

Table H: D0<D2>D4>D6

Gene Title	Gene Identifier	D2/D0	D4/D0	D6/D0	P
Ankyrin repeat and IBR domain containing 1, mRNA (cDNA clone MGC:179082 IMAGE:9054074)	AK014523	1.05	-2.04	-2.17	0.004
Ankyrin repeat domain 44, mRNA (cDNA clone MGC:198807 IMAGE:9054388)	AI315673	1.53	1.29	-2.13	0.009
Annexin A13 (Anxa13), mRNA	BC013521	1.16	-1.05	-2.5	0.001
ATPase, Ca ⁺⁺ transporting, type 2C, member 2, mRNA (cDNA clone IMAGE:4984604)	AK007419	1.16	-1.09	-2.08	0.001
CDNA clone IMAGE:9053535	BB335888	1.36	1.02	-4.17	0.007
CDNA fis, clone TRACH3029665, weakly similar to Zinc finger protein 84	AI852969	1.03	-1.3	-2.38	0.004
Chemokine (C-C motif) receptor-like 1 (Ccr11), mRNA	AF306532	1.2	1.1	-2.04	0.006
Clone 1.5A LB9 mRNA, 3' UTR	U20265	4.57	1.8	1.1	0.006
Cytochrome P450, family 2, subfamily c, polypeptide 68, mRNA (cDNA clone MGC:28459 IMAGE:4160492)	AI265721	1.2	1.13	-4.76	0.001
D site albumin promoter binding protein, mRNA (cDNA clone MGC:18803 IMAGE:4195116)	BC018323	3.47	2.3	-1.49	0.001
DNA segment, Chr 2, ERATO Doi 239, expressed	C79214	2.76	1.96	-1.35	0.006
EST	AV034993	3.36	2.34	2.09	0.006
ESTs	BB442009	6.9	5.87	3.12	0.007
ESTs	BB732233	2.47	2.11	-1.09	0.001
ESTs	BB481761	3.45	1.18	-1.67	0.008
ESTs, Moderately similar to GNMSLL retrovirus-related reverse transcriptase homolog - mouse retrotransposon (M.musculus)	BI076611	1.4	-1.3	-2.78	0.002
ESTs, Weakly similar to TYROSINE-PROTEIN KINASE JAK3 (M.musculus)	BB796950	1.16	-1.3	-2.17	0.001
Golgi phosphoprotein 3-like (Golp31), mRNA	BC027194	1.05	-1.37	-2	0.001
Guanylate cyclase 2g (Gucy2g), mRNA	BB427699	2.01	1.73	-2.94	0.001

Hexaribonucleotide binding protein 3 (Hrnbp3)	BB386621	4.09	3.05	2.85	0.001
Hydroxyacid oxidase (glycolate oxidase) 3, mRNA (cDNA clone MGC:29258 IMAGE:5064100)	NM_019545	1.01	-1.05	-2.44	0.001
Indian hedgehog (Ihh), mRNA	NM_010544	1.07	-1.45	-2.22	0.001
Kallikrein 1-related peptidase b9 (Klk1b9), mRNA	M17962	1.03	-1.3	-3.85	0.001
Mouse, glutathione transferase GT9.3 mRNA, 3' end	J03953	1.11	-1.54	-2.33	0.001
Mu opioid receptor variant A	AF260307	2.78	1.81	-1.14	0.008
Myelin protein zero-like 2, mRNA (cDNA clone MGC:13828 IMAGE:4188907)	BC015076	1	-1.54	-2.08	0.001
Nephroblastoma overexpressed gene, mRNA (cDNA clone MGC:5979 IMAGE:3494896)	X96585	1.1	-1.37	-2.44	0.001
Nuclear receptor subfamily 1, group I, member 3, mRNA (cDNA clone IMAGE:5052215)	AF009328	1.53	-1.09	-3.57	0.005
Phosphatidylserine synthase 2, mRNA (cDNA clone IMAGE:3583796)	AI596401	1.31	-4.35	-4.35	0.001
Poliovirus receptor-related 3 (Pvrl3), transcript variant gamma, mRNA	NM_021496	1.09	-1.92	-2.33	0.008
Potassium channel, subfamily K, member 5 (Kcnk5), mRNA	AF319542	1	-1.22	-2.22	0.001
PR domain containing 14 (Prdm14), mRNA	BB283625	1.32	1.21	-2.63	0.005
PREDICTED: Mus musculus predicted gene, EG665934 (EG665934), mRNA	AV378858	2.53	2.43	-1.75	0.004
RASD family, member 2, mRNA (cDNA clone IMAGE:4988981)	BC026377	1.14	-1.52	-3.13	0.001
RIKEN cDNA 2210020M01 gene, mRNA (cDNA clone IMAGE:1179778)	BE852759	1.06	-1.03	-2.17	0.002
RIKEN cDNA 2310044G17 gene, mRNA (cDNA clone MGC:31149 IMAGE:4168224)	AK009800	1.13	-1.43	-2.27	0.001
RIKEN cDNA 9130414P19 gene	AK018677	1.13	1.02	-4.35	0.001
Solute carrier family 20, member 1, mRNA (cDNA clone MGC:14029 IMAGE:4015575)	NM_015747	1.1	-1.16	-3.03	0.001

sphingosine kinase 2	BB550183	3.81	2.73	-1.25	0.001
sulfotransferase family 1A, phenol-preferring, member 1	AK002700	1.02	1.01	-2.04	0.001
T-box 4 (Tbx4)	AY075134	3.72	1.28	-1.67	0.003
Transcribed locus	BB097972	1	-1.01	-2.5	0.007
Transcribed locus	AV356118	1.05	-1.54	-2.04	0.001
Transcribed locus	AK016405	4.43	1.9	1.68	0.007
Transcribed locus	BB667588	4.45	3.39	1.17	0.004
Transcribed locus	BB283300	1.09	-1.14	-5.26	0.001
Transcribed locus	BB465699	1.05	-2	-3.33	0.001
Transcribed locus	BM120390	2.48	1.71	-3.13	0.009
Transcribed locus	BB440150	2.49	1.74	-1.89	0.008
Transcribed locus, strongly similar to NP_037393.1 peroxisome proliferator-activated receptor gamma, coactivator 1 alpha [Hom	BM120569	1.03	-1.05	-2.5	0.001
Transmembrane protease, serine 2, mRNA (cDNA clone MGC:6821 IMAGE:2648878)	BM250339	1.26	1	-2.44	0.003
Tubulin tyrosine ligase-like family, member 7, mRNA (cDNA clone MGC:169898 IMAGE:8861293)	AK014905	1.3	-1.04	-2.63	0.009
Uroporphyrinogen III synthase (Uros), mRNA	AI415298	2.61	1.8	-1.05	0.002
Zinc finger protein 365, mRNA (cDNA clone MGC:189953 IMAGE:9088140)	BB277790	1.05	-3.03	-9.09	0.001
Zonadhesin (Zan), mRNA	NM_011741	1	-1.01	-4.76	0.001

(-) represents fold down regulation from day 0 control.