

Id	GenBank	UniGene	Description	Symbol	EG	upregulated in EG1	downregulated in EG1	upregulated in EG2	downregulated in EG2	EG1 p-value	EG1 q-value (Storey)	EG2 p-value	EG2 q-value (Storey)
219282_s_at	NM_015930	Hs.279746	transient receptor potential cation channel, subfamily	TRPV2	EG1	4	10	5	1	2,15E-11	0,0029338410,0147492140,111110101		
1555996_s_at	AI332397	Hs.478553	eukaryotic translation initiation factor 4A, isoform 2	EIF4A2	EG1	10	3	6	2	6,10E-10	0,0029338410,0005786180,011359908		
201468_s_at	NM_000903	Hs.406515	NAD(P)H dehydrogenase, quinone 1	NQO1	EG1	9	4	3	2	6,10E-10	0,0029338410,0540847830,274766231		
1555989_at	AA890373	Hs.19156	dishevelled associated activator of morphogenesis 1	DAAM1	EG1	8	4	5	3	1,28E-08	0,0029338410,0005786180,011359908		
218252_at	NM_018204	Hs.444028	cytoskeleton associated protein 2	CKAP2	EG1	11	1	4	4	1,28E-08	0,0029338410,0005786180,011359908		
201123_s_at	NM_001970	Hs.534314	eukaryotic translation initiation factor 5A	EIF5A	EG1	11	1	4	2	1,28E-08	0,0029338410,0147492140,111110101		
203028_s_at	NM_000101	Hs.513803	cytochrome b-245, alpha polypeptide	CYBA	EG1	3	9	5	1	1,28E-08	0,0029338410,0147492140,111110101		
212366_at	BF223237	Hs.485892	zinc finger protein 292	ZNF292	EG1	8	4	3	3	1,28E-08	0,0029338410,0147492140,111110101		
225191_at	AL565767	Hs.501309	cold inducible RNA binding protein	CIRBP	EG1	8	4	3	3	1,28E-08	0,0029338410,0147492140,111110101		
225767_at	AI825833	Hs.370699		LOC284801	EG1	7	5	1	5	1,28E-08	0,0029338410,0147492140,111110101		
228273_at	BG165011	Hs.274448		FLJ11029	EG1	5	7	2	4	1,28E-08	0,0029338410,0147492140,111110101		
229390_at	AV734646	Hs.381220		LOC441168	EG1	5	7	3	3	1,28E-08	0,0029338410,0147492140,111110101		
239577_at	AV699781				EG1	10	2	5	1	1,28E-08	0,0029338410,0147492140,111110101		
235419_at	AW612461	Hs.11169	ERBB receptor feedback inhibitor 1	ERRFI1	EG1	7	5	3	1	1,28E-08	0,0029338410,1581335050,559565828		
201310_s_at	NM_004772	Hs.483067	chromosome 5 open reading frame 13	C5orf13	EG1	4	7	5	3	2,08E-07	0,0029338410,0005786180,011359908		
202589_at	NM_001071	Hs.369762	thymidylate synthetase	TYMS	EG1	4	7	2	6	2,08E-07	0,0029338410,0005786180,011359908		
204127_at	BC000149	Hs.115474	replication factor C (activator 1) 3, 38kDa	RFC3	EG1	10	1	3	5	2,08E-07	0,0029338410,0005786180,011359908		
205193_at	NM_012323	Hs.517617	v-maf musculoaponeurotic fibrosarcoma oncogene ho	MAFF	EG1	8	3	5	3	2,08E-07	0,0029338410,0005786180,011359908		
209211_at	AF132818	Hs.508234	Kruppel-like factor 5 (intestinal)	KLF5	EG1	7	4	2	6	2,08E-07	0,0029338410,0005786180,011359908		
219049_at	NM_018371	Hs.387794		ChGn	EG1	2	9	2	6	2,08E-07	0,0029338410,0005786180,011359908		
223255_at	AA642341	Hs.509008	KIAA1333	KIAA1333	EG1	11	0	4	4	2,08E-07	0,0029338410,0005786180,011359908		
228974_at	AI816281	Hs.20506	zinc finger protein 677	ZNF677	EG1	8	3	5	3	2,08E-07	0,0029338410,0005786180,011359908		
AFFX-PheX-M	AFFX-PHEX-M				EG1	5	6	5	3	2,08E-07	0,0029338410,0005786180,011359908		
216237_s_at	AA807529	Hs.517582	MCM5 minichromosome maintenance deficient 5, cell	MCM5	EG1	5	6	2	5	2,08E-07	0,0029338410,0032420520,038332332		
224407_s_at	AF344882	Hs.444247		MASK	EG1	10	1	4	3	2,08E-07	0,0029338410,0032420520,038332332		
201792_at	NM_001129	Hs.439463	AE binding protein 1	AEBP1	EG1	3	8	2	4	2,08E-07	0,0029338410,0147492140,111110101		
202581_at	NM_005346	Hs.274402	heat shock 70kDa protein 1B	HSPA1B	EG1	10	1	3	3	2,08E-07	0,0029338410,0147492140,111110101		
202688_at	NM_003810	Hs.478275	tumor necrosis factor (ligand) superfamily, member 1	TNFSF10	EG1	2	9	1	5	2,08E-07	0,0029338410,0147492140,111110101		
214132_at	BG232034	Hs.271135	ATP synthase, H+ transporting, mitochondrial F1 comp	ATP5C1	EG1	8	3	5	1	2,08E-07	0,0029338410,0147492140,111110101		
214820_at	AJ002572	Hs.314338	bromodomain and WD repeat domain containing 1	BRWD1	EG1	9	2	2	4	2,08E-07	0,0029338410,0147492140,111110101		
226998_at	AL556909	Hs.555985	NMDA receptor regulated 1	NARG1	EG1	11	0	4	2	2,08E-07	0,0029338410,0147492140,111110101		
228334_x_at	AI633734	Hs.555989	KIAA1712	KIAA1712	EG1	9	2	3	3	2,08E-07	0,0029338410,0147492140,111110101		
229391_s_at	AV734646	Hs.381220		LOC441168	EG1	4	7	3	3	2,08E-07	0,0029338410,0147492140,111110101		
235412_at	AI040887	Hs.508738	Rho guanine nucleotide exchange factor (GEF) 7	ARHGEF7	EG1	6	5	4	2	2,08E-07	0,0029338410,0147492140,111110101		
236114_at	AI798118	Hs.149261,	H runt-related transcription factor 1 (acute myeloid leuk	RUNX1	EG1	0	11	6	0	2,08E-07	0,0029338410,0147492140,111110101		
AFFX-PheX-3_	AFFX-PHEX-3				EG1	5	6	3	3	2,08E-07	0,0029338410,0147492140,111110101		
201211_s_at	AF061337	Hs.380774	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked	DDX3X	EG1	9	2	3	2	2,08E-07	0,0029338410,0540847830,274766231		
205092_x_at	NM_014950	Hs.400802	zinc finger and BTB domain containing 1	ZBTB1	EG1	8	3	2	3	2,08E-07	0,0029338410,0540847830,274766231		
207838_x_at	NM_020524	Hs.505806	pre-B-cell leukemia transcription factor interacting pr	PBXIP1	EG1	5	6	1	4	2,08E-07	0,0029338410,0540847830,274766231		
209679_s_at	BC003379	Hs.206501		LOC57228	EG1	5	6	3	2	2,08E-07	0,0029338410,0540847830,274766231		
221848_at	AL121845	Hs.11900	zinc finger, CCCH-type with G patch domain	ZGPAT	EG1	2	9	4	1	2,08E-07	0,0029338410,0540847830,274766231		
226613_at	AI742029	Hs.444950	TBC1 domain family, member 10A	TBC1D10A	EG1	2	9	1	4	2,08E-07	0,0029338410,0540847830,274766231		
228328_at	AI982758				EG1	9	2	2	3	2,08E-07	0,0029338410,0540847830,274766231		

233014_at	AK022980	Hs.283652	isopentenyl-diphosphate delta isomerase 1	IDI1	EG1	8	3	3	2	2,08E-07	0,0029338410,0540847830,274766231
231975_s_at	AK002183	Hs.419195		FLJ35954	EG1	11	0	3	1	2,08E-07	0,0029338410,1581335050,559565828
241242_at	BE503118	Hs.372003	chromosome 9 open reading frame 10	C9orf10	EG1	3	8	4	0	2,08E-07	0,0029338410,1581335050,559565828
207247_s_at	NM_003411	Hs.522845	zinc finger protein, Y-linked	ZFY	EG1	10	1	3	0	2,08E-07	0,0029338410,3636349230,911710613
211538_s_at	U56725	Hs.432648	heat shock 70kDa protein 2	HSPA2	EG1	7	4	2	1	2,08E-07	0,0029338410,3636349230,911710613
221135_s_at	NM_014065	Hs.100878		HT001	EG1	9	2	2	0	2,08E-07	0,0029338410,6490161710,911710613
1552701_a_at	NM_052889	Hs.348365		COP1	EG1	1	9	7	1	2,66E-06	0,0029338410,0005786180,011359908
1554609_at	BC019340	Hs.292791		MGC12965	EG1	3	7	1	7	2,66E-06	0,0029338410,0005786180,011359908
204092_s_at	NM_003600	Hs.250822	serine/threonine kinase 6	STK6	EG1	8	2	3	5	2,66E-06	0,0029338410,0005786180,011359908
204244_s_at	NM_006716	Hs.489037		ASK	EG1	10	0	3	5	2,66E-06	0,0029338410,0005786180,011359908
204709_s_at	NM_004856	Hs.270845	kinesin family member 23	KIF23	EG1	7	3	3	5	2,66E-06	0,0029338410,0005786180,011359908
207165_at	NM_012485	Hs.72550	hyaluronan-mediated motility receptor (RHAMM)	HMMR	EG1	7	3	3	5	2,66E-06	0,0029338410,0005786180,011359908
211368_s_at	U13700	Hs.2490	caspase 1, apoptosis-related cysteine peptidase (inter	CASP1	EG1	2	8	6	2	2,66E-06	0,0029338410,0005786180,011359908
212977_at	AI817041	Hs.471751	chemokine orphan receptor 1	CMKOR1	EG1	0	10	3	5	2,66E-06	0,0029338410,0005786180,011359908
217924_at	AL523965	Hs.485162	chromosome 6 open reading frame 106	C6orf106	EG1	7	3	8	0	2,66E-06	0,0029338410,0005786180,011359908
222040_at	AI144007	Hs.210850	H heterogeneous nuclear ribonucleoprotein A1	HNRPA1	EG1	9	1	4	4	2,66E-06	0,0029338410,0005786180,011359908
222088_s_at	AA778684	Hs.210227	solute carrier family 2 (facilitated glucose transporter)	SLC2A14	EG1	5	5	4	4	2,66E-06	0,0029338410,0005786180,011359908
225902_at	AW194716	Hs.470544	peptidyl-prolyl isomerase G (cyclophilin G)	PPIG	EG1	9	1	1	7	2,66E-06	0,0029338410,0005786180,011359908
228751_at	AW975057	Hs.406557	CDC-like kinase 4	CLK4	EG1	7	3	5	3	2,66E-06	0,0029338410,0005786180,011359908
230461_s_at	AI860326	Hs.515016	melanoma associated antigen (mutated) 1	MUM1	EG1	7	3	4	4	2,66E-06	0,0029338410,0005786180,011359908
232797_at	AU144005	Hs.436873	integrin, alpha V (vitronectin receptor, alpha polypept	ITGAV	EG1	4	6	8	0	2,66E-06	0,0029338410,0005786180,011359908
233899_x_at	AK024296	Hs.205742	zinc finger and BTB domain containing 10	ZBTB10	EG1	8	2	3	5	2,66E-06	0,0029338410,0005786180,011359908
235152_at	AW293849				EG1	1	9	7	1	2,66E-06	0,0029338410,0005786180,011359908
235167_at	BE972419	Hs.528462, Hs.531664, Hs.547315, Hs.556385		DKFZp547E087	EG1	8	2	5	3	2,66E-06	0,0029338410,0005786180,011359908
236402_at	AW184034	Hs.490366	v-raf murine sarcoma viral oncogene homolog B1	BRAF	EG1	6	4	4	4	2,66E-06	0,0029338410,0005786180,011359908
236752_at	AA913146	Hs.407580	plakophilin 4	PKP4	EG1	6	4	8	0	2,66E-06	0,0029338410,0005786180,011359908
238712_at	BF801735				EG1	1	9	6	2	2,66E-06	0,0029338410,0005786180,011359908
243648_at	AA280627	Hs.532399	zinc finger CCCH-type containing 11A	ZC3H11A	EG1	3	7	5	3	2,66E-06	0,0029338410,0005786180,011359908
AFFX-r2-Bs-th	AFFX-R2-BS-THR-3				EG1	5	5	6	2	2,66E-06	0,0029338410,0005786180,011359908
201774_s_at	AK022511	Hs.5719		CNAP1	EG1	5	5	2	5	2,66E-06	0,0029338410,0032420520,038332332
202338_at	NM_003258	Hs.515122	thymidine kinase 1, soluble	TK1	EG1	5	5	2	5	2,66E-06	0,0029338410,0032420520,038332332
203109_at	NM_003969	Hs.406068	ubiquitin-conjugating enzyme E2M (UBC12 homolog, y	UBE2M	EG1	7	3	4	3	2,66E-06	0,0029338410,0032420520,038332332
203422_at	NM_002691	Hs.279413	polymerase (DNA directed), delta 1, catalytic subunit	POLD1	EG1	3	7	2	5	2,66E-06	0,0029338410,0032420520,038332332
206917_at	NM_006572	Hs.515018	guanine nucleotide binding protein (G protein), alpha	GNA13	EG1	9	1	7	0	2,66E-06	0,0029338410,0032420520,038332332
209093_s_at	K02920	Hs.282997, H	glucosidase, beta, acid (includes glucosylceramidase)	GBA	EG1	2	8	4	3	2,66E-06	0,0029338410,0032420520,038332332
210006_at	BC002571	Hs.534400	abhydrolase domain containing 14A	ABHD14A	EG1	2	8	4	3	2,66E-06	0,0029338410,0032420520,038332332
211366_x_at	U13698	Hs.2490	caspase 1, apoptosis-related cysteine peptidase (inter	CASP1	EG1	2	8	6	1	2,66E-06	0,0029338410,0032420520,038332332
212187_x_at	NM_000954	Hs.446429	prostaglandin D2 synthase 21kDa (brain)	PTGDS	EG1	6	4	5	2	2,66E-06	0,0029338410,0032420520,038332332
213023_at	NM_007124	Hs.133135	utrophin (homologous to dystrophin)	UTRN	EG1	8	2	4	3	2,66E-06	0,0029338410,0032420520,038332332
214877_at	BE794663	Hs.484844	CDK5 regulatory subunit associated protein 1-like 1	CDKAL1	EG1	5	5	4	3	2,66E-06	0,0029338410,0032420520,038332332
219004_s_at	NM_018944	Hs.190518	chromosome 21 open reading frame 45	C21orf45	EG1	5	5	2	5	2,66E-06	0,0029338410,0032420520,038332332
219334_s_at	NM_022837	Hs.118183		FLJ22833	EG1	5	5	3	4	2,66E-06	0,0029338410,0032420520,038332332
221843_s_at	AA195017	Hs.288274		KIAA1609	EG1	2	8	2	5	2,66E-06	0,0029338410,0032420520,038332332
225762_x_at	AL531683	Hs.370699		LOC284801	EG1	7	3	2	5	2,66E-06	0,0029338410,0032420520,038332332
225786_at	AI440495	Hs.411490, H	family with sequence similarity 36, member A	FAM36A	EG1	8	2	2	5	2,66E-06	0,0029338410,0032420520,038332332

227198_at	AW085505	Hs.444414	AF4/FMR2 family, member 3	AFF3	EG1	5	5	1	6	2,66E-06	0,0029338410,0032420520,038332332
228561_at	BF510563	Hs.493361	CDC37 cell division cycle 37 homolog (<i>S. cerevisiae</i>)-li	CDC37L1	EG1	9	1	4	3	2,66E-06	0,0029338410,0032420520,038332332
231069_at	AI768895				EG1	6	4	5	2	2,66E-06	0,0029338410,0032420520,038332332
231152_at	AW452971				EG1	9	1	3	4	2,66E-06	0,0029338410,0032420520,038332332
242868_at	T70087	Hs.468410	endothelial PAS domain protein 1	EPAS1	EG1	6	4	6	1	2,66E-06	0,0029338410,0032420520,038332332
243829_at	AW613053	Hs.490366	v-raf murine sarcoma viral oncogene homolog B1	BRAF	EG1	9	1	6	1	2,66E-06	0,0029338410,0032420520,038332332
36711_at	AL021977	Hs.517617	v-maf musculoaponeurotic fibrosarcoma oncogene ho	MAFF	EG1	7	3	4	3	2,66E-06	0,0029338410,0032420520,038332332
AFFX-r2-Bs-ph	AFFX-R2-BS-PHE-M				EG1	4	6	3	4	2,66E-06	0,0029338410,0032420520,038332332
1552302_at	NM_145041	Hs.446690		MGC20235	EG1	3	7	3	3	2,66E-06	0,0029338410,0147492140,111110101
1552703_s_at	NM_052889	Hs.348365		COP1	EG1	1	9	4	2	2,66E-06	0,0029338410,0147492140,111110101
1555892_s_at	BF476087	Hs.131417, Hs.193725		LOC253039	EG1	6	4	2	4	2,66E-06	0,0029338410,0147492140,111110101
202107_s_at	NM_004526	Hs.477481	MCM2 minichromosome maintenance deficient 2, mit	MCM2	EG1	4	6	2	4	2,66E-06	0,0029338410,0147492140,111110101
202295_s_at	NM_004390	Hs.148641	cathepsin H	CTSH	EG1	2	8	2	4	2,66E-06	0,0029338410,0147492140,111110101
206548_at	NM_024880	Hs.501080		FLJ23556	EG1	5	5	5	1	2,66E-06	0,0029338410,0147492140,111110101
207030_s_at	NM_001321	Hs.530904	cysteine and glycine-rich protein 2	CSRP2	EG1	4	6	3	3	2,66E-06	0,0029338410,0147492140,111110101
208881_x_at	BC005247	Hs.283652	isopentenyl-diphosphate delta isomerase 1	IDI1	EG1	9	1	6	0	2,66E-06	0,0029338410,0147492140,111110101
210589_s_at	D13287	Hs.282997, H	glucosidase, beta, acid (includes glucosylceramidase)	GBA	EG1	1	9	4	2	2,66E-06	0,0029338410,0147492140,111110101
210845_s_at	U08839	Hs.466871	plasminogen activator, urokinase receptor	PLAUR	EG1	7	3	4	2	2,66E-06	0,0029338410,0147492140,111110101
211676_s_at	AF056979	Hs.520414	interferon gamma receptor 1	IFNGR1	EG1	9	1	5	1	2,66E-06	0,0029338410,0147492140,111110101
213302_at	AL044326	Hs.88139	phosphoribosylformylglycinamide synthase (FGAR a	PFAS	EG1	5	5	3	3	2,66E-06	0,0029338410,0147492140,111110101
219854_at	NM_021030	Hs.466261	zinc finger protein 14 (KOX 6)	ZNF14	EG1	7	3	4	2	2,66E-06	0,0029338410,0147492140,111110101
224558_s_at	AI446756	Hs.187199, H	metastasis associated lung adenocarcinoma transcript	MALAT1	EG1	8	2	5	1	2,66E-06	0,0029338410,0147492140,111110101
224973_at	AL078599	Hs.10784	family with sequence similarity 46, member A	FAM46A	EG1	7	3	5	1	2,66E-06	0,0029338410,0147492140,111110101
225227_at	AW294869				EG1	8	2	3	3	2,66E-06	0,0029338410,0147492140,111110101
227265_at	AW135176				EG1	5	5	5	1	2,66E-06	0,0029338410,0147492140,111110101
234995_at	AA668779	Hs.477144		LOC152185	EG1	4	6	3	3	2,66E-06	0,0029338410,0147492140,111110101
240572_s_at	BF436632	Hs.396637		LOC374443	EG1	3	7	4	2	2,66E-06	0,0029338410,0147492140,111110101
1553984_s_at	AF258562	Hs.471873	deoxythymidylate kinase (thymidylate kinase)	DTYMK	EG1	4	6	2	3	2,66E-06	0,0029338410,0540847830,274766231
1562836_at	AK021715	Hs.408461	DEAD (Asp-Glu-Ala-Asp) box polypeptide 6	DDX6	EG1	8	2	2	3	2,66E-06	0,0029338410,0540847830,274766231
1563469_at	AL832681	Hs.535297	AT rich interactive domain 5B (MRF1-like)	ARID5B	EG1	2	8	2	3	2,66E-06	0,0029338410,0540847830,274766231
202687_s_at	U57059	Hs.478275	tumor necrosis factor (ligand) superfamily, member 1	TNFSF10	EG1	2	8	1	4	2,66E-06	0,0029338410,0540847830,274766231
204170_s_at	NM_001827	Hs.83758	CDC28 protein kinase regulatory subunit 2	CKS2	EG1	9	1	2	3	2,66E-06	0,0029338410,0540847830,274766231
212514_x_at	R60068	Hs.380774	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked	DDX3X	EG1	10	0	3	2	2,66E-06	0,0029338410,0540847830,274766231
226444_at	AI700476	Hs.413434	solute carrier family 39 (zinc transporter), member 10	SLC39A10	EG1	8	2	5	0	2,66E-06	0,0029338410,0540847830,274766231
228742_at	AA745978				EG1	2	8	1	4	2,66E-06	0,0029338410,0540847830,274766231
229735_s_at	W15435	Hs.523442	NIPA-like domain containing 3	NPAL3	EG1	4	6	1	4	2,66E-06	0,0029338410,0540847830,274766231
230036_at	BE669858	Hs.489118	sterile alpha motif domain containing 9-like	SAMD9L	EG1	3	7	1	4	2,66E-06	0,0029338410,0540847830,274766231
231269_at	AU153330	Hs.486031		DJ467N11.1	EG1	9	1	4	1	2,66E-06	0,0029338410,0540847830,274766231
232978_at	AF339810	Hs.125715	muscleblind-like 2 (<i>Drosophila</i>)	MBNL2	EG1	7	3	5	0	2,66E-06	0,0029338410,0540847830,274766231
233849_s_at	AK023014	Hs.525287	Rho GTPase activating protein 5	ARHGAP5	EG1	8	2	0	5	2,66E-06	0,0029338410,0540847830,274766231
236974_at	AA808018	Hs.518827	cyclin I	CCNI	EG1	8	2	5	0	2,66E-06	0,0029338410,0540847830,274766231
241627_x_at	AI640434	Hs.35125		FLJ10357	EG1	6	4	3	2	2,66E-06	0,0029338410,0540847830,274766231
1557813_at	BF724621	Hs.102735	single-stranded DNA binding protein 2	SSBP2	EG1	8	2	3	1	2,66E-06	0,0029338410,1581335050,559565828
205449_at	NM_013299	Hs.23642	SAC3 domain containing 1	SAC3D1	EG1	4	6	1	3	2,66E-06	0,0029338410,1581335050,559565828
214787_at	BE268538	Hs.513817	c-myc promoter binding protein	MYCPBP	EG1	10	0	1	3	2,66E-06	0,0029338410,1581335050,559565828

221428_s_at	NM_030921	Hs.438970	transducin (beta)-like 1X-linked receptor 1	TBL1XR1	EG1	7	3	1	3	2,66E-06	0,0029338410,1581335050,559565828
221763_at	AI694023	Hs.413416,	H jumonji domain containing 1C	JMJD1C	EG1	9	1	2	2	2,66E-06	0,0029338410,1581335050,559565828
244811_at	AI561173	Hs.511817	pleckstrin homology domain interacting protein	PHIP	EG1	9	1	2	2	2,66E-06	0,0029338410,1581335050,559565828
1557522_x_at	AA653638	Hs.472651	bladder cancer associated protein	BLCAP	EG1	7	3	1	2	2,66E-06	0,0029338410,3636349230,911710613
204868_at	NM_001545	Hs.407955	immature colon carcinoma transcript 1	ICT1	EG1	3	7	1	2	2,66E-06	0,0029338410,3636349230,911710613
211752_s_at	BC005954	Hs.211914	NADH dehydrogenase (ubiquinone) Fe-S protein 7, 20	NDUFS7	EG1	2	8	1	2	2,66E-06	0,0029338410,3636349230,911710613
212037_at	BF508848	Hs.409965	pinin, desmosome associated protein	PNN	EG1	10	0	1	2	2,66E-06	0,0029338410,3636349230,911710613
218166_s_at	NM_016578	Hs.420229	hepatitis B virus x associated protein	HBXAP	EG1	9	1	2	1	2,66E-06	0,0029338410,3636349230,911710613
219470_x_at	NM_019084	Hs.281348	cyclin J	CCNJ	EG1	9	1	3	0	2,66E-06	0,0029338410,3636349230,911710613
228758_at	AW264036	Hs.478589		LOC389185	EG1	7	3	3	0	2,66E-06	0,0029338410,3636349230,911710613
232382_s_at	BE150929	Hs.308480		LOC115294	EG1	4	6	3	0	2,66E-06	0,0029338410,3636349230,911710613
234411_x_at	U94903	Hs.502328	CD44 antigen (homing function and Indian blood grou	CD44	EG1	9	1	2	0	2,66E-06	0,0029338410,6490161710,911710613
238465_at	AI265933	Hs.85950		MGC33648	EG1	5	5	1	1	2,66E-06	0,0029338410,6490161710,911710613
239493_at	AI984074	Hs.421257	ribosomal protein L7	RPL7	EG1	9	1	1	1	2,66E-06	0,0029338410,6490161710,911710613
234881_at	L33988				EG1	3	7	1	0	2,66E-06	0,0029338410,8976818490,911710613
238456_at	AI567893				EG1	9	1	0	1	2,66E-06	0,0029338410,8976818490,911710613
1558401_at	AW953994				EG1	5	4	4	4	2,72E-05	0,0029338410,0005786180,011359908
1569157_s_at	BC037359	Hs.546512		LOC162993	EG1	2	7	2	6	2,72E-05	0,0029338410,0005786180,011359908
201136_at	NM_002668	Hs.77422	proteolipid protein 2 (colonic epithelium-enriched)	PLP2	EG1	2	7	0	8	2,72E-05	0,0029338410,0005786180,011359908
201294_s_at	N24643	Hs.446017	WD repeat and SOCS box-containing 1	WSB1	EG1	5	4	4	4	2,72E-05	0,0029338410,0005786180,011359908
203022_at	NM_006397	Hs.532851	ribonuclease H2, large subunit	RNASEH2A	EG1	4	5	4	4	2,72E-05	0,0029338410,0005786180,011359908
206632_s_at	NM_004900	Hs.226307	apolipoprotein B mRNA editing enzyme, catalytic poly	APOBEC3B	EG1	5	4	2	6	2,72E-05	0,0029338410,0005786180,011359908
208116_s_at	NM_005907	Hs.102788	mannosidase, alpha, class 1A, member 1	MAN1A1	EG1	7	2	5	3	2,72E-05	0,0029338410,0005786180,011359908
208733_at	AW301641	Hs.369017	RAB2, member RAS oncogene family	RAB2	EG1	4	5	4	4	2,72E-05	0,0029338410,0005786180,011359908
209164_s_at	BC002976	Hs.355264	cytochrome b-561	CYB561	EG1	2	7	3	5	2,72E-05	0,0029338410,0005786180,011359908
210172_at	D26121	Hs.502829	splicing factor 1	SF1	EG1	4	5	4	4	2,72E-05	0,0029338410,0005786180,011359908
212583_at	AB011132	Hs.510958	aquarius homolog (mouse)	AQR	EG1	4	5	7	1	2,72E-05	0,0029338410,0005786180,011359908
212847_at	AL036840	Hs.269099	far upstream element (FUSE) binding protein 1	FUBP1	EG1	8	1	5	3	2,72E-05	0,0029338410,0005786180,011359908
212907_at	AI972416	Hs.519469	solute carrier family 30 (zinc transporter), member 1	SLC30A1	EG1	9	0	8	0	2,72E-05	0,0029338410,0005786180,011359908
213182_x_at	R78668	Hs.106070	cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	EG1	3	6	4	4	2,72E-05	0,0029338410,0005786180,011359908
219215_s_at	NM_017767	Hs.521934	solute carrier family 39 (zinc transporter), member 4	SLC39A4	EG1	2	7	4	4	2,72E-05	0,0029338410,0005786180,011359908
221555_x_at	AU145941	Hs.40582	CDC14 cell division cycle 14 homolog B (S. cerevisiae)	CDC14B	EG1	3	6	4	4	2,72E-05	0,0029338410,0005786180,011359908
222316_at	AW973253	Hs.292689		VDP	EG1	7	2	6	2	2,72E-05	0,0029338410,0005786180,011359908
222662_at	W60806	Hs.458513		LOC286044	EG1	6	3	6	2	2,72E-05	0,0029338410,0005786180,011359908
223103_at	AF151810	Hs.188606	START domain containing 10	STARD10	EG1	3	6	0	8	2,72E-05	0,0029338410,0005786180,011359908
225834_at	AL135396	Hs.534958		LOC440687	EG1	4	5	3	5	2,72E-05	0,0029338410,0005786180,011359908
226682_at	AW006185	Hs.512943		LOC283666	EG1	4	5	5	3	2,72E-05	0,0029338410,0005786180,011359908
228153_at	AI953847	Hs.148741	IBR domain containing 2	IBRDC2	EG1	4	5	5	3	2,72E-05	0,0029338410,0005786180,011359908
229256_at	AV724329	Hs.26612	phosphoglucomutase 2-like 1	PGM2L1	EG1	6	3	3	5	2,72E-05	0,0029338410,0005786180,011359908
232573_at	AL389956	Hs.403933	F-box protein 32	FBXO32	EG1	7	2	6	2	2,72E-05	0,0029338410,0005786180,011359908
235440_at	BE780878	Hs.268668		FLJ39441	EG1	8	1	6	2	2,72E-05	0,0029338410,0005786180,011359908
235592_at	AW960145	Hs.192221,	H elongation factor, RNA polymerase II, 2	ELL2	EG1	6	3	4	4	2,72E-05	0,0029338410,0005786180,011359908
238909_at	BF126155	Hs.143873	S100 calcium binding protein A10 (annexin II ligand, ca	S100A10	EG1	1	8	6	2	2,72E-05	0,0029338410,0005786180,011359908
242691_at	AA829017				EG1	4	5	3	5	2,72E-05	0,0029338410,0005786180,011359908
244353_s_at	AI675682	Hs.486508	solute carrier family 2 (facilitated glucose transporter)	SLC2A12	EG1	4	5	1	7	2,72E-05	0,0029338410,0005786180,011359908

AFFX-r2-Bs-ph	AFFX-R2-BS-PHE-3				EG1	5	4	5	3	2,72E-05	0,0029338410,0005786180,011359908
1558102_at	AK055438	Hs.513094	transmembrane 6 superfamily member 1	TM6SF1	EG1	5	4	5	2	2,72E-05	0,0029338410,0032420520,038332332
1566901_at	AL832409	Hs.373550	TGFB-induced factor (TALE family homeobox)	TGIF	EG1	8	1	7	0	2,72E-05	0,0029338410,0032420520,038332332
201626_at	BG292233	Hs.520819	insulin induced gene 1	INSIG1	EG1	7	2	4	3	2,72E-05	0,0029338410,0032420520,038332332
201627_s_at	NM_005542	Hs.520819	insulin induced gene 1	INSIG1	EG1	7	2	4	3	2,72E-05	0,0029338410,0032420520,038332332
201702_s_at	AI492873	Hs.106019	protein phosphatase 1, regulatory subunit 10	PPP1R10	EG1	4	5	2	5	2,72E-05	0,0029338410,0032420520,038332332
202708_s_at	NM_003528	Hs.2178	histone 2, H2be	HIST2H2BE	EG1	8	1	4	3	2,72E-05	0,0029338410,0032420520,038332332
202869_at	NM_016816	Hs.524760	2',5'-oligoadenylate synthetase 1, 40/46kDa	OAS1	EG1	3	6	6	1	2,72E-05	0,0029338410,0032420520,038332332
202979_s_at	NM_021212	Hs.535319		ZF	EG1	8	1	5	2	2,72E-05	0,0029338410,0032420520,038332332
203828_s_at	NM_004221	Hs.943	interleukin 32	IL32	EG1	2	7	1	6	2,72E-05	0,0029338410,0032420520,038332332
204502_at	NM_015474	Hs.472630	SAM domain and HD domain 1	SAMHD1	EG1	5	4	2	5	2,72E-05	0,0029338410,0032420520,038332332
205681_at	NM_004049	Hs.227817	BCL2-related protein A1	BCL2A1	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
205798_at	NM_002185	Hs.362807	interleukin 7 receptor	IL7R	EG1	4	5	2	5	2,72E-05	0,0029338410,0032420520,038332332
208859_s_at	AI650257	Hs.533526	H alpha thalassemia/mental retardation syndrome X-lin	ATRX	EG1	2	7	5	2	2,72E-05	0,0029338410,0032420520,038332332
209417_s_at	BC001356	Hs.50842	interferon-induced protein 35	IFI35	EG1	2	7	3	4	2,72E-05	0,0029338410,0032420520,038332332
209512_at	BC004331	Hs.59486	hydroxysteroid dehydrogenase like 2	HSDL2	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
209681_at	AF153330	Hs.30246	solute carrier family 19 (thiamine transporter), memb	SLC19A2	EG1	9	0	4	3	2,72E-05	0,0029338410,0032420520,038332332
210778_s_at	BC002713	Hs.102402	MAX dimerization protein 4	MXD4	EG1	3	6	5	2	2,72E-05	0,0029338410,0032420520,038332332
212944_at	AK024896	Hs.302742	mitochondrial ribosomal protein S6	MRPS6	EG1	5	4	7	0	2,72E-05	0,0029338410,0032420520,038332332
213372_at	AW173157	Hs.556026	progesterin and adipoQ receptor family member III	PAQR3	EG1	8	1	6	1	2,72E-05	0,0029338410,0032420520,038332332
216766_at	AK025152	Hs.97432	protein kinase C, epsilon	PRKCE	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
218585_s_at	NM_016448	Hs.126774	denticleless homolog (Drosophila)	DTL	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
218918_at	NM_020379	Hs.197043	mannosidase, alpha, class 1C, member 1	MAN1C1	EG1	3	6	2	5	2,72E-05	0,0029338410,0032420520,038332332
219622_at	NM_017817	Hs.508720	RAB20, member RAS oncogene family	RAB20	EG1	4	5	0	7	2,72E-05	0,0029338410,0032420520,038332332
219978_s_at	NM_018454	Hs.511093	nucleolar and spindle associated protein 1	NUSAP1	EG1	6	3	2	5	2,72E-05	0,0029338410,0032420520,038332332
220941_s_at	NM_017447	Hs.293811	chromosome 21 open reading frame 91	C21orf91	EG1	6	3	6	1	2,72E-05	0,0029338410,0032420520,038332332
221276_s_at	NM_030786	Hs.16003	syncoilin, intermediate filament 1	SYNC1	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
221985_at	AW006750	Hs.407709	kelch-like 24 (Drosophila)	KLHL24	EG1	7	2	3	4	2,72E-05	0,0029338410,0032420520,038332332
222592_s_at	AW173691	Hs.11638	acyl-CoA synthetase long-chain family member 5	ACSL5	EG1	1	8	3	4	2,72E-05	0,0029338410,0032420520,038332332
225107_at	AI963008	Hs.535028		LOC442518	EG1	8	1	3	4	2,72E-05	0,0029338410,0032420520,038332332
225876_at	T84558	Hs.523442	NIPA-like domain containing 3	NPAL3	EG1	2	7	3	4	2,72E-05	0,0029338410,0032420520,038332332
226158_at	AL120021	Hs.407709	kelch-like 24 (Drosophila)	KLHL24	EG1	7	2	3	4	2,72E-05	0,0029338410,0032420520,038332332
226390_at	AA628398	Hs.93842	START domain containing 4, sterol regulated	STARD4	EG1	7	2	6	1	2,72E-05	0,0029338410,0032420520,038332332
226542_at	AI300571				EG1	5	4	1	6	2,72E-05	0,0029338410,0032420520,038332332
227040_at	AI655763	Hs.507783		LOC387921	EG1	4	5	2	5	2,72E-05	0,0029338410,0032420520,038332332
228160_at	AI433706	Hs.552607, Hs.553799		LOC400642	EG1	5	4	3	4	2,72E-05	0,0029338410,0032420520,038332332
228468_at	BF108964	Hs.276905	microtubule associated serine/threonine kinase-like	MASTL	EG1	8	1	2	5	2,72E-05	0,0029338410,0032420520,038332332
228674_s_at	AA524507	Hs.432438	echinoderm microtubule associated protein like 4	EML4	EG1	4	5	2	5	2,72E-05	0,0029338410,0032420520,038332332
229574_at	AI268231	Hs.445652		TRA2A	EG1	5	4	3	4	2,72E-05	0,0029338410,0032420520,038332332
229685_at	AI743207	Hs.518611	TBC1 domain family, member 14	TBC1D14	EG1	2	7	1	6	2,72E-05	0,0029338410,0032420520,038332332
230082_at	AW137053	Hs.475319	leucine rich repeat (in FLII) interacting protein 2	LRRFIP2	EG1	8	1	7	0	2,72E-05	0,0029338410,0032420520,038332332
234973_at	BG325630	Hs.195155	solute carrier family 38, member 5	SLC38A5	EG1	3	6	5	2	2,72E-05	0,0029338410,0032420520,038332332
235190_at	BF591288	Hs.282410	H calmodulin 2 (phosphorylase kinase, delta)	CALM2	EG1	8	1	7	0	2,72E-05	0,0029338410,0032420520,038332332
236072_at	N64578	Hs.435948	ATPase family, AAA domain containing 1	ATAD1	EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
236989_at	AW293012	Hs.405590	eukaryotic translation initiation factor 3, subunit 6 48k	EIF3S6	EG1	7	2	3	4	2,72E-05	0,0029338410,0032420520,038332332

238452_at	AI393356	Hs.517422	Fc receptor-like and mucin-like 2	FCRLM2	EG1	2	7	4	3	2,72E-05	0,0029338410,0032420520,038332332
238781_at	AI859076	Hs.519347	splicing factor, arginine/serine-rich 12	SFRS12	EG1	7	2	4	3	2,72E-05	0,0029338410,0032420520,038332332
242836_at	AI800470				EG1	5	4	4	3	2,72E-05	0,0029338410,0032420520,038332332
243016_at	AW271958	Hs.369762	thymidylate synthetase	TYMS	EG1	7	2	4	3	2,72E-05	0,0029338410,0032420520,038332332
1565703_at	AL832789	Hs.75862	SMAD, mothers against DPP homolog 4 (Drosophila)	SMAD4	EG1	6	3	4	2	2,72E-05	0,0029338410,0147492140,111110101
1565951_s_at	AJ293392	Hs.170129	choroideremia-like (Rab escort protein 2)	CHML	EG1	9	0	3	3	2,72E-05	0,0029338410,0147492140,111110101
201755_at	NM_006739	Hs.517582	MCM5 minichromosome maintenance deficient 5, cell	MCM5	EG1	3	6	2	4	2,72E-05	0,0029338410,0147492140,111110101
202464_s_at	NM_004566	Hs.195471	6-phosphofructo-2-kinase/fructose-2,6-biphosphatas	PFKFB3	EG1	4	5	3	3	2,72E-05	0,0029338410,0147492140,111110101
202779_s_at	NM_014501	Hs.396393	ubiquitin-conjugating enzyme E2S	UBE2S	EG1	7	2	3	3	2,72E-05	0,0029338410,0147492140,111110101
203836_s_at	D84476	Hs.186486	mitogen-activated protein kinase kinase kinase 5	MAP3K5	EG1	4	5	4	2	2,72E-05	0,0029338410,0147492140,111110101
204194_at	NM_001186	Hs.154276	BTB and CNC homology 1, basic leucine zipper transcri	BACH1	EG1	7	2	3	3	2,72E-05	0,0029338410,0147492140,111110101
205139_s_at	NM_005715	Hs.309250	uronyl-2-sulfotransferase	UST	EG1	7	2	1	5	2,72E-05	0,0029338410,0147492140,111110101
205436_s_at	NM_002105	Hs.477879	H2A histone family, member X	H2AFX	EG1	7	2	3	3	2,72E-05	0,0029338410,0147492140,111110101
208744_x_at	BG403660	Hs.36927	heat shock 105kDa/110kDa protein 1	HSPH1	EG1	9	0	4	2	2,72E-05	0,0029338410,0147492140,111110101
209163_at	AL514271	Hs.355264	cytochrome b-561	CYB561	EG1	4	5	3	3	2,72E-05	0,0029338410,0147492140,111110101
209421_at	U04045	Hs.156519	mutS homolog 2, colon cancer, nonpolyposis type 1 (E.	MSH2	EG1	3	6	2	4	2,72E-05	0,0029338410,0147492140,111110101
209840_s_at	AI221950	Hs.3781	leucine rich repeat neuronal 3	LRRN3	EG1	3	6	1	5	2,72E-05	0,0029338410,0147492140,111110101
210424_s_at	AF163441	Hs.182982	H golgi autoantigen, golgin subfamily a, 8A	GOLGA8A	EG1	6	3	5	1	2,72E-05	0,0029338410,0147492140,111110101
210793_s_at	U41815	Hs.524750	nucleoporin 98kDa	NUP98	EG1	7	2	5	1	2,72E-05	0,0029338410,0147492140,111110101
211323_s_at	L38019	Hs.374613	inositol 1,4,5-triphosphate receptor, type 1	ITPR1	EG1	7	2	2	4	2,72E-05	0,0029338410,0147492140,111110101
211748_x_at	BC005939	Hs.446429	prostaglandin D2 synthase 21kDa (brain)	PTGDS	EG1	4	5	4	2	2,72E-05	0,0029338410,0147492140,111110101
212746_s_at	AA126789	Hs.408293	H KIAA0470	KIAA0470	EG1	6	3	2	4	2,72E-05	0,0029338410,0147492140,111110101
212890_at	BF663461	Hs.352240		MGC15523	EG1	0	9	4	2	2,72E-05	0,0029338410,0147492140,111110101
215011_at	AJ006835	Hs.469723	RNA, U17D small nucleolar	RNU17D	EG1	8	1	2	4	2,72E-05	0,0029338410,0147492140,111110101
217949_s_at	NM_024006	Hs.324844	vitamin K epoxide reductase complex, subunit 1	VKORC1	EG1	0	9	4	2	2,72E-05	0,0029338410,0147492140,111110101
218317_x_at	NM_024044	Hs.460558	GIY-YIG domain containing 2	GIYD2	EG1	1	8	1	5	2,72E-05	0,0029338410,0147492140,111110101
220148_at	NM_022568	Hs.486520	aldehyde dehydrogenase 8 family, member A1	ALDH8A1	EG1	7	2	1	5	2,72E-05	0,0029338410,0147492140,111110101
222429_at	AF077353	Hs.436500	drebrin-like	DBNL	EG1	2	7	2	4	2,72E-05	0,0029338410,0147492140,111110101
223276_at	AF313413	Hs.29444		NID67	EG1	2	7	5	1	2,72E-05	0,0029338410,0147492140,111110101
223407_at	AL136786	Hs.307084	chromosome 16 open reading frame 48	C16orf48	EG1	1	8	0	6	2,72E-05	0,0029338410,0147492140,111110101
223577_x_at	AA827878	Hs.187199	H metastasis associated lung adenocarcinoma transcript	MALAT1	EG1	6	3	3	3	2,72E-05	0,0029338410,0147492140,111110101
225541_at	BE274422	Hs.380933	ribosomal protein L22-like 1	RPL22L1	EG1	5	4	3	3	2,72E-05	0,0029338410,0147492140,111110101
225899_x_at	AL040396	Hs.484396		LOC440395	EG1	5	4	2	4	2,72E-05	0,0029338410,0147492140,111110101
226922_at	BF215302	Hs.199561	RAN binding protein 2	RANBP2	EG1	8	1	5	1	2,72E-05	0,0029338410,0147492140,111110101
226923_at	AW205790	Hs.302287	sec1 family domain containing 2	SCFD2	EG1	3	6	3	3	2,72E-05	0,0029338410,0147492140,111110101
228193_s_at	AI744499	Hs.507866		RGC32	EG1	7	2	0	6	2,72E-05	0,0029338410,0147492140,111110101
228570_at	BF510581	Hs.271272	H BTB (POZ) domain containing 11	BTBD11	EG1	5	4	1	5	2,72E-05	0,0029338410,0147492140,111110101
229075_at	AI754871				EG1	6	3	2	4	2,72E-05	0,0029338410,0147492140,111110101
229393_at	AA088346	Hs.486466	l(3)mbt-like 3 (Drosophila)	L3MBTL3	EG1	7	2	2	4	2,72E-05	0,0029338410,0147492140,111110101
233292_s_at	AK024189	Hs.434219	ankyrin repeat and KH domain containing 1	ANKHD1	EG1	8	1	4	2	2,72E-05	0,0029338410,0147492140,111110101
235787_at	T65394	Hs.493361	CDC37 cell division cycle 37 homolog (S. cerevisiae)-li	CDC37L1	EG1	7	2	3	3	2,72E-05	0,0029338410,0147492140,111110101
235926_at	AI312527	Hs.7101	anaphase promoting complex subunit 5	ANAPC5	EG1	5	4	5	1	2,72E-05	0,0029338410,0147492140,111110101
236703_at	BF114733				EG1	5	4	4	2	2,72E-05	0,0029338410,0147492140,111110101
239228_at	AI298887	Hs.446484	H casein kinase 2, alpha 1 polypeptide	CSNK2A1	EG1	4	5	5	1	2,72E-05	0,0029338410,0147492140,111110101
242110_at	AV735587	Hs.525287	Rho GTPase activating protein 5	ARHGAP5	EG1	6	3	4	2	2,72E-05	0,0029338410,0147492140,111110101

244054_at	AI760298	Hs.445725	SKI-like	SKIL	EG1	7	2	4	2	2,72E-05	0,0029338410,0147492140,111110101
1555247_a_at	AF394782	Hs.483329	Rap guanine nucleotide exchange factor (GEF) 6	RAPGEF6	EG1	6	3	0	5	2,72E-05	0,0029338410,0540847830,274766231
1558837_a_at	BQ024490				EG1	3	6	4	1	2,72E-05	0,0029338410,0540847830,274766231
1559401_a_at	BI052176	Hs.170498	zinc finger protein 609	ZNF609	EG1	4	5	1	4	2,72E-05	0,0029338410,0540847830,274766231
1570021_at	BC036226	Hs.434961, Hs.558004		LOC360030	EG1	1	8	4	1	2,72E-05	0,0029338410,0540847830,274766231
200832_s_at	AB032261	Hs.368641	stearoyl-CoA desaturase (delta-9-desaturase)	SCD	EG1	6	3	1	4	2,72E-05	0,0029338410,0540847830,274766231
201435_s_at	AW268640	Hs.249718	eukaryotic translation initiation factor 4E	EIF4E	EG1	7	2	3	2	2,72E-05	0,0029338410,0540847830,274766231
201601_x_at	NM_003641	Hs.458414	interferon induced transmembrane protein 1 (9-27)	IFITM1	EG1	1	8	3	2	2,72E-05	0,0029338410,0540847830,274766231
201673_s_at	NM_002103	Hs.386225	glycogen synthase 1 (muscle)	GYS1	EG1	2	7	2	3	2,72E-05	0,0029338410,0540847830,274766231
201712_s_at	NM_006267	Hs.199561	RAN binding protein 2	RANBP2	EG1	9	0	4	1	2,72E-05	0,0029338410,0540847830,274766231
203209_at	BC001866	Hs.506989	replication factor C (activator 1) 5, 36.5kDa	RFC5	EG1	4	5	2	3	2,72E-05	0,0029338410,0540847830,274766231
204243_at	NM_012421	Hs.205627	rearranged L-myc fusion	RLF	EG1	6	3	2	3	2,72E-05	0,0029338410,0540847830,274766231
204270_at	AI568728	Hs.555981		FLJ13941	EG1	3	6	2	3	2,72E-05	0,0029338410,0540847830,274766231
204331_s_at	NM_021107	Hs.411125	mitochondrial ribosomal protein S12	MRPS12	EG1	2	7	3	2	2,72E-05	0,0029338410,0540847830,274766231
204545_at	NM_000287	Hs.107410	peroxisomal biogenesis factor 6	PEX6	EG1	3	6	0	5	2,72E-05	0,0029338410,0540847830,274766231
204857_at	NM_003550	Hs.209128	MAD1 mitotic arrest deficient-like 1 (yeast)	MAD1L1	EG1	1	8	1	4	2,72E-05	0,0029338410,0540847830,274766231
207713_s_at	NM_006462	Hs.247280	chromosome 20 open reading frame 18	C20orf18	EG1	1	8	2	3	2,72E-05	0,0029338410,0540847830,274766231
210048_at	BC001889	Hs.464622	N-ethylmaleimide-sensitive factor attachment protei	NAPG	EG1	6	3	2	3	2,72E-05	0,0029338410,0540847830,274766231
217370_x_at	S75762	Hs.513522	fusion (involved in t(12, 16) in malignant liposarcoma)	FUS	EG1	7	2	3	2	2,72E-05	0,0029338410,0540847830,274766231
217936_at	AW044631	Hs.525287	Rho GTPase activating protein 5	ARHGAP5	EG1	7	2	0	5	2,72E-05	0,0029338410,0540847830,274766231
220934_s_at	NM_024084	Hs.530463		MGC3196	EG1	2	7	3	2	2,72E-05	0,0029338410,0540847830,274766231
221919_at	AW450929	Hs.210850, H	heterogeneous nuclear ribonucleoprotein A1	HNRPA1	EG1	9	0	4	1	2,72E-05	0,0029338410,0540847830,274766231
222872_x_at	AU157541	Hs.118183		FLJ22833	EG1	6	3	3	2	2,72E-05	0,0029338410,0540847830,274766231
223025_s_at	AF290613	Hs.71040	adaptor-related protein complex 1, mu 1 subunit	AP1M1	EG1	4	5	2	3	2,72E-05	0,0029338410,0540847830,274766231
223839_s_at	AF132203	Hs.368641	stearoyl-CoA desaturase (delta-9-desaturase)	SCD	EG1	3	6	1	4	2,72E-05	0,0029338410,0540847830,274766231
224315_at	AL133598	Hs.485810	DEAD (Asp-Glu-Ala-Asp) box polypeptide 20	DDX20	EG1	8	1	3	2	2,72E-05	0,0029338410,0540847830,274766231
225539_at	AP001745	Hs.434947	zinc finger protein 295	ZNF295	EG1	9	0	4	1	2,72E-05	0,0029338410,0540847830,274766231
225747_at	AL521634	Hs.4096		FLJ32452	EG1	6	3	4	1	2,72E-05	0,0029338410,0540847830,274766231
225766_s_at	AA527296	Hs.482497	transportin 1	TNPO1	EG1	7	2	2	3	2,72E-05	0,0029338410,0540847830,274766231
226018_at	W73230	Hs.200100		Ells1	EG1	6	3	2	3	2,72E-05	0,0029338410,0540847830,274766231
226541_at	AI808182	Hs.421095	F-box protein 30	FBXO30	EG1	9	0	4	1	2,72E-05	0,0029338410,0540847830,274766231
228812_at	AI652899				EG1	7	2	3	2	2,72E-05	0,0029338410,0540847830,274766231
229151_at	BE673587	Hs.101307	solute carrier family 14 (urea transporter), member 1 (SLC14A1	EG1	6	3	4	1	2,72E-05	0,0029338410,0540847830,274766231
231252_at	AI680874	Hs.122934		FLJ23861	EG1	6	3	3	2	2,72E-05	0,0029338410,0540847830,274766231
231793_s_at	AA448956	Hs.144114	calcium/calmodulin-dependent protein kinase (CaM k	CAMK2D	EG1	7	2	3	2	2,72E-05	0,0029338410,0540847830,274766231
232092_at	AI268054	Hs.46791	mitochondrial carrier triple repeat 1	MCART1	EG1	8	1	3	2	2,72E-05	0,0029338410,0540847830,274766231
233406_at	AK022100	Hs.9997		KIAA0256	EG1	2	7	3	2	2,72E-05	0,0029338410,0540847830,274766231
235067_at	D81987	Hs.44693, Hs	muskelin 1, intracellular mediator containing kelch m	MKLN1	EG1	7	2	3	2	2,72E-05	0,0029338410,0540847830,274766231
235848_x_at	N35250				EG1	6	3	1	4	2,72E-05	0,0029338410,0540847830,274766231
239334_at	AI349167	Hs.490795	family with sequence similarity 62 (C2 domain contain	FAM62B	EG1	5	4	4	1	2,72E-05	0,0029338410,0540847830,274766231
239630_at	BF516583				EG1	4	5	3	2	2,72E-05	0,0029338410,0540847830,274766231
239835_at	AA669114	Hs.116665	kelch repeat and BTB (POZ) domain containing 8	KBTD8	EG1	5	4	4	1	2,72E-05	0,0029338410,0540847830,274766231
239979_at	BE645480	Hs.546467	epithelial stromal interaction 1 (breast)	EPSTI1	EG1	3	6	3	2	2,72E-05	0,0029338410,0540847830,274766231
240141_at	BF062399				EG1	4	5	4	1	2,72E-05	0,0029338410,0540847830,274766231
241388_at	AL567118				EG1	6	3	2	3	2,72E-05	0,0029338410,0540847830,274766231

241837_at	AI289774	Hs.535297	AT rich interactive domain 5B (MRF1-like)	ARID5B	EG1	4	5	3	2	2,72E-05	0,0029338410,0540847830,274766231
242539_at	AW665509	Hs.471637		MGC42174	EG1	5	4	2	3	2,72E-05	0,0029338410,0540847830,274766231
243659_at	N63876	Hs.192221,	H elongation factor, RNA polymerase II, 2	ELL2	EG1	5	4	3	2	2,72E-05	0,0029338410,0540847830,274766231
AFFX-PheX-5_	AFFX-PHEX-5				EG1	5	4	2	3	2,72E-05	0,0029338410,0540847830,274766231
1555878_at	AK094613	Hs.356794	ribosomal protein S24	RPS24	EG1	8	1	3	1	2,72E-05	0,0029338410,1581335050,559565828
1561640_at	BC008637	Hs.193725	proteasome (prosome, macropain) 26S subunit, non-A	PSMD5	EG1	6	3	2	2	2,72E-05	0,0029338410,1581335050,559565828
201660_at	AL525798	Hs.471461	acyl-CoA synthetase long-chain family member 3	ACSL3	EG1	9	0	1	3	2,72E-05	0,0029338410,1581335050,559565828
204824_at	NM_004435	Hs.420106	endonuclease G	ENDOG	EG1	2	7	1	3	2,72E-05	0,0029338410,1581335050,559565828
205241_at	NM_005138	Hs.549099	SCO cytochrome oxidase deficient homolog 2 (yeast)	SCO2	EG1	2	7	3	1	2,72E-05	0,0029338410,1581335050,559565828
205249_at	NM_000399	Hs.1395	early growth response 2 (Krox-20 homolog, Drosophila)	EGR2	EG1	3	6	3	1	2,72E-05	0,0029338410,1581335050,559565828
206613_s_at	NM_005681	Hs.153088	TATA box binding protein (TBP)-associated factor, RN	TAF1A	EG1	7	2	2	2	2,72E-05	0,0029338410,1581335050,559565828
209482_at	BC001430	Hs.416994	processing of precursor 7, ribonuclease P subunit (S. c	POP7	EG1	3	6	2	2	2,72E-05	0,0029338410,1581335050,559565828
209919_x_at	L20490	Hs.287505,	H gamma-glutamyltransferase 1	GGT1	EG1	3	6	3	1	2,72E-05	0,0029338410,1581335050,559565828
210075_at	AF151074	Hs.445113	membrane-associated ring finger (C3HC4) 2	MARCH2	EG1	3	6	2	2	2,72E-05	0,0029338410,1581335050,559565828
210716_s_at	M97501	Hs.524809	restin (Reed-Steinberg cell-expressed intermediate fi	RSN	EG1	9	0	3	1	2,72E-05	0,0029338410,1581335050,559565828
212930_at	AW576457	Hs.506276	ATPase, Ca++ transporting, plasma membrane 1	ATP2B1	EG1	8	1	1	3	2,72E-05	0,0029338410,1581335050,559565828
215043_s_at	X83301	Hs.519366,	Hs.553981	SMA5	EG1	2	7	2	2	2,72E-05	0,0029338410,1581335050,559565828
215567_at	AU144919	Hs.343173	chromosome 14 open reading frame 111	C14orf111	EG1	6	3	3	1	2,72E-05	0,0029338410,1581335050,559565828
217954_s_at	NM_015153	Hs.348921	PHD finger protein 3	PHF3	EG1	9	0	1	3	2,72E-05	0,0029338410,1581335050,559565828
218070_s_at	NM_013335	Hs.27059	GDP-mannose pyrophosphorylase A	GMPPA	EG1	1	8	2	2	2,72E-05	0,0029338410,1581335050,559565828
218399_s_at	NM_017955	Hs.34045	cell division cycle associated 4	CDCA4	EG1	5	4	2	2	2,72E-05	0,0029338410,1581335050,559565828
218529_at	NM_016579	Hs.333427	CD320 antigen	CD320	EG1	3	6	3	1	2,72E-05	0,0029338410,1581335050,559565828
219164_s_at	NM_018036	Hs.168241	chromosome 14 open reading frame 103	C14orf103	EG1	7	2	2	2	2,72E-05	0,0029338410,1581335050,559565828
220368_s_at	NM_017936	Hs.533887	KIAA2010	KIAA2010	EG1	9	0	3	1	2,72E-05	0,0029338410,1581335050,559565828
221826_at	BE671941	Hs.157078	angel homolog 2 (Drosophila)	ANGEL2	EG1	8	1	1	3	2,72E-05	0,0029338410,1581335050,559565828
224582_s_at	H09085	Hs.213061	nuclear casein kinase and cyclin-dependent kinase su	NUCKS1	EG1	8	1	3	1	2,72E-05	0,0029338410,1581335050,559565828
225197_at	W58461	Hs.221497		PRO0149	EG1	8	1	0	4	2,72E-05	0,0029338410,1581335050,559565828
225870_s_at	BF569208	Hs.432413	trafficking protein particle complex 5	TRAPPC5	EG1	2	7	1	3	2,72E-05	0,0029338410,1581335050,559565828
225898_at	AK023015	Hs.554894	WD repeat domain 54	WDR54	EG1	1	8	4	0	2,72E-05	0,0029338410,1581335050,559565828
227031_at	AV681975	Hs.487648	sorting nexin 13	SNX13	EG1	8	1	2	2	2,72E-05	0,0029338410,1581335050,559565828
227751_at	AI817145	Hs.443831,	H programmed cell death 5	PDCD5	EG1	5	4	3	1	2,72E-05	0,0029338410,1581335050,559565828
230850_at	AI652647	Hs.179838	formin-like 3	FMNL3	EG1	8	1	3	1	2,72E-05	0,0029338410,1581335050,559565828
230937_at	AW008207	Hs.130838		LOC442200	EG1	2	7	1	3	2,72E-05	0,0029338410,1581335050,559565828
235698_at	AA771779	Hs.461074	zinc finger protein 90 homolog (mouse)	ZFP90	EG1	4	5	3	1	2,72E-05	0,0029338410,1581335050,559565828
235791_x_at	AU155298	Hs.519474	chromodomain helicase DNA binding protein 1	CHD1	EG1	8	1	2	2	2,72E-05	0,0029338410,1581335050,559565828
236059_at	AW168168	Hs.146079		FLJ23342	EG1	8	1	2	2	2,72E-05	0,0029338410,1581335050,559565828
239106_at	AI801563	Hs.532326	carbonic anhydrase VB-like	CA5BL	EG1	9	0	3	1	2,72E-05	0,0029338410,1581335050,559565828
239437_at	AI915827				EG1	8	1	3	1	2,72E-05	0,0029338410,1581335050,559565828
240600_at	AI342146	Hs.532091	adaptor-related protein complex 3, beta 1 subunit	AP3B1	EG1	2	7	4	0	2,72E-05	0,0029338410,1581335050,559565828
243709_at	BG054799	Hs.128696		FLJ90709	EG1	3	6	2	2	2,72E-05	0,0029338410,1581335050,559565828
243788_at	AA789293	Hs.369039	PHD finger protein 11	PHF11	EG1	1	8	2	2	2,72E-05	0,0029338410,1581335050,559565828
1554229_at	AY174896	Hs.484195	adult retina protein	LOC153222	EG1	4	5	2	1	2,72E-05	0,0029338410,3636349230,911710613
1559025_at	AL833770	Hs.440932	septin 9	SEPT9	EG1	7	2	3	0	2,72E-05	0,0029338410,3636349230,911710613
1563497_at	AL833500	Hs.473370	ubiquitin specific peptidase 25	USP25	EG1	5	4	2	1	2,72E-05	0,0029338410,3636349230,911710613
201284_s_at	NM_001640	Hs.517969	N-acylaminoacyl-peptide hydrolase	APEH	EG1	1	8	1	2	2,72E-05	0,0029338410,3636349230,911710613

202331_at	NM_000709	Hs.433307	branched chain keto acid dehydrogenase E1, alpha pol	BCKDHA	EG1	0	9	1	2	2,72E-05	0,0029338410,3636349230,911710613
202928_s_at	NM_024165	Hs.166204	PHD finger protein 1	PHF1	EG1	5	4	1	2	2,72E-05	0,0029338410,3636349230,911710613
203189_s_at	NM_002496	Hs.90443	NADH dehydrogenase (ubiquinone) Fe-S protein 8, 23	NDUFS8	EG1	2	7	0	3	2,72E-05	0,0029338410,3636349230,911710613
206976_s_at	NM_006644	Hs.36927	heat shock 105kDa/110kDa protein 1	HSPH1	EG1	9	0	2	1	2,72E-05	0,0029338410,3636349230,911710613
210950_s_at	BC003573	Hs.546253	farnesyl-diphosphate farnesyltransferase 1	FDFT1	EG1	2	7	1	2	2,72E-05	0,0029338410,3636349230,911710613
222569_at	AU153746	Hs.34180	UDP-glucose ceramide glucosyltransferase-like 1	UGCGL1	EG1	7	2	0	3	2,72E-05	0,0029338410,3636349230,911710613
223109_at	BC001457	Hs.555929	TruB pseudouridine (psi) synthase homolog 2 (E. coli)	TRUB2	EG1	1	8	2	1	2,72E-05	0,0029338410,3636349230,911710613
223585_x_at	AF151831	Hs.372541,	kelch repeat and BTB (POZ) domain containing 2	KBTBD2	EG1	8	1	3	0	2,72E-05	0,0029338410,3636349230,911710613
224489_at	BC006271	Hs.405514		LOC284058	EG1	5	4	2	1	2,72E-05	0,0029338410,3636349230,911710613
225830_at	AI052247	Hs.501149	PDZ domain containing 8	PDZK8	EG1	8	1	3	0	2,72E-05	0,0029338410,3636349230,911710613
225878_at	AL135264	Hs.97858	kinesin family member 1B	KIF1B	EG1	9	0	2	1	2,72E-05	0,0029338410,3636349230,911710613
226370_at	BG149487	Hs.495854	kelch-like 15 (Drosophila)	KLHL15	EG1	8	1	1	2	2,72E-05	0,0029338410,3636349230,911710613
226942_at	AI742668	Hs.304362	PHD finger protein 20-like 1	PHF20L1	EG1	8	1	2	1	2,72E-05	0,0029338410,3636349230,911710613
228655_at	BE466077				EG1	6	3	2	1	2,72E-05	0,0029338410,3636349230,911710613
230970_at	AA975530	Hs.335205	slingshot homolog 2 (Drosophila)	SSH2	EG1	4	5	1	2	2,72E-05	0,0029338410,3636349230,911710613
232628_at	AI740629	Hs.97270,	Hs family with sequence similarity 13, member A1	FAM13A1	EG1	4	5	2	1	2,72E-05	0,0029338410,3636349230,911710613
235390_at	AA398321	Hs.519343		P18SRP	EG1	9	0	3	0	2,72E-05	0,0029338410,3636349230,911710613
235557_at	AW082827	Hs.348629		LOC150763	EG1	5	4	0	3	2,72E-05	0,0029338410,3636349230,911710613
235860_at	AA719526	Hs.345694	potassium channel modulatory factor 1	KCMF1	EG1	8	1	3	0	2,72E-05	0,0029338410,3636349230,911710613
236310_at	BE645334	Hs.278064,	zinc finger protein 37b (KOX 21)	ZNF37B	EG1	8	1	2	1	2,72E-05	0,0029338410,3636349230,911710613
238032_at	T68858	Hs.289347	dehydrogenase/reductase (SDR family) member 3	DHRS3	EG1	4	5	0	3	2,72E-05	0,0029338410,3636349230,911710613
238496_at	AA741074	Hs.32099	Wolf-Hirschhorn syndrome candidate 1-like 1	WHSC1L1	EG1	6	3	1	2	2,72E-05	0,0029338410,3636349230,911710613
239070_at	AI434790				EG1	2	7	0	3	2,72E-05	0,0029338410,3636349230,911710613
AFFX-LysX-3_a	AFFX-LYSX-3				EG1	4	5	2	1	2,72E-05	0,0029338410,3636349230,911710613
1559946_s_at	AI860098	Hs.515846	RuvB-like 2 (E. coli)	RUVBL2	EG1	2	7	1	1	2,72E-05	0,0029338410,6490161710,911710613
200822_x_at	NM_000365	Hs.524219	triosephosphate isomerase 1	TPI1	EG1	1	8	0	2	2,72E-05	0,0029338410,6490161710,911710613
204080_at	NM_025077	Hs.288198	target of EGR1, member 1 (nuclear)	TOE1	EG1	4	5	2	0	2,72E-05	0,0029338410,6490161710,911710613
212436_at	AI967961	Hs.26837	tripartite motif-containing 33	TRIM33	EG1	7	2	2	0	2,72E-05	0,0029338410,6490161710,911710613
213672_at	AA621558	Hs.355867	methionine-tRNA synthetase	MARS	EG1	7	2	2	0	2,72E-05	0,0029338410,6490161710,911710613
216960_s_at	AL049646	Hs.472221	zinc finger protein 133 (clone pHZ-13)	ZNF133	EG1	0	9	2	0	2,72E-05	0,0029338410,6490161710,911710613
219433_at	NM_017745	Hs.186424	BCL6 co-repressor	BCOR	EG1	7	2	2	0	2,72E-05	0,0029338410,6490161710,911710613
222133_s_at	AK022280	Hs.304362	PHD finger protein 20-like 1	PHF20L1	EG1	8	1	1	1	2,72E-05	0,0029338410,6490161710,911710613
222580_at	AK023596	Hs.173001	zinc finger protein 644	ZNF644	EG1	8	1	1	1	2,72E-05	0,0029338410,6490161710,911710613
223515_s_at	AL136726	Hs.299208	coenzyme Q3 homolog, methyltransferase (yeast)	COQ3	EG1	7	2	1	1	2,72E-05	0,0029338410,6490161710,911710613
224956_at	AI743689	Hs.462598		182-FIP	EG1	9	0	2	0	2,72E-05	0,0029338410,6490161710,911710613
225565_at	AA769455	Hs.516646	cAMP responsive element binding protein 1	CREB1	EG1	1	8	1	1	2,72E-05	0,0029338410,6490161710,911710613
225743_at	AI125996	Hs.33719	RNA pseudouridylate synthase domain containing 3	RPUSD3	EG1	3	6	2	0	2,72E-05	0,0029338410,6490161710,911710613
225821_s_at	AI744726	Hs.106204		KIAA1327	EG1	8	1	1	1	2,72E-05	0,0029338410,6490161710,911710613
226934_at	AU149663	Hs.369606	cleavage and polyadenylation specific factor 6, 68kDa	CPSF6	EG1	8	1	0	2	2,72E-05	0,0029338410,6490161710,911710613
228764_s_at	AI813313	Hs.220963		MGC5987	EG1	2	7	0	2	2,72E-05	0,0029338410,6490161710,911710613
230120_s_at	AI088455	Hs.528525		LOC400966	EG1	7	2	1	1	2,72E-05	0,0029338410,6490161710,911710613
235689_at	AL563572	Hs.531615	mitochondrial methionyl-tRNA formyltransferase	MTFMT	EG1	3	6	1	1	2,72E-05	0,0029338410,6490161710,911710613
239946_at	AA776723	Hs.205572		KIAA0922	EG1	7	2	0	2	2,72E-05	0,0029338410,6490161710,911710613
200776_s_at	AL518328	Hs.529231		LOC151579	EG1	9	0	1	0	2,72E-05	0,0029338410,8976818490,911710613
229914_at	AU148141	Hs.485420		FLJ38717	EG1	8	1	0	1	2,72E-05	0,0029338410,8976818490,911710613

231319_x_at	AI657069	Hs.373947	kinesin family member 9	KIF9	EG1	8	1	0	1	2,72E-05	0,002933841	0,897681849	0,911710613
232709_at	AU156167				EG1	9	0	1	0	2,72E-05	0,002933841	0,897681849	0,911710613
236629_at	AW243177	Hs.237017	chromosome 1 open reading frame 69	C1orf69	EG1	6	3	1	0	2,72E-05	0,002933841	0,897681849	0,911710613
1555858_at	CA430188				EG1 + EG2	12	3	7	2	5,34E-13	0,002933841	8,42E-05	0,002977773
202503_s_at	NM_014736	Hs.81892	KIAA0101	KIAA0101	EG1 + EG2	7	7	6	7	2,15E-11	0,002933841	4,56E-09	0,001562433
209773_s_at	BC001886	Hs.226390	ribonucleotide reductase M2 polypeptide	RRM2	EG1 + EG2	8	6	3	7	2,15E-11	0,002933841	9,97E-06	0,001562433
210567_s_at	BC001441	Hs.23348	S-phase kinase-associated protein 2 (p45)	SKP2	EG1 + EG2	5	9	3	7	2,15E-11	0,002933841	9,97E-06	0,001562433
1554020_at	BC010091	Hs.505202	bicaudal D homolog 1 (Drosophila)	BICD1	EG1 + EG2	10	3	7	4	6,10E-10	0,002933841	9,59E-07	0,001562433
201890_at	BE966236	Hs.226390	ribonucleotide reductase M2 polypeptide	RRM2	EG1 + EG2	7	5	4	9	1,28E-08	0,002933841	4,56E-09	0,001562433
242245_at	N90719	Hs.533853	synapse defective 1, Rho GTPase, homolog 2 (C. elega	SYDE2	EG1 + EG2	7	5	5	8	1,28E-08	0,002933841	4,56E-09	0,001562433
205822_s_at	NM_002130	Hs.397729	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (s	HMGCS1	EG1 + EG2	9	3	9	3	1,28E-08	0,002933841	7,43E-08	0,001562433
208078_s_at	NM_030751	Hs.124503	transcription factor 8 (represses interleukin 2 expressi	TCF8	EG1 + EG2	6	6	6	6	1,28E-08	0,002933841	7,43E-08	0,001562433
225687_at	BC001068	Hs.472716	chromosome 20 open reading frame 129	C20orf129	EG1 + EG2	11	1	7	4	1,28E-08	0,002933841	9,59E-07	0,001562433
238542_at	AA831769	Hs.250618	UL16 binding protein 2	ULBP2	EG1 + EG2	10	2	7	4	1,28E-08	0,002933841	9,59E-07	0,001562433
202705_at	NM_004701	Hs.194698	cyclin B2	CCNB2	EG1 + EG2	5	7	3	7	1,28E-08	0,002933841	9,97E-06	0,001562433
211367_s_at	U13699	Hs.2490	caspase 1, apoptosis-related cysteine peptidase (inter	CASP1	EG1 + EG2	2	10	7	3	1,28E-08	0,002933841	9,97E-06	0,001562433
212143_s_at	BF340228	Hs.450230	insulin-like growth factor binding protein 3	IGFBP3	EG1 + EG2	4	8	7	3	1,28E-08	0,002933841	9,97E-06	0,001562433
222118_at	AK023669	Hs.283532		BM039	EG1 + EG2	10	2	3	7	1,28E-08	0,002933841	9,97E-06	0,001562433
1559096_x_at	AK095307	Hs.216653	F-box protein 9	FBX09	EG1 + EG2	8	4	8	1	1,28E-08	0,002933841	8,42E-05	0,002977773
201694_s_at	NM_001964	Hs.326035	early growth response 1	EGR1	EG1 + EG2	6	6	7	2	1,28E-08	0,002933841	8,42E-05	0,002977773
206102_at	NM_021067	Hs.360033		PSF1	EG1 + EG2	5	7	3	6	1,28E-08	0,002933841	8,42E-05	0,002977773
222846_at	AB038995	Hs.389733	RAB8B, member RAS oncogene family	RAB8B	EG1 + EG2	10	2	6	3	1,28E-08	0,002933841	8,42E-05	0,002977773
201008_s_at	AA812232	Hs.533977	thioredoxin interacting protein	TXNIP	EG1 + EG2	2	9	7	7	2,08E-07	0,002933841	2,17E-10	0,001562433
201009_s_at	AI439556	Hs.533977	thioredoxin interacting protein	TXNIP	EG1 + EG2	2	9	7	7	2,08E-07	0,002933841	2,17E-10	0,001562433
201170_s_at	NM_003670	Hs.171825	basic helix-loop-helix domain containing, class B, 2	BHLHB2	EG1 + EG2	5	6	10	3	2,08E-07	0,002933841	4,56E-09	0,001562433
216607_s_at	U40053	Hs.417077	cytochrome P450, family 51, subfamily A, polypeptide	CYP51A1	EG1 + EG2	10	1	10	3	2,08E-07	0,002933841	4,56E-09	0,001562433
1555938_x_at	AI520969	Hs.533317	vimentin	VIM	EG1 + EG2	8	3	10	2	2,08E-07	0,002933841	7,43E-08	0,001562433
1557527_at	BU789637	Hs.149261,	H runt-related transcription factor 1 (acute myeloid leuk	RUNX1	EG1 + EG2	2	9	9	3	2,08E-07	0,002933841	7,43E-08	0,001562433
203799_at	NM_014880	Hs.130014	CD302 antigen	CD302	EG1 + EG2	7	4	6	6	2,08E-07	0,002933841	7,43E-08	0,001562433
1556821_x_at	H48516	Hs.508041	deleted in lymphocytic leukemia, 2	DLEU2	EG1 + EG2	7	4	5	6	2,08E-07	0,002933841	9,59E-07	0,001562433
213348_at	N33167	Hs.106070	cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	EG1 + EG2	5	6	5	6	2,08E-07	0,002933841	9,59E-07	0,001562433
214657_s_at	AU134977	Hs.523789		TncRNA	EG1 + EG2	8	3	6	5	2,08E-07	0,002933841	9,59E-07	0,001562433
218009_s_at	NM_003981	Hs.459362	protein regulator of cytokinesis 1	PRC1	EG1 + EG2	10	1	5	6	2,08E-07	0,002933841	9,59E-07	0,001562433
223079_s_at	AI828035	Hs.116448	glutaminase	GLS	EG1 + EG2	10	1	7	4	2,08E-07	0,002933841	9,59E-07	0,001562433
228069_at	AL138828	Hs.121536	family with sequence similarity 54, member A	FAM54A	EG1 + EG2	9	2	2	9	2,08E-07	0,002933841	9,59E-07	0,001562433
230494_at	AI671885	Hs.187946	solute carrier family 20 (phosphate transporter), mem	SLC20A1	EG1 + EG2	7	4	8	3	2,08E-07	0,002933841	9,59E-07	0,001562433
201663_s_at	NM_005496	Hs.58992	SMC4 structural maintenance of chromosomes 4-like 1	SMC4L1	EG1 + EG2	10	1	5	5	2,08E-07	0,002933841	9,97E-06	0,001562433
201790_s_at	AW150953	Hs.503134	7-dehydrocholesterol reductase	DHCR7	EG1 + EG2	7	4	7	3	2,08E-07	0,002933841	9,97E-06	0,001562433
203362_s_at	NM_002358	Hs.509523,	H MAD2 mitotic arrest deficient-like 1 (yeast)	MAD2L1	EG1 + EG2	7	4	3	7	2,08E-07	0,002933841	9,97E-06	0,001562433
205321_at	NM_001415	Hs.539684	eukaryotic translation initiation factor 2, subunit 3 ga	EIF2S3	EG1 + EG2	10	1	3	7	2,08E-07	0,002933841	9,97E-06	0,001562433
208079_s_at	NM_003158	Hs.250822	serine/threonine kinase 6	STK6	EG1 + EG2	9	2	3	7	2,08E-07	0,002933841	9,97E-06	0,001562433
217591_at	BF725121	Hs.445725	SKI-like	SKIL	EG1 + EG2	9	2	7	3	2,08E-07	0,002933841	9,97E-06	0,001562433
218039_at	NM_016359	Hs.511093	nucleolar and spindle associated protein 1	NUSAP1	EG1 + EG2	6	5	3	7	2,08E-07	0,002933841	9,97E-06	0,001562433
226568_at	AI478747	Hs.200230		LOC284611	EG1 + EG2	9	2	5	5	2,08E-07	0,002933841	9,97E-06	0,001562433
226713_at	AI247881	Hs.478682	chromosome 3 open reading frame 6	C3orf6	EG1 + EG2	5	6	9	1	2,08E-07	0,002933841	9,97E-06	0,001562433

202095_s_at	NM_001168	Hs.514527	baculoviral IAP repeat-containing 5 (survivin)	BIRC5	EG1 + EG2	6	5	3	6	2,08E-07	0,002933841	8,42E-05	0,002977773
209408_at	U63743	Hs.69360	kinesin family member 2C	KIF2C	EG1 + EG2	5	6	3	6	2,08E-07	0,002933841	8,42E-05	0,002977773
218723_s_at	NM_014059	Hs.507866		RGC32	EG1 + EG2	8	3	1	8	2,08E-07	0,002933841	8,42E-05	0,002977773
220789_s_at	NM_004749	Hs.231411	transforming growth factor beta regulator 4	TBRG4	EG1 + EG2	2	9	3	6	2,08E-07	0,002933841	8,42E-05	0,002977773
223229_at	AB032931	Hs.5199	ubiquitin-conjugating enzyme E2T (putative)	UBE2T	EG1 + EG2	5	6	2	7	2,08E-07	0,002933841	8,42E-05	0,002977773
224566_at	AI042152	Hs.523789		TncRNA	EG1 + EG2	7	4	7	2	2,08E-07	0,002933841	8,42E-05	0,002977773
227044_at	AV705507				EG1 + EG2	6	5	5	4	2,08E-07	0,002933841	8,42E-05	0,002977773
229715_at	AW006182				EG1 + EG2	8	3	8	1	2,08E-07	0,002933841	8,42E-05	0,002977773
232338_at	AU144815	Hs.285657	zinc finger protein 431	ZNF431	EG1 + EG2	8	3	5	4	2,08E-07	0,002933841	8,42E-05	0,002977773
235117_at	AI191897	Hs.40763	ankyrin repeat and SOCS box-containing 3	ASB3	EG1 + EG2	9	2	7	2	2,08E-07	0,002933841	8,42E-05	0,002977773
204205_at	NM_021822	Hs.474853	apolipoprotein B mRNA editing enzyme, catalytic poly	APOBEC3G	EG1 + EG2	0	10	8	8	2,66E-06	0,002933841	1,94E-13	0,001562433
204439_at	NM_006820	Hs.389724	interferon-induced protein 44-like	IFI44L	EG1 + EG2	3	7	8	5	2,66E-06	0,002933841	4,56E-09	0,001562433
226388_at	AI675780	Hs.148105	H transcription elongation factor A (SII), 3	TCEA3	EG1 + EG2	2	8	4	9	2,66E-06	0,002933841	4,56E-09	0,001562433
227236_at	AI743596	Hs.310458	tetraspanin 2	TSPAN2	EG1 + EG2	4	6	7	6	2,66E-06	0,002933841	4,56E-09	0,001562433
203973_s_at	NM_005195	Hs.440829	CCAAT/enhancer binding protein (C/EBP), delta	CEBPD	EG1 + EG2	1	9	3	9	2,66E-06	0,002933841	7,43E-08	0,001562433
204026_s_at	NM_007057	Hs.42650	ZW10 interactor	ZWINT	EG1 + EG2	5	5	4	8	2,66E-06	0,002933841	7,43E-08	0,001562433
221521_s_at	BC003186	Hs.433180		Pfs2	EG1 + EG2	4	6	2	10	2,66E-06	0,002933841	7,43E-08	0,001562433
224559_at	AF001540	Hs.187199	H metastasis associated lung adenocarcinoma transcript	MALAT1	EG1 + EG2	6	4	5	7	2,66E-06	0,002933841	7,43E-08	0,001562433
227099_s_at	AW276078	Hs.530443		LOC387763	EG1 + EG2	7	3	9	3	2,66E-06	0,002933841	7,43E-08	0,001562433
228003_at	AW294640	Hs.40758	RAB30, member RAS oncogene family	RAB30	EG1 + EG2	10	0	5	7	2,66E-06	0,002933841	7,43E-08	0,001562433
230664_at	H09657	Hs.496530		MGC39900	EG1 + EG2	6	4	2	10	2,66E-06	0,002933841	7,43E-08	0,001562433
205659_at	NM_014707	Hs.196054	histone deacetylase 9	HDAC9	EG1 + EG2	4	6	7	4	2,66E-06	0,002933841	9,59E-07	0,001562433
214710_s_at	BE407516	Hs.23960	cyclin B1	CCNB1	EG1 + EG2	5	5	4	7	2,66E-06	0,002933841	9,59E-07	0,001562433
227609_at	AA633203	Hs.546467	epithelial stromal interaction 1 (breast)	EPSTI1	EG1 + EG2	2	8	2	9	2,66E-06	0,002933841	9,59E-07	0,001562433
238013_at	BF347859	Hs.369123	pleckstrin homology domain containing, family A (pho	PLEKHA2	EG1 + EG2	4	6	5	6	2,66E-06	0,002933841	9,59E-07	0,001562433
1553994_at	BC015940	Hs.153952	5'-nucleotidase, ecto (CD73)	NT5E	EG1 + EG2	5	5	5	5	2,66E-06	0,002933841	9,97E-06	0,001562433
201250_s_at	NM_006516	Hs.473721	solute carrier family 2 (facilitated glucose transporter)	SLC2A1	EG1 + EG2	6	4	7	3	2,66E-06	0,002933841	9,97E-06	0,001562433
201291_s_at	AU159942	Hs.156346	topoisomerase (DNA) II alpha 170kDa	TOP2A	EG1 + EG2	6	4	3	7	2,66E-06	0,002933841	9,97E-06	0,001562433
202450_s_at	NM_000396	Hs.523594	cathepsin K (pyncnodysostosis)	CTSK	EG1 + EG2	0	10	9	1	2,66E-06	0,002933841	9,97E-06	0,001562433
207564_x_at	NM_003605	Hs.405410	O-linked N-acetylglucosamine (GlcNAc) transferase (U	OGT	EG1 + EG2	9	1	7	3	2,66E-06	0,002933841	9,97E-06	0,001562433
213158_at	AA045174				EG1 + EG2	8	2	3	7	2,66E-06	0,002933841	9,97E-06	0,001562433
213527_s_at	AI095896	Hs.301463	zinc finger protein 688	ZNF688	EG1 + EG2	2	8	6	4	2,66E-06	0,002933841	9,97E-06	0,001562433
227404_s_at	AI459194	Hs.326035	early growth response 1	EGR1	EG1 + EG2	5	5	7	3	2,66E-06	0,002933841	9,97E-06	0,001562433
227510_x_at	AL037917	Hs.187199	H metastasis associated lung adenocarcinoma transcript	MALAT1	EG1 + EG2	7	3	5	5	2,66E-06	0,002933841	9,97E-06	0,001562433
205967_at	NM_003542	Hs.46423	histone 1, H4c	HIST1H4C	EG1 + EG2	3	7	3	6	2,66E-06	0,002933841	8,42E-05	0,002977773
209970_x_at	M87507	Hs.2490	caspase 1, apoptosis-related cysteine peptidase (inter	CASP1	EG1 + EG2	2	8	7	2	2,66E-06	0,002933841	8,42E-05	0,002977773
210052_s_at	AF098158	Hs.244580	TPX2, microtubule-associated, homolog (Xenopus lae	TPX2	EG1 + EG2	5	5	3	6	2,66E-06	0,002933841	8,42E-05	0,002977773
213645_at	AF305057	Hs.180433	enolase superfamily member 1	ENOSF1	EG1 + EG2	8	2	6	3	2,66E-06	0,002933841	8,42E-05	0,002977773
215629_s_at	AA905286	Hs.269768		KIAA1799	EG1 + EG2	6	4	4	5	2,66E-06	0,002933841	8,42E-05	0,002977773
226750_at	AI767732	Hs.480712	La ribonucleoprotein domain family, member 2	LARP2	EG1 + EG2	9	1	7	2	2,66E-06	0,002933841	8,42E-05	0,002977773
227164_at	AL521786	Hs.68714	splicing factor, arginine/serine-rich 1 (splicing factor 2	SFRS1	EG1 + EG2	10	0	3	6	2,66E-06	0,002933841	8,42E-05	0,002977773
227961_at	AA130998	Hs.520898	cathepsin B	CTSB	EG1 + EG2	3	7	4	5	2,66E-06	0,002933841	8,42E-05	0,002977773
228964_at	AI692659	Hs.436023	PR domain containing 1, with ZNF domain	PRDM1	EG1 + EG2	0	10	4	5	2,66E-06	0,002933841	8,42E-05	0,002977773
237400_at	AI308101	Hs.438489	ATP synthase, H+ transporting, mitochondrial F0 comp	ATP5S	EG1 + EG2	4	6	5	4	2,66E-06	0,002933841	8,42E-05	0,002977773
242727_at	BG032269	Hs.25362	ADP-ribosylation factor-like 8	ARL8	EG1 + EG2	9	1	4	5	2,66E-06	0,002933841	8,42E-05	0,002977773

202016_at	NM_002402	Hs.270978	mesoderm specific transcript homolog (mouse)	MEST	EG1 + EG2	4	5	8	8	2,72E-05	0,002933841	1,94E-13	0,001562433
219304_s_at	NM_025208	Hs.352298	platelet derived growth factor D	PDGFD	EG1 + EG2	5	4	6	8	2,72E-05	0,002933841	2,17E-10	0,001562433
201010_s_at	NM_006472	Hs.533977	thioredoxin interacting protein	TXNIP	EG1 + EG2	2	7	7	6	2,72E-05	0,002933841	4,56E-09	0,001562433
201195_s_at	AB018009	Hs.513797	solute carrier family 7 (cationic amino acid transporter	SLC7A5	EG1 + EG2	7	2	9	4	2,72E-05	0,002933841	4,56E-09	0,001562433
219529_at	NM_004669	Hs.64746	chloride intracellular channel 3	CLIC3	EG1 + EG2	0	9	7	6	2,72E-05	0,002933841	4,56E-09	0,001562433
223220_s_at	AF307338	Hs.518200	poly (ADP-ribose) polymerase family, member 9	PARP9	EG1 + EG2	1	8	4	9	2,72E-05	0,002933841	4,56E-09	0,001562433
225606_at	AI949179	Hs.469658	BCL2-like 11 (apoptosis facilitator)	BCL2L11	EG1 + EG2	5	4	7	6	2,72E-05	0,002933841	4,56E-09	0,001562433
228582_x_at	AI475544	Hs.187199, H	metastasis associated lung adenocarcinoma transcript	MALAT1	EG1 + EG2	6	3	4	9	2,72E-05	0,002933841	4,56E-09	0,001562433
201596_x_at	NM_000224	Hs.406013	keratin 18	KRT18	EG1 + EG2	2	7	4	8	2,72E-05	0,002933841	7,43E-08	0,001562433
201688_s_at	BG389015	Hs.368433	tumor protein D52	TPD52	EG1 + EG2	1	8	9	3	2,72E-05	0,002933841	7,43E-08	0,001562433
205807_s_at	NM_020127	Hs.489922	tuftelin 1	TUFT1	EG1 + EG2	6	3	4	8	2,72E-05	0,002933841	7,43E-08	0,001562433
206085_s_at	NM_001902	Hs.19904	cystathionase (cystathionine gamma-lyase)	CTH	EG1 + EG2	7	2	8	4	2,72E-05	0,002933841	7,43E-08	0,001562433
209765_at	Y13786	Hs.483944	ADAM metallopeptidase domain 19 (meltrin beta)	ADAM19	EG1 + EG2	2	7	3	9	2,72E-05	0,002933841	7,43E-08	0,001562433
221841_s_at	BF514079	Hs.376206	Kruppel-like factor 4 (gut)	KLF4	EG1 + EG2	3	6	7	5	2,72E-05	0,002933841	7,43E-08	0,001562433
223044_at	AL136944	Hs.529285	solute carrier family 40 (iron-regulated transporter),	SLC40A1	EG1 + EG2	2	7	3	9	2,72E-05	0,002933841	7,43E-08	0,001562433
235593_at	AL546529	Hs.34871	zinc finger homeobox 1b	ZFXH1B	EG1 + EG2	8	1	7	5	2,72E-05	0,002933841	7,43E-08	0,001562433
1555860_x_at	CA430188				EG1 + EG2	8	1	8	3	2,72E-05	0,002933841	9,59E-07	0,001562433
1556035_s_at	AI201248	Hs.500775	zinc finger protein 207	ZNF207	EG1 + EG2	7	2	8	3	2,72E-05	0,002933841	9,59E-07	0,001562433
204334_at	AA488672	Hs.471221	Kruppel-like factor 7 (ubiquitous)	KLF7	EG1 + EG2	3	6	3	8	2,72E-05	0,002933841	9,59E-07	0,001562433
204825_at	NM_014791	Hs.184339	maternal embryonic leucine zipper kinase	MELK	EG1 + EG2	5	4	3	8	2,72E-05	0,002933841	9,59E-07	0,001562433
204908_s_at	NM_005178	Hs.31210	B-cell CLL/lymphoma 3	BCL3	EG1 + EG2	7	2	4	7	2,72E-05	0,002933841	9,59E-07	0,001562433
205282_at	NM_004631	Hs.444637	low density lipoprotein receptor-related protein 8, ap	LRP8	EG1 + EG2	7	2	5	6	2,72E-05	0,002933841	9,59E-07	0,001562433
206108_s_at	NM_006275	Hs.6891	splicing factor, arginine/serine-rich 6	SFRS6	EG1 + EG2	6	3	1	10	2,72E-05	0,002933841	9,59E-07	0,001562433
206385_s_at	NM_020987	Hs.499725	ankyrin 3, node of Ranvier (ankyrin G)	ANK3	EG1 + EG2	8	1	3	8	2,72E-05	0,002933841	9,59E-07	0,001562433
207563_s_at	U77413	Hs.405410	O-linked N-acetylglucosamine (GlcNAc) transferase (U	OGT	EG1 + EG2	7	2	8	3	2,72E-05	0,002933841	9,59E-07	0,001562433
212446_s_at	AI658534	Hs.506829	LAG1 longevity assurance homolog 6 (S. cerevisiae)	LASS6	EG1 + EG2	6	3	5	6	2,72E-05	0,002933841	9,59E-07	0,001562433
218782_s_at	NM_014109	Hs.370834	ATPase family, AAA domain containing 2	ATAD2	EG1 + EG2	8	1	4	7	2,72E-05	0,002933841	9,59E-07	0,001562433
223062_s_at	BC004863	Hs.494261	phosphoserine aminotransferase 1	PSAT1	EG1 + EG2	7	2	5	6	2,72E-05	0,002933841	9,59E-07	0,001562433
227478_at	BF739885	Hs.435458	SET binding protein 1	SETBP1	EG1 + EG2	3	6	6	5	2,72E-05	0,002933841	9,59E-07	0,001562433
227618_at	AI250910	Hs.496303		FLJ44635	EG1 + EG2	4	5	8	3	2,72E-05	0,002933841	9,59E-07	0,001562433
228390_at	AA489100				EG1 + EG2	7	2	2	9	2,72E-05	0,002933841	9,59E-07	0,001562433
235476_at	AW182459	Hs.212957	tripartite motif-containing 59	TRIM59	EG1 + EG2	3	6	4	7	2,72E-05	0,002933841	9,59E-07	0,001562433
201292_at	AL561834	Hs.156346	topoisomerase (DNA) II alpha 170kDa	TOP2A	EG1 + EG2	6	3	4	6	2,72E-05	0,002933841	9,97E-06	0,001562433
201650_at	NM_002276	Hs.514167	keratin 19	KRT19	EG1 + EG2	4	5	5	5	2,72E-05	0,002933841	9,97E-06	0,001562433
203158_s_at	AF097493	Hs.116448	glutaminase	GLS	EG1 + EG2	7	2	7	3	2,72E-05	0,002933841	9,97E-06	0,001562433
203186_s_at	NM_002961	Hs.81256	S100 calcium binding protein A4 (calcium protein, calv	S100A4	EG1 + EG2	1	8	5	5	2,72E-05	0,002933841	9,97E-06	0,001562433
203554_x_at	NM_004219	Hs.350966	pituitary tumor-transforming 1	PTTG1	EG1 + EG2	4	5	6	4	2,72E-05	0,002933841	9,97E-06	0,001562433
206114_at	NM_004438	Hs.371218	EPH receptor A4	EPHA4	EG1 + EG2	3	6	2	8	2,72E-05	0,002933841	9,97E-06	0,001562433
208891_at	BC003143	Hs.298654	dual specificity phosphatase 6	DUSP6	EG1 + EG2	6	3	8	2	2,72E-05	0,002933841	9,97E-06	0,001562433
212706_at	AB011110	Hs.520599	RAS p21 protein activator 4	RASA4	EG1 + EG2	1	8	4	6	2,72E-05	0,002933841	9,97E-06	0,001562433
213629_x_at	BF246115	Hs.513626	metallothionein 1F (functional)	MT1F	EG1 + EG2	2	7	6	4	2,72E-05	0,002933841	9,97E-06	0,001562433
215252_at	AW814026	Hs.273621, H	DnaJ (Hsp40) homolog, subfamily C, member 7	DNAJC7	EG1 + EG2	7	2	6	4	2,72E-05	0,002933841	9,97E-06	0,001562433
216252_x_at	Z70519	Hs.244139	Fas (TNF receptor superfamily, member 6)	FAS	EG1 + EG2	4	5	8	2	2,72E-05	0,002933841	9,97E-06	0,001562433
218894_s_at	NM_018048	Hs.104650		FLJ10292	EG1 + EG2	4	5	3	7	2,72E-05	0,002933841	9,97E-06	0,001562433
218921_at	NM_021805	Hs.501624		SIGIRR	EG1 + EG2	0	9	1	9	2,72E-05	0,002933841	9,97E-06	0,001562433

218983_at	NM_016546	Hs.525264	complement component 1, r subcomponent-like	C1RL	EG1 + EG2	1	8	2	8	2,72E-05	0,002933841	9,97E-06	0,001562433
221892_at	AK024548	Hs.463511	hexose-6-phosphate dehydrogenase (glucose 1-dehy	H6PD	EG1 + EG2	0	9	3	7	2,72E-05	0,002933841	9,97E-06	0,001562433
222217_s_at	BC003654	Hs.438723	solute carrier family 27 (fatty acid transporter), memb	SLC27A3	EG1 + EG2	2	7	4	6	2,72E-05	0,002933841	9,97E-06	0,001562433
222740_at	AI925583	Hs.370834	ATPase family, AAA domain containing 2	ATAD2	EG1 + EG2	9	0	3	7	2,72E-05	0,002933841	9,97E-06	0,001562433
225341_at	BF697312	Hs.5009	MTERF domain containing 3	MTERFD3	EG1 + EG2	8	1	5	5	2,72E-05	0,002933841	9,97E-06	0,001562433
225485_at	AJ278890	Hs.368315	testis specific, 14	TSGA14	EG1 + EG2	5	4	3	7	2,72E-05	0,002933841	9,97E-06	0,001562433
226666_at	BG434703	Hs.19156	dishevelled associated activator of morphogenesis 1	DAAM1	EG1 + EG2	4	5	3	7	2,72E-05	0,002933841	9,97E-06	0,001562433
227212_s_at	BE544837	Hs.460124	PHD finger protein 19	PHF19	EG1 + EG2	6	3	4	6	2,72E-05	0,002933841	9,97E-06	0,001562433
228793_at	BF002296	Hs.413416,	H jumonji domain containing 1C	JMJD1C	EG1 + EG2	7	2	7	3	2,72E-05	0,002933841	9,97E-06	0,001562433
229007_at	N50864			LOC283788	EG1 + EG2	5	4	6	4	2,72E-05	0,002933841	9,97E-06	0,001562433
229221_at	BE467023	Hs.502328	CD44 antigen (homing function and Indian blood grou	CD44	EG1 + EG2	7	2	5	5	2,72E-05	0,002933841	9,97E-06	0,001562433
232369_at	AF339768	Hs.125715	muscleblind-like 2 (Drosophila)	MBNL2	EG1 + EG2	3	6	9	1	2,72E-05	0,002933841	9,97E-06	0,001562433
239487_at	AI743261	Hs.468140		DKFZP564F0522	EG1 + EG2	4	5	7	3	2,72E-05	0,002933841	9,97E-06	0,001562433
241755_at	AI961429	Hs.528803	ubiquinol-cytochrome c reductase core protein II	UQCRC2	EG1 + EG2	5	4	4	6	2,72E-05	0,002933841	9,97E-06	0,001562433
244871_s_at	AW268357	Hs.132868	ubiquitin specific peptidase 32	USP32	EG1 + EG2	3	6	7	3	2,72E-05	0,002933841	9,97E-06	0,001562433
1555419_a_at	BC016828	Hs.527412	N-acylsphingosine amidohydrolase (acid ceramidase)	ASAH1	EG1 + EG2	5	4	7	2	2,72E-05	0,002933841	8,42E-05	0,002977773
1560814_a_at	AK098781	Hs.146520		MGC20481	EG1 + EG2	5	4	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773
201631_s_at	NM_003897	Hs.76095	immediate early response 3	IER3	EG1 + EG2	4	5	8	1	2,72E-05	0,002933841	8,42E-05	0,002977773
201664_at	AL136877	Hs.58992	SMC4 structural maintenance of chromosomes 4-like 1	SMC4L1	EG1 + EG2	7	2	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
202022_at	NM_005165	Hs.155247	aldolase C, fructose-bisphosphate	ALDOC	EG1 + EG2	1	8	5	4	2,72E-05	0,002933841	8,42E-05	0,002977773
202668_at	BF001670	Hs.149239	ephrin-B2	EFNB2	EG1 + EG2	7	2	2	7	2,72E-05	0,002933841	8,42E-05	0,002977773
203868_s_at	NM_001078	Hs.109225	vascular cell adhesion molecule 1	VCAM1	EG1 + EG2	6	3	5	4	2,72E-05	0,002933841	8,42E-05	0,002977773
204768_s_at	NM_004111	Hs.409065	flap structure-specific endonuclease 1	FEN1	EG1 + EG2	4	5	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
205691_at	NM_004209	Hs.435277	synaptogyrin 3	SYNGR3	EG1 + EG2	2	7	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773
205842_s_at	AF001362	Hs.434374	Janus kinase 2 (a protein tyrosine kinase)	JAK2	EG1 + EG2	6	3	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
210461_s_at	BC002448	Hs.438236,	H actin binding LIM protein 1	ABLIM1	EG1 + EG2	2	7	2	7	2,72E-05	0,002933841	8,42E-05	0,002977773
213693_s_at	AI610869	Hs.89603	mucin 1, transmembrane	MUC1	EG1 + EG2	1	8	0	9	2,72E-05	0,002933841	8,42E-05	0,002977773
214290_s_at	AI313324	Hs.530461	histone 2, H2aa	HIST2H2AA	EG1 + EG2	4	5	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
218793_s_at	NM_006746	Hs.109655	sex comb on midleg-like 1 (Drosophila)	SCML1	EG1 + EG2	9	0	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
219787_s_at	NM_018098	Hs.518299	epithelial cell transforming sequence 2 oncogene	ECT2	EG1 + EG2	6	3	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
221750_at	BG035985	Hs.397729	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (s	HMGCS1	EG1 + EG2	7	2	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773
221823_at	AL565741	Hs.482976		LOC90355	EG1 + EG2	9	0	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773
222077_s_at	AU153848	Hs.505469	Rac GTPase activating protein 1	RACGAP1	EG1 + EG2	4	5	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
224753_at	BE614410	Hs.434886	cell division cycle associated 5	CDCA5	EG1 + EG2	5	4	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
225521_at	AL137586	Hs.529280	anaphase promoting complex subunit 7	ANAPC7	EG1 + EG2	4	5	5	4	2,72E-05	0,002933841	8,42E-05	0,002977773
226333_at	AV700030	Hs.135087	interleukin 6 receptor	IL6R	EG1 + EG2	3	6	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
226350_at	AU155565	Hs.409081,	H opsin 3 (encephalopsin, panopsin)	OPN3	EG1 + EG2	7	2	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
226936_at	BG492359	Hs.486401	chromosome 6 open reading frame 173	C6orf173	EG1 + EG2	4	5	2	7	2,72E-05	0,002933841	8,42E-05	0,002977773
227606_s_at	AI638611	Hs.16229		AMSH-LP	EG1 + EG2	5	4	5	4	2,72E-05	0,002933841	8,42E-05	0,002977773
228729_at	N90191	Hs.23960	cyclin B1	CCNB1	EG1 + EG2	6	3	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
229704_at	BF111072	Hs.369263,	H androgen-induced proliferation inhibitor	APRIN	EG1 + EG2	6	3	7	2	2,72E-05	0,002933841	8,42E-05	0,002977773
230505_at	AU158212	Hs.85077		LOC145474	EG1 + EG2	4	5	7	2	2,72E-05	0,002933841	8,42E-05	0,002977773
231769_at	AF129536	Hs.464419	F-box protein 6	FBXO6	EG1 + EG2	3	6	4	5	2,72E-05	0,002933841	8,42E-05	0,002977773
232194_at	AA764787	Hs.126888	methyltransferase like 4	METTL4	EG1 + EG2	7	2	3	6	2,72E-05	0,002933841	8,42E-05	0,002977773
235693_at	D81004				EG1 + EG2	0	9	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773

236364_at	AA156297				EG1 + EG2	2	7	5	4	2,72E-05	0,002933841	8,42E-05	0,002977773
241769_at	AW962458	Hs.436873	integrin, alpha V (vitronectin receptor, alpha polypept	ITGAV	EG1 + EG2	3	6	6	3	2,72E-05	0,002933841	8,42E-05	0,002977773
242471_at	AI916641				EG1 + EG2	5	4	7	2	2,72E-05	0,002933841	8,42E-05	0,002977773
204748_at	NM_000963	Hs.196384	prostaglandin-endoperoxide synthase 2 (prostaglandi	PTGS2	EG2	1	7	8	6	0,0002234990,008816118	2,17E-10	0,001562433	
201340_s_at	AF010314	Hs.104925	ectodermal-neural cortex (with BTB-like domain)	ENC1	EG2	6	2	9	4	0,0002234990,008816118	4,56E-09	0,001562433	
201867_s_at	AW968555	Hs.495656	transducin (beta)-like 1X-linked	TBL1X	EG2	2	6	6	7	0,0002234990,008816118	4,56E-09	0,001562433	
204529_s_at	AI961231	Hs.491805		TOX	EG2	6	2	4	9	0,0002234990,008816118	4,56E-09	0,001562433	
204612_at	NM_006823	Hs.433700	protein kinase (cAMP-dependent, catalytic) inhibitor	PKIA	EG2	4	4	8	5	0,0002234990,008816118	4,56E-09	0,001562433	
210299_s_at	AF063002	Hs.435369	four and a half LIM domains 1	FHL1	EG2	3	5	1	12	0,0002234990,008816118	4,56E-09	0,001562433	
226610_at	AA722878	Hs.433422	proline rich 6	PRR6	EG2	7	1	5	8	0,0002234990,008816118	4,56E-09	0,001562433	
238599_at	AI168124	Hs.556018	interleukin-1 receptor-associated kinase 1 binding pro	IRAK1BP1	EG2	3	5	3	10	0,0002234990,008816118	4,56E-09	0,001562433	
201313_at	NM_001975	Hs.511915	enolase 2 (gamma, neuronal)	ENO2	EG2	2	6	5	7	0,0002234990,008816118	7,43E-08	0,001562433	
202086_at	NM_002462	Hs.517307	myxovirus (influenza virus) resistance 1, interferon-in	MX1	EG2	1	7	2	10	0,0002234990,008816118	7,43E-08	0,001562433	
202291_s_at	NM_000900	Hs.365706	matrix Gla protein	MGP	EG2	2	6	4	8	0,0002234990,008816118	7,43E-08	0,001562433	
204279_at	NM_002800	Hs.381081	proteasome (prosome, macropain) subunit, beta type	PSMB9	EG2	3	5	4	8	0,0002234990,008816118	7,43E-08	0,001562433	
204457_s_at	NM_002048	Hs.65029	growth arrest-specific 1	GAS1	EG2	3	5	6	6	0,0002234990,008816118	7,43E-08	0,001562433	
207826_s_at	NM_002167	Hs.76884	inhibitor of DNA binding 3, dominant negative helix-l	ID3	EG2	6	2	6	6	0,0002234990,008816118	7,43E-08	0,001562433	
208442_s_at	NM_000051	Hs.435561	ataxia telangiectasia mutated (includes complementa	ATM	EG2	7	1	7	5	0,0002234990,008816118	7,43E-08	0,001562433	
210095_s_at	M31159	Hs.450230	insulin-like growth factor binding protein 3	IGFBP3	EG2	4	4	8	4	0,0002234990,008816118	7,43E-08	0,001562433	
214731_at	AB037854	Hs.485899	CTTNBP2 N-terminal like	CTTNBP2NL	EG2	5	3	11	1	0,0002234990,008816118	7,43E-08	0,001562433	
218486_at	AA149594	Hs.12229	Kruppel-like factor 11	KLF11	EG2	7	1	7	5	0,0002234990,008816118	7,43E-08	0,001562433	
223578_x_at	AF113016	Hs.187199, Hs.556259		PRO1073	EG2	5	3	7	5	0,0002234990,008816118	7,43E-08	0,001562433	
227449_at	AI799018	Hs.371218	EPH receptor A4	EPHA4	EG2	4	4	4	8	0,0002234990,008816118	7,43E-08	0,001562433	
229899_s_at	AI625235	Hs.356766, Hs.401929		HSUP1	EG2	8	0	4	8	0,0002234990,008816118	7,43E-08	0,001562433	
242273_at	AA747287				EG2	5	3	3	9	0,0002234990,008816118	7,43E-08	0,001562433	
203628_at	H05812	Hs.20573	insulin-like growth factor 1 receptor	IGF1R	EG2	3	5	1	10	0,0002234990,008816118	9,59E-07	0,001562433	
203919_at	NM_003195	Hs.505004	transcription elongation factor A (SII), 2	TCEA2	EG2	0	8	2	9	0,0002234990,008816118	9,59E-07	0,001562433	
204415_at	NM_022873	Hs.523847	interferon, alpha-inducible protein (clone IFI-6-16)	G1P3	EG2	0	8	3	8	0,0002234990,008816118	9,59E-07	0,001562433	
209031_at	AL519710	Hs.370510	immunoglobulin superfamily, member 4	IGSF4	EG2	3	5	9	2	0,0002234990,008816118	9,59E-07	0,001562433	
213164_at	AI867198	Hs.302742	mitochondrial ribosomal protein S6	MRPS6	EG2	6	2	11	0	0,0002234990,008816118	9,59E-07	0,001562433	
213459_at	AU155515	Hs.433701	ribosomal protein L37a	RPL37A	EG2	7	1	5	6	0,0002234990,008816118	9,59E-07	0,001562433	
214995_s_at	BF508948	Hs.24656	KIAA0907	KIAA0907	EG2	2	6	5	6	0,0002234990,008816118	9,59E-07	0,001562433	
224724_at	AL133001	Hs.162016	sulfatase 2	SULF2	EG2	0	8	7	4	0,0002234990,008816118	9,59E-07	0,001562433	
226876_at	AI961778	Hs.345588		MGC45871	EG2	1	7	6	5	0,0002234990,008816118	9,59E-07	0,001562433	
228846_at	AW071793	Hs.468908	MAX dimerization protein 1	MXD1	EG2	7	1	8	3	0,0002234990,008816118	9,59E-07	0,001562433	
229072_at	BF968097				EG2	6	2	1	10	0,0002234990,008816118	9,59E-07	0,001562433	
232914_s_at	AB046817	Hs.369520	synaptotagmin-like 2	SYTL2	EG2	4	4	6	5	0,0002234990,008816118	9,59E-07	0,001562433	
238931_at	AL045793	Hs.224187		LOC284009	EG2	5	3	5	6	0,0002234990,008816118	9,59E-07	0,001562433	
244766_at	BG180003	Hs.444600, Hs.460558		LAT1-3TM	EG2	6	2	8	3	0,0002234990,008816118	9,59E-07	0,001562433	
201169_s_at	BG326045	Hs.171825	basic helix-loop-helix domain containing, class B, 2	BHLHB2	EG2	5	3	7	3	0,0002234990,008816118	9,97E-06	0,001562433	
201925_s_at	NM_000574	Hs.527653	decay accelerating factor for complement (CD55, Cro	DAF	EG2	5	3	4	6	0,0002234990,008816118	9,97E-06	0,001562433	
202479_s_at	BC002637	Hs.467751	tribbles homolog 2 (Drosophila)	TRIB2	EG2	3	5	5	5	0,0002234990,008816118	9,97E-06	0,001562433	
202748_at	NM_004120	Hs.386567	guanylate binding protein 2, interferon-inducible	GBP2	EG2	0	8	1	9	0,0002234990,008816118	9,97E-06	0,001562433	
203716_s_at	M80536	Hs.368912	dipeptidylpeptidase 4 (CD26, adenosine deaminase c	DPP4	EG2	4	4	3	7	0,0002234990,008816118	9,97E-06	0,001562433	
203764_at	NM_014750	Hs.77695	discs, large homolog 7 (Drosophila)	DLG7	EG2	5	3	3	7	0,0002234990,008816118	9,97E-06	0,001562433	

204784_s_at	NM_022443	Hs.85195	myeloid leukemia factor 1	MLF1	EG2	6	2	8	2	0,0002234990,008816118	9,97E-06	0,001562433
206448_at	NM_014951	Hs.22653	zinc finger protein 365	ZNF365	EG2	3	5	5	5	0,0002234990,008816118	9,97E-06	0,001562433
208892_s_at	BC003143	Hs.298654	dual specificity phosphatase 6	DUSP6	EG2	5	3	9	1	0,0002234990,008816118	9,97E-06	0,001562433
209184_s_at	BF700086	Hs.442344	insulin receptor substrate 2	IRS2	EG2	6	2	2	8	0,0002234990,008816118	9,97E-06	0,001562433
209238_at	BE966922	Hs.530733	syntaxin 3A	STX3A	EG2	4	4	9	1	0,0002234990,008816118	9,97E-06	0,001562433
210512_s_at	AF022375	Hs.73793	vascular endothelial growth factor	VEGF	EG2	4	4	9	1	0,0002234990,008816118	9,97E-06	0,001562433
213222_at	AL049593	Hs.431173	phospholipase C, beta 1 (phosphoinositide-specific)	PLCB1	EG2	5	3	3	7	0,0002234990,008816118	9,97E-06	0,001562433
213226_at	AI346350	Hs.91728	exosome component 9	EXOSC9	EG2	6	2	4	6	0,0002234990,008816118	9,97E-06	0,001562433
216041_x_at	AK023348	Hs.514220	granulin	GRN	EG2	0	8	8	2	0,0002234990,008816118	9,97E-06	0,001562433
216894_x_at	D64137	Hs.106070	cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	EG2	6	2	4	6	0,0002234990,008816118	9,97E-06	0,001562433
219024_at	NM_021622	Hs.287830	pleckstrin homology domain containing, family A (pho	PLEKHA1	EG2	4	4	8	2	0,0002234990,008816118	9,97E-06	0,001562433
220739_s_at	NM_017623	Hs.150895	cyclin M3	CNNM3	EG2	0	8	7	3	0,0002234990,008816118	9,97E-06	0,001562433
221484_at	BF691447	Hs.370487	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, p	B4GALT5	EG2	5	3	7	3	0,0002234990,008816118	9,97E-06	0,001562433
224428_s_at	AY029179	Hs.470654	cell division cycle associated 7	CDCA7	EG2	5	3	3	7	0,0002234990,008816118	9,97E-06	0,001562433
225123_at	BE883841	Hs.120633	sestrin 3	SESN3	EG2	6	2	6	4	0,0002234990,008816118	9,97E-06	0,001562433
227004_at	AI611074	Hs.435570	cyclin-dependent kinase-like 5	CDKL5	EG2	5	3	4	6	0,0002234990,008816118	9,97E-06	0,001562433
227856_at	AI110850	Hs.23439		FLJ39370	EG2	5	3	6	4	0,0002234990,008816118	9,97E-06	0,001562433
228157_at	AI125646	Hs.500775	zinc finger protein 207	ZNF207	EG2	7	1	6	4	0,0002234990,008816118	9,97E-06	0,001562433
235197_s_at	AV713913	Hs.226780	osteopetrosis associated transmembrane protein 1	OSTM1	EG2	7	1	8	2	0,0002234990,008816118	9,97E-06	0,001562433
238623_at	AI633559	Hs.390428	mitogen-activated protein kinase kinase kinase 4	MAP3K4	EG2	7	1	9	1	0,0002234990,008816118	9,97E-06	0,001562433
241036_at	AA215451	Hs.477898	Hermansky-Pudlak syndrome 3	HPS3	EG2	6	2	8	2	0,0002234990,008816118	9,97E-06	0,001562433
242134_at	AI733194				EG2	8	0	3	7	0,0002234990,008816118	9,97E-06	0,001562433
45714_at	AA436930	Hs.412304,	H host cell factor C1 regulator 1 (XPO1 dependant)	HCFC1R1	EG2	0	8	4	6	0,0002234990,008816118	9,97E-06	0,001562433
1552619_a_at	NM_018685	Hs.62180	anillin, actin binding protein (scraps homolog, Drosop	ANLN	EG2	6	2	3	6	0,0002234990,008816118	8,42E-05	0,002977773
1555486_a_at	BC008922	Hs.19987		FLJ14213	EG2	0	8	7	2	0,0002234990,008816118	8,42E-05	0,002977773
1555978_s_at	AW467415	Hs.233936		LOC440476	EG2	5	3	4	5	0,0002234990,008816118	8,42E-05	0,002977773
1559094_at	AK095307	Hs.216653	F-box protein 9	FBXO9	EG2	6	2	6	3	0,0002234990,008816118	8,42E-05	0,002977773
1559190_s_at	AL833150	Hs.327631	retinol dehydrogenase 13 (all-trans and 9-cis)	RDH13	EG2	3	5	1	8	0,0002234990,008816118	8,42E-05	0,002977773
200632_s_at	NM_006096	Hs.372914	N-myc downstream regulated gene 1	NDRG1	EG2	1	7	5	4	0,0002234990,008816118	8,42E-05	0,002977773
200678_x_at	NM_002087	Hs.514220	granulin	GRN	EG2	0	8	7	2	0,0002234990,008816118	8,42E-05	0,002977773
201282_at	NM_002541	Hs.488181	oxoglutarate (alpha-ketoglutarate) dehydrogenase (li	OGDH	EG2	4	4	5	4	0,0002234990,008816118	8,42E-05	0,002977773
202499_s_at	NM_006931	Hs.210227,	H solute carrier family 2 (facilitated glucose transporter)	SLC2A3	EG2	4	4	5	4	0,0002234990,008816118	8,42E-05	0,002977773
203017_s_at	R52678	Hs.22587	synovial sarcoma, X breakpoint 2 interacting protein	SSX2IP	EG2	7	1	7	2	0,0002234990,008816118	8,42E-05	0,002977773
203037_s_at	NM_014751	Hs.336994	metastasis suppressor 1	MTSS1	EG2	2	6	8	1	0,0002234990,008816118	8,42E-05	0,002977773
203213_at	AL524035	Hs.334562	cell division cycle 2, G1 to S and G2 to M	CDC2	EG2	5	3	3	6	0,0002234990,008816118	8,42E-05	0,002977773
203276_at	NM_005573	Hs.89497	lamin B1	LMNB1	EG2	6	2	2	7	0,0002234990,008816118	8,42E-05	0,002977773
203755_at	NM_001211	Hs.36708	BUB1 budding uninhibited by benzimidazoles 1 homol	BUB1B	EG2	5	3	3	6	0,0002234990,008816118	8,42E-05	0,002977773
204507_s_at	NM_000945	Hs.280604	protein phosphatase 3 (formerly 2B), regulatory subu	PPP3R1	EG2	5	3	5	4	0,0002234990,008816118	8,42E-05	0,002977773
204510_at	NM_003503	Hs.533573	CDC7 cell division cycle 7 (S. cerevisiae)	CDC7	EG2	5	3	4	5	0,0002234990,008816118	8,42E-05	0,002977773
205171_at	NM_002830	Hs.469809	protein tyrosine phosphatase, non-receptor type 4 (m	PTPN4	EG2	5	3	7	2	0,0002234990,008816118	8,42E-05	0,002977773
205339_at	NM_003035	Hs.525198	TAL1 (SCL) interrupting locus	SIL	EG2	5	3	3	6	0,0002234990,008816118	8,42E-05	0,002977773
206011_at	AI719655	Hs.2490	caspase 1, apoptosis-related cysteine peptidase (inter	CASP1	EG2	2	6	7	2	0,0002234990,008816118	8,42E-05	0,002977773
208070_s_at	NM_002912	Hs.232021	REV3-like, catalytic subunit of DNA polymerase zeta (y	REV3L	EG2	6	2	2	7	0,0002234990,008816118	8,42E-05	0,002977773
209267_s_at	AB040120	Hs.288034	solute carrier family 39 (zinc transporter), member 8	SLC39A8	EG2	2	6	1	8	0,0002234990,008816118	8,42E-05	0,002977773
210559_s_at	D88357	Hs.334562	cell division cycle 2, G1 to S and G2 to M	CDC2	EG2	6	2	3	6	0,0002234990,008816118	8,42E-05	0,002977773

211991_s_at	M27487	Hs.347270	major histocompatibility complex, class II, DP alpha 1	HLA-DPA1	EG2	2	6	7	2	0,0002234990,008816118	8,42E-05	0,002977773
212141_at	AA604621	Hs.460184	MCM4 minichromosome maintenance deficient 4 (S. c	MCM4	EG2	4	4	4	5	0,0002234990,008816118	8,42E-05	0,002977773
212188_at	AA551075	Hs.109438	potassium channel tetramerisation domain containing	KCTD12	EG2	4	4	2	7	0,0002234990,008816118	8,42E-05	0,002977773
212822_at	AA121502	Hs.477420	HEG homolog 1 (zebrafish)	HEG1	EG2	3	5	7	2	0,0002234990,008816118	8,42E-05	0,002977773
213046_at	AI130920	Hs.117176	poly(A) binding protein, nuclear 1	PABPN1	EG2	7	1	5	4	0,0002234990,008816118	8,42E-05	0,002977773
213906_at	AW592266	Hs.445898	v-myb myeloblastosis viral oncogene homolog (avian)	MYBL1	EG2	6	2	4	5	0,0002234990,008816118	8,42E-05	0,002977773
214012_at	BE551138	Hs.436186, Hs.440961		ARTS-1	EG2	5	3	2	7	0,0002234990,008816118	8,42E-05	0,002977773
215388_s_at	X56210	Hs.154224	complement factor H-related 1 pseudogene	CFHL1P	EG2	1	7	4	5	0,0002234990,008816118	8,42E-05	0,002977773
217627_at	BE515346	Hs.290703	zinc finger protein 573	ZNF573	EG2	4	4	2	7	0,0002234990,008816118	8,42E-05	0,002977773
217966_s_at	NM_022083	Hs.518662	chromosome 1 open reading frame 24	C1orf24	EG2	3	5	5	4	0,0002234990,008816118	8,42E-05	0,002977773
218308_at	NM_006342	Hs.104019	transforming, acidic coiled-coil containing protein 3	TACC3	EG2	4	4	2	7	0,0002234990,008816118	8,42E-05	0,002977773
218755_at	NM_005733	Hs.73625	kinesin family member 20A	KIF20A	EG2	5	3	2	7	0,0002234990,008816118	8,42E-05	0,002977773
220987_s_at	NM_030952	Hs.497512	NUAK family, SNF1-like kinase, 2	NUAK2	EG2	7	1	4	5	0,0002234990,008816118	8,42E-05	0,002977773
222439_s_at	BE967048	Hs.160211	thyroid hormone receptor associated protein 3	THRAP3	EG2	1	7	5	4	0,0002234990,008816118	8,42E-05	0,002977773
222608_s_at	AK023208	Hs.62180	anillin, actin binding protein (scraps homolog, Drosop	ANLN	EG2	6	2	2	7	0,0002234990,008816118	8,42E-05	0,002977773
223307_at	BC002551	Hs.524216	cell division cycle associated 3	CDCA3	EG2	4	4	2	7	0,0002234990,008816118	8,42E-05	0,002977773
223940_x_at	AF132202	Hs.187199, H	metastasis associated lung adenocarcinoma transcript	MALAT1	EG2	6	2	5	4	0,0002234990,008816118	8,42E-05	0,002977773
225875_s_at	T84558	Hs.523442	NIPA-like domain containing 3	NPAL3	EG2	2	6	4	5	0,0002234990,008816118	8,42E-05	0,002977773
227100_at	AI569766	Hs.13205		B3GTL	EG2	4	4	6	3	0,0002234990,008816118	8,42E-05	0,002977773
227517_s_at	AI056992	Hs.531856	growth arrest-specific 5	GAS5	EG2	8	0	5	4	0,0002234990,008816118	8,42E-05	0,002977773
228049_x_at	AA523172				EG2	6	2	3	6	0,0002234990,008816118	8,42E-05	0,002977773
228937_at	AI659800	Hs.210586		FLJ38725	EG2	5	3	3	6	0,0002234990,008816118	8,42E-05	0,002977773
229943_at	BF939833	Hs.436922	ret finger protein 2	RFP2	EG2	3	5	5	4	0,0002234990,008816118	8,42E-05	0,002977773
230180_at	AA521056	Hs.528305	DEAD (Asp-Glu-Ala-Asp) box polypeptide 17	DDX17	EG2	4	4	3	6	0,0002234990,008816118	8,42E-05	0,002977773
230560_at	N21096	Hs.508958	syntaxin binding protein 6 (amisyn)	STXBP6	EG2	1	7	3	6	0,0002234990,008816118	8,42E-05	0,002977773
231042_s_at	AA809487	Hs.144114	calcium/calmodulin-dependent protein kinase (CaM k	CAMK2D	EG2	7	1	3	6	0,0002234990,008816118	8,42E-05	0,002977773
231817_at	H25097	Hs.431081	ubiquitin specific peptidase 53	USP53	EG2	8	0	2	7	0,0002234990,008816118	8,42E-05	0,002977773
231863_at	AF161419	Hs.489811	inhibitor of growth family, member 3	ING3	EG2	6	2	6	3	0,0002234990,008816118	8,42E-05	0,002977773
242059_at	AW976631	Hs.240056	ethanolamine kinase 1	ETNK1	EG2	6	2	6	3	0,0002234990,008816118	8,42E-05	0,002977773
242389_at	BE887449	Hs.130293, Hs.534387		CROP	EG2	5	3	6	3	0,0002234990,008816118	8,42E-05	0,002977773
244447_at	AW292830	Hs.435001	Kruppel-like factor 10	KLF10	EG2	5	3	9	0	0,0002234990,008816118	8,42E-05	0,002977773
39248_at	N74607	Hs.234642	aquaporin 3	AQP3	EG2	2	6	2	7	0,0002234990,008816118	8,42E-05	0,002977773
212810_s_at	W72527	Hs.323878	solute carrier family 1 (glutamate/neutral amino acid t	SLC1A4	EG2	6	1	11	6	0,0014790570,028294149	3,06E-15	0,001562433
225647_s_at	AI246687	Hs.128065	cathepsin C	CTSC	EG2	2	5	10	5	0,0014790570,028294149	7,74E-12	0,001562433
201397_at	NM_006623	Hs.487296	phosphoglycerate dehydrogenase	PHGDH	EG2	2	5	4	10	0,0014790570,028294149	2,17E-10	0,001562433
220576_at	NM_024989	Hs.229988		PGAP1	EG2	6	1	8	6	0,0014790570,028294149	2,17E-10	0,001562433
225842_at	AK026181	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	6	1	10	4	0,0014790570,028294149	2,17E-10	0,001562433
229450_at	AI075407	Hs.47338	interferon-induced protein with tetratricopeptide rep	IFIT3	EG2	2	5	7	7	0,0014790570,028294149	2,17E-10	0,001562433
202934_at	AI761561	Hs.76090, Hs	hexokinase 2	HK2	EG2	6	1	7	6	0,0014790570,028294149	4,56E-09	0,001562433
209189_at	BC004490	Hs.25647	v-fos FBJ murine osteosarcoma viral oncogene homol	FOS	EG2	1	6	6	7	0,0014790570,028294149	4,56E-09	0,001562433
215177_s_at	AV733308	Hs.133397	integrin, alpha 6	ITGA6	EG2	6	1	6	7	0,0014790570,028294149	4,56E-09	0,001562433
221011_s_at	NM_030915	Hs.468044		LBH	EG2	3	4	5	8	0,0014790570,028294149	4,56E-09	0,001562433
201079_at	NM_004710	Hs.464210	synaptogyrin 2	SYNGR2	EG2	2	5	6	6	0,0014790570,028294149	7,43E-08	0,001562433
202528_at	NM_000403	Hs.76057	UDP-galactose-4-epimerase	GALE	EG2	2	5	11	1	0,0014790570,028294149	7,43E-08	0,001562433
202847_at	NM_004563	Hs.75812	phosphoenolpyruvate carboxykinase 2 (mitochondrial	PCK2	EG2	0	7	4	8	0,0014790570,028294149	7,43E-08	0,001562433

203139_at	NM_004938	Hs.380277	death-associated protein kinase 1	DAPK1	EG2	2	5	7	5	0,0014790570,028294149	7,43E-08	0,001562433
203358_s_at	NM_004456	Hs.444082	enhancer of zeste homolog 2 (Drosophila)	EZH2	EG2	5	2	4	8	0,0014790570,028294149	7,43E-08	0,001562433
203498_at	NM_005822	Hs.440168	Down syndrome critical region gene 1-like 1	DSCR1L1	EG2	5	2	5	7	0,0014790570,028294149	7,43E-08	0,001562433
204011_at	NM_005842	Hs.18676	sprouty homolog 2 (Drosophila)	SPRY2	EG2	4	3	10	2	0,0014790570,028294149	7,43E-08	0,001562433
204589_at	NM_014840	Hs.524692	NUAK family, SNF1-like kinase, 1	NUAK1	EG2	3	4	8	4	0,0014790570,028294149	7,43E-08	0,001562433
210001_s_at	AB005043	Hs.50640	suppressor of cytokine signaling 1	SOCS1	EG2	4	3	2	10	0,0014790570,028294149	7,43E-08	0,001562433
228702_at	AL569506	Hs.150556		FLJ43663	EG2	6	1	11	1	0,0014790570,028294149	7,43E-08	0,001562433
229744_at	AL556611	Hs.196983	sperm specific antigen 2	SSFA2	EG2	6	1	9	3	0,0014790570,028294149	7,43E-08	0,001562433
230179_at	N52572	Hs.435694		LOC285812	EG2	5	2	11	1	0,0014790570,028294149	7,43E-08	0,001562433
230930_at	AI435839	Hs.558072		LOC338620	EG2	3	4	7	5	0,0014790570,028294149	7,43E-08	0,001562433
231775_at	W65310	Hs.401745	tumor necrosis factor receptor superfamily, member 1	TNFRSF10A	EG2	4	3	6	6	0,0014790570,028294149	7,43E-08	0,001562433
233020_at	AU154125	Hs.534212	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	SEC22L1	EG2	3	4	9	3	0,0014790570,028294149	7,43E-08	0,001562433
241763_s_at	BF244402	Hs.403933	F-box protein 32	FBXO32	EG2	5	2	4	8	0,0014790570,028294149	7,43E-08	0,001562433
1553995_a_at	BC015940	Hs.153952	5'-nucleotidase, ecto (CD73)	NT5E	EG2	5	2	7	4	0,0014790570,028294149	9,59E-07	0,001562433
1555827_at	AY034790	Hs.4859	cyclin L1	CCNL1	EG2	6	1	4	7	0,0014790570,028294149	9,59E-07	0,001562433
1555847_a_at	BU617052	Hs.436426		LOC284454	EG2	6	1	10	1	0,0014790570,028294149	9,59E-07	0,001562433
200962_at	AI348010	Hs.442657	TBC1 domain family, member 8 (with GRAM domain)	TBC1D8	EG2	7	0	3	8	0,0014790570,028294149	9,59E-07	0,001562433
203404_at	NM_014782	Hs.48924	armadillo repeat containing, X-linked 2	ARMCX2	EG2	3	4	5	6	0,0014790570,028294149	9,59E-07	0,001562433
204162_at	NM_006101	Hs.414407	kinetochore associated 2	KNTC2	EG2	4	3	4	7	0,0014790570,028294149	9,59E-07	0,001562433
205081_at	NM_001311	Hs.70327	cysteine-rich protein 1 (intestinal)	CRIP1	EG2	0	7	4	7	0,0014790570,028294149	9,59E-07	0,001562433
208690_s_at	BC000915	Hs.368525	PDZ and LIM domain 1 (elfin)	PDLIM1	EG2	2	5	5	6	0,0014790570,028294149	9,59E-07	0,001562433
209016_s_at	BC002700	Hs.411501	keratin 7	KRT7	EG2	3	4	5	6	0,0014790570,028294149	9,59E-07	0,001562433
209891_at	AF225416	Hs.421956	spindle pole body component 25 homolog (S. cerevisi	SPBC25	EG2	4	3	2	9	0,0014790570,028294149	9,59E-07	0,001562433
209911_x_at	BC002842	Hs.130853	histone 1, H2bd	HIST1H2BD	EG2	5	2	6	5	0,0014790570,028294149	9,59E-07	0,001562433
210524_x_at	AF078844				EG2	1	6	7	4	0,0014790570,028294149	9,59E-07	0,001562433
213865_at	AI378788	Hs.203691	discoïdin, CUB and LCCL domain containing 2	DCBLD2	EG2	6	1	10	1	0,0014790570,028294149	9,59E-07	0,001562433
215071_s_at	AL353759	Hs.484950	histone 1, H2ac	HIST1H2AC	EG2	5	2	5	6	0,0014790570,028294149	9,59E-07	0,001562433
217979_at	NM_014399	Hs.364544	tetraspanin 13	TSPAN13	EG2	6	1	9	2	0,0014790570,028294149	9,59E-07	0,001562433
221778_at	BE217882	Hs.308710		KIAA1718	EG2	5	2	6	5	0,0014790570,028294149	9,59E-07	0,001562433
225415_at	AA577672	Hs.518201	deltex 3-like (Drosophila)	DTX3L	EG2	2	5	4	7	0,0014790570,028294149	9,59E-07	0,001562433
227223_at	BE466173	Hs.282901	RNA-binding region (RNP1, RRM) containing 2	RNPC2	EG2	1	6	6	5	0,0014790570,028294149	9,59E-07	0,001562433
227467_at	AV697515	Hs.244940	retinol dehydrogenase 10 (all-trans)	RDH10	EG2	2	5	5	6	0,0014790570,028294149	9,59E-07	0,001562433
228948_at	T15545	Hs.371218	EPH receptor A4	EPHA4	EG2	5	2	3	8	0,0014790570,028294149	9,59E-07	0,001562433
230746_s_at	AW003173	Hs.25590	stanniocalcin 1	STC1	EG2	3	4	9	2	0,0014790570,028294149	9,59E-07	0,001562433
235652_at	AI431345	Hs.109655	sex comb on midleg-like 1 (Drosophila)	SCML1	EG2	5	2	5	6	0,0014790570,028294149	9,59E-07	0,001562433
235803_at	AA843122	Hs.370168	cytokine receptor-like factor 3	CRLF3	EG2	4	3	6	5	0,0014790570,028294149	9,59E-07	0,001562433
236769_at	BE504242	Hs.153799		LOC158402	EG2	3	4	4	7	0,0014790570,028294149	9,59E-07	0,001562433
239108_at	H16791	Hs.298851	male sterility domain containing 1	MLSTD1	EG2	1	6	1	10	0,0014790570,028294149	9,59E-07	0,001562433
1553185_at	NM_152573	Hs.129136	RAS and EF-hand domain containing	RASEF	EG2	2	5	5	5	0,0014790570,028294149	9,97E-06	0,001562433
1559883_s_at	AF147427	Hs.472630	SAM domain and HD domain 1	SAMHD1	EG2	2	5	4	6	0,0014790570,028294149	9,97E-06	0,001562433
200974_at	NM_001613	Hs.500483	actin, alpha 2, smooth muscle, aorta	ACTA2	EG2	0	7	6	4	0,0014790570,028294149	9,97E-06	0,001562433
201392_s_at	BG031974	Hs.487062	insulin-like growth factor 2 receptor	IGF2R	EG2	1	6	8	2	0,0014790570,028294149	9,97E-06	0,001562433
201752_s_at	AI763123	Hs.501012	adducin 3 (gamma)	ADD3	EG2	2	5	4	6	0,0014790570,028294149	9,97E-06	0,001562433
202393_s_at	NM_005655	Hs.435001	Kruppel-like factor 10	KLF10	EG2	5	2	9	1	0,0014790570,028294149	9,97E-06	0,001562433
202481_at	NM_004753	Hs.289347	dehydrogenase/reductase (SDR family) member 3	DHRS3	EG2	2	5	2	8	0,0014790570,028294149	9,97E-06	0,001562433

204444_at	NM_004523	Hs.8878	kinesin family member 11	KIF11	EG2	4	3	3	7	0,0014790570,028294149	9,97E-06	0,001562433
204793_at	NM_014710	Hs.522730	G protein-coupled receptor associated sorting protein	GPRASP1	EG2	6	1	1	9	0,0014790570,028294149	9,97E-06	0,001562433
205884_at	NM_000885	Hs.555880	integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VL	ITGA4	EG2	5	2	1	9	0,0014790570,028294149	9,97E-06	0,001562433
209255_at	D87454	Hs.520710		KIAA0265	EG2	4	3	7	3	0,0014790570,028294149	9,97E-06	0,001562433
209304_x_at	AF087853	Hs.110571	growth arrest and DNA-damage-inducible, beta	GADD45B	EG2	3	4	5	5	0,0014790570,028294149	9,97E-06	0,001562433
210868_s_at	BC001305	Hs.412939	ELOVL family member 6, elongation of long chain fatty	ELOVL6	EG2	6	1	4	6	0,0014790570,028294149	9,97E-06	0,001562433
212099_at	AI263909	Hs.502876	ras homolog gene family, member B	RHOB	EG2	4	3	7	3	0,0014790570,028294149	9,97E-06	0,001562433
212307_s_at	BF001665	Hs.405410	O-linked N-acetylglucosamine (GlcNAc) transferase (U	OGT	EG2	6	1	7	3	0,0014790570,028294149	9,97E-06	0,001562433
215191_at	AW836210	Hs.124147	F-box and leucine-rich repeat protein 11	FBXL11	EG2	4	3	6	4	0,0014790570,028294149	9,97E-06	0,001562433
215719_x_at	X83493	Hs.244139	Fas (TNF receptor superfamily, member 6)	FAS	EG2	3	4	8	2	0,0014790570,028294149	9,97E-06	0,001562433
217997_at	AI795908	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	6	1	8	2	0,0014790570,028294149	9,97E-06	0,001562433
218273_s_at	NM_018444	Hs.22265	protein phosphatase 2C, magnesium-dependent, cata	PPM2C	EG2	4	3	4	6	0,0014790570,028294149	9,97E-06	0,001562433
218724_s_at	NM_021809	Hs.292281	TGFB-induced factor 2 (TALE family homeobox)	TGIF2	EG2	5	2	1	9	0,0014790570,028294149	9,97E-06	0,001562433
219038_at	NM_024657	Hs.496544	MORC family CW-type zinc finger 4	MORC4	EG2	4	3	9	1	0,0014790570,028294149	9,97E-06	0,001562433
221766_s_at	AW246673	Hs.10784	family with sequence similarity 46, member A	FAM46A	EG2	5	2	7	3	0,0014790570,028294149	9,97E-06	0,001562433
222696_at	BF684446	Hs.156527	axin 2 (conductin, axil)	AXIN2	EG2	3	4	1	9	0,0014790570,028294149	9,97E-06	0,001562433
226269_at	BF002104	Hs.168950	ganglioside-induced differentiation-associated protei	GDAP1	EG2	5	2	2	8	0,0014790570,028294149	9,97E-06	0,001562433
229881_at	R41200	Hs.373857	Kruppel-like factor 12	KLF12	EG2	6	1	4	6	0,0014790570,028294149	9,97E-06	0,001562433
230958_s_at	BE670797				EG2	6	1	5	5	0,0014790570,028294149	9,97E-06	0,001562433
234989_at	AV699657	Hs.523789		TncRNA	EG2	5	2	7	3	0,0014790570,028294149	9,97E-06	0,001562433
235061_at	AV706522	Hs.291000	protein phosphatase 1K (PP2C domain containing)	PPM1K	EG2	7	0	5	5	0,0014790570,028294149	9,97E-06	0,001562433
235463_s_at	AI081356	Hs.506829	LAG1 longevity assurance homolog 6 (S. cerevisiae)	LASS6	EG2	5	2	4	6	0,0014790570,028294149	9,97E-06	0,001562433
236600_at	AI651603	Hs.440414	spastic paraplegia 20, spartin (Troyer syndrome)	SPG20	EG2	6	1	4	6	0,0014790570,028294149	9,97E-06	0,001562433
238002_at	BF342391	Hs.143600	golgi phosphoprotein 4	GOLPH4	EG2	7	0	7	3	0,0014790570,028294149	9,97E-06	0,001562433
239251_at	AW963634	Hs.429581	reticulon 4	RTN4	EG2	4	3	8	2	0,0014790570,028294149	9,97E-06	0,001562433
241722_x_at	BF724558				EG2	3	4	7	3	0,0014790570,028294149	9,97E-06	0,001562433
AFFX-HUMRG	AFFX-HUMRGE/M10098_5				EG2	3	4	2	8	0,0014790570,028294149	9,97E-06	0,001562433
1553575_at	NM_173714				EG2	1	6	3	6	0,0014790570,028294149	8,42E-05	0,002977773
1554408_a_at	BC007986	Hs.515122	thymidine kinase 1, soluble	TK1	EG2	3	4	3	6	0,0014790570,028294149	8,42E-05	0,002977773
1554478_a_at	BC033077	Hs.313917		FLJ20718	EG2	5	2	2	7	0,0014790570,028294149	8,42E-05	0,002977773
1555852_at	AI375915	Hs.352018	transporter 1, ATP-binding cassette, sub-family B (MD	TAP1	EG2	2	5	2	7	0,0014790570,028294149	8,42E-05	0,002977773
201666_at	NM_003254	Hs.522632	TIMP metalloproteinase inhibitor 1	TIMP1	EG2	0	7	6	3	0,0014790570,028294149	8,42E-05	0,002977773
202269_x_at	BC002666	Hs.62661, Hs	guanylate binding protein 1, interferon-inducible, 67k	GBP1	EG2	1	6	2	7	0,0014790570,028294149	8,42E-05	0,002977773
202270_at	NM_002053	Hs.62661, Hs	guanylate binding protein 1, interferon-inducible, 67k	GBP1	EG2	1	6	2	7	0,0014790570,028294149	8,42E-05	0,002977773
202534_x_at	NM_000791	Hs.83765, Hs	dihydrofolate reductase	DHFR	EG2	3	4	2	7	0,0014790570,028294149	8,42E-05	0,002977773
203373_at	NM_003877	Hs.485572	suppressor of cytokine signaling 2	SOCS2	EG2	2	5	6	3	0,0014790570,028294149	8,42E-05	0,002977773
203586_s_at	NM_001661	Hs.183153	ADP-ribosylation factor 4-like	ARF4L	EG2	6	1	5	4	0,0014790570,028294149	8,42E-05	0,002977773
203665_at	NM_002133	Hs.517581	heme oxygenase (decycling) 1	HMOX1	EG2	4	3	8	1	0,0014790570,028294149	8,42E-05	0,002977773
204780_s_at	AA164751	Hs.244139	Fas (TNF receptor superfamily, member 6)	FAS	EG2	2	5	7	2	0,0014790570,028294149	8,42E-05	0,002977773
205443_at	NM_003082	Hs.179312	small nuclear RNA activating complex, polypeptide 1,	SNAPC1	EG2	4	3	4	5	0,0014790570,028294149	8,42E-05	0,002977773
205518_s_at	NM_003570	Hs.484918	cytidine monophosphate-N-acetylneuraminic acid hy	CMAH	EG2	3	4	2	7	0,0014790570,028294149	8,42E-05	0,002977773
205608_s_at	U83508	Hs.369675	angiopoietin 1	ANGPT1	EG2	7	0	4	5	0,0014790570,028294149	8,42E-05	0,002977773
206561_s_at	NM_020299	Hs.116724	aldo-keto reductase family 1, member B10 (aldose red	AKR1B10	EG2	6	1	5	4	0,0014790570,028294149	8,42E-05	0,002977773
208960_s_at	BE675435	Hs.4055	Kruppel-like factor 6	KLF6	EG2	2	5	5	4	0,0014790570,028294149	8,42E-05	0,002977773
209147_s_at	AB000888	Hs.435122	phosphatidic acid phosphatase type 2A	PPAP2A	EG2	2	5	4	5	0,0014790570,028294149	8,42E-05	0,002977773

209193_at	M24779	Hs.81170	pim-1 oncogene	PIM1	EG2	2	5	1	8	0,0014790570,028294149	8,42E-05	0,002977773
209581_at	BC001387	Hs.502775	HRAS-like suppressor 3	HRASLS3	EG2	4	3	6	3	0,0014790570,028294149	8,42E-05	0,002977773
209744_x_at	AB056663	Hs.472509	itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	EG2	4	3	6	3	0,0014790570,028294149	8,42E-05	0,002977773
209806_at	BC000893	Hs.437275	histone 1, H2bk	HIST1H2BK	EG2	4	3	3	6	0,0014790570,028294149	8,42E-05	0,002977773
209862_s_at	BC001233	Hs.101014		PIG8	EG2	5	2	5	4	0,0014790570,028294149	8,42E-05	0,002977773
212196_at	AW242916	Hs.454699,	H interleukin 6 signal transducer (gp130, oncostatin M re	IL6ST	EG2	7	0	2	7	0,0014790570,028294149	8,42E-05	0,002977773
212730_at	AK026420	Hs.207106	desmuslin	DMN	EG2	4	3	5	4	0,0014790570,028294149	8,42E-05	0,002977773
212841_s_at	AI692180	Hs.16019	PTPRF interacting protein, binding protein 2 (liprin be	PPFIBP2	EG2	4	3	2	7	0,0014790570,028294149	8,42E-05	0,002977773
216028_at	AL049980	Hs.184216		DKFZP564C152	EG2	4	3	5	4	0,0014790570,028294149	8,42E-05	0,002977773
218485_s_at	NM_018389	Hs.12211	solute carrier family 35, member C1	SLC35C1	EG2	1	6	3	6	0,0014790570,028294149	8,42E-05	0,002977773
218662_s_at	NM_022346	Hs.446201,	Hs.479270	HCAP-G	EG2	4	3	3	6	0,0014790570,028294149	8,42E-05	0,002977773
218795_at	NM_016361	Hs.528084	acid phosphatase 6, lysophosphatidic	ACP6	EG2	0	7	3	6	0,0014790570,028294149	8,42E-05	0,002977773
218935_at	NM_014600	Hs.368808	EH-domain containing 3	EHD3	EG2	1	6	6	3	0,0014790570,028294149	8,42E-05	0,002977773
219000_s_at	NM_024094	Hs.315167		DCC1	EG2	3	4	3	6	0,0014790570,028294149	8,42E-05	0,002977773
219117_s_at	NM_016594	Hs.438695	FK506 binding protein 11, 19 kDa	FKBP11	EG2	6	1	7	2	0,0014790570,028294149	8,42E-05	0,002977773
219148_at	NM_018492	Hs.104741	PDZ binding kinase	PBK	EG2	5	2	3	6	0,0014790570,028294149	8,42E-05	0,002977773
220940_at	NM_025190	Hs.503490,	H KIAA1641	KIAA1641	EG2	3	4	5	4	0,0014790570,028294149	8,42E-05	0,002977773
222640_at	N26002	Hs.515840	DNA (cytosine-5-)-methyltransferase 3 alpha	DNMT3A	EG2	1	6	2	7	0,0014790570,028294149	8,42E-05	0,002977773
223222_at	BC001075	Hs.514470	solute carrier family 25 (mitochondrial deoxynucleoti	SLC25A19	EG2	4	3	5	4	0,0014790570,028294149	8,42E-05	0,002977773
223319_at	AF272663	Hs.208765	gephyrin	GPHN	EG2	6	1	3	6	0,0014790570,028294149	8,42E-05	0,002977773
223464_at	AL136918	Hs.436166	oxysterol binding protein-like 5	OSBPL5	EG2	4	3	5	4	0,0014790570,028294149	8,42E-05	0,002977773
224836_at	AL109824	Hs.516994	tumor protein p53 inducible nuclear protein 2	TP53INP2	EG2	6	1	5	4	0,0014790570,028294149	8,42E-05	0,002977773
225129_at	AW170571	Hs.339809	copine II	CPNE2	EG2	1	6	0	9	0,0014790570,028294149	8,42E-05	0,002977773
225613_at	AI096389	Hs.133539	microtubule associated serine/threonine kinase famil	MAST4	EG2	3	4	2	7	0,0014790570,028294149	8,42E-05	0,002977773
226583_at	AL560962	Hs.44817		FLJ40142	EG2	3	4	4	5	0,0014790570,028294149	8,42E-05	0,002977773
226718_at	AA001423	Hs.12264	adhesion molecule with Ig-like domain 1	AMIGO1	EG2	1	6	0	9	0,0014790570,028294149	8,42E-05	0,002977773
228370_at	BF114870	Hs.525700	small nuclear ribonucleoprotein polypeptide N	SNRPN	EG2	7	0	5	4	0,0014790570,028294149	8,42E-05	0,002977773
228667_at	AI733330	Hs.353175	1-acylglycerol-3-phosphate O-acyltransferase 4 (lysop	AGPAT4	EG2	3	4	9	0	0,0014790570,028294149	8,42E-05	0,002977773
228829_at	AI279868	Hs.12286	activating transcription factor 7	ATF7	EG2	2	5	9	0	0,0014790570,028294149	8,42E-05	0,002977773
229189_s_at	BF672306				EG2	0	7	5	4	0,0014790570,028294149	8,42E-05	0,002977773
229193_at	AA005430	Hs.130293,	Hs.534387	CROP	EG2	6	1	6	3	0,0014790570,028294149	8,42E-05	0,002977773
229490_s_at	AW271106	Hs.133294	IQ motif containing GTPase activating protein 3	IQGAP3	EG2	4	3	2	7	0,0014790570,028294149	8,42E-05	0,002977773
229553_at	AA736452	Hs.26612	phosphoglucomutase 2-like 1	PGM2L1	EG2	5	2	2	7	0,0014790570,028294149	8,42E-05	0,002977773
229748_x_at	AI380156				EG2	1	6	5	4	0,0014790570,028294149	8,42E-05	0,002977773
230083_at	AW188464	Hs.431081	ubiquitin specific peptidase 53	USP53	EG2	5	2	2	7	0,0014790570,028294149	8,42E-05	0,002977773
230097_at	AI207338	Hs.473648	phosphoribosylglycinamide formyltransferase, phosp	GART	EG2	6	1	6	3	0,0014790570,028294149	8,42E-05	0,002977773
230300_at	N22849				EG2	4	3	6	3	0,0014790570,028294149	8,42E-05	0,002977773
230305_at	AW000995				EG2	6	1	5	4	0,0014790570,028294149	8,42E-05	0,002977773
230906_at	AK024931	Hs.34421,	Hs UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-	GALNT10	EG2	4	3	8	1	0,0014790570,028294149	8,42E-05	0,002977773
231577_s_at	AW014593	Hs.62661,	Hs guanylate binding protein 1, interferon-inducible, 67k	GBP1	EG2	1	6	2	7	0,0014790570,028294149	8,42E-05	0,002977773
235113_at	AA742244	Hs.556013	peptidylprolyl isomerase (cyclophilin)-like 5	PPIL5	EG2	4	3	5	4	0,0014790570,028294149	8,42E-05	0,002977773
235315_at	AI809519	Hs.450972,	H TSC22 domain family, member 1	TSC22D1	EG2	7	0	3	6	0,0014790570,028294149	8,42E-05	0,002977773
238902_at	T85248	Hs.308480		LOC115294	EG2	5	2	4	5	0,0014790570,028294149	8,42E-05	0,002977773
239208_s_at	AW945538				EG2	2	5	2	7	0,0014790570,028294149	8,42E-05	0,002977773
240165_at	AI678013				EG2	4	3	8	1	0,0014790570,028294149	8,42E-05	0,002977773

242907_at	BF509371	Hs.386567	guanylate binding protein 2, interferon-inducible	GBP2	EG2	1	6	0	9	0,0014790570,028294149	8,42E-05	0,002977773
52940_at	AA085764	Hs.501624		SIGIRR	EG2	0	7	1	8	0,0014790570,028294149	8,42E-05	0,002977773
AFFX-r2-Bs-lys	AFFX-R2-BS-LYS-5				EG2	2	5	2	7	0,0014790570,028294149	8,42E-05	0,002977773
209610_s_at	BF340083	Hs.323878	solute carrier family 1 (glutamate/neutral amino acid t	SLC1A4	EG2	5	1	11	7	0,0078675760,079901528	2,28E-17	0,001562433
207761_s_at	NM_014033	Hs.288771		DKFZP586A0522	EG2	3	3	10	5	0,0078675760,079901528	7,74E-12	0,001562433
226142_at	AV682252	Hs.553516	GLI pathogenesis-related 1 (glioma)	GLIPR1	EG2	5	1	7	8	0,0078675760,079901528	7,74E-12	0,001562433
228708_at	BF438386	Hs.25318	RAB27B, member RAS oncogene family	RAB27B	EG2	3	3	11	4	0,0078675760,079901528	7,74E-12	0,001562433
226225_at	BE967311	Hs.483104	mutated in colorectal cancers	MCC	EG2	3	3	3	11	0,0078675760,079901528	2,17E-10	0,001562433
201487_at	NM_001814	Hs.128065	cathepsin C	CTSC	EG2	1	5	8	5	0,0078675760,079901528	4,56E-09	0,001562433
204256_at	NM_024090	Hs.412939	ELOVL family member 6, elongation of long chain fatty	ELOVL6	EG2	5	1	7	6	0,0078675760,079901528	4,56E-09	0,001562433
204749_at	NM_004538	Hs.21365	nucleosome assembly protein 1-like 3	NAP1L3	EG2	5	1	9	4	0,0078675760,079901528	4,56E-09	0,001562433
208926_at	U84246	Hs.520037	sialidase 1 (lysosomal sialidase)	NEU1	EG2	4	2	11	2	0,0078675760,079901528	4,56E-09	0,001562433
216236_s_at	AL110298	Hs.210227	solute carrier family 2 (facilitated glucose transporter)	SLC2A3	EG2	4	2	7	6	0,0078675760,079901528	4,56E-09	0,001562433
217996_at	AA576961	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	4	2	8	5	0,0078675760,079901528	4,56E-09	0,001562433
222862_s_at	BG169832	Hs.18268	adenylate kinase 5	AK5	EG2	4	2	6	7	0,0078675760,079901528	4,56E-09	0,001562433
227224_at	AW003297	Hs.496222	Ral GEF with PH domain and SH3 binding motif 2	RALGPS2	EG2	4	2	3	10	0,0078675760,079901528	4,56E-09	0,001562433
227486_at	AI086864	Hs.153952	5'-nucleotidase, ecto (CD73)	NT5E	EG2	4	2	9	4	0,0078675760,079901528	4,56E-09	0,001562433
228033_at	AI341146	Hs.416375	E2F transcription factor 7	E2F7	EG2	3	3	11	2	0,0078675760,079901528	4,56E-09	0,001562433
239614_x_at	AW173003	Hs.116448	glutaminase	GLS	EG2	4	2	6	7	0,0078675760,079901528	4,56E-09	0,001562433
242051_at	AI695695				EG2	3	3	6	7	0,0078675760,079901528	4,56E-09	0,001562433
201341_at	NM_003633	Hs.104925	ectodermal-neural cortex (with BTB-like domain)	ENC1	EG2	5	1	8	4	0,0078675760,079901528	7,43E-08	0,001562433
201540_at	NM_001449	Hs.435369	four and a half LIM domains 1	FHL1	EG2	1	5	1	11	0,0078675760,079901528	7,43E-08	0,001562433
201739_at	NM_005627	Hs.296323	serum/glucocorticoid regulated kinase	SGK	EG2	4	2	4	8	0,0078675760,079901528	7,43E-08	0,001562433
202887_s_at	NM_019058	Hs.523012	DNA-damage-inducible transcript 4	DDIT4	EG2	0	6	4	8	0,0078675760,079901528	7,43E-08	0,001562433
202912_at	NM_001124	Hs.441047	adrenomedullin	ADM	EG2	5	1	5	7	0,0078675760,079901528	7,43E-08	0,001562433
203851_at	NM_002178	Hs.274313	insulin-like growth factor binding protein 6	IGFBP6	EG2	1	5	4	8	0,0078675760,079901528	7,43E-08	0,001562433
204596_s_at	U46768	Hs.25590	stanniocalcin 1	STC1	EG2	3	3	10	2	0,0078675760,079901528	7,43E-08	0,001562433
205803_s_at	NM_003304	Hs.250687	transient receptor potential cation channel, subfamily	TRPC1	EG2	2	4	4	8	0,0078675760,079901528	7,43E-08	0,001562433
209383_at	BC003637	Hs.505777	DNA-damage-inducible transcript 3	DDIT3	EG2	5	1	10	2	0,0078675760,079901528	7,43E-08	0,001562433
217967_s_at	AF288391	Hs.518662	chromosome 1 open reading frame 24	C1orf24	EG2	2	4	6	6	0,0078675760,079901528	7,43E-08	0,001562433
219118_at	NM_016594	Hs.438695	FK506 binding protein 11, 19 kDa	FKBP11	EG2	3	3	8	4	0,0078675760,079901528	7,43E-08	0,001562433
221973_at	AI983904				EG2	4	2	7	5	0,0078675760,079901528	7,43E-08	0,001562433
221974_at	AW770748	Hs.525700	small nuclear ribonucleoprotein polypeptide N	SNRPN	EG2	5	1	6	6	0,0078675760,079901528	7,43E-08	0,001562433
222931_s_at	AI809864	Hs.555980	threonine synthase-like 1 (bacterial)	THNSL1	EG2	3	3	6	6	0,0078675760,079901528	7,43E-08	0,001562433
223196_s_at	AL136551	Hs.469543	sestrin 2	SESN2	EG2	5	1	10	2	0,0078675760,079901528	7,43E-08	0,001562433
225496_s_at	N21426	Hs.369520	synaptotagmin-like 2	SYTL2	EG2	2	4	7	5	0,0078675760,079901528	7,43E-08	0,001562433
227326_at	BE966768	Hs.250723	matrix-remodelling associated 7	MXRA7	EG2	3	3	8	4	0,0078675760,079901528	7,43E-08	0,001562433
227345_at	AI738556	Hs.213467	tumor necrosis factor receptor superfamily, member 1	TNFRSF10D	EG2	4	2	5	7	0,0078675760,079901528	7,43E-08	0,001562433
228754_at	BG150485	Hs.517819	glutamate receptor interacting protein 2	GRIP2	EG2	3	3	9	3	0,0078675760,079901528	7,43E-08	0,001562433
228774_at	AW268594	Hs.374421	chromosome 9 open reading frame 81	C9orf81	EG2	4	2	10	2	0,0078675760,079901528	7,43E-08	0,001562433
231798_at	AL575177	Hs.248201	noggin	NOG	EG2	1	5	6	6	0,0078675760,079901528	7,43E-08	0,001562433
232344_at	AK021812	Hs.553501	RAS p21 protein activator (GTPase activating protein) 1	RASA1	EG2	2	4	5	7	0,0078675760,079901528	7,43E-08	0,001562433
242751_at	N55072	Hs.120	peroxiredoxin 6	PRDX6	EG2	3	3	12	0	0,0078675760,079901528	7,43E-08	0,001562433
1552622_s_at	BQ613856	Hs.535030, Hs.556000		MGC13098	EG2	2	4	4	7	0,0078675760,079901528	9,59E-07	0,001562433
202600_s_at	AI824012	Hs.155017	nuclear receptor interacting protein 1	NRIP1	EG2	5	1	1	10	0,0078675760,079901528	9,59E-07	0,001562433

203989_x_at	NM_001992	Hs.482562	coagulation factor II (thrombin) receptor	F2R	EG2	1	5	6	5	0,0078675760,079901528	9,59E-07	0,001562433
204497_at	AB011092	Hs.391860	adenylate cyclase 9	ADCY9	EG2	1	5	4	7	0,0078675760,079901528	9,59E-07	0,001562433
204620_s_at	NM_004385	Hs.443681	chondroitin sulfate proteoglycan 2 (versican)	CSPG2	EG2	3	3	6	5	0,0078675760,079901528	9,59E-07	0,001562433
204863_s_at	BE856546	Hs.454699	H interleukin 6 signal transducer (gp130, oncostatin M re	IL6ST	EG2	6	0	2	9	0,0078675760,079901528	9,59E-07	0,001562433
205771_s_at	AL137063	Hs.486483	A kinase (PRKA) anchor protein 7	AKAP7	EG2	4	2	3	8	0,0078675760,079901528	9,59E-07	0,001562433
206744_s_at	NM_014242	Hs.530988	zinc finger, MYM-type 5	ZMYM5	EG2	1	5	7	4	0,0078675760,079901528	9,59E-07	0,001562433
208003_s_at	NM_006599	Hs.371987	nuclear factor of activated T-cells 5, tonicity-responsiv	NFAT5	EG2	2	4	5	6	0,0078675760,079901528	9,59E-07	0,001562433
208998_at	U94592	Hs.80658	uncoupling protein 2 (mitochondrial, proton carrier)	UCP2	EG2	3	3	4	7	0,0078675760,079901528	9,59E-07	0,001562433
209706_at	AF247704	Hs.55999	NK3 transcription factor related, locus 1 (Drosophila)	NKX3-1	EG2	5	1	5	6	0,0078675760,079901528	9,59E-07	0,001562433
212624_s_at	BF339445	Hs.380138	chimerin (chimaerin) 1	CHN1	EG2	3	3	3	8	0,0078675760,079901528	9,59E-07	0,001562433
213506_at	BE965369	Hs.154299	coagulation factor II (thrombin) receptor-like 1	F2RL1	EG2	0	6	2	9	0,0078675760,079901528	9,59E-07	0,001562433
214196_s_at	AA602532	Hs.523454	tripeptidyl peptidase I	TPP1	EG2	5	1	10	1	0,0078675760,079901528	9,59E-07	0,001562433
219587_at	NM_017868	Hs.288772	tetratricopeptide repeat domain 12	TTC12	EG2	3	3	4	7	0,0078675760,079901528	9,59E-07	0,001562433
221731_x_at	BF218922	Hs.443681	chondroitin sulfate proteoglycan 2 (versican)	CSPG2	EG2	3	3	6	5	0,0078675760,079901528	9,59E-07	0,001562433
223125_s_at	AL563236	Hs.497159	H chromosome 1 open reading frame 21	C1orf21	EG2	4	2	4	7	0,0078675760,079901528	9,59E-07	0,001562433
223199_at	AA404592	Hs.515032	MAP kinase interacting serine/threonine kinase 2	MKMK2	EG2	5	1	6	5	0,0078675760,079901528	9,59E-07	0,001562433
223484_at	AF228422	Hs.112242		NMES1	EG2	2	4	6	5	0,0078675760,079901528	9,59E-07	0,001562433
225081_s_at	AK022955	Hs.520245	cell division cycle associated 7-like	CDCA7L	EG2	3	3	5	6	0,0078675760,079901528	9,59E-07	0,001562433
225421_at	AI654133	Hs.356247	aminoacylase 1-like 2	ACY1L2	EG2	4	2	2	9	0,0078675760,079901528	9,59E-07	0,001562433
226085_at	AA181060	Hs.349283	chromobox homolog 5 (HP1 alpha homolog, Drosophil	CBX5	EG2	4	2	3	8	0,0078675760,079901528	9,59E-07	0,001562433
226707_at	AI925574	Hs.493164	nicotinate phosphoribosyltransferase domain contain	NAPRT1	EG2	0	6	3	8	0,0078675760,079901528	9,59E-07	0,001562433
227917_at	AW192692				EG2	0	6	2	9	0,0078675760,079901528	9,59E-07	0,001562433
228497_at	AI279062	Hs.125482	solute carrier family 22 (organic cation transporter), m	SLC22A15	EG2	1	5	10	1	0,0078675760,079901528	9,59E-07	0,001562433
229515_at	AA632147	Hs.406074	PRKC, apoptosis, WT1, regulator	PAWR	EG2	5	1	4	7	0,0078675760,079901528	9,59E-07	0,001562433
235028_at	BG288330				EG2	5	1	6	5	0,0078675760,079901528	9,59E-07	0,001562433
235106_at	AW847318	Hs.428214	mastermind-like 2 (Drosophila)	MAML2	EG2	4	2	1	10	0,0078675760,079901528	9,59E-07	0,001562433
238021_s_at	AA954994	Hs.237396		LOC388279	EG2	5	1	7	4	0,0078675760,079901528	9,59E-07	0,001562433
240690_at	AI494291	Hs.29068		DKFZp761P0423	EG2	3	3	5	6	0,0078675760,079901528	9,59E-07	0,001562433
242467_at	BF433200	Hs.529862	casein kinase 1, alpha 1	CSNK1A1	EG2	5	1	8	3	0,0078675760,079901528	9,59E-07	0,001562433
1552733_at	NM_172193	Hs.509258	kelch domain containing 1	KLHDC1	EG2	4	2	5	5	0,0078675760,079901528	9,97E-06	0,001562433
1557270_at	AA632049				EG2	6	0	3	7	0,0078675760,079901528	9,97E-06	0,001562433
1564796_at	BC017854	Hs.436298	epithelial membrane protein 1	EMP1	EG2	3	3	8	2	0,0078675760,079901528	9,97E-06	0,001562433
1569312_at	BE383308	Hs.301819	zinc finger protein 146	ZNF146	EG2	5	1	7	3	0,0078675760,079901528	9,97E-06	0,001562433
201422_at	NM_006332	Hs.14623	interferon, gamma-inducible protein 30	IFI30	EG2	1	5	8	2	0,0078675760,079901528	9,97E-06	0,001562433
201427_s_at	NM_005410	Hs.275775	selenoprotein P, plasma, 1	SEPP1	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
202497_x_at	AI631159	Hs.210227	H solute carrier family 2 (facilitated glucose transporter)	SLC2A3	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
202498_s_at	BE550486	Hs.210227	H solute carrier family 2 (facilitated glucose transporter)	SLC2A3	EG2	4	2	5	5	0,0078675760,079901528	9,97E-06	0,001562433
203408_s_at	NM_002971	Hs.517717	special AT-rich sequence binding protein 1 (binds to n	SATB1	EG2	2	4	4	6	0,0078675760,079901528	9,97E-06	0,001562433
203946_s_at	U75667	Hs.553485	arginase, type II	ARG2	EG2	5	1	6	4	0,0078675760,079901528	9,97E-06	0,001562433
204783_at	AI911434	Hs.85195	myeloid leukemia factor 1	MLF1	EG2	5	1	7	3	0,0078675760,079901528	9,97E-06	0,001562433
205128_x_at	NM_000962	Hs.201978	prostaglandin-endoperoxide synthase 1 (prostaglandi	PTGS1	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
205756_s_at	NM_000132	Hs.413083	coagulation factor VIII, procoagulant component (hem	F8	EG2	2	4	5	5	0,0078675760,079901528	9,97E-06	0,001562433
207231_at	NM_014648	Hs.409210		DZIP3	EG2	2	4	3	7	0,0078675760,079901528	9,97E-06	0,001562433
207813_s_at	NM_004110	Hs.69745	ferredoxin reductase	FDXR	EG2	1	5	4	6	0,0078675760,079901528	9,97E-06	0,001562433
207980_s_at	NM_006079	Hs.82071	Cbp/p300-interacting transactivator, with Glu/Asp-rich	CITED2	EG2	1	5	3	7	0,0078675760,079901528	9,97E-06	0,001562433

208190_s_at	NM_015925	Hs.466507		LISCH7	EG2	3	3	2	8	0,0078675760,079901528	9,97E-06	0,001562433
209754_s_at	AF113682	Hs.11355	thymopoietin	TMPO	EG2	3	3	3	7	0,0078675760,079901528	9,97E-06	0,001562433
209969_s_at	BC002704	Hs.470943	signal transducer and activator of transcription 1, 91kD	STAT1	EG2	1	5	3	7	0,0078675760,079901528	9,97E-06	0,001562433
210538_s_at	U37546	Hs.127799	baculoviral IAP repeat-containing 3	BIRC3	EG2	6	0	3	7	0,0078675760,079901528	9,97E-06	0,001562433
211085_s_at	Z25430	Hs.472838	serine/threonine kinase 4	STK4	EG2	1	5	3	7	0,0078675760,079901528	9,97E-06	0,001562433
214085_x_at	AI912583	Hs.205558	HIV-1 rev binding protein 2	HRB2	EG2	6	0	5	5	0,0078675760,079901528	9,97E-06	0,001562433
214252_s_at	AV700514	Hs.30213	ceroid-lipofuscinosis, neuronal 5	CLN5	EG2	3	3	4	6	0,0078675760,079901528	9,97E-06	0,001562433
215646_s_at	R94644	Hs.443681	chondroitin sulfate proteoglycan 2 (versican)	CSPG2	EG2	4	2	5	5	0,0078675760,079901528	9,97E-06	0,001562433
216870_x_at	AF264787	Hs.508041	deleted in lymphocytic leukemia, 2	DLEU2	EG2	5	1	4	6	0,0078675760,079901528	9,97E-06	0,001562433
217127_at	AL354872	Hs.19904	cystathionase (cystathionine gamma-lyase)	CTH	EG2	6	0	7	3	0,0078675760,079901528	9,97E-06	0,001562433
217165_x_at	M10943	Hs.513626	metallothionein 1F (functional)	MT1F	EG2	2	4	5	5	0,0078675760,079901528	9,97E-06	0,001562433
217208_s_at	AL121981	Hs.292549	discs, large homolog 1 (Drosophila)	DLG1	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
218701_at	NM_016027	Hs.118554	lactamase, beta 2	LACTB2	EG2	3	3	4	6	0,0078675760,079901528	9,97E-06	0,001562433
218726_at	NM_018410	Hs.532968		DKFZp762E1312	EG2	5	1	3	7	0,0078675760,079901528	9,97E-06	0,001562433
219209_at	NM_022168	Hs.163173	interferon induced with helicase C domain 1	IFIH1	EG2	2	4	2	8	0,0078675760,079901528	9,97E-06	0,001562433
219250_s_at	NM_013281	Hs.41296	fibronectin leucine rich transmembrane protein 3	FLRT3	EG2	0	6	4	6	0,0078675760,079901528	9,97E-06	0,001562433
219493_at	NM_024745	Hs.123253	SHC SH2-domain binding protein 1	SHCBP1	EG2	5	1	3	7	0,0078675760,079901528	9,97E-06	0,001562433
219684_at	NM_022147	Hs.43388		IFRG28	EG2	0	6	0	10	0,0078675760,079901528	9,97E-06	0,001562433
221207_s_at	NM_015678	Hs.491172	neurobeachin	NBEA	EG2	4	2	2	8	0,0078675760,079901528	9,97E-06	0,001562433
221922_at	AW195581	Hs.489353	G-protein signalling modulator 2 (AGS3-like, C. elegan	GPSM2	EG2	3	3	4	6	0,0078675760,079901528	9,97E-06	0,001562433
222787_s_at	AV705186	Hs.396358		FLJ11273	EG2	2	4	2	8	0,0078675760,079901528	9,97E-06	0,001562433
223681_s_at	AB044807	Hs.478125	InaD-like (Drosophila)	INADL	EG2	2	4	0	10	0,0078675760,079901528	9,97E-06	0,001562433
225626_at	AK000680	Hs.266175,	H phosphoprotein associated with glycosphingolipid mi	PAG1	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
226218_at	BE217880	Hs.362807	interleukin 7 receptor	IL7R	EG2	4	2	4	6	0,0078675760,079901528	9,97E-06	0,001562433
226239_at	AL573722	Hs.356187		FLJ90024	EG2	0	6	8	2	0,0078675760,079901528	9,97E-06	0,001562433
226939_at	AI202327	Hs.374216	cytoplasmic polyadenylation element binding protein	CPEB2	EG2	6	0	10	0	0,0078675760,079901528	9,97E-06	0,001562433
227124_at	AI870473				EG2	6	0	8	2	0,0078675760,079901528	9,97E-06	0,001562433
227156_at	AK025872	Hs.3327	trinucleotide repeat containing 8	TNRC8	EG2	4	2	1	9	0,0078675760,079901528	9,97E-06	0,001562433
227458_at	AI608902	Hs.521989	CD274 antigen	CD274	EG2	4	2	9	1	0,0078675760,079901528	9,97E-06	0,001562433
227462_at	BE889628	Hs.482910		LRAP	EG2	3	3	3	7	0,0078675760,079901528	9,97E-06	0,001562433
228171_s_at	AI056683	Hs.188781	pleckstrin homology domain containing, family G (wit	PLEKHG4	EG2	2	4	4	6	0,0078675760,079901528	9,97E-06	0,001562433
228480_at	AW296039	Hs.165195	VAMP (vesicle-associated membrane protein)-associa	VAPA	EG2	1	5	5	5	0,0078675760,079901528	9,97E-06	0,001562433
232235_at	AK021539	Hs.124673	chromosome 18 open reading frame 4	C18orf4	EG2	4	2	2	8	0,0078675760,079901528	9,97E-06	0,001562433
234994_at	AA088177	Hs.172870	KIAA1913	KIAA1913	EG2	3	3	5	5	0,0078675760,079901528	9,97E-06	0,001562433
235274_at	AA740632				EG2	5	1	3	7	0,0078675760,079901528	9,97E-06	0,001562433
235547_at	BG548427	Hs.507680		PFAAP5	EG2	5	1	5	5	0,0078675760,079901528	9,97E-06	0,001562433
236046_at	AA781880	Hs.293833		FLJ44896	EG2	2	4	4	6	0,0078675760,079901528	9,97E-06	0,001562433
238458_at	AI868167	Hs.403594	EF-hand domain family, member A2	EFHA2	EG2	2	4	3	7	0,0078675760,079901528	9,97E-06	0,001562433
238587_at	AI927919	Hs.444075		STS-1	EG2	3	3	4	6	0,0078675760,079901528	9,97E-06	0,001562433
239413_at	AI130715	Hs.443005		Cep152	EG2	6	0	5	5	0,0078675760,079901528	9,97E-06	0,001562433
239866_at	AA705933				EG2	2	4	4	6	0,0078675760,079901528	9,97E-06	0,001562433
240146_at	AW418562	Hs.446123	capping protein (actin filament) muscle Z-line, alpha 2	CAPZA2	EG2	4	2	6	4	0,0078675760,079901528	9,97E-06	0,001562433
240307_at	N54783	Hs.269099	far upstream element (FUSE) binding protein 1	FUBP1	EG2	2	4	6	4	0,0078675760,079901528	9,97E-06	0,001562433
241879_at	AW511222				EG2	6	0	5	5	0,0078675760,079901528	9,97E-06	0,001562433
1557081_at	AA580691	Hs.531106	RNA binding motif protein 25	RBM25	EG2	3	3	4	5	0,0078675760,079901528	8,42E-05	0,002977773

1558522_at	BC013077	Hs.533355	membrane protein, palmitoylated 6 (MAGUK p55 subf	MPP6	EG2	4	2	4	5	0,0078675760,079901528	8,42E-05	0,002977773
200897_s_at	NM_016081	Hs.151220		KIAA0992	EG2	4	2	5	4	0,0078675760,079901528	8,42E-05	0,002977773
201389_at	NM_002205	Hs.505654	integrin, alpha 5 (fibronectin receptor, alpha polypept	ITGA5	EG2	3	3	5	4	0,0078675760,079901528	8,42E-05	0,002977773
201810_s_at	AL562152	Hs.257761	SH3-domain binding protein 5 (BTK-associated)	SH3BP5	EG2	0	6	5	4	0,0078675760,079901528	8,42E-05	0,002977773
202284_s_at	NM_000389	Hs.370771	cyclin-dependent kinase inhibitor 1A (p21, Cip1)	CDKN1A	EG2	6	0	6	3	0,0078675760,079901528	8,42E-05	0,002977773
202409_at	X07868	Hs.523414, Hs.556244		LOC492304	EG2	2	4	2	7	0,0078675760,079901528	8,42E-05	0,002977773
202870_s_at	NM_001255	Hs.524947	CDC20 cell division cycle 20 homolog (S. cerevisiae)	CDC20	EG2	3	3	3	6	0,0078675760,079901528	8,42E-05	0,002977773
202973_x_at	NM_014883	Hs.97270, Hs	family with sequence similarity 13, member A1	FAM13A1	EG2	2	4	5	4	0,0078675760,079901528	8,42E-05	0,002977773
203016_s_at	AK001710	Hs.22587	synovial sarcoma, X breakpoint 2 interacting protein	SSX2IP	EG2	5	1	7	2	0,0078675760,079901528	8,42E-05	0,002977773
203603_s_at	NM_014795	Hs.34871	zinc finger homeobox 1b	ZFXH1B	EG2	5	1	6	3	0,0078675760,079901528	8,42E-05	0,002977773
203680_at	NM_002736	Hs.433068	protein kinase, cAMP-dependent, regulatory, type II,	PRKAR2B	EG2	5	1	6	3	0,0078675760,079901528	8,42E-05	0,002977773
203867_s_at	NM_018096	Hs.85570	notchless homolog 1 (Drosophila)	NLE1	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
204015_s_at	BC002671	Hs.417962	dual specificity phosphatase 4	DUSP4	EG2	2	4	3	6	0,0078675760,079901528	8,42E-05	0,002977773
204066_s_at	NM_014914	Hs.435039	centaurin, gamma 2	CENTG2	EG2	1	5	6	3	0,0078675760,079901528	8,42E-05	0,002977773
204506_at	AL544951	Hs.280604	protein phosphatase 3 (formerly 2B), regulatory subu	PPP3R1	EG2	5	1	2	7	0,0078675760,079901528	8,42E-05	0,002977773
204686_at	NM_005544	Hs.471508	insulin receptor substrate 1	IRS1	EG2	3	3	3	6	0,0078675760,079901528	8,42E-05	0,002977773
204796_at	AI825937	Hs.12451	echinoderm microtubule associated protein like 1	EML1	EG2	3	3	5	4	0,0078675760,079901528	8,42E-05	0,002977773
204900_x_at	NM_003864	Hs.413835	sin3-associated polypeptide, 30kDa	SAP30	EG2	3	3	6	3	0,0078675760,079901528	8,42E-05	0,002977773
205259_at	NM_000901	Hs.163924	nuclear receptor subfamily 3, group C, member 2	NR3C2	EG2	3	3	2	7	0,0078675760,079901528	8,42E-05	0,002977773
205398_s_at	NM_005902	Hs.36915, Hs	SMAD, mothers against DPP homolog 3 (Drosophila)	SMAD3	EG2	1	5	5	4	0,0078675760,079901528	8,42E-05	0,002977773
205802_at	NM_003304	Hs.250687	transient receptor potential cation channel, subfamily	TRPC1	EG2	3	3	2	7	0,0078675760,079901528	8,42E-05	0,002977773
207029_at	NM_000899	Hs.1048	KIT ligand	KITLG	EG2	5	1	4	5	0,0078675760,079901528	8,42E-05	0,002977773
207417_s_at	NM_003451	Hs.172979	zinc finger protein 177	ZNF177	EG2	1	5	4	5	0,0078675760,079901528	8,42E-05	0,002977773
207426_s_at	NM_003326	Hs.181097	tumor necrosis factor (ligand) superfamily, member 4	TNFSF4	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
208534_s_at	NM_006989	Hs.520599	RAS p21 protein activator 4	RASA4	EG2	1	5	4	5	0,0078675760,079901528	8,42E-05	0,002977773
208663_s_at	AI652848	Hs.368214	tetratricopeptide repeat domain 3	TTC3	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
208710_s_at	AI424923	Hs.512815	adaptor-related protein complex 3, delta 1 subunit	AP3D1	EG2	0	6	3	6	0,0078675760,079901528	8,42E-05	0,002977773
209278_s_at	L27624	Hs.438231	tissue factor pathway inhibitor 2	TFPI2	EG2	2	4	4	5	0,0078675760,079901528	8,42E-05	0,002977773
209714_s_at	AF213033	Hs.84113	cyclin-dependent kinase inhibitor 3 (CDK2-associated)	CDKN3	EG2	4	2	3	6	0,0078675760,079901528	8,42E-05	0,002977773
209782_s_at	U79283	Hs.414480	D site of albumin promoter (albumin D-box) binding p	DBP	EG2	0	6	0	9	0,0078675760,079901528	8,42E-05	0,002977773
210785_s_at	AB035482	Hs.10649	chromosome 1 open reading frame 38	C1orf38	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
211115_x_at	AB037703	Hs.533862	survival of motor neuron protein interacting protein 1	SIP1	EG2	5	1	6	3	0,0078675760,079901528	8,42E-05	0,002977773
211302_s_at	L20966	Hs.198072	phosphodiesterase 4B, cAMP-specific (phosphodieste	PDE4B	EG2	4	2	5	4	0,0078675760,079901528	8,42E-05	0,002977773
211343_s_at	M33653	Hs.211933	collagen, type XIII, alpha 1	COL13A1	EG2	3	3	6	3	0,0078675760,079901528	8,42E-05	0,002977773
211596_s_at	AB050468	Hs.518055	leucine-rich repeats and immunoglobulin-like domain	LRIG1	EG2	1	5	2	7	0,0078675760,079901528	8,42E-05	0,002977773
211602_s_at	U31110	Hs.250687	transient receptor potential cation channel, subfamily	TRPC1	EG2	4	2	5	4	0,0078675760,079901528	8,42E-05	0,002977773
211653_x_at	M33376	Hs.460260, H	aldo-keto reductase family 1, member C2 (dihydrodiol	AKR1C2	EG2	3	3	5	4	0,0078675760,079901528	8,42E-05	0,002977773
212022_s_at	BF001806	Hs.80976	antigen identified by monoclonal antibody Ki-67	MKI67	EG2	4	2	3	6	0,0078675760,079901528	8,42E-05	0,002977773
212543_at	U83115	Hs.486074	absent in melanoma 1	AIM1	EG2	4	2	3	6	0,0078675760,079901528	8,42E-05	0,002977773
212758_s_at	AI373166	Hs.124503	transcription factor 8 (represses interleukin 2 expressi	TCF8	EG2	4	2	2	7	0,0078675760,079901528	8,42E-05	0,002977773
212812_at	AI700633				EG2	5	1	2	7	0,0078675760,079901528	8,42E-05	0,002977773
212913_at	BE674960	Hs.371225	mutS homolog 5 (E. coli)	MSH5	EG2	6	0	4	5	0,0078675760,079901528	8,42E-05	0,002977773
213006_at	AV655640	Hs.440829	CCAAT/enhancer binding protein (C/EBP), delta	CEBPD	EG2	1	5	3	6	0,0078675760,079901528	8,42E-05	0,002977773
213035_at	AI081194	Hs.335239	ankyrin repeat domain 28	ANKRD28	EG2	3	3	7	2	0,0078675760,079901528	8,42E-05	0,002977773
213338_at	BF062629	Hs.35861		RIS1	EG2	4	2	8	1	0,0078675760,079901528	8,42E-05	0,002977773

213908_at	AI824078	Hs.212670, Hs.513595		LOC339005	EG2	4	2	4	5	0,0078675760,079901528	8,42E-05	0,002977773
214255_at	N35112	Hs.128041	ATPase, Class V, type 10A	ATP10A	EG2	1	5	4	5	0,0078675760,079901528	8,42E-05	0,002977773
214933_at	AA769818	Hs.408449	calcium channel, voltage-dependent, P/Q type, alpha	CACNA1A	EG2	1	5	6	3	0,0078675760,079901528	8,42E-05	0,002977773
217838_s_at	NM_016337	Hs.125867	Enah/Vasp-like	EVL	EG2	0	6	2	7	0,0078675760,079901528	8,42E-05	0,002977773
218475_at	NM_022727	Hs.552590		HTF9C	EG2	0	6	4	5	0,0078675760,079901528	8,42E-05	0,002977773
218729_at	NM_020169	Hs.478067	latexin	LXN	EG2	0	6	2	7	0,0078675760,079901528	8,42E-05	0,002977773
218747_s_at	NM_018009	Hs.504597	TAP binding protein-like	TAPBPL	EG2	0	6	6	3	0,0078675760,079901528	8,42E-05	0,002977773
219918_s_at	NM_018123	Hs.121028	asp (abnormal spindle)-like, microcephaly associated	ASPM	EG2	3	3	3	6	0,0078675760,079901528	8,42E-05	0,002977773
221561_at	L21934	Hs.496383	sterol O-acyltransferase (acyl-Coenzyme A: cholester	SOAT1	EG2	4	2	3	6	0,0078675760,079901528	8,42E-05	0,002977773
222680_s_at	AK001261	Hs.126774	denticleless homolog (Drosophila)	DTL	EG2	3	3	4	5	0,0078675760,079901528	8,42E-05	0,002977773
223194_s_at	AL512737	Hs.132340	chromosome 6 open reading frame 85	C6orf85	EG2	3	3	1	8	0,0078675760,079901528	8,42E-05	0,002977773
223257_at	AI823905	Hs.509008	KIAA1333	KIAA1333	EG2	6	0	5	4	0,0078675760,079901528	8,42E-05	0,002977773
224828_at	AV704132	Hs.127126	cytoplasmic polyadenylation element binding protein	CPEB4	EG2	4	2	8	1	0,0078675760,079901528	8,42E-05	0,002977773
224840_at	AI753747	Hs.407190	FK506 binding protein 5	FKBP5	EG2	5	1	3	6	0,0078675760,079901528	8,42E-05	0,002977773
224860_at	AL137489	Hs.7517	chromosome 9 open reading frame 123	C9orf123	EG2	2	4	5	4	0,0078675760,079901528	8,42E-05	0,002977773
224901_at	AL571375	Hs.379191	stearoyl-CoA desaturase 5	SCD5	EG2	2	4	7	2	0,0078675760,079901528	8,42E-05	0,002977773
225177_at	AA143793	Hs.191179	RAB11 family interacting protein 1 (class I)	RAB11FIP1	EG2	3	3	2	7	0,0078675760,079901528	8,42E-05	0,002977773
225214_at	AI762915	Hs.213470	proteasome (prosome, macropain) subunit, beta type	PSMB7	EG2	4	2	7	2	0,0078675760,079901528	8,42E-05	0,002977773
225582_at	AA425726	Hs.523252	KIAA1754	KIAA1754	EG2	4	2	8	1	0,0078675760,079901528	8,42E-05	0,002977773
226200_at	BE222664	Hs.102910	valyl-tRNA synthetase like	VARSL	EG2	0	6	5	4	0,0078675760,079901528	8,42E-05	0,002977773
226275_at	AI188653	Hs.468908	MAX dimerization protein 1	MXD1	EG2	5	1	8	1	0,0078675760,079901528	8,42E-05	0,002977773
226456_at	AW138157	Hs.347524		MGC24665	EG2	3	3	2	7	0,0078675760,079901528	8,42E-05	0,002977773
226905_at	BG036514	Hs.345588		MGC45871	EG2	3	3	5	4	0,0078675760,079901528	8,42E-05	0,002977773
226982_at	AI745624	Hs.192221,	H elongation factor, RNA polymerase II, 2	ELL2	EG2	6	0	5	4	0,0078675760,079901528	8,42E-05	0,002977773
227020_at	BE502982	Hs.463613	yippee-like 2 (Drosophila)	YPEL2	EG2	1	5	1	8	0,0078675760,079901528	8,42E-05	0,002977773
227062_at	AU155361	Hs.523789		TncRNA	EG2	3	3	4	5	0,0078675760,079901528	8,42E-05	0,002977773
227176_at	AL565362	Hs.249829		FLJ40126	EG2	5	1	6	3	0,0078675760,079901528	8,42E-05	0,002977773
227501_at	AI377135	Hs.446017	WD repeat and SOCS box-containing 1	WSB1	EG2	4	2	6	3	0,0078675760,079901528	8,42E-05	0,002977773
227755_at	AA042983				EG2	2	4	7	2	0,0078675760,079901528	8,42E-05	0,002977773
227793_at	AA969238	Hs.213065		LOC158257	EG2	5	1	5	4	0,0078675760,079901528	8,42E-05	0,002977773
227868_at	AI928764	Hs.511547, Hs.532236		LOC154761	EG2	0	6	3	6	0,0078675760,079901528	8,42E-05	0,002977773
228287_at	BG054893	Hs.471876,	H inhibitor of growth family, member 5	ING5	EG2	4	2	4	5	0,0078675760,079901528	8,42E-05	0,002977773
228393_s_at	BF508739	Hs.436350	zinc finger protein 302	ZNF302	EG2	3	3	5	4	0,0078675760,079901528	8,42E-05	0,002977773
229800_at	AI129626	Hs.507755	doublecortin and CaM kinase-like 1	DCAMKL1	EG2	3	3	4	5	0,0078675760,079901528	8,42E-05	0,002977773
230629_s_at	AI809582	Hs.507307	E1A binding protein p400	EP400	EG2	1	5	3	6	0,0078675760,079901528	8,42E-05	0,002977773
230733_at	H98113	Hs.190086		MRCL3	EG2	4	2	9	0	0,0078675760,079901528	8,42E-05	0,002977773
230793_at	BE671038	Hs.116470	leucine rich repeat containing 16	LRRC16	EG2	5	1	3	6	0,0078675760,079901528	8,42E-05	0,002977773
231035_s_at	AI697976	Hs.499042	OTU domain containing 1	OTUD1	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
231727_s_at	NM_020679	Hs.325631		AD023	EG2	3	3	7	2	0,0078675760,079901528	8,42E-05	0,002977773
232021_at	AI864273	Hs.259347		LOC283464	EG2	4	2	6	3	0,0078675760,079901528	8,42E-05	0,002977773
232535_at	AL133570				EG2	5	1	3	6	0,0078675760,079901528	8,42E-05	0,002977773
233401_at	BF723605	Hs.21422	neuronal cell adhesion molecule	NRCAM	EG2	4	2	4	5	0,0078675760,079901528	8,42E-05	0,002977773
235085_at	BF739767	Hs.29068		DKFZp761P0423	EG2	1	5	4	5	0,0078675760,079901528	8,42E-05	0,002977773
235289_at	AV747725	Hs.164144	eukaryotic translation initiation factor 5A2	EIF5A2	EG2	5	1	7	2	0,0078675760,079901528	8,42E-05	0,002977773
236075_s_at	AI693543	Hs.279914	zinc finger protein 232	ZNF232	EG2	3	3	3	6	0,0078675760,079901528	8,42E-05	0,002977773

236599_at	AA703280	Hs.38449	serpin peptidase inhibitor, clade E (nexin, plasminoge	SERPINE2	EG2	3	3	9	0	0,0078675760,079901528	8,42E-05	0,002977773
236834_at	AI025103	Hs.302287	sec1 family domain containing 2	SCFD2	EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
239944_at	AA431379				EG2	2	4	6	3	0,0078675760,079901528	8,42E-05	0,002977773
240282_at	AW770902	Hs.128548	WD repeat domain 1	WDR1	EG2	3	3	7	2	0,0078675760,079901528	8,42E-05	0,002977773
242029_at	N32832	Hs.159430	fibronectin type III domain containing 3B	FNDC3B	EG2	5	1	8	1	0,0078675760,079901528	8,42E-05	0,002977773
242463_x_at	AI620827	Hs.166312	zinc finger protein 600	ZNF600	EG2	5	1	8	1	0,0078675760,079901528	8,42E-05	0,002977773
243747_at	AI222019	Hs.368375	zinc finger protein 599	ZNF599	EG2	3	3	3	6	0,0078675760,079901528	8,42E-05	0,002977773
244433_at	AI950023				EG2	3	3	8	1	0,0078675760,079901528	8,42E-05	0,002977773
244623_at	BF513800				EG2	4	2	7	2	0,0078675760,079901528	8,42E-05	0,002977773
37152_at	L07592	Hs.485196	peroxisome proliferative activated receptor, delta	PPARD	EG2	0	6	0	9	0,0078675760,079901528	8,42E-05	0,002977773
AFFX-r2-Bs-lys	AFFX-R2-BS-LYS-M				EG2	3	3	4	5	0,0078675760,079901528	8,42E-05	0,002977773
222912_at	BE207758	Hs.503284	arrestin, beta 1	ARRB1	EG2	0	5	8	9	0,03340802 0,201243287	3,06E-15	0,001562433
212811_x_at	AI889380	Hs.323878	solute carrier family 1 (glutamate/neutral amino acid t	SLC1A4	EG2	4	1	9	7	0,03340802 0,201243287	1,94E-13	0,001562433
218145_at	NM_021158	Hs.516826	tribbles homolog 3 (Drosophila)	TRIB3	EG2	0	5	10	5	0,03340802 0,201243287	7,74E-12	0,001562433
203542_s_at	AI690205	Hs.150557	Kruppel-like factor 9	KLF9	EG2	3	2	3	11	0,03340802 0,201243287	2,17E-10	0,001562433
212675_s_at	AB011154	Hs.146007	KIAA0582	KIAA0582	EG2	1	4	3	11	0,03340802 0,201243287	2,17E-10	0,001562433
213524_s_at	NM_015714	Hs.432132	G0/G1switch 2	G0S2	EG2	2	3	9	4	0,03340802 0,201243287	4,56E-09	0,001562433
225646_at	AI246687	Hs.128065	cathepsin C	CTSC	EG2	2	3	8	5	0,03340802 0,201243287	4,56E-09	0,001562433
235408_x_at	AW058673	Hs.250693	zinc finger protein 117 (HPF9)	ZNF117	EG2	3	2	3	10	0,03340802 0,201243287	4,56E-09	0,001562433
1555950_a_at	CA448665	Hs.527653	decay accelerating factor for complement (CD55, Cro	DAF	EG2	3	2	6	6	0,03340802 0,201243287	7,43E-08	0,001562433
200907_s_at	AU157932	Hs.151220		KIAA0992	EG2	4	1	6	6	0,03340802 0,201243287	7,43E-08	0,001562433
202388_at	NM_002923	Hs.78944	regulator of G-protein signalling 2, 24kDa	RGS2	EG2	1	4	7	5	0,03340802 0,201243287	7,43E-08	0,001562433
204472_at	NM_005261	Hs.345139	GTP binding protein overexpressed in skeletal muscle	GEM	EG2	5	0	12	0	0,03340802 0,201243287	7,43E-08	0,001562433
204595_s_at	AI300520	Hs.25590	stanniocalcin 1	STC1	EG2	2	3	10	2	0,03340802 0,201243287	7,43E-08	0,001562433
205505_at	NM_001490	Hs.521568	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6	GCNT1	EG2	1	4	6	6	0,03340802 0,201243287	7,43E-08	0,001562433
213112_s_at	N30649	Hs.529892	sequestosome 1	SQSTM1	EG2	5	0	10	2	0,03340802 0,201243287	7,43E-08	0,001562433
213850_s_at	AI984932	Hs.210367	splicing factor, arginine/serine-rich 2, interacting prot	SFRS2IP	EG2	2	3	7	5	0,03340802 0,201243287	7,43E-08	0,001562433
226136_at	N32834				EG2	5	0	7	5	0,03340802 0,201243287	7,43E-08	0,001562433
227354_at	BF589359	Hs.266175,	H phosphoprotein associated with glycosphingolipid mi	PAG1	EG2	3	2	7	5	0,03340802 0,201243287	7,43E-08	0,001562433
229374_at	AI758962	Hs.371218	EPH receptor A4	EPHA4	EG2	4	1	3	9	0,03340802 0,201243287	7,43E-08	0,001562433
1558173_a_at	AK093016	Hs.257900,	H leucine zipper protein 1	LUZP1	EG2	1	4	6	5	0,03340802 0,201243287	9,59E-07	0,001562433
201212_at	D55696	Hs.18069	legumain	LGMN	EG2	4	1	4	7	0,03340802 0,201243287	9,59E-07	0,001562433
202206_at	AW450363	Hs.111554	ADP-ribosylation factor-like 7	ARL7	EG2	2	3	7	4	0,03340802 0,201243287	9,59E-07	0,001562433
203387_s_at	NM_014832	Hs.210891	TBC1 domain family, member 4	TBC1D4	EG2	2	3	4	7	0,03340802 0,201243287	9,59E-07	0,001562433
204014_at	NM_001394	Hs.417962	dual specificity phosphatase 4	DUSP4	EG2	2	3	3	8	0,03340802 0,201243287	9,59E-07	0,001562433
205027_s_at	NM_005204	Hs.432453	mitogen-activated protein kinase kinase kinase 8	MAP3K8	EG2	1	4	3	8	0,03340802 0,201243287	9,59E-07	0,001562433
206693_at	NM_000880	Hs.536926	interleukin 7	IL7	EG2	0	5	7	4	0,03340802 0,201243287	9,59E-07	0,001562433
208893_s_at	BC005047	Hs.298654	dual specificity phosphatase 6	DUSP6	EG2	3	2	9	2	0,03340802 0,201243287	9,59E-07	0,001562433
210715_s_at	AF027205	Hs.31439	serine peptidase inhibitor, Kunitz type, 2	SPINT2	EG2	3	2	3	8	0,03340802 0,201243287	9,59E-07	0,001562433
211668_s_at	K03226	Hs.77274	plasminogen activator, urokinase	PLAU	EG2	2	3	10	1	0,03340802 0,201243287	9,59E-07	0,001562433
212190_at	AL541302	Hs.38449	serpin peptidase inhibitor, clade E (nexin, plasminoge	SERPINE2	EG2	2	3	7	4	0,03340802 0,201243287	9,59E-07	0,001562433
212444_at	AA156240				EG2	3	2	4	7	0,03340802 0,201243287	9,59E-07	0,001562433
216598_s_at	S69738	Hs.303649	chemokine (C-C motif) ligand 2	CCL2	EG2	3	2	5	6	0,03340802 0,201243287	9,59E-07	0,001562433
218409_s_at	NM_022365	Hs.499000	DnaJ (Hsp40) homolog, subfamily C, member 1	DNAJC1	EG2	0	5	9	2	0,03340802 0,201243287	9,59E-07	0,001562433
219352_at	NM_017912	Hs.529317	hect domain and RLD 6	HERC6	EG2	0	5	1	10	0,03340802 0,201243287	9,59E-07	0,001562433

221757_at	BE042976	Hs.26670		MGC17330	EG2	1	4	4	7	0,03340802	0,201243287	9,59E-07	0,001562433
221865_at	BF969986	Hs.522357	chromosome 9 open reading frame 91	C9orf91	EG2	2	3	4	7	0,03340802	0,201243287	9,59E-07	0,001562433
223195_s_at	BF131886	Hs.469543	sestrin 2	SESN2	EG2	4	1	9	2	0,03340802	0,201243287	9,59E-07	0,001562433
223712_at	AL136721	Hs.527213	6-pyruvoyl-tetrahydropterin synthase/dimerization c	PCBD2	EG2	2	3	6	5	0,03340802	0,201243287	9,59E-07	0,001562433
226348_at	AK026764				EG2	0	5	9	2	0,03340802	0,201243287	9,59E-07	0,001562433
228423_at	AI887898	Hs.549331		LOC389237	EG2	5	0	4	7	0,03340802	0,201243287	9,59E-07	0,001562433
229071_at	AI681419	Hs.262480		LOC388327	EG2	2	3	3	8	0,03340802	0,201243287	9,59E-07	0,001562433
229307_at	N32051	Hs.335239	ankyrin repeat domain 28	ANKRD28	EG2	2	3	6	5	0,03340802	0,201243287	9,59E-07	0,001562433
230147_at	AI378647	Hs.42502	coagulation factor II (thrombin) receptor-like 2	F2RL2	EG2	1	4	8	3	0,03340802	0,201243287	9,59E-07	0,001562433
230508_at	AL569601	Hs.292156	dickkopf homolog 3 (<i>Xenopus laevis</i>)	DKK3	EG2	2	3	6	5	0,03340802	0,201243287	9,59E-07	0,001562433
235266_at	AI139629	Hs.370834	ATPase family, AAA domain containing 2	ATAD2	EG2	2	3	3	8	0,03340802	0,201243287	9,59E-07	0,001562433
235836_at	N51413	Hs.250723	H matrix-remodelling associated 7	MXRA7	EG2	2	3	9	2	0,03340802	0,201243287	9,59E-07	0,001562433
236545_at	AA532718	Hs.435512	protein phosphatase 3 (formerly 2B), catalytic subunit	PPP3CA	EG2	2	3	9	2	0,03340802	0,201243287	9,59E-07	0,001562433
236907_at	AI760366	Hs.387804	poly(A) binding protein, cytoplasmic 1	PABPC1	EG2	3	2	3	8	0,03340802	0,201243287	9,59E-07	0,001562433
239049_at	BF514509				EG2	1	4	5	6	0,03340802	0,201243287	9,59E-07	0,001562433
242121_at	AW973232	Hs.122121	ring finger protein 12	RNF12	EG2	2	3	5	6	0,03340802	0,201243287	9,59E-07	0,001562433
244341_at	AA827728	Hs.269528	Mak3 homolog (<i>S. cerevisiae</i>)	MAK3	EG2	2	3	7	4	0,03340802	0,201243287	9,59E-07	0,001562433
244753_at	BF000430	Hs.270291	actinin, alpha 4	ACTN4	EG2	3	2	10	1	0,03340802	0,201243287	9,59E-07	0,001562433
1568611_at	CA418310	Hs.519568	procollagen-proline, 2-oxoglutarate 4-dioxygenase (p	P4HA2	EG2	0	5	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
200906_s_at	AK025843	Hs.151220		KIAA0992	EG2	4	1	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
202205_at	NM_003370	Hs.515469	vasodilator-stimulated phosphoprotein	VASP	EG2	4	1	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
202350_s_at	NM_002380	Hs.189445	matrilin 2	MATN2	EG2	1	4	2	8	0,03340802	0,201243287	9,97E-06	0,001562433
202859_x_at	NM_000584	Hs.624	interleukin 8	IL8	EG2	3	2	9	1	0,03340802	0,201243287	9,97E-06	0,001562433
203397_s_at	BF063271	Hs.170986	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-	GALNT3	EG2	4	1	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
203823_at	NM_021106	Hs.494875	regulator of G-protein signalling 3	RGS3	EG2	0	5	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
204221_x_at	U16307	Hs.553516	GLI pathogenesis-related 1 (glioma)	GLIPR1	EG2	5	0	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
204326_x_at	NM_002450	Hs.374950	metallothionein 1X	MT1X	EG2	2	3	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
204407_at	AF080255	Hs.486818	transcription termination factor, RNA polymerase II	TTF2	EG2	3	2	1	9	0,03340802	0,201243287	9,97E-06	0,001562433
204597_x_at	NM_003155	Hs.25590	stanniocalcin 1	STC1	EG2	2	3	9	1	0,03340802	0,201243287	9,97E-06	0,001562433
204638_at	NM_001611	Hs.1211	acid phosphatase 5, tartrate resistant	ACP5	EG2	2	3	7	3	0,03340802	0,201243287	9,97E-06	0,001562433
205200_at	NM_003278	Hs.476092	C-type lectin domain family 3, member B	CLEC3B	EG2	0	5	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
205559_s_at	NM_006200	Hs.368542	proprotein convertase subtilisin/kexin type 5	PCSK5	EG2	3	2	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
206133_at	NM_017523	Hs.441975		BIRC4BP	EG2	2	3	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
206554_x_at	NM_006515	Hs.475300	SET domain and mariner transposase fusion gene	SETMAR	EG2	2	3	1	9	0,03340802	0,201243287	9,97E-06	0,001562433
206584_at	NM_015364	Hs.69328	lymphocyte antigen 96	LY96	EG2	2	3	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
206686_at	NM_002610	Hs.470633	pyruvate dehydrogenase kinase, isoenzyme 1	PDK1	EG2	0	5	7	3	0,03340802	0,201243287	9,97E-06	0,001562433
206756_at	NM_019886	Hs.138155	carbohydrate (N-acetylglucosamine 6-O) sulfotransfer	CHST7	EG2	0	5	9	1	0,03340802	0,201243287	9,97E-06	0,001562433
208527_x_at	NM_003523	Hs.534369	histone 1, H2be	HIST1H2BE	EG2	3	2	2	8	0,03340802	0,201243287	9,97E-06	0,001562433
208961_s_at	AB017493	Hs.4055	Kruppel-like factor 6	KLF6	EG2	1	4	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
209258_s_at	AI373676	Hs.24485	chondroitin sulfate proteoglycan 6 (bamacan)	CSPG6	EG2	1	4	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
209276_s_at	AF162769	Hs.28988	glutaredoxin (thioltransferase)	GLRX	EG2	1	4	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
209822_s_at	L22431	Hs.370422	very low density lipoprotein receptor	VLDLR	EG2	0	5	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
212621_at	AB006624	Hs.533787		KIAA0286	EG2	5	0	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
214169_at	BE615699	Hs.438072	unc-84 homolog A (<i>C. elegans</i>)	UNC84A	EG2	2	3	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
215092_s_at	AJ005683	Hs.371987	nuclear factor of activated T-cells 5, tonicity-responsiv	NFAT5	EG2	3	2	4	6	0,03340802	0,201243287	9,97E-06	0,001562433

218383_at	NM_017815	Hs.442782	chromosome 14 open reading frame 94	C14orf94	EG2	0	5	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
219315_s_at	NM_024600	Hs.459652	chromosome 16 open reading frame 30	C16orf30	EG2	0	5	0	10	0,03340802	0,201243287	9,97E-06	0,001562433
219806_s_at	NM_020179	Hs.438064		FN5	EG2	1	4	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
219938_s_at	NM_024430	Hs.368623	proline-serine-threonine phosphatase interacting pro	PSTPIP2	EG2	2	3	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
221833_at	AI971258	Hs.295923	seven in absentia homolog 1 (Drosophila)	SIAH1	EG2	0	5	2	8	0,03340802	0,201243287	9,97E-06	0,001562433
221986_s_at	AW006750	Hs.407709	kelch-like 24 (Drosophila)	KLHL24	EG2	5	0	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
222736_s_at	BC000049	Hs.411925	transmembrane protein 38B	TMEM38B	EG2	5	0	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
224496_s_at	BC006292	Hs.513933		MGC10744	EG2	3	2	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
224707_at	AL522667	Hs.483561		ORF1-FL49	EG2	5	0	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
225412_at	AA761169	Hs.469690		FLJ14681	EG2	3	2	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
225611_at	AI922968	Hs.133539	microtubule associated serine/threonine kinase famil	MAST4	EG2	3	2	0	10	0,03340802	0,201243287	9,97E-06	0,001562433
225855_at	AB046768	Hs.369232	erythrocyte membrane protein band 4.1 like 5	EPB41L5	EG2	4	1	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
226051_at	BF973568	Hs.55940		SELM	EG2	1	4	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
226264_at	AL137432	Hs.494827	sushi domain containing 1	SUSD1	EG2	0	5	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
227674_at	AA910852	Hs.390568	zinc finger protein 585A	ZNF585A	EG2	1	4	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
227722_at	AW043594	Hs.527193	ribosomal protein S23	RPS23	EG2	3	2	8	2	0,03340802	0,201243287	9,97E-06	0,001562433
227911_at	AI935647	Hs.183114	Rho GTPase activating protein 28	ARHGAP28	EG2	1	4	1	9	0,03340802	0,201243287	9,97E-06	0,001562433
228490_at	AW292816	Hs.549108	abhydrolase domain containing 2	ABHD2	EG2	3	2	10	0	0,03340802	0,201243287	9,97E-06	0,001562433
229963_at	AV726956	Hs.47209	NGFRAP1-like 1	NGFRAP1L1	EG2	4	1	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
230206_at	AI692645	Hs.195403	dedicator of cytokinesis 5	DOCK5	EG2	2	3	9	1	0,03340802	0,201243287	9,97E-06	0,001562433
230256_at	AW009436	Hs.131810	chromosome 1 open reading frame 104	C1orf104	EG2	0	5	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
230292_at	AA868809	Hs.25447	regulator of chromosome condensation (RCC1) and BT	RCBTB2	EG2	3	2	5	5	0,03340802	0,201243287	9,97E-06	0,001562433
231862_at	AK023520	Hs.349283	chromobox homolog 5 (HP1 alpha homolog, Drosophil	CBX5	EG2	4	1	4	6	0,03340802	0,201243287	9,97E-06	0,001562433
233303_at	AL110175	Hs.472031,	H ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog	UBE2D3	EG2	4	1	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
235171_at	AI354636				EG2	2	3	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
235609_at	BF056791	Hs.532799	BRCA1 interacting protein C-terminal helicase 1	BRIP1	EG2	4	1	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
236524_at	AA737437	Hs.276876	TM2 domain containing 1	TM2D1	EG2	2	3	6	4	0,03340802	0,201243287	9,97E-06	0,001562433
237563_s_at	AI286239	Hs.181156		LOC440731	EG2	2	3	3	7	0,03340802	0,201243287	9,97E-06	0,001562433
1554153_a_at	BC015714	Hs.502458	PHD finger protein 21A	PHF21A	EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
1555976_s_at	BU676221	Hs.233936		LOC440476	EG2	5	0	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
1556346_at	AJ227860	Hs.289092	coactosin-like 1 (Dictyostelium)	COTL1	EG2	2	3	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
1558290_a_at	BG200951	Hs.133107		LOC441378	EG2	0	5	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
1559232_a_at	BC029450	Hs.478031	solute carrier family 33 (acetyl-CoA transporter), mem	SLC33A1	EG2	4	1	9	0	0,03340802	0,201243287	8,42E-05	0,002977773
201171_at	NM_003945	Hs.484188	ATPase, H+ transporting, lysosomal 9kDa, V0 subunit e	ATP6V0E	EG2	1	4	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
201482_at	NM_002826	Hs.518374	quiescin Q6	QSCN6	EG2	1	4	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
201506_at	NM_000358	Hs.369397	transforming growth factor, beta-induced, 68kDa	TGFBI	EG2	2	3	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
201795_at	NM_002296	Hs.435166	lamin B receptor	LBR	EG2	3	2	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
202241_at	NM_025195	Hs.444947	tribbles homolog 1 (Drosophila)	TRIB1	EG2	1	4	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
202263_at	NM_016243	Hs.334832	cytochrome b5 reductase 1	CYB5R1	EG2	1	4	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
202419_at	NM_002035	Hs.74050	follicular lymphoma variant translocation 1	FVT1	EG2	4	1	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
202458_at	NM_007173	Hs.25338	protease, serine, 23	PRSS23	EG2	1	4	9	0	0,03340802	0,201243287	8,42E-05	0,002977773
202478_at	NM_021643	Hs.467751	tribbles homolog 2 (Drosophila)	TRIB2	EG2	1	4	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
203216_s_at	NM_004999	Hs.149387	myosin VI	MYO6	EG2	5	0	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
203231_s_at	AW235612	Hs.434961	ataxin 1	ATXN1	EG2	2	3	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
203414_at	NM_012329	Hs.463483	monocyte to macrophage differentiation-associated	MMD	EG2	4	1	3	6	0,03340802	0,201243287	8,42E-05	0,002977773

203865_s_at	NM_015833	Hs.474018	adenosine deaminase, RNA-specific, B1 (RED1 homolo	ADARB1	EG2	0	5	0	9	0,03340802	0,201243287	8,42E-05	0,002977773
203881_s_at	NM_004010	Hs.495912	dystrophin (muscular dystrophy, Duchenne and Becke	DMD	EG2	1	4	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
204160_s_at	AW194947	Hs.54037	ectonucleotide pyrophosphatase/phosphodiesterase	ENPP4	EG2	4	1	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
204396_s_at	NM_005308	Hs.524625	G protein-coupled receptor kinase 5	GRK5	EG2	1	4	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
204418_x_at	NM_000848	Hs.279837	glutathione S-transferase M2 (muscle)	GSTM2	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
204759_at	NM_001268	Hs.25447	regulator of chromosome condensation (RCC1) and BT	RCBTB2	EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
204774_at	NM_014210	Hs.553491	ecotropic viral integration site 2A	EVI2A	EG2	4	1	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
204971_at	NM_005213	Hs.518198	cystatin A (stefin A)	CSTA	EG2	2	3	9	0	0,03340802	0,201243287	8,42E-05	0,002977773
206307_s_at	NM_004472	Hs.519385	forkhead box D1	FOXD1	EG2	2	3	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
207992_s_at	NM_000480	Hs.501890	adenosine monophosphate deaminase (isoform E)	AMPD3	EG2	2	3	0	9	0,03340802	0,201243287	8,42E-05	0,002977773
208025_s_at	NM_003483	Hs.505924	high mobility group AT-hook 2	HMGA2	EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
208579_x_at	NM_017445	Hs.473961	H2B histone family, member S	H2BFS	EG2	3	2	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
209447_at	AF043290	Hs.12967	spectrin repeat containing, nuclear envelope 1	SYNE1	EG2	2	3	9	0	0,03340802	0,201243287	8,42E-05	0,002977773
211000_s_at	AB015706	Hs.454699,	interleukin 6 signal transducer (gp130, oncostatin M re	IL6ST	EG2	4	1	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
211571_s_at	D32039	Hs.443681	chondroitin sulfate proteoglycan 2 (versican)	CSPG2	EG2	4	1	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
212224_at	NM_000689	Hs.76392	aldehyde dehydrogenase 1 family, member A1	ALDH1A1	EG2	3	2	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
212442_s_at	BG289001	Hs.506829	LAG1 longevity assurance homolog 6 (S. cerevisiae)	LASS6	EG2	4	1	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
212828_at	AA191573	Hs.434494	synaptojanin 2	SYNJ2	EG2	3	2	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
212989_at	AI377497	Hs.386215	transmembrane protein 23	TMEM23	EG2	4	1	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
213029_at	BG478428	Hs.370359	nuclear factor I/B	NFIB	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
213286_at	BF445199	Hs.435231	zinc finger RNA binding protein	ZFR	EG2	4	1	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
213328_at	AI936517	Hs.481181	NIMA (never in mitosis gene a)-related kinase 1	NEK1	EG2	1	4	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
214305_s_at	AW003030	Hs.471011	splicing factor 3b, subunit 1, 155kDa	SF3B1	EG2	2	3	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
214375_at	AI962377	Hs.172445	PTPRF interacting protein, binding protein 1 (liprin be	PPFIBP1	EG2	1	4	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
216250_s_at	X77598	Hs.125474	leupaxin	LPXN	EG2	2	3	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
218128_at	AU151875	Hs.84928	nuclear transcription factor Y, beta	NFYB	EG2	1	4	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
218170_at	NM_016048	Hs.483296	isochorismatase domain containing 1	ISOC1	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
218522_s_at	NM_018174	Hs.66048	BPY2 interacting protein 1	BPY2IP1	EG2	0	5	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
218694_at	NM_016608	Hs.9728	armadillo repeat containing, X-linked 1	ARMCX1	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
218772_x_at	NM_018112	Hs.411925	transmembrane protein 38B	TMEM38B	EG2	5	0	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
219014_at	NM_016619	Hs.546392	placenta-specific 8	PLAC8	EG2	1	4	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
219104_at	NM_016422	Hs.44685	ring finger protein 141	RNF141	EG2	2	3	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
219131_at	NM_013319	Hs.522933	UbiA prenyltransferase domain containing 1	UBIAD1	EG2	2	3	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
219132_at	NM_021255	Hs.105103	pellino homolog 2 (Drosophila)	PELI2	EG2	4	1	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
219308_s_at	NM_012093	Hs.18268	adenylate kinase 5	AK5	EG2	4	1	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
221593_s_at	BC001663	Hs.469473,	ribosomal protein L31	RPL31	EG2	4	1	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
221756_at	AL540260	Hs.26670		MGC17330	EG2	0	5	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
222587_s_at	BF699855	Hs.127407	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-	GALNT7	EG2	2	3	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
223339_at	BC001867	Hs.241336	ATPase inhibitory factor 1	ATPIF1	EG2	4	1	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
223622_s_at	AF284751	Hs.473855	hydroxypyruvate isomerase homolog (E. coli)	HYI	EG2	1	4	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
223672_at	AL136561	Hs.132121	SH3-domain GRB2-like (endophilin) interacting protei	SGIP1	EG2	4	1	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
223769_x_at	AF284750	Hs.473855	hydroxypyruvate isomerase homolog (E. coli)	HYI	EG2	3	2	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
224944_at	AL566034	Hs.11355	thymopoietin	TMPO	EG2	4	1	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
225330_at	AL044092	Hs.20573	insulin-like growth factor 1 receptor	IGF1R	EG2	3	2	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
225592_at	D81048	Hs.519993	nurim (nuclear envelope membrane protein)	NRM	EG2	2	3	5	4	0,03340802	0,201243287	8,42E-05	0,002977773

225816_at	AW138134	Hs.12420	PHD finger protein 17	PHF17	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
226021_at	AW150720	Hs.244940	retinol dehydrogenase 10 (all-trans)	RDH10	EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
226037_s_at	AL049589	Hs.171723,	H TAF9-like RNA polymerase II, TATA box binding protei	TAF9L	EG2	5	0	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
226272_at	N25986				EG2	3	2	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
226384_at	BE858787	Hs.437179	phosphatidic acid phosphatase type 2 domain containi	PPAPDC1B	EG2	4	1	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
226550_at	AI672159				EG2	4	1	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
227114_at	BG435876	Hs.380110		DKFZp547C195	EG2	0	5	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
227607_at	AI638611	Hs.16229		AMSH-LP	EG2	2	3	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
228333_at	AI912571				EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
228955_at	AL041761				EG2	4	1	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
229359_at	BE552208				EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
229491_at	BF433180	Hs.546482		LOC133308	EG2	3	2	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
229538_s_at	AW271106	Hs.133294	IQ motif containing GTPase activating protein 3	IQGAP3	EG2	3	2	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
229692_at	AW135003	Hs.435771	apoptosis inhibitor 5	API5	EG2	4	1	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
229850_at	N30152				EG2	5	0	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
229865_at	AW058617	Hs.159430	fibronectin type III domain containing 3B	FNDC3B	EG2	4	1	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
230345_at	AI654547				EG2	1	4	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
231929_at	AI458439				EG2	3	2	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
232164_s_at	AL137725	Hs.200412	epiplakin 1	EPPK1	EG2	3	2	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
232889_at	AU147591			LOC389281	EG2	1	4	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
233543_s_at	AK021582	Hs.334772		FLJ13614	EG2	3	2	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
235046_at	AA456099				EG2	5	0	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
235276_at	AA781795	Hs.546467	epithelial stromal interaction 1 (breast)	EPSTI1	EG2	0	5	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
235309_at	AA126311				EG2	3	2	6	3	0,03340802	0,201243287	8,42E-05	0,002977773
235653_s_at	BF685315	Hs.479971	THAP domain containing 6	THAP6	EG2	0	5	4	5	0,03340802	0,201243287	8,42E-05	0,002977773
236449_at	AI885390	Hs.695	cystatin B (stefin B)	CSTB	EG2	2	3	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
237459_at	AA811257	Hs.506415	PCTAIRE protein kinase 2	PCTK2	EG2	1	4	9	0	0,03340802	0,201243287	8,42E-05	0,002977773
238320_at	AV659198	Hs.523789		TncRNA	EG2	2	3	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
238462_at	AI418293	Hs.444075		STS-1	EG2	4	1	3	6	0,03340802	0,201243287	8,42E-05	0,002977773
238673_at	AV724325				EG2	4	1	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
239147_at	AI243677	Hs.163914		DKFZp313G1735	EG2	1	4	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
239331_at	AW954199				EG2	3	2	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
241017_at	AA779927	Hs.469473,	H ribosomal protein L31	RPL31	EG2	3	2	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
241359_at	AA729232				EG2	1	4	2	7	0,03340802	0,201243287	8,42E-05	0,002977773
241838_at	AA699443				EG2	3	2	1	8	0,03340802	0,201243287	8,42E-05	0,002977773
242201_at	AI418253	Hs.397073	postmeiotic segregation increased 2-like 5	PMS2L5	EG2	1	4	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
242429_at	BF435183	Hs.412517	zinc finger protein 567	ZNF567	EG2	0	5	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
242750_at	AA678047	Hs.117747	methylmalonic aciduria (cobalamin deficiency) cblA ty	MMAA	EG2	4	1	7	2	0,03340802	0,201243287	8,42E-05	0,002977773
242856_at	AI291804				EG2	2	3	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
244025_at	BF590917				EG2	4	1	8	1	0,03340802	0,201243287	8,42E-05	0,002977773
91703_at	AA149545	Hs.502867	EH domain binding protein 1-like 1	EHBP1L1	EG2	1	4	5	4	0,03340802	0,201243287	8,42E-05	0,002977773
200742_s_at	BG231932	Hs.523454	tripeptidyl peptidase I	TPP1	EG2	0	4	11	3	0,1119520740,440114659		2,17E-10	0,001562433
221539_at	AB044548	Hs.411641	eukaryotic translation initiation factor 4E binding prot	EIF4EBP1	EG2	0	4	8	6	0,1119520740,440114659		2,17E-10	0,001562433
221911_at	BE881590	Hs.22634	ets variant gene 1	ETV1	EG2	2	2	9	5	0,1119520740,440114659		2,17E-10	0,001562433
1553613_s_at	NM_001453	Hs.348883	forkhead box C1	FOXC1	EG2	1	3	3	10	0,1119520740,440114659		4,56E-09	0,001562433

202679_at	NM_000271	Hs.464779	Niemann-Pick disease, type C1	NPC1	EG2	1	3	12	1	0,1119520740,440114659	4,56E-09	0,001562433
204070_at	NM_004585	Hs.17466	retinoic acid receptor responder (tazarotene induced)	RARRES3	EG2	2	2	4	9	0,1119520740,440114659	4,56E-09	0,001562433
209459_s_at	AF237813	Hs.336768	4-aminobutyrate aminotransferase	ABAT	EG2	2	2	6	7	0,1119520740,440114659	4,56E-09	0,001562433
213067_at	AI382123	Hs.16355	myosin, heavy polypeptide 10, non-muscle	MYH10	EG2	1	3	2	11	0,1119520740,440114659	4,56E-09	0,001562433
224851_at	AW274756	Hs.119882	cyclin-dependent kinase 6	CDK6	EG2	2	2	5	8	0,1119520740,440114659	4,56E-09	0,001562433
226757_at	AA131041	Hs.437609	interferon-induced protein with tetratricopeptide rep	IFIT2	EG2	2	2	8	5	0,1119520740,440114659	4,56E-09	0,001562433
242560_at	AA579890	Hs.208388	Fanconi anemia, complementation group D2	FANCD2	EG2	3	1	3	10	0,1119520740,440114659	4,56E-09	0,001562433
203386_at	AI650848	Hs.210891	TBC1 domain family, member 4	TBC1D4	EG2	2	2	5	7	0,1119520740,440114659	7,43E-08	0,001562433
207057_at	NM_004731	Hs.439643	solute carrier family 16 (monocarboxylic acid transpor	SLC16A7	EG2	2	2	3	9	0,1119520740,440114659	7,43E-08	0,001562433
209160_at	AB018580	Hs.78183	aldo-keto reductase family 1, member C3 (3-alpha hyd	AKR1C3	EG2	1	3	6	6	0,1119520740,440114659	7,43E-08	0,001562433
217998_at	NM_007350	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	2	2	10	2	0,1119520740,440114659	7,43E-08	0,001562433
217999_s_at	NM_007350	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	3	1	10	2	0,1119520740,440114659	7,43E-08	0,001562433
1554016_a_at	BC006291	Hs.408702		FLJ13154	EG2	0	4	7	4	0,1119520740,440114659	9,59E-07	0,001562433
200879_s_at	NM_001430	Hs.468410	endothelial PAS domain protein 1	EPAS1	EG2	4	0	8	3	0,1119520740,440114659	9,59E-07	0,001562433
201107_s_at	AI812030	Hs.164226	thrombospondin 1	THBS1	EG2	2	2	5	6	0,1119520740,440114659	9,59E-07	0,001562433
201580_s_at	AL544094	Hs.169358	thioredoxin domain containing 13	TXNDC13	EG2	0	4	7	4	0,1119520740,440114659	9,59E-07	0,001562433
201858_s_at	J03223	Hs.1908	proteoglycan 1, secretory granule	PRG1	EG2	3	1	6	5	0,1119520740,440114659	9,59E-07	0,001562433
202669_s_at	U16797	Hs.149239	ephrin-B2	EFNB2	EG2	2	2	0	11	0,1119520740,440114659	9,59E-07	0,001562433
202838_at	NM_000147	Hs.370858	fucosidase, alpha-L- 1, tissue	FUCA1	EG2	1	3	9	2	0,1119520740,440114659	9,59E-07	0,001562433
204224_s_at	NM_000161	Hs.86724	GTP cyclohydrolase 1 (dopa-responsive dystonia)	GCH1	EG2	2	2	7	4	0,1119520740,440114659	9,59E-07	0,001562433
204338_s_at	NM_005613	Hs.386726	regulator of G-protein signalling 4	RGS4	EG2	3	1	9	2	0,1119520740,440114659	9,59E-07	0,001562433
205047_s_at	NM_001673	Hs.489207	asparagine synthetase	ASNS	EG2	4	0	5	6	0,1119520740,440114659	9,59E-07	0,001562433
206237_s_at	NM_013957	Hs.453951	neuregulin 1	NRG1	EG2	2	2	9	2	0,1119520740,440114659	9,59E-07	0,001562433
212979_s_at	AW293343	Hs.406492		KIAA0738	EG2	2	2	4	7	0,1119520740,440114659	9,59E-07	0,001562433
215813_s_at	S36219	Hs.201978	prostaglandin-endoperoxide synthase 1 (prostaglandi	PTGS1	EG2	1	3	6	5	0,1119520740,440114659	9,59E-07	0,001562433
218590_at	NM_021830	Hs.22678	progressive external ophthalmoplegia 1	PEO1	EG2	1	3	4	7	0,1119520740,440114659	9,59E-07	0,001562433
219326_s_at	NM_006577	Hs.173203	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltr	B3GNT1	EG2	4	0	6	5	0,1119520740,440114659	9,59E-07	0,001562433
219675_s_at	NM_025076	Hs.469561	UDP-glucuronate decarboxylase 1	UXS1	EG2	2	2	4	7	0,1119520740,440114659	9,59E-07	0,001562433
221864_at	AW517464	Hs.460617		MGC13024	EG2	0	4	6	5	0,1119520740,440114659	9,59E-07	0,001562433
222572_at	BG542521	Hs.22265	protein phosphatase 2C, magnesium-dependent, cata	PPM2C	EG2	4	0	4	7	0,1119520740,440114659	9,59E-07	0,001562433
223217_s_at	BE646573	Hs.319171	nuclear factor of kappa light polypeptide gene enhanc	NFKBIZ	EG2	0	4	4	7	0,1119520740,440114659	9,59E-07	0,001562433
224480_s_at	BC006236	Hs.99196		MGC11324	EG2	3	1	5	6	0,1119520740,440114659	9,59E-07	0,001562433
225798_at	AL047908	Hs.368944		JAZF1	EG2	1	3	6	5	0,1119520740,440114659	9,59E-07	0,001562433
225929_s_at	AA233374	Hs.195642	chromosome 17 open reading frame 27	C17orf27	EG2	0	4	4	7	0,1119520740,440114659	9,59E-07	0,001562433
226460_at	AB040883	Hs.556658		KIAA1450	EG2	2	2	6	5	0,1119520740,440114659	9,59E-07	0,001562433
228244_at	BF062383	Hs.103902	biogenesis of lysosome-related organelles complex-1	BLOC1S3	EG2	0	4	7	4	0,1119520740,440114659	9,59E-07	0,001562433
228416_at	AI149508	Hs.470174	origin recognition complex, subunit 4-like (yeast)	ORC4L	EG2	2	2	7	4	0,1119520740,440114659	9,59E-07	0,001562433
228461_at	AL566989	Hs.535157		LOC442040	EG2	3	1	0	11	0,1119520740,440114659	9,59E-07	0,001562433
228574_at	AI862551	Hs.444240		DKFZp762A217	EG2	1	3	0	11	0,1119520740,440114659	9,59E-07	0,001562433
230109_at	AI638433	Hs.126980	phosphodiesterase 7B	PDE7B	EG2	2	2	2	9	0,1119520740,440114659	9,59E-07	0,001562433
232031_s_at	AK023817	Hs.514843	KIAA1632	KIAA1632	EG2	3	1	11	0	0,1119520740,440114659	9,59E-07	0,001562433
235405_at	N79662	Hs.485557	glutathione S-transferase A4	GSTA4	EG2	3	1	5	6	0,1119520740,440114659	9,59E-07	0,001562433
235542_at	BF675754	Hs.516107		MGC22014	EG2	2	2	3	8	0,1119520740,440114659	9,59E-07	0,001562433
236356_at	AI808395	Hs.471207	NADH dehydrogenase (ubiquinone) Fe-S protein 1, 75	NDUFS1	EG2	2	2	6	5	0,1119520740,440114659	9,59E-07	0,001562433
238940_at	BE328128	Hs.373857	Kruppel-like factor 12	KLF12	EG2	4	0	4	7	0,1119520740,440114659	9,59E-07	0,001562433

240383_at	AI239832	Hs.472031,	Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog)	UBE2D3	EG2	2	2	6	5	0,1119520740,440114659	9,59E-07	0,001562433
242767_at	N95437	Hs.475353	LIM and cysteine-rich domains 1	LMCD1	EG2	2	2	5	6	0,1119520740,440114659	9,59E-07	0,001562433
244546_at	AI760495	Hs.437060	cytochrome c, somatic	CYCS	EG2	1	3	7	4	0,1119520740,440114659	9,59E-07	0,001562433
1557852_at	AW418842	Hs.524271	polyhomeotic-like 2 (Drosophila)	PHC2	EG2	1	3	8	2	0,1119520740,440114659	9,97E-06	0,001562433
1559332_at	BC016339	Hs.332847	cysteine rich transmembrane BMP regulator 1 (chordi	CRIM1	EG2	1	3	8	2	0,1119520740,440114659	9,97E-06	0,001562433
1565651_at	BI868311				EG2	0	4	9	1	0,1119520740,440114659	9,97E-06	0,001562433
203505_at	AF285167	Hs.429294	ATP-binding cassette, sub-family A (ABC1), member 1	ABCA1	EG2	4	0	3	7	0,1119520740,440114659	9,97E-06	0,001562433
203725_at	NM_001924	Hs.80409	growth arrest and DNA-damage-inducible, alpha	GADD45A	EG2	4	0	5	5	0,1119520740,440114659	9,97E-06	0,001562433
203817_at	W93728	Hs.77890	guanylate cyclase 1, soluble, beta 3	GUCY1B3	EG2	2	2	3	7	0,1119520740,440114659	9,97E-06	0,001562433
204352_at	NM_004619	Hs.523930	TNF receptor-associated factor 5	TRAF5	EG2	1	3	2	8	0,1119520740,440114659	9,97E-06	0,001562433
204646_at	NM_000110	Hs.335034	dihydropyrimidine dehydrogenase	DPYD	EG2	2	2	2	8	0,1119520740,440114659	9,97E-06	0,001562433
205885_s_at	L12002	Hs.555880	integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VL	ITGA4	EG2	2	2	3	7	0,1119520740,440114659	9,97E-06	0,001562433
205891_at	NM_000676	Hs.167046	adenosine A2b receptor	ADORA2B	EG2	3	1	6	4	0,1119520740,440114659	9,97E-06	0,001562433
205966_at	NM_005645	Hs.502508	TAF13 RNA polymerase II, TATA box binding protein (T	TAF13	EG2	4	0	3	7	0,1119520740,440114659	9,97E-06	0,001562433
207574_s_at	NM_015675	Hs.110571	growth arrest and DNA-damage-inducible, beta	GADD45B	EG2	2	2	5	5	0,1119520740,440114659	9,97E-06	0,001562433
209305_s_at	AF078077	Hs.110571	growth arrest and DNA-damage-inducible, beta	GADD45B	EG2	2	2	7	3	0,1119520740,440114659	9,97E-06	0,001562433
209561_at	L38969	Hs.169875	thrombospondin 3	THBS3	EG2	1	3	4	6	0,1119520740,440114659	9,97E-06	0,001562433
210298_x_at	AF098518	Hs.435369	four and a half LIM domains 1	FHL1	EG2	1	3	1	9	0,1119520740,440114659	9,97E-06	0,001562433
211317_s_at	AF041461	Hs.390736	CASP8 and FADD-like apoptosis regulator	CFLAR	EG2	0	4	5	5	0,1119520740,440114659	9,97E-06	0,001562433
212092_at	BE858180	Hs.147492	paternally expressed 10	PEG10	EG2	3	1	4	6	0,1119520740,440114659	9,97E-06	0,001562433
213416_at	BG532690	Hs.440955	ceramide kinase-like	CERKL	EG2	1	3	0	10	0,1119520740,440114659	9,97E-06	0,001562433
213836_s_at	AW052084	Hs.463964		WIPI49	EG2	2	2	7	3	0,1119520740,440114659	9,97E-06	0,001562433
218543_s_at	NM_022750	Hs.12646	poly (ADP-ribose) polymerase family, member 12	PARP12	EG2	2	2	2	8	0,1119520740,440114659	9,97E-06	0,001562433
218613_at	NM_018422	Hs.434255	pleckstrin and Sec7 domain containing 3	PSD3	EG2	2	2	4	6	0,1119520740,440114659	9,97E-06	0,001562433
219073_s_at	NM_017784	Hs.150122	oxysterol binding protein-like 10	OSBPL10	EG2	1	3	4	6	0,1119520740,440114659	9,97E-06	0,001562433
219279_at	NM_017718	Hs.46578	dedicator of cytokinesis 10	DOCK10	EG2	4	0	3	7	0,1119520740,440114659	9,97E-06	0,001562433
220260_at	NM_018317	Hs.479403	TBC1 domain family, member 19	TBC1D19	EG2	3	1	6	4	0,1119520740,440114659	9,97E-06	0,001562433
220892_s_at	NM_021154	Hs.494261	phosphoserine aminotransferase 1	PSAT1	EG2	4	0	5	5	0,1119520740,440114659	9,97E-06	0,001562433
224847_at	AW051349	Hs.119882	cyclin-dependent kinase 6	CDK6	EG2	3	1	5	5	0,1119520740,440114659	9,97E-06	0,001562433
225373_at	BE271644	Hs.47382, Hs.546421		C10orf54	EG2	0	4	4	6	0,1119520740,440114659	9,97E-06	0,001562433
225529_at	AI492175	Hs.535257	centaurin, beta 5	CENTB5	EG2	1	3	6	4	0,1119520740,440114659	9,97E-06	0,001562433
225931_s_at	AI954660	Hs.195642	chromosome 17 open reading frame 27	C17orf27	EG2	0	4	5	5	0,1119520740,440114659	9,97E-06	0,001562433
226279_at	AW471145	Hs.25338	protease, serine, 23	PRSS23	EG2	3	1	10	0	0,1119520740,440114659	9,97E-06	0,001562433
226419_s_at	AA046439	Hs.68714	splicing factor, arginine/serine-rich 1 (splicing factor 2	SFRS1	EG2	4	0	6	4	0,1119520740,440114659	9,97E-06	0,001562433
226558_at	BE856637				EG2	2	2	5	5	0,1119520740,440114659	9,97E-06	0,001562433
227059_at	AI651255	Hs.444329	glypican 6	GPC6	EG2	2	2	5	5	0,1119520740,440114659	9,97E-06	0,001562433
227566_at	AW085558	Hs.504352		HNT	EG2	2	2	8	2	0,1119520740,440114659	9,97E-06	0,001562433
228617_at	AA142842	Hs.441975		BIRC4BP	EG2	1	3	5	5	0,1119520740,440114659	9,97E-06	0,001562433
228762_at	AW151924	Hs.159142	lunatic fringe homolog (Drosophila)	LFNG	EG2	1	3	6	4	0,1119520740,440114659	9,97E-06	0,001562433
229371_at	BF940010	Hs.491611	chromosome 8 open reading frame 40	C8orf40	EG2	4	0	5	5	0,1119520740,440114659	9,97E-06	0,001562433
230703_at	AA001543	Hs.437831	chromosome 14 open reading frame 32	C14orf32	EG2	3	1	5	5	0,1119520740,440114659	9,97E-06	0,001562433
230886_at	AW069499				EG2	3	1	5	5	0,1119520740,440114659	9,97E-06	0,001562433
231057_at	AU144266	Hs.181326	myotubularin related protein 2	MTMR2	EG2	3	1	7	3	0,1119520740,440114659	9,97E-06	0,001562433
231890_at	AK022804				EG2	3	1	7	3	0,1119520740,440114659	9,97E-06	0,001562433
232412_at	AK024690	Hs.462946	F-box and leucine-rich repeat protein 20	FBXL20	EG2	2	2	4	6	0,1119520740,440114659	9,97E-06	0,001562433

235919_at	AA227879			EG2	4	0	7	3	0,1119520740,440114659	9,97E-06	0,001562433
236229_at	AW014345	Hs.467370		LOC400721	EG2	2	2	5	0,1119520740,440114659	9,97E-06	0,001562433
236492_at	AI934447	Hs.146339	protein phosphatase 2 (formerly 2A), regulatory subunit	PPP2R2A	EG2	4	0	8	0,1119520740,440114659	9,97E-06	0,001562433
238449_at	BG534511	Hs.528462, Hs.531664, Hs.547315, Hs.556385		DKFZp547E087	EG2	4	0	5	0,1119520740,440114659	9,97E-06	0,001562433
238824_at	BF844863				EG2	2	2	5	0,1119520740,440114659	9,97E-06	0,001562433
242191_at	AI701905	Hs.445080, Hs.454852		AE01	EG2	3	1	6	0,1119520740,440114659	9,97E-06	0,001562433
1552643_at	NM_145297	Hs.50158	zinc finger protein 626	ZNF626	EG2	2	2	5	0,1119520740,440114659	8,42E-05	0,002977773
1557685_at	AK054840	Hs.106015	development and differentiation enhancing factor 1	DDEF1	EG2	2	2	3	0,1119520740,440114659	8,42E-05	0,002977773
1558143_a_at	AK027160	Hs.469658	BCL2-like 11 (apoptosis facilitator)	BCL2L11	EG2	4	0	7	0,1119520740,440114659	8,42E-05	0,002977773
1566482_at	AL833114	Hs.368986		LOC400047	EG2	2	2	5	0,1119520740,440114659	8,42E-05	0,002977773
1566557_at	AK096609	Hs.448889		FLJ90757	EG2	0	4	0	0,1119520740,440114659	8,42E-05	0,002977773
1569022_a_at	BC040952	Hs.175343	phosphoinositide-3-kinase, class 2, alpha polypeptide	PIK3C2A	EG2	3	1	6	0,1119520740,440114659	8,42E-05	0,002977773
200965_s_at	NM_006720	Hs.438236	actin binding LIM protein 1	ABLIM1	EG2	2	2	2	0,1119520740,440114659	8,42E-05	0,002977773
202345_s_at	NM_001444	Hs.408061	fatty acid binding protein 5 (psoriasis-associated)	FABP5	EG2	4	0	4	0,1119520740,440114659	8,42E-05	0,002977773
202742_s_at	NM_002731	Hs.487325	protein kinase, cAMP-dependent, catalytic, beta	PRKACB	EG2	4	0	5	0,1119520740,440114659	8,42E-05	0,002977773
202761_s_at	NM_015180	Hs.525392	spectrin repeat containing, nuclear envelope 2	SYNE2	EG2	4	0	3	0,1119520740,440114659	8,42E-05	0,002977773
202897_at	AB023430	Hs.128846	protein tyrosine phosphatase, non-receptor type subunit	PTPNS1	EG2	1	3	5	0,1119520740,440114659	8,42E-05	0,002977773
203128_at	NM_004863	Hs.435661	serine palmitoyltransferase, long chain base subunit 2	SPTLC2	EG2	0	4	7	0,1119520740,440114659	8,42E-05	0,002977773
203492_x_at	AA918224	Hs.101014		PIG8	EG2	3	1	4	0,1119520740,440114659	8,42E-05	0,002977773
203493_s_at	AL525206	Hs.101014		PIG8	EG2	3	1	4	0,1119520740,440114659	8,42E-05	0,002977773
203753_at	NM_003199	Hs.200285	transcription factor 4	TCF4	EG2	0	4	4	0,1119520740,440114659	8,42E-05	0,002977773
203821_at	NM_001945	Hs.799	heparin-binding EGF-like growth factor	HBEGF	EG2	4	0	9	0,1119520740,440114659	8,42E-05	0,002977773
203921_at	NM_004267	Hs.8786	carbohydrate (N-acetylglucosamine-6-O) sulfotransferase	CHST2	EG2	2	2	7	0,1119520740,440114659	8,42E-05	0,002977773
204088_at	NM_002560	Hs.321709	purinergic receptor P2X, ligand-gated ion channel, 4	P2RX4	EG2	1	3	3	0,1119520740,440114659	8,42E-05	0,002977773
204363_at	NM_001993	Hs.62192	coagulation factor III (thromboplastin, tissue factor)	F3	EG2	1	3	2	0,1119520740,440114659	8,42E-05	0,002977773
204663_at	NM_006680	Hs.199743	malic enzyme 3, NADP(+)-dependent, mitochondrial	ME3	EG2	0	4	8	0,1119520740,440114659	8,42E-05	0,002977773
204720_s_at	AV729634	Hs.536063	DnaJ (Hsp40) homolog, subfamily C, member 6	DNAJC6	EG2	3	1	6	0,1119520740,440114659	8,42E-05	0,002977773
204862_s_at	NM_002513	Hs.514065	non-metastatic cells 3, protein expressed in	NME3	EG2	1	3	5	0,1119520740,440114659	8,42E-05	0,002977773
205379_at	NM_001236	Hs.154510	carbonyl reductase 3	CBR3	EG2	2	2	1	0,1119520740,440114659	8,42E-05	0,002977773
205648_at	NM_003391	Hs.128553	wingless-type MMTV integration site family member 2	WNT2	EG2	0	4	1	0,1119520740,440114659	8,42E-05	0,002977773
206025_s_at	AW188198	Hs.437322	tumor necrosis factor, alpha-induced protein 6	TNFAIP6	EG2	2	2	8	0,1119520740,440114659	8,42E-05	0,002977773
207304_at	NM_003425	Hs.381285	zinc finger protein 45	ZNF45	EG2	0	4	7	0,1119520740,440114659	8,42E-05	0,002977773
209015_s_at	BC002446	Hs.490745	DnaJ (Hsp40) homolog, subfamily B, member 6	DNAJB6	EG2	4	0	3	0,1119520740,440114659	8,42E-05	0,002977773
209292_at	AL022726	Hs.519601	inhibitor of DNA binding 4, dominant negative helix-loop	ID4	EG2	3	1	8	0,1119520740,440114659	8,42E-05	0,002977773
209829_at	AB002384	Hs.484915	chromosome 6 open reading frame 32	C6orf32	EG2	3	1	0	0,1119520740,440114659	8,42E-05	0,002977773
211981_at	NM_001845	Hs.17441	collagen, type IV, alpha 1	COL4A1	EG2	2	2	4	0,1119520740,440114659	8,42E-05	0,002977773
212070_at	AL554008	Hs.513633	G protein-coupled receptor 56	GPR56	EG2	0	4	9	0,1119520740,440114659	8,42E-05	0,002977773
212136_at	AW517686	Hs.343522	ATPase, Ca++ transporting, plasma membrane 4	ATP2B4	EG2	2	2	5	0,1119520740,440114659	8,42E-05	0,002977773
212607_at	N32526	Hs.498292	v-akt murine thymoma viral oncogene homolog 3 (pro	AKT3	EG2	4	0	3	0,1119520740,440114659	8,42E-05	0,002977773
213069_at	AI148659	Hs.477420	HEG homolog 1 (zebrafish)	HEG1	EG2	1	3	6	0,1119520740,440114659	8,42E-05	0,002977773
213435_at	AB028957	Hs.516617	SATB family member 2	SATB2	EG2	3	1	7	0,1119520740,440114659	8,42E-05	0,002977773
213449_at	D31765	Hs.252828	processing of precursor 1, ribonuclease P/MRP subunit	POP1	EG2	2	2	6	0,1119520740,440114659	8,42E-05	0,002977773
213652_at	AU152579	Hs.368542	proprotein convertase subtilisin/kexin type 5	PCSK5	EG2	3	1	3	0,1119520740,440114659	8,42E-05	0,002977773
216035_x_at	AV721430	Hs.501080	transcription factor 7-like 2 (T-cell specific, HMG-box)	TCF7L2	EG2	2	2	6	0,1119520740,440114659	8,42E-05	0,002977773
217452_s_at	Y15014	Hs.518834	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide	B3GALT2	EG2	3	1	3	0,1119520740,440114659	8,42E-05	0,002977773

218537_at	NM_017885	Hs.412304,	H host cell factor C1 regulator 1 (XPO1 dependant)	HCFC1R1	EG2	0	4	5	4	0,1119520740,440114659	8,42E-05	0,002977773
219737_s_at	AI524125	Hs.407643	protocadherin 9	PCDH9	EG2	2	2	7	2	0,1119520740,440114659	8,42E-05	0,002977773
220651_s_at	NM_018518	Hs.198363	MCM10 minichromosome maintenance deficient 10 (S	MCM10	EG2	3	1	4	5	0,1119520740,440114659	8,42E-05	0,002977773
221127_s_at	NM_006394	Hs.292156		RIG	EG2	0	4	3	6	0,1119520740,440114659	8,42E-05	0,002977773
221423_s_at	NM_030799	Hs.372050	Yip1 domain family, member 5	YIPF5	EG2	3	1	5	4	0,1119520740,440114659	8,42E-05	0,002977773
222044_at	AI199589	Hs.472856	chromosome 20 open reading frame 67	C20orf67	EG2	2	2	6	3	0,1119520740,440114659	8,42E-05	0,002977773
222557_at	AL353715	Hs.348326	stathmin-like 3	STMN3	EG2	0	4	3	6	0,1119520740,440114659	8,42E-05	0,002977773
222853_at	N71923	Hs.41296	fibronectin leucine rich transmembrane protein 3	FLRT3	EG2	0	4	4	5	0,1119520740,440114659	8,42E-05	0,002977773
223315_at	AF278532	Hs.201034	netrin 4	NTN4	EG2	2	2	1	8	0,1119520740,440114659	8,42E-05	0,002977773
223614_at	AL136588	Hs.492187		DKFZp761D112	EG2	2	2	7	2	0,1119520740,440114659	8,42E-05	0,002977773
223805_at	AF323728	Hs.318775	oxysterol binding protein-like 6	OSBPL6	EG2	3	1	6	3	0,1119520740,440114659	8,42E-05	0,002977773
225724_at	AW136120	Hs.531089	proteasome (prosome, macropain) subunit, alpha typ	PSMA3	EG2	3	1	5	4	0,1119520740,440114659	8,42E-05	0,002977773
225879_at	BF345244	Hs.378501	tRNA splicing endonuclease 54 homolog (SEN54, S. cer	TSEN54	EG2	1	3	6	3	0,1119520740,440114659	8,42E-05	0,002977773
225957_at	AI307750	Hs.484195	adult retina protein	LOC153222	EG2	2	2	5	4	0,1119520740,440114659	8,42E-05	0,002977773
226150_at	BF111651	Hs.437179	phosphatidic acid phosphatase type 2 domain containi	PPAPDC1B	EG2	3	1	8	1	0,1119520740,440114659	8,42E-05	0,002977773
226157_at	AI569747	Hs.379018	transcription factor Dp-2 (E2F dimerization partner 2)	TFDP2	EG2	2	2	6	3	0,1119520740,440114659	8,42E-05	0,002977773
226210_s_at	AI291123	Hs.525589,	H maternally expressed 3	MEG3	EG2	0	4	3	6	0,1119520740,440114659	8,42E-05	0,002977773
226670_s_at	AL109839	Hs.411022	chromosome 20 open reading frame 119	C20orf119	EG2	2	2	6	3	0,1119520740,440114659	8,42E-05	0,002977773
226869_at	AI655611				EG2	1	3	1	8	0,1119520740,440114659	8,42E-05	0,002977773
227051_at	AU157716				EG2	2	2	8	1	0,1119520740,440114659	8,42E-05	0,002977773
227088_at	BF221547	Hs.370661	phosphodiesterase 5A, cGMP-specific	PDE5A	EG2	3	1	2	7	0,1119520740,440114659	8,42E-05	0,002977773
227249_at	AI857685	Hs.460107	nudE nuclear distribution gene E homolog 1 (A. nidula	NDE1	EG2	2	2	4	5	0,1119520740,440114659	8,42E-05	0,002977773
227605_at	AI762870	Hs.480465	small inducible cytokine subfamily E, member 1 (endo	SCYE1	EG2	3	1	3	6	0,1119520740,440114659	8,42E-05	0,002977773
227649_s_at	AU144000	Hs.497575	SLIT-ROBO Rho GTPase activating protein 2	SRGAP2	EG2	2	2	6	3	0,1119520740,440114659	8,42E-05	0,002977773
228062_at	AW025330	Hs.12554	nucleosome assembly protein 1-like 5	NAP1L5	EG2	3	1	7	2	0,1119520740,440114659	8,42E-05	0,002977773
228885_at	AI862120	Hs.127386	MAM domain containing 2	MAMDC2	EG2	1	3	6	3	0,1119520740,440114659	8,42E-05	0,002977773
228927_at	AW291411	Hs.464896	zinc finger protein 397	ZNF397	EG2	4	0	3	6	0,1119520740,440114659	8,42E-05	0,002977773
229549_at	AA868461	Hs.550465	opsin 1 (cone pigments), short-wave-sensitive (color	OPN1SW	EG2	4	0	7	2	0,1119520740,440114659	8,42E-05	0,002977773
229693_at	AI952836	Hs.462230		LOC388335	EG2	1	3	2	7	0,1119520740,440114659	8,42E-05	0,002977773
229886_at	AA502768	Hs.448041		FLJ32363	EG2	3	1	3	6	0,1119520740,440114659	8,42E-05	0,002977773
230218_at	BF476403				EG2	2	2	2	7	0,1119520740,440114659	8,42E-05	0,002977773
231234_at	AV699565	Hs.128065	cathepsin C	CTSC	EG2	2	2	5	4	0,1119520740,440114659	8,42E-05	0,002977773
231513_at	BF111326	Hs.1547	potassium inwardly-rectifying channel, subfamily J, m	KCNJ2	EG2	3	1	4	5	0,1119520740,440114659	8,42E-05	0,002977773
231779_at	AI246590	Hs.449207	interleukin-1 receptor-associated kinase 2	IRAK2	EG2	3	1	8	1	0,1119520740,440114659	8,42E-05	0,002977773
231807_at	AL157473	Hs.445885	KIAA1217	KIAA1217	EG2	1	3	1	8	0,1119520740,440114659	8,42E-05	0,002977773
231956_at	AA976354	Hs.514554	KIAA1618	KIAA1618	EG2	1	3	4	5	0,1119520740,440114659	8,42E-05	0,002977773
232038_at	AI052103	Hs.121396	chromosome 6 open reading frame 170	C6orf170	EG2	3	1	4	5	0,1119520740,440114659	8,42E-05	0,002977773
232238_at	AK001380	Hs.121028	asp (abnormal spindle)-like, microcephaly associated	ASPM	EG2	2	2	2	7	0,1119520740,440114659	8,42E-05	0,002977773
232589_at	AK021551				EG2	2	2	3	6	0,1119520740,440114659	8,42E-05	0,002977773
232865_at	N59653	Hs.519313	AF4/FMR2 family, member 4	AFF4	EG2	2	2	7	2	0,1119520740,440114659	8,42E-05	0,002977773
233223_at	AK000850	Hs.37982	neural precursor cell expressed, developmentally do	NEDD9	EG2	1	3	8	1	0,1119520740,440114659	8,42E-05	0,002977773
236009_at	AI767250				EG2	3	1	7	2	0,1119520740,440114659	8,42E-05	0,002977773
236016_at	AI702962	Hs.463010	SWI/SNF related, matrix associated, actin dependent r	SMARCE1	EG2	0	4	5	4	0,1119520740,440114659	8,42E-05	0,002977773
238469_at	BE620374	Hs.368337	chromosome 6 open reading frame 155	C6orf155	EG2	4	0	7	2	0,1119520740,440114659	8,42E-05	0,002977773
238860_at	AA398043	Hs.227457	chromosome 6 open reading frame 130	C6orf130	EG2	1	3	2	7	0,1119520740,440114659	8,42E-05	0,002977773

239043_at	AA084273	Hs.76561	zinc finger protein 404	ZNF404	EG2	1	3	6	3	0,1119520740,440114659	8,42E-05	0,002977773
239231_at	BE464819				EG2	3	1	3	6	0,1119520740,440114659	8,42E-05	0,002977773
239587_at	AI686890				EG2	2	2	7	2	0,1119520740,440114659	8,42E-05	0,002977773
240145_at	AW628059				EG2	1	3	3	6	0,1119520740,440114659	8,42E-05	0,002977773
241484_x_at	AI668696				EG2	2	2	7	2	0,1119520740,440114659	8,42E-05	0,002977773
241820_at	BF666241	Hs.536537	RAP1 interacting factor homolog (yeast)	RIF1	EG2	2	2	5	4	0,1119520740,440114659	8,42E-05	0,002977773
242052_at	AW979272	Hs.505202	bicaudal D homolog 1 (Drosophila)	BICD1	EG2	0	4	6	3	0,1119520740,440114659	8,42E-05	0,002977773
37892_at	J04177	Hs.523446	collagen, type XI, alpha 1	COL11A1	EG2	0	4	3	6	0,1119520740,440114659	8,42E-05	0,002977773
229872_s_at	AA532655	Hs.549388		LOC440667	EG2	0	3	3	11	0,2913855570,799386187	2,17E-10	0,001562433
231907_at	AK025877	Hs.159472	v-abl Abelson murine leukemia viral oncogene homol	ABL2	EG2	2	1	9	5	0,2913855570,799386187	2,17E-10	0,001562433
1555788_a_at	AF250311	Hs.516826	tribbles homolog 3 (Drosophila)	TRIB3	EG2	1	2	8	5	0,2913855570,799386187	4,56E-09	0,001562433
204222_s_at	NM_006851	Hs.553516	GLI pathogenesis-related 1 (glioma)	GLIPR1	EG2	3	0	7	6	0,2913855570,799386187	4,56E-09	0,001562433
211272_s_at	AF064771	Hs.524488	diacylglycerol kinase, alpha 80kDa	DGKA	EG2	2	1	5	8	0,2913855570,799386187	4,56E-09	0,001562433
215307_at	AL109722				EG2	2	1	6	7	0,2913855570,799386187	4,56E-09	0,001562433
217914_at	NM_017901	Hs.524763	two pore segment channel 1	TPCN1	EG2	1	2	5	8	0,2913855570,799386187	4,56E-09	0,001562433
218953_s_at	NM_024028	Hs.483796		MGC3265	EG2	2	1	6	7	0,2913855570,799386187	4,56E-09	0,001562433
227530_at	BF511276	Hs.371240	A kinase (PRKA) anchor protein (gravin) 12	AKAP12	EG2	1	2	7	6	0,2913855570,799386187	4,56E-09	0,001562433
228726_at	AW512196	Hs.381167	serpin peptidase inhibitor, clade B (ovalbumin), mem	SERPINB1	EG2	2	1	5	8	0,2913855570,799386187	4,56E-09	0,001562433
231993_at	AK026784				EG2	2	1	4	9	0,2913855570,799386187	4,56E-09	0,001562433
203157_s_at	AB020645	Hs.116448	glutaminase	GLS	EG2	3	0	8	4	0,2913855570,799386187	7,43E-08	0,001562433
203708_at	NM_002600	Hs.198072	phosphodiesterase 4B, cAMP-specific (phosphodieste	PDE4B	EG2	1	2	5	7	0,2913855570,799386187	7,43E-08	0,001562433
206342_x_at	NM_006123	Hs.460960	iduronate 2-sulfatase (Hunter syndrome)	IDS	EG2	2	1	8	4	0,2913855570,799386187	7,43E-08	0,001562433
210511_s_at	M13436	Hs.28792	inhibin, beta A (activin A, activin AB alpha polypeptid	INHBA	EG2	2	1	9	3	0,2913855570,799386187	7,43E-08	0,001562433
212311_at	AA522514	Hs.479384		KIAA0746	EG2	1	2	9	3	0,2913855570,799386187	7,43E-08	0,001562433
219773_at	NM_016931	Hs.371036	NADPH oxidase 4	NOX4	EG2	1	2	4	8	0,2913855570,799386187	7,43E-08	0,001562433
221019_s_at	NM_030781	Hs.464422	collectin sub-family member 12	COLEC12	EG2	0	3	2	10	0,2913855570,799386187	7,43E-08	0,001562433
225919_s_at	AI832598	Hs.493639	chromosome 9 open reading frame 72	C9orf72	EG2	3	0	8	4	0,2913855570,799386187	7,43E-08	0,001562433
235629_at	AI333596	Hs.203717	fibronectin 1	FN1	EG2	2	1	6	6	0,2913855570,799386187	7,43E-08	0,001562433
235944_at	BF446673	Hs.58877	hemicentin 1	HMCN1	EG2	2	1	5	7	0,2913855570,799386187	7,43E-08	0,001562433
238622_at	AW269686	Hs.98643	RAP2B, member of RAS oncogene family	RAP2B	EG2	2	1	4	8	0,2913855570,799386187	7,43E-08	0,001562433
201141_at	NM_002510	Hs.190495	glycoprotein (transmembrane) nmb	GPNMB	EG2	1	2	9	2	0,2913855570,799386187	9,59E-07	0,001562433
201360_at	NM_000099	Hs.304682	cystatin C (amyloid angiopathy and cerebral hemorrha	CST3	EG2	0	3	5	6	0,2913855570,799386187	9,59E-07	0,001562433
203108_at	NM_003979	Hs.546325	G protein-coupled receptor, family C, group 5, membe	GPRC5A	EG2	1	2	4	7	0,2913855570,799386187	9,59E-07	0,001562433
203908_at	NM_003759	Hs.5462	solute carrier family 4, sodium bicarbonate cotranspor	SLC4A4	EG2	2	1	9	2	0,2913855570,799386187	9,59E-07	0,001562433
204678_s_at	U90065	Hs.208544	potassium channel, subfamily K, member 1	KCNK1	EG2	1	2	9	2	0,2913855570,799386187	9,59E-07	0,001562433
210534_s_at	BC002944	Hs.462445		EPPB9	EG2	1	2	8	3	0,2913855570,799386187	9,59E-07	0,001562433
213032_at	AI186739	Hs.370359	nuclear factor I/B	NFIB	EG2	2	1	3	8	0,2913855570,799386187	9,59E-07	0,001562433
213703_at	W95043	Hs.531822		LOC150759	EG2	3	0	9	2	0,2913855570,799386187	9,59E-07	0,001562433
214701_s_at	AJ276395	Hs.203717	fibronectin 1	FN1	EG2	2	1	5	6	0,2913855570,799386187	9,59E-07	0,001562433
226492_at	AL036088	Hs.511265	sema domain, transmembrane domain (TM), and cyto	SEMA6D	EG2	0	3	8	3	0,2913855570,799386187	9,59E-07	0,001562433
227955_s_at	BE670307				EG2	2	1	6	5	0,2913855570,799386187	9,59E-07	0,001562433
228057_at	AA528140	Hs.480378	DNA-damage-inducible transcript 4-like	DDIT4L	EG2	2	1	9	2	0,2913855570,799386187	9,59E-07	0,001562433
235457_at	AI769569	Hs.428214	mastermind-like 2 (Drosophila)	MAML2	EG2	3	0	2	9	0,2913855570,799386187	9,59E-07	0,001562433
237411_at	N71063	Hs.344757,	HADAM metalloproteinase with thrombospondin type 1	ADAMTS6	EG2	1	2	10	1	0,2913855570,799386187	9,59E-07	0,001562433
1555526_a_at	AF403061	Hs.496666	septin 6	SEPT6	EG2	3	0	3	7	0,2913855570,799386187	9,97E-06	0,001562433

1558199_at	W73431	Hs.203717	fibronectin 1	FN1	EG2	3	0	6	4	0,2913855570,799386187	9,97E-06	0,001562433
1568609_s_at	AL118843	Hs.381235,	H phosphodiesterase 4D interacting protein (myomegali	PDE4DIP	EG2	1	2	5	5	0,2913855570,799386187	9,97E-06	0,001562433
1569472_s_at	BC026260	Hs.368214	tetratricopeptide repeat domain 3	TTC3	EG2	2	1	6	4	0,2913855570,799386187	9,97E-06	0,001562433
205347_s_at	NM_021992	Hs.56145	thymosin-like 8	TMSL8	EG2	3	0	4	6	0,2913855570,799386187	9,97E-06	0,001562433
205381_at	NM_005824	Hs.550454	leucine rich repeat containing 17	LRRC17	EG2	0	3	3	7	0,2913855570,799386187	9,97E-06	0,001562433
205397_x_at	U76622	Hs.36915,	Hs SMAD, mothers against DPP homolog 3 (Drosophila)	SMAD3	EG2	1	2	4	6	0,2913855570,799386187	9,97E-06	0,001562433
207357_s_at	NM_017540	Hs.34421,	Hs UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-	GALNT10	EG2	1	2	8	2	0,2913855570,799386187	9,97E-06	0,001562433
209406_at	AF095192	Hs.55220	BCL2-associated athanogene 2	BAG2	EG2	2	1	5	5	0,2913855570,799386187	9,97E-06	0,001562433
209598_at	AB020690	Hs.521466	paraneoplastic antigen MA2	PNMA2	EG2	0	3	4	6	0,2913855570,799386187	9,97E-06	0,001562433
209760_at	AL136932	Hs.205572		KIAA0922	EG2	2	1	1	9	0,2913855570,799386187	9,97E-06	0,001562433
210102_at	BC001234	Hs.152944	loss of heterozygosity, 11, chromosomal region 2, gen	LOH11CR2A	EG2	3	0	4	6	0,2913855570,799386187	9,97E-06	0,001562433
210869_s_at	M29277	Hs.511397	melanoma cell adhesion molecule	MCAM	EG2	2	1	3	7	0,2913855570,799386187	9,97E-06	0,001562433
211376_s_at	BC005212	Hs.258798	chromosome 10 open reading frame 86	C10orf86	EG2	1	2	3	7	0,2913855570,799386187	9,97E-06	0,001562433
214587_at	BE877796	Hs.134830	collagen, type VIII, alpha 1	COL8A1	EG2	2	1	5	5	0,2913855570,799386187	9,97E-06	0,001562433
216048_s_at	AK023621	Hs.445030	Rho-related BTB domain containing 3	RHOBTB3	EG2	2	1	5	5	0,2913855570,799386187	9,97E-06	0,001562433
218717_s_at	NM_018192	Hs.374191	leprecan-like 1	LEPREL1	EG2	2	1	8	2	0,2913855570,799386187	9,97E-06	0,001562433
219506_at	NM_024579	Hs.91283	chromosome 1 open reading frame 54	C1orf54	EG2	0	3	5	5	0,2913855570,799386187	9,97E-06	0,001562433
221679_s_at	AF225418	Hs.476454	abhydrolase domain containing 6	ABHD6	EG2	2	1	8	2	0,2913855570,799386187	9,97E-06	0,001562433
225583_at	AL573637	Hs.469561	UDP-glucuronate decarboxylase 1	UXS1	EG2	2	1	3	7	0,2913855570,799386187	9,97E-06	0,001562433
227074_at	AA524669	Hs.348418	down-regulator of transcription 1, TBP-binding (negati	DR1	EG2	1	2	3	7	0,2913855570,799386187	9,97E-06	0,001562433
227290_at	AU157881				EG2	2	1	5	5	0,2913855570,799386187	9,97E-06	0,001562433
227415_at	BF109303	Hs.527910	diacylglycerol kinase, eta	DGKH	EG2	3	0	3	7	0,2913855570,799386187	9,97E-06	0,001562433
227487_s_at	AI359165	Hs.38449	serpin peptidase inhibitor, clade E (nexin, plasminoge	SERPINE2	EG2	1	2	9	1	0,2913855570,799386187	9,97E-06	0,001562433
227802_at	AI075999	Hs.518801		LOC441022	EG2	2	1	5	5	0,2913855570,799386187	9,97E-06	0,001562433
228158_at	AI623211				EG2	2	1	4	6	0,2913855570,799386187	9,97E-06	0,001562433
228191_at	AA206016				EG2	3	0	7	3	0,2913855570,799386187	9,97E-06	0,001562433
228242_at	BF055201				EG2	3	0	3	7	0,2913855570,799386187	9,97E-06	0,001562433
228566_at	AA778694	Hs.464912		P15RS	EG2	3	0	7	3	0,2913855570,799386187	9,97E-06	0,001562433
229285_at	AI669749	Hs.518545	ribonuclease L (2',5'-oligoadenylate synthetase-de	RNASEL	EG2	1	2	2	8	0,2913855570,799386187	9,97E-06	0,001562433
229838_at	AI377271	Hs.128686	nucleobindin 2	NUCB2	EG2	3	0	3	7	0,2913855570,799386187	9,97E-06	0,001562433
230722_at	AI377043	Hs.435309	basonuclin 2	BNC2	EG2	3	0	6	4	0,2913855570,799386187	9,97E-06	0,001562433
232060_at	AK000776				EG2	2	1	2	8	0,2913855570,799386187	9,97E-06	0,001562433
235088_at	AI276663	Hs.380920		LOC201725	EG2	3	0	5	5	0,2913855570,799386187	9,97E-06	0,001562433
235320_at	AL138043	Hs.373801	ADP-ribosylation factor-like 6	ARL6	EG2	2	1	7	3	0,2913855570,799386187	9,97E-06	0,001562433
239336_at	BF109732	Hs.164226	thrombospondin 1	THBS1	EG2	1	2	4	6	0,2913855570,799386187	9,97E-06	0,001562433
1553962_s_at	BI668074	Hs.502876	ras homolog gene family, member B	RHOB	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
1554250_s_at	BC033812	Hs.511015,	H tripartite motif-containing 50B	TRIM50B	EG2	1	2	2	7	0,2913855570,799386187	8,42E-05	0,002977773
1555870_at	AK056658	Hs.444729	chromosome 1 open reading frame 188	C1orf188	EG2	1	2	2	7	0,2913855570,799386187	8,42E-05	0,002977773
1555920_at	BU683892	Hs.381189	chromobox homolog 3 (HP1 gamma homolog, Drosop	CBX3	EG2	1	2	5	4	0,2913855570,799386187	8,42E-05	0,002977773
1556641_at	AK094547	Hs.42834		FLJ37228	EG2	0	3	0	9	0,2913855570,799386187	8,42E-05	0,002977773
1557905_s_at	AL552534	Hs.502328	CD44 antigen (homing function and Indian blood grou	CD44	EG2	1	2	5	4	0,2913855570,799386187	8,42E-05	0,002977773
1560296_at	AV702787	Hs.485616	dystonin	DST	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
1567224_at	U29113	Hs.505924	high mobility group AT-hook 2	HMGA2	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
1568619_s_at	BC015343	Hs.530899		LOC162073	EG2	0	3	6	3	0,2913855570,799386187	8,42E-05	0,002977773
200766_at	NM_001909	Hs.121575,	H cathepsin D (lysosomal aspartyl peptidase)	CTSD	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773

201262_s_at	NM_001711	Hs.821	biglycan	BGN	EG2	1	2	5	4	0,2913855570,799386187	8,42E-05	0,002977773
201325_s_at	NM_001423	Hs.436298	epithelial membrane protein 1	EMP1	EG2	3	0	5	4	0,2913855570,799386187	8,42E-05	0,002977773
201425_at	NM_000690	Hs.436437	aldehyde dehydrogenase 2 family (mitochondrial)	ALDH2	EG2	0	3	5	4	0,2913855570,799386187	8,42E-05	0,002977773
201539_s_at	U29538	Hs.435369	four and a half LIM domains 1	FHL1	EG2	1	2	1	8	0,2913855570,799386187	8,42E-05	0,002977773
201581_at	BF572868	Hs.169358	thioredoxin domain containing 13	TXNDC13	EG2	3	0	5	4	0,2913855570,799386187	8,42E-05	0,002977773
201818_at	NM_024830	Hs.368853		FLJ12443	EG2	1	2	8	1	0,2913855570,799386187	8,42E-05	0,002977773
201869_s_at	BF593932	Hs.495656	transducin (beta)-like 1X-linked	TBL1X	EG2	2	1	5	4	0,2913855570,799386187	8,42E-05	0,002977773
202341_s_at	AA149745	Hs.435711	tripartite motif-containing 2	TRIM2	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
202382_s_at	NM_005471	Hs.278500	glucosamine-6-phosphate deaminase 1	GNPDA1	EG2	1	2	6	3	0,2913855570,799386187	8,42E-05	0,002977773
203153_at	NM_001548	Hs.20315	interferon-induced protein with tetratricopeptide rep	IFIT1	EG2	1	2	5	4	0,2913855570,799386187	8,42E-05	0,002977773
203217_s_at	NM_003896	Hs.415117	ST3 beta-galactoside alpha-2,3-sialyltransferase 5	ST3GAL5	EG2	1	2	6	3	0,2913855570,799386187	8,42E-05	0,002977773
204084_s_at	AI911687	Hs.30213	ceroid-lipofuscinosis, neuronal 5	CLN5	EG2	2	1	3	6	0,2913855570,799386187	8,42E-05	0,002977773
204137_at	NM_003272	Hs.498160	transmembrane 7 superfamily member 1 (upregulate	TM7SF1	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
204140_at	NM_003596	Hs.421194	tyrosylprotein sulfotransferase 1	TPST1	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
204493_at	NM_001196	Hs.474150	BH3 interacting domain death agonist	BID	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
205407_at	NM_021111	Hs.388918	reversion-inducing-cysteine-rich protein with kazal m	RECK	EG2	1	2	2	7	0,2913855570,799386187	8,42E-05	0,002977773
205540_s_at	NM_016656	Hs.50282	Ras-related GTP binding B	RRAGB	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
206777_s_at	NM_000496	Hs.373074	crystallin, beta B2	CRYBB2	EG2	0	3	8	1	0,2913855570,799386187	8,42E-05	0,002977773
206907_at	NM_003811	Hs.1524	tumor necrosis factor (ligand) superfamily, member 9	TNFSF9	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
208230_s_at	NM_013960	Hs.453951	neuregulin 1	NRG1	EG2	2	1	9	0	0,2913855570,799386187	8,42E-05	0,002977773
209035_at	M69148	Hs.82045	midkine (neurite growth-promoting factor 2)	MDK	EG2	0	3	1	8	0,2913855570,799386187	8,42E-05	0,002977773
209146_at	AV704962	Hs.105269	sterol-C4-methyl oxidase-like	SC4MOL	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
209277_at	AL574096	Hs.438231	tissue factor pathway inhibitor 2	TFPI2	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
209373_at	BC003179	Hs.185055	mal, T-cell differentiation protein-like	MALL	EG2	2	1	5	4	0,2913855570,799386187	8,42E-05	0,002977773
209409_at	D86962	Hs.164060	growth factor receptor-bound protein 10	GRB10	EG2	2	1	3	6	0,2913855570,799386187	8,42E-05	0,002977773
209537_at	AF000416	Hs.357637	exostoses (multiple)-like 2	EXTL2	EG2	3	0	9	0	0,2913855570,799386187	8,42E-05	0,002977773
209758_s_at	U37283	Hs.512842	microfibrillar associated protein 5	MFAP5	EG2	0	3	3	6	0,2913855570,799386187	8,42E-05	0,002977773
210059_s_at	BC000433	Hs.178695	mitogen-activated protein kinase 13	MAPK13	EG2	0	3	7	2	0,2913855570,799386187	8,42E-05	0,002977773
210732_s_at	AF342816	Hs.4082	lectin, galactoside-binding, soluble, 8 (galectin 8)	LGALS8	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
210916_s_at	AF098641	Hs.502328	CD44 antigen (homing function and Indian blood grou	CD44	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
211208_s_at	AB039327	Hs.495984	calcium/calmodulin-dependent serine protein kinase	CASK	EG2	2	1	1	8	0,2913855570,799386187	8,42E-05	0,002977773
211825_s_at	AF327066	Hs.374477	Ewing sarcoma breakpoint region 1	EWSR1	EG2	3	0	5	4	0,2913855570,799386187	8,42E-05	0,002977773
211980_at	AI922605	Hs.17441	collagen, type IV, alpha 1	COL4A1	EG2	1	2	4	5	0,2913855570,799386187	8,42E-05	0,002977773
212959_s_at	AK001821	Hs.46850	N-acetylglucosamine-1-phosphate transferase, alpha	GNPTAB	EG2	2	1	8	1	0,2913855570,799386187	8,42E-05	0,002977773
213764_s_at	AW665892	Hs.512842	microfibrillar associated protein 5	MFAP5	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
214250_at	AI337584	Hs.523873	nuclear mitotic apparatus protein 1	NUMA1	EG2	1	2	2	7	0,2913855570,799386187	8,42E-05	0,002977773
214505_s_at	AF220153	Hs.435369	four and a half LIM domains 1	FHL1	EG2	1	2	1	8	0,2913855570,799386187	8,42E-05	0,002977773
214702_at	AJ276395	Hs.203717	fibronectin 1	FN1	EG2	1	2	4	5	0,2913855570,799386187	8,42E-05	0,002977773
214723_x_at	AB046861	Hs.503490, H	KIAA1641	KIAA1641	EG2	1	2	6	3	0,2913855570,799386187	8,42E-05	0,002977773
214927_at	AL359052	Hs.508597	integrin, beta-like 1 (with EGF-like repeat domains)	ITGBL1	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
215775_at	BF084105	Hs.164226	thrombospondin 1	THBS1	EG2	0	3	3	6	0,2913855570,799386187	8,42E-05	0,002977773
217785_s_at	NM_006555	Hs.520794		YKT6	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
218051_s_at	NM_022908	Hs.84753		FLJ12442	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
218934_s_at	NM_014424	Hs.502612	heat shock 27kDa protein family, member 7 (cardiovas	HSPB7	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
219501_at	NM_017993	Hs.128258		FLJ10094	EG2	3	0	7	2	0,2913855570,799386187	8,42E-05	0,002977773

220615_s_at	NM_018099	Hs.298851	male sterility domain containing 1	MLSTD1	EG2	1	2	1	8	0,2913855570,799386187	8,42E-05	0,002977773
221142_s_at	NM_018441	Hs.281680	peroxisomal trans-2-enoyl-CoA reductase	PECR	EG2	0	3	2	7	0,2913855570,799386187	8,42E-05	0,002977773
221882_s_at	AI636233	Hs.288940	transmembrane protein 8 (five membrane-spanning d	TMEM8	EG2	0	3	4	5	0,2913855570,799386187	8,42E-05	0,002977773
223271_s_at	AF161543	Hs.497967	CTD (carboxy-terminal domain, RNA polymerase II, po	CTDSPL2	EG2	3	0	4	5	0,2913855570,799386187	8,42E-05	0,002977773
223287_s_at	AF146696	Hs.431498	forkhead box P1	FOXP1	EG2	0	3	4	5	0,2913855570,799386187	8,42E-05	0,002977773
223374_s_at	AF154848	Hs.418062	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, p	B3GALT3	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
223395_at	AB056106	Hs.477015	ABI gene family, member 3 (NESH) binding protein	ABI3BP	EG2	0	3	7	2	0,2913855570,799386187	8,42E-05	0,002977773
224755_at	BE621524	Hs.500674		SMBP	EG2	3	0	4	5	0,2913855570,799386187	8,42E-05	0,002977773
225140_at	BF438116	Hs.298658	Kruppel-like factor 3 (basic)	KLF3	EG2	2	1	8	1	0,2913855570,799386187	8,42E-05	0,002977773
225579_at	AL516202	Hs.274415	PQ loop repeat containing 3	PQLC3	EG2	0	3	4	5	0,2913855570,799386187	8,42E-05	0,002977773
225640_at	AA875998	Hs.446271		LOC401504	EG2	1	2	6	3	0,2913855570,799386187	8,42E-05	0,002977773
225742_at	W93501	Hs.497492	Mdm4, transformed 3T3 cell double minute 4, p53 bin	MDM4	EG2	2	1	5	4	0,2913855570,799386187	8,42E-05	0,002977773
225871_at	BF680588	Hs.489051	six transmembrane epithelial antigen of the prostate	STEAP2	EG2	1	2	7	2	0,2913855570,799386187	8,42E-05	0,002977773
226534_at	AI446414	Hs.1048	KIT ligand	KITLG	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
226908_at	AI627704	Hs.253736	leucine-rich repeats and immunoglobulin-like domain	LRIG3	EG2	2	1	3	6	0,2913855570,799386187	8,42E-05	0,002977773
226912_at	AW003367	Hs.21902	zinc finger, DHHC-type containing 23	ZDHHC23	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
227408_s_at	AW008976	Hs.369091	sorting nexin 25	SNX25	EG2	3	0	4	5	0,2913855570,799386187	8,42E-05	0,002977773
227525_at	AA058770	Hs.131673	glucocorticoid induced transcript 1	GLCCI1	EG2	3	0	2	7	0,2913855570,799386187	8,42E-05	0,002977773
228314_at	BE877357				EG2	3	0	5	4	0,2913855570,799386187	8,42E-05	0,002977773
228613_at	BF183535	Hs.531642	RAB11 family interacting protein 3 (class II)	RAB11FIP3	EG2	3	0	5	4	0,2913855570,799386187	8,42E-05	0,002977773
229010_at	AI807026	Hs.504096	Cas-Br-M (murine) ecotropic retroviral transforming s	CBL	EG2	2	1	2	7	0,2913855570,799386187	8,42E-05	0,002977773
229512_at	BE464337				EG2	2	1	3	6	0,2913855570,799386187	8,42E-05	0,002977773
230000_at	AI523817	Hs.195642	chromosome 17 open reading frame 27	C17orf27	EG2	1	2	4	5	0,2913855570,799386187	8,42E-05	0,002977773
230165_at	N31731	Hs.44269	shugoshin-like 2 (S. pombe)	SGOL2	EG2	2	1	4	5	0,2913855570,799386187	8,42E-05	0,002977773
230424_at	AU144860	Hs.483067	chromosome 5 open reading frame 13	C5orf13	EG2	0	3	6	3	0,2913855570,799386187	8,42E-05	0,002977773
234863_x_at	AK026197	Hs.520506	F-box protein 5	FBXO5	EG2	2	1	6	3	0,2913855570,799386187	8,42E-05	0,002977773
235147_at	R56118	Hs.528187		LOC440928	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
235287_at	AW192700	Hs.119882	cyclin-dependent kinase 6	CDK6	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
235746_s_at	BE048919	Hs.410477	phospholipase A2 receptor 1, 180kDa	PLA2R1	EG2	0	3	2	7	0,2913855570,799386187	8,42E-05	0,002977773
238722_x_at	AI460037	Hs.324271		NAPE-PLD	EG2	1	2	3	6	0,2913855570,799386187	8,42E-05	0,002977773
241893_at	BE927766				EG2	2	1	3	6	0,2913855570,799386187	8,42E-05	0,002977773
243507_s_at	AV715251	Hs.529340		FLJ25067	EG2	2	1	2	7	0,2913855570,799386187	8,42E-05	0,002977773
243982_at	AA455180	Hs.174682	BTB (POZ) domain containing 5	BTBD5	EG2	2	1	7	2	0,2913855570,799386187	8,42E-05	0,002977773
244669_at	T16443	Hs.292457		LOC441164	EG2	3	0	2	7	0,2913855570,799386187	8,42E-05	0,002977773
212677_s_at	BG530481	Hs.310645	RAB1A, member RAS oncogene family	RAB1A	EG2	0	2	4	10	0,5783259580,896050827	2,17E-10	0,001562433
209217_s_at	BC000464	Hs.156701	WD repeat domain 45	WDR45	EG2	1	1	8	5	0,5783259580,896050827	4,56E-09	0,001562433
227140_at	AI343467	Hs.28792	inhibin, beta A (activin A, activin AB alpha polypeptid	INHBA	EG2	1	1	9	4	0,5783259580,896050827	4,56E-09	0,001562433
202625_at	AI356412	Hs.491767	v-yes-1 Yamaguchi sarcoma viral related oncogene ho	LYN	EG2	2	0	6	6	0,5783259580,896050827	7,43E-08	0,001562433
205463_s_at	NM_002607	Hs.376032	H platelet-derived growth factor alpha polypeptide	PDGFA	EG2	1	1	2	10	0,5783259580,896050827	7,43E-08	0,001562433
1557242_at	AI769729	Hs.130838		LOC285835	EG2	1	1	7	4	0,5783259580,896050827	9,59E-07	0,001562433
201578_at	NM_005397	Hs.16426	podocalyxin-like	PODXL	EG2	0	2	6	5	0,5783259580,896050827	9,59E-07	0,001562433
202975_s_at	N21138	Hs.445030	Rho-related BTB domain containing 3	RHOBTB3	EG2	1	1	7	4	0,5783259580,896050827	9,59E-07	0,001562433
204797_s_at	NM_004434	Hs.12451	echinoderm microtubule associated protein like 1	EML1	EG2	2	0	5	6	0,5783259580,896050827	9,59E-07	0,001562433
205634_x_at	NM_024298	Hs.467279	leukocyte receptor cluster (LRC) member 4	LENG4	EG2	1	1	8	3	0,5783259580,896050827	9,59E-07	0,001562433
206059_at	NM_003430	Hs.531632	zinc finger protein 91 (HPF7, HTF10)	ZNF91	EG2	2	0	7	4	0,5783259580,896050827	9,59E-07	0,001562433

209940_at	AF083068	Hs.271742	poly (ADP-ribose) polymerase family, member 3	PARP3	EG2	0	2	4	7	0,5783259580,896050827	9,59E-07	0,001562433
213260_at	AU145890	Hs.348883	forkhead box C1	FOXC1	EG2	2	0	1	10	0,5783259580,896050827	9,59E-07	0,001562433
213572_s_at	AI554300	Hs.381167	serpin peptidase inhibitor, clade B (ovalbumin), mem	SERPINB1	EG2	0	2	5	6	0,5783259580,896050827	9,59E-07	0,001562433
219210_s_at	NM_016530	Hs.389733	RAB8B, member RAS oncogene family	RAB8B	EG2	2	0	9	2	0,5783259580,896050827	9,59E-07	0,001562433
238478_at	H97386	Hs.435309	basonuclin 2	BNC2	EG2	2	0	5	6	0,5783259580,896050827	9,59E-07	0,001562433
1554097_a_at	BC021861	Hs.458096		LOC554202	EG2	1	1	6	4	0,5783259580,896050827	9,97E-06	0,001562433
201538_s_at	NM_004090	Hs.181046	dual specificity phosphatase 3 (vaccinia virus phospho	DUSP3	EG2	1	1	9	1	0,5783259580,896050827	9,97E-06	0,001562433
202037_s_at	NM_003012	Hs.213424	secreted frizzled-related protein 1	SFRP1	EG2	1	1	7	3	0,5783259580,896050827	9,97E-06	0,001562433
202510_s_at	NM_006291	Hs.525607	tumor necrosis factor, alpha-induced protein 2	TNFAIP2	EG2	0	2	5	5	0,5783259580,896050827	9,97E-06	0,001562433
202627_s_at	AL574210	Hs.414795	serpin peptidase inhibitor, clade E (nexin, plasminoge	SERPINE1	EG2	2	0	7	3	0,5783259580,896050827	9,97E-06	0,001562433
203385_at	NM_001345	Hs.524488	diacylglycerol kinase, alpha 80kDa	DGKA	EG2	0	2	3	7	0,5783259580,896050827	9,97E-06	0,001562433
203504_s_at	NM_005502	Hs.429294	ATP-binding cassette, sub-family A (ABC1), member 1	ABCA1	EG2	1	1	3	7	0,5783259580,896050827	9,97E-06	0,001562433
203579_s_at	AI660619	Hs.334848	solute carrier family 7, member 6 opposite strand	SLC7A6OS	EG2	1	1	5	5	0,5783259580,896050827	9,97E-06	0,001562433
203741_s_at	NM_001114	Hs.513578	adenylate cyclase 7	ADCY7	EG2	0	2	4	6	0,5783259580,896050827	9,97E-06	0,001562433
203951_at	NM_001299	Hs.465929	calponin 1, basic, smooth muscle	CNN1	EG2	1	1	2	8	0,5783259580,896050827	9,97E-06	0,001562433
204925_at	NM_004937	Hs.187667	cystinosis, nephropathic	CTNS	EG2	0	2	7	3	0,5783259580,896050827	9,97E-06	0,001562433
206396_at	NM_004170	Hs.444915	solute carrier family 1 (neuronal/epithelial high affini	SLC1A1	EG2	1	1	7	3	0,5783259580,896050827	9,97E-06	0,001562433
218000_s_at	NM_007350	Hs.484885	pleckstrin homology-like domain, family A, member 1	PHLDA1	EG2	2	0	9	1	0,5783259580,896050827	9,97E-06	0,001562433
221577_x_at	AF003934	Hs.515258	growth differentiation factor 15	GDF15	EG2	2	0	9	1	0,5783259580,896050827	9,97E-06	0,001562433
222719_s_at	AB033831	Hs.148162	platelet derived growth factor C	PDGFC	EG2	2	0	8	2	0,5783259580,896050827	9,97E-06	0,001562433
223734_at	AF329088	Hs.154140		OSAP	EG2	1	1	4	6	0,5783259580,896050827	9,97E-06	0,001562433
223789_s_at	AF116627	Hs.485449	GTP binding protein 2	GTPBP2	EG2	1	1	5	5	0,5783259580,896050827	9,97E-06	0,001562433
224942_at	BG434272	Hs.494928	pregnancy-associated plasma protein A, pappalysin 1	PAPPA	EG2	2	0	7	3	0,5783259580,896050827	9,97E-06	0,001562433
224964_s_at	AK026424	Hs.187772	guanine nucleotide binding protein (G protein), gam	GNG2	EG2	2	0	5	5	0,5783259580,896050827	9,97E-06	0,001562433
226278_at	AI150224	Hs.349096		DKFZp313A2432	EG2	0	2	4	6	0,5783259580,896050827	9,97E-06	0,001562433
226810_at	BE500942	Hs.368337	chromosome 6 open reading frame 155	C6orf155	EG2	2	0	8	2	0,5783259580,896050827	9,97E-06	0,001562433
227396_at	AI631833				EG2	2	0	7	3	0,5783259580,896050827	9,97E-06	0,001562433
229011_at	AA150501				EG2	2	0	6	4	0,5783259580,896050827	9,97E-06	0,001562433
231067_s_at	BF114967	Hs.371240	A kinase (PRKA) anchor protein (gravin) 12	AKAP12	EG2	1	1	4	6	0,5783259580,896050827	9,97E-06	0,001562433
232790_at	AI744580				EG2	1	1	3	7	0,5783259580,896050827	9,97E-06	0,001562433
235311_at	BG110260	Hs.390838	FK506 binding protein 14, 22 kDa	FKBP14	EG2	2	0	8	2	0,5783259580,896050827	9,97E-06	0,001562433
1557080_s_at	AI753143	Hs.508597	integrin, beta-like 1 (with EGF-like repeat domains)	ITGBL1	EG2	1	1	3	6	0,5783259580,896050827	8,42E-05	0,002977773
201185_at	NM_002775	Hs.501280	HtrA serine peptidase 1	HTRA1	EG2	0	2	5	4	0,5783259580,896050827	8,42E-05	0,002977773
201615_x_at	AI685060	Hs.490203	caldesmon 1	CALD1	EG2	2	0	3	6	0,5783259580,896050827	8,42E-05	0,002977773
201668_x_at	AW163148	Hs.519909	myristoylated alanine-rich protein kinase C substrate	MARCKS	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
202948_at	NM_000877	Hs.516249	interleukin 1 receptor, type I	IL1R1	EG2	0	2	2	7	0,5783259580,896050827	8,42E-05	0,002977773
203354_s_at	AW117368	Hs.434255	pleckstrin and Sec7 domain containing 3	PSD3	EG2	2	0	2	7	0,5783259580,896050827	8,42E-05	0,002977773
203424_s_at	AW157548	Hs.369982,	H insulin-like growth factor binding protein 5	IGFBP5	EG2	2	0	3	6	0,5783259580,896050827	8,42E-05	0,002977773
203827_at	NM_017983	Hs.463964		WIPI49	EG2	0	2	6	3	0,5783259580,896050827	8,42E-05	0,002977773
203879_at	U86453	Hs.518451	phosphoinositide-3-kinase, catalytic, delta polypeptid	PIK3CD	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
204807_at	BF224146	Hs.216386	transmembrane protein 5	TMEM5	EG2	1	1	3	6	0,5783259580,896050827	8,42E-05	0,002977773
204979_s_at	NM_007341	Hs.473847	SH3 domain binding glutamic acid-rich protein	SH3BGR	EG2	1	1	8	1	0,5783259580,896050827	8,42E-05	0,002977773
205234_at	NM_004696	Hs.351306	solute carrier family 16 (monocarboxylic acid transpor	SLC16A4	EG2	0	2	4	5	0,5783259580,896050827	8,42E-05	0,002977773
205741_s_at	NM_001392	Hs.58919	dystrobrevin, alpha	DTNA	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
205925_s_at	NM_002867	Hs.123072	RAB3B, member RAS oncogene family	RAB3B	EG2	2	0	6	3	0,5783259580,896050827	8,42E-05	0,002977773

207264_at	NM_016657	Hs.554798	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protei	KDELR3	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
208107_s_at	NM_030941	Hs.177926		LOC81691	EG2	1	1	5	4	0,5783259580,896050827	8,42E-05	0,002977773
208191_x_at	NM_002780	Hs.466860	pregnancy specific beta-1-glycoprotein 4	PSG4	EG2	0	2	6	3	0,5783259580,896050827	8,42E-05	0,002977773
209216_at	BC000464	Hs.156701	WD repeat domain 45	WDR45	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
211340_s_at	M28882	Hs.511397	melanoma cell adhesion molecule	MCAM	EG2	0	2	2	7	0,5783259580,896050827	8,42E-05	0,002977773
211713_x_at	BC005832	Hs.81892	KIAA0101	KIAA0101	EG2	2	0	3	6	0,5783259580,896050827	8,42E-05	0,002977773
212386_at	BF592782	Hs.200285	transcription factor 4	TCF4	EG2	1	1	4	5	0,5783259580,896050827	8,42E-05	0,002977773
213206_at	AW149492	Hs.463278	golgi SNAP receptor complex member 2	GOSR2	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
213765_at	AW665892	Hs.512842	microfibrillar associated protein 5	MFAP5	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
213959_s_at	BF515597	Hs.298382		KIAA1005	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
214151_s_at	AU144243	Hs.126115	phosphatidylinositol glycan, class B	PIGB	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
214581_x_at	BE568134	Hs.443577	tumor necrosis factor receptor superfamily, member 2	TNFRSF21	EG2	2	0	4	5	0,5783259580,896050827	8,42E-05	0,002977773
214724_at	AF070621	Hs.116796	DIX domain containing 1	DIXDC1	EG2	1	1	3	6	0,5783259580,896050827	8,42E-05	0,002977773
217502_at	BE888744	Hs.437609	interferon-induced protein with tetratricopeptide rep	IFIT2	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
218284_at	NM_015400	Hs.36915, Hs	SMAD, mothers against DPP homolog 3 (Drosophila)	SMAD3	EG2	0	2	4	5	0,5783259580,896050827	8,42E-05	0,002977773
218665_at	NM_012193	Hs.19545	frizzled homolog 4 (Drosophila)	FZD4	EG2	0	2	3	6	0,5783259580,896050827	8,42E-05	0,002977773
218986_s_at	NM_017631	Hs.481141		FLJ20035	EG2	0	2	2	7	0,5783259580,896050827	8,42E-05	0,002977773
219703_at	NM_018365	Hs.444483	meiosis-specific nuclear structural 1	MNS1	EG2	2	0	4	5	0,5783259580,896050827	8,42E-05	0,002977773
221012_s_at	NM_030912	Hs.336810	tripartite motif-containing 8	TRIM8	EG2	0	2	4	5	0,5783259580,896050827	8,42E-05	0,002977773
221139_s_at	NM_015989	Hs.279815	cysteine sulfinic acid decarboxylase	CSAD	EG2	0	2	4	5	0,5783259580,896050827	8,42E-05	0,002977773
221489_s_at	W48843	Hs.323308	sprouty homolog 4 (Drosophila)	SPRY4	EG2	1	1	7	2	0,5783259580,896050827	8,42E-05	0,002977773
222020_s_at	AW117456	Hs.504352		HNT	EG2	1	1	6	3	0,5783259580,896050827	8,42E-05	0,002977773
222621_at	BF591419	Hs.499000	DnaJ (Hsp40) homolog, subfamily C, member 1	DNAJC1	EG2	0	2	9	0	0,5783259580,896050827	8,42E-05	0,002977773
222802_at	J05008				EG2	2	0	3	6	0,5783259580,896050827	8,42E-05	0,002977773
224839_s_at	BG328998	Hs.460693	glutamic pyruvate transaminase (alanine aminotransf	GPT2	EG2	0	2	6	3	0,5783259580,896050827	8,42E-05	0,002977773
224959_at	AI718385	Hs.302738	solute carrier family 26 (sulfate transporter), member	SLC26A2	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773
225411_at	AI769794	Hs.469690		FLJ14681	EG2	1	1	3	6	0,5783259580,896050827	8,42E-05	0,002977773
225571_at	AA701657	Hs.133421	leukemia inhibitory factor receptor	LIFR	EG2	0	2	2	7	0,5783259580,896050827	8,42E-05	0,002977773
226833_at	AI921877	Hs.27475		FLJ32499	EG2	2	0	7	2	0,5783259580,896050827	8,42E-05	0,002977773
227037_at	AA029500	Hs.31652		LOC201164	EG2	1	1	5	4	0,5783259580,896050827	8,42E-05	0,002977773
227070_at	W63754	Hs.159993	glycosyltransferase 8 domain containing 2	GLT8D2	EG2	0	2	5	4	0,5783259580,896050827	8,42E-05	0,002977773
227347_x_at	NM_021170	Hs.154029	hairy and enhancer of split 4 (Drosophila)	HES4	EG2	1	1	3	6	0,5783259580,896050827	8,42E-05	0,002977773
227657_at	AA722069	Hs.480825	ring finger protein 150	RNF150	EG2	0	2	1	8	0,5783259580,896050827	8,42E-05	0,002977773
227897_at	N20927	Hs.98643	RAP2B, member of RAS oncogene family	RAP2B	EG2	1	1	5	4	0,5783259580,896050827	8,42E-05	0,002977773
228946_at	AW150229	Hs.391481	PDZ domain containing 6	PDZK6	EG2	2	0	3	6	0,5783259580,896050827	8,42E-05	0,002977773
228998_at	AV693653	Hs.372082	trinucleotide repeat containing 6B	TNRC6B	EG2	0	2	2	7	0,5783259580,896050827	8,42E-05	0,002977773
229893_at	BF589413	Hs.127535	FERM domain containing 3	FRMD3	EG2	1	1	6	3	0,5783259580,896050827	8,42E-05	0,002977773
230425_at	AI674183	Hs.116092	EPH receptor B1	EPHB1	EG2	0	2	8	1	0,5783259580,896050827	8,42E-05	0,002977773
230559_x_at	AI277617	Hs.117835	FYVE, RhoGEF and PH domain containing 4	FGD4	EG2	0	2	0	9	0,5783259580,896050827	8,42E-05	0,002977773
230645_at	BF110588	Hs.127535	FERM domain containing 3	FRMD3	EG2	1	1	6	3	0,5783259580,896050827	8,42E-05	0,002977773
231904_at	AU122448	Hs.365116	U2(RNU2) small nuclear RNA auxiliary factor 1	U2AF1	EG2	1	1	9	0	0,5783259580,896050827	8,42E-05	0,002977773
235202_x_at	BG498328	Hs.252543		IKIP	EG2	2	0	7	2	0,5783259580,896050827	8,42E-05	0,002977773
235706_at	AW663908	Hs.334873	carboxypeptidase M	CPM	EG2	1	1	0	9	0,5783259580,896050827	8,42E-05	0,002977773
238841_at	AA651920	Hs.437943	protein tyrosine phosphatase domain containing 1	PTPDC1	EG2	1	1	8	1	0,5783259580,896050827	8,42E-05	0,002977773
239761_at	AI088120	Hs.521568	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6	GCNT1	EG2	2	0	5	4	0,5783259580,896050827	8,42E-05	0,002977773

242343_x_at	H57111	Hs.147895	zinc finger protein 518	ZNF518	EG2	0	2	8	1	0,5783259580,896050827	8,42E-05	0,002977773
1555812_a_at	AF498927	Hs.504877	Rho GDP dissociation inhibitor (GDI) beta	ARHGDI3	EG2	0	1	3	9	0,8651132690,896050827	7,43E-08	0,001562433
200743_s_at	NM_000391	Hs.523454	tripeptidyl peptidase I	TPP1	EG2	0	1	10	2	0,8651132690,896050827	7,43E-08	0,001562433
227529_s_at	BF511276	Hs.371240	A kinase (PRKA) anchor protein (gravin) 12	AKAP12	EG2	1	0	5	7	0,8651132690,896050827	7,43E-08	0,001562433
203571_s_at	NM_006829	Hs.74120	chromosome 10 open reading frame 116	C10orf116	EG2	0	1	5	6	0,8651132690,896050827	9,59E-07	0,001562433
204484_at	NM_002646	Hs.497487	phosphoinositide-3-kinase, class 2, beta polypeptide	PIK3C2B	EG2	0	1	4	7	0,8651132690,896050827	9,59E-07	0,001562433
227611_at	AA442856	Hs.6918	threonyl-tRNA synthetase-like 2	TARSL2	EG2	0	1	5	6	0,8651132690,896050827	9,59E-07	0,001562433
228128_x_at	AI110886	Hs.494928	pregnancy-associated plasma protein A, pappalysin 1	PAPPA	EG2	1	0	8	3	0,8651132690,896050827	9,59E-07	0,001562433
229638_at	AI681917	Hs.499205	iroquois homeobox protein 3	IRX3	EG2	1	0	4	7	0,8651132690,896050827	9,59E-07	0,001562433
201981_at	AA148534	Hs.494928	pregnancy-associated plasma protein A, pappalysin 1	PAPPA	EG2	1	0	8	2	0,8651132690,896050827	9,97E-06	0,001562433
202036_s_at	AF017987	Hs.213424	secreted frizzled-related protein 1	SFRP1	EG2	0	1	7	3	0,8651132690,896050827	9,97E-06	0,001562433
206172_at	NM_000640	Hs.336046	interleukin 13 receptor, alpha 2	IL13RA2	EG2	1	0	7	3	0,8651132690,896050827	9,97E-06	0,001562433
206371_at	NM_000804	Hs.352	folate receptor 3 (gamma)	FOLR3	EG2	0	1	7	3	0,8651132690,896050827	9,97E-06	0,001562433
209460_at	AF237813	Hs.336768	4-aminobutyrate aminotransferase	ABAT	EG2	1	0	6	4	0,8651132690,896050827	9,97E-06	0,001562433
210517_s_at	AB003476	Hs.371240	A kinase (PRKA) anchor protein (gravin) 12	AKAP12	EG2	1	0	4	6	0,8651132690,896050827	9,97E-06	0,001562433
217564_s_at	W80357	Hs.149252	carbonyl-phosphate synthetase 1, mitochondrial	CPS1	EG2	1	0	5	5	0,8651132690,896050827	9,97E-06	0,001562433
219441_s_at	NM_024652	Hs.513207	leucine-rich repeat kinase 1	LRRK1	EG2	0	1	0	10	0,8651132690,896050827	9,97E-06	0,001562433
222150_s_at	AK026747	Hs.186649, Hs.549342		LOC54103	EG2	1	0	5	5	0,8651132690,896050827	9,97E-06	0,001562433
224940_s_at	BF107618	Hs.494928	pregnancy-associated plasma protein A, pappalysin 1	PAPPA	EG2	1	0	7	3	0,8651132690,896050827	9,97E-06	0,001562433
224941_at	BF107618	Hs.494928	pregnancy-associated plasma protein A, pappalysin 1	PAPPA	EG2	1	0	7	3	0,8651132690,896050827	9,97E-06	0,001562433
227997_at	AW007080	Hs.150725	interleukin 17 receptor D	IL17RD	EG2	0	1	5	5	0,8651132690,896050827	9,97E-06	0,001562433
229441_at	AI569872	Hs.19545	frizzled homolog 4 (Drosophila)	FZD4	EG2	1	0	9	1	0,8651132690,896050827	9,97E-06	0,001562433
230831_at	AW294986	Hs.368399	FERM domain containing 5	FRMD5	EG2	1	0	7	3	0,8651132690,896050827	9,97E-06	0,001562433
235299_at	AI769269				EG2	0	1	7	3	0,8651132690,896050827	9,97E-06	0,001562433
239629_at	AI634046	Hs.390736	CASP8 and FADD-like apoptosis regulator	CFLAR	EG2	1	0	7	3	0,8651132690,896050827	9,97E-06	0,001562433
1552812_a_at	NM_014554	Hs.371957	SUMO1/sentrin specific peptidase 1	SENP1	EG2	1	0	5	4	0,8651132690,896050827	8,42E-05	0,002977773
1557275_a_at	BF726849				EG2	0	1	0	9	0,8651132690,896050827	8,42E-05	0,002977773
200706_s_at	NM_004862	Hs.459940	lipopolysaccharide-induced TNF factor	LITAF	EG2	0	1	5	4	0,8651132690,896050827	8,42E-05	0,002977773
201645_at	NM_002160	Hs.143250	tenascin C (hexabrachion)	TNC	EG2	1	0	8	1	0,8651132690,896050827	8,42E-05	0,002977773
201801_s_at	AF079117	Hs.25450	solute carrier family 29 (nucleoside transporters), me	SLC29A1	EG2	0	1	8	1	0,8651132690,896050827	8,42E-05	0,002977773
202741_at	AA130247	Hs.487325	protein kinase, cAMP-dependent, catalytic, beta	PRKACB	EG2	1	0	4	5	0,8651132690,896050827	8,42E-05	0,002977773
205542_at	NM_012449	Hs.61635	six transmembrane epithelial antigen of the prostate	STEAP1	EG2	0	1	7	2	0,8651132690,896050827	8,42E-05	0,002977773
205920_at	NM_003043	Hs.529488	solute carrier family 6 (neurotransmitter transporter, t	SLC6A6	EG2	1	0	7	2	0,8651132690,896050827	8,42E-05	0,002977773
206942_s_at	NM_002674	Hs.2182	pro-melanin-concentrating hormone	PMCH	EG2	0	1	2	7	0,8651132690,896050827	8,42E-05	0,002977773
209505_at	AI951185	Hs.519445	nuclear receptor subfamily 2, group F, member 1	NR2F1	EG2	1	0	3	6	0,8651132690,896050827	8,42E-05	0,002977773
211527_x_at	M27281	Hs.73793	vascular endothelial growth factor	VEGF	EG2	0	1	8	1	0,8651132690,896050827	8,42E-05	0,002977773
212382_at	BF433429	Hs.200285	transcription factor 4	TCF4	EG2	0	1	4	5	0,8651132690,896050827	8,42E-05	0,002977773
212521_s_at	BE568219	Hs.9333	phosphodiesterase 8A	PDE8A	EG2	0	1	3	6	0,8651132690,896050827	8,42E-05	0,002977773
213295_at	AA555096	Hs.432993	cylindromatosis (turban tumor syndrome)	CYLD	EG2	1	0	4	5	0,8651132690,896050827	8,42E-05	0,002977773
214023_x_at	AL533838	Hs.300701		TUBB-PARALOG	EG2	0	1	1	8	0,8651132690,896050827	8,42E-05	0,002977773
222668_at	W73820	Hs.221873	potassium channel tetramerisation domain containing	KCTD15	EG2	0	1	6	3	0,8651132690,896050827	8,42E-05	0,002977773
222717_at	BF982174	Hs.26530	serum deprivation response (phosphatidylserine bind	SDPR	EG2	1	0	5	4	0,8651132690,896050827	8,42E-05	0,002977773
224002_s_at	AF100751	Hs.410378	FK506 binding protein 7	FKBP7	EG2	0	1	7	2	0,8651132690,896050827	8,42E-05	0,002977773
224719_s_at	BG339653	Hs.405913		GRCC10	EG2	0	1	5	4	0,8651132690,896050827	8,42E-05	0,002977773
224817_at	W93554	Hs.549085	neuralized-like (Drosophila)	NEURL	EG2	1	0	4	5	0,8651132690,896050827	8,42E-05	0,002977773

224837_at	AW080845	Hs.431498	forkhead box P1	FOXP1	EG2	0	1	4	5	0,8651132690,896050827	8,42E-05	0,002977773
225653_at	AV755269	Hs.446350	transforming growth factor, beta receptor associated	TGFBRAP1	EG2	0	1	5	4	0,8651132690,896050827	8,42E-05	0,002977773
226719_at	AI868039				EG2	0	1	9	0	0,8651132690,896050827	8,42E-05	0,002977773
226771_at	AB032963	Hs.435700	ATPase, Class I, type 8B, member 2	ATP8B2	EG2	0	1	4	5	0,8651132690,896050827	8,42E-05	0,002977773
229830_at	AW205919	Hs.376032,	H platelet-derived growth factor alpha polypeptide	PDGFA	EG2	0	1	2	7	0,8651132690,896050827	8,42E-05	0,002977773
230748_at	AI873273	Hs.42645	solute carrier family 16 (monocarboxylic acid transpor	SLC16A6	EG2	0	1	7	2	0,8651132690,896050827	8,42E-05	0,002977773
231403_at	N21108	Hs.130031	triple functional domain (PTPRF interacting)	TRIO	EG2	0	1	6	3	0,8651132690,896050827	8,42E-05	0,002977773
235508_at	AW291023	Hs.526464	promyelocytic leukemia	PML	EG2	1	0	8	1	0,8651132690,896050827	8,42E-05	0,002977773
237169_at	AI266750	Hs.143250	tenascin C (hexabrachion)	TNC	EG2	1	0	9	0	0,8651132690,896050827	8,42E-05	0,002977773
242979_at	AI474666				EG2	1	0	3	6	0,8651132690,896050827	8,42E-05	0,002977773
217853_at	NM_022748	Hs.520814	tensin 3	TNS3	EG2	0	0	5	7	1 0,896050827	7,43E-08	0,001562433
204203_at	NM_001806	Hs.429666	CCAAT/enhancer binding protein (C/EBP), gamma	CEBPG	EG2	0	0	8	3	1 0,896050827	9,59E-07	0,001562433
209199_s_at	N22468	Hs.444409	MADS box transcription enhancer factor 2, polypeptid	MEF2C	EG2	0	0	8	3	1 0,896050827	9,59E-07	0,001562433
210041_s_at	BC001258	Hs.553496	phosphoglucosyltransferase 3	PGM3	EG2	0	0	6	4	1 0,896050827	9,97E-06	0,001562433
213664_at	AW235061	Hs.444915	solute carrier family 1 (neuronal/epithelial high affini	SLC1A1	EG2	0	0	7	3	1 0,896050827	9,97E-06	0,001562433
213895_at	BF445047	Hs.436298	epithelial membrane protein 1	EMP1	EG2	0	0	6	4	1 0,896050827	9,97E-06	0,001562433
224838_at	AK026898	Hs.431498	forkhead box P1	FOXP1	EG2	0	0	4	6	1 0,896050827	9,97E-06	0,001562433
225362_at	AI341165	Hs.404706		LOC159090	EG2	0	0	5	5	1 0,896050827	9,97E-06	0,001562433
228335_at	AW264204	Hs.31595	claudin 11 (oligodendrocyte transmembrane protein)	CLDN11	EG2	0	0	3	7	1 0,896050827	9,97E-06	0,001562433
39582_at	AL050166	Hs.432993	cylindromatosis (turban tumor syndrome)	CYLD	EG2	0	0	5	5	1 0,896050827	9,97E-06	0,001562433
1554795_a_at	BC019895	Hs.530101	filamin binding LIM protein 1	FBLIM1	EG2	0	0	1	8	1 0,896050827	8,42E-05	0,002977773
207265_s_at	NM_016657	Hs.554798	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protei	KDEL3	EG2	0	0	6	3	1 0,896050827	8,42E-05	0,002977773
210026_s_at	AY028896	Hs.57973	caspase recruitment domain family, member 10	CARD10	EG2	0	0	1	8	1 0,896050827	8,42E-05	0,002977773
210138_at	AF074979	Hs.368733	regulator of G-protein signalling 20	RGS20	EG2	0	0	7	2	1 0,896050827	8,42E-05	0,002977773
212816_s_at	BE613178	Hs.533013	cystathionine-beta-synthase	CBS	EG2	0	0	4	5	1 0,896050827	8,42E-05	0,002977773
213358_at	AB018345	Hs.464563	KIAA0802	KIAA0802	EG2	0	0	3	6	1 0,896050827	8,42E-05	0,002977773
220161_s_at	NM_019114	Hs.269180	erythrocyte membrane protein band 4.1 like 4B	EPB41L4B	EG2	0	0	1	8	1 0,896050827	8,42E-05	0,002977773
224204_x_at	AF231339	Hs.434269	aryl hydrocarbon receptor nuclear translocator-like 2	ARNTL2	EG2	0	0	2	7	1 0,896050827	8,42E-05	0,002977773
230121_at	BF508679				EG2	0	0	2	7	1 0,896050827	8,42E-05	0,002977773
237466_s_at	AW444502	Hs.507991	hedgehog interacting protein	HHIP	EG2	0	0	0	9	1 0,896050827	8,42E-05	0,002977773
241387_at	AW276701	Hs.395482	PTK2 protein tyrosine kinase 2	PTK2	EG2	0	0	3	6	1 0,896050827	8,42E-05	0,002977773