamily G, member 1
1.15	0.05	1.10	BC013885	Hs.26395	FLJ10989	hypothetical protein FLJ10989
1.15	0.92	0.23	BC013775	Hs.26954	GNB2L1	guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
1.15	-0.52	1.67	BC000382	Hs.75117	SERPINB1	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1
1.15	1.12	0.02	BC012857	Hs.31130	C21orf33	chromosome 21 open reading frame 33
1.14	0.75	0.39	BC014071	Hs.70500	NAPG	N-ethylmaleimide-sensitive factor attachment protein, gamma
1.14	0.78	0.37	BC015790	Hs.164464	HSPCB	heat shock 90kDa protein 1, beta
1.14	0.66	0.48	BC009821	Hs.375217	MLCB	myosin, light polypeptide, regulatory, non-sarcomeric (20kD)
1.14	0.83	0.31	BC010084	Hs.44592	DKFZP586M1523	DKFZP586M1523 protein
1.14	1.09	0.05	BC000799	Hs.151536	RAB13	RAB13, member RAS oncogene family
1.14	0.80	0.34	BC020498	Data not found	TALDO1	transaldolase 1

1.13	-0.31	1.44	BC000170	Hs.173611	NDUFS2	NADH dehydrogenase (ubiquinone) Fe-S protein 2, 49kDa (NADH-coenzyme Q reductase)
1.13	0.31	0.83	BC002732	Hs.347349	APBB1	amyloid beta (A4) precursor protein-binding, family B, member 1 (Fe65)
1.13	-0.37	1.50	BC012133	Hs.109281	BZW2	basic leucine zipper and W2 domains 2
1.13	-0.61	1.74	BC004136	Hs.9329	PLAT	plasminogen activator, tissue
1.13	-0.57	1.70	BC005279	Hs.2879	HSPB1	heat shock 27kDa protein 1
1.13	-0.01	1.14	BC000341	Hs.74564	VPS33A	vacuolar protein sorting 33A (yeast)
1.12	0.45	0.67	BC005072	Hs.326732	ZNF43	zinc finger protein 43 (HTF6)
1.12	-0.09	1.21	BC006121	Hs.117062	DKFZP586M1019	DKFZP586M1019 protein
1.12	0.41	0.72	BC010423	Hs.61460	PCTK1	PCTAIRE protein kinase 1
1.12	-0.14	1.26	BC017400	Hs.273387	BIRC5	baculoviral IAP repeat-containing 5 (survivin)
1.12	0.87	0.25	BC000671	Hs.5372	PSMD9	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9
1.12	0.70	0.42	BC009548	Hs.16269	TUBB2	tubulin, beta, 2
1.12	0.84	0.29	BC009353	Hs.737	MAGEH1	APR-1 protein
1.12	0.11	1.01	BC009868	Hs.1608	SCMH1	sex comb on midleg homolog 1 (Drosophila)
1.12	0.11	0.01	BC011904	Hs.130558	LOC91661	hypothetical protein BC001610
1.12	-0.84	1.96	BC009418	Hs.283725	LETM1	leucine zipper-EF-hand containing transmembrane protein 1
1.11	-0.37	1.49	BC005069	Hs.61268	ARHGDI	Rho GDP dissociation inhibitor (GDI) alpha
1.11	-0.41	1.53	BC000683	Hs.10784	SH3GLB1	SH3-domain GRB2-like endophilin B1
1.11	-0.92	2.03	BC011379	Hs.17936	FSCN1	fascin homolog 1, actin-bundling protein (Strongylocentrotus purpuratus)
1.11	0.65	0.46	BC012170	Data not found	DKFZP564C103	DKFZP564C103 protein
1.11	0.88	0.23	BC003083	Hs.177596	THEA	thioesterase, adipose associated
1.10	-0.36	1.46	BC002640	Data not found	LOC55974	stromal cell protein
1.10	0.18	0.92	BC009832	Hs.234546	NINJ2	ninjurin 2
1.10	-0.30	1.40	BC009386	Hs.79410	DOM3Z	dom-3 homolog Z (C. elegans)
1.10	0.76	0.34	BC013414	Hs.75090	NA 4-14-03 345	
1.10	0.52	0.58	BC018993	Hs.356360	FLJ20312	hypothetical protein FLJ20312
1.10	0.67	0.43	BC011561	Hs.31720	VT11B	vesicle transport through interaction with t-SNAREs homolog 1B (yeast)
1.10	-0.14	1.24	BC000179	Hs.79474	YWHAE	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide
1.10	0.78	0.31	BC005307	Hs.171995	HTATIP2	HIV-1 Tat interactive protein 2, 30kDa
1.10	0.97	0.13	BC014538	Hs.161427	ZNF215	zinc finger protein 215
1.10	0.56	0.54	BC001622	Hs.30488	MT1H	metallothionein 1H
1.09	0.19	0.91	BC012729	Hs.288697	DYT1	dystonia 1, torsion (autosomal dominant; torsin A)
1.09	-0.08	1.18	BC009460	Hs.351872	MGC23244	hypothetical protein MGC23244
1.09	-0.07	1.16	BC003176	Hs.273230	FLJ10607	glucosamine-phosphate N-acetyltransferase 1
1.09	-0.42	1.52	BC009964	Hs.7345	DKFZP586D0919	hepatocellularcarcinoma-associated antigen HCA557a
1.09	-1.65	2.74	BC012593	Hs.275215	HSD11B1	hydroxysteroid (11-beta) dehydrogenase 1
1.09	0.66	0.43	BC006405	Hs.74346	RetSDR2	retinal short-chain dehydrogenase/reductase 2
1.09	0.34	0.75	BC008898	Hs.56828	TNRC5	trinucleotide repeat containing 5
1.08	0.37	0.71	BC011701	Hs.446459	OAT	ornithine aminotransferase (gyrate atrophy)
1.08	0.77	0.31	BC007765	Hs.459	NA 4-14-03 12	
1.08	1.02	0.06	BC010703	Hs.160483	NONO	
1.08	0.59	0.49	BC008789	Hs.93574	HOXD3	homeo box D3
1.08	0.83	0.24	BC019231	Data not found	NA 4-14-03 375	
1.08	0.80	0.27	BC014863	Hs.20478	EG1	
1.07	0.45	0.63	BC005060	Hs.8935	SH3GLB2	SH3-domain GRB2-like endophilin B2
1.07	0.17	0.90	BC012060	Hs.120565	MEL	mel transforming oncogene (derived from cell line NK14)- RAB8 homolog
1.07	0.96	0.11	BC011772	Hs.75648	LOC116211	hypothetical protein BC013113
1.07	-0.59	1.66	BC018949	Hs.77492	VT11A	vesicle transport through interaction with t-SNAREs homolog 1A (yeast)
1.06	0.01	1.05	BC008928	Hs.4291	CFL1	cofilin 1 (non-muscle)
1.06	0.70	0.36	BC004101	Hs.193163	FLJ10520	hypothetical protein FLJ10520
1.06	-1.71	2.77	BC014514	Hs.213289	POT1	protection of telomeres 1
1.06	-0.19	1.25	BC007287	Hs.115284	LOC90736	hypothetical protein BC000919
1.06	0.77	0.29	BC007835	Hs.21396	DKFZP564J157	DKFZp564J157 protein

1.06	1.03	0.02	BC015936	Hs.6856	RPS25	ribosomal protein S25
1.06	-1.63	2.69	BC015179	Hs.12102	MED8	mediator of RNA polymerase II transcription, subunit 8 homolog (yeast)
1.05	-0.56	1.62	BC006006	Hs.333526	GALK1	galactokinase 1
1.05	0.57	0.48	BC017716	Hs.95655	MGC20533	similar to RIKEN cDNA 2410004L22 gene (M. musculus)
1.05	0.44	0.61	BC003627	Hs.80624	MGC2560	hypothetical protein MGC2560
1.05	-0.58	1.63	BC000965	Hs.97996	MTERF	transcription termination factor, mitochondrial
1.05	1.03	0.02	BC019348	Data not found	NA 4-14-03 435	
1.05	0.04	1.01	BC002449	Hs.24391	CRMP5	collapsin response mediator protein-5; CRMP3-associated molecule
1.05	0.25	0.79	BC005126	Hs.284202	BGN	biglycan
1.04	0.82	0.22	BC019096	Hs.21811	DGCR8	DiGeorge syndrome critical region gene 8
1.04	0.54	0.50	BC001748	Hs.217493	LOC112817	hypothetical protein BC011916
1.04	-0.59	1.63	BC004219	Data not found	ALDH9A1	aldehyde dehydrogenase 9 family, member A1
1.04	0.01	1.03	BC000032	Hs.8047	HNOEL-iso	HNOEL-iso protein
1.04	-0.68	1.72	BC022845	Hs.6353	MRG15	mortality factor 4 like 1
1.04	0.93	0.11	BC013038	Hs.108183	NUDT3	nudix (nucleoside diphosphate linked moiety X)-type motif 3
1.04	0.65	0.39	BC006544	Hs.226396	PX19	px19-like protein
1.04	0.19	0.85	BC004938	Hs.184640	KEAP1	Kelch-like ECH-associated protein 1
1.04	-0.83	1.87	BC011870	Hs.116056	ZNF271	zinc finger protein 271
1.04	-0.16	1.20	BC002630	Hs.355722	GPR87	G protein-coupled receptor 87
1.04	-0.36	1.40	BC002858	Hs.178471	KIAA0701	KIAA0701 protein
1.04	0.38	0.66	BC010931	Hs.78881	MEF2B	MADS box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B)
1.04	0.32	0.71	BC017172	Hs.79259	FAIM2	Fas apoptotic inhibitory molecule 2
1.04	0.46	0.58	BC008919	Hs.155829	CBR1	carbonyl reductase 1
1.04	0.11	0.92	BC010907	Hs.52256	FLJ20624	hypothetical protein FLJ20624
1.04	-0.24	1.28	BC002843	Hs.433469	HSPC043	HSPC043 protein
1.04	-0.19	1.23	BC018633	Hs.256583	TETRAN	tetracycline transporter-like protein
1.04	-1.42	2.46	BC016706	Hs.106673	SPAG9	sperm associated antigen 9
1.03	0.69	0.34	BC004250	Hs.400124	KIAA0063	KIAA0063 gene product
1.03	0.39	0.65	BC004931	Hs.421608	EIF3S7	eukaryotic translation initiation factor 3, subunit 7 zeta, 66/67kDa
1.03	0.64	0.39	AB058759	Hs.381163	TUBA6	tubulin alpha 6
1.03	0.27	0.76	BC000423	Hs.7476	ATP6V0B	ATPase, H+ transporting, lysosomal 21kDa, V0 subunit c''
1.03	-0.46	1.49	BC007624	Hs.322852	CCNH	cyclin H
1.03	0.53	0.50	BC014101	Hs.273357	ALDH6A1	aldehyde dehydrogenase 6 family, member A1
1.03	-1.25	2.28	BC018130	Hs.154299	KRT15	keratin 15
1.03	0.85	0.18	BC011940	Data not found	FLJ21432	hypothetical protein FLJ21432
1.03	0.40	0.64	BC013155	Hs.330208	SMUG1	single-strand selective monofunctional uracil DNA glycosylase
1.03	0.16	0.87	BC007303	Hs.334586	HRB2	HIV-1 rev binding protein 2
1.03	0.84	0.19	BC012571	Hs.50889	HUMMLC2B	myosin light chain 2
1.03	0.03	1.00	BC000241	Hs.71787	FLJ21839	hypothetical protein FLJ21839
1.03	-0.31	1.33	BC015625	Hs.50441	LOC57038	similar to arginyl-tRNA synthetase
1.03	0.86	0.17	BC011746	Hs.26267	APOBEC3D	apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 3D
1.03	-0.32	1.35	BC013577	Hs.31595	CTSS	cathepsin S
1.03	-0.88	1.91	BC020965	Hs.287850	FLJ10432	hypothetical protein FLJ10432
1.02	0.02	1.00	BC004248	Hs.170195	SLC25A11	solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11
1.02	-0.36	1.38	BC002663	Hs.298089	KRAS2	v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog
1.02	-0.66	1.68	BC015571	Hs.50868	MGC16037	hypothetical protein MGC16037
1.02	-0.73	1.75	BC014271	Hs.76753	FLJ10979	hypothetical protein FLJ10979
1.02	0.06	0.96	BC016700	Hs.9265	DKFZP434J154	DKFZP434J154 protein
1.02	-0.22	1.24	BC011519	Hs.3697	EIF3S7	eukaryotic translation initiation factor 3, subunit 7 zeta, 66/67kDa
1.02	0.12	0.90	BC009047	Hs.18953	PRCC	papillary renal cell carcinoma (translocation-associated)
1.02	-0.42	1.44	BC013876	Hs.278898	EKI1	ethanolamine kinase
1.02	0.21	0.81	BC004540	Hs.194286	DNAJA2	DnaJ (Hsp40) homolog, subfamily A, member 2
1.02	0.37	0.65	BC002475	Hs.408943	MGC2396	hypothetical zinc finger protein MGC2396



1.02	0.76	0.25	BC013816	Hs.46	RPS28	ribosomal protein S28
1.02	0.09	0.93	BC000148	Hs.184376	SNAP23	synaptosomal-associated protein, 23kDa
1.02	-0.29	1.30	BC011927	Hs.65234	NA 4-14-03 141	
1.01	0.51	0.51	BC016879	Hs.29857	GLUL	glutamate-ammonia ligase (glutamine synthase)
1.01	0.29	0.73	BC011453	Hs.380833	HNRPD	heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kDa)
1.01	0.08	0.93	BC001692	Hs.33084	SLC2A5	solute carrier family 2 (facilitated glucose/fructose transporter), member 5
1.01	0.25	0.76	BC009980	Hs.343245	AGTRAP	angiotensin II receptor-associated protein
1.01	-0.70	1.71	BC008122	Data not found	FLJ23306	hypothetical protein FLJ23306
1.01	-0.25	1.26	BC008092	Hs.406511		
1.01	0.68	0.33	BC012942	Hs.180378	LSM4	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae)
1.01	-0.15	1.16	BC001400	Hs.206501	LOC57228	hypothetical protein from clone 643
1.01	0.49	0.52	BC014781	Hs.348401	LIPT1	lipoyltransferase
1.01	0.51	0.49	BC009935	Hs.181202	CLK3	CDC-like kinase 3
1.01	0.72	0.29	BC006236	Hs.99196	MGC11324	hypothetical protein MGC11324
1.00	0.19	0.81	BC004530	Hs.15580	PAFAH2	platelet-activating factor acetylhydrolase 2, 40kDa
1.00	-0.29	1.29	BC001697	Hs.433406	RPS15A	ribosomal protein S15a
1.00	0.54	0.46	BC007073	Hs.5085	FABP5	fatty acid binding protein 5 (psoriasis-associated)
1.00	0.28	0.72	BC007696	Hs.334604	TIP47	cargo selection protein (mannose 6 phosphate receptor binding protein)
1.00	-0.73	1.73	BC011231	Data not found	NA 4-14-03 392	
1.00	0.33	0.67	BC005253	Hs.106290	KIAA0107	KIAA0107 gene product
1.00	-0.20	1.20	BC014315	Hs.30174	E2-EPF	ubiquitin carrier protein
1.00	-0.68	1.68	BC001280	Hs.250822	STK6	serine/threonine kinase 6
0.99	-0.10	1.09	BC010356	Hs.70333	WAC	WW domain-containing adapter with a coiled-coil region
0.99	-0.07	1.07	BC016147	Hs.1119	NSPC1	likely ortholog of mouse nervous system polycomb 1
0.99	-1.61	2.61	BC013083	Hs.304682	DSCR5	Down syndrome critical region gene 5
0.99	0.79	0.21	BC012179	Hs.27931	FLJ10830	cytosolic nonspecific dipeptidase (EC 3.4.13.18)
0.99	0.86	0.13	BC004143	Hs.69771	BF	B-factor, properdin
0.99	0.90	0.09	BC007613	Hs.155392	TSLP	thymic stromal lymphopoietin
0.99	0.71	0.28	BC018346	Hs.16165	ZNF258	zinc finger protein 258
0.99	0.69	0.30	BC000694	Hs.250493	SCGN	secretagogin, EF-hand calcium binding protein
0.99	0.96	0.03	BC020566	Hs.289271	MGC12518	similar to ZINC FINGER PROTEIN 257 (BONE MARROW ZINC FINGER 4)
(BMZF-4)						
0.99	0.01	0.98	BC009921	Hs.130965	SH2D3A	SH2 domain containing 3A
0.99	0.31	0.68	BC009416	Hs.194864	FLJ12448	hypothetical protein FLJ12448
0.99	0.16	0.83	BC018514	Hs.30246	COMT	catechol-O-methyltransferase
0.99	0.38	0.61	BC012863	Hs.239	CGI-105	CGI-105 protein
0.99	0.34	0.65	BC014225	Hs.13775	C9orf16	chromosome 9 open reading frame 16
0.99	-0.93	1.92	BC017423	Hs.222746	FLJ12949	hypothetical protein FLJ12949
0.99	-0.74	1.73	BC001886	Hs.75319	NA 4-14-03 82	
0.99	-0.55	1.54	BC010854	Hs.3763	CPR2	cell cycle progression 2 protein
0.99	-0.56	1.54	BC006541	Hs.149846	CED-6	CED-6 protein
0.99	0.19	0.79	BC009071	Hs.42514	C9orf9	chromosome 9 open reading frame 9
0.99	0.32	0.67	BC009710	Hs.100651	NRN1	neuritin 1
0.98	0.43	0.55	BC003608	Hs.80248	MRPS21	mitochondrial ribosomal protein S21
0.98	-1.69	2.67	BC000150	Hs.161279	FLJ23360	hypothetical protein FLJ23360
0.98	0.34	0.64	BC011973	Hs.4932	SMP1	small membrane protein 1
0.98	0.70	0.28	BC009699	Data not found	NXT1	NTF2-like export factor 1
0.98	-0.27	1.24	BC014649	Hs.17958	KIAA1160	KIAA1160 protein
0.98	-0.78	1.75	BC005033	Hs.182485	MSLN	mesothelin
0.98	-0.70	1.67	BC015943	Hs.348350	MGC10911	hypothetical protein MGC10911
0.98	0.96	0.01	BC003079	Hs.180859	LOC51142	16.7Kd protein
0.98	-0.90	1.88	BC004818	Hs.4104	HSPC210	chromosome 14 open reading frame 129
0.98	-0.07	1.05	BC012181	Hs.59242	VAT1	vesicle amine transport protein 1 homolog (T californica)

0.97	0.86	0.11	BC006134	Hs.303653	TPI1	triosephosphate isomerase 1
0.97	-0.26	1.23	BC011399	Hs.74101	MYOD1	myogenic factor 3
0.97	0.40	0.57	BC013278	Hs.40342	PFC	properdin P factor, complement
0.97	0.92	0.05	BC021620	Hs.433387	MGC10646	hypothetical protein MGC10646
0.97	0.27	0.70	BC009923	Hs.69235	HEI10	enhancer of invasion 10
0.97	-2.74	3.71	BC001745	Hs.79404	MGC5178	hypothetical protein MGC5178
0.97	-0.52	1.49	BC010904	Hs.23495	FLJ11252	HBxAg transactivated protein 1
0.97	0.63	0.33	BC014191	Hs.379380	AKR1B10	aldo-keto reductase family 1, member B10 (aldose reductase)
0.97	-0.44	1.41	BC001687	Hs.427212	NARS	asparaginyl-tRNA synthetase
0.97	-0.14	1.10	BC003595	Hs.9732	MGC4368	hypothetical protein MGC4368
0.97	0.26	0.70	BC002509	Hs.288217	MGC2941	hypothetical protein MGC2941
0.96	-0.05	1.02	BC009885	Hs.83848	NA 4-14-03 554	
0.96	0.84	0.13	BC015735	Hs.154899	NEK6	NIMA (never in mitosis gene a)-related kinase 6
0.96	0.85	0.12	BC006496	Hs.15456	K-ALPHA-1	tubulin, alpha, ubiquitous
0.96	-0.04	1.01	BC011935	Hs.240013	FAM14B	family with sequence similarity 14, member B
0.96	0.37	0.59	BC010858	Hs.77256	PTD015	PTD015 protein
0.96	-0.06	1.01	BC001821	Hs.323949	KAI1	kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))
0.95	-0.22	1.17	BC007235	Hs.320834	HSPC177	hypothetical protein HSPC177
0.95	0.73	0.22	BC004498	Hs.283639	FLJ12671	hypothetical protein FLJ12671
0.95	0.08	0.87	BC015786	Data not found	PPP1R8	protein phosphatase 1, regulatory (inhibitor) subunit 8
0.95	-0.68	1.64	BC005864	Hs.95577	CDK4	cyclin-dependent kinase 4
0.95	-0.69	1.64	BC003164	Hs.78768	C6orf29	chromosome 6 open reading frame 29
0.95	0.61	0.34	BC002506	Hs.28866	PDCD10	programmed cell death 10
0.95	-0.63	1.58	BC015356	Hs.82159	IL18	interleukin 18 (interferon-gamma-inducing factor)
0.95	-2.34	3.29	BC019341	Hs.211612	SEC24A	SEC24 related gene family, member A (S. cerevisiae)
0.95	-0.76	1.71	BC000517	Data not found	PPP2R5D	protein phosphatase 2, regulatory subunit B (B56), delta isoform
0.95	-0.38	1.33	BC009073	Hs.112654	MID2	midline 2
0.95	0.09	0.85	BC009311	Hs.75227	CDC25C	cell division cycle 25C
0.94	0.37	0.58	BC009383	Hs.73452	ASGR2	asialoglycoprotein receptor 2
0.94	0.84	0.10	BC008873	Hs.288912	MGC17528	
0.94	0.53	0.42	BC016051	Hs.287863	NA 4-14-03 447	
0.94	-0.04	0.99	BC012359	Hs.106671	TCTA	T-cell leukemia translocation altered gene
0.94	0.26	0.68	BC004367	Hs.240395	RAB7L1	RAB7, member RAS oncogene family-like 1
0.94	0.39	0.54	BC011593	Hs.344097	NALP2	NALP2 protein
0.94	-0.01	0.95	BC016931	Hs.30985	FLJ20489	hypothetical protein FLJ20489
0.94	-0.37	1.31	BC005828	Hs.6262		
0.94	0.24	0.69	BC013879	Hs.3838	PLTP	phospholipid transfer protein
0.94	0.08	0.86	BC010647	Hs.84389	MADH4	MAD, mothers against decapentaplegic homolog 4 (Drosophila)
0.94	0.70	0.24	BC005321	Hs.332405	CCBP2	chemokine binding protein 2
0.94	0.10	0.83	BC014001	Hs.76591	MGC2654	hypothetical protein MGC2654
0.93	-0.19	1.13	BC015814	Hs.106614	FLJ22215	hypothetical protein FLJ22215
0.93	-0.33	1.27	BC007323	Hs.193313	ADSL	adenylosuccinate lyase
0.93	-0.46	1.40	BC016855	Hs.181243	ELA3A	elastase 3A, pancreatic (protease E)
0.93	0.17	0.77	BC009567	Hs.159440	MGC12679	hypothetical protein MGC12679
0.93	-0.92	1.86	BC005324	Hs.78224	SMBP	SM-11044 binding protein
0.93	-0.06	0.99	BC003523	Hs.181173	LIMR	lipocalin-interacting membrane receptor
0.93	0.75	0.17	BC008917	Hs.42806	MT3	metallothionein 3 (growth inhibitory factor (neurotrophic))
0.93	0.43	0.50	BC009993	Data not found	FLJ23186	hypothetical protein FLJ23186
0.93	0.60	0.33	BC010649	Hs.3945	ATP6V0B	ATPase, H <sup>+</sup> transporting, lysosomal 21kDa, V0 subunit c''
0.93	0.43	0.49	BC017032	Hs.194710	RPLP2	
0.92	0.83	0.09	BC014656	Hs.15259	HPRP3P	U4/U6-associated RNA splicing factor
0.92	0.89	0.03	BC009829	Hs.405885	NA 4-14-03 550	

0.92	-1.61	2.54	BC009962	Hs.279939	TRIP4	thyroid hormone receptor interactor 4
0.92	0.24	0.68	BC004244	Hs.821	MGC10200	hypothetical protein MGC:10200
0.92	-0.34	1.26	BC001777	Hs.288654	PSMB7	proteasome (prosome, macropain) subunit, beta type, 7
0.92	0.43	0.49	BC004366	Hs.302446	LOC51042	zinc finger protein
0.92	-0.01	0.93	BC001022	Hs.184011	CRSP8	
0.92	-0.93	1.85	BC013903	Hs.155191	ILVBL	ilvB (bacterial acetolactate synthase)-like
0.92	-1.25	2.17	BC006122	Hs.13024	NA 4-14-03 433	
0.92	-0.69	1.61	BC007750	Hs.85524	MGC3262	hypothetical protein MGC3262
0.92	0.61	0.31	BC007052	Hs.418901	BBX	bobby sox homolog (Drosophila)
0.92	0.37	0.55	BC003552	Hs.155560	DDX1	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 1
0.92	0.60	0.32	BC009876	Hs.184519	MI-ER1	mesoderm induction early response 1
0.92	-0.44	1.35	BC007932	Hs.26270	GOLGA2	golgi autoantigen, golgin subfamily a, 2
0.91	0.58	0.33	BC013003	Hs.84264	CKS1B	CDC28 protein kinase regulatory subunit 1B
0.91	0.73	0.19	BC008071	Hs.169531	GNB2L1	guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
0.91	0.33	0.58	BC007330	Hs.108258	NA 4-14-03 221	
0.91	-0.76	1.66	BC015566	Hs.380447	NFE2	nuclear factor (erythroid-derived 2), 45kDa
0.91	0.18	0.73	BC015805	Data not found	DKFZP547E1010	DKFZP547E1010 protein
0.91	0.47	0.44	BC006214	Hs.26110	CGI-128	CGI-128 protein
0.91	0.73	0.17	BC011372	Hs.154576	RPS13	ribosomal protein S13
0.90	-0.01	0.91	BC015866	Hs.348334		
0.90	0.41	0.49	BC003152	Hs.7854	EGFR-RS	likely ortholog of mouse epidermal growth factor receptor, related sequence
0.90	-0.12	1.02	BC006403	Hs.54589	GTPBG3	mitochondrial GTP binding protein
0.90	0.36	0.54	BC015816	Hs.101474	ELKS	ELKS protein
0.90	-0.33	1.23	BC002342	Hs.17377	BCAN	chondroitin sulfate proteoglycan BEHAB/brevican
0.90	0.67	0.23	BC006404	Hs.21263	POLR2D	polymerase (RNA) II (DNA directed) polypeptide D
0.90	0.48	0.42	BC013280	Hs.298553	C20orf140	chromosome 20 open reading frame 140
0.90	0.52	0.37	BC007221	Hs.22880	KIAA1165	Nedd4 family interacting protein 2
0.90	-0.78	1.68	BC016161	Hs.134185	MGC15476	thymus expressed gene 3-like
0.89	-0.47	1.36	BC000135	Hs.78769	THOP1	thimet oligopeptidase 1
0.89	-0.52	1.41	BC014546	Hs.393201	ACTR2	ARP2 actin-related protein 2 homolog (yeast)
0.89	0.76	0.14	BC001392	Hs.311640	RPS27A	ribosomal protein S27a
0.89	0.36	0.53	BC002477	Hs.271586	MGC4093	hypothetical protein MGC4093
0.89	-0.26	1.15	BC006143	Hs.333389	SYNGR3	synaptogyrin 3
0.89	0.76	0.13	BC000905	Hs.227327	RAB1A	RAB1A, member RAS oncogene family
0.89	-0.06	0.95	BC004373	In multiple clusters	FLJ10504	misato
0.89	-1.09	1.98	BC007310	Hs.332563	KIAA1143	KIAA1143 protein
0.89	0.51	0.37	BC012125	Hs.284205	RPS24	ribosomal protein S24
0.88	-0.44	1.33	BC008910	Hs.1690	NA 4-14-03 226	
0.88	0.28	0.60	BC007996	Hs.286	HNRPF	heterogeneous nuclear ribonucleoprotein F
0.88	0.21	0.67	BC000184	Hs.11776	KIAA1169	two-pore channel 1, homolog
0.88	0.45	0.43	BC015923	Hs.347964	DKFZP566F0546	DKFZP566F0546 protein
0.88	0.16	0.73	BC004935	Hs.102910	FLJ20695	hypothetical protein FLJ20695
0.88	0.66	0.22	BC013998	Hs.92282	FLJ14107	hypothetical protein FLJ14107
0.88	-0.28	1.17	BC014157	Hs.348316	FLJ22795	hypothetical protein FLJ22795
0.88	-0.24	1.12	BC001254	Hs.342849	ARL5	ADP-ribosylation factor-like 5
0.88	0.58	0.30	BC001174	Hs.184161	HINT1	histidine triad nucleotide binding protein 1
0.88	0.04	0.84	BC012078	Hs.30783	DPH2L2	diphtheria toxin resistance protein required for diphthamide biosynthesis-like 2 (S. cerevisiae)
0.88	0.61	0.26	BC007646	Hs.177926	LOC51142	16.7Kd protein
0.88	0.30	0.57	BC011027	Hs.23581	HSOBRGRP	leptin receptor gene-related protein
0.88	0.20	0.68	BC006528	Hs.74107	MGC13008	hypothetical protein MGC13008
0.88	0.03	0.85	BC012328	Hs.132748	FLJ10307	hypothetical protein FLJ10307
0.87	-0.17	1.04	BC007373	Hs.334714	FLJ14639	hypothetical protein FLJ14639
0.87	0.00	0.87	BC013007	Hs.68257	ZNF31	zinc finger protein 31 (KOX 29)

0.87	-1.12	1.99	BC011696	Hs.155020	GPRC5C	G protein-coupled receptor, family C, group 5, member C
0.87	-0.01	0.88	BC000494	Hs.170001	EIF2B2	eukaryotic translation initiation factor 2B, subunit 2 beta, 39kDa
0.87	-1.74	2.61	BC000699	Hs.4975	EXO1	exonuclease 1
0.87	-0.52	1.39	BC001230	Hs.108408	APH-1A	likely ortholog of <i>C. elegans</i> anterior pharynx defective 1A
0.87	-0.80	1.67	BC000296	Hs.15519	SPOP	speckle-type POZ protein
0.86	-0.01	0.88	BC013323	Hs.78950	FLJ10241	hypothetical protein FLJ10241
0.86	0.03	0.84	BC011571	Hs.98074	COPE	coatamer protein complex, subunit epsilon
0.86	-0.21	1.07	BC020079	Hs.152931	NRBF-2	nuclear receptor binding factor-2
0.86	0.10	0.77	BC008369	Hs.96870	HYAL3	hyaluronoglucosaminidase 3
0.86	0.82	0.04	BC012176	Hs.240951	MGC4614	hypothetical protein MGC4614
0.86	-0.12	0.98	BC001164	Hs.78466	HMOX1	heme oxygenase (decycling) 1
0.86	-0.22	1.08	BC000177	Hs.181195	DNAJB6	DnaJ (Hsp40) homolog, subfamily B, member 6
0.86	0.57	0.29	NM_006779	Hs.12289	GTF2H1	general transcription factor IIH, polypeptide 1, 62kDa
0.86	-0.41	1.27	BC009594	Hs.7917	HIG1	likely ortholog of mouse hypoxia induced gene 1
0.86	-0.24	1.10	BC004156	Hs.426460	RPL9	ribosomal protein L9
0.86	-0.50	1.36	BC014520	Hs.80132	FUSIP1	FUS interacting protein (serine-arginine rich) 1
0.86	0.48	0.38	BC007387	Hs.288716	FLJ13614	hypothetical protein FLJ13614
0.86	0.09	0.77	BC000017	Hs.152894	ATP5F1	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex, subunit b, isoform 1
0.86	0.85	0.01	BC007776	Hs.233765	K-ALPHA-1	tubulin, alpha, ubiquitous
0.86	-0.73	1.59	BC008230	Hs.405042	NA 4-14-03 60	
0.86	0.00	0.86	BC013981	Hs.180895	RNGTT	RNA guanylyltransferase and 5'-phosphatase
0.86	0.38	0.48	BC016774	Hs.217493	SLC25A6	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 6
0.86	0.55	0.31	BC008360	Hs.258494	C21orf51	chromosome 21 open reading frame 51
0.86	-0.58	1.44	BC000515	Hs.26009	LOC113026	similar to phospholipase C, delta
0.86	0.78	0.07	BC011054	Hs.334878	PCM1	pericentriolar material 1
0.86	-0.17	1.02	BC010640	Hs.166684	EIF3S2	eukaryotic translation initiation factor 3, subunit 2 beta, 36kDa
0.86	-0.63	1.49	BC004260	Hs.112218	FLJ20457	hypothetical protein FLJ20457
0.86	0.67	0.18	BC006464	Hs.425808	CALM2	calmodulin 2 (phosphorylase kinase, delta)
0.85	-0.07	0.93	BC021668	Hs.355719	LZIC	leucine zipper and CTNNBIP1 domain containing
0.85	0.05	0.80	BC005143	Hs.92384	GTPBG3	mitochondrial GTP binding protein
0.85	0.42	0.43	BC013437	Hs.182280	FLJ10346	hypothetical protein FLJ10346
0.85	-0.53	1.39	NM_005950	Hs.433391	COX7B	cytochrome c oxidase subunit VIIb
0.85	-0.94	1.80	BC004383	Hs.172550	VAT1	vesicle amine transport protein 1 homolog (T californica)
0.85	-0.29	1.14	BC011621	Hs.250752	MGC13204	hypothetical protein MGC13204
0.85	0.49	0.37	BC011674	Hs.153357	RAB7	RAB7, member RAS oncogene family
0.85	0.42	0.44	BC003690	Hs.61790	FLJ23338	importin 4
0.85	0.36	0.49	BC005010	Hs.78909	HLA-E	major histocompatibility complex, class I, E
0.85	0.59	0.26	BC009824	Hs.68090	PWP2H	PWP2 periodic tryptophan protein homolog (yeast)
0.85	-0.23	1.08	BC001709	Hs.220324	BAIAP2	BAII-associated protein 2
0.85	0.03	0.81	BC010661	Hs.48403	MGC4707	hypothetical protein MGC4707
0.85	-0.15	1.00	BC006570	Hs.110837	INVS	inversin
0.85	-0.22	1.07	BC013768	Hs.63788	GGCX	gamma-glutamyl carboxylase
0.85	-0.89	1.73	BC000806	Hs.351475	POLR2K	polymerase (RNA) II (DNA directed) polypeptide K, 7.0kDa
0.84	-0.22	1.06	BC016926	Hs.74552	NA 4-14-03 111	
0.84	-1.10	1.94	BC011679	Hs.36574	NA 4-14-03 242	
0.84	0.11	0.74	BC009327	Hs.374534	FLJ00026	hypothetical protein FLJ00026
0.84	-0.12	0.96	BC008602	Hs.129758	EIF3S3	eukaryotic translation initiation factor 3, subunit 3 gamma, 40kDa
0.84	-1.32	2.16	BC001531	Hs.315167	MGC3329	hypothetical protein MGC3329
0.84	-0.33	1.18	BC000992	Hs.10526	CSRP2	cysteine and glycine-rich protein 2
0.84	-0.44	1.28	BC012395	Hs.89606	DKC1	dyskeratosis congenita 1, dyskerin
0.84	-0.35	1.19	BC005884	Data not found	NA 4-14-03 336	
0.84	-0.09	0.93	BC007639	Hs.104938	C15orf15	chromosome 15 open reading frame 15

0.84	-0.21	1.05	BC001688	Hs.352415	SLC39A4	solute carrier family 39 (zinc transporter), member 4
0.84	0.26	0.58	BC012162	Hs.7165	POLE3	polymerase (DNA directed), epsilon 3 (p17 subunit)
0.84	0.07	0.77	BC010956	Hs.374988	FGF7	fibroblast growth factor 7 (keratinocyte growth factor)
0.84	0.11	0.73	BC004884	Hs.326744	NA 4-14-03 271	
0.84	-0.11	0.95	BC010850	Hs.272688	PPARD	peroxisome proliferative activated receptor, delta
0.84	0.62	0.22	BC017114	Hs.118183	C11orf10	chromosome 11 open reading frame 10
0.84	0.59	0.25	BC013782	Hs.78935	MGC5466	hypothetical protein MGC5466
0.83	-0.13	0.96	BC004858	Hs.430121	ITM2B	integral membrane protein 2B
0.83	-1.32	2.15	BC005064	Hs.13972	ANKH	ankylosis, progressive homolog (mouse)
0.83	-0.31	1.14	BC014424	Data not found	MGC4618	hypothetical protein MGC4618
0.83	0.38	0.45	BC009906	Hs.173103	FLJ20303	hypothetical protein FLJ20303
0.83	-0.14	0.98	BC009441	Hs.180903	CTSG	cathepsin G
0.83	-0.71	1.54	BC000654	Hs.242463	PSMD11	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11
0.83	0.39	0.45	BC009698	Hs.268571	DUSP4	dual specificity phosphatase 4
0.83	-0.28	1.12	BC000481	Hs.198281	PKM2	pyruvate kinase, muscle
0.83	0.62	0.21	BC011776	Hs.300772	HLA-DOA	major histocompatibility complex, class II, DO alpha
0.83	-0.35	1.18	BC005932	Hs.82159	PCTK1	PCTAIRE protein kinase 1
0.83	0.64	0.19	BC008718	Hs.1578	GRIM19	cell death-regulatory protein GRIM19
0.83	-0.38	1.21	BC017301	Hs.5345	HTF9C	HpaII tiny fragments locus 9C
0.83	0.54	0.28	BC013172	Data not found	MGC3222	hypothetical protein MGC3222
0.83	0.18	0.65	BC012081	Hs.238270	MIC2	CD99 antigen
0.83	0.16	0.67	BC015801	Hs.64016	RFC2	replication factor C (activator 1) 2, 40kDa
0.83	0.55	0.28	BC004922	Hs.420106	MGC15619	hypothetical protein MGC15619
0.83	-0.34	1.17	BC003093	Hs.377992	MGC10715	novel C2H2 type zinc finger protein
0.82	0.27	0.56	BC014300	Data not found	C22orf20	chromosome 22 open reading frame 20
0.82	-0.12	0.95	BC001857	Hs.95867	LAGY	homeodomain-only protein
0.82	-0.84	1.67	BC015156	Hs.418650	IL13RA1	interleukin 13 receptor, alpha 1
0.82	-0.59	1.41	BC004271	Hs.217112	TAPBP-R	TAP binding protein related
0.82	0.38	0.45	BC021285	Hs.424414	PXMP2	peroxisomal membrane protein 2, 22kDa
0.82	-0.18	1.00	BC004202	Hs.20295	LCMT	leucine carboxyl methyltransferase 1
0.82	-0.95	1.77	BC014573	Hs.154938	CALR	calreticulin
0.82	-0.07	0.89	BC006358	Hs.66048	BAT1	HLA-B associated transcript 1
0.82	-0.37	1.19	BC012815	Hs.94395	C20orf154	chromosome 20 open reading frame 154
0.82	0.72	0.10	BC016281	Hs.227817	TUBA6	tubulin alpha 6
0.82	0.02	0.80	BC007424	Hs.374973	PRPF4	PRP4 pre-mRNA processing factor 4 homolog (yeast)
0.82	0.67	0.15	BC009713	Hs.380938	FLJ20337	hypothetical protein FLJ20337
0.82	-0.61	1.43	BC011130	Hs.254105	LOC51256	hypothetical protein LOC51256
0.82	0.05	0.76	BC011713	Hs.9925	KIAA1811	KIAA1811 protein
0.82	-0.12	0.93	BC000004	Hs.146329	RPS2	ribosomal protein S2
0.81	0.42	0.39	BC000907	Hs.99598	MGC5338	hypothetical protein MGC5338
0.81	0.53	0.28	BC005344	Hs.284162	CGI-141	CGI-141 protein
0.81	0.21	0.61	BC003564	Hs.90336	DKFZP564K2062	DKFZP564K2062 protein
0.81	0.24	0.57	BC009299	Hs.7236	VIL1	villin 1
0.81	0.24	0.57	BC015655	Hs.263928	SORT1	sortilin 1
0.81	0.47	0.34	BC006377	Hs.18079	MGC4730	similar to RIKEN cDNA 3110001D03 gene (M. musculus)
0.81	0.02	0.79	BC006218	Hs.28088	LOC51308	SGC32445 protein
0.81	0.00	0.81	BC010362	In multiple clusters	VPS35	
0.81	0.31	0.50	BC000979	Hs.143187	FLJ10432	hypothetical protein FLJ10432
0.81	0.32	0.49	BC011400	Hs.265960	PEX16	peroxisomal biogenesis factor 16
0.81	-0.54	1.35	BC002577	Hs.82159	HCA127	hepatocellular carcinoma-associated antigen 127
0.81	-1.51	2.32	BC002334	Hs.75416	HAND1	heart and neural crest derivatives expressed 1
0.81	-0.20	1.00	BC015385	Hs.1191	GDBR1	putative glioblastoma cell differentiation-related
0.81	-0.07	0.88	BC016710	Hs.79300	LOC51219	clone FLB5214

0.81	-0.03	0.83	BC003177	Hs.156667		NA 4-14-03 43	
0.81	-0.07	0.87	BC014967	Hs.5637	RTN4	reticulon 4	
0.80	-0.76	1.56	BC006457	Hs.380749	CLTB	clathrin, light polypeptide (Lcb)	
0.80	0.47	0.34	BC015365	Hs.13740	JTV1	JTV1 gene	
0.80	0.23	0.57	BC006190	Hs.171834	PCTK1	PCTAIRE protein kinase 1	
0.80	0.29	0.52	BC006494	Hs.79110	NCL	nucleolin	
0.80	-0.45	1.25	BC001487	Hs.193989	HSPC142	HSPC142 protein	
0.80	0.14	0.66	BC006320	Hs.30318	RBMS1	RNA binding motif, single stranded interacting protein 1	
0.80	0.61	0.19	BC005265	Hs.346918	MAGOH	mago-nashi homolog, proliferation-associated (Drosophila)	
0.80	-0.22	1.02	BC007289	Hs.135411	HCS	cytochrome c, somatic	
0.80	-0.04	0.84	BC011577	Hs.334775	GYS1	glycogen synthase 1 (muscle)	
0.80	0.56	0.25	BC007094	Hs.302740		NA 4-14-03 102	
0.80	-1.08	1.88	BC002811	Data not found	AP1M1	adaptor-related protein complex 1, mu 1 subunit	
0.80	0.58	0.22	BC000018	Hs.315164	EIF2C2	eukaryotic translation initiation factor 2C, 2	
0.80	-0.19	0.99	BC015206	Hs.433898	RBMS1	RNA binding motif, single stranded interacting protein 1	
0.80	0.36	0.44	BC002499	Hs.250282	STK6	serine/threonine kinase 6	
0.80	0.17	0.62	BC012184	Hs.67619	ARF6	ADP-ribosylation factor 6	
0.80	-0.01	0.81	BC020500	Hs.22879	FLJ39653	hypothetical protein FLJ39653	
0.80	0.60	0.19	NM_005762	Hs.228059	SLC9A1	solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (antiporter, Na <sup>+</sup> /H <sup>+</sup> , amiloride sensitive)	
0.80	-0.98	1.77	BC007452	Hs.231840		WBP2	WW domain binding protein 2
0.80	0.11	0.69	BC016284	Hs.15164	KIAA0998	KIAA0998 protein	
0.79	0.59	0.20	BC007722	Hs.293885		MGC4677	hypothetical protein MGC4677
0.79	-0.37	1.17	BC014264	Hs.348553		EEG1	EEG1
0.79	-0.22	1.01	BC002921	Hs.20447	SLC5A6	solute carrier family 5 (sodium-dependent vitamin transporter), member 6	
0.79	-0.04	0.83	BC006180	Hs.153261		IGHM	immunoglobulin heavy constant mu
0.79	0.34	0.45	BC015813	Hs.256184		RAC1	ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)
0.79	0.09	0.70	BC000813	Hs.83916	NDUFA5	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5, 13kDa	
0.79	-1.16	1.95	BC010433	Hs.411818		FLJ22582	hypothetical protein FLJ22582
0.79	0.33	0.46	BC008901	Hs.111092		MGC23401	hypothetical protein MGC23401
0.79	-0.22	1.01	BC007601	Hs.71941	LOC51759	hepatocellular carcinoma-associated antigen 59	
0.79	-0.46	1.25	BC011708	Hs.274313		PDCD6IP	programmed cell death 6 interacting protein
0.79	0.05	0.74	BC017329	Hs.278502		ACTC	actin, alpha, cardiac muscle
0.79	0.59	0.19	BC016180	Hs.274691		MDH2	malate dehydrogenase 2, NAD (mitochondrial)
0.78	0.37	0.42	BC012527	Hs.197540		NORE1	NORE1 protein
0.78	-1.09	1.88	BC002649	Hs.7644	COPS7B	COP9 constitutive photomorphogenic homolog subunit 7B (Arabidopsis)	
0.78	0.31	0.47	BC012057	Data not found		CCNI	cyclin I
0.78	0.18	0.60	BC006994	Hs.125565		C20orf111	chromosome 20 open reading frame 111
0.78	0.27	0.51	BC005984	Hs.44296	IL11RA	interleukin 11 receptor, alpha	
0.78	0.74	0.04	BC000299	Hs.43505	SNAP23	synaptosomal-associated protein, 23kDa	
0.78	-0.45	1.23	BC000697	Hs.184352		DATF1	death associated transcription factor 1
0.78	-0.07	0.85	BC011670	Data not found		SFXN1	sideroflexin 1
0.78	0.13	0.64	BC008023	Hs.62112	HIF1AN	hypoxia-inducible factor 1, alpha subunit inhibitor	
0.78	0.53	0.24	BC008607	Hs.23410	D21S2056E	DNA segment on chromosome 21 (unique) 2056 expressed sequence	
0.78	-0.52	1.30	BC015059	Hs.16606		CGI-32	CGI-32 protein
0.78	0.28	0.50	BC014067	Hs.73986		FKBP4	FK506 binding protein 4, 59kDa
0.78	0.17	0.60	BC010671	Hs.301305		BCS1L	BCS1-like (yeast)
0.78	-0.57	1.35	BC009857	Data not found		SNAPC1	small nuclear RNA activating complex, polypeptide 1, 43kDa
0.78	0.36	0.41	BC001864	Hs.323432		ICAMI	intercellular adhesion molecule 1 (CD54), human rhinovirus receptor
0.77	0.24	0.54	BC007632	Hs.65746	MGC24447	chromosome 14 open reading frame 147	
0.77	0.07	0.70	BC008618	Data not found		SPAG5	sperm associated antigen 5
0.77	0.26	0.51	BC007352	Hs.6980	PXN	paxillin	
0.77	0.51	0.26	BC001701	Hs.378162		HSPC018	HSPC018 protein

0.77	-0.20	0.97	BC002792	Data not found	RAB39B	RAB39B, member RAS oncogene family
0.77	0.39	0.38	BC009771	Hs.279862	BCCIP	BRCA2 and CDKN1A interacting protein
0.77	-0.37	1.14	BC005336	Hs.110853		Homo sapiens, clone IMAGE:4152387, mRNA
0.77	-1.41	2.19	BC015542	Hs.171844	JKK	STE20-like kinase
0.77	-0.04	0.81	BC006315	Hs.314956	TSPAN-3	tetraspan 3
0.77	0.48	0.29	BC004256	Hs.7165	FLJ12827	zinc finger protein 408
0.77	0.65	0.12	BC016288	Hs.379557	FLJ10719	hypothetical protein FLJ10719
0.77	0.43	0.34	BC001225	Hs.20815	HPRP8BP	U5 snRNP-specific 40 kDa protein (hPrp8-binding)
0.77	0.34	0.43	BC006301	Hs.7101	PSMB3	proteasome (prosome, macropain) subunit, beta type, 3
0.77	0.63	0.14	BC009116	Hs.284141	TRNT1	tRNA nucleotidyl transferase, CCA-adding, 1
0.77	-1.40	2.17	BC002831	Hs.280135	EZH1	enhancer of zeste homolog 1 (Drosophila)
0.77	-0.21	0.98	BC009969	Hs.9805	MRPS33	mitochondrial ribosomal protein S33
0.77	0.30	0.47	BC004346	Hs.14927	FAH	fumarylacetoacetate hydrolase (fumarylacetoacetase)
0.77	-1.21	1.97	BC017728	Hs.8578	NA 4-14-03 73	
0.77	0.07	0.70	BC004157	Hs.31895	KIAA1630	KIAA1630 protein
0.77	-0.78	1.55	BC007631	Hs.11849	CRIP1	postsynaptic protein CRIP1
0.77	0.16	0.60	BC016844	Hs.29106	FLJ10759	hypothetical protein FLJ10759
0.77	-1.16	1.93	BC010493	Hs.107381	MGC16142	hypothetical protein MGC16142
0.77	-0.61	1.38	BC010409	Hs.182265		Homo sapiens, clone IMAGE:4877271, mRNA
0.76	0.63	0.13	BC004140	Hs.233950	SPINT1	serine protease inhibitor, Kunitz type 1
0.76	-0.05	0.81	BC000644	Hs.276770	MGC11271	hypothetical protein MGC11271
0.76	-1.63	2.39	BC010541	Data not found	BLP2	BBP-like protein 2
0.76	0.20	0.56	BC008054	Hs.323193	ARL6IP4	ADP-ribosylation-like factor 6 interacting protein 4
0.76	0.24	0.53	BC014535	Hs.44077	PARVA	parvin, alpha
0.76	-0.64	1.40	BC012620	Hs.8868	GOSR1	golgi SNAP receptor complex member 1
0.76	-0.20	0.96	BC010061	Hs.20558	MAGEB1	melanoma antigen, family B, 1
0.76	0.12	0.64	BC018361	Hs.25887	LOC113386	similar to envelope protein
0.76	0.44	0.32	BC010397	Data not found	LYPLA1	
0.76	-0.52	1.28	BC007405	Hs.82240	KPNB1	karyopherin (importin) beta 1
0.76	0.42	0.34	BC010163	Hs.303125	HLA-DPB1	major histocompatibility complex, class II, DP beta 1
0.76	0.31	0.45	BC019068	Hs.413760	APH-1A	likely ortholog of C. elegans anterior pharynx defective 1A
0.76	-1.99	2.75	BC002697	Hs.21537	LOC114928	hypothetical protein BC013576
0.76	-0.20	0.96	BC014659	Hs.334514	LENG4	leukocyte receptor cluster (LRC) member 4
0.76	0.23	0.53	BC016992	Hs.143253	FLJ13386	hypothetical protein FLJ13386
0.76	0.30	0.45	BC001419	Hs.19513	FLJ12436	hypothetical protein FLJ12436
0.76	-0.55	1.30	BC000715	Hs.236516	TSHR	thyroid stimulating hormone receptor
0.76	-1.39	2.15	BC004993	Hs.300711	TAF10	TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 30kDa
0.75	-0.14	0.89	BC012351	Hs.154029	humNRDR	dehydrogenase/reductase (SDR family) member 4
0.75	0.59	0.16	BC015521	Hs.171581	C11orf24	chromosome 11 open reading frame 24
0.75	-0.99	1.74	BC007874	Hs.153636	MGC20785	hypothetical protein MGC20785
0.75	-0.40	1.15	BC012138	Hs.77269	NICE-4	NICE-4 protein
0.75	-1.30	2.05	BC011783	Hs.101742	RP42	RP42 homolog
0.75	-0.04	0.79	BC001255	Hs.240770	APEX1	APEX nuclease (multifunctional DNA repair enzyme) 1
0.75	-0.39	1.15	BC013360	Hs.356590	NA 4-14-03 546	
0.75	-0.56	1.32	BC000556	Hs.178011	Rpo1-2	similar to DNA-directed RNA polymerase I (135 kDa)
0.75	0.10	0.65	BC001396	Hs.284250	AD-003	AD-003 protein
0.75	-0.16	0.91	BC008983	Hs.8986	APH-1A	likely ortholog of C. elegans anterior pharynx defective 1A
0.75	0.49	0.26	BC005122	Hs.13014	S100A13	S100 calcium binding protein A13
0.75	0.28	0.47	BC006410	Hs.283739	C1orf6	chromosome 1 open reading frame 6
0.75	0.34	0.41	BC001043	Hs.15075	DKFZP434E2216	splicing factor 4
0.75	-0.25	1.00	BC008827	Hs.195041	GTF2F1	general transcription factor IIF, polypeptide 1, 74kDa
0.75	-0.11	0.86	BC009727	Hs.282990	CDC42	cell division cycle 42 (GTP binding protein, 25kDa)
0.75	0.19	0.55	BC021196	Hs.13377	AF053356_CDS3	hypothetical protein AF053356_CDS3

0.75	0.22	0.52	BC021161	Hs.267527	FLJ14668	hypothetical protein FLJ14668
0.74	0.45	0.29	BC001456	Hs.173611	NDUFS2	NADH dehydrogenase (ubiquinone) Fe-S protein 2, 49kDa (NADH-coenzyme Q reductase)
0.74	0.19	0.56	BC021909	Data not found	JFC1	NADPH oxidase-related, C2 domain-containing protein
0.74	-0.17	0.91	BC005057	Hs.278712	TACSTD2	tumor-associated calcium signal transducer 2
0.74	0.63	0.12	BC010488	Hs.63168	SSX2	
0.74	0.53	0.21	BC009219	Hs.77039	FLJ10986	hypothetical protein FLJ10986
0.74	0.73	0.01	BC007559	Hs.283781	TPI1	triosephosphate isomerase 1
0.74	0.41	0.33	BC021670	Hs.430150	TIP47	cargo selection protein (mannose 6 phosphate receptor binding protein)
0.74	-0.80	1.54	BC009819	Hs.288158	DDX27	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 27
0.74	0.16	0.58	BC007782	Hs.102950	CAPN1	
0.74	-0.64	1.38	BC007576	Hs.4113	NA 4-14-03 521	
0.74	-0.48	1.22	BC000882	Hs.75187	TOMM20-PENDING	translocase of outer mitochondrial membrane 20 (yeast) homolog
0.74	-0.05	0.79	BC010187	Hs.182740	ID3	inhibitor of DNA binding 3, dominant negative helix-loop-helix protein
0.74	-0.60	1.34	BC010614	Hs.285673	GNPAT	glyceronephosphate O-acyltransferase
0.74	-0.40	1.14	BC000003	Hs.127496	ATP6V1G1	ATPase, H <sup>+</sup> transporting, lysosomal 13kDa, V1 subunit G isoform 1
0.74	-0.21	0.95	BC001038	Data not found	76P	gamma tubulin ring complex protein (76p gene)
0.74	0.60	0.13	BC007405	Hs.82240	STX3A	syntaxin 3A
0.74	0.22	0.51	BC001155	Hs.63525	TPM3	
0.74	-0.20	0.94	BC018142	Hs.306227	NA 4-14-03 97	
0.74	-0.96	1.69	BC009110	Hs.75730	MGC12262	hypothetical protein MGC12262
0.73	-0.99	1.72	BC012141	In multiple clusters	MRPL39	mitochondrial ribosomal protein L39
0.73	0.11	0.62	BC009864	Data not found	FLJ13912	hypothetical protein FLJ13912
0.73	-0.13	0.86	BC008015	Hs.118638	G1P3	interferon, alpha-inducible protein (clone IFI-6-16)
0.73	-0.24	0.98	BC000457	Hs.1575	SNRPD3	small nuclear ribonucleoprotein D3 polypeptide 18kDa
0.73	-0.15	0.88	BC000759	Hs.24563	NXT1	NTF2-like export factor 1
0.73	-0.30	1.04	BC000465	Hs.9701	KIAA0014	KIAA0014 gene product
0.73	0.01	0.72	BC013114	Hs.130712	ARHC	ras homolog gene family, member C
0.73	-1.52	2.25	BC002525	Hs.62515	KIAA0494	KIAA0494 gene product
0.73	-1.19	1.92	BC002530	Hs.399736	CD44	CD44 antigen (homing function and Indian blood group system)
0.73	-0.42	1.15	BC000516	Hs.1276	BN51T	BN51 (BHK21) temperature sensitivity complementing
0.73	0.56	0.17	BC012136	Hs.214397	BAK1	BCL2-antagonist/killer 1
0.73	0.23	0.51	BC020515	Hs.431584	CLECSF2	C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced)
0.73	0.45	0.28	BC014259	Hs.310640	MRPS21	mitochondrial ribosomal protein S21
0.73	-0.26	0.99	BC012612	Hs.78056	CTSL	cathepsin L
0.73	0.43	0.30	BC009257	Hs.59425	NDUFA9	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9, 39kDa
0.73	0.30	0.43	BC012536	Data not found	NA 4-14-03 20	
0.73	-0.18	0.91	BC001484	Hs.75375	SYNJ2BP	synaptojanin 2 binding protein
0.73	0.24	0.48	BC011046	Hs.179600	GDI1	GDP dissociation inhibitor 1
0.73	-1.54	2.27	BC007633	Hs.193053	PVRL3	poliovirus receptor-related 3
0.73	0.31	0.41	BC001500	Hs.75693	ERCC3	excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing)
0.73	0.55	0.17	BC011907	Data not found	NA 4-14-03 90	
0.72	-0.73	1.45	BC014019	Hs.78713	BFAR	bifunctional apoptosis regulator
0.72	0.63	0.10	BC015397	Hs.122055	NA 4-14-03 193	
0.72	0.34	0.38	BC009358	Hs.18851	NDST1	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
0.72	-0.31	1.03	BC004896	Hs.387358	NONO	
0.72	-0.21	0.94	BC002678	Hs.110364	NID67	putative small membrane protein NID67
0.72	0.59	0.13	BC009317	Hs.13222	GPRK6	G protein-coupled receptor kinase 6
0.72	-0.29	1.01	BC000442	Hs.180655		
0.72	-1.16	1.88	BC006172	Hs.269914	C12orf22	chromosome 12 open reading frame 22
0.72	-0.33	1.05	BC001435	Hs.247362	DDAH2	dimethylarginine dimethylaminohydrolase 2
0.72	0.35	0.36	BC011545	Data not found	ACY1	aminoacylase 1



0.72	-0.03	0.75	BC003094	Hs.12570	LSM5	LSM5 homolog, U6 small nuclear RNA associated (S. cerevisiae)
0.72	0.31	0.41	BC016613	Hs.121190	KIAA0303	KIAA0303 protein
0.72	0.64	0.08	BC001616	Hs.81361	HNRPAB	heterogeneous nuclear ribonucleoprotein A/B
0.72	-1.78	2.50	BC003537	Hs.409158	TBPL1	TBP-like 1
0.72	0.06	0.65	BC016843	Hs.351582	NA 4-14-03 144	
0.72	-2.11	2.82	BC004240	Hs.40403	ALEX2	armadillo repeat protein ALEX2
0.72	-0.14	0.85	BC008692	Hs.22868	DXF68S1E	family with sequence similarity 16, member A, X-linked
0.71	0.00	0.72	BC020494	Data not found	MGC4171	hypothetical protein MGC4171
0.71	0.44	0.27	BC012065	Hs.24371	ARF5	ADP-ribosylation factor 5
0.71	-0.15	0.87	BC010996	Hs.23019	MGC27466	hypothetical protein MGC27466
0.71	0.42	0.29	BC007338	Hs.283690	MGC17515	hypothetical protein MGC17515
0.71	0.10	0.62	BC000569	Hs.284142	MAX	MAX protein
0.71	0.09	0.62	BC013293	Hs.76930	CLN8	ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)
0.71	0.22	0.49	BC003380	Hs.71331	LANPL	leucine-rich acidic nuclear protein like
0.71	0.24	0.47	BC001125	Hs.394389	PP1B	peptidylprolyl isomerase B (cyclophilin B)
0.71	0.24	0.47	BC017337	Hs.286088	NA 4-14-03 27	
0.71	-0.61	1.32	BC011897	Hs.29169	UROD	uroporphyrinogen decarboxylase
0.71	-0.38	1.09	BC012850	Hs.13228	DKFZP564B147	DKFZP564B147 protein
0.71	0.12	0.59	BC009774	Hs.91393	LIN7C	lin-7 homolog C (C. elegans)
0.71	0.07	0.64	BC001060	Hs.73149	PAX8	paired box gene 8
0.71	-0.49	1.20	BC014973	Hs.100448	CFL1	cofilin 1 (non-muscle)
0.71	-0.14	0.85	BC017502	Hs.2007	CBCIP2	CAP-binding protein complex interacting protein 2
0.71	-0.76	1.47	BC007334	Hs.283532	CBS	cystathionine-beta-synthase
0.71	-0.38	1.08	BC008884	Hs.153937	ISL2	ISL2 transcription factor, LIM/homeodomain, (islet-2)
0.71	-1.52	2.22	BC005200	Hs.7870	ARHGEF2	rho/rac guanine nucleotide exchange factor (GEF) 2
0.71	0.68	0.03	BC010357	Hs.191381	LOC51319	hypothetical protein LOC51319
0.70	0.41	0.30	BC007637	Hs.194610	NPM1	nucleophosmin (nucleolar phosphoprotein B23, numatrin)
0.70	0.18	0.52	BC007308	Hs.389933	CCNH	cyclin H
0.70	-0.25	0.95	BC001382	Hs.25450	SLC29A1	solute carrier family 29 (nucleoside transporters), member 1
0.70	-0.57	1.27	BC001798	Hs.284291	UBE2C	ubiquitin-conjugating enzyme E2C
0.70	0.20	0.50	BC020953	Hs.180817	COTL1	coactosin-like 1 (Dictyostelium)
0.70	0.43	0.27	BC016996	Hs.17211	MGC19947	core 1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase 2
0.70	-0.17	0.87	BC003030	Hs.79037	HLA-DQB1	major histocompatibility complex, class II, DQ beta 1
0.70	0.54	0.16	BC014129	Hs.296398	CRI1	CREBBP/EP300 inhibitory protein 1
0.70	-0.29	0.99	BC016949	Hs.270956	FXSD2	FXSD domain containing ion transport regulator 2
0.70	-0.39	1.09	BC009530	Data not found	FLJ20045	hypothetical protein FLJ20045
0.70	-1.45	2.15	BC014092	Hs.300463	FLJ20625	hypothetical protein FLJ20625
0.70	0.24	0.46	BC014460	Hs.100764	384D8-2	hypothetical protein 384D8_6
0.70	-0.25	0.94	BC005965	Data not found	MRPL34	mitochondrial ribosomal protein L34
0.70	-0.75	1.45	BC001107	Hs.3530	HBB	hemoglobin, beta
0.70	-0.13	0.83	BC002551	Hs.30114	GRCC8	likely ortholog of mouse gene rich cluster, C8 gene
0.70	-0.33	1.03	BC010175	Hs.379466	PB1	polybromo 1
0.70	-0.26	0.96	BC013289	Hs.19520	NGLY1	N-glycanase 1
0.70	-0.82	1.51	BC001248	Hs.26090	FLJ20272	hypothetical protein FLJ20272
0.70	-0.26	0.95	BC015164	Hs.408943	HNRPAB	heterogeneous nuclear ribonucleoprotein A/B
0.70	-1.02	1.71	BC010365	Hs.32356	FLJ20628	hypothetical protein FLJ20628
0.69	-0.42	1.11	BC001708	Hs.77039	NA 4-14-03 26	
0.69	0.62	0.07	BC003618	Hs.40342	HRIHFB2122	Tara-like protein
0.69	0.37	0.33	BC013357	Hs.15470	HCK	hemopoietic cell kinase
0.69	0.08	0.62	BC006494	Hs.79110	CGI-41	CGI-41 protein
0.69	-0.34	1.03	BC011781	Hs.22164	DKFZP434B168	DKFZP434B168 protein
0.69	0.24	0.45	BC000658	Hs.155223	G6PT1	glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1
0.69	0.63	0.06	BC012075	Hs.85570	MGC3413	hypothetical protein MGC3413

0.69	-0.53	1.22	BC008640	Hs.167927	KCNN4	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4
0.69	0.02	0.67	BC004445	Hs.288800	CCNF	cyclin F
0.69	-0.31	1.00	BC014445	Hs.325650	GAGED3	G antigen, family D, 3
0.69	-0.52	1.21	BC015886	Hs.271277	GRLF1	glucocorticoid receptor DNA binding factor 1
0.69	-0.74	1.43	BC000564	Hs.25817	DKFZP564L2423	hypothetical protein DKFZp564L2423
0.69	0.46	0.22	BC010576	Hs.3005	SNRPD3	small nuclear ribonucleoprotein D3 polypeptide 18kDa
0.69	-2.13	2.82	BC006199	Hs.182579	HIG1	likely ortholog of mouse hypoxia induced gene 1
0.69	-0.80	1.48	BC000396	Hs.75355	AMOTL2	angiomotin like 2
0.68	0.42	0.27	BC008716	Hs.75562	KIAA0738	KIAA0738 gene product
0.68	-0.80	1.49	BC014950	Hs.63287	LOC151568	hypothetical protein BC009491
0.68	-0.81	1.49	BC010183	Hs.110445	LAMP2	lysosomal-associated membrane protein 2
0.68	0.20	0.48	BC001044	Hs.67991	POLR2L	polymerase (RNA) II (DNA directed) polypeptide L, 7.6kDa
0.68	0.05	0.63	BC003355	Hs.11669	LAMA5	laminin, alpha 5
0.68	-0.93	1.61	BC009712	Hs.76781	KRT14	keratin 14 (epidermolysis bullosa simplex, Dowling-Meara, Koebner)
0.68	0.01	0.68	BC015394	Hs.44131	HSD-3.1	spermatogenesis associated 7
0.68	-0.02	0.71	BC014010	Hs.10101	ATP5G1	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1
0.68	0.31	0.38	BC002737	Hs.433827	FLJ12681	hypothetical protein FLJ12681
0.68	-0.30	0.98	BC005024	Hs.16089	FOXO3A	
0.68	0.01	0.68	BC003621	Hs.103804	HNRPU	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)
0.68	-0.18	0.86	BC000916	Hs.74637	TEGT	testis enhanced gene transcript (BAX inhibitor 1)
0.68	-2.32	3.00	BC010042	Hs.91579	RAF1	v-raf-1 murine leukemia viral oncogene homolog 1
0.68	0.24	0.44	BC001271	Hs.8867	CYR61	cysteine-rich, angiogenic inducer, 61
0.68	-0.31	0.99	BC008784	Hs.111279	FLJ10404	hypothetical protein FLJ10404
0.68	-0.47	1.15	BC003067	Hs.76556	CD68	CD68 antigen
0.68	0.35	0.33	BC000487	Hs.296380	FLJ23119	hypothetical protein FLJ23119
0.68	-0.16	0.84	BC013867	Hs.194431	XRN2	5'-3' exoribonuclease 2
0.68	-0.10	0.78	BC013425	Hs.42251	HSPC009	HSPC009 protein
0.68	-0.48	1.16	BC016715	Hs.430150	MGC15875	hypothetical protein MGC15875
0.68	0.20	0.48	BC014146	Hs.28298	FLJ13102	dehydrodolichyl diphosphate synthase
0.68	-0.03	0.71	BC000082	Hs.111801	HEXB	hexosaminidase B (beta polypeptide)
0.68	0.64	0.04	BC015484	Hs.106857	XRCC4	X-ray repair complementing defective repair in Chinese hamster cells 4
0.68	-0.12	0.80	BC016292	Hs.93002	MGC10765	hypothetical protein MGC10765
0.68	0.30	0.38	BC000142	Hs.77171	MCM5	MCM5 minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)
0.68	-1.28	1.96	BC013408	Hs.149997	NAB2	NGFI-A binding protein 2 (EGR1 binding protein 2)
0.68	-0.37	1.04	BC018151	Hs.356019	CDA11	CDA11 protein
0.68	0.07	0.60	BC019313	Hs.172330	MGC2705	hypothetical protein MGC2705
0.67	-0.54	1.21	BC009580	Hs.168669	OGDH	oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide)
0.67	-0.68	1.36	BC018466	Hs.325530	MAP2K2	mitogen-activated protein kinase kinase 2
0.67	0.08	0.60	BC001411	Hs.272630	APOL2	apolipoprotein L, 2
0.67	-0.24	0.92	BC001194	Hs.336780	SPTLC1	serine palmitoyltransferase, long chain base subunit 1
0.67	0.11	0.56	BC004176	Hs.29173	SSH-3	slingshot 3
0.67	-1.97	2.64	BC013966	Hs.86211	TMEM9	transmembrane protein 9
0.67	-0.20	0.87	BC000651	Hs.104637	MRPL9	mitochondrial ribosomal protein L9
0.67	-0.37	1.05	BC001463	Hs.24220	SCOTIN	scotin
0.67	0.66	0.01	BC008188	Hs.10958	DJ-1	Parkinson disease (autosomal recessive, early onset) 7
0.67	-0.89	1.57	BC007656	Hs.93002	SNX6	sorting nexin 6
0.67	-0.55	1.23	BC001801	Data not found	AF1Q	ALL1-fused gene from chromosome 1q
0.67	0.48	0.19	BC001149	Hs.271926	NA 4-14-03 411	
0.67	-0.14	0.81	BC003155	Hs.10326	ITCH	itchy homolog E3 ubiquitin protein ligase (mouse)
0.67	-0.64	1.32	BC001519	Hs.286192	RARRES2	retinoic acid receptor responder (tazarotene induced) 2
0.67	0.48	0.19	BC000586	Hs.285306	MED6	mediator of RNA polymerase II transcription, subunit 6 homolog (yeast)
0.67	0.30	0.37	BC013291	Hs.21431	BCL2L12	BCL2-like 12 (proline rich)
0.67	-0.17	0.84	BC014068	Hs.154196	SSRP1	structure specific recognition protein 1

0.67	0.31	0.36	BC015015	Hs.272838	MGC10771	hypothetical protein MGC10771
0.67	-0.05	0.72	BC015316	Hs.337078	YR-29	hypothetical protein YR-29
0.67	0.58	0.09	BC005202	Hs.380421	PPIL4	peptidylprolyl isomerase (cyclophilin)-like 4
0.67	0.38	0.29	BC000533	Hs.4835	EIF3S8	eukaryotic translation initiation factor 3, subunit 8, 110kDa
0.67	-0.23	0.90	BC016617	Hs.432714	SSR2	signal sequence receptor, beta (translocon-associated protein beta)
0.67	-0.44	1.11	BC017217	Hs.46458	DCTN1	dynactin 1 (p150, glued homolog, Drosophila)
0.67	-1.14	1.81	BC014050	Hs.300876	PDCD7	programmed cell death 7
0.67	0.11	0.56	BC000939	Hs.355888	PLCB2	phospholipase C, beta 2
0.67	-0.18	0.85	BC002532	Hs.433222	PSME2	proteasome (prosome, macropain) activator subunit 2 (PA28 beta)
0.67	0.29	0.37	BC014878	Hs.274430	NOSIP	nitric oxide synthase interacting protein
0.67	-0.03	0.70	BC017712	Hs.693	NA 4-14-03 194	
0.67	0.46	0.21	BC000266	Hs.74823	KLK5	kallikrein 5
0.67	0.54	0.13	BC004188	Hs.251653	TUBB2	tubulin, beta, 2
0.67	-0.26	0.92	BC001011	Hs.172865	CSTF1	cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kDa
0.67	0.03	0.64	BC011580	Data not found	CDT6	angiopoietin-like factor
0.67	0.15	0.51	BC007737	Hs.69285	CDK5RAP3	CDK5 regulatory subunit associated protein 3
0.67	-0.93	1.60	BC010048	Hs.153768	CLIPR-59	CLIP-170-related protein
0.66	-0.29	0.96	BC000725	Hs.99899	GRIPAP1	GRIP1 associated protein 1
0.66	-1.81	2.48	BC005356	Hs.433404	LOC113655	hypothetical protein BC011982
0.66	-1.50	2.16	BC012596	Hs.323342	ARPC4	actin related protein 2/3 complex, subunit 4, 20kDa
0.66	-1.15	1.82	BC002632	Hs.386741	EMP2	epithelial membrane protein 2
0.66	0.02	0.64	BC001731	Hs.278672	C15orf15	chromosome 15 open reading frame 15
0.66	-0.06	0.73	BC001540	Hs.279832	CPSF6	cleavage and polyadenylation specific factor 6, 68kDa
0.66	0.40	0.26	BC002507	Hs.12142	WDR13	WD repeat domain 13
0.66	0.04	0.62	BC012205	Hs.107139	FLJ11149	hypothetical protein FLJ11149
0.66	-0.71	1.37	BC010545	Hs.60300	GNAS	GNAS complex locus
0.66	0.32	0.34	BC008226	Hs.336898	PPP2CB	protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform
0.66	0.31	0.35	BC011057	Hs.24835	NIPSNAP1	nipsnap homolog 1 (C. elegans)
0.66	0.29	0.37	BC015891	Hs.279704	SSX2	
0.66	-0.06	0.72	BC008727	In multiple clusters	RARA	
0.66	0.34	0.31	BC010881	Hs.84045	FLJ13154	hypothetical protein FLJ13154
0.66	-1.75	2.40	BC012149	Hs.426460	DKFZP564C186	DKFZP564C186 protein
0.66	-0.53	1.18	BC014414	Hs.187378	PAI-RBP1	PAI-1 mRNA-binding protein
0.66	0.20	0.45	BC000482	Hs.26002	LDB1	LIM domain binding 1
0.65	-0.31	0.96	BC004144	Hs.9850	MGC1842	hypothetical protein MGC1842
0.65	0.26	0.39	BC008868	Hs.173611	CSTB	cystatin B (stefin B)
0.65	-1.67	2.33	BC006510	Hs.23960	GRWD	glutamate rich WD repeat protein GRWD
0.65	0.09	0.56	BC008929	Hs.78305	CTSH	cathepsin H
0.65	-0.14	0.79	BC008770	Hs.6527	HSPC216	hypothetical protein HSPC216
0.65	0.22	0.43	BC000413	Hs.192023	STK3	serine/threonine kinase 3 (STE20 homolog, yeast)
0.65	-0.18	0.84	BC001043	Hs.15075	DKFZP434E2216	splicing factor 4
0.65	-0.42	1.07	BC007070	Hs.410016	EVIN1	epidermodysplasia verruciformis 1
0.65	0.30	0.35	BC001450	Hs.89563	NCBP1	nuclear cap binding protein subunit 1, 80kDa
0.65	-1.59	2.25	BC011819	Hs.380774	KIAA1327	KIAA1327 protein
0.65	0.43	0.22	BC000824	Hs.1384	MGMT	O-6-methylguanine-DNA methyltransferase
0.65	0.53	0.12	BC016513	Hs.26204	RPS27L	ribosomal protein S27-like
0.65	0.41	0.24	BC000327	Hs.6780	FLJ14154	hypothetical protein FLJ14154
0.65	0.40	0.25	BC000045	Hs.9030	IFIT1	interferon-induced protein with tetratricopeptide repeats 1
0.65	-0.10	0.74	BC005252	Hs.55498	MAL2	mal, T-cell differentiation protein 2
0.65	0.21	0.44	BC019270	Hs.260024	AICDA	activation-induced cytidine deaminase
0.65	0.23	0.42	BC007565	Hs.75648	STMN1	stathmin 1/oncoprotein 18
0.65	0.63	0.02	BC008900	Hs.288679	C20orf60	chromosome 20 open reading frame 60
0.65	0.15	0.49	BC017094	Hs.130829	NCF1	neutrophil cytosolic factor 1 (47kDa, chronic granulomatous disease, autosomal 1)

0.65	0.31	0.33	BC007219	Hs.180638	MGC22825	hypothetical protein MGC22825
0.65	-0.18	0.83	BC011906	Hs.9908	DJ465N24.2.1	hypothetical protein DJ465N24.2.1
0.64	0.10	0.54	BC008019	Hs.1708	TRIM16	tripartite motif-containing 16
0.64	-0.95	1.59	BC001101	Hs.119503	L3MBTL2	l(3)mbt-like 2 (Drosophila)
0.64	-0.01	0.65	BC001021	Hs.94576	MGC11256	hypothetical protein MGC11256
0.64	0.07	0.58	BC012189	Hs.324127		
0.64	0.14	0.50	BC006175	Hs.78465	GPR30	G protein-coupled receptor 30
0.64	-0.64	1.28	BC000193	Hs.22199	FLJ12687	hypothetical protein FLJ12687
0.64	-0.35	0.99	BC012088	Hs.89657	ANXA5	annexin A5
0.64	0.59	0.05	BC007866	Hs.122655	DKFZP586D0824	immunity associated protein 2
0.64	0.35	0.29	BC000124	Hs.283844	FLJ10535	pyridoxine-5'-phosphate oxidase
0.64	0.33	0.31	BC000185	Hs.259785	Di-Ras2	DIRAS family, GTP-binding RAS-like 2
0.64	0.36	0.28	BC016836	Hs.8185	MMS19L	MMS19-like (MET18 homolog, S. cerevisiae)
0.64	0.41	0.23	BC007037	Hs.82401	MGC4677	hypothetical protein MGC4677
0.64	-0.66	1.30	BC003170	Hs.8127	GNAI2	guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 2
0.64	-0.23	0.86	BC014653	Hs.286	PLK	polo-like kinase (Drosophila)
0.64	0.18	0.46	BC022091	Hs.44070	MGAT2	mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
0.64	-1.80	2.43	BC001139	Hs.380413	FLJ21939	hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2
0.64	-0.94	1.57	BC005966	Hs.173048	SNX5	sorting nexin 5
0.64	-0.48	1.12	BC006337	Hs.119591	KIAA0116	KIAA0116 protein
0.64	-0.05	0.68	BC013907	Data not found	RBSK	ribokinase
0.63	-0.95	1.58	BC000121	Hs.72132	YAP	YY1 associated protein
0.63	0.28	0.36	NM_152561	Hs.11607	LILRA2	leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2
0.63	-0.36	0.99	BC001889	Hs.7181	KIAA0370	exportin 6
0.63	-0.13	0.76	BF343213	Hs.350552	FLJ22875	hypothetical protein FLJ22875
0.63	0.17	0.46	BC010030	Hs.303666	LOC51193	zinc finger protein ANC_2H01
0.63	-0.36	0.99	BC000128	Hs.15725	IER5	immediate early response 5
0.63	0.15	0.48	BC014327	Hs.12150	MGC14353	hypothetical protein MGC14353
0.63	0.52	0.11	BC010492	Hs.7019	N4WBP5	Nedd4 family interacting protein 1
0.63	0.35	0.29	BC011636	Hs.25127	NA 4-14-03 80	
0.63	0.19	0.44	BC004226	Hs.6454	MRPL40	mitochondrial ribosomal protein L40
0.63	-1.52	2.15	BC014494	Hs.352252	ARHGEF5	Rho guanine nucleotide exchange factor (GEF) 5
0.63	-0.54	1.17	BC005951	Hs.153261	SEC61B	protein translocation complex beta
0.63	-0.15	0.78	BC008039	Hs.5464	MGC861	hypothetical protein MGC861
0.63	0.54	0.09	BC009446	Hs.6166	TH1L	TH1-like (Drosophila)
0.63	0.15	0.48	BC008383	Hs.181289	ATF4	activating transcription factor 4 (tax-responsive enhancer element B67)
0.63	0.59	0.04	BC014026	Hs.180455	CDK5	cyclin-dependent kinase 5
0.63	0.28	0.35	BC010041	Hs.12084	FLJ22955	hypothetical protein FLJ22955
0.63	0.50	0.13	BC013905	Data not found	TTL1	tubulin tyrosine ligase-like 1
0.63	-1.15	1.78	BC002412	Hs.78504	IMMT	inner membrane protein, mitochondrial (mitofilin)
0.63	-0.71	1.33	BC001388	Hs.217493	ANXA2	annexin A2
0.63	-1.42	2.05	BC009489	Hs.65377	MTF1	metal-regulatory transcription factor 1
0.62	0.54	0.08	BC012887	Hs.279905	KRTHA3B	keratin, hair, acidic, 3B
0.62	-0.78	1.40	BC002326	Hs.324178	MAPKAP1	mitogen-activated protein kinase associated protein 1
0.62	0.00	0.62	BC006249	Hs.3764	COVA1	cytosolic ovarian carcinoma antigen 1
0.62	0.35	0.27	BC009190	Hs.333823	PLSCR3	phospholipid scramblase 3
0.62	0.06	0.57	BC004875	Hs.66309	HMP19	HMP19 protein
0.62	0.52	0.11	BC004179	Hs.326560	MGC2780	hypothetical protein MGC2780
0.62	0.19	0.43	BC000881	Hs.1594	CENPA	centromere protein A, 17kDa
0.62	-0.79	1.41	BC006108	Hs.55014	RAN	RAN, member RAS oncogene family
0.62	0.54	0.08	BC007953	Hs.332381	MGC12518	similar to ZINC FINGER PROTEIN 257 (BONE MARROW ZINC FINGER 4)
(BMZF-4)						
0.62	-0.47	1.09	BC008325	Hs.379091	FLJ22002	hypothetical protein FLJ22002

0.62	0.06	0.56	BC004471	Hs.74120	MMP2	matrix metalloproteinase 2 (gelatinase A, 72kDa gelatinase, 72kDa type IV collagenase)
0.62	0.25	0.37	BC007348	Hs.77573	DKFZp434N0650	hypothetical protein DKFZp434N0650
0.62	0.44	0.18	BC004292	Hs.9729	NXF2	nuclear RNA export factor 2
0.62	-0.60	1.22	BC002637	Hs.155418	PLAU	plasminogen activator, urokinase
0.62	-0.40	1.02	BC000427	Hs.84285	FLJ20516	hypothetical protein FLJ20516
0.62	-0.14	0.76	BC015547	Hs.9597	PAIP1	polyadenylate binding protein-interacting protein 1
0.62	-0.78	1.40	BC008069	Hs.281648		
0.62	0.58	0.04	BC016727	Hs.82829	VPS4A	vacuolar protein sorting 4A (yeast)
0.62	-1.78	2.40	BC008348	Hs.42287	MGC23280	hypothetical protein MGC23280
0.62	-0.32	0.93	BC014908	Hs.75257	LOC51128	GTP-binding protein Sara
0.62	-0.43	1.04	BC014021	Hs.91813	VAMP8	vesicle-associated membrane protein 8 (endobrevin)
0.61	-0.60	1.21	BC011764	Hs.301266	CYB5	cytochrome b-5
0.61	-1.62	2.23	BC005322	Hs.433989	FLJ21080	hypothetical protein FLJ21080
0.61	-0.90	1.52	BC011359	Hs.418650	SDF2	stromal cell-derived factor 2
0.61	0.40	0.21	BC015797	Hs.285829	S100A2	S100 calcium binding protein A2
0.61	0.45	0.16	BC008741	Hs.154103	RCV1	recoverin
0.61	-0.48	1.09	BC002939	Hs.93121	TUBG1	tubulin, gamma 1
0.61	0.23	0.39	BC016851	Hs.375023	NA 4-14-03 120	
0.61	-2.46	3.08	BC016054	Hs.23467	C20orf158	chromosome 20 open reading frame 158
0.61	-0.90	1.51	BC007297	Hs.10824	CHEK1	CHK1 checkpoint homolog (S. pombe)
0.61	0.08	0.53	BC007391	Hs.381200	ORF1-FL49	putative nuclear protein ORF1-FL49
0.61	-0.07	0.67	BC015616	Hs.289271	JUNB	jun B proto-oncogene
0.61	-0.82	1.42	BC000844	Hs.76095	IER3	immediate early response 3
0.61	-0.33	0.94	BC002823	Hs.7446	NCKAP1	NCK-associated protein 1
0.61	-1.03	1.64	BC007641	Hs.17126	NA 4-14-03 22	
0.61	-1.30	1.91	BC012560	Hs.380887	MRPS10	mitochondrial ribosomal protein S10
0.61	-1.23	1.84	BC011537	Data not found	KPNA1	karyopherin alpha 1 (importin alpha 5)
0.61	-0.93	1.54	BC010088	Hs.89545	NA 4-14-03 89	
0.60	0.49	0.12	BC006309	Hs.78768	F23149_1	hypothetical protein F23149_1
0.60	-0.04	0.65	BC013047	Hs.322456	FKBP14	FK506 binding protein 14, 22 kDa
0.60	0.12	0.49	BC017572	Hs.21201	EIF2C2	eukaryotic translation initiation factor 2C, 2
0.60	-1.96	2.56	BC001507	Hs.334360	FLJ21827	hypothetical protein FLJ21827
0.60	-1.46	2.06	BC003525	Hs.42712	WSB2	likely ortholog of mouse WD-40-repeat-containing protein with a SOCS box 2
0.60	0.28	0.32	BC000627	Hs.321677	SLC25A19	solute carrier family 25 (mitochondrial deoxynucleotide carrier), member 19
0.60	-0.62	1.22	BC016984	Hs.11065	ZD52F10	hypothetical gene ZD52F10
0.60	-0.42	1.02	BC015921	Hs.396266	IGHM	immunoglobulin heavy constant mu
0.60	0.50	0.10	BC009842	Hs.323664	MGC29643	hypothetical protein MGC29643
0.60	-0.81	1.41	BC000619	Hs.21635	MCLC	Mid-1-related chloride channel 1
0.60	-0.81	1.41	BC006234	Hs.7788	NPD007	NPD007 protein
0.60	0.09	0.51	BC014431	In multiple clusters	FLJ11151	hypothetical protein FLJ11151
0.60	-0.30	0.90	BC004548	In multiple clusters	GRINL1A	glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A
0.60	-1.21	1.81	BC001782	Hs.322852	EFG2	mitochondrial elongation factor G2
0.60	0.11	0.49	BC015619	Hs.126783	DKFZP434A0131	DKFZp434A0131 protein
0.60	-1.17	1.77	BC009804	Hs.7179	RAD1	RAD1 homolog (S. pombe)
0.60	-0.47	1.07	AK096655	Hs.6983	ATP6V0C	ATPase, H+ transporting, lysosomal 16kDa, V0 subunit c
0.60	-0.08	0.68	BC011403	Hs.24969	PRKAB1	protein kinase, AMP-activated, beta 1 non-catalytic subunit
0.60	0.19	0.41	BC009427	Hs.380843	PRO2013	hypothetical protein PRO2013
0.60	-0.08	0.68	BC012340	Hs.24792	ARL4	ADP-ribosylation factor-like 4
0.60	-0.59	1.19	BC011952	Hs.406737	APOA1	apolipoprotein A-I
0.60	0.01	0.58	BC009074	Hs.118821	RGNEF	hypothetical protein FLJ21817 similar to Rhoip2
0.60	-0.36	0.96	BC004505	Hs.18627	BTBD2	BTB (POZ) domain containing 2
0.60	0.23	0.37	BC003663	Hs.9877	NUP62	nucleoporin 62kDa
0.60	-0.86	1.45	BC009468	Hs.84153	MGC2306	hypothetical protein MGC2306

0.59	0.01	0.59	BC018896	Hs.80598	MGC13057	hypothetical protein MGC13057
0.59	0.34	0.26	BC008316	Hs.10824	SEP15	15 kDa selenoprotein
0.59	0.32	0.27	BC000898	Hs.79026	MLF2	myeloid leukemia factor 2
0.59	0.45	0.14	BC008921	Data not found	SEMA3B	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin)
3B						
0.59	-0.97	1.57	BC012087	Hs.6682	MGC4400	hypothetical protein MGC4400
0.59	0.06	0.53	BC008487	Hs.99960	SLC7A7	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7
0.59	0.53	0.07	BC014580	Hs.12102		
0.59	0.30	0.30	BC000374	Hs.409634	LOC51064	glutathione S-transferase subunit 13 homolog
0.59	0.05	0.54	BC003565	Data not found	ASB13	ankyrin repeat and SOCS box-containing 13
0.59	-0.41	1.01	BC017054	Hs.154672	HSPC155	hypothetical protein HSPC155
0.59	0.43	0.16	BC007830	Hs.4877	ARSA	arylsulfatase A
0.59	0.19	0.40	BC001308	Hs.279905	ANKT	nucleolar protein ANKT
0.59	-0.32	0.91	BC000250	Hs.81687	PPT1	palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile)
0.59	0.04	0.55	BC010435	Data not found	LOC89941	mitochondrial Rho 2
0.59	-0.19	0.78	BC021195	Hs.73893	MGC13098	hypothetical protein MGC13098
0.59	-1.00	1.59	BC012728	Data not found	FLJ12770	hypothetical protein FLJ12770
0.59	0.18	0.41	BC002716	Hs.5472	HDAC11	histone deacetylase 11
0.59	-0.37	0.96	BC015052	Hs.83419	H2BFB	histone 1, H2bd
0.59	-0.62	1.21	BC005870	Hs.334626	MGC4238	hypothetical protein MGC4238
0.59	-0.40	0.98	BC012187	Hs.192375	TERA	TERA protein
0.59	0.33	0.26	BC013432	Hs.7535	LOC55871	COBW-like protein
0.59	-0.05	0.64	BC012265	Hs.180370	EIF1AY	eukaryotic translation initiation factor 1A, Y chromosome
0.59	0.34	0.25	BC001822	Hs.29656	PITPNB	phosphatidylinositol transfer protein, beta
0.59	-0.29	0.88	BC020507	Hs.71746	NA 4-14-03 140	
0.59	-0.15	0.73	BC007964	Hs.66052	PPIA	peptidylprolyl isomerase A (cyclophilin A)
0.58	-0.42	1.01	BC011634	Hs.113207	JUN	v-jun sarcoma virus 17 oncogene homolog (avian)
0.58	0.50	0.08	BC004462	Hs.180780	LOC112885	hypothetical protein BC012187
0.58	-0.36	0.94	BC007606	Hs.1578	MANBAL	mannosidase, beta A, lysosomal-like
0.58	0.04	0.54	BC013374	Data not found	IGLJ3	immunoglobulin lambda joining 3
0.58	0.25	0.33	BC013956	Data not found	RRM2	ribonucleotide reductase M2 polypeptide
0.58	-0.96	1.54	BC015480	Hs.343748	PREB	prolactin regulatory element binding
0.58	-0.80	1.39	BC001416	Hs.2903	FLJ11560	hypothetical protein FLJ11560
0.58	-0.88	1.46	BC002668	Hs.15250	PRKCSH	protein kinase C substrate 80K-H
0.58	-0.99	1.57	BC006401	Hs.58831	C20orf45	chromosome 20 open reading frame 45
0.58	0.52	0.06	BC004269	Hs.18449	SRP	stresscopin-related peptide
0.58	-0.03	0.61	BC006390	Hs.172195	SFXN2	sideroflexin 2
0.58	-1.11	1.68	BC011562	Hs.77617	ATP6V1E1	ATPase, H+ transporting, lysosomal 31kDa, V1 subunit E isoform 1
0.58	-0.05	0.62	BC004134	Hs.333257	AVIL	advillin
0.58	0.57	0.01	BC020548	Hs.101766	NMB	neuromedin B
0.58	-0.13	0.70	BC000638	Data not found	NRLN1	likely ortholog of mouse neuralin 1
0.58	-0.69	1.27	BC007803	Hs.273357	KEAP1	Kelch-like ECH-associated protein 1
0.58	-0.40	0.97	BC015096	Hs.12622	CALM2	calmodulin 2 (phosphorylase kinase, delta)
0.57	-0.99	1.56	BC005362	Hs.194695	MGC10765	hypothetical protein MGC10765
0.57	-1.68	2.26	BC017314	Hs.18063	RPLP0	ribosomal protein, large, P0
0.57	-5.39	5.97	BC007100	Hs.406720	ATF5	activating transcription factor 5
0.57	0.46	0.12	BC000380	Hs.110757	TIMM13	translocase of inner mitochondrial membrane 13 homolog (yeast)
0.57	-2.52	3.09	BC015924	Hs.37482	THG-1	TSC-22-like
0.57	-1.19	1.76	BC006280	Hs.176067	MGC11386	hypothetical protein MGC11386
0.57	-0.21	0.78	BC012161	Hs.388456	PSMD2	proteasome (prosome, macropain) 26S subunit, non-ATPase, 2
0.57	0.37	0.20	BC021257	Hs.79411	FLJ22501	Hermansky-Pudlak syndrome-6 protein
0.57	-1.22	1.79	BC013082	Hs.20191	FLJ12886	hypothetical protein FLJ12886
0.57	0.46	0.11	BC001706	Hs.109773	ACO2	aconitase 2, mitochondrial

0.57	-0.06	0.63	BC012455	Hs.62275	C15orf15	chromosome 15 open reading frame 15
0.57	0.48	0.08	BC011671	Hs.155090	PAX6	paired box gene 6 (aniridia, keratitis)
0.57	0.21	0.36	BC008466	Hs.5085	R3HDM	R3H domain (binds single-stranded nucleic acids) containing
0.57	-1.33	1.90	BC009882	Hs.20084	IRAK1	interleukin-1 receptor-associated kinase 1
0.57	-1.00	1.57	BC011741	Hs.74376	RNASE4	ribonuclease, RNase A family, 4
0.57	-0.39	0.95	BC000464	Hs.21753	VPS45A	vacuolar protein sorting 45A (yeast)
0.57	0.00	0.57	BC019087	Hs.267566	MGC16703	alpha tubulin-like
0.57	-0.58	1.15	BC014766	Hs.7771	RPA40	RNA polymerase I subunit
0.57	0.08	0.48	BC009480	Hs.9950	ECH1	enoyl Coenzyme A hydratase 1, peroxisomal
0.57	-0.36	0.92	BC009355	Hs.6294	S100A3	S100 calcium binding protein A3
0.57	0.44	0.13	BC012166	Hs.283667	MGC4399	mitochondrial carrier protein
0.56	0.35	0.21	BC002560	Hs.157237	DNMT3L	DNA (cytosine-5-)-methyltransferase 3-like
0.56	-0.06	0.62	BC005918	Hs.425790	VILL	villin-like
0.56	0.51	0.05	BC007615	Hs.298262	ST7	suppression of tumorigenicity
0.56	-0.16	0.72	BC016699	Hs.134846	ARMET	arginine-rich, mutated in early stage tumors
0.56	-0.36	0.92	BC010385	Hs.966	COIL	coilin
0.56	0.33	0.23	BC000680	Hs.266935	CBFA2T1	core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related
0.56	-0.28	0.85	BC002440	Hs.218842	CCNB1	cyclin B1
0.56	-0.11	0.67	BC009903	Hs.115232	SARS	seryl-tRNA synthetase
0.56	0.48	0.08	BC008011	Hs.111779	PGRMC2	progesterone receptor membrane component 2
0.56	0.24	0.32	BC007872	Hs.105097	FLJ14466	hypothetical protein FLJ14466
0.56	-0.36	0.92	BC014654	Hs.195464	GPAA1	GPAA1P anchor attachment protein 1 homolog (yeast)
0.56	-0.42	0.98	BC015650	Hs.132760	MPST	mercaptopyruvate sulfurtransferase
0.56	0.51	0.05	BC011392	Hs.3833	USP11	ubiquitin specific protease 11
0.56	-0.17	0.73	BC003508	Hs.248267	EIF4A2	eukaryotic translation initiation factor 4A, isoform 2
0.56	-0.07	0.63	BC000722	Hs.118721	FLJ20150	hypothetical protein FLJ20150
0.56	-0.64	1.20	BC010518	Data not found	MACF1	microtubule-actin crosslinking factor 1
0.56	0.09	0.47	BC012812	Hs.235390	FLJ12895	hypothetical protein FLJ12895
0.56	0.09	0.47	BC013568	Hs.26994	H11	protein kinase H11
0.56	0.14	0.41	BC001668	Hs.31439	SPINT2	serine protease inhibitor, Kunitz type, 2
0.56	-1.74	2.29	BC015473	Data not found	MRE11A	MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
0.55	0.45	0.10	BC009681	Hs.315463	TOMM70A	translocase of outer mitochondrial membrane 70 homolog A (yeast)
0.55	0.22	0.34	BC001013	Hs.102696	MGC10715	novel C2H2 type zinc finger protein
0.55	0.06	0.49	BC010018	Hs.172673	SB52	glucosamine-6-phosphate isomerase
0.55	0.10	0.45	BC010696	Hs.32978	MRPL12	mitochondrial ribosomal protein L12
0.55	-0.26	0.81	BC007510	Hs.119503	EIF3S61P	eukaryotic translation initiation factor 3, subunit 6 interacting protein
0.55	-1.28	1.83	AK096306	Hs.300383	SSNA1	Sjogren's syndrome nuclear autoantigen 1
0.55	-0.60	1.15	BC009708	Hs.409634		
0.55	-0.37	0.93	BC001095	Hs.118244	PRPF4	PRP4 pre-mRNA processing factor 4 homolog (yeast)
0.55	0.04	0.51	BC017367	Hs.91566	NDP52	nuclear domain 10 protein
0.55	-0.03	0.58	BC014980	Hs.129548	WAS	Wiskott-Aldrich syndrome (eczema-thrombocytopenia)
0.55	-0.78	1.33	BC013916	Hs.71618	E2-230K	likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K
0.55	-0.74	1.30	BC011988	Hs.1244	ZBPB	zona pellucida binding protein
0.55	-0.25	0.80	BC009780	Hs.195614	SF3B3	splicing factor 3b, subunit 3, 130kDa
0.55	-0.17	0.73	BC007817	Hs.159309	HADHB	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit
0.55	-0.51	1.06	BC003643	Hs.225160	AP4B1	adaptor-related protein complex 4, beta 1 subunit
0.55	-1.46	2.01	BC018163	Hs.348024	P2RY6	pyrimidineric receptor P2Y, G-protein coupled, 6
0.55	0.52	0.03	BC003102	Hs.19561	NA 4-14-03 490	
0.55	-0.83	1.38	BC021989	Hs.79428	KRT19	keratin 19
0.55	-0.08	0.63	BC012531	Hs.227933	ALG5	Alg5, S. cerevisiae, homolog of
0.55	0.19	0.36	BC017081	Hs.351484	AWP1	protein associated with PRK1
0.55	0.50	0.05	BC014039	Hs.184339	WBCSR5	Williams-Beuren syndrome chromosome region 5

0.55	0.22	0.33	BC003033	Hs.301342	RAD51L3	RAD51-like 3 (S. cerevisiae)
0.55	-0.82	1.37	BC008305	Hs.289080	NA 4-14-03 100	
0.55	-1.31	1.86	BC008966	Hs.173739	GNRPX	likely ortholog of mouse guanine nucleotide releasing protein x
0.55	0.10	0.45	BC004386	Hs.79768	PIGT	phosphatidyl inositol glycan class T
0.55	-0.09	0.64	BC000978	Data not found	FLJ20259	
0.55	0.54	0.01	BC001950	Hs.7101	ANAPC5	anaphase promoting complex subunit 5
0.55	0.38	0.16	BC008687	Hs.248149	DKFZP434D146	DKFZP434D146 protein
0.55	0.02	0.52	BC000521	Hs.118400	DKFZP434H132	DKFZP434H132 protein
0.54	0.00	0.54	BC007024	Hs.201967	LOC153516	hypothetical protein BC010563
0.54	-1.12	1.66	BC021241	Hs.32952	RAGD	Rag D protein
0.54	-1.75	2.29	BC008298	Hs.183253	PHEMX	pan-hematopoietic expression
0.54	-1.21	1.76	BC003046	Hs.63304	NA 4-14-03 489	
0.54	-0.61	1.15	BC016155	In multiple clusters	MGC11303	hypothetical protein MGC11303 similar to Zink transporter 2
0.54	0.49	0.06	BC002902	Hs.90297	DKFZP586O0120	DKFZP586O0120 protein
0.54	-2.68	3.23	BC016737	Hs.248267	MYBL2	v-myb myeloblastosis viral oncogene homolog (avian)-like 2
0.54	-0.31	0.85	NM_153611	Hs.22546	MAGEA4	melanoma antigen, family A, 4
0.54	-0.39	0.93	BC020523	Hs.48604	LOC85026	AD038
0.54	0.00	0.54	BC001936	Hs.41714	SLC25A3	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3
0.54	0.18	0.36	BC009841	Hs.182215	FLJ23322	hypothetical protein FLJ23322
0.54	-0.11	0.65	BC021965	Hs.8526	MGC3262	hypothetical protein MGC3262
0.54	0.03	0.51	BC011564	Hs.268490	CDKN1A	cyclin-dependent kinase inhibitor 1A (p21, Cip1)
0.54	0.12	0.42	BC002894	Hs.28029	MGC3062	likely ortholog of mouse phosducin-like 2
0.54	0.47	0.07	BC007720	Hs.121482	HSPCB	heat shock 90kDa protein 1, beta
0.54	-0.26	0.80	BC010636	Hs.79194	LAPTM4A	lysosomal-associated protein transmembrane 4 alpha
0.54	0.25	0.29	BC002485	Hs.4992	MGC2848	hypothetical protein MGC2848
0.54	-0.58	1.12	BC012117	Hs.23557	HTR3A	5-hydroxytryptamine (serotonin) receptor 3A
0.54	0.09	0.44	BC000314	Hs.99947	RTN1	reticulon 1
0.54	0.02	0.52	BC007026	Hs.1955	CENPH	centromere protein H
0.54	-0.62	1.16	BC009714	Hs.24970	NA 4-14-03 306	
0.54	-0.70	1.24	BC005934	Hs.288106	ASMT	acetylserotonin O-methyltransferase
0.54	0.13	0.41	BC009998	Hs.283722	IDH1	isocitrate dehydrogenase 1 (NADP+), soluble
0.54	-2.79	3.32	BC000343	Hs.211594	FLJ12428	hypothetical protein FLJ12428
0.54	-0.17	0.70	BC007305	Hs.83190	FLJ20130	hypothetical protein FLJ20130
0.54	-0.02	0.56	BC010659	Hs.15830	MGC4701	hypothetical protein MGC4701
0.54	0.34	0.19	BM906317	Hs.55336	FLJ14346	hypothetical protein FLJ14346
0.53	-0.59	1.12	BC008918	Hs.89474	UCC1	upregulated in colorectal cancer gene 1
0.53	0.43	0.11	BC009228	Data not found	MGC27171	hypothetical protein MGC27171
0.53	0.49	0.04	BC010632	Hs.54617	BAT1	HLA-B associated transcript 1
0.53	0.20	0.33	BC010892	Hs.279836	HSPC166	HSPC166 protein
0.53	0.29	0.24	BC000112	Hs.179817	VPS18	vacuolar protein sorting protein 18
0.53	-2.06	2.59	BC005907	Data not found	NAGA	N-acetylgalactosaminidase, alpha-
0.53	0.15	0.38	BC009229	Hs.9100	SERPINB4	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 4
0.53	-0.30	0.83	BC012362	Hs.239527	NA 4-14-03 28	
0.53	-0.86	1.39	BC011905	Hs.17936	RFP2	ret finger protein 2
0.53	0.12	0.41	BC017247	Hs.118962	TCOF1	Treacher Collins-Franceschetti syndrome 1
0.53	0.47	0.06	BC007080	Hs.273077	MGC2477	hypothetical protein MGC2477
0.53	0.27	0.26	BC016856	Hs.166173	EEF1E1	eukaryotic translation elongation factor 1 epsilon 1
0.53	-1.62	2.15	BC007667	Hs.63489	GAS7	growth arrest-specific 7
0.53	-0.04	0.57	BC015946	Hs.418650	IGFBP6	insulin-like growth factor binding protein 6
0.53	0.37	0.16	BC006238	Hs.37936	SUV39H1	suppressor of variegation 3-9 homolog 1 (Drosophila)
0.53	0.02	0.51	BC000688	Hs.300631	TNRC5	trinucleotide repeat containing 5
0.53	0.02	0.51	BC000368	Hs.392837	LOC51290	CDA14
0.53	0.17	0.36	BC002662	Hs.22777	LOC219738	hypothetical protein BC013587



0.53	-0.01	0.54	BC001712	Hs.182625	DPP3	dipeptidylpeptidase 3
0.53	-0.24	0.77	BC004536	Hs.333120	CDV-1	carnitine deficiency-associated gene expressed in ventricle 1
0.53	-0.41	0.94	BC013899	Hs.153640	NA 4-14-03 555	
0.53	-0.09	0.61	BC001274	Hs.59498	CDC2L5	cell division cycle 2-like 5 (cholinesterase-related cell division controller)
0.53	0.42	0.11	BC017486	Hs.89545	BHD	folliculin
0.53	-0.01	0.54	BC004187	Hs.47649	MCCC1	methylcrotonoyl-Coenzyme A carboxylase 1 (alpha)
0.53	0.00	0.53	BC015239	Hs.129837	RNASEH2A	ribonuclease H2, large subunit
0.53	0.50	0.03	BC006508	Hs.184877	SLC25A11	solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11
0.53	-2.51	3.04	BC009867	Hs.406511	ETS1	v-ets erythroblastosis virus E26 oncogene homolog 1 (avian)
0.53	0.20	0.32	BC008656	Hs.177011	DNAJC9	DnaJ (Hsp40) homolog, subfamily C, member 9
0.52	-0.02	0.55	BC003684	Hs.79187	CXADR	coxsackie virus and adenovirus receptor
0.52	-0.91	1.44	BC014252	Hs.124010	DKFZp586C1924	hypothetical protein DKFZp586C1924
0.52	-1.20	1.72	BC013682	Hs.37034	TNFRSF21	tumor necrosis factor receptor superfamily, member 21
0.52	-0.65	1.17	BC014438	Hs.172035	FLJ14431	hypothetical protein FLJ14431
0.52	-0.07	0.59	BC015329	Hs.169330	FLJ11016	hypothetical protein FLJ11016
0.52	-0.20	0.72	BC017558	Hs.180877	SLC25A3	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3
0.52	-1.08	1.60	BC013687	Hs.421654	UQCRB	ubiquinol-cytochrome c reductase binding protein
0.52	-1.02	1.54	BC009607	Hs.169248	HCS	cytochrome c, somatic
0.52	-0.61	1.13	BC000062	Hs.183556	MDM2	Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)
0.52	-0.42	0.94	BC010389	Hs.351622	PDX1	E3-binding protein
0.52	0.21	0.31	BC003660	Data not found	NA 4-14-03 139	
0.52	0.28	0.24	BC009758	Hs.166254	PCDHGC3	protocadherin gamma subfamily C, 3
0.52	0.38	0.14	BC016040	Hs.120996	PAFAH1B3	platelet-activating factor acetylhydrolase, isoform Ib, gamma subunit 29kDa
0.52	0.37	0.15	BC002947	Hs.73769	RPL8	ribosomal protein L8
0.52	0.01	0.51	BC001420	Hs.109706	PTDSS1	phosphatidylserine synthase 1
0.52	-0.95	1.47	BC007049	Hs.78944	NA 4-14-03 148	
0.52	-0.50	1.02	BC009582	Hs.169248	HCS	cytochrome c, somatic
0.52	0.44	0.08	BC003698	Hs.64211	MGC5601	hypothetical protein MGC5601
0.52	-0.12	0.63	BC001848	Hs.70582	MGC2835	DEAD box helicase 97 kDa
0.52	-0.85	1.36	BC001501	Hs.82932	MGC11316	
0.52	-2.12	2.64	BC007458	Hs.288193	ZDHHC3	zinc finger, DHHC domain containing 3
0.52	0.13	0.39	BC006479	Hs.108847	MGC2749	hypothetical protein MGC2749
0.52	0.12	0.40	BC007797	Hs.21907	ELMO2	engulfment and cell motility 2 (ced-12 homolog, C. elegans)
0.52	0.17	0.34	BC006818	Hs.240534	AGPAT1	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)
0.52	-0.16	0.68	BC006425	Hs.337251	IL1RN	interleukin 1 receptor antagonist
0.51	-1.17	1.69	BC009894	Hs.274230	FLJ10634	hypothetical protein FLJ10634
0.51	-0.29	0.80	BC008404	Hs.811	COQ7	coenzyme Q7 homolog, ubiquinone (yeast)
0.51	-0.39	0.91	BC016866	Hs.91096	CRIPT	postsynaptic protein CRIPT
0.51	0.18	0.34	BC000840	Hs.155482	HAGH	hydroxyacyl glutathione hydrolase
0.51	-0.19	0.70	BC009808	Hs.277721	FLJ10493	hypothetical protein FLJ10493
0.51	-2.55	3.06	BC004913	Hs.9629	PDE9A	phosphodiesterase 9A
0.51	-0.07	0.58	BC004494	Hs.32018	PSMC2	proteasome (prosome, macropain) 26S subunit, ATPase, 2
0.51	-1.37	1.88	BC004324	Hs.397609	TYROBP	TYRO protein tyrosine kinase binding protein
0.51	0.02	0.49	BC000693	Hs.153961	WBSCR20A	Williams Beuren syndrome chromosome region 20A
0.51	-1.66	2.17	BC004468	Hs.40834	MGC3038	hypothetical protein similar to actin related protein 2/3 complex, subunit 5
0.51	0.24	0.27	BC003545	Hs.236510	MINA53	myc-induced nuclear antigen, 53 kDa
0.51	0.33	0.18	BC008783	Hs.247478	BAT4	HLA-B associated transcript 4
0.51	0.01	0.50	BC004565	Hs.247812	CD7	CD7 antigen (p41)
0.51	0.14	0.37	BC001660	Hs.8297	RNASE6PL	ribonuclease 6 precursor
0.51	0.28	0.23	BC013979	Hs.77719	PCCB	propionyl Coenzyme A carboxylase, beta polypeptide
0.51	-0.36	0.87	BC001915	Hs.5105	KRT18	keratin 18
0.51	-0.95	1.45	BC002765	Hs.279784	FLJ20445	hypothetical protein FLJ20445

0.51	-0.56	1.06	BC009873	Hs.334712	CGI-150	CGI-150 protein
0.51	-0.40	0.91	BC009015	Hs.183583	ILF2	interleukin enhancer binding factor 2, 45kDa
0.51	0.27	0.24	BC000117	Hs.105435	DAD1	defender against cell death 1
0.51	0.12	0.38	BC007628	Hs.182265	BNIP3	BCL2/adenovirus E1B 19kDa interacting protein 3
0.51	0.35	0.16	BC015536	Hs.118944	LOC51659	HSPC037 protein
0.51	-0.84	1.35	BC009524	Hs.178761	CCRK	cell cycle related kinase
0.51	0.20	0.31	BC003110	Hs.64310	C20orf7	chromosome 20 open reading frame 7
0.51	0.34	0.17	BC009031	Hs.155182	GOT1	glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)
0.51	-0.23	0.74	BC010145	Data not found	ZNF297	zinc finger protein 297
0.51	0.13	0.38	BC005134	Hs.235422	PEX10	peroxisome biogenesis factor 10
0.51	-1.04	1.55	BC010014	Hs.26516	PSMD10	proteasome (prosome, macropain) 26S subunit, non-ATPase, 10
0.51	0.49	0.02	BC008145	Hs.73722	NOL4	nucleolar protein 4
0.51	-0.26	0.77	BC020969	Data not found	NDUFA6	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6, 14kDa
0.51	-0.45	0.96	BC001946	Hs.41641	BRUNOL4	bruno-like 4, RNA binding protein (Drosophila)
0.51	-1.23	1.73	BC002850	Hs.379039	PTD013	PTD013 protein
0.50	0.09	0.41	BC012944	Hs.74579	ENC1	ectodermal-neural cortex (with BTB-like domain)
0.50	-1.88	2.39	BC009566	Hs.171723	FLJ12748	hypothetical protein FLJ12748
0.50	-1.05	1.55	BC004424	Hs.250745	P2RY2	purinergic receptor P2Y, G-protein coupled, 2
0.50	-0.52	1.03	BC014221	Hs.256583	NA 4-14-03 54	
0.50	-0.59	1.09	BC011234	Hs.79968	DKFZp434K1210	hypothetical protein DKFZp434K1210
0.50	0.27	0.23	BC017123	Hs.77039	RAB4A	RAB4A, member RAS oncogene family
0.50	-0.23	0.73	BC000552	Data not found	MGC11141	fucosyltransferase 10 (alpha (1,3) fucosyltransferase)
0.50	-0.19	0.70	BC007460	Hs.1852	HRIHFB2072	short coiled coil protein
0.50	0.49	0.01	BC004111	Hs.6980	ACY1	aminoacylase 1
0.50	-0.55	1.05	BC001028	Hs.30057	FLJ12949	hypothetical protein FLJ12949
0.50	-0.41	0.92	BC008937	Hs.250005	CBX8	chromobox homolog 8 (Pc class homolog, Drosophila)
0.50	-4.53	5.03	BC001744	Hs.74592	ADM	adrenomedullin
0.50	0.33	0.17	BC009529	Hs.77558	MGC4161	hypothetical protein MGC4161
0.50	-0.76	1.26	BC012992	Hs.241567	ING4	inhibitor of growth family, member 4
0.50	-0.22	0.71	BC002650	Hs.259599	SERPINE1	serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1
0.50	-0.09	0.59	BC004518	Hs.26971	MRPS30	mitochondrial ribosomal protein S30
0.50	0.03	0.47	BC017238	Hs.17283	MEOX2	mesenchyme homeo box 2 (growth arrest-specific homeo box)
0.50	0.38	0.11	BC001861	Hs.97978	P24B	integral type I protein
0.50	0.11	0.39	BC003560	Hs.406532	FZD7	frizzled homolog 7 (Drosophila)
0.50	-1.46	1.95	BC002836	Hs.1422	PRO1855	hypothetical protein PRO1855
0.50	0.07	0.43	BC011685	Hs.79353	LOC132158	similar to CG9886 gene product
0.50	-0.38	0.87	BC004301	Hs.285313	DNAJA2	DnaJ (Hsp40) homolog, subfamily A, member 2
0.50	-0.56	1.05	BC020981	Hs.75782	ZNF237	zinc finger protein 237
0.50	0.15	0.34	BC013402	Hs.152894	LENG4	leukocyte receptor cluster (LRC) member 4
0.49	-0.02	0.52	BC007339	Hs.272499	MCM10	MCM10 minichromosome maintenance deficient 10 (S. cerevisiae)
0.49	0.21	0.28	BC000261	Hs.724	UBE2D1	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)
0.49	-0.77	1.26	Data not found	FLJ20331	FLJ20331	hypothetical protein FLJ20331
0.49	-0.11	0.60	BC000375	Hs.2281	VIP32	hypothetical protein PP5395
0.49	0.41	0.08	BC004864	Hs.75206	STK16	serine/threonine kinase 16
0.49	-2.37	2.87	BC017472	Hs.24654	NRGN	neurogranin (protein kinase C substrate, RC3)
0.49	-0.29	0.78	BC003642	Hs.375620	ASM3A	acid sphingomyelinase-like phosphodiesterase
0.49	-0.37	0.87	BC014074	Hs.30743	CLPP	ClpP caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli)
0.49	0.15	0.34	BC015765	Hs.315103	ARHGDI5	Rho GDP dissociation inhibitor (GDI) beta
0.49	-0.48	0.97	BC011700	Hs.170171	TEAD4	TEA domain family member 4
0.49	0.28	0.22	BC009788	Hs.247324	MRPS14	mitochondrial ribosomal protein S14
0.49	0.31	0.18	BC007072	Hs.246759	FBXO26	F-box only protein 26
0.49	0.28	0.21	BC011550	Hs.182729	POLR2E	polymerase (RNA) II (DNA directed) polypeptide E, 25kDa

0.49	-0.26	0.75	BC013106	Hs.206097	FLJ11294	hypothetical protein FLJ11294
0.49	-0.12	0.61	BC006811	Hs.100724	PPARG	peroxisome proliferative activated receptor, gamma
0.49	0.31	0.18	BC016137	Hs.2471	LOC51255	hypothetical protein LOC51255
0.49	0.13	0.36	BC014051	Hs.333513	RECQL5	RecQ protein-like 5
0.49	-1.44	1.92	BC007013	Hs.152983	FKBPL	FK506 binding protein like
0.49	-1.13	1.62	BC000948	Hs.31082	FLJ10525	hypothetical protein FLJ10525
0.49	0.15	0.34	BC001648	Hs.325321	WDR18	WD repeat domain 18
0.49	-0.20	0.69	BC000007	Hs.279529	FLJ11126	hypothetical protein FLJ11126
0.49	-0.08	0.56	BC007734	Hs.68055	KPTN	kaptin (actin binding protein)
0.49	0.03	0.45	BC019324	Hs.317403	MGC2744	hypothetical protein MGC2744
0.49	-0.73	1.22	BC015708	Hs.121590	ZNF259	zinc finger protein 259
0.49	-0.07	0.56	BC015576	Hs.194657	HCA112	hepatocellular carcinoma-associated antigen 112
0.49	0.08	0.41	BC006498	Hs.2934	CGI-38	brain specific protein
0.49	-0.80	1.29	BC001818	Hs.7570	ATPIF1	ATPase inhibitory factor 1
0.49	-0.13	0.61	BC020546	Hs.68879	FLJ14627	hypothetical protein FLJ14627
0.49	0.39	0.10	BC000964	Hs.75485	OAT	ornithine aminotransferase (gyrate atrophy)
0.48	0.29	0.20	BC008936	Hs.254099	PFN1	profilin 1
0.48	-1.74	2.23	BC017272	Hs.76297	FLJ21963	hypothetical protein FLJ21963
0.48	0.38	0.11	BC000138	Hs.79024	HNRPM	heterogeneous nuclear ribonucleoprotein M
0.48	-1.72	2.20	BC014474	Hs.166160	MGC15419	MGC15419 protein
0.48	-0.83	1.31	BC014392	Hs.433503	DUSP12	dual specificity phosphatase 12
0.48	-0.01	0.49	BC010668	Hs.380094	UBCE7IP5	likely ortholog of mouse ubiquitin conjugating enzyme 7 interacting protein 5
0.48	-0.53	1.01	BC004960	Hs.326739	SILV	silver homolog (mouse)
0.48	0.08	0.41	BC012386	Hs.8179	NFKBIE	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon
0.48	0.11	0.37	BC015739	Hs.90708	SLC4A2	solute carrier family 4, anion exchanger, member 2 (erythrocyte membrane protein band 3-like 1)
0.48	0.46	0.03	BC003084	Hs.173739	GNRPX	likely ortholog of mouse guanine nucleotide releasing protein x
0.48	-0.57	1.05	BC017029	Hs.356463	FLJ11218	hypothetical protein FLJ11218
0.48	0.35	0.13	BC014187	Hs.121830	EIF4G1	eukaryotic translation initiation factor 4 gamma, 1
0.48	0.06	0.42	BC011689	Hs.15921	MKPX	mitogen-activated protein kinase phosphatase x
0.48	0.45	0.03	BC001219	Hs.20768	DKFZP434B0335	DKFZP434B0335 protein
0.48	0.22	0.26	BC010016	Hs.313342	C3orf4	chromosome 3 open reading frame 4
0.48	-0.95	1.43	BC012542	Hs.36232	KIAA0186	KIAA0186 gene product
0.48	-0.14	0.62	BC021085	Hs.878	ZDHHC9	zinc finger, DHHC domain containing 9
0.48	0.29	0.19	BC017059	Hs.155530	FUSIP1	FUS interacting protein (serine-arginine rich) 1
0.48	-0.54	1.02	BC000921	Hs.273063	HSPC133	HSPC133 protein
0.48	0.16	0.32	BC011748	Hs.25292	MGC17919	hypothetical protein MGC17919
0.48	-0.64	1.12	BC012389	Hs.121068	NA 4-14-03 401	
0.48	-0.10	0.57	BC012801	Hs.20621	NA 4-14-03 127	
0.48	-0.92	1.40	BC013724	Hs.418650	SPP1	secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1)
0.48	-0.16	0.64	BC002993	Hs.272822	NVL	nuclear VCP-like
0.48	-1.37	1.84	BC008330	Data not found	HCLA-ISO	cyclin L2
0.48	0.07	0.40	BC005936	Hs.44163	LMOD1	leiomodrin 1 (smooth muscle)
0.47	-0.25	0.73	BC001604	Hs.132834	NA 4-14-03 493	
0.47	0.15	0.32	BC019016	Hs.9070	COP55	COP9 constitutive photomorphogenic homolog subunit 5 (Arabidopsis)
0.47	0.24	0.24	BC012761	Hs.9625	NR1	NADPH-dependent FMN and FAD containing oxidoreductase
0.47	-0.07	0.54	BC013586	Hs.1432	PECI	peroxisomal D3,D2-enoyl-CoA isomerase
0.47	0.02	0.45	BC015931	Hs.94308	MS4A1	membrane-spanning 4-domains, subfamily A, member 1
0.47	-1.10	1.57	BC009748	Hs.380681	MGC12904	hypothetical protein MGC12904
0.47	0.35	0.12	BC009376	Hs.250005	PDEF	prostate epithelium-specific Ets transcription factor
0.47	-1.15	1.63	BC007682	Hs.12479	AMSH	associated molecule with the SH3 domain of STAM
0.47	0.45	0.02	BC010086	Hs.30376	MAP3K7	mitogen-activated protein kinase kinase kinase 7
0.47	-1.51	1.98	BC016312	Hs.284162	MGC15613	hypothetical protein MGC15613
0.47	-0.06	0.53	BC015127	Hs.356190	MGC14436	hypothetical protein MGC14436

0.47	-0.19	0.67	BC009889	Hs.143954		CBX1	chromobox homolog 1 (HP1 beta homolog Drosophila )
0.47	-2.15	2.62	BC013589	Hs.82116	ET		hypothetical protein ET
0.47	-0.59	1.06	BC014478	Hs.193725		HSPB1	heat shock 27kDa protein 1
0.47	-0.88	1.35	BE971258	Hs.352254		CRIP2	
0.47	-0.47	0.94	BC005212	Hs.258798		C1QTNF6	C1q and tumor necrosis factor related protein 6
0.47	-0.49	0.96	BC017453	Hs.10554	NA 4-14-03 109		
0.47	-0.39	0.86	BC005818	Hs.140452		TIP47	cargo selection protein (mannose 6 phosphate receptor binding protein)
0.47	-0.02	0.49	BC000917	Hs.79070	MYC		v-myc myelocytomatosis viral oncogene homolog (avian)
0.47	0.42	0.05	BC008217	Hs.91684	LOC92906		hypothetical protein BC008217
0.47	-1.50	1.97	BC011761	Hs.78880	FLJ12903		hypothetical protein FLJ12903
0.47	-0.73	1.20	BC001045	Hs.226499		SARCOSIN	sarcomeric muscle protein
0.47	0.46	0.01	BC002984	Hs.15098	GLUL		glutamate-ammonia ligase (glutamine synthase)
0.47	-0.28	0.75	BC008495	Hs.355719		NGFRAP1	nerve growth factor receptor (TNFRSF16) associated protein 1
0.47	0.16	0.31	BC016294	Hs.184877		BMP7	bone morphogenetic protein 7 (osteogenic protein 1)
0.47	0.11	0.35	BC003125	Hs.323193		ARG2	arginase, type II
0.47	-1.22	1.69	BC016703	Hs.122939		PTMS	
0.47	-0.45	0.91	BC011821	Hs.7393	YARS		tyrosyl-tRNA synthetase
0.47	0.09	0.38	BC019094	Hs.3426	SDS-RS1		likely ortholog of mouse serine dehydratase related sequence 1
0.47	0.46	0.01	BC010152	Hs.3439	NA 4-14-03 246		
0.47	-0.33	0.80	BC001175	Hs.118244		CALM2	calmodulin 2 (phosphorylase kinase, delta)
0.47	0.31	0.16	BC000443	Hs.433813		GDAP1L1	ganglioside-induced differentiation-associated protein 1-like 1
0.46	-0.13	0.59	BC000383	Hs.119	AAAS		achalasia, adrenocortical insufficiency, alacrimia (Allgrove, triple-A)
0.46	0.18	0.29	BC000607	Data not found		RING1	ring finger protein 1
0.46	0.08	0.38	BC008934	Hs.247902		ABLIM1	actin binding LIM protein 1
0.46	-0.55	1.02	BC006795	Hs.332938		MGC5370	hypothetical protein MGC5370
0.46	-0.68	1.14	BC019102	Hs.171118		NXN	nucleoredoxin
0.46	-1.96	2.42	BC001966	Hs.102447		THG-1	TSC-22-like
0.46	-0.62	1.08	BC018991	Hs.89525	MDS024		MDS024 protein
0.46	-0.25	0.72	BC002841	Hs.125819		TIA1	TIA1 cytotoxic granule-associated RNA binding protein
0.46	0.03	0.44	BC015497	Hs.94865	GLUL		glutamate-ammonia ligase (glutamine synthase)
0.46	-0.16	0.62	BC017258	Hs.57101	FLJ13902		hypothetical protein FLJ13902
0.46	-0.95	1.41	BC005850	Hs.31551	SECP43		tRNA selenocysteine associated protein
0.46	0.43	0.03	BC008675	Hs.278619		PVT1	Pvt1 oncogene homolog, MYC activator (mouse)
0.46	-0.08	0.54	BC005402	Hs.284292		G22P1	thyroid autoantigen 70kDa (Ku antigen)
0.46	-0.06	0.52	BC001472	Hs.173162		NOC4	neighbor of COX4
0.46	-0.36	0.82	BC014774	Hs.41714	RPS27A		ribosomal protein S27a
0.46	0.09	0.37	BC017444	Hs.75653	NA 4-14-03 63		
0.46	-0.04	0.50	BC017369	Hs.184167		FSTL1	follistatin-like 1
0.46	-0.31	0.77	BC000686	Hs.46721	H3F3B		H3 histone, family 3B (H3.3B)
0.46	-0.04	0.50	BC008430	Hs.87295	RPL34		ribosomal protein L34
0.46	-0.07	0.53	BC008830	Hs.88959	ABCD4		ATP-binding cassette, sub-family D (ALD), member 4
0.46	-0.39	0.85	BC015981	Hs.192371			
0.46	-0.96	1.42	BC000345	Hs.406511	NA 4-14-03 65		
0.46	-0.51	0.97	BC000532	Hs.76228	OS-9		amplified in osteosarcoma
0.46	-0.27	0.72	BC001114	Hs.180062		PSMB8	proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)
0.46	-0.63	1.09	BC006250	Hs.181035		FHL2	four and a half LIM domains 2
0.46	0.38	0.07	BC000071	Hs.424961	NA 4-14-03 416		
0.46	0.33	0.12	BC021898	Hs.348666		PIASY	protein inhibitor of activated STAT protein PIASy
0.46	-0.14	0.60	BC014269	Hs.77324	POP4		POP4 (processing of precursor , S. cerevisiae) homolog
0.46	0.22	0.24	BC017033	Hs.71465	CTRB1		chymotrypsinogen B1
0.46	0.38	0.08	BC001088	Hs.82906	CDC20		CDC20 cell division cycle 20 homolog (S. cerevisiae)
0.46	-0.20	0.66	BC014572	Hs.146812		UPK1A	uroplakin 1A
0.46	0.20	0.26	BC002529	Hs.281680		PECR	peroxisomal trans 2-enoyl CoA reductase

0.46	0.25	0.21	BC009621	Hs.292154	NA 4-14-03 15	
0.45	0.02	0.44	BC004555	Hs.279852	PTTG1IP	pituitary tumor-transforming 1 interacting protein
0.45	-1.11	1.57	NM_032380	Hs.334718	GAS2L1	growth arrest-specific 2 like 1
0.45	0.31	0.14	BC005236	Hs.334672	SRPR	signal recognition particle receptor ('docking protein')
0.45	-1.27	1.72	BC017046	Hs.118796	SLC7A7	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7
0.45	-0.50	0.95	BC002488	Hs.165998	PAI-RBP1	PAI-1 mRNA-binding protein
0.45	-0.17	0.62	BC012946	Hs.33254	CGI-62	CGI-62 protein
0.45	0.35	0.10	BC020499	Hs.29203	G22P1	thyroid autoantigen 70kDa (Ku antigen)
0.45	0.32	0.13	BC000456	Hs.114360	PL6	PL6 protein
0.45	-0.36	0.81	BC000463	Hs.195614	ZNF22	zinc finger protein 22 (KOX 15)
0.45	-0.48	0.93	BC002513	Hs.151777	EIF2S1	eukaryotic translation initiation factor 2, subunit 1 alpha, 35kDa
0.45	0.39	0.06	BC017211	Hs.54320	SPHK2	sphingosine kinase 2
0.45	-0.33	0.78	BC008053	Hs.8008	CKS1B	CDC28 protein kinase regulatory subunit 1B
0.45	-0.05	0.50	BC008307	Hs.279529	NDUFV1	NADH dehydrogenase (ubiquinone) flavoprotein 1, 51kDa
0.45	-0.43	0.88	BC007097	Hs.5831	FLJ10597	hypothetical protein FLJ10597
0.45	-1.65	2.10	BC007655	In multiple clusters	PPP1R2	
0.45	0.42	0.03	BC007729	Hs.12845	TNNT1	troponin T1, skeletal, slow
0.45	-0.60	1.05	BC018189	Hs.349506	FLJ22175	hypothetical protein FLJ22175
0.45	0.36	0.09	BC002825	Hs.38681	HMG20A	high-mobility group 20A
0.45	-1.55	1.99	BC008720	Hs.9383	CDT1	DNA replication factor
0.45	-0.17	0.62	BC012766	Hs.180428	XPA	xeroderma pigmentosum, complementation group A
0.45	-0.39	0.83	BC004479	Hs.37106	ITM2C	integral membrane protein 2C
0.45	-0.68	1.12	BC017091	Hs.78547	MGP	matrix Gla protein
0.45	0.42	0.02	BC001108	Hs.5245	FLJ20643	hypothetical protein FLJ20643
0.45	-0.98	1.43	BC015701	Hs.170096	NFKB2	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)
0.44	-0.81	1.25	BC002815	Hs.324784	MGC4342	hypothetical protein MGC4342
0.44	0.17	0.27	BC005186	Hs.3385	SERPINA1	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
0.44	-0.60	1.04	BC007220	Hs.381043	TOP3B	topoisomerase (DNA) III beta
0.44	-0.47	0.92	BC006295	Hs.127337	SDCCAG3	serologically defined colon cancer antigen 3
0.44	-0.52	0.96	BC006291	Hs.25303	FLJ20288	FLJ20288 protein
0.44	-0.61	1.05	BC013353	Hs.14070	TAGAP	T-cell activation GTPase activating protein
0.44	-0.36	0.80	BC002588	Hs.405528	PPP1CA	protein phosphatase 1, catalytic subunit, alpha isoform
0.44	-0.81	1.26	BC001331	Hs.79672	KIAA0652	KIAA0652 gene product
0.44	-0.33	0.78	BC001459	Hs.343807	RAD51	RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)
0.44	0.30	0.14	BC015391	Hs.433898	RTDR1	rhabdoid tumor deletion region gene 1
0.44	-0.10	0.54	BC018109	Hs.272673	FLJ10983	hypothetical protein FLJ10983
0.44	-0.23	0.67	BC010188	Data not found	FMNL	formin-like
0.44	-1.16	1.60	BC015355	Hs.200395	SAA4	serum amyloid A4, constitutive
0.44	-1.43	1.87	BC008734	Hs.111903	FCGRT	Fc fragment of IgG, receptor, transporter, alpha
0.44	-1.08	1.52	BC007300	Hs.84746	ASAHI	N-acylsphingosine amidohydrolase (acid ceramidase) 1
0.44	-0.17	0.61	BC015809	Hs.21189	COPEB	core promoter element binding protein
0.44	0.37	0.07	BC009731	Hs.334637	ENDOG	endonuclease G
0.44	-0.62	1.06	BC005848	Hs.70582	DDX32	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 32
0.44	-0.10	0.54	BC016835	Hs.80919	KIAA1049	KIAA1049 protein
0.44	0.29	0.15	BC014449	Hs.266738	NA 4-14-03 142	
0.44	-0.40	0.83	BC006290	Hs.348407	PTD011	PTD011 protein
0.44	-1.19	1.63	BC004971	Hs.433411	NDUFS1	NADH dehydrogenase (ubiquinone) Fe-S protein 1, 75kDa (NADH-coenzyme Q reductase)
0.44	0.24	0.20	BC003161	Hs.6780	ARR3	arrestin 3, retinal (X-arrestin)
0.44	-0.29	0.72	BC018509	Hs.179881	PTK7	
0.44	0.08	0.36	BC012056	Hs.99603	HGS	
0.44	0.00	0.44	BC002414	In multiple clusters	MGC10701	hypothetical protein MGC10701
0.44	0.16	0.27	BC010629	Hs.129055	MGC4692	hypothetical protein MGC4692
0.44	0.14	0.30	BC014204	Hs.311100	TUBB2	tubulin, beta, 2

0.44	-0.10	0.54	BC003373	Hs.288856	PFDN5	prefoldin 5
0.44	-0.93	1.37	BC010241	Hs.159651	NA 4-14-03 37	
0.44	-0.79	1.23	BC001957	Data not found	DPP7	
0.44	-0.54	0.97	BC000745	Hs.42053	MXD3	MAX dimerization protein 3
0.44	-0.33	0.77	BC004372	Hs.169610	ARL2	ADP-ribosylation factor-like 2
0.44	-0.40	0.84	BC001613	Hs.8719	POLR3K	polymerase (RNA) III (DNA directed) polypeptide K, 12.3 kDa
0.44	-0.21	0.64	BC011238	Hs.236438	MGC15397	similar to RIKEN cDNA 5730578N08 gene
0.44	0.33	0.10	BC004371	Hs.279518	NDUFB7	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7, 18kDa
0.44	0.10	0.33	BC017655	Hs.8859	SFRS4	splicing factor, arginine/serine-rich 4
0.44	-0.07	0.50	BC007268	Hs.279529	ITGB3BP	integrin beta 3 binding protein (beta3-endonexin)
0.43	-0.76	1.19	BC001176	Hs.84232	ACOX1	acyl-Coenzyme A oxidase 1, palmitoyl
0.43	-1.09	1.52	BC010995	Hs.26110	NT5C3	5'-nucleotidase, cytosolic III
0.43	0.23	0.20	BC000024	Hs.180780	C15orf15	chromosome 15 open reading frame 15
0.43	0.00	0.43	BC015387	Data not found	DT1P1A10	hypothetical protein DT1P1A10
0.43	-0.21	0.64	BC004139	Hs.334450	IRF5	interferon regulatory factor 5
0.43	-0.17	0.60	BC018645	Hs.72241	GOA	tripartite motif-containing 47
0.43	0.19	0.24	BC016158	Hs.288623	GNG7	guanine nucleotide binding protein (G protein), gamma 7
0.43	0.05	0.38	BC009199	Hs.8867	MLPH	melanophilin
0.43	-0.67	1.10	BC000158	Hs.374608	MGC:5244	hypothetical protein MGC5244
0.43	0.08	0.35	BC010137	Hs.127496	MGC17943	hypothetical protein MGC17943
0.43	0.01	0.42	BC009680	Hs.77290	LYL1	lymphoblastic leukemia derived sequence 1
0.43	-0.07	0.50	BC010868	Hs.100861	FLJ10324	hypothetical protein FLJ10324
0.43	-0.48	0.90	BC017287	Hs.5662	LOXL1	lysyl oxidase-like 1
0.43	-0.41	0.84	BC012137	Hs.302300	DUSP6	dual specificity phosphatase 6
0.43	-0.53	0.96	BC008865	Hs.406548		Homo sapiens, clone IMAGE:4151631, mRNA
0.43	-0.92	1.35	BC017455	Hs.77385	CYLN2	cytoplasmic linker 2
0.43	-0.52	0.95	BC000253	Hs.75527	NDUFC2	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2, 14.5kDa
0.43	0.07	0.35	BC014108	Hs.1416	RAPSN	receptor-associated protein of the synapse, 43kD
0.43	0.15	0.28	BC015417	Hs.131987	LOC113444	hypothetical protein BC011880
0.43	0.38	0.05	BC004105	Hs.18048	MGLL	monoglyceride lipase
0.43	-1.10	1.53	BC005954	Hs.211914	PCDHA2	protocadherin alpha 2
0.43	-0.04	0.47	BC014551	Hs.7277	PEX3	peroxisomal biogenesis factor 3
0.43	-0.41	0.83	BC015466	Hs.159651	MGC19604	similar to RIKEN cDNA B230118G17 gene
0.43	-0.39	0.82	BC000740	Hs.203	MGC14817	hypothetical protein MGC14817
0.43	-0.50	0.93	BC011855	Hs.6216	RBM5	RNA binding motif protein 5
0.43	-0.62	1.05	BC010046	Hs.334	H2AFC	histone 1, H2ai
0.43	-0.29	0.72	BC003611	Hs.279868	SAE1	SUMO-1 activating enzyme subunit 1
0.43	-1.32	1.74	AB067510	Hs.85969	MGC4175	hypothetical protein MGC4175
0.42	-0.01	0.44	BC010405	Hs.57100	DKFZP547N043	hypothetical protein DKFZp547N043
0.42	0.21	0.21	BC014563	Hs.334562	AGPAT2	1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
0.42	-0.95	1.37	BC003183	Hs.3826	C3IP1	kelch-like protein C3IP1
0.42	0.40	0.02	BC010915	Hs.226795	GSTP1	glutathione S-transferase pi
0.42	-0.54	0.97	BC000010	Hs.44396	HOXB13	homeo box B13
0.42	-0.73	1.15	BC001100	Hs.279554	H2AFL	histone 1, H2ac
0.42	0.22	0.20	BC011586	Hs.6980	ENO1	enolase 1, (alpha)
0.42	0.14	0.28	BC019034	Data not found	GPSN2	glycoprotein, synaptic 2
0.42	-0.99	1.41	BC003382	Hs.11183	SNX2	sorting nexin 2
0.42	-0.23	0.66	BC002436	Hs.83734	DSCR1	Down syndrome critical region gene 1
0.42	-0.29	0.71	BC011845	Hs.406444	SH120	putative G-protein coupled receptor
0.42	0.33	0.09	BC007213	Hs.22353	ATP5L	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g
0.42	0.07	0.35	BC002444	Hs.15106	ERCC1	excision repair cross-complementing rodent repair deficiency, complementation group 1 (includes overlapping antisense sequence)

0.42	0.41	0.01	BC010860	Hs.82085	KIAA1622	HEAT-like repeat-containing protein
0.42	-0.23	0.65	BC012751	Data not found	HOXD1	homeo box D1
0.42	0.11	0.31	BC008717	Hs.283738	CSNK1A1	casein kinase 1, alpha 1
0.42	-0.83	1.25	BC012448	Hs.116784	MTCH1	mitochondrial carrier homolog 1
0.42	0.33	0.09	BC007538	Data not found	SLC16A4	solute carrier family 16 (monocarboxylic acid transporters), member 4
0.42	-0.81	1.24	BC014260	Hs.271742	SMAP-5	golgi membrane protein SB140
0.42	-0.79	1.21	BC007441	Hs.23348	SKP2	S-phase kinase-associated protein 2 (p45)
0.42	-0.34	0.76	BC002912	Hs.380081	MGC2198	hypothetical protein MGC2198
0.42	-1.18	1.60	BC017366	Hs.26703	FLJ10565	hypothetical protein FLJ10565
0.42	-0.03	0.45	BC009025	Hs.94891	FLJ12606	hypothetical protein FLJ12606
0.42	-0.88	1.30	BC007781	Hs.108183	RBMS1	RNA binding motif, single stranded interacting protein 1
0.42	-0.50	0.92	BC012600	Hs.84673	TNNI1	troponin I, skeletal, slow
0.42	0.19	0.23	BC001767	Hs.425311	CD24	CD24 antigen (small cell lung carcinoma cluster 4 antigen)
0.42	-0.43	0.84	BC000795	Hs.194385	FLJ20234	hypothetical protein FLJ20234
0.42	0.18	0.24	BC000707	Hs.46735	UBXD1	UBX domain containing 1
0.42	-0.34	0.76	BC009722	Hs.105887	PLP1	proteolipid protein 1 (Pelizaeus-Merzbacher disease, spastic paraplegia 2, uncomplicated)
0.42	-0.60	1.01	BC003558	Hs.75852	HRIHFB2206	HRIHFB2206 protein
0.42	0.35	0.07	BC012977	Hs.283724	LGALS3BP	lectin, galactoside-binding, soluble, 3 binding protein
0.42	-0.11	0.52	BC015930	Hs.186961	TRIP6	thyroid hormone receptor interactor 6
0.42	0.18	0.24	BC001756	Hs.1869	PIGC	phosphatidylinositol glycan, class C
0.42	-0.28	0.70	BC010950	Hs.99949	PIP	prolactin-induced protein
0.42	-0.88	1.30	BC002347	Data not found	PXR1	peroxisome receptor 1
0.42	-0.53	0.95	BC009701	Hs.33455	SNX5	sorting nexin 5
0.42	-0.11	0.53	BC008017	Hs.20776	SERPINA1	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
0.42	-0.85	1.27	BC002448	Hs.158203	CLDN6	claudin 6
0.42	-0.46	0.88	BC020490	Hs.247302	CDK5RAP3	CDK5 regulatory subunit associated protein 3
0.42	-0.89	1.31	BC015490	Hs.431175	PIPE	peptidylprolyl isomerase E (cyclophilin E)
0.41	-1.01	1.42	Data not found		NDRG2	NDRG family member 2
0.41	0.36	0.06	BC002460	Hs.28555	LOC51760	B/K protein
0.41	0.18	0.23	BC013566	Hs.234642	ZNF146	zinc finger protein 146
0.41	-1.86	2.28	BC006097	Hs.1802	BMI1	B lymphoma Mo-MLV insertion region (mouse)
0.41	0.10	0.32	BC012510	Hs.64639	FLJ20010	hypothetical protein FLJ20010
0.41	-0.61	1.02	BC017382	Hs.101799	PPARD	peroxisome proliferative activated receptor, delta
0.41	-1.11	1.53	BC007023	Hs.194382	LOC133957	
0.41	-0.74	1.15	BC006969	Hs.7627	CRB3	crumbs homolog 3 (Drosophila)
0.41	-2.39	2.80	BC002586	Hs.250745	APP	amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)
0.41	0.38	0.03	BC015990	Hs.423444	D4S234E	DNA segment on chromosome 4 (unique) 234 expressed sequence
0.41	-2.50	2.91	BC002564	Hs.155218	E1B-AP5	E1B-55kDa-associated protein 5
0.41	-1.37	1.78	BC012321	Hs.40888	FLJ20721	hypothetical protein FLJ20721
0.41	-0.04	0.45	BC007254	Hs.76228	ID1I	isopentenyl-diphosphate delta isomerase
0.41	-0.01	0.42	BC012381	Hs.6823	HIBCH	3-hydroxyisobutyryl-Coenzyme A hydrolase
0.41	-0.08	0.49	BC020983	Hs.111126	NHP2L1	NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)
0.41	-1.85	2.25	BC002364	In multiple clusters	STOM	stomatins
0.41	0.37	0.04	BC002690	Hs.355214	ABCD3	ATP-binding cassette, sub-family D (ALD), member 3
0.41	0.22	0.19	BC005139	Hs.3759	TETRAN	tetracycline transporter-like protein
0.41	-0.98	1.39	BC000344	Hs.279868	FLJ23436	hypothetical protein FLJ23436
0.41	-0.53	0.94	BC001244	Data not found	KIAA0174	KIAA0174 gene product
0.41	-0.34	0.75	BC005333	Hs.180866	TNFRSF10A	tumor necrosis factor receptor superfamily, member 10a
0.41	-0.74	1.15	BC010512	Hs.93678	MGC16212	hypothetical protein MGC16212
0.41	-0.55	0.96	BC007860	Hs.22393	MGC24302	hypothetical protein MGC24302
0.41	-0.89	1.30	BC000329	Hs.184510	SFN	stratifin
0.41	-0.76	1.17	BC009893	Hs.170027	SLC1A5	solute carrier family 1 (neutral amino acid transporter), member 5
0.41	-0.30	0.71	BC011557	Hs.172740	SIRT2	sirtuin (silent mating type information regulation 2 homolog) 2 (S. cerevisiae)

0.41	0.03	0.38	BC009196	Hs.432840	NA 4-14-03 130	
0.41	-0.82	1.23	BC003640	Hs.30661	ACAS2	acetyl-Coenzyme A synthetase 2 (ADP forming)
0.41	-0.16	0.57	BC007034	Hs.118786	NA 4-14-03 75	
0.41	0.18	0.22	BC008887	Hs.95867	PIM2	pim-2 oncogene
0.41	-0.84	1.25	BC019019	Hs.75850	MGC4645	hypothetical protein MGC4645
0.41	-0.82	1.23	BC018119	Hs.349650	MGC19606	similar to HYPOTHETICAL 34.0 KDA PROTEIN ZK795.3 IN CHROMOSOME IV
0.41	-0.09	0.49	BC009872	Hs.20644	ALP	asparaginase like 1
0.41	0.26	0.15	BC000391	Hs.66196	KIAA1796	KIAA1796 protein
0.41	-0.58	0.98	BC015037	Hs.51305	LAMA4	laminin, alpha 4
0.41	-0.54	0.94	BC015749	Hs.239356	FLJ22728	hypothetical protein FLJ22728
0.41	0.05	0.35	BC009743	Hs.227782	AMID	apoptosis-inducing factor (AIF)-homologous mitochondrion-associated inducer of death
0.41	0.15	0.25	BC008844	Hs.6294	MRPL20	mitochondrial ribosomal protein L20
0.41	0.10	0.30	BC006402	Hs.413826	DNAJB2	DnaJ (Hsp40) homolog, subfamily B, member 2
0.40	-0.18	0.58	BC012050	Hs.14376	CGI-20	CGI-20 protein
0.40	-0.74	1.14	BC001672	Hs.352065	C20orf162	chromosome 20 open reading frame 162
0.40	-0.63	1.03	BC001693	Hs.382367	LGALS1	lectin, galactoside-binding, soluble, 1 (galectin 1)
0.40	0.36	0.04	BC016282	Hs.155212	DKFZP586I2223	intermediate filament-like MGC:2625
0.40	0.27	0.13	BC016841	Hs.301853	RPL23	ribosomal protein L23
0.40	-0.37	0.77	BC002491	Hs.271614	CGI-112	CGI-112 protein
0.40	-0.35	0.75	BC002712	Hs.25960	ZFP36	zinc finger protein 36, C3H type, homolog (mouse)
0.40	-0.10	0.51	BC011681	Hs.4877	AASDHPPT	aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase
0.40	-0.60	1.00	BC021701	Hs.353161	Cab45	calcium binding protein Cab45 precursor
0.40	-0.68	1.09	BC005946	Hs.406578	BART1	binder of Arl Two
0.40	0.03	0.37	BC001738	Hs.192853	NA 4-14-03 112	
0.40	0.15	0.25	BC001482	Hs.8128	MGC2650	hypothetical protein MGC2650
0.40	-0.23	0.63	BC000370	Hs.91448	NA 4-14-03 125	
0.40	-0.15	0.55	BC009920	Hs.9315	FANCG	Fanconi anemia, complementation group G
0.40	-0.36	0.76	BC010374	Hs.96448	ZNF193	zinc finger protein 193
0.40	-0.02	0.42	BC014961	Hs.11463	FLJ21977	hypothetical protein FLJ21977
0.40	-0.10	0.50	BC003353	Hs.13885	MGC5309	hypothetical protein MGC5309
0.40	-0.72	1.12	BC014459	Hs.419463	NPD002	NPD002 protein
0.40	-0.41	0.81	BC011702	Hs.381200	RABL2B	RAB, member of RAS oncogene family-like 2B
0.40	0.30	0.10	BC011901	Hs.2785	ZNF24	zinc finger protein 24 (KOX 17)
0.40	-0.20	0.60	BC010100	Hs.28935	HSPC023	HSPC023 protein
0.40	-0.65	1.05	BC010879	Hs.3123	MGC10744	hypothetical protein MGC10744
0.40	-1.29	1.69	BC006110	Hs.7980	TLE2	transducin-like enhancer of split 2 (E(sp1) homolog, Drosophila)
0.40	-0.38	0.78	BC007889	Hs.251653	FLJ20211	hypothetical protein FLJ20211
0.40	-0.33	0.72	BC004552	Hs.330986	MIC2	CD99 antigen
0.40	-0.57	0.97	BC012629	Hs.30212	TRIP15	thyroid receptor interacting protein 15
0.40	-0.20	0.60	BC007381	Hs.119756	FLJ12787	hypothetical protein FLJ12787
0.40	-0.02	0.41	BC000397	Hs.202384	ASIC4	putative acid-sensing ion channel
0.40	-0.81	1.20	BC001956	Hs.125489	CALM1	calmodulin 1 (phosphorylase kinase, delta)
0.39	-1.44	1.83	BC007562	Hs.106357	MGC23937	hypothetical protein MGC23937 similar to CG4798
0.39	0.20	0.19	BC001711	Hs.434034	SGK2	serum/glucocorticoid regulated kinase 2
0.39	0.31	0.09	BC005028	Hs.25625	NA 4-14-03 83	
0.39	-4.03	4.42	BC017016	Hs.149227	RPP30	ribonuclease P (30kD)
0.39	-0.48	0.88	BC014978	Hs.380417	NA 4-14-03 455	
0.39	-0.23	0.62	BC001841	Hs.250624	KIAA1838	KIAA1838 protein
0.39	-0.83	1.22	BC015013	Hs.283716	FLJ22191	hypothetical protein FLJ22191
0.39	-0.83	1.22	BC008115	Hs.25245	MGC16063	hypothetical protein MGC16063
0.39	0.31	0.08	BC014940	Hs.180933	NA 4-14-03 143	
0.39	0.36	0.02	BC010153	Hs.95577	RPA1	replication protein A1, 70kDa
0.39	-0.21	0.60	BC007092	Hs.66731	CORO2A	coronin, actin binding protein, 2A



0.39	0.02	0.36	BC004146	Hs.261927	PSMB5	proteasome (prosome, macropain) subunit, beta type, 5
0.39	-0.29	0.68	BC012074	Hs.8966	FLJ20424	chromosome 14 open reading frame 94
0.39	-1.32	1.71	BC000924	Hs.321170	NBR2	NBR2
0.39	0.29	0.10	BC007346	Hs.417710	RSU1	Ras suppressor protein 1
0.39	-1.33	1.72	BC021209	Hs.110953	MGC13102	hypothetical protein MGC13102
0.39	-1.16	1.54	BC001352	Hs.336780	TUBB	tubulin, beta polypeptide
0.39	-0.60	0.99	BC001321	Hs.301404	RBM3	RNA binding motif protein 3
0.39	-0.23	0.62	BC004222	Hs.19002	DKFZp434H247	hypothetical protein DKFZp434H247
0.39	-1.80	2.19	BC016026	Hs.42438	HMG20B	high-mobility group 20B
0.39	0.36	0.03	BC005181	Hs.109253	RBM8A	RNA binding motif protein 8A
0.39	-0.34	0.73	BC001006	Hs.289271	C1orf2	chromosome 1 open reading frame 2
0.39	-1.41	1.80	BC014476	Hs.36752	STARD7	START domain containing 7
0.39	0.33	0.06	BC021985	Hs.283109	NOLA2	nucleolar protein family A, member 2 (H/ACA small nucleolar RNPs)
0.39	-0.44	0.83	BC017175	Hs.323396	ARF4L	ADP-ribosylation factor 4-like
0.39	0.27	0.11	BC016028	Data not found	SDCBP2	syndecan binding protein (syntenin) 2
0.39	0.28	0.11	BC000664	Hs.300423	EIF4A1	eukaryotic translation initiation factor 4A, isoform 1
0.39	-0.48	0.87	BC009321	Data not found	NA 4-14-03 76	
0.38	-0.28	0.66	BC006463	Hs.100641	CASP9	caspase 9, apoptosis-related cysteine protease
0.38	-0.31	0.69	BC005235	Hs.15164	AGA	aspartylglucosaminidase
0.38	-0.09	0.48	BC019305	Hs.406444	TA-NFKBH	T-cell activation NFKB-like protein
0.38	-1.26	1.64	BC001106	Hs.181444	FLJ10156	hypothetical protein FLJ10156
0.38	-0.26	0.64	BC014006	Hs.100071	C20orf4	chromosome 20 open reading frame 4
0.38	-0.08	0.47	BC015313	Hs.349905	ACY-3	aspartoacylase-3
0.38	-0.20	0.59	BC008151	Hs.5422	ST13	suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)
0.38	0.23	0.15	BC007621	Hs.225795	LOC115950	hypothetical protein BC016816
0.38	0.32	0.07	BC007349	Hs.60679	NNMT	nicotinamide N-methyltransferase
0.38	-0.34	0.72	BC019077	Hs.17958	FLJ11637	hypothetical protein FLJ11637
0.38	-0.42	0.81	BC001968	Hs.150423	CDK9	cyclin-dependent kinase 9 (CDC2-related kinase)
0.38	0.12	0.26	BC001676	Hs.315177	IFRD2	interferon-related developmental regulator 2
0.38	0.32	0.06	BC015932	Hs.75470	HSU84971	fetal hypothetical protein
0.38	-0.57	0.95	BC000022	Hs.93523	ZNF274	zinc finger protein 274
0.38	0.23	0.15	BC003361	Hs.184276	SLC9A3R1	solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulatory factor 1
0.38	0.00	0.38	BC010178	Hs.78881	POLL	polymerase (DNA directed), lambda
0.38	-1.46	1.84	BC002665	Hs.1787	LOC124220	similar to common salivary protein 1
0.38	0.02	0.35	BC009934	Hs.68583	LOC56961	hypothetical protein from EUROIMAGE 2168212
0.38	-0.66	1.04	BC008909	Hs.10177	TCN2	transcobalamin II; macrocytic anemia
0.38	-0.19	0.57	BC013201	Hs.301920	RAB3A	RAB3A, member RAS oncogene family
0.38	-0.70	1.07	BC014672	Hs.6126	HH114	hypothetical protein HH114
0.38	-0.77	1.14	BC008699	Data not found	NA 4-14-03 333	
0.38	-0.73	1.11	BC015009	Hs.78877	FLJ14909	hypothetical protein FLJ14909
0.37	-0.32	0.69	BC006113	Hs.127762	EPHX2	epoxide hydrolase 2, cytoplasmic
0.37	-0.14	0.51	BC003168	Hs.321622	KIAA1042	OGT(O-Glc-NAc transferase)-interacting protein 106 KDa
0.37	0.26	0.11	BC017449	Hs.165998	MIR16	membrane interacting protein of RGS16
0.37	0.28	0.09	BC010899	Hs.36069	HSPC230	HSPC230 gene
0.37	0.05	0.33	BC015817	Hs.9452	RTBDN	retbindin
0.37	-0.33	0.70	BC002855	Hs.8008	REQ	requiem, apoptosis response zinc finger gene
0.37	-0.87	1.24	BC005377	Hs.79158	ENTPD5	ectonucleoside triphosphate diphosphohydrolase 5
0.37	-0.78	1.16	BC008958	Hs.78040	KDELRL1	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1
0.37	-0.60	0.97	Data not found		JUP	junction plakoglobin
0.37	-0.23	0.60	Data not found		MGC4342	hypothetical protein MGC4342
0.37	0.05	0.32	BC003559	Hs.234726	DKFZP564D0478	hypothetical protein DKFZp564D0478
0.37	0.18	0.19	BC001671	Hs.32826	CGI-130	CGI-130 protein
0.37	-0.15	0.52	BC000716	Hs.144063	SF3A2	splicing factor 3a, subunit 2, 66kDa

0.37	0.32	0.05	BC014645	Hs.86172	NDUFS5	NADH dehydrogenase (ubiquinone) Fe-S protein 5, 15kDa (NADH-coenzyme Q reductase)
0.37	0.00	0.37	BC010146	Hs.33787	FLJ20574	hypothetical protein FLJ20574
0.37	-0.55	0.92	BC009084	Hs.334841	LOC90268	hypothetical protein BC007706
0.37	-0.22	0.59	BC003571	Hs.416629	SRC	v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)
0.37	-0.15	0.52	BC003130	Hs.8015	IGFBP4	insulin-like growth factor binding protein 4
0.37	-0.25	0.61	BC001453	Hs.26403	GSTZ1	glutathione transferase zeta 1 (maleylacetoacetate isomerase)
0.37	0.29	0.07	BC000076	Hs.82932	LYPLA2	lysophospholipase II
0.37	-0.23	0.60	BC012443	Hs.30792	HSPC126	HSPC126 protein
0.37	-0.13	0.50	BC010444	Hs.19368	UCHL1	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)
0.37	-0.33	0.69	BC004440	Hs.11342	LOC114987	similar to spermatid WD-repeat protein
0.37	0.05	0.31	BC010466	In multiple clusters	FLJ20274	hypothetical protein FLJ20274
0.37	-0.61	0.97	BC001961	Hs.10499	AF1Q	ALL1-fused gene from chromosome 1q
0.37	0.19	0.17	BC013900	Hs.268189	POLH	polymerase (DNA directed), eta
0.37	-0.48	0.84	BC006306	Hs.108779	IK	IK cytokine, down-regulator of HLA II
0.37	-0.09	0.46	BC002970	Hs.10729	MSI2	musashi homolog 2 (Drosophila)
0.36	-0.02	0.39	Data not found		COCH	coagulation factor C homolog, cochlin (Limulus polyphemus)
0.36	-0.66	1.02	BC017088	Hs.10283	NAT5	N-acetyltransferase 5 (ARD1 homolog, S. cerevisiae)
0.36	-0.49	0.86	BC016790	Hs.173042	MGC15716	hypothetical protein MGC15716
0.36	-0.56	0.92	BC014974	Data not found	MAPKAPK3	mitogen-activated protein kinase-activated protein kinase 3
0.36	-0.04	0.40	BC006181	Data not found	NA 4-14-03 21	
0.36	-1.34	1.70	BC001937	Hs.180903	AIPL1	aryl hydrocarbon receptor interacting protein-like 1
0.36	0.02	0.35	BC005387	Hs.75313	CXorf12	chromosome X open reading frame 12
0.36	-1.26	1.62	BC015904	Hs.347535	PURB	purine-rich element binding protein B
0.36	-0.54	0.90	BC006128	Hs.98328	BLNK	B-cell linker
0.36	-0.79	1.15	BC006399	Hs.155839	LOC201853	similar to hypothetical protein (L1H 3 region) - human
0.36	-1.36	1.72	BC012561	Hs.241531	PEF	PEF protein with a long N-terminal hydrophobic domain (peffin)
0.36	-0.04	0.40	BC007778	Data not found	MRPL20	mitochondrial ribosomal protein L20
0.36	-0.11	0.47	BC009610	Hs.349506	PC4	activated RNA polymerase II transcription cofactor 4
0.36	-0.17	0.53	BC014534	Hs.94125	FLJ11113	methyl-CpG binding domain protein 5
0.36	-0.33	0.69	BC015773	Hs.192822	PLAT	plasminogen activator, tissue
0.36	-1.53	1.89	BC002960	Hs.69745	FLJ11184	hypothetical protein FLJ11184
0.36	-0.20	0.56	BC015191	Hs.104741	TOPK	T-LAK cell-originated protein kinase
0.36	-0.54	0.90	BC002536	Hs.99910	MGC10993	hypothetical protein MGC10993
0.36	-0.23	0.59	BC002372	Hs.62112	ALDH9A1	aldehyde dehydrogenase 9 family, member A1
0.36	-0.60	0.95	BC009424	Hs.352397	LOC122618	hypothetical protein BC015003
0.36	-1.02	1.37	BC009944	Hs.157144	MGC2731	hypothetical protein MGC2731
0.36	-0.17	0.53	BC010363	Hs.105633	FLJ10583	WINS1 protein with Drosophila Lines (Lin) homologous domain
0.36	-0.67	1.03	BC012535	Hs.62354	LRBA	LPS-responsive vesicle trafficking, beach and anchor containing
0.36	0.20	0.16	BC009928	Hs.4747	HSPB7	heat shock 27kDa protein family, member 7 (cardiovascular)
0.36	-1.35	1.70	BC001261	Hs.140944	D6S2654E	DNA segment on chromosome 6(unique) 2654 expressed sequence
0.36	-1.40	1.75	BC017574	Hs.24371	MGC15668	hypothetical protein MGC15668
0.36	-1.52	1.88	BC001167	Hs.79372	EIF3S6	eukaryotic translation initiation factor 3, subunit 6 48kDa
0.36	-0.69	1.05	BC001364	Hs.50785	SEC22L1	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)
0.36	-0.02	0.38	BC017188	Hs.226275		
0.35	0.19	0.16	BC012358	Hs.30570	PRC1	protein regulator of cytokinesis 1
0.35	0.21	0.15	BC009046	Hs.72981	LCP2	lymphocyte cytosolic protein 2 (SH2 domain containing leukocyte protein of 76kDa)
0.35	0.26	0.10	BC010900	Hs.21275	FLJ11011	hypothetical protein FLJ11011
0.35	-0.34	0.69	BC005045	Hs.283007	SNK	serum-inducible kinase
0.35	-1.03	1.38	BC002772	Hs.274416	NA 4-14-03 211	
0.35	0.33	0.02	BC006231	Hs.226573	IKBKB	inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta
0.35	-0.61	0.96	BC000635	Hs.166160	GSTA2	glutathione S-transferase A2
0.35	-0.25	0.60	BC009769	Hs.3576	C20orf30	chromosome 20 open reading frame 30
0.35	0.08	0.28	BC012083	Hs.21068	DDX37	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 37

0.35	0.10	0.25	BC004341	Hs.69485	CGI-63	CGI-63 protein
0.35	0.15	0.20	BC005385	Hs.74502	SQLE	squalene epoxidase
0.35	-0.28	0.63	BC013643	Hs.323512	MGC15523	hypothetical protein MGC15523
0.35	-0.53	0.88	BC001462	Hs.75799	PRSS8	protease, serine, 8 (prostasin)
0.35	-0.70	1.05	BC007315	Hs.380963	PDXK	pyridoxal (pyridoxine, vitamin B6) kinase
0.35	-0.25	0.60	BC000105	Hs.239760	IMPA2	inositol(myo)-1(or 4)-monophosphatase 2
0.35	0.33	0.02	BC017339	Hs.293691	LAMR1	laminin receptor 1 (ribosomal protein SA, 67kDa)
0.35	0.33	0.02	BC007821	Hs.334525	UBB	ubiquitin B
0.35	0.16	0.19	BC000408	Hs.278544	C5orf3	chromosome 5 open reading frame 3
0.35	-0.31	0.66	BC011606	Hs.172673	SYNGR1	synaptogyrin 1
0.35	-0.19	0.54	BC017049	Hs.249163	ORC4L	origin recognition complex, subunit 4-like (yeast)
0.35	0.20	0.15	BC006996	Hs.21756	FAM3D	family with sequence similarity 3, member D
0.35	-0.24	0.59	BC007012	Hs.64746	LOC51061	hypothetical protein LOC51061
0.35	-0.99	1.34	BC001778	Hs.78601	HSPBAP1	HSPB (heat shock 27kDa) associated protein 1
0.35	-1.14	1.49	BC014272	Hs.57101	ARHGAP9	Rho GTPase activating protein 9
0.35	-0.76	1.10	BC005860	Hs.326	TARBP2	TAR (HIV) RNA binding protein 2
0.35	-0.05	0.40	BC002899	Hs.161362	MGC1203	hypothetical protein MGC1203
0.35	-0.43	0.78	BC012926	Hs.5366	NDUFB8	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8, 19kDa
0.35	-1.95	2.30	BC007965	Hs.425808	LOC132948	LOC132948
0.35	-1.00	1.34	BC005924	Hs.390278	KIAA0153	KIAA0153 protein
0.35	-0.51	0.86	BC016009	Hs.418650	KCNK13	potassium channel, subfamily K, member 13
0.35	-0.75	1.10	BC010235	Hs.30773	SSBP3	
0.35	0.32	0.02	BC000867	Hs.6451	PRO0659	PRO0659 protein
0.35	-1.09	1.43	BC011396	Hs.98806	PCYT2	phosphate cytidyltransferase 2, ethanolamine
0.35	-0.55	0.90	BC001594	Hs.5298	TRIAD3	TRIAD3 protein
0.34	-0.08	0.42	BC000369	Hs.107003	HEI10	enhancer of invasion 10
0.34	-0.36	0.71	BC003008	Hs.20930	PCBP4	poly(rC) binding protein 4
0.34	-0.05	0.40	BC015109	Hs.283693	DKFZp727A071	similar to tRNA synthetase class II
0.34	-0.16	0.50	BC015823	Hs.105938	LTF	lactotransferrin
0.34	-0.09	0.43	BC001220	Hs.128764	PSAP	prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy)
0.34	-1.32	1.67	BC008070	Hs.30026	DGCR6L	DiGeorge syndrome critical region gene 6-like
0.34	-0.42	0.77	BC007856	Hs.380388		Homo sapiens, clone IMAGE:4310290, mRNA
0.34	-0.27	0.61	BC004137	Hs.3631	IGBP1	immunoglobulin (CD79A) binding protein 1
0.34	-0.23	0.57	BC002818	In multiple clusters	CHRAC1	chromatin accessibility complex 1
0.34	0.16	0.18	BC014513	Hs.101813	WNT7A	wingless-type MMTV integration site family, member 7A
0.34	-0.03	0.37	BC002934	Hs.289108	HKE4	HLA class II region expressed gene KE4
0.34	0.18	0.16	BC003412	Hs.9880	PIIH	peptidyl prolyl isomerase H (cyclophilin H)
0.34	-0.29	0.63	BC016952	Hs.8867	TRAP1	heat shock protein 75
0.34	-0.25	0.59	BC011924	Hs.118261	QARS	glutaminyl-tRNA synthetase
0.34	-0.19	0.53	BC012976	Hs.44856	POF1B	premature ovarian failure 1B
0.34	-0.53	0.88	BC006371	Hs.47438	KDELR2	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2
0.34	-1.52	1.86	BC009312	Hs.20725	ADRM1	adhesion regulating molecule 1
0.34	-0.57	0.91	BC009601	Hs.75454	PRDX3	peroxiredoxin 3
0.34	0.11	0.23	BC007495	Hs.208912	MGC861	hypothetical protein MGC861
0.34	-0.02	0.36	BC012327	Hs.252229	PA2G4	
0.34	-2.61	2.95	BC004973	Hs.181015	COL6A2	collagen, type VI, alpha 2
0.34	-0.32	0.66	BC005825	Hs.38041	GSTTLp28	glutathione-S-transferase like; glutathione transferase omega
0.34	0.25	0.09	BC006838	Hs.122579	ECT2	epithelial cell transforming sequence 2 oncogene
0.34	0.04	0.30	BC004291	Hs.380439	DNCI2	dynein, cytoplasmic, intermediate polypeptide 2
0.34	-0.26	0.59	BC018702	Hs.233950	SPINT1	serine protease inhibitor, Kunitz type 1
0.34	-2.00	2.34	BC011539	Hs.17908	PSME1	proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)
0.34	0.29	0.04	BC008729	Hs.348262	KIAA1309	KIAA1309 protein
0.34	-0.71	1.05	BC001179	Hs.177516	FPGS	folylpolyglutamate synthase

0.34	-0.83	1.17	BC009340	Data not found	C20orf108	chromosome 20 open reading frame 108
0.34	-1.07	1.40	BC006298	Hs.77039	CLTA	clathrin, light polypeptide (Lca)
0.34	-0.42	0.76	BC013115	Hs.351478	PSEN1	presenilin 1 (Alzheimer disease 3)
0.34	-0.83	1.17	BC001256	Hs.272299	HEI10	enhancer of invasion 10
0.34	-0.53	0.87	BC010959	Hs.77572	BNIP1	BCL2/adenovirus E1B 19kDa interacting protein 1
0.33	-0.48	0.81	BC001621	Hs.334811	CIP29	cytokine induced protein 29 kDa
0.33	0.28	0.05	BC004457	Hs.84045	SLC1A7	solute carrier family 1 (glutamate transporter), member 7
0.33	0.13	0.20	BC014894	Data not found	FLJ23093	chromosome 14 open reading frame 140
0.33	-0.66	0.99	BC000397	Hs.202384	T54	T54 protein
0.33	0.05	0.28	BC017061	Hs.3363	FLJ10450	hypothetical protein FLJ10450
0.33	0.13	0.20	BC014665	Hs.78989	NA 4-14-03 448	
0.33	0.28	0.06	BC015474	Hs.238039	IFNAR2	interferon (alpha, beta and omega) receptor 2
0.33	-1.15	1.48	BC006475	Hs.39504	MGC4308	hypothetical protein MGC4308
0.33	0.31	0.02	BC014455	Hs.155595	MESDC2	mesoderm development candidate 2
0.33	-1.50	1.83	BC008374	Hs.432325	PR48	protein phosphatase 2A 48 kDa regulatory subunit
0.33	-0.73	1.06	BC013973	Hs.184325	TSSC3	tumor suppressing subtransferable candidate 3
0.33	0.21	0.12	BC011361	Hs.169743	PSMB7	proteasome (prosome, macropain) subunit, beta type, 7
0.33	0.18	0.15	BC004361	Hs.303091	CCND1	cyclin D1 (PRAD1: parathyroid adenomatosis 1)
0.33	-0.21	0.54	BC000274	Hs.324618	SRP9	signal recognition particle 9kDa
0.33	0.26	0.07	BC000342	Hs.174051	SEMA4B	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B
0.33	0.05	0.28	BC000504	Hs.7043	HDAC6	histone deacetylase 6
0.33	0.18	0.15	BC019098	Hs.1497	MAP4	microtubule-associated protein 4
0.33	-0.79	1.12	BC002402	Hs.240534	AGPAT1	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)
0.33	-0.31	0.64	BC013426	Hs.111217	ASNA1	arsA arsenite transporter, ATP-binding, homolog 1 (bacterial)
0.33	-0.84	1.17	BC012846	Hs.11223	STARD7	START domain containing 7
0.33	-0.16	0.49	BC017401	Hs.123035	LOC90120	hypothetical gene supported by AK023162
0.33	0.00	0.33	BC000021	Hs.355264	NA 4-14-03 422	
0.33	-0.18	0.51	BC017264	Hs.124962	SERF2	small EDRK-rich factor 2
0.33	-0.08	0.41	BC007714	Hs.111244	RAB6KIFL	kinesin family member 20A
0.33	-0.41	0.74	BC000777	Hs.5301	HYPK	Huntingtin interacting protein K
0.33	-0.53	0.86	BC003356	Hs.282997	GBA	glucosidase, beta; acid (includes glucosylceramidase)
0.33	-0.64	0.97	BC010992	Hs.18878	FLJ20605	hypothetical protein FLJ20605
0.33	-0.27	0.60	BC007783	Hs.73817	IGFBP7	insulin-like growth factor binding protein 7
0.33	-0.36	0.68	BC000785	Hs.274411	FLJ22471	limkain beta 2
0.33	0.23	0.09	BC008781	Hs.18029	FLJ10204	hypothetical protein FLJ10204
0.33	0.03	0.29	BC000395	Hs.75884	SMARCC2	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2
0.33	-0.44	0.77	BC002482	Hs.286261	LGALS3BP	lectin, galactoside-binding, soluble, 3 binding protein
0.33	0.02	0.31	BC001690	Hs.177530	ATP5E	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, epsilon subunit
0.33	-0.76	1.08	BC006460	Hs.15299	HIS1	HMBA-inducible
0.32	-0.26	0.58	BC015066	Hs.153934	CBFA2T2	core-binding factor, runt domain, alpha subunit 2; translocated to, 2
0.32	-2.26	2.58	BC000182	Hs.422986	MGC2555	hypothetical protein MGC2555
0.32	-0.75	1.08	BC020107	Hs.12457	MGC19825	similar to RIKEN cDNA 5630401J11
0.32	0.15	0.18	BC018992	Hs.170171	CARKL	carbohydrate kinase-like
0.32	-0.59	0.92	BC000349	Hs.282997	LOC51693	unknown
0.32	-0.05	0.38	BC014469	Hs.72901	ANKH	ankylosis, progressive homolog (mouse)
0.32	-2.63	2.96	BC018370	Hs.12523	CREB3	cAMP responsive element binding protein 3 (luman)
0.32	0.06	0.26	BC014175	Hs.159428	DKFZP434A043	DKFZP434A043 protein
0.32	-1.05	1.37	BC009304	Hs.65234	PAM	peptidylglycine alpha-amidating monooxygenase
0.32	-1.26	1.58	BC021214	Hs.180900	MGC13114	hypothetical protein MGC13114
0.32	-1.80	2.12	BC000414	Hs.99843	CBS	cystathionine-beta-synthase

0.32	-0.02	0.34	BC007449	Hs.127685	KIAA1627	KIAA1627 protein
0.32	0.12	0.20	BC010117	Hs.158515	BTC	betacellulin
0.32	-0.47	0.79	BC001334	Hs.82520	POLD4	polymerase (DNA-directed), delta 4
0.32	-1.25	1.56	BC008493	Hs.1852	Apg4B	KIAA0943 protein
0.32	-0.90	1.22	BC002338	Hs.73722	NCPB2	nuclear cap binding protein subunit 2, 20kDa
0.32	0.02	0.29	BC000286	Hs.150748	EIF3S4	eukaryotic translation initiation factor 3, subunit 4 delta, 44kDa
0.32	-0.25	0.57	BC013033	Hs.184691	CPZ	carboxypeptidase Z
0.32	-0.15	0.47	BC014416	Hs.285176	MLC-B	myosin regulatory light chain
0.32	-0.33	0.64	NM_018383	Hs.107000	FLJ11294	hypothetical protein FLJ11294
0.32	-1.05	1.37	BC020521	Hs.250705	GADD45A	growth arrest and DNA-damage-inducible, alpha
0.32	-0.31	0.63	BC003366	Hs.92198	CARHSP1	calcium regulated heat stable protein 1, 24kDa
0.32	-1.49	1.81	BC005155	Hs.252280	TREX1	three prime repair exonuclease 1
0.32	-1.05	1.37	BC007957	Hs.28393	LOC115131	similar to POSSIBLE GUSTATORY RECEPTOR CLONE PTE01
0.32	-0.13	0.45	BC007677	Hs.83530	FLJ20811	hypothetical protein FLJ20811
0.32	-0.70	1.01	BC004152	Hs.5199	HSPC150	HSPC150 protein similar to ubiquitin-conjugating enzyme
0.31	-0.52	0.83	BC006534	Hs.50550	NESCA	RUN and SH3 domain containing 1
0.31	-0.74	1.05	BC011812	Hs.7960	RPPN	ropporin, rhophilin associated protein 1
0.31	-0.80	1.12	BC004873	Hs.79507	CNOT2	CCR4-NOT transcription complex, subunit 2
0.31	-1.89	2.20	BC005827	Hs.2178	GNAT2	guanine nucleotide binding protein (G protein), alpha transducing activity polypeptide 2
0.31	0.26	0.06	BC016039	Hs.279914	UQCRH	ubiquinol-cytochrome c reductase hinge protein
0.31	-1.05	1.36	BC003055	Hs.238928	BTD	biotinidase
0.31	-0.15	0.47	BC018503	Hs.2025	PTPN2	protein tyrosine phosphatase, non-receptor type 2
0.31	0.16	0.15	BC008191	Hs.7984	PSCD3	pleckstrin homology, Sec7 and coiled/coil domains 3
0.31	0.00	0.31	BC001401	Hs.115175	ZAK	sterile alpha motif and leucine zipper containing kinase AZK
0.31	-1.07	1.38	BC014889	Hs.13495	DMAP1	DNA methyltransferase 1-associated protein 1
0.31	-0.90	1.21	BC004141	Hs.75335	GATM	glycine amidinotransferase (L-arginine:glycine amidinotransferase)
0.31	-0.58	0.90	BC020967	Hs.150555	ATP5E	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, epsilon subunit
0.31	-1.12	1.43	BC014310	Hs.308332	FLJ10305	hypothetical protein FLJ10305
0.31	0.00	0.31	BC017565	Hs.119177	ATP1B1	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 1 polypeptide
0.31	0.10	0.21	BC022881	Hs.369550	PSMB4	proteasome (prosome, macropain) subunit, beta type, 4
0.31	-0.10	0.41	BC015775	Hs.75852	MAGEA8	melanoma antigen, family A, 8
0.31	-0.48	0.79	BC011786	Data not found	ARP3BETA	actin-related protein 3-beta
0.31	-0.85	1.16	BC000151	Hs.10729	HSPC117	hypothetical protein HSPC117
0.31	0.15	0.16	BC009264	Hs.192728	KIAA1720	KIAA1720 protein
0.31	-0.55	0.86	BC021217	Hs.59745	NDUFV3	NADH dehydrogenase (ubiquinone) flavoprotein 3, 10kDa
0.31	-0.59	0.90	BC005013	Hs.106004	FLJ10350	hypothetical protein FLJ10350
0.31	-0.31	0.61	BC006107	Hs.19807	MCM2	MCM2 minichromosome maintenance deficient 2, mitotin (S. cerevisiae)
0.31	-0.28	0.58	BC010737	Data not found	C22orf3	chromosome 22 open reading frame 3
0.31	0.06	0.25	BC000075	Hs.251064	CKB	creatine kinase, brain
0.31	-0.71	1.01	BC007654	Hs.21943	NIF3L1	NIF3 NGG1 interacting factor 3-like 1 (S. pombe)
0.31	0.17	0.13	BC011889	Hs.334639	RNF32	ring finger protein 32
0.31	-0.64	0.95	BC004537	Hs.389107	ATP6V0C	ATPase, H <sup>+</sup> transporting, lysosomal 16kDa, V0 subunit c
0.31	-0.56	0.87	BC001871	Hs.184062	EFEMP1	EGF-containing fibulin-like extracellular matrix protein 1
0.31	-0.02	0.33	BC021165	Hs.158210	HSPCB	heat shock 90kDa protein 1, beta
0.31	-1.12	1.43	BC011983	Hs.381179	UBL5	ubiquitin-like 5
0.31	-0.40	0.71	BC018891	Data not found	MGC13071	hypothetical protein MGC13071
0.31	-0.52	0.82	BC005185	Hs.234896	PP2447	hypothetical protein PP2447
0.30	-0.86	1.16	BC000759	Hs.24563	NXT1	NTF2-like export factor 1
0.30	-1.05	1.36	BC014356	Hs.283693	PELO	pelota homolog (Drosophila)
0.30	-0.34	0.65	BC015926	Hs.48924	CITED1	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 1
0.30	0.22	0.08	BC012826	Hs.274351	SNRPN	small nuclear ribonucleoprotein polypeptide N
0.30	0.26	0.05	BC002454	Hs.71642	PSA	phosphoserine aminotransferase
0.30	-0.53	0.83	BC012613	Hs.646	CPA3	carboxypeptidase A3 (mast cell)

0.30	0.02	0.28	BC010922	Hs.142245	HHLA3	HERV-H LTR-associating 3
0.30	0.24	0.07	BC021226	Hs.19122	EIF4EL3	eukaryotic translation initiation factor 4E-like 3
0.30	0.20	0.10	BC000944	Hs.15580	LENG5	leukocyte receptor cluster (LRC) member 5
0.30	0.09	0.21	BC019040	Hs.103724	GEMIN7	gem (nuclear organelle) associated protein 7
0.30	-1.05	1.35	BC000819	Hs.180378	LOC56267	hypothetical protein 669
0.30	-0.96	1.26	BC009614	Hs.6783	MXD4	MAX dimerization protein 4
0.30	-0.23	0.54	BC009051	Data not found	CSTF2	cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kDa
0.30	-1.29	1.59	BC005158	Hs.89072	NA 4-14-03 137	
0.30	0.11	0.19	BC014666	Hs.187520	NRP2	neuropilin 2
0.30	0.26	0.04	BC014457	Hs.54609	LOC151194	similar to hepatocellular carcinoma-associated antigen HCA557b
0.30	-1.02	1.33	BC005050	Hs.191460	HSD17B12	hydroxysteroid (17-beta) dehydrogenase 12
0.30	-0.23	0.53	Data not found		CGI-18	CGI-18 protein
0.30	-1.84	2.14	BC003061	Hs.18069	MDS032	uncharacterized hematopoietic stem/progenitor cells protein MDS032
0.30	-0.47	0.77	BC015947	Hs.47584	BCAT2	branched chain aminotransferase 2, mitochondrial
0.30	-0.30	0.60	BC000706	Hs.109798	FLJ20847	hypothetical protein FLJ20847
0.30	-0.72	1.02	BC008849	Data not found	ACY1	aminoacylase 1
0.30	0.22	0.08	BC012051	Hs.234282	MUTYH	mutY homolog (E. coli)
0.30	0.19	0.11	BC000373	Hs.279518	SPUVE	protease, serine, 23
0.30	0.15	0.14	BC001210	Hs.12851	PPIA	peptidylprolyl isomerase A (cyclophilin A)
0.30	-0.46	0.76	BC001340	Hs.12293	FLJ21103	hypothetical protein FLJ21103
0.30	-1.04	1.34	BC000234	Hs.364345	TAF9	TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 32kDa
0.30	-0.27	0.57	BC000430	Hs.75824	NA 4-14-03 177	
0.30	-0.71	1.00	BC004368	Hs.433810	NPC2	Niemann-Pick disease, type C2
0.30	-2.62	2.92	BC004217	Hs.105547	LOC112970	hypothetical protein BC012173
0.30	-0.27	0.57	BC002820	Hs.272018	ALTE	Ac-like transposable element
0.30	-2.82	3.12	BC008095	Hs.333417	MGC21636	hypothetical protein MGC21636
0.30	-0.13	0.42	BC012305	Hs.333417	MARK4	MAP/microtubule affinity-regulating kinase 4
0.30	0.16	0.14	BC009470	Hs.18571	SCAMP3	secretory carrier membrane protein 3
0.30	-1.31	1.60	BC008337	Hs.272838	C11orf10	chromosome 11 open reading frame 10
0.30	-0.17	0.47	BC007116	Hs.334568	P3	Protein P3
0.30	-0.10	0.40	BC005210	Hs.26570	CABYR	calcium-binding tyrosine-(Y)-phosphorylation regulated (fibrousheathin 2)
0.29	-2.07	2.36	BC011405	Hs.23650	PDHA1	pyruvate dehydrogenase (lipoamide) alpha 1
0.29	0.00	0.30	BC008767	Hs.379991	TCN2	transcobalamin II; macrocytic anemia
0.29	0.06	0.23	BC018409	Hs.150319	CGI-60	CGI-60 protein
0.29	0.19	0.11	BC008323	Hs.173724	LOC51234	hypothetical protein LOC51234
0.29	-0.43	0.73	BC006566	Hs.9599	ACTR1B	ARP1 actin-related protein 1 homolog B, cetractin beta (yeast)
0.29	-1.19	1.49	BC009388	Data not found	IL6	interleukin 6 (interferon, beta 2)
0.29	-0.47	0.77	BC014501	Data not found	MTA1	metastasis associated 1
0.29	-1.13	1.42	BC015768	Hs.285115	STARD7	START domain containing 7
0.29	0.19	0.10	BC006114	Hs.89605	CSRP2BP	CSRP2 binding protein
0.29	0.13	0.16	BC009471	Hs.128749	HNRPAB	heterogeneous nuclear ribonucleoprotein A/B
0.29	-0.11	0.40	BC007674	Hs.375108	LSM1	LSM1 homolog, U6 small nuclear RNA associated (S. cerevisiae)
0.29	0.13	0.16	BC016325	Hs.75290	HADH2	hydroxyacyl-Coenzyme A dehydrogenase, type II
0.29	-0.65	0.94	BC010965	Hs.28988	GLRX	glutaredoxin (thioltransferase)
0.29	-0.79	1.08	BF313143	Hs.318456	IDH3B	isocitrate dehydrogenase 3 (NAD+) beta
0.29	-0.17	0.46	BC008364	Hs.418901	HNRPC	heterogeneous nuclear ribonucleoprotein C (C1/C2)
0.29	0.18	0.11	BC003385	Hs.129951	SPOP	speckle-type POZ protein
0.29	-0.77	1.06	BC000590	Hs.83583	SF1	splicing factor 1
0.29	-0.47	0.76	BC009396	Hs.288891	SQRDL	sulfide quinone reductase-like (yeast)
0.29	0.16	0.13	BC020584	Hs.25307	PIGO	phosphatidylinositol glycan class O
0.29	-1.58	1.87	BC001959	Hs.7862	RPS2	ribosomal protein S2
0.29	-0.40	0.69	BC000498	Hs.597	KIAA1036	KIAA1036 protein
0.29	-0.46	0.74	BC014844	Hs.118640	UQCR	ubiquinol-cytochrome c reductase (6.4kD) subunit

0.29	-0.42	0.70	BC009856	Hs.12045	CEACAM1	carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
0.29	-0.68	0.97	BC012855	Hs.125201	ALB	albumin
0.29	-0.74	1.03	BC005233	Hs.73923	CRADD	CASP2 and RIPK1 domain containing adaptor with death domain
0.29	0.04	0.25	BC005397	Hs.55408	SNX7	sorting nexin 7
0.29	-0.76	1.05	BC005920	Data not found		
0.29	0.05	0.24	BC006294	Data not found	DLG7	discs, large homolog 7 (Drosophila)
0.29	0.09	0.19	BC007408	Hs.143641	MGC3121	hypothetical protein MGC3121
0.29	-0.46	0.74	BC008417	In multiple clusters	NALP2	NALP2 protein
0.29	0.24	0.04	BC004973	Hs.181015	LAT	linker for activation of T cells
0.29	-0.68	0.97	BC001251	Hs.80426	BRE	brain and reproductive organ-expressed (TNFRSF1A modulator)
0.29	-1.66	1.95	BC019246	Hs.195155	NR1H3	nuclear receptor subfamily 1, group H, member 3
0.29	0.07	0.22	BC015998	Hs.82906	CDC34	cell division cycle 34
0.29	-1.15	1.43	BC009439	Hs.182986	PACE-1	ezrin-binding partner PACE-1
0.29	-0.57	0.86	BC007704	Hs.381183	MDH1	malate dehydrogenase 1, NAD (soluble)
0.29	-1.51	1.79	BC001716	Hs.396644	ZNF323	zinc finger protein 323
0.29	-0.32	0.61	NM_018474	Hs.173515	FLJ14779	hypothetical protein FLJ14779
0.29	-0.76	1.05	BC010953	Hs.91161	PFDN4	prefoldin 4
0.29	-0.34	0.62	BC021224	In multiple clusters	Magmas	mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction
0.28	-0.16	0.44	BC002851	Hs.429437	FLJ32915	hypothetical protein FLJ32915
0.28	-1.64	1.92	BC015517	Hs.302008	SLC27A4	solute carrier family 27 (fatty acid transporter), member 4
0.28	-0.15	0.43	BC017066	Hs.6614	RNF40	ring finger protein 40
0.28	-0.41	0.69	BC009348	Hs.151001	CES1	carboxylesterase 1 (monocyte/macrophage serine esterase 1)
0.28	-1.52	1.80	BC004885	Hs.272299	ABCB6	ATP-binding cassette, sub-family B (MDR/TAP), member 6
0.28	-0.77	1.06	BC000320	Hs.5790	SCAMP-4	secretory carrier membrane protein 4
0.28	-0.05	0.33	BC020987	Hs.78442	IMPA1	inositol(myo)-1(or 4)-monophosphatase 1
0.28	-0.64	0.93	BC000049	Hs.279610	M17S2	membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)
0.28	-0.25	0.53	BC002781	Hs.76122	SHAPY	Ca <sup>2+</sup> -dependent endoplasmic reticulum nucleoside diphosphatase
0.28	-0.51	0.79	BC017352	Hs.82237	CLCN7	chloride channel 7
0.28	-0.04	0.33	BC007591	Hs.227789	XRCC4	X-ray repair complementing defective repair in Chinese hamster cells 4
0.28	0.02	0.26	BC018145	Hs.19523	TPMT	thiopurine S-methyltransferase
0.28	0.00	0.28	BC002711	Hs.146409	C20orf64	chromosome 20 open reading frame 64
0.28	-0.14	0.42	BC007065	Hs.77490	LOC90693	alpha-1,3(6)-mannosylglycoprotein beta-1,6-N-acetyl-glucosaminyltransferase-like
0.28	0.12	0.16	BC002918	Hs.25204	PEA15	phosphoprotein enriched in astrocytes 15
0.28	-0.54	0.82	BC005388	Hs.55055	FAAH	fatty acid hydroxylase
0.28	-0.03	0.31	BC010147	Hs.389107	MGC21874	hypothetical protein MGC21874
0.28	0.08	0.20	BC017270	Hs.250175	UAP1	UDP-N-acetylglucosamine pyrophosphorylase 1
0.28	-0.88	1.16	BC006849	Hs.19574	MGC5469	THO complex 3
0.28	-1.07	1.35	BC016850	Hs.261023	MCM7	MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)
0.28	-0.22	0.49	BC013311	Hs.25682	ZAP128	peroxisomal long-chain acyl-coA thioesterase
0.28	-2.33	2.61	BC002568	Hs.4900	LOC57019	hypothetical protein LOC57019
0.28	-0.61	0.89	BC008982	Hs.2161	MAGEA9	melanoma antigen, family A, 9
0.28	0.06	0.21	BC009179	Hs.15277	UBE2D1	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)
0.28	-0.43	0.70	BC008380	Hs.286145	NMU	neuromedin U
0.28	0.14	0.14	BC002403	Hs.79339	CHI3L2	chitinase 3-like 2
0.28	-0.08	0.36	BC002905	Hs.24998	RPL39	ribosomal protein L39
0.28	-0.12	0.40	BC001493	Hs.241489	LOC90678	hypothetical protein BC009239
0.28	-0.27	0.54	BC005084	Hs.29874	QPRT	quinolate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating))
0.28	-0.65	0.92	BC011180	Hs.124942	SH3YL1	likely ortholog of mouse Sh3 domain YSC-like 1
0.27	-0.10	0.37	BC014014	Hs.169992	MCM3	MCM3 minichromosome maintenance deficient 3 (S. cerevisiae)
0.27	-0.16	0.43	BC009217	Hs.337557	CGI-10	coenzyme Q6 homolog (yeast)
0.27	-0.49	0.76	BC012787	Hs.180895	SIMP	source of immunodominant MHC-associated peptides
0.27	0.02	0.25	BC003000	Hs.79748	BIRC5	baculoviral IAP repeat-containing 5 (survivin)

0.27	-0.87	1.14	BC015831	Hs.88646	COG4	component of oligomeric golgi complex 4
0.27	0.06	0.21	BC009078	Data not found	LOC136895	hypothetical protein BC015397
0.27	0.18	0.09	BC011981	Hs.110407	COL4A6	collagen, type IV, alpha 6
0.27	-0.60	0.87	BC009428	Hs.423968	MGC10200	hypothetical protein MGC:10200
0.27	-0.36	0.63	BC005271	Hs.19236	NAT1	N-acetyltransferase 1 (arylamine N-acetyltransferase)
0.27	-0.32	0.59	BC014946	Hs.321653	VIT1	
0.27	-0.69	0.96	BC004553	Hs.103755	DKFZP566C243	DKFZP566C243 protein
0.27	-0.56	0.83	BC001023	Hs.318720	HSPC219	Nef associated protein 1
0.27	-0.89	1.16	BC011566	Hs.198298	POU2F1	POU domain, class 2, transcription factor 1
0.27	-0.28	0.55	BC000306	Hs.8110	ATP6V1E2	ATPase, H <sup>+</sup> transporting, lysosomal 31kDa, V1 subunit E isoform 2
0.27	-1.19	1.46	BC001635	Hs.429621	PPP1CC	protein phosphatase 1, catalytic subunit, gamma isoform
0.27	-0.85	1.12	BC017193	Hs.283655	CCND1	cyclin D1 (PRAD1: parathyroid adenomatosis 1)
0.27	-0.88	1.15	BC000052	Hs.998	C1orf15	chromosome 1 open reading frame 15
0.27	-0.83	1.10	BC004561	Hs.81361	AMACR	alpha-methylacyl-CoA racemase
0.27	-0.72	0.99	BC014785	Hs.692	NME7	non-metastatic cells 7, protein expressed in (nucleoside-diphosphate kinase)
0.27	-0.05	0.32	BC011579	Hs.321214	SERPINA3	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 3
0.27	-1.03	1.30	BC010708	Hs.241503	384D8-2	hypothetical protein 384D8_6
0.27	-0.41	0.68	BC001899	Hs.46328	PMPCB	peptidase (mitochondrial processing) beta
0.27	-0.77	1.04	BC005312	Hs.318720	STAF42	SPT3-associated factor 42
0.27	-0.47	0.73	BC010221	Hs.16986	STRAIT11499	hypothetical protein STRAIT11499
0.27	-1.20	1.46	BC011375	Hs.23106	RGS19IP1	regulator of G-protein signalling 19 interacting protein 1
0.27	-0.82	1.08	BC007873	Hs.278426		Homo sapiens, clone IMAGE:4829271, mRNA
0.27	-0.42	0.68	BC005002	Hs.152738	ZNF358	zinc finger protein 358
0.27	-0.90	1.16	BC008198	Hs.129445	SLC30A5	solute carrier family 30 (zinc transporter), member 5
0.27	-0.34	0.61	BC002791	Hs.356137	FLJ13855	hypothetical protein FLJ13855
0.27	0.14	0.13	BC012741	Hs.7104	FLJ12783	ionized calcium binding adapter molecule 2
0.27	-0.22	0.49	BC012797	Hs.16232	BIN3	bridging integrator 3
0.27	0.23	0.04	BC013327	Hs.17287	KCNJ15	potassium inwardly-rectifying channel, subfamily J, member 15
0.27	0.09	0.17	BC000165	Hs.78902	VDAC2	voltage-dependent anion channel 2
0.26	-0.45	0.71	BC012191	Hs.375206	MRPS12	mitochondrial ribosomal protein S12
0.26	-0.09	0.35	BC015764	Hs.25895	NA 4-14-03 8	
0.26	-1.81	2.08	NM_001715	Hs.2243	VDAC3	voltage-dependent anion channel 3
0.26	0.16	0.10	BC005349	Hs.2265	BDH	3-hydroxybutyrate dehydrogenase (heart, mitochondrial)
0.26	-1.08	1.34	BC008608	Hs.14144	ARD1	ARD1 homolog, N-acetyltransferase ( <i>S. cerevisiae</i> )
0.26	-1.12	1.38	BC003097	Hs.433303	MDS032	uncharacterized hematopoietic stem/progenitor cells protein MDS032
0.26	-0.52	0.78	BC015562	Hs.94814	BAG3	BCL2-associated athanogene 3
0.26	-0.60	0.86	BC001934	Hs.73818	ZNF232	zinc finger protein 232
0.26	0.11	0.16	BC019235	Hs.153261	EMD	emerin (Emery-Dreifuss muscular dystrophy)
0.26	0.23	0.04	BC000437	Hs.90744	KRT8	keratin 8
0.26	-0.99	1.26	BC007061	Hs.1340	NA 4-14-03 235	
0.26	-1.08	1.34	BC010846	Hs.16229	MGC4170	MGC4170 protein
0.26	0.18	0.09	BC013002	Hs.73999	EXTL3	exostoses (multiple)-like 3
0.26	-0.88	1.14	BC016148	Hs.239625	FAAH	fatty acid hydroxylase
0.26	0.06	0.21	BC016927	Hs.12835	MGC17337	similar to RIKEN cDNA 5730528L13 gene
0.26	-1.02	1.28	BC000887	Hs.26813	PGD	phosphogluconate dehydrogenase
0.26	-1.81	2.07	BC002787	Hs.227782	LOC147632	hypothetical protein BC010734
0.26	-0.21	0.47	BC005843	Hs.30002	CINP	cyclin-dependent kinase 2-interacting protein
0.26	0.18	0.08	BC016452	Hs.6780	NA 4-14-03 395	
0.26	-0.76	1.02	BC007726	Hs.26516	DKFZP586M1523	DKFZP586M1523 protein
0.26	-0.35	0.61	BC012398	Data not found	DKFZP564G2022	DKFZP564G2022 protein
0.26	-0.95	1.21	BC015883	Hs.334842	WAC	WW domain-containing adapter with a coiled-coil region
0.26	0.25	0.01	empty	empty	empty	empty



0.26	0.10	0.16	BC000392	Hs.75139	SYT11	synaptotagmin XI
0.26	0.15	0.10	BC012539	Hs.27289	CGI-125	CGI-125 protein
0.26	-0.36	0.62	BC001019	Hs.300141	MGC10471	hypothetical protein MGC10471
0.26	-0.11	0.37	BC011006	Hs.9527	3-Apr	apoptosis related protein APR-3
0.26	-0.36	0.62	BC011747	Hs.306019	FLJ14466	hypothetical protein FLJ14466
0.26	-0.95	1.21	BC008403	Hs.308026	ASB6	ankyrin repeat and SOCS box-containing 6
0.26	-0.37	0.63	BC004390	Hs.77329	HN1	hematological and neurological expressed 1
0.26	-1.82	2.08	BC014477	Hs.83465	NA 4-14-03 449	
0.26	0.00	0.26	BC000046	Hs.108287	KIAA0121	KIAA0121 gene product
0.26	-0.31	0.56	BC016057	Hs.132945	MGC16384	hypothetical protein MGC16384
0.26	-0.10	0.36	BC001765	Hs.8217	ETV5	ets variant gene 5 (ets-related molecule)
0.26	-0.27	0.53	BC011706	Hs.124005	FLJ11259	hypothetical protein FLJ11259
0.26	-0.44	0.69	BC005404	Hs.173497	SEC23B	Sec23 homolog B (S. cerevisiae)
0.26	-0.20	0.45	BC008719	Hs.173980	CDC20	CDC20 cell division cycle 20 homolog (S. cerevisiae)
0.26	0.19	0.06	BC016720	Hs.128051	CRMP1	collapsin response mediator protein 1
0.26	-0.60	0.85	BC003569	Data not found	NME1	non-metastatic cells 1, protein (NM23A) expressed in
0.26	-0.62	0.87	BC002655	Hs.26322	POH1	26S proteasome-associated pad1 homolog
0.25	-1.93	2.18	BC015541	Hs.159546	KIAA0431	KIAA0431 protein
0.25	-0.93	1.18	BC000809	Hs.95243	TCEAL1	transcription elongation factor A (SII)-like 1
0.25	0.16	0.09	BC009508	Hs.432419	GARS	glycyl-tRNA synthetase
0.25	-0.34	0.60	BC000376	Hs.173518	ZFR	zinc finger RNA binding protein
0.25	-0.63	0.89	BC010894	Hs.5324	CL25022	hypothetical protein CL25022
0.25	0.06	0.19	BC017487	Hs.51299	RNF25	ring finger protein 25
0.25	-0.35	0.60	BC008477	Hs.169248	FLJ10759	hypothetical protein FLJ10759
0.25	-1.47	1.73	BC007333	Hs.43697	STAG2	stromal antigen 2
0.25	-0.23	0.49	BC021229	Hs.127243	KIAA1724	KIAA1724 protein
0.25	-0.01	0.26	BC005172	Hs.274336	ALDH3B1	aldehyde dehydrogenase 3 family, member B1
0.25	0.24	0.01	Data not found		KIAA0152	KIAA0152 gene product
0.25	-0.19	0.44	BC007382	Hs.55405	P5326	hypothetical protein p5326
0.25	-1.38	1.63	BC000866	Hs.5831	TIMP1	tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor)
0.25	-0.25	0.50	BC012891	Hs.422118	TGFA	transforming growth factor, alpha
0.25	-0.05	0.30	BC007415	Hs.433334	MRPS7	mitochondrial ribosomal protein S7
0.25	-0.15	0.40	BC002801	Hs.12865	FLJ10545	hypothetical protein FLJ10545
0.25	-0.57	0.81	BC006195	Hs.174140	FLJ20519	hypothetical protein FLJ20519
0.25	-0.19	0.44	BC002810	Hs.13128	RB1CC1	RB1-inducible coiled-coil 1
0.25	-0.55	0.80	BC008984	Hs.406462	OSR2	odd-skipped-related 2A protein
0.25	-0.48	0.73	BC020988	Hs.37747	INPP4B	inositol polyphosphate-4-phosphatase, type II, 105kDa
0.25	-0.67	0.92	BC017370	Hs.279919	RBX1	ring-box 1
0.25	-0.49	0.73	BC012764	Hs.11056	POLK	polymerase (DNA directed) kappa
0.25	0.14	0.10	BC010130	Hs.289112	HSPA8	heat shock 70kDa protein 8
0.25	-0.75	0.99	BC004288	Hs.115659	FYN	FYN oncogene related to SRC, FGR, YES
0.25	-1.63	1.88	BC004380	Hs.15159	SDF2	stromal cell-derived factor 2
0.25	-0.37	0.62	BC017053	Hs.12773	GNB2	guanine nucleotide binding protein (G protein), beta polypeptide 2
0.25	0.12	0.12	BC001399	Hs.418650	FTH1	ferritin, heavy polypeptide 1
0.25	-0.58	0.83	BC004550	Hs.272801	HGS	
0.25	0.18	0.07	BC007423	Hs.76927	TOMM34	translocase of outer mitochondrial membrane 34
0.24	-0.59	0.83	BC002496	Hs.30376	HSPC194	hypothetical protein HSPC194
0.24	-1.33	1.58	BC001763	Hs.76927	TMEPAI	transmembrane, prostate androgen induced RNA
0.24	-0.24	0.49	BC002656	Hs.85092	IDS	iduronate 2-sulfatase (Hunter syndrome)
0.24	-0.29	0.53	BC015913	Hs.198726	CAPN3	calpain 3, (p94)
0.24	-0.11	0.36	BC004180	Hs.307016	KRTAP4-12	keratin associated protein 4-12
0.24	-0.27	0.51	BC012737	Hs.80768	TRIM29	tripartite motif-containing 29
0.24	-1.97	2.21	BC006428	Hs.15093	DKFZP434N014	DKFZP434N014 protein

0.24	-0.20	0.44	BC004491	Hs.89862	MTCP1	mature T-cell proliferation 1
0.24	-0.05	0.29	BC004364	Hs.25544	WDR8	WD repeat domain 8
0.24	-0.26	0.51	BC015405	Hs.356019	TAZ	transcriptional co-activator with PDZ-binding motif (TAZ)
0.24	-1.42	1.66	BC006793	Hs.169946	GATA3	GATA binding protein 3
0.24	-1.08	1.32	BC014901	Hs.10842	C20orf3	chromosome 20 open reading frame 3
0.24	-1.09	1.33	BC020568	Hs.132569	LOC51136	PTD016 protein
0.24	-0.94	1.18	BC021213	Hs.29595	SEC22C	SEC22 vesicle trafficking protein-like 3 (S. cerevisiae)
0.24	0.19	0.05	BC000496	Hs.168670	KRT18	keratin 18
0.24	-0.04	0.28	BC012984	Data not found	HINT3	histidine triad nucleotide binding protein 3
0.24	-1.26	1.50	BC000468	Hs.75875	SC4MOL	sterol-C4-methyl oxidase-like
0.24	-0.71	0.95	BC021275	Hs.409563	RAGA	Ras-related GTP-binding protein
0.24	-1.52	1.76	BC006101	Hs.1101	LAT1-3TM	LAT1-3TM protein
0.24	-0.39	0.62	BC001407	Hs.380359	SLC25A16	solute carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16
0.24	-0.11	0.35	BC007555	Hs.433506	MGC3077	hypothetical protein MGC3077
0.24	-0.47	0.71	BC012033	Hs.164989	TBCE	tubulin-specific chaperone e
0.24	-0.07	0.31	BC002898	Hs.50748	LEPROTL1	leptin receptor overlapping transcript-like 1
0.24	-0.24	0.48	BC003182	Hs.772	MGC20255	hypothetical protein MGC20255
0.23	-0.22	0.46	BC015482	Hs.62771	MIZF	MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein
0.23	-0.67	0.90	BC016172	Hs.92023	NA 4-14-03 310	
0.23	-0.53	0.77	BC010025	Hs.356742	NA 4-14-03 247	
0.23	0.12	0.11	BC000794	Hs.155244	PRPF18	PRP18 pre-mRNA processing factor 18 homolog (yeast)
0.23	-0.46	0.70	BC001620	Hs.169609	FLJ21168	hypothetical protein FLJ21168
0.23	-0.48	0.71	BC010893	Hs.279761	HSPC134	HSPC134 protein
0.23	-1.16	1.40	BC002614	Hs.77643	NME4	non-metastatic cells 4, protein expressed in
0.23	-0.31	0.55	BC002542	Hs.23413	KIAA1273	KIAA1273 protein
0.23	-0.68	0.92	BC011963	Hs.432876	SGT	small glutamine-rich tetratricopeptide repeat (TPR)-containing
0.23	0.05	0.18	BC019042	Hs.71746	MGC14151	hypothetical protein MGC14151
0.23	-0.88	1.11	BC007949	Hs.256184	LOC115548	hypothetical protein BC014311
0.23	-0.19	0.42	BC008386	Hs.43104	CYC1	cytochrome c-1
0.23	-1.09	1.33	BC008733	Hs.154583	RBM10	RNA binding motif protein 10
0.23	-0.57	0.80	BC005902	Hs.81029	KIAA0410	nucleoporin like 1
0.23	0.04	0.19	BC007407	Hs.83321	NME2	non-metastatic cells 2, protein (NM23B) expressed in
0.23	-1.41	1.65	Data not found		FRAT2	frequently rearranged in advanced T-cell lymphomas 2
0.23	-1.88	2.11	BC011744	Hs.417816	OSBPL3	oxysterol binding protein-like 3
0.23	-0.39	0.62	BC014480	Data not found	IGFBP2	insulin-like growth factor binding protein 2, 36kDa
0.23	-0.45	0.68	BC000019	Hs.32963	KIAA0841	KIAA0841 protein
0.23	-0.77	1.00	BC017660	Hs.109798	C6orf48	chromosome 6 open reading frame 48
0.23	0.06	0.17	BC014253	Hs.5672	BDG-29	BDG-29 proten
0.23	-2.05	2.28	BC015647	Hs.352221	C16orf44	chromosome 16 open reading frame 44
0.23	-1.05	1.28	BC006435	Hs.423216	DRD2	dopamine receptor D2
0.23	-1.40	1.63	BC002677	Hs.10700	FLJ20139	hypothetical protein FLJ20139
0.23	-0.46	0.68	BC013788	Hs.48712	BASP1	brain abundant, membrane attached signal protein 1
0.23	-0.94	1.17	BC013961	Hs.171075	C11orf23	chromosome 11 open reading frame 23
0.23	-0.15	0.38	BC005237	Hs.107942	KRT18	keratin 18
0.23	-0.14	0.36	BC014791	Hs.25723	EPB41L3	erythrocyte membrane protein band 4.1-like 3
0.23	-0.47	0.69	BC016654	Hs.10842	NA 4-14-03 481	
0.23	-0.77	0.99	BC006191	Hs.79672	KIAA0652	KIAA0652 gene product
0.23	0.13	0.10	BC014178	Hs.154868	FLJ23441	hypothetical protein FLJ23441
0.23	-0.20	0.42	BC003540	Hs.423968	GLO1	glyoxalase I
0.23	-2.32	2.54	BC014640	Hs.403836	COL14A1	collagen, type XIV, alpha 1 (undulin)
0.23	0.06	0.16	BC016961	Data not found	LOC51668	HSPCO34 protein
0.23	-0.46	0.68	BC010469	Hs.194673	UBC	ubiquitin C

0.23	-1.06	1.28	BC006808	Hs.101660	MGC5391	hypothetical protein MGC5391
0.22	-2.06	2.28	BC008286	Hs.181046	DUSP3	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)
0.22	0.17	0.05	BC002740	Hs.288991	ACTA2	actin, alpha 2, smooth muscle, aorta
0.22	0.12	0.11	Data not found	CA8	carbonic anhydrase VIII	
0.22	-0.06	0.28	BC010896	Data not found	NA 4-14-03 436	
0.22	-0.01	0.23	BC009357	Hs.406504	RAI16	retinoic acid induced 16
0.22	-0.15	0.37	BC015153	Hs.263812	CDC42EP1	CDC42 effector protein (Rho GTPase binding) 1
0.22	-1.85	2.07	BC016668	Hs.110950	CKIP-1	CK2 interacting protein 1; HQ0024c protein
0.22	-0.14	0.36	BC007453	Hs.4990	KIAA1089	KIAA1089 protein
0.22	-0.53	0.75	BC007862	Hs.167641	NA 4-14-03 515	
0.22	0.19	0.03	BC008469	Hs.28081	MLYCD	malonyl-CoA decarboxylase
0.22	-0.95	1.17	BC000731	Hs.6139	AHCY	S-adenosylhomocysteine hydrolase
0.22	-0.20	0.42	BC017050	Hs.198365	NA 4-14-03 25	
0.22	-0.89	1.11	BC002681	Hs.87435	C22orf5	chromosome 22 open reading frame 5
0.22	-0.93	1.15	BC011832	Hs.274248	ATPIF1	ATPase inhibitory factor 1
0.22	0.18	0.04	BC011377	Hs.155650	GADD45G	growth arrest and DNA-damage-inducible, gamma
0.22	-0.34	0.56	BC012062	Hs.65588	POLR2F	polymerase (RNA) II (DNA directed) polypeptide F
0.22	-0.02	0.24	BC011834	Hs.282410	KIAA1961	KIAA1961 protein
0.22	-0.06	0.27	BC003138	Hs.344037	PQBP1	polyglutamine binding protein 1
0.22	-1.97	2.19	BC009898	Hs.89718	PFN2	profilin 2
0.21	-0.19	0.41	BC003672	Hs.391561	FLJ20635	hypothetical protein FLJ20635
0.21	0.06	0.15	BC000169	Hs.155020	WBSR22	Williams Beuren syndrome chromosome region 22
0.21	0.05	0.16	BC018646	Hs.75648	RPL36AL	ribosomal protein L36a-like
0.21	0.17	0.05	BC002597	Hs.239720	KIAA0582	KIAA0582 protein
0.21	0.04	0.18	BC006454	Hs.226133	GAS7	growth arrest-specific 7
0.21	-0.30	0.51	BC009765	Hs.169992	SDBCAG84	serologically defined breast cancer antigen 84
0.21	-0.22	0.43	BC009991	Hs.339837	FLJ20371	chromosome 14 open reading frame 58
0.21	-0.38	0.59	BC000026	Hs.209119	CDC2	cell division cycle 2, G1 to S and G2 to M
0.21	-0.37	0.58	NM_005505	Hs.180616	LOC146542	similar to hypothetical protein MGC13138
0.21	-0.90	1.12	NM_152221	Hs.79658	TXNRD2	thioredoxin reductase 2
0.21	-0.42	0.64	BC007230	Hs.21016		
0.21	-0.21	0.42	BC017321	Data not found	CSNK2A1	casein kinase 2, alpha 1 polypeptide
0.21	-0.62	0.83	BC008726	Hs.95220	OSR1	oxidative-stress responsive 1
0.21	-0.39	0.60	BC012349	Hs.1973	FLJ22688	hypothetical protein FLJ22688
0.21	-0.24	0.45	BC003409	Hs.132071	OCIA	ovarian carcinoma immunoreactive antigen
0.21	0.12	0.09	BC002400	Hs.132760	GFAP	glial fibrillary acidic protein
0.21	-0.75	0.96	BC011574	Hs.78713	BAG1	BCL2-associated athanogene
0.21	-0.31	0.51	BC006383	Hs.4742	CGI-57	hypothetical protein CGI-57
0.21	-0.30	0.51	BC001142	Hs.166887	MAGP2	Microfibril-associated glycoprotein-2
0.21	-0.43	0.64	BC012348	Hs.91299	VAMP3	vesicle-associated membrane protein 3 (cellubrevin)
0.21	0.02	0.18	BC018953	Hs.250581	MIZF	MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein
0.21	-0.65	0.85	BC003100	Hs.1697	MGC14799	hypothetical protein MGC14799
0.21	-0.29	0.49	BC014213	Hs.159505	RNF34	ring finger protein 34
0.20	-1.63	1.84	BC000264	Hs.250570	HCS	cytochrome c, somatic
0.20	0.11	0.10	BC000530	Hs.426977	SLU7	step II splicing factor SLU7
0.20	-0.82	1.03	BC004381	Hs.95697	GCDH	glutaryl-Coenzyme A dehydrogenase
0.20	-0.56	0.76	BC009574	Hs.168289	SDHD	succinate dehydrogenase complex, subunit D, integral membrane protein
0.20	-0.13	0.34	BC009823	Hs.61809	IGKC	immunoglobulin kappa constant
0.20	-0.20	0.41	BC010067	Hs.84746	USP3	ubiquitin specific protease 3
0.20	-1.00	1.20	BC009691	Hs.183435	MKNK1	MAP kinase-interacting serine/threonine kinase 1
0.20	-0.11	0.31	BC015030	Hs.9933	FLJ20013	hypothetical protein FLJ20013
0.20	0.14	0.06	BC004274	Hs.143554	MRPL10	mitochondrial ribosomal protein L10
0.20	-1.38	1.58	BC014679	Hs.76917	CXCL11	chemokine (C-X-C motif) ligand 11

0.20	0.14	0.06	BC010073	Hs.91299	ACOX3	acyl-Coenzyme A oxidase 3, pristanoyl
0.20	-0.16	0.36	BC006319	Hs.56874	DKC1	dyskeratosis congenita 1, dyskerin
0.20	0.07	0.13	BC018951	Hs.241567	GNG10	guanine nucleotide binding protein (G protein), gamma 10
0.20	-0.62	0.82	BC003402	Hs.25516	FLJ10290	hypothetical protein FLJ10290
0.20	-0.47	0.67	BC016276	Hs.77695	NA 4-14-03 339	
0.20	-2.00	2.20	BC007321	Hs.273234	MRPS23	mitochondrial ribosomal protein S23
0.20	-0.98	1.18	BC014568	Data not found	FYCO1	FYVE and coiled-coil domain containing 1
0.20	-0.42	0.62	BC021892	Hs.323567	NSEPI	nuclease sensitive element binding protein 1
0.20	0.05	0.15	BC012098	Hs.1691	DNPEP	aspartyl aminopeptidase
0.20	0.13	0.07	BC013014	Hs.323213	RNF29	ring finger protein 29
0.20	-0.53	0.72	BC000285	Hs.274319	LOC55871	COBW-like protein
0.20	-0.04	0.24	BC002447	Hs.123637	PSMC3	proteasome (prosome, macropain) 26S subunit, ATPase, 3
0.20	-0.13	0.33	BC007250	Hs.10326	KIAA0103	KIAA0103 gene product
0.20	-0.17	0.37	BC003553	Hs.5148	SLC22A5	solute carrier family 22 (organic cation transporter), member 5
0.20	-0.14	0.34	BC014148	Hs.201677	MGC4659	hypothetical protein MGC4659
0.20	-0.21	0.41	BC001327	Hs.315177	IFRD2	interferon-related developmental regulator 2
0.20	-1.47	1.66	BC006551	Hs.76084	MAPK14	mitogen-activated protein kinase 14
0.20	-0.44	0.64	BC017594	Hs.104058	NA 4-14-03 88	
0.20	-0.40	0.60	BC001958	Hs.103561	FLJ10759	hypothetical protein FLJ10759
0.20	0.05	0.14	BC006363	Hs.9018	TRIP10	thyroid hormone receptor interactor 10
0.20	-0.74	0.94	Data not found		SNX3	sorting nexin 3
0.19	-0.11	0.30	BC007717	Hs.296365	PUM1	pumilio homolog 1 (Drosophila)
0.19	0.04	0.15	BC000894	Hs.9676	FASTK	FAST kinase
0.19	-1.35	1.54	BC012602	Hs.32970	SLAM	signaling lymphocytic activation molecule
0.19	-0.47	0.67	BC021998	Hs.421349	GFPT2	glutamine-fructose-6-phosphate transaminase 2
0.19	0.08	0.11	BC007833	Hs.149255	CHST8	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8
0.19	-0.05	0.24	BC008456	Hs.81029	MGC3162	hypothetical protein MGC3162
0.19	0.16	0.04	BC006486	Hs.429607	MOG1	homolog of yeast MOG1
0.19	-0.35	0.54	BC005891	Hs.123536	MGC3103	hypothetical protein MGC3103
0.19	-1.20	1.39	BC009963	Hs.3080	PEX10	peroxisome biogenesis factor 10
0.19	-0.24	0.43	BC011765	Hs.303025	UCHL5	ubiquitin carboxyl-terminal hydrolase L5
0.19	-0.23	0.42	BC004902	Hs.5070	PEX11A	peroxisomal biogenesis factor 11A
0.19	-1.41	1.61	BC010913	Hs.283108	HBG1	hemoglobin, gamma A
0.19	-0.88	1.07	BC008679	Hs.33922	GABPB2	GA binding protein transcription factor, beta subunit 2, 47kDa
0.19	-1.48	1.67	BC011959	Hs.317589	ARHI	ras homolog gene family, member 1
0.19	-1.58	1.77	BC013923	Hs.816	HSGT1	suppressor of S. cerevisiae gcr2
0.19	-0.60	0.79	BC002734	Hs.251573	HEBP1	heme binding protein 1
0.19	-0.06	0.25	BC013436	Hs.39619	MGC2494	hypothetical protein MGC2494
0.19	0.12	0.07	BC013069	Hs.46634	PRO2389	WD repeat domain 23
0.19	0.00	0.19	BC008771	Hs.157199	TFRC	transferrin receptor (p90, CD71)
0.19	-1.18	1.37	BC000837	Hs.351327	FLJ20452	hypothetical protein FLJ20452
0.19	-1.92	2.10	BC017072	Hs.332382	TTR	transthyretin (prealbumin, amyloidosis type I)
0.19	-0.35	0.54	BC007978	Hs.6449	TREX2	three prime repair exonuclease 2
0.19	-1.16	1.34	BC015534	Hs.16258	CSNK1G1	casein kinase 1, gamma 1
0.19	-0.10	0.28	BC016824	Hs.355719	CSR2BP	CSR2 binding protein
0.19	-2.02	2.21	BC002687	Hs.24563	NA 4-14-03 31	
0.19	0.17	0.02	BC012865	Hs.299148	TAGLN2	transgelin 2
0.19	-0.02	0.21	BC021667	Hs.154387	COPE	coatamer protein complex, subunit epsilon
0.19	-0.90	1.08	BC015632	Hs.236547	GMMPR2	guanosine monophosphate reductase 2
0.19	-1.18	1.37	BC016043	Hs.94814	C6.1A	c6.1A
0.19	0.06	0.13	BC002469	Hs.79457	ACLY	ATP citrate lyase
0.19	-0.11	0.29	BC009733	Hs.422118	MGC4840	hypothetical protein MGC4840
0.19	-1.32	1.51	BC001204	Hs.309161	UCP3	uncoupling protein 3 (mitochondrial, proton carrier)

0.18	-1.23	1.42	BC015722	Hs.23241	CERK	ceramide kinase
0.18	-0.67	0.86	BC009957	Hs.20157	TSG	twisted gastrulation homolog 1 (Drosophila)
0.18	-0.50	0.69	BC008472	Hs.82159	EIF3S9	eukaryotic translation initiation factor 3, subunit 9 eta, 116kDa
0.18	-0.77	0.95	BC000836	Hs.184542	CGI-127	yippee protein
0.18	-0.83	1.02	BC002489	Hs.85100	WDR1	WD repeat domain 1
0.18	-0.48	0.66	BC009568	Hs.6467	MGC13090	hypothetical protein MGC13090
0.18	-1.68	1.87	BC007705	Hs.31439	LOC51042	zinc finger protein
0.18	-3.97	4.15	BC004290	Hs.282960	CASP7	caspase 7, apoptosis-related cysteine protease
0.18	-1.13	1.32	BC013306	Hs.172772	LOC51204	clone HQ0477 PRO0477p
0.18	-0.36	0.54	BC000526	Hs.3447	PX19	px19-like protein
0.18	0.13	0.05	BC008320	Hs.146393	RPE	ribulose-5-phosphate-3-epimerase
0.18	-0.79	0.97	BC013996	Hs.6980	C12orf4	chromosome 12 open reading frame 4
0.18	-0.64	0.82	BC006271	Data not found	AGPAT2	1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
0.18	-0.02	0.20	BC015548	Data not found	PIN4	protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)
0.18	0.09	0.09	BC017378	Hs.51043	ARS2	arsenate resistance protein ARS2
0.18	-5.08	5.26	BC009763	Hs.83761	PPIL2	peptidylprolyl isomerase (cyclophilin)-like 2
0.18	-0.34	0.52	BC014320	Hs.240767	ETV4	ets variant gene 4 (E1A enhancer binding protein, E1AF)
0.18	0.09	0.09	BC002619	Hs.70669	MGC11061	hypothetical protein MGC11061
0.18	-1.59	1.77	BC004817	Hs.129953	EWSR1	Ewing sarcoma breakpoint region 1
0.18	0.13	0.05	BC010218	Hs.262823	GK001	GK001 protein
0.18	-0.23	0.40	BC013705	Hs.23744	RISC	likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase
0.18	-0.34	0.52	BC001177	Hs.271687	KIAA0010	ubiquitin-protein isopeptide ligase (E3)
0.18	-0.43	0.61	BC008863	Hs.5409	REA	repressor of estrogen receptor activity
0.18	-0.05	0.23	BC008831	Hs.429777	PHB	prohibitin
0.18	0.11	0.07	BC000398	Hs.93354	EXO70	likely ortholog of mouse exocyst component protein 70 kDa homolog (S. cerevisiae) Exo70: exocyst component protein 70 kDa homolog (S. cerevisiae)
0.18	-0.24	0.41	BC005249	Hs.9222	MLC1SA	myosin light chain 1 slow a
0.18	-1.79	1.97	BC007403	Hs.82890	GMDS	GDP-mannose 4,6-dehydratase
0.17	-0.83	1.01	BC018926	Hs.279746	AKAP12	A kinase (PRKA) anchor protein (gravin) 12
0.17	-0.40	0.57	BC016767	Hs.30724	MGC4796	hypothetical protein MGC4796
0.17	-0.49	0.66	BC015563	Hs.179915	ZNF339	zinc finger protein 339
0.17	0.14	0.03	BC001346	Hs.22142	CYB5R2	cytochrome b5 reductase b5R.2
0.17	0.12	0.06	BC007279	Hs.5486	UBE2V2	ubiquitin-conjugating enzyme E2 variant 2
0.17	-0.10	0.28	BC003135	Hs.157	SPN	sialophorin (gpL115, leukosialin, CD43)
0.17	-1.80	1.97	BC000793	Hs.4310	EIF1A	eukaryotic translation initiation factor 1A
0.17	-0.39	0.56	BC013571	Hs.289043	SDR1	short-chain dehydrogenase/reductase 1
0.17	-0.77	0.95	BC015711	Hs.8261	MGC10818	hypothetical protein MGC10818
0.17	-0.11	0.28	BC011755	Hs.5198	E2F3	E2F transcription factor 3
0.17	-0.16	0.34	BC018938	Hs.284295	NSE1	non-SMC (structural maintenance of chromosomes) element 1 protein
0.17	-0.15	0.32	BC009363	Hs.285005	DEC2	2,4-dienoyl CoA reductase 2, peroxisomal
0.17	-1.38	1.55	BC012303	Hs.271473	DKFZP434O1427	dactylidin
0.17	-0.75	0.92	BC007008	Hs.408767	SPINT2	serine protease inhibitor, Kunitz type, 2
0.17	-0.52	0.69	BC012384	Hs.80019	PSMC6	proteasome (prosome, macropain) 26S subunit, ATPase, 6
0.17	-0.62	0.79	BC001211	Hs.23131	CAPNS2	calpain small subunit 2
0.17	0.05	0.12	BC001165	Hs.75932	IGFBP2	insulin-like growth factor binding protein 2, 36kDa
0.17	-1.04	1.21	BC016641	Hs.61628	FLJ40452	hypothetical protein FLJ40452
0.17	-0.15	0.32	BC007814	Hs.277477	MRPL51	mitochondrial ribosomal protein L51
0.17	-0.80	0.97	BC015602	Hs.302131	LOC57228	hypothetical protein from clone 643
0.17	-0.93	1.10	BC000875	Hs.279609	ZNF9	zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)
0.17	-0.86	1.03	BC018543	Hs.108947	REC14	recombination protein REC14
0.17	-0.37	0.54	BC008631	Hs.334913	RBM4	RNA binding motif protein 4
0.17	0.02	0.15	BF797140	Hs.446643	NA 4-14-03 147	

0.17	-0.93	1.10	BC015039	Hs.389137	LOC90346	hypothetical protein BC004930
0.17	-1.19	1.36	BC001429	Hs.300711	CITED2	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
0.17	-1.83	1.99	BC008882	Hs.288672	CDC42	cell division cycle 42 (GTP binding protein, 25kDa)
0.17	-1.85	2.02	BC009702	Hs.349177	HSPC132	hypothetical protein HSPC132
0.16	-0.82	0.98	BC009297	Data not found	FLJ21128	hypothetical protein FLJ21128
0.16	0.10	0.06	BC016759	Hs.77054	SLC3A2	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
0.16	-0.04	0.21	BC011778	Hs.11039	DDX30	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 30
0.16	-0.43	0.59	BC012616	Hs.80756	BHMT	betaine-homocysteine methyltransferase
0.16	-0.59	0.75	BC008442	Hs.351316	AP1S1	adaptor-related protein complex 1, sigma 1 subunit
0.16	-1.28	1.44	BC005009	Hs.271599	UREB1	upstream regulatory element binding protein 1
0.16	-1.72	1.88	BC011020	Hs.152720	MPHOSPH6	M-phase phosphoprotein 6
0.16	-0.33	0.49	BC021276	Hs.413826	HSPA8	heat shock 70kDa protein 8
0.16	-1.79	1.95	BC014454	Hs.211581	LOC92979	hypothetical protein BC009489
0.16	-0.27	0.43	BC004302	Hs.6671	TOMM34	translocase of outer mitochondrial membrane 34
0.16	-0.12	0.28	BC008475	Hs.169248	MGC3180	hypothetical protein MGC3180
0.16	-0.12	0.28	BC015858	Hs.54488	LOC113521	similar to avenaII
0.16	-0.43	0.59	BC007340	Hs.106880	GDAP1L1	ganglioside-induced differentiation-associated protein 1-like 1
0.16	-0.83	0.99	BC006328	Hs.279946	LOC114926	hypothetical protein BC013035
0.16	-0.83	0.99	BC015727	Hs.235782	PAGE-5	PAGE-5 protein
0.16	-1.43	1.59	BC002968	Hs.182595	MYCL1	v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)
0.16	-0.08	0.24	BC006367	Hs.118695	KIAA0638	KIAA0638 protein
0.16	-0.21	0.36	BC001367	Hs.84162	HUMAGCGB	chromosome 3p21.1 gene sequence
0.16	-0.21	0.36	BC000739	Hs.432574	NA 4-14-03 48	
0.16	-0.65	0.81	BC018302	Hs.158947	SPHK1	sphingosine kinase 1
0.16	-1.43	1.59	BC004955	Data not found	CGBP	CpG binding protein
0.16	-1.04	1.19	BC011640	Hs.6770	SNX14	sorting nexin 14
0.16	-0.31	0.46	BC004449	Hs.78881	HES2	hairly and enhancer of split (Drosophila) homolog 2
0.16	-2.05	2.21	BC015651	Hs.125379	FLJ21820	hypothetical protein FLJ21820
0.15	-0.88	1.03	BC001222	Hs.15921	HCS	cytochrome c, somatic
0.15	-1.61	1.77	BC007991	Hs.6686	DKFZP564O1664	hypothetical protein DKFZp564O1664
0.15	-0.13	0.29	BC011568	Hs.297875	NFKBIA	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha
0.15	-0.11	0.26	BC014230	Data not found	PPP1R16A	protein phosphatase 1, regulatory (inhibitor) subunit 16A
0.15	-0.16	0.31	BC008933	Hs.433307	BCKDHA	branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease)
0.15	-2.00	2.15	BC004261	Hs.130316	SNAI2	snail homolog 2 (Drosophila)
0.15	-1.63	1.78	BC002628	Hs.71622	Dlc2	dynein light chain 2
0.15	-0.04	0.19	BC004861	Hs.128231	LOC51234	hypothetical protein LOC51234
0.15	-0.96	1.11	BC010623	Hs.83469	MARS	methionine-tRNA synthetase
0.15	-0.80	0.95	BC000807	Hs.206882	KR18	KRAB zinc finger protein KR18
0.15	-0.30	0.45	BC001239	Hs.5268	CD63	CD63 antigen (melanoma 1 antigen)
0.15	-0.77	0.92	BC019291	Hs.73722	PCOLCE2	procollagen C-endopeptidase enhancer 2
0.15	-0.04	0.19	BC004310	Hs.240534	FLJ20220	hypothetical protein FLJ20220
0.15	-0.07	0.22	BC000891	Hs.18349	MRPL15	mitochondrial ribosomal protein L15
0.15	0.14	0.01	BC006997	Hs.220689	FLJ22386	hypothetical protein FLJ22386
0.15	-0.77	0.92	BC011549	Hs.56155	MGAT1	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
0.15	-0.37	0.52	BC001800	Hs.98258	CSRP3	cysteine and glycine-rich protein 3 (cardiac LIM protein)
0.15	-0.87	1.02	BC009770	Hs.3576	C20orf30	chromosome 20 open reading frame 30
0.15	-1.44	1.59	BC001793	Hs.406503	C1orf2	chromosome 1 open reading frame 2
0.15	-0.14	0.28	BC008769	Hs.417068	ALG12	dolichyl-p-mannose:Man7GlcNAc2-PP-dolichyl mannosyltransferase
0.15	-0.10	0.25	BC011792	Hs.196176	IFI27	interferon, alpha-inducible protein 27
0.15	-0.21	0.36	BC001725	Hs.47313	BACH	brain acyl-CoA hydrolase
0.15	-0.53	0.67	BC014664	Hs.347991	GTF2H4	general transcription factor IIH, polypeptide 4, 52kDa
0.15	-0.94	1.09	BC004967	Hs.9194	NA 4-14-03 99	

0.15	-1.43	1.57	BC000933	Hs.75253	IDH3G	isocitrate dehydrogenase 3 (NAD+) gamma
0.15	-0.74	0.88	BC014123	Hs.288959		
0.14	0.06	0.09	BC009055	Hs.20537	LFNG	lunatic fringe homolog (Drosophila)
0.14	-0.06	0.20	BC005141	Hs.129228	DGCR14	DiGeorge syndrome critical region gene 14
0.14	-1.22	1.36	BC001042	Hs.19306	SIRT3	sirtuin (silent mating type information regulation 2 homolog) 3 (S. cerevisiae)
0.14	-0.21	0.35	BC006272	Hs.82906	C20orf4	chromosome 20 open reading frame 4
0.14	-0.55	0.69	BC001292	Hs.20017	C22orf4	chromosome 22 open reading frame 4
0.14	-2.99	3.13	BC001012	Hs.279916	DKFZP566B183	DKFZP566B183 protein
0.14	-0.28	0.42	BC016946	Hs.11039	TXNDC4	thioredoxin domain containing 4 (endoplasmic reticulum)
0.14	-1.51	1.65	BC011949	Hs.155097	REG1A	regenerating islet-derived 1 alpha (pancreatic stone protein, pancreatic thread protein)
0.14	-1.10	1.24	BC009747	Hs.334684	MGC13005	hypothetical protein MGC13005
0.14	-0.48	0.63	BC001643	Hs.76090	TNFAIP1	tumor necrosis factor, alpha-induced protein 1 (endothelial)
0.14	-0.36	0.50	NM_005488	Hs.9482	JAM1	F11 receptor
0.14	-0.76	0.91	BC006242	Hs.98798	PYCR1	pyrroline-5-carboxylate reductase 1
0.14	-0.62	0.76	BC013044	Hs.21189	MGC11287	ribosomal protein S6 kinase-like 1
0.14	-0.07	0.21	BC009596	Hs.381154	TPMT	thiopurine S-methyltransferase
0.14	-0.02	0.16	BC017441	Hs.129445	SLC30A5	solute carrier family 30 (zinc transporter), member 5
0.14	-0.85	0.99	BC008282	Hs.97858	SH3BP1	SH3-domain binding protein 1
0.14	-0.37	0.51	BC005280	Hs.514	C20orf72	chromosome 20 open reading frame 72
0.14	0.13	0.01	BC016944	Hs.48855	IL18	interleukin 18 (interferon-gamma-inducing factor)
0.14	-0.39	0.53	BC015928	Hs.24684	CBR3	carbonyl reductase 3
0.14	-1.11	1.25	BC005149	Hs.12909	G1P3	interferon, alpha-inducible protein (clone IFI-6-16)
0.14	-0.17	0.30	BC003154	Hs.236218	PCSK5	proprotein convertase subtilisin/kexin type 5
0.14	-0.67	0.81	BC001628	Hs.14394	WDF2	WD40 and FYVE domain containing 2
0.14	-0.01	0.15	BC004463	Hs.107382	DJ667H12.2	hypothetical protein DJ667H12.2
0.14	-0.13	0.27	BC000555	Hs.296290	MGC11102	hypothetical protein MGC11102
0.14	-0.62	0.76	BC013017	Hs.433996	EPB49	erythrocyte membrane protein band 4.9 (dematin)
0.14	-0.01	0.15	BC021968	Hs.289072	RYBP	RING1 and YY1 binding protein
0.14	-0.88	1.02	BC014515	Hs.94300	AXIN2	axin 2 (conductin, axil)
0.14	-0.52	0.65	BC002675	Hs.177776	FLJ21916	chromosome 14 open reading frame 137
0.14	-1.05	1.18	BC009016	Hs.94953	API5	apoptosis inhibitor 5
0.13	-0.26	0.40	BC009194	Hs.30541	NAT2	N-acetyltransferase 2 (arylamine N-acetyltransferase)
0.13	-0.16	0.30	BC009670	Hs.279669	AAK1	adaptor-associated kinase 1
0.13	0.11	0.02	BC001721	Hs.86948	S100A8	S100 calcium binding protein A8 (calgranulin A)
0.13	-0.64	0.77	BC009107	Hs.14014	HNRPDL	heterogeneous nuclear ribonucleoprotein D-like
0.13	-1.45	1.59	BC001217	Hs.268053	DPM1	dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic subunit
0.13	-0.23	0.36	BC000449	Hs.183704	PEA15	phosphoprotein enriched in astrocytes 15
0.13	-0.15	0.28	BC005152	Hs.279818	AKR7A2	aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)
0.13	-0.51	0.64	BC012341	Hs.79530	SEPX1	selenoprotein X, 1
0.13	-1.97	2.10	BC000996	Hs.6396	JTB	jumping translocation breakpoint
0.13	0.05	0.08	BC000941	Hs.124027	SPS	selenophosphate synthetase
0.13	-0.15	0.28	BC006419	Hs.334504	BAAT	bile acid Coenzyme A: amino acid N-acyltransferase (glycine N-choyltransferase)
0.13	-0.53	0.66	BC010010	Hs.334641	PTGS2	prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)
0.13	-0.29	0.42	BC019346	Hs.131190	MGC13034	hypothetical protein MGC13034
0.13	-0.83	0.96	BC010897	Hs.95420	GAGEC1	G antigen, family C, 1
0.13	-0.31	0.44	BC004333	Hs.273415	WWP2	Nedd-4-like ubiquitin-protein ligase
0.13	-1.80	1.93	BC007846	Hs.132898	LOC115861	
0.13	-0.30	0.43	BC015126	Hs.40919	HLA-DPB1	major histocompatibility complex, class II, DP beta 1
0.13	0.00	0.13	BC007419	Hs.26440	HPRP3P	U4/U6-associated RNA splicing factor
0.13	-0.57	0.70	BC010021	Hs.180900	LOC90557	hypothetical protein BC016861
0.13	-0.28	0.41	BC022062	Hs.282804	FLJ10035	hypothetical protein FLJ10035
0.13	-0.22	0.35	BC014324	Hs.44423	MGC15438	hypothetical protein MGC15438
0.13	-0.53	0.66	Data not found		HRMT1L2	HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae)

0.13	-0.61	0.73	BC004207	Hs.146329	FLJ25070	hypothetical protein FLJ25070
0.13	-0.61	0.73	BC014994	Hs.6150	ACOX1	acyl-Coenzyme A oxidase 1, palmitoyl
0.12	-0.21	0.34	BC014349	Hs.355719	PKM2	pyruvate kinase, muscle
0.12	-0.01	0.13	BC003563	Hs.424138	KIAA0101	KIAA0101 gene product
0.12	-0.25	0.37	BC011025	Hs.422340	SRI	sorcini
0.12	-3.05	3.17	BC017693	Hs.171012	DHRS1	dehydrogenase/reductase SDR family member 1
0.12	-0.40	0.53	BC011797	Hs.96247	TPD52	tumor protein D52
0.12	-0.61	0.73	BC015984	Hs.356795	FLJ20374	hypothetical protein FLJ20374
0.12	-0.76	0.88	BC016319	Hs.433898	APBA3	amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)
0.12	0.01	0.11	BC002523	Hs.63489	RPS10	
0.12	-1.24	1.36	BC004289	Hs.7482	MATR3	matrin 3
0.12	-0.22	0.34	BC018198	Hs.247220	MAPK7	mitogen-activated protein kinase 7
0.12	0.11	0.01	BC016842	Hs.8258	RCL	putative c-Myc-responsive
0.12	-0.47	0.59	BC000358	Hs.3136	FLJ22390	hypothetical protein FLJ22390
0.12	-0.42	0.54	BC008765	Hs.82109	SDC1	syndecan 1
0.12	-0.63	0.75	BC014075	Hs.283677	CDK5RAP1	CDK5 regulatory subunit associated protein 1
0.12	-0.25	0.37	BC014145	Hs.261134	MGC14128	hypothetical protein MGC14128
0.12	-1.86	1.98	BC013420	Hs.223014	WNT5B	wingless-type MMTV integration site family, member 5B
0.12	-0.47	0.59	BC008293	Hs.293659	FOXM1	forkhead box M1
0.12	-0.54	0.66	BC017242	Hs.104637	PELO	pelota homolog (Drosophila)
0.12	0.01	0.11	BC007527	Hs.183138	ENDOG	endonuclease G
0.12	0.03	0.09	BC015128	Hs.107476	ZDHHC6	zinc finger, DHHC domain containing 6
0.12	0.09	0.03	BC006550	Hs.146381	RBMX	RNA binding motif protein, X chromosome
0.12	-1.94	2.06	BC012069	Hs.355399	TCFL1	transcription factor-like 1
0.12	-1.61	1.73	BC013162	Hs.235356	STEAP	six transmembrane epithelial antigen of the prostate
0.12	-1.00	1.11	BC000439	Hs.979	PDHB	pyruvate dehydrogenase (lipoamide) beta
0.12	-1.16	1.27	BC020956	Hs.7187	BCAS2	breast carcinoma amplified sequence 2
0.12	-1.37	1.49	BC004270	Hs.238730	MKRN2	makorin, ring finger protein, 2
0.12	-0.37	0.49	BC007412	Hs.265317	FLJ13352	hypothetical protein FLJ13352
0.12	-0.97	1.08	BC011034	Hs.9667	BBOX1	butyrobetaine (gamma), 2-oxoglutarate dioxygenase (gamma-butyrobetaine hydroxylase) 1
0.12	-1.76	1.88	BC017729	Hs.9222	RPA3	replication protein A3, 14kDa
0.12	-0.34	0.46	BC000404	Hs.6566	PCTK3	PCTAIRE protein kinase 3
0.12	-0.38	0.50	BC013302	Hs.75564		
0.12	-0.58	0.70	BC007868	Hs.334541	PSMA1	proteasome (prosome, macropain) subunit, alpha type, 1
0.11	0.08	0.03	BC018105	Hs.127241	CAPNS2	calpain small subunit 2
0.11	-0.69	0.80	BC001380	Hs.469	SDHA	succinate dehydrogenase complex, subunit A, flavoprotein (Fp)
0.11	-0.69	0.81	BC008800	Hs.99890	POLD1	polymerase (DNA directed), delta 1, catalytic subunit 125kDa
0.11	-1.11	1.23	BC000718	Hs.84153	PSG4	pregnancy specific beta-1-glycoprotein 4
0.11	-0.35	0.46	BC007286	Hs.120363	Spir-1	Spir-1 protein
0.11	-0.56	0.67	BC002857	Hs.46659	MUT	methylmalonyl Coenzyme A mutase
0.11	-0.61	0.72	BC009238	Hs.75318	MGC1314	hypothetical protein MGC1314 similar to fucosidase, alpha-L- 1, tissue
0.11	-0.43	0.54	BC006281	Hs.268541	ZBRK1	zinc-finger protein ZBRK1
0.11	-0.48	0.59	BC021264	Hs.352211	ZNF2	zinc finger protein 2 (A1-5)
0.11	0.08	0.03	BC008215	Hs.334928	ARHE	ras homolog gene family, member E
0.11	-0.49	0.60	BC012811	Hs.198281	STMN2	stathmin-like 2
0.11	-0.11	0.22	BC001410	Hs.417004	MGC10974	hypothetical protein MGC10974
0.11	-0.17	0.28	BC009302	Hs.9275	DKFZP564G0222	unc-50 related
0.11	-1.08	1.19	BC006769	Hs.75337	NOLC1	nucleolar and coiled-body phosphoprotein 1
0.11	-0.42	0.53	BC002933	Hs.283009	DKFZP547E2110	anti-silencing function 1A
0.11	0.00	0.11	BC011675	Hs.26333	FLJ10074	hypothetical protein FLJ10074
0.11	-0.04	0.14	BC000490	Hs.7811	GALE	galactose-4-epimerase, UDP-
0.11	0.01	0.09	BC006229	Hs.1342	COX5B	cytochrome c oxidase subunit Vb
0.11	-1.14	1.25	BC014301	Hs.433413	PXMP4	peroxisomal membrane protein 4, 24kDa



0.11	-0.06	0.17	BC001344	Hs.237971	MGC5627	hypothetical protein MGC5627
0.11	-0.31	0.42	BC009346	Data not found	MGC20781	hypothetical protein MGC20781
0.11	-0.62	0.73	BC005919	Hs.76422	EIF3S4	eukaryotic translation initiation factor 3, subunit 4 delta, 44kDa
0.11	-1.74	1.85	BC001951	In multiple clusters	MAFG	v-maf musculoaponeurotic fibrosarcoma oncogene homolog G (avian)
0.11	-0.07	0.17	BC020512	Hs.374608	VAMP4	vesicle-associated membrane protein 4
0.11	-2.73	2.84	BC002839	Hs.153591	LOC51315	hypothetical protein LOC51315
0.11	-0.17	0.28	BC015401	Hs.107968	HLA-DRB4	major histocompatibility complex, class II, DR beta 4
0.11	-0.02	0.13	BC003519	Data not found	H2AFY2	H2A histone family, member Y2
0.11	0.09	0.02	BC003112	Hs.902	ATP1B3	ATPase, Na <sup>+</sup> /K <sup>+</sup> transporting, beta 3 polypeptide
0.10	-0.76	0.87	BC000120	Hs.68257	GLB1	galactosidase, beta 1
0.10	-0.46	0.56	BC008365	Hs.43621	MBC3205	hypothetical protein MBC3205
0.10	-1.11	1.21	BC015523	Hs.169907	TIMM17B	translocase of inner mitochondrial membrane 17 homolog B (yeast)
0.10	-0.57	0.67	BC018143	Hs.423615	IRF3	interferon regulatory factor 3
0.10	-0.48	0.59	NM_016448	Hs.126774	RAMP	RA-regulated nuclear matrix-associated protein
0.10	-0.21	0.31	BC013027	Hs.83135	ATF3	activating transcription factor 3
0.10	-0.02	0.12	BC003115	Hs.8719	NY-REN-41	NY-REN-41 antigen
0.10	-0.39	0.49	BC014794	Hs.99514	SCAM-1	vinexin beta (SH3-containing adaptor molecule-1)
0.10	-0.43	0.54	BC015138	Hs.381074	DENR	density-regulated protein
0.10	-2.35	2.45	BC016956	Hs.324830	CYFIP2	cytoplasmic FMR1 interacting protein 2
0.10	-0.92	1.02	BC014372	Hs.42644	FLJ10349	hypothetical protein FLJ10349
0.10	-1.71	1.82	BC003103	Hs.78748	8D6A	8D6 antigen
0.10	-0.90	1.00	BC016845	Hs.104938	TMPRSS4	transmembrane protease, serine 4
0.10	0.04	0.06	BC005298	Hs.184298	TNFRSF5	tumor necrosis factor receptor superfamily, member 5
0.10	-1.66	1.76	BC013740	Hs.244378	ERO1L	ERO1-like (S. cerevisiae)
0.10	-0.52	0.62	BC018647	Hs.65238	MGC3251	STARD3 N-terminal like
0.10	-0.70	0.80	BC001123	Hs.279929	HSGP25L2G	gp25L2 protein
0.10	-0.78	0.87	BC002809	Hs.16697	SLBP	stem-loop (histone) binding protein
0.10	-0.02	0.12	BC008481	Hs.296341	CALM2	calmodulin 2 (phosphorylase kinase, delta)
0.10	-1.43	1.53	BC002866	Hs.301547	TTK	TTK protein kinase
0.10	-1.62	1.71	BC007708	Hs.113	BCL2L1	BCL2-like 1
0.10	-0.59	0.68	BC020560	Hs.18616	ELA3B	elastase 3B, pancreatic
0.10	-0.66	0.75	BC010122	Hs.6048	MRPS2	mitochondrial ribosomal protein S2
0.10	0.08	0.02	BC002490	Hs.15093	HSPC195	hypothetical protein HSPC195
0.10	-0.48	0.58	BC006240	Data not found	CCND1	cyclin D1 (PRAD1: parathyroid adenomatosis 1)
0.10	-1.41	1.50	BC015615	Data not found	FH	fumarate hydratase
0.10	-0.25	0.35	BC013150	Hs.348334	CDC20	CDC20 cell division cycle 20 homolog (S. cerevisiae)
0.10	-1.06	1.15	BC000180	Hs.406013	KRT18	keratin 18
0.10	-0.32	0.42	BC016340	Hs.302754		
0.09	-0.12	0.22	BC005831	Hs.2484	NOL1	nucleolar protein 1, 120kDa
0.09	-0.16	0.25	BC000611	Hs.48375	ZDHHC9	zinc finger, DHHC domain containing 9
0.09	-0.95	1.04	BC011835	Hs.76941	NF2	neurofibromin 2 (bilateral acoustic neuroma)
0.09	-1.02	1.11	BC021676	Hs.160786	MLLT6	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6
0.09	-0.51	0.60	BC000589	Hs.279574	BIRC5	baculoviral IAP repeat-containing 5 (survivin)
0.09	-1.23	1.32	BC015045	Hs.75641	PEPD	peptidase D
0.09	-0.29	0.38	BC012604	Hs.27076	RTCD1	RTC domain containing 1
0.09	-0.20	0.29	BC005853	Hs.343236	MGC2487	hypothetical protein MGC2487
0.09	-1.22	1.31	BC003388	Hs.146847	TANK	TRAF family member-associated NFKB activator
0.09	-0.19	0.28	BC007552	Data not found	NA 4-14-03 520	
0.09	-0.24	0.33	BC007007	Hs.83077	FLJ10468	hypothetical protein FLJ10468
0.09	-1.14	1.23	BC017498	Hs.133034	APEX1	APEX nuclease (multifunctional DNA repair enzyme) 1
0.09	-0.02	0.11	BC009256	Hs.374608	ANXA5	annexin A5
0.09	-1.44	1.53	BC014366	Hs.65454	MGC14433	hypothetical protein MGC14433
0.09	-0.95	1.04	BC008686	Hs.78221	ACADVL	acyl-Coenzyme A dehydrogenase, very long chain

0.09	-1.70	1.80	BC003186	Hs.433180	VDAC1	voltage-dependent anion channel 1
0.09	-0.15	0.24	BC001873	Hs.234434	TFG	TRK-fused gene
0.09	-0.01	0.10	BC012603	Hs.168289	SDHD	succinate dehydrogenase complex, subunit D, integral membrane protein
0.09	-0.48	0.57	BC015692	Hs.334668	MGC13272	hypothetical protein MGC13272
0.09	-0.05	0.14	BC021664	Hs.351306	NA 4-14-03 376	
0.09	-0.12	0.21	BC003570	Hs.66708	GNB2	guanine nucleotide binding protein (G protein), beta polypeptide 2
0.09	-0.62	0.70	BC009930	Hs.133968	PCDHB5	protocadherin beta 5
0.09	-1.15	1.24	BC016058	Hs.83942	FLJ20288	FLJ20288 protein
0.09	-0.18	0.27	BC009679	Hs.11411	C20orf154	chromosome 20 open reading frame 154
0.09	-0.41	0.50	BC009583	Hs.7917	HIG1	likely ortholog of mouse hypoxia induced gene 1
0.09	-1.76	1.85	BC000595	Hs.349121	MGC11349	hypothetical protein MGC11349
0.09	-0.06	0.15	BC009423	Data not found	CGBP	CpG binding protein
0.09	-0.85	0.94	BC003547	Hs.375214	LOC91862	similar to RIKEN cDNA 1810006A16 gene
0.09	-0.32	0.41	BC009642	Hs.333418	HSPC002	HSPC002 protein
0.09	-0.58	0.67	BC002403	Hs.79339	FLJ20195	hypothetical protein FLJ20195
0.09	-0.08	0.16	BC000666	Hs.279861	APH2	Abl-philin 2
0.09	-0.27	0.35	BC003641	Hs.332381	WASF1	WAS protein family, member 1
0.09	-1.00	1.09	BC004560	Hs.277704	HNRPA0	heterogeneous nuclear ribonucleoprotein A0
0.09	-1.44	1.53	BC014109	Hs.143774	NPD002	NPD002 protein
0.09	-0.65	0.73	BC002923	Hs.31968	LDLR	low density lipoprotein receptor (familial hypercholesterolemia)
0.09	-0.05	0.14	BC013928	Hs.430150	HLA-DRB3	major histocompatibility complex, class II, DR beta 3
0.09	0.00	0.09	BC000632	Hs.432009	ARFGAP3	ADP-ribosylation factor GTPase activating protein 3
0.08	-0.63	0.71	BC009925	Hs.405913	RBM15	RNA binding motif protein 15
0.08	-0.80	0.89	BC005250	Hs.279902	APG12L	APG12 autophagy 12-like (S. cerevisiae)
0.08	-0.40	0.49	BC014900	Hs.5320	COPZ1	CGI-120 protein
0.08	-0.11	0.20	BC002931	Hs.43299	HSF1	heat shock transcription factor 1
0.08	-0.87	0.95	BC009875	Hs.422118	SDHD	succinate dehydrogenase complex, subunit D, integral membrane protein
0.08	-0.79	0.87	BC007813	Hs.417157	PNLIP	pancreatic lipase
0.08	-1.97	2.05	BC012325	Hs.15813	FLN29	FLN29 gene product
0.08	-0.52	0.60	BC003398	Hs.196437	C2orf6	chromosome 2 open reading frame 6
0.08	-0.80	0.88	BC017205	Hs.194638	SKD3	suppressor of potassium transport defect 3
0.08	-1.05	1.13	BC007317	Hs.54702	CPLX1	complexin 1
0.08	-0.93	1.02	BC000336	Hs.116428	ZNF219	zinc finger protein 219
0.08	-0.13	0.21	BC012886	Hs.272822	RAB6IP1	RAB6 interacting protein 1
0.08	-0.01	0.09	BC005930	Hs.75626	GPAA1	GPAA1P anchor attachment protein 1 homolog (yeast)
0.08	-0.72	0.80	BC018272	Hs.25413	MIF	macrophage migration inhibitory factor (glycosylation-inhibiting factor)
0.08	-0.29	0.37	BC008488	Hs.401703	TOLLIP	Toll-interacting protein
0.08	-0.38	0.46	BC004436	Hs.299629	MLANA	melan-A
0.08	0.03	0.05	BC007848	Hs.80305		Homo sapiens, clone IMAGE:4620009, mRNA
0.08	-1.07	1.15	BC001030	Hs.324874	TNFRSF9	tumor necrosis factor receptor superfamily, member 9
0.08	-0.17	0.25	BC012806	Hs.321579	PAK4	p21(CDKN1A)-activated kinase 4
0.08	0.00	0.08	BC008794	Hs.295112	POM121	POM121 membrane glycoprotein (rat)
0.08	-1.36	1.44	BC002432	Hs.194685	FLJ12118	hypothetical protein FLJ12118
0.08	-0.15	0.23	BC002645	Hs.302300	CAVI	caveolin 1, caveolae protein, 22kDa
0.08	-0.19	0.27	BC000717	Hs.119537	FLJ22318	hypothetical protein FLJ22318
0.08	-0.37	0.44	BC007545	Hs.25197	RetSDR2	retinal short-chain dehydrogenase/reductase 2
0.08	-0.65	0.73	BC003607	Hs.44423	FLJ20514	hypothetical protein FLJ20514
0.08	-0.20	0.27	BC002671	Hs.2359	APOC1	apolipoprotein C-I
0.08	-0.30	0.37	BC004952	Hs.316750	NR4A1	nuclear receptor subfamily 4, group A, member 1
0.08	-0.23	0.31	BC007208	Hs.9822	LOC120379	hypothetical protein BC019238
0.08	-0.12	0.20	BC000335	Hs.172108	FLJ21802	hypothetical protein FLJ21802
0.07	-0.43	0.50	BC010504	Hs.26450	MGC16153	hypothetical protein MGC16153
0.07	-0.47	0.54	BC001205	Data not found	SP2	Sp2 transcription factor

0.07	-0.71	0.78	BC006525	Hs.267963	BA108L7.2	similar to rat tricarboxylate carrier-like protein
0.07	-0.35	0.42	BC013433	Hs.158196	AP3S2	adaptor-related protein complex 3, sigma 2 subunit
0.07	-1.05	1.12	BC004427	Hs.233952	MRPL53	mitochondrial ribosomal protein L53
0.07	-0.20	0.27	BC014367	Data not found		
0.07	0.03	0.05	BC017101	Hs.296380	HNRPU	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)
0.07	-0.21	0.29	BC018931	Hs.74451	CAPNS1	calpain, small subunit 1
0.07	-0.80	0.87	BC013772	Hs.73021	FLJ20345	hypothetical protein FLJ20345
0.07	-3.34	3.41	BC012146	Hs.119598	NA 4-14-03 294	
0.07	-0.74	0.81	BC007740	Hs.247917	KIAA1228	KIAA1228 protein
0.07	-0.21	0.28	BC010076	Hs.424751	LOC115811	hypothetical protein BC013151
0.07	0.01	0.06	BC008078	Hs.22129	DJ1042K10.2	hypothetical protein DJ1042K10.2
0.07	0.05	0.02	BC015038	Hs.66881	SYT11	synaptotagmin XI
0.07	-1.57	1.64	BC009247	Hs.74571	MGC12916	hypothetical protein MGC12916
0.07	-0.50	0.57	BC010112	Hs.79037	SYN47	homer homolog 1 (Drosophila)
0.07	-0.45	0.52	BC006333	Hs.179703	KIAA1951	KIAA1951 protein
0.07	-0.46	0.53	BC000319	Hs.15961	NA 4-14-03 201	
0.07	-0.11	0.18	BC009278	Hs.325520	POU2F2	POU domain, class 2, transcription factor 2
0.07	-0.17	0.24	BC015663	Hs.21858	MGC16279	hypothetical protein MGC16279
0.07	-1.85	1.92	BC014453	Hs.271411	MGC15396	hypothetical protein MGC15396
0.07	-0.29	0.35	BC012421	Hs.266902	FLJ20502	hypothetical protein FLJ20502
0.07	-0.01	0.08	BC000942	Hs.7733	TTC1	tetratricopeptide repeat domain 1
0.07	0.03	0.04	BC009914	Hs.3764	MVD	mevalonate (diphospho) decarboxylase
0.07	-0.50	0.56	BC003157	Hs.95583	NET-7	transmembrane 4 superfamily member tetraspan NET-7
0.07	-0.06	0.13	BC016680	Hs.77031	SEDL	
0.07	-1.04	1.11	BC011873	Hs.70488	MO25	MO25 protein
0.07	-0.95	1.02	BC021087	Hs.146332	P24B	integral type I protein
0.07	-1.79	1.86	BC010930	Hs.423336	MGC13523	hypothetical protein MGC13523
0.06	-1.17	1.24	BC009787	Hs.31218	SCAMP1	secretory carrier membrane protein 1
0.06	-0.54	0.60	BC003187	Hs.16085	ITGB4BP	integrin beta 4 binding protein
0.06	-0.61	0.68	BC002480	Hs.22972	MGC2562	Similar to RIKEN cDNA 2810002N01 gene
0.06	-1.12	1.19	BC021190	Hs.152531	DAZAP2	DAZ associated protein 2
0.06	0.02	0.05	BC004889	Hs.3759	MGC3040	hypothetical protein MGC3040
0.06	-0.32	0.39	BC002758	Hs.188661	ZNF9	zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)
0.06	-0.83	0.89	BC010923	Hs.41683	CART1	cartilage paired-class homeoprotein 1
0.06	-1.47	1.53	BC019103	Hs.406648	CAPN1	
0.06	-0.37	0.43	NM_005100	Hs.788	TRPV2	transient receptor potential cation channel, subfamily V, member 2
0.06	-0.99	1.05	BC019045	Hs.98628	FUBP3	far upstream element (FUSE) binding protein 3
0.06	0.03	0.04	BC001689	Hs.13845	SLC25A20	solute carrier family 25 (carnitine/acylcarnitine translocase), member 20
0.06	-0.30	0.36	BC005991	Hs.213248	MGC3232	hypothetical protein MGC3232
0.06	-0.39	0.45	BC000048	Hs.334475	PAPSS2	3'-phosphoadenosine 5'-phosphosulfate synthase 2
0.06	-0.15	0.21	BC017327	Hs.75260	FLJ23375	hypothetical protein FLJ23375
0.06	-1.33	1.39	BC000371	Hs.181418	FLJ20265	hypothetical protein FLJ20265
0.06	-0.37	0.43	BC010740	Hs.15898	MGC4796	hypothetical protein MGC4796
0.06	0.04	0.02	BC013151	Hs.254936	MRPS6	mitochondrial ribosomal protein S6
0.06	-0.35	0.41	BC013039	Hs.83135	UBE1L	ubiquitin-activating enzyme E1-like
0.06	0.05	0.01	BC007082	Hs.332848	LMO4	LIM domain only 4
0.06	-0.19	0.25	BC020571	Hs.306422	SERPINF1	serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 1
0.06	-0.72	0.78	BC014621	Hs.75692	FLJ10439	hypothetical protein FLJ10439
0.06	-1.62	1.68	BC021992	Hs.177574		
0.06	-0.26	0.32	BC013913	Hs.273077	SPIB	Spi-B transcription factor (Spi-1/PU.1 related)
0.06	-0.15	0.20	BC007272	Hs.380969	SMAP	small acidic protein
0.06	-0.26	0.32	BC012317	Hs.149219	LOC93082	hypothetical protein BC012317

0.06	-1.27	1.33	BC022198	Hs.76719	VAX2	ventral anterior homeobox 2
0.06	-1.02	1.07	BC009794	Hs.53973	VIP	vasoactive intestinal peptide
0.05	-0.13	0.18	BC017713	Hs.153546	PCQAP	PC2 (positive cofactor 2, multiprotein complex) glutamine/Q-rich-associated protein
0.05	-0.04	0.10	BC005975	Hs.27258	DKFZP586I2223	intermediate filament-like MGC:2625
0.05	-0.90	0.95	BC010407	Hs.227789	NA 4-14-03 34	
0.05	-0.23	0.28	BC016389	Data not found	SLN	sarcolipin
0.05	-0.70	0.76	BC011986	Hs.421646	ZYX	zyxin
0.05	-1.49	1.55	BC006221	Hs.197764	RAB11A	RAB11A, member RAS oncogene family
0.05	-1.28	1.33	BC009686	Hs.415877	PRMT6	protein arginine N-methyltransferase 6
0.05	-1.36	1.41	BC012543	Hs.13144	ORMDL2	ORM1-like 2 (S. cerevisiae)
0.05	-0.59	0.65	BC020517	Hs.348389	NA 4-14-03 474	
0.05	-1.94	2.00	BC006146	Hs.334371	ASB3	ankyrin repeat and SOCS box-containing 3
0.05	-0.96	1.01	BC014077	Hs.83575	PRO1580	hypothetical protein PRO1580
0.05	-0.83	0.88	BC000029	Hs.154846	C20orf20	chromosome 20 open reading frame 20
0.05	-0.53	0.58	BC000435	Hs.78614	DCTD	dCMP deaminase
0.05	-0.03	0.08	BC015122	Hs.234489	RPL9	ribosomal protein L9
0.05	-0.06	0.11	BC002704	Hs.21486	TM9SF1	transmembrane 9 superfamily member 1
0.05	-0.52	0.57	BC014523	Data not found	CYFIP1	cytoplasmic FMR1 interacting protein 1
0.05	-1.05	1.10	BC016157	Hs.334721	FLJ13391	hypothetical protein FLJ13391
0.05	-1.23	1.28	BC014891	Hs.153293	KIAA0798	KIAA0798 gene product
0.05	-0.08	0.13	BC010167	Hs.339834	CBARA1	calcium binding atopy-related autoantigen 1
0.05	-0.13	0.18	BC015491	Hs.380718	BOP1	block of proliferation 1
0.05	-1.38	1.43	BC000103	Hs.101695	MGC955	hypothetical protein MGC955
0.05	-0.37	0.41	BC006081	Hs.5807	FLJ10687	hypothetical protein FLJ10687
0.05	-0.07	0.11	BC000949	Hs.274350	BAF53A	BAF53
0.05	-0.67	0.71	BC007676	Hs.407995	MIF	macrophage migration inhibitory factor (glycosylation-inhibiting factor)
0.05	-0.75	0.80	BC004358	Hs.190161	MGC20398	hypothetical protein MGC20398
0.05	-0.07	0.11	BC000055	Hs.433622	SFRS7	splicing factor, arginine/serine-rich 7, 35kDa
0.05	-0.12	0.17	BC018160	Hs.73946	C20orf67	chromosome 20 open reading frame 67
0.05	-0.70	0.75	BC016728	Hs.64595	RERG	RAS-like, estrogen-regulated, growth-inhibitor
0.04	-0.03	0.07	BC008299	Hs.41502	MGC24447	chromosome 14 open reading frame 147
0.04	-0.96	1.01	BC001169	Hs.82193	ITGB3BP	integrin beta 3 binding protein (beta3-endonexin)
0.04	-2.45	2.50	BC006472	Hs.83097	B4GALT2	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2
0.04	-0.44	0.48	BC015842	Hs.173912	NUP160	nucleoporin 160kDa
0.04	-0.08	0.12	BC008500	Hs.274549	NA 4-14-03 283	
0.04	-0.60	0.64	BC020979	Hs.42650	CPB2	carboxypeptidase B2 (plasma, carboxypeptidase U)
0.04	-0.18	0.22	BC011856	Hs.112188	BCAT2	branched chain aminotransferase 2, mitochondrial
0.04	-0.12	0.16	BC013035	Hs.10018	MARS	methionine-tRNA synthetase
0.04	-0.20	0.24	BC002725	Hs.105191	MADH5	MAD, mothers against decapentaplegic homolog 5 (Drosophila)
0.04	-1.25	1.29	BC012452	Hs.288057	MGC15407	similar to RIKEN cDNA 4931428D14 gene
0.04	-0.43	0.47	BC004100	Hs.177	METTL1	methyltransferase-like 1
0.04	-0.37	0.41	BC006552	Hs.29647	MGC3248	dynactin 4
0.04	-0.84	0.88	BC013181	Data not found	LOC51244	hypothetical protein LOC51244
0.04	-0.31	0.35	BC006980	Hs.39733	TRIM31	tripartite motif-containing 31
0.04	-0.40	0.45	BC014340	Hs.348540	RXRA	retinoid X receptor, alpha
0.04	-0.16	0.20	BC017328	Hs.12854	LOC113730	hypothetical protein BC009980
0.04	-0.22	0.26	BC000582	Hs.178292	CAPN3	calpain 3, (p94)
0.04	-0.17	0.21	BC021194	Hs.575	FLJ20291	hypothetical protein FLJ20291
0.04	-1.81	1.85	BC013915	Hs.401787	CD38	CD38 antigen (p45)
0.04	-0.06	0.10	BC014545	Hs.321618	FLJ12525	hypothetical protein FLJ12525
0.04	-0.30	0.34	BC008835	Hs.32017	IVD	isovaleryl Coenzyme A dehydrogenase
0.04	-1.73	1.77	BC007689	Hs.262958	DKFZP434B044	hypothetical protein DKFZp434B044
0.04	-0.73	0.77	BC000808	Hs.129445	SLC30A5	solute carrier family 30 (zinc transporter), member 5

0.04	-0.59	0.63	BC009255	Hs.120824	NBP	nucleotide binding protein
0.04	-1.04	1.07	BC007038	Hs.79914	NIP30	NEFA-interacting nuclear protein NIP30
0.04	-1.03	1.06	BC007312	Hs.145729	SRP9	signal recognition particle 9kDa
0.04	-0.25	0.29	BC017196	Hs.104125	WWP2	Nedd-4-like ubiquitin-protein ligase
0.04	-0.29	0.32	BC005234	Hs.11774	FLJ22548	
0.04	-0.07	0.11	BC011875	Hs.380145	SLC26A6	solute carrier family 26, member 6
0.04	-0.72	0.75	BC012925	Hs.285681	EIF3S7	eukaryotic translation initiation factor 3, subunit 7 zeta, 66/67kDa
0.04	-0.33	0.37	BC010116	Hs.343214	MAT1A	methionine adenosyltransferase I, alpha
0.04	-0.17	0.20	BC009445	Data not found	PPP2R5C	protein phosphatase 2, regulatory subunit B (B56), gamma isoform
0.04	-1.57	1.61	BC015514	Hs.170279	UQCRCF1	ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1
0.04	-0.41	0.44	BC009457	Hs.182874	OAI	ocular albinism 1 (Nettleship-Falls)
0.04	-0.89	0.92	BC021697	Hs.90606	NMT1	N-myristoyltransferase 1
0.03	-0.44	0.48	BC000540	Hs.17936	FLJ14525	hypothetical protein FLJ14525
0.03	-1.56	1.59	BC002952	Hs.89474	MY014	sorting nexin 27
0.03	-2.69	2.73	BC001434	Hs.9061	MGC2477	hypothetical protein MGC2477
0.03	-0.98	1.01	BC008485	Hs.149176	GYG	glycogenin
0.03	-0.39	0.42	BC012022	Hs.80350	TRAP25	TRAP/Mediator complex component
0.03	-0.04	0.08	BC017089	Hs.105484	PTD002	PTD002 protein
0.03	-0.58	0.61	BC014879	Hs.355874	KLK10	kallikrein 10
0.03	-0.62	0.66	BC009411	Hs.11039	FLJ20859	hypothetical protein FLJ20859
0.03	-0.19	0.22	BC003020	Hs.250821	RPA2	replication protein A2, 32kDa
0.03	-0.46	0.49	BC016285	Hs.87773	MGC10812	hypothetical protein MGC10812
0.03	-0.26	0.29	BC000494	Hs.170001		
0.03	-0.10	0.13	BC002515	Hs.74294	ALDH7A1	aldehyde dehydrogenase 7 family, member A1
0.03	-1.58	1.61	BC005352	Hs.17839	ADIR	ATP-dependant interferon responsive
0.03	-0.50	0.53	BC014471	Hs.274614	NA 4-14-03 40	
0.03	-0.58	0.61	Data not found		TTRAP	TRAF and TNF receptor-associated protein
0.03	-2.22	2.25	BC018677	Hs.425808	BSCL2	Bernardinelli-Seip congenital lipodystrophy 2 (seipin)
0.03	-0.02	0.05	BC013187	Hs.77961	PLSCR3	phospholipid scramblase 3
0.03	-0.78	0.80	BC016825	Hs.16758	MGC15634	hypothetical protein MGC15634
0.03	-0.04	0.07	BC011551	Hs.13525	MGC4189	hypothetical protein MGC4189
0.03	-1.07	1.10	BC011033	Hs.264482	APG12L	APG12 autophagy 12-like (S. cerevisiae)
0.03	-0.91	0.94	BC005326	Hs.76064	AHCYL1	S-adenosylhomocysteine hydrolase-like 1
0.03	-0.06	0.09	BC015476	Hs.274317	FOXA2	forkhead box A2
0.03	-0.80	0.82	BC017377	Hs.349589	STXBP1	syntaxin binding protein 1
0.03	-0.04	0.07	BC001682	Hs.323467	NEUGRN	mesenchymal stem cell protein DSC92
0.03	-1.31	1.34	BC014668	Hs.135150	MKPX	mitogen-activated protein kinase phosphatase x
0.03	-0.72	0.75	BC008413	Hs.292566	AKR1B1	aldo-keto reductase family 1, member B1 (aldose reductase)
0.03	-1.24	1.26	BC008818	Hs.78781	VEGFB	vascular endothelial growth factor B
0.03	-0.65	0.67	BC018201	Data not found	KIAA0354	KIAA0354 gene product
0.03	-0.24	0.26	BC008788	Hs.126704	FLJ20584	hypothetical protein FLJ20584
0.02	-0.61	0.63	BC000616	Hs.153026	MGC10731	hypothetical protein MGC10731
0.02	-1.97	1.99	BC005003	Hs.168157	C15orf15	chromosome 15 open reading frame 15
0.02	-0.54	0.57	BC008954	Hs.25450	SLC29A1	solute carrier family 29 (nucleoside transporters), member 1
0.02	-0.30	0.32	BC016331	Hs.284162	IARS	isoleucine-tRNA synthetase
0.02	-0.23	0.25	BC003541	Hs.279832	FLJ13855	hypothetical protein FLJ13855
0.02	-1.34	1.36	BC000983	Hs.247280	C20orf18	chromosome 20 open reading frame 18
0.02	-0.13	0.15	BC005961	Hs.89626	H105E3	NAD(P) dependent steroid dehydrogenase-like; H105e3
0.02	-0.94	0.97	BC000251	Hs.78802	BAP29	B-cell receptor-associated protein BAP29
0.02	-0.06	0.09	BC016847	Hs.1189	DSCR2	Down syndrome critical region gene 2
0.02	-0.01	0.03	BC009538	Hs.112208	GLA	galactosidase, alpha
0.02	-0.83	0.85	BC008091	Data not found	CASP3	caspase 3, apoptosis-related cysteine protease
0.02	-1.31	1.34	BC016005	Hs.356547	ACTG1	actin, gamma 1

0.02	-0.46	0.48	BC005937	Hs.267288	MGC2628	hypothetical protein MGC2628
0.02	-0.31	0.33	BC008502	Data not found	SDCCAG16	serologically defined colon cancer antigen 16
0.02	-1.21	1.23	BC009576	Hs.82159	PSMA1	proteasome (prosome, macropain) subunit, alpha type, 1
0.02	-3.30	3.32	BC000544	Hs.102308	ENTH	enthoprotin
0.02	-0.27	0.29	BC015848	Hs.279061	PPP1R15B	protein phosphatase 1, regulatory (inhibitor) subunit 15B
0.02	-1.13	1.15	BC004974	Hs.111779	BC002942	hypothetical protein BC002942
0.02	-0.90	0.92	BC013413	Hs.268384	GARS	glycyl-tRNA synthetase
0.02	-0.40	0.42	BC001000	Data not found	MGC5363	
0.02	-1.73	1.75	BC009512	Hs.334842	TFPT	TCF3 (E2A) fusion partner (in childhood Leukemia)
0.02	-0.69	0.70	BC012444	Hs.25035	HDCMA18P	HDCMA18P protein
0.02	-0.16	0.17	BC001137	Hs.306225	MGC2875	hypothetical protein MGC2875
0.02	-0.35	0.36	BC000542	Hs.301789	HOXD3	homeo box D3
0.02	-0.90	0.92	BC015531	Hs.250502		
0.02	-1.39	1.40	BC008202	Hs.55781	FLJ20604	hypothetical protein FLJ20604
0.01	-0.81	0.83	BC011444	Hs.180141	CFL2	cofilin 2 (muscle)
0.01	-1.07	1.09	BC004856	Hs.301760	PSK-1	type I transmembrane receptor (seizure-related protein)
0.01	-0.23	0.25	BC005022	Hs.7905	PTGES2	prostaglandin E synthase 2
0.01	-0.18	0.19	BC015781	Hs.405961	DKFZP434N178	kinesin family member 26A
0.01	-1.35	1.36	BC007122	Hs.183655	SEC22A	SEC22 vesicle trafficking protein-like 2 (S. cerevisiae)
0.01	-0.04	0.05	BC015958	Hs.172182	TRIM5	tripartite motif-containing 5
0.01	-1.05	1.06	BC013728	Hs.356386	PLOD3	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3
0.01	-1.46	1.47	BC012196	Hs.274422	HEY2	hair/enhancer-of-split related with YRPW motif 2
0.01	-0.47	0.48	BC000972	Hs.26333	JM1	JM1 protein
0.01	-1.01	1.02	BC016919	Hs.351335	HPGD	hydroxyprostaglandin dehydrogenase 15-(NAD)
0.01	-0.52	0.53	BC014885	Hs.301005	NUBP2	nucleotide binding protein 2 (MinD homolog, E. coli)
0.01	-0.64	0.65	BC013380	Hs.426930	KIAA0212	KIAA0212 gene product
0.01	-0.26	0.27	BC015528	Hs.9731	PRO2730	hypothetical protein PRO2730
0.01	-0.51	0.52	BC010914	Hs.283108	HBG1	hemoglobin, gamma A
0.01	-0.03	0.04	BC001350	Hs.172851	SLC29A2	solute carrier family 29 (nucleoside transporters), member 2
0.01	-0.54	0.55	BC013101	Hs.40094	ARHGEF1	Rho guanine nucleotide exchange factor (GEF) 1
0.01	-1.09	1.11	BC014861	Hs.426930	RPS6KA2	ribosomal protein S6 kinase, 90kDa, polypeptide 2
0.01	-0.66	0.67	BC000971	Hs.286049	PSA	phosphoserine aminotransferase
0.01	-0.41	0.42	BC007857	Hs.180920	CGI-87	CGI-87 protein
0.01	-0.18	0.18	BC016052	Hs.102929	FLJ12089	hypothetical protein FLJ12089
0.01	-0.35	0.36	BC015912	Hs.70877	ATP6V1G1	ATPase, H <sup>+</sup> transporting, lysosomal 13kDa, V1 subunit G isoform 1
0.01	-0.46	0.47	BC017336	Data not found	NA 4-14-03 468	
0.01	-0.71	0.72	BC004399	Hs.125783	NA 4-14-03 135	
0.01	-1.04	1.04	NM_006370	Hs.169206	SYNGR3	synaptogyrin 3
0.01	-0.64	0.65	BC009106	Hs.293650	KNS2	kinesin 2 60/70kDa
0.01	-0.11	0.12	BC013935	Data not found	MGC14407	hypothetical protein MGC14407
0.01	-1.25	1.26	BC013773	Hs.184465	MGC19780	hypothetical protein MGC19780
0.01	-0.36	0.37	BC007019	Hs.6651	MGC:5244	hypothetical protein MGC5244
0.01	-1.05	1.06	BC006999	Hs.76038	ICAP-1A	integrin cytoplasmic domain-associated protein 1
0.01	-0.37	0.38	BC010951	Hs.99949	PIP	prolactin-induced protein