

Supplementary Table 1. Proteins differentially expressed between right and left hemisphere following cilostazol treatment (p-value ≤ 0.05)

Accession	Description	A8: 115/113	B8: 115/113	A8: 116/113	B8: 116/113	A8: 117/113	B8: 117/113	A8: 118/114	B8: 118/114	A8: 119/114	B8: 119/114	A8: 121/114	B8: 121/114	T-Test
Q9ET80	Junctophilin-1 OS=Mus musculus GN=Jph1 PE=1 SV=1 - [JPH1_MOUSE]	0.10	0.53	0.44	0.50	0.44	0.51	-0.24	-0.16	-0.08	-0.45	-0.37	-0.26	0.00
O88428	Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthase 2 OS=Mus musculus GN=Paps2 PE=1 SV=2 - [PAPS2_MOUSE]	-0.36	-0.21	-0.33	-0.14	-0.36	-0.03	0.31	0.09	0.34	0.30	0.22	0.17	0.00
Q3UIL6-6	Isoform 6 of Pleckstrin homology domain-containing family A member 7 OS=Mus musculus GN=Plekha7 - [PKHA7_MOUSE]	0.55	0.72	0.26	0.54	0.47	0.62	0.03	0.12	0.07	-0.24	0.01	-0.02	0.00
Q8C3X8	Lipase maturation factor 2 OS=Mus musculus GN=Lmf2 PE=2 SV=1 - [LMF2_MOUSE]	-1.27	-0.67	-0.84	-0.50	-0.61	-0.36	0.51	0.04	0.73	0.66	0.10	0.72	0.00
P14231	Sodium/potassium-transporting ATPase subunit beta-2 OS=Mus musculus GN=Atp1b2 PE=1 SV=2 - [AT1B2_MOUSE]	-0.28	-0.28	-0.07	-0.18	-0.18	-0.23	0.16	0.01	0.17	0.14	0.12	-0.03	0.00
Q5I012-4	Isoform 4 of Putative sodium-coupled neutral amino acid transporter 10 OS=Mus musculus GN=Slc38a10 - [S38AA_MOUSE]	0.08	-0.06	0.09	-0.04	0.18	0.14	0.84	1.15	0.89	0.67	0.80	1.29	0.00
Q7TSG1-2	Isoform 2 of Centrosomal protein of 120 kDa OS=Mus musculus GN=Cep120 - [CE120_MOUSE]	0.38	1.54	1.01	1.49	0.59	1.41	-0.40	-0.21	-0.10	-0.77	-1.09	-0.49	0.00
E9Q3C1	Protein C2cd2 OS=Mus musculus GN=C2cd2 PE=2 SV=1 - [E9Q3C1_MOUSE]	1.40	0.83	0.89	0.63	0.38	0.33	-0.70	-0.54	-0.75	-0.94	0.04	-0.39	0.00
Q6NZN0-5	Isoform 5 of RNA-binding protein 26 OS=Mus musculus GN=Rbm26 - [RBM26_MOUSE]	0.06	0.39	0.08	0.26	0.15	0.35	-0.25	-0.04	-0.19	-0.41	-0.21	-0.29	0.00
Q9D2P4	Ubiquitin-related modifier 1 homolog OS=Mus musculus GN=Urm1 PE=1 SV=1 - [URM1_MOUSE]	-0.20	-0.37	-0.45	-0.51	0.00	-0.26	0.41	0.55	0.49	0.46	-0.01	0.68	0.00
Q99KG3-3	Isoform 3 of RNA-binding protein 10 OS=Mus musculus GN=Rbm10 - [RBM10_MOUSE]	0.14	0.10	0.02	0.05	0.10	0.06	-0.28	-0.05	-0.24	-0.17	-0.13	-0.20	0.00
Q9Z1L5	Voltage-dependent calcium channel subunit alpha-2/delta-3 OS=Mus musculus GN=Cacna2d3 PE=1 SV=1 - [CA2D3_MOUSE]	0.00	-0.02	-0.07	-0.15	0.01	-0.06	0.18	0.21	0.04	0.11	0.19	0.20	0.00

P51910	Apolipoprotein D OS=Mus musculus GN=Apod PE=2 SV=1 - [APOD_MOUSE]	-0.26	-0.08	-0.27	-0.03	-0.14	-0.06	0.06	0.35	0.26	0.35	0.16	0.48	0.00
P00397	Cytochrome c oxidase subunit 1 OS=Mus musculus GN=Mtco1 PE=3 SV=1 - [COX1_MOUSE]	-0.44	-0.61	-0.32	-0.27	-0.35	-0.15	0.51	0.19	0.08	0.02	0.06	0.08	0.00
D3YXK0	MCG4624, isoform CRA_c OS=Mus musculus GN=Jakmp2 PE=4 SV=2 - [D3YXK0_MOUSE]	-0.12	0.21	0.23	-0.01	0.12	0.00	-0.28	-0.13	-0.45	-0.29	-0.37	-0.52	0.00
B2RY56	RNA-binding protein 25 OS=Mus musculus GN=Rbm25 PE=1 SV=2 - [RBM25_MOUSE]	-0.04	0.07	0.14	-0.01	0.03	0.01	-0.11	-0.30	-0.09	-0.20	-0.19	-0.21	0.00
Q9ESN4	Complement C1q-like protein 3 OS=Mus musculus GN=C1q3 PE=1 SV=1 - [C1QL3_MOUSE]	-0.27	-0.31	0.07	-0.41	-0.11	-0.15	0.13	0.41	0.45	0.52	0.24	0.11	0.00
Q91V14	Solute carrier family 12 member 5 OS=Mus musculus GN=S1c12a5 PE=1 SV=2 - [S12A5_MOUSE]	-0.19	-0.15	-0.07	-0.16	-0.13	-0.14	0.03	0.00	-0.02	0.05	-0.04	-0.08	0.00
A2A6T1	Cerebellar degeneration-related protein 2-like OS=Mus musculus GN=Cdr2l PE=2 SV=1 - [CDR2L_MOUSE]	0.28	0.13	0.33	0.31	0.18	0.19	0.11	-0.02	-0.11	0.02	0.04	-0.13	0.00
Q3UJH1	TNF receptor- associated factor 3 OS=Mus musculus GN=Traf3 PE=2 SV=1 - [Q3UJH1_MOUSE]	-0.02	-0.17	-0.11	-0.22	-0.02	-0.21	0.10	0.07	0.15	0.20	0.02	0.14	0.00
Q7TQG1	Pleckstrin homology domain-containing family A member 6 OS=Mus musculus GN=Plekha6 PE=1 SV=1 - [PKHA6_MOUSE]	-0.06	0.23	0.06	0.03	0.08	0.01	-0.31	-0.27	-0.11	-0.22	-0.09	-0.28	0.00
Q6NXK7	Inactive dipeptidyl peptidase 10 OS=Mus musculus GN=Dpp10 PE=2 SV=1 - [DPP10_MOUSE]	-0.12	-0.12	-0.10	0.01	-0.05	-0.12	0.02	0.02	0.10	0.01	0.06	0.11	0.00
Q8R395	COMM domain- containing protein 5 OS=Mus musculus GN=Comm5 PE=2 SV=1 - [COMD5_MOUSE]	-0.09	-0.19	-0.18	-0.14	-0.29	-0.17	0.25	0.13	0.15	0.35	-0.04	0.15	0.00
P51175	Protoporphyrinogen oxidase OS=Mus musculus GN=Ppox PE=2 SV=1 - [PPOX_MOUSE]	0.22	0.32	0.06	0.30	0.22	0.50	-0.06	-0.13	-0.06	-0.44	-0.02	-0.18	0.00
B1AXV0	DOMON domain- containing protein FRRS1L OS=Mus musculus GN=Frs1l PE=1 SV=1 - [FRS1L_MOUSE]	0.27	0.27	0.13	0.29	0.11	0.27	0.04	-0.17	-0.05	-0.12	0.05	-0.27	0.00

Q99JG2	Endothelin B receptor-like protein 2 OS=Mus musculus GN=Gpr3711 PE=1 SV=2 - [ETBR2_MOUSE]	-0.60	-0.63	-0.47	-0.41	-0.24	-0.66	-0.30	0.10	0.32	0.12	0.10	-0.12	0.00
Q9CQD1	Ras-related protein Rab-5A OS=Mus musculus GN=Rab5a PE=1 SV=1 - [RAB5A_MOUSE]	-0.20	-0.45	-0.33	-0.26	-0.06	-0.47	0.17	0.24	-0.13	0.23	0.03	0.16	0.00
P35831	Tyrosine-protein phosphatase non-receptor type 12 OS=Mus musculus GN=Ptpn12 PE=1 SV=3 - [PTN12_MOUSE]	0.13	0.32	0.12	0.42	0.03	0.31	-0.38	-0.24	-0.12	0.02	-0.17	-0.41	0.00
D3Z1T2	MCG15614 OS=Mus musculus GN=2310009B15Rik PE=4 SV=1 - [D3Z1T2_MOUSE]	0.91	0.52	0.96	0.43	1.39	0.66	-0.40	-0.23	0.34	0.13	-0.03	-0.31	0.00
P35283	Ras-related protein Rab-12 OS=Mus musculus GN=Rab12 PE=1 SV=3 - [RAB12_MOUSE]	-0.20	-0.17	-0.06	-0.32	-0.06	-0.10	0.16	0.17	0.05	-0.02	0.06	0.21	0.00
E9Q5D5	WW domain-containing adapter protein with coiled-coil OS=Mus musculus GN=Wac PE=2 SV=1 - [E9Q5D5_MOUSE]	0.07	0.14	0.33	0.06	0.10	0.35	0.02	-0.16	-0.16	-0.09	-0.26	-0.19	0.00
P98200	Probable phospholipid-transporting ATPase IB OS=Mus musculus GN=Atp8a2 PE=1 SV=1 - [AT8A2_MOUSE]	-0.15	0.51	0.03	0.17	-0.06	0.15	-0.38	-0.50	-0.22	-0.57	-0.64	-0.37	0.00
Q6P9Q6	FK506-binding protein 15 OS=Mus musculus GN=Fkbp15 PE=1 SV=2 - [FKB15_MOUSE]	0.03	0.00	0.07	-0.01	-0.03	0.10	-0.13	-0.12	-0.14	-0.02	-0.13	-0.20	0.00
P35922-9	Isoform ISO9 of Fragile X mental retardation protein 1 homolog OS=Mus musculus GN=Fmr1 - [FMR1_MOUSE]	0.16	0.12	0.10	0.19	0.06	0.23	-0.04	-0.21	-0.17	-0.16	-0.21	-0.48	0.00
Q8C5W0-4	Isoform 4 of Calmin OS=Mus musculus GN=Clnn - [CLMN_MOUSE]	0.23	0.13	0.21	0.07	0.10	0.10	-0.09	-0.07	-0.20	0.03	-0.08	0.02	0.00
Q3USH1	Protein FAM196A OS=Mus musculus GN=Fam196a PE=2 SV=1 - [F196A_MOUSE]	1.10	1.37	0.52	1.47	0.37	1.63	-0.45	0.38	-0.51	-0.45	-0.24	0.28	0.00
F7B0R9	Protein 2010300C02Rik (Fragment) OS=Mus musculus GN=2010300C02Rik PE=4 SV=1 - [F7B0R9_MOUSE]	0.27	-0.09	0.24	0.52	0.29	-0.14	-0.78	-0.83	-0.33	-0.92	-0.20	-0.32	0.00
Q9EQQ2	Protein YIPF5 OS=Mus musculus GN=Yipf5 PE=2 SV=1 - [YIPF5_MOUSE]	-0.50	-0.49	-0.23	-0.24	-0.36	-0.28	0.17	0.02	-0.05	-0.03	-0.25	0.07	0.00

O35136	Neural cell adhesion molecule 2 OS=Mus musculus GN=Ncam2 PE=1 SV=1 - [NCAM2_MOUSE]	-0.07	-0.07	-0.07	-0.05	-0.12	-0.18	-0.01	0.08	0.15	0.00	0.02	0.11	0.00
Q8BYH8-2	Isoform 2 of Chromodomain-helicase-DNA-binding protein 9 OS=Mus musculus GN=Chd9 - [CHD9_MOUSE]	0.62	0.22	0.70	0.03	0.20	-0.14	-1.06	-0.95	-0.29	-0.38	-0.21	-0.75	0.00
Q99JP7	Gamma-glutamyltransferase 7 OS=Mus musculus GN=Ggt7 PE=1 SV=2 - [GGT7_MOUSE]	-0.09	-0.12	-0.16	0.03	-0.21	-0.23	0.13	-0.01	0.16	0.09	0.06	0.06	0.00
Q62277	Synaptophysin OS=Mus musculus GN=Syp PE=1 SV=2 - [SYPH_MOUSE]	-0.06	-0.05	-0.09	-0.04	-0.10	-0.02	0.09	0.07	0.10	-0.04	0.05	0.12	0.00
Q8BQ30	Phostensin OS=Mus musculus GN=Ppp1r18 PE=1 SV=1 - [PPR18_MOUSE]	0.02	0.33	0.19	0.33	0.25	0.26	-0.17	-0.04	0.09	-0.03	-0.06	-0.29	0.00
Q7TMR0	Lysosomal Pro-X carboxypeptidase OS=Mus musculus GN=Prpc PE=2 SV=2 - [PCP_MOUSE]	0.00	-0.42	-0.32	-0.54	-0.49	-0.39	0.51	-0.17	0.04	0.16	0.36	0.24	0.00
A2AWR5	Protein Gm14085 OS=Mus musculus GN=Gm14085 PE=4 SV=1 - [A2AWR5_MOUSE]	0.49	-0.56	0.31	-0.52	-0.07	-0.91	2.35	0.39	2.06	1.01	1.66	1.15	0.00
Q80V42	Carboxypeptidase M OS=Mus musculus GN=Cpm PE=2 SV=2 - [CBPM_MOUSE]	-0.06	-0.17	0.06	-0.10	-0.15	-0.04	0.08	0.08	0.14	0.02	0.14	0.18	0.00
P33174	Chromosome-associated kinesin KIF4 OS=Mus musculus GN=Kif4 PE=1 SV=3 - [KIF4_MOUSE]	0.00	0.75	-0.15	0.06	0.19	0.27	-1.00	-0.89	-0.10	-0.66	-0.37	-0.69	0.00
P97350	Plakophilin-1 OS=Mus musculus GN=Pkp1 PE=1 SV=1 - [PKP1_MOUSE]	0.33	0.77	0.39	0.33	0.61	0.49	-0.03	-0.43	0.03	-0.57	0.00	-0.83	0.00
Q6R5P0	Toll-like receptor 11 OS=Mus musculus GN=Tlr11 PE=1 SV=1 - [TLR11_MOUSE]	0.16	0.01	0.34	0.33	0.49	-0.11	1.22	0.49	1.44	1.33	1.16	0.66	0.00
Q8K2F0	Bromodomain-containing protein 3 OS=Mus musculus GN=Brd3 PE=1 SV=2 - [BRD3_MOUSE]	-0.03	0.15	0.15	0.29	0.40	0.36	-0.18	-0.71	-0.12	-0.58	-0.09	-0.56	0.00
P59481	VIP36-like protein OS=Mus musculus GN=Lman2l PE=2 SV=1 - [LMA2L_MOUSE]	0.25	0.39	0.02	-0.07	0.06	0.10	-0.77	-0.60	-0.16	-0.12	-0.61	-0.74	0.00
E9QP09	Cullin-9 OS=Mus musculus GN=Cul9 PE=3 SV=1 - [E9QP09_MOUSE]	0.87	0.25	0.36	0.11	0.23	0.21	-0.43	-0.44	0.04	-0.43	-0.04	-0.33	0.00
A3KGB4	TBC1 domain family member 8B OS=Mus musculus GN=Tbc1d8b PE=2 SV=1 - [TBC8B_MOUSE]	1.64	0.41	1.02	0.43	0.56	0.13	-0.59	-0.29	-1.20	0.03	-1.01	-0.26	0.00

Q9CY73	39S ribosomal protein L44, mitochondrial OS=Mus musculus GN=Mrlp44 PE=1 SV=3 - [RM44_MOUSE]	1.12	0.97	1.19	0.08	1.46	0.53	-0.24	-0.31	-0.47	-0.19	-0.31	-0.33	0.00
F6ZKC7	Cleavage stimulation factor subunit 2 (Fragment) OS=Mus musculus GN=Cstf2 PE=4 SV=1 - [F6ZKC7_MOUSE]	1.05	0.39	0.23	0.69	0.34	0.29	-0.10	-0.81	0.01	-0.47	0.00	-0.74	0.00
O35409	Glutamate carboxypeptidase 2 OS=Mus musculus GN=Folh1 PE=2 SV=2 - [FOLH1_MOUSE]	-0.23	-0.24	-0.16	-0.26	-0.05	-0.28	-0.06	-0.06	0.17	0.02	0.20	0.00	0.00
Q3TC93	HCLS1-binding protein 3 OS=Mus musculus GN=Hs1bp3 PE=1 SV=2 - [H1BP3_MOUSE]	-0.10	0.30	0.15	0.01	0.25	0.16	-0.43	-0.42	0.03	-0.22	-0.23	-0.32	0.00
A2AL79	Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1 - [A2AL79_MOUSE]	-0.49	-0.18	0.02	-0.42	-0.30	-0.15	0.45	-0.02	0.14	0.34	0.21	0.02	0.00
P35802	Neuronal membrane glycoprotein M6-a OS=Mus musculus GN=Gpm6a PE=1 SV=1 - [GPM6A_MOUSE]	-0.34	-0.17	-0.13	-0.08	-0.23	-0.13	-0.10	-0.02	0.03	0.06	0.13	0.12	0.00
Q9D0V7	Receptor-binding cancer antigen expressed on SiSo cells OS=Mus musculus GN=Ebag9 PE=1 SV=2 - [RCAS1_MOUSE]	0.79	0.51	0.18	0.25	0.59	0.34	0.03	-0.11	0.09	-0.30	-0.08	0.09	0.00
P62827	GTP-binding nuclear protein Ran OS=Mus musculus GN=Ran PE=1 SV=3 - [RAN_MOUSE]	0.73	0.58	0.46	0.50	0.40	0.30	0.38	0.13	-0.05	-0.13	0.12	0.16	0.00
Q69ZS6	Synaptic vesicle glycoprotein 2C OS=Mus musculus GN=Sv2c PE=1 SV=2 - [SV2C_MOUSE]	-0.23	-0.37	-0.03	-0.34	0.03	-0.47	0.09	0.10	0.16	0.28	0.08	0.35	0.00
P60904	DnaJ homolog subfamily C member 5 OS=Mus musculus GN=Dnajc5 PE=1 SV=1 - [DNJC5_MOUSE]	0.00	-0.21	-0.08	-0.09	-0.23	-0.24	-0.01	0.24	0.22	-0.04	0.09	0.17	0.00
Q9R0L6	Pericentriolar material 1 protein OS=Mus musculus GN=Pcm1 PE=1 SV=2 - [PCM1_MOUSE]	0.16	-0.04	0.03	0.07	0.09	0.10	-0.25	-0.41	-0.04	-0.22	-0.09	-0.25	0.00
Q6PIE5	Sodium/potassium-transporting ATPase subunit alpha-2 OS=Mus musculus GN=Atp1a2 PE=1 SV=1 - [AT1A2_MOUSE]	-0.15	-0.17	0.03	-0.05	-0.15	-0.17	0.13	0.03	0.14	0.01	0.00	0.08	0.00
B1AXN3	MCG56945 OS=Mus musculus GN=Gm382 PE=4 SV=1 - [B1AXN3_MOUSE]	-0.19	0.34	0.17	0.13	0.13	0.09	-0.10	-0.26	-0.47	-0.10	-0.37	-0.29	0.00

O08581	Potassium channel subfamily K member 1 OS=Mus musculus GN=Kcnk1 PE=1 SV=2 - [KCNK1_MOUSE]	-0.12	-1.46	-0.33	-0.64	-1.40	-0.23	0.34	0.74	0.53	1.70	-0.07	0.97	0.00
B2KF50	Protein Uhrf1bp1 OS=Mus musculus GN=Uhrf1bp1 PE=2 SV=1 - [B2KF50_MOUSE]	-0.18	-0.04	0.05	-0.04	-0.01	-0.15	0.16	0.17	0.02	0.34	0.11	0.16	0.00
B2RUS7	ATP-binding cassette, sub-family C (CFTR/MRP), member 8 OS=Mus musculus GN=Abcc8 PE=2 SV=1 - [B2RUS7_MOUSE]	-0.02	-0.14	-0.08	-0.03	-0.22	-0.22	0.03	0.08	0.04	0.16	-0.03	0.12	0.00
E9PX70	Collagen alpha-1(XII) chain OS=Mus musculus GN=Col12a1 PE=2 SV=1 - [E9PX70_MOUSE]	0.44	0.03	0.31	0.13	0.09	-0.15	0.03	-0.33	-0.44	-0.50	-0.25	-0.38	0.00
A1A535	Ventricular zone-expressed PH domain-containing protein 1 OS=Mus musculus GN=Veph1 PE=2 SV=2 - [MELT_MOUSE]	-0.66	-0.90	-0.76	-0.55	-0.94	-0.72	-0.08	-0.12	-0.43	0.30	-0.54	-0.09	0.00
P35278	Ras-related protein Rab-5C OS=Mus musculus GN=Rab5c PE=1 SV=2 - [RAB5C_MOUSE]	-0.09	-0.05	-0.12	-0.15	0.03	-0.21	0.02	0.14	0.23	0.03	0.21	0.03	0.00
P49817	Caveolin-1 OS=Mus musculus GN=Cav1 PE=1 SV=1 - [CAV1_MOUSE]	-0.49	-0.18	-0.16	-0.04	-0.13	-0.33	0.08	-0.04	0.17	0.19	0.08	0.12	0.00
O55100-2	Isoform 1B of Synaptogyrin-1 OS=Mus musculus GN=Syng1 - [SNG1_MOUSE]	-0.26	-0.25	-0.07	-0.11	-0.38	-0.25	0.11	0.01	-0.08	0.09	-0.05	-0.03	0.00
Q5FWH7-2	Isoform 2 of Zinc transporter ZIP12 OS=Mus musculus GN=Slc39a12 - [S39AC_MOUSE]	-0.22	-0.34	-0.18	-0.11	-0.13	-0.31	0.06	-0.15	0.18	0.03	0.25	0.04	0.00
E9QAQ7	AT-rich interactive domain-containing protein 1A OS=Mus musculus GN=Arid1a PE=2 SV=1 - [E9QAQ7_MOUSE]	0.18	0.11	0.14	0.06	0.05	0.17	-0.35	-0.50	-0.01	-0.51	-0.17	-0.23	0.00
Q64337	Sequestosome-1 OS=Mus musculus GN=Sqstm1 PE=1 SV=1 - [SQSTM_MOUSE]	0.48	0.01	-0.04	-0.23	-0.11	0.35	-0.50	-0.58	-0.42	-0.65	-0.28	-0.47	0.00
Q61885	Myelin-oligodendrocyte glycoprotein OS=Mus musculus GN=Mog PE=1 SV=1 - [MOG_MOUSE]	-0.06	-0.07	0.02	0.14	-0.04	-0.16	0.09	0.31	0.12	0.22	0.13	0.27	0.00
Q3UYC0	Protein phosphatase 1H OS=Mus musculus GN=Ppm1h PE=1 SV=1 - [PPM1H_MOUSE]	0.24	0.24	0.23	0.21	0.05	0.15	0.07	0.12	-0.14	-0.01	-0.12	-0.06	0.00
Q8VDD8	WAS protein family homolog 1 OS=Mus musculus GN=Wash1 PE=2 SV=1 - [WASH1_MOUSE]	0.01	0.07	0.22	0.25	0.03	0.36	-0.10	-0.20	-0.06	0.03	-0.24	-0.21	0.00

Q61771	Kinesin-like protein KIF3B OS=Mus musculus GN=Kif3b PE=1 SV=1 - [KIF3B_MOUSE]	-0.03	0.16	0.17	-0.02	0.02	0.09	0.05	-0.32	-0.24	-0.22	-0.16	-0.21	0.00
Q8R238	Serine dehydratase- like OS=Mus musculus GN=Sdsl PE=2 SV=1 - [SDSL_MOUSE]	-0.50	-0.03	-0.41	-0.24	-0.34	-0.28	0.12	-0.10	0.48	-0.09	0.29	0.33	0.00
P84228	Histone H3.2 OS=Mus musculus GN=Hist1h3b PE=1 SV=2 - [H32_MOUSE]	-0.96	-0.54	-1.06	-0.44	-0.03	-0.38	-0.19	0.29	0.06	0.29	0.84	0.16	0.00
Q8C8R3-7	Isoform 7 of Ankyrin-2 OS=Mus musculus GN=Ank2 - [ANK2_MOUSE]	-0.23	-0.33	0.07	-0.38	-0.42	-0.71	0.46	0.15	0.05	0.41	-0.08	0.13	0.00
P03958	Adenosine deaminase OS=Mus musculus GN=Ada PE=1 SV=3 - [ADA_MOUSE]	-0.13	0.09	-0.31	-0.10	-0.23	0.03	0.17	0.03	0.27	0.31	0.27	0.10	0.00
O55074	A-kinase anchor protein 7 isoform alpha OS=Mus musculus GN=Akap7 PE=1 SV=4 - [AKA7A_MOUSE]	-0.48	-0.10	-0.25	-0.01	-0.19	-0.36	0.01	0.03	0.33	0.43	0.09	0.04	0.00
D3Z2W1	Protein Dgki OS=Mus musculus GN=Dgki PE=2 SV=1 - [D3Z2W1_MOUSE]	0.03	-0.04	0.08	0.09	0.03	0.08	-0.09	-0.25	-0.12	-0.10	-0.02	-0.25	0.00
E9QPM7	Guanine nucleotide exchange factor DBS OS=Mus musculus GN=Mcf2l PE=2 SV=1 - [E9QPM7_MOUSE]	0.61	0.38	0.08	0.35	0.14	0.24	-0.18	-0.64	-0.09	-0.61	0.13	-0.33	0.00
Q9CZ70	MCG129801, isoform CRA_a OS=Mus musculus GN=311000122Rik PE=2 SV=1 - [Q9CZ70_MOUSE]	-0.37	-0.18	-0.39	-1.11	-0.12	-0.94	-0.40	0.64	0.36	0.53	0.43	0.55	0.00
E0CXD4	Protein Pcdh1 OS=Mus musculus GN=Pcdh1 PE=2 SV=1 - [E0CXD4_MOUSE]	0.14	0.15	0.16	0.09	0.05	0.07	-0.34	-0.08	-0.07	-0.06	-0.15	-0.01	0.00
Q6UY53	HN1-like protein OS=Mus musculus GN=AY358078 PE=2 SV=1 - [Q6UY53_MOUSE]	0.52	-0.64	-1.18	-0.35	-0.05	-1.11	0.74	0.32	0.75	0.89	1.08	1.00	0.00
Q05CL8	La-related protein 7 OS=Mus musculus GN=Larp7 PE=1 SV=2 - [LARP7_MOUSE]	0.17	0.30	0.10	0.24	0.34	0.47	-0.10	-0.06	0.09	-0.03	0.18	-0.07	0.00
F6ZGI7	Molybdopterin synthase catalytic subunit OS=Mus musculus GN=Mocs2 PE=4 SV=1 - [F6ZGI7_MOUSE]	0.27	0.00	-0.04	0.03	0.24	-0.08	-0.38	-0.25	-0.17	-0.30	-0.03	-0.25	0.00
P97300	Neuroplastin OS=Mus musculus GN=Nptn PE=1 SV=3 - [NPTN_MOUSE]	-0.09	-0.07	-0.12	-0.05	-0.06	-0.05	-0.02	0.14	0.02	0.07	-0.01	0.08	0.00
Q62469	Integrin alpha-2 OS=Mus musculus GN=Itga2 PE=1 SV=2 - [ITA2_MOUSE]	-0.70	0.00	-0.32	-0.19	-0.20	-0.47	-0.19	0.63	0.28	0.32	0.14	0.58	0.00

Q91Z67	SLIT-ROBO Rho GTPase-activating protein 2 OS=Mus musculus GN=Srgap2 PE=1 SV=2 - [SRGP2_MOUSE]	0.20	0.02	0.21	0.04	0.07	0.10	0.06	-0.08	-0.08	-0.04	-0.12	-0.05	0.00
P62717	60S ribosomal protein L18a OS=Mus musculus GN=Rpl18a PE=1 SV=1 - [RL18A_MOUSE]	-0.30	-0.20	-0.28	-0.17	-0.01	-0.03	0.02	0.21	0.06	-0.02	0.03	0.13	0.00
Q9JI59	Junctional adhesion molecule B OS=Mus musculus GN=Jam2 PE=1 SV=1 - [JAM2_MOUSE]	-0.33	-0.13	-0.09	-0.08	-0.04	-0.13	-0.09	0.23	0.31	0.25	0.11	0.42	0.00
P98191	Phosphatidate cytidyltransferase 1 OS=Mus musculus GN=Cds1 PE=1 SV=2 - [CDS1_MOUSE]	-0.03	-0.27	0.09	-0.46	-0.03	-0.47	0.27	0.22	0.42	0.16	0.42	0.07	0.00
Q920Q8	Influenza virus NS1A-binding protein homolog OS=Mus musculus GN=Ivns1abp PE=1 SV=2 - [NS1BP_MOUSE]	0.61	0.57	0.15	0.30	0.44	0.49	0.12	-0.12	-0.11	0.01	0.37	-0.15	0.00
Q8BLE7	Vesicular glutamate transporter 2 OS=Mus musculus GN=Slc17a6 PE=1 SV=1 - [VGLU2_MOUSE]	-0.09	-0.10	-0.23	-0.21	-0.06	-0.11	0.13	-0.11	0.13	0.07	0.00	0.14	0.00
P61294	Ras-related protein Rab-6B OS=Mus musculus GN=Rab6b PE=1 SV=1 - [RAB6B_MOUSE]	-0.24	-0.01	-0.07	-0.17	-0.10	-0.30	0.01	0.08	0.19	0.03	0.06	0.01	0.00
Q9DAM5	Mitochondrial thiamine pyrophosphate carrier OS=Mus musculus GN=Slc25a19 PE=2 SV=1 - [TPC_MOUSE]	0.64	0.73	-0.15	0.41	0.31	1.18	0.13	-0.17	-1.09	-0.26	-0.46	-0.65	0.00
Q3U5C7	Prickle-like protein 1 OS=Mus musculus GN=Prickle1 PE=1 SV=1 - [PRIC1_MOUSE]	0.30	0.38	0.14	0.06	-0.02	-0.04	-0.60	-0.35	0.00	-0.13	-0.44	-0.22	0.00
Q91WT4	DnaJ homolog subfamily C member 17 OS=Mus musculus GN=Dnajc17 PE=2 SV=2 - [DJC17_MOUSE]	-0.06	-0.26	-0.12	-0.12	-0.22	-0.28	0.19	0.05	0.63	0.13	0.42	0.09	0.00
Q9WU84	Copper chaperone for superoxide dismutase OS=Mus musculus GN=Ccs PE=1 SV=1 - [CCS_MOUSE]	0.22	0.28	0.28	0.23	0.00	0.31	0.16	-0.34	-0.16	-0.30	0.01	-0.29	0.00
P97494	Glutamate-cysteine ligase catalytic subunit OS=Mus musculus GN=Gclc PE=2 SV=4 - [GSH1_MOUSE]	-0.30	0.01	-0.03	-0.02	-0.16	-0.14	0.19	0.04	0.18	0.01	0.22	0.07	0.00

Q9DC51	Guanine nucleotide-binding protein G(k) subunit alpha OS=Mus musculus GN=Gnai3 PE=1 SV=3 - [GNAI3_MOUSE]	-0.11	-0.38	-0.15	-0.12	0.01	-0.33	0.15	0.10	0.13	-0.06	0.01	0.27	0.00
Q80XU8	Leucine-rich repeat and fibronectin type-III domain-containing protein 4 OS=Mus musculus GN=Lrfn4 PE=1 SV=1 - [LRFN4_MOUSE]	-0.17	-0.28	-0.06	-0.29	-0.11	-0.40	-0.06	-0.11	0.03	0.18	0.07	0.05	0.00
Q60973	Histone-binding protein RBBP7 OS=Mus musculus GN=Rbbp7 PE=1 SV=1 - [RBBP7_MOUSE]	-0.39	-0.42	0.08	-0.18	-0.15	-0.26	0.20	0.27	-0.08	0.20	-0.02	0.26	0.00
Q9WUC3	Lymphocyte antigen 6H OS=Mus musculus GN=Ly6h PE=2 SV=2 - [LY6H_MOUSE]	-1.63	-0.81	-1.89	-0.95	-0.17	-0.30	-0.10	0.68	-0.01	0.56	0.17	0.60	0.00
P84244	Histone H3.3 OS=Mus musculus GN=H3f3a PE=1 SV=2 - [H33_MOUSE]	-0.74	-0.42	-0.56	-0.24	-0.17	-0.14	-0.39	0.55	0.27	0.33	0.26	0.31	0.00
Q8C6G1	Protein C21orf2 homolog OS=Mus musculus PE=2 SV=1 - [CU002_MOUSE]	0.92	1.33	0.21	1.04	0.47	0.26	-0.48	-0.25	0.21	-0.25	0.15	-0.16	0.00
A1L341	Dual-specificity tyrosine-phosphorylation-regulated kinase 1A OS=Mus musculus GN=Dyrk1a PE=2 SV=1 - [A1L341_MOUSE]	-0.13	-0.45	-0.11	-0.20	-0.20	-0.38	0.29	-0.02	-0.10	0.01	0.02	0.04	0.00
Q9CQ62	2,4-dienoyl-CoA reductase, mitochondrial OS=Mus musculus GN=Decr1 PE=1 SV=1 - [DECR_MOUSE]	0.35	0.18	0.15	0.13	0.10	0.10	0.09	-0.17	-0.02	0.02	0.01	-0.11	0.00
Q7TNF0	Double C2-like domain-containing protein alpha OS=Mus musculus GN=Doc2a PE=2 SV=1 - [DOC2A_MOUSE]	-0.49	-0.01	-0.14	-0.12	-0.23	-0.35	0.02	0.25	0.20	-0.13	0.10	0.33	0.00
O08638	Myosin-11 OS=Mus musculus GN=Myh11 PE=1 SV=1 - [MYH11_MOUSE]	0.36	0.85	0.16	0.64	-0.12	0.41	0.02	-1.08	-0.05	-0.92	-0.24	-0.91	0.00
G3UYF9	Prefoldin subunit 6 OS=Mus musculus GN=H2-Ke2 PE=2 SV=1 - [G3UYF9_MOUSE]	-0.08	0.30	-0.27	0.13	0.12	0.17	-0.72	-0.31	-0.37	-0.11	-0.27	-0.35	0.00
Q9CYN9	Renin receptor OS=Mus musculus GN=Atp6ap2 PE=2 SV=2 - [RENK_MOUSE]	-0.55	-0.26	-0.07	-0.51	-0.49	-0.60	-0.08	0.02	0.62	-0.10	0.07	0.01	0.00
Q80ZX8-2	Isoform 2 of Sperm-associated antigen 1 OS=Mus musculus GN=Spag1 - [SPAG1_MOUSE]	-0.14	-0.41	-0.05	-0.53	-0.48	-0.50	0.13	-0.03	0.46	0.03	-0.11	-0.03	0.00

F7CUT1	Interleukin-1 receptor-associated kinase 1 (Fragment) OS=Mus musculus GN=Irak1 PE=4 SV=1 - [F7CUT1_MOUSE]	1.34	0.81	0.22	0.52	0.05	0.85	0.02	-0.08	-0.25	-0.55	-0.23	-0.22	0.00
Q68EF0	Rab-3A-interacting protein OS=Mus musculus GN=Rab3ip PE=1 SV=1 - [RAB3I_MOUSE]	-0.28	-0.47	-0.10	-0.14	-0.03	-0.48	-0.06	0.16	-0.02	0.35	0.06	0.22	0.01
D3Z2Y6	Protein S100a16 (Fragment) OS=Mus musculus GN=S100a16 PE=2 SV=1 - [D3Z2Y6_MOUSE]	0.21	0.44	-0.01	0.35	-0.02	0.20	-0.05	-0.40	0.03	-0.43	-0.05	-0.36	0.01
Q3V3N7	Protein Bbs1 OS=Mus musculus GN=Bbs1 PE=2 SV=1 - [Q3V3N7_MOUSE]	1.12	-0.09	0.48	0.16	0.43	0.55	0.18	-0.93	-0.14	-0.72	-0.21	-0.62	0.01
Q6PHN9	Ras-related protein Rab-35 OS=Mus musculus GN=Rab35 PE=1 SV=1 - [RAB35_MOUSE]	-0.26	-0.18	-0.01	-0.16	-0.27	-0.12	0.12	0.30	0.00	0.31	-0.11	0.18	0.01
Q8K0B2-3	Isoform 3 of Probable lysosomal cobalamin transporter OS=Mus musculus GN=Lmbd1 - [LMBD1_MOUSE]	-0.07	-0.34	-0.15	-0.54	0.04	-0.63	-0.13	0.16	0.50	0.24	0.16	0.35	0.01
E9PUC5	PH and SEC7 domain-containing protein 3 OS=Mus musculus GN=Psd3 PE=2 SV=1 - [E9PUC5_MOUSE]	-0.10	-0.03	-0.34	0.05	-0.15	0.01	0.15	0.11	0.23	0.02	0.35	0.15	0.01
Q91W39	Nuclear receptor coactivator 5 OS=Mus musculus GN=Ncoa5 PE=1 SV=1 - [NCOA5_MOUSE]	0.11	0.06	0.27	0.12	0.03	0.05	0.06	-0.19	0.01	-0.20	-0.10	-0.13	0.01
Q6NXJ0	Protein WWC2 OS=Mus musculus GN=Wwc2 PE=2 SV=1 - [WWC2_MOUSE]	0.45	0.16	-0.05	-0.03	0.11	0.12	0.65	0.44	0.23	0.67	0.36	0.73	0.01
O54941	SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily E member 1 OS=Mus musculus GN=Smarca1 PE=1 SV=1 - [SMCE1_MOUSE]	0.15	0.07	-0.04	0.10	0.17	0.36	-0.14	-0.08	-0.04	-0.31	-0.07	-0.03	0.01
Q60709	Amyloid-like protein 2 OS=Mus musculus GN=App2 PE=2 SV=1 - [Q60709_MOUSE]	-0.98	-0.54	-0.64	-0.58	0.04	-0.32	-0.03	0.08	0.22	0.15	0.23	0.00	0.01
Q9CR13	UPF0562 protein C7orf5 homolog OS=Mus musculus GN=C7orf5 PE=3 SV=1 - [CG055_MOUSE]	0.84	0.29	0.32	0.51	0.00	-0.10	-0.01	-0.16	-0.80	-0.51	-0.40	-0.14	0.01
Q9D4C9	Clavesin-1 OS=Mus musculus GN=Civs1 PE=2 SV=1 - [CLVS1_MOUSE]	0.12	0.05	0.21	-0.06	0.08	-0.08	0.00	-0.22	-0.12	-0.41	-0.22	-0.21	0.01

Q8VE38-2	Isoform 2 of Oxidoreductase NAD-binding domain-containing protein 1 OS=Mus musculus GN=Oxnad1 - [OXND1_MOUSE]	-0.24	0.07	-0.20	0.20	0.03	0.05	-0.51	-0.57	-0.09	-0.51	-0.18	-0.46	0.01
Q9EQK5	Major vault protein OS=Mus musculus GN=Mvp PE=1 SV=4 - [MVP_MOUSE]	-0.03	-0.06	0.03	-0.08	-0.16	-0.25	0.12	0.37	-0.02	0.21	0.11	0.09	0.01
A2A473	Protein Tox2 OS=Mus musculus GN=Tox2 PE=2 SV=1 - [A2A473_MOUSE]	-0.55	-0.82	-0.51	-0.44	-0.02	-0.97	-0.49	0.13	0.21	0.10	0.22	0.25	0.01
Q9R0M5-2	Isoform 2 of Thiamin pyrophosphokinase 1 OS=Mus musculus GN=Tpk1 - [TPK1_MOUSE]	-0.38	0.01	-0.39	0.14	-0.44	-0.75	0.21	-0.03	0.24	0.73	0.43	0.21	0.01
Q3UKC1	Tax1-binding protein 1 homolog OS=Mus musculus GN=Tax1bp1 PE=1 SV=2 - [TAXB1_MOUSE]	0.09	0.13	0.11	0.09	0.09	-0.10	-0.14	-0.03	-0.06	-0.09	-0.12	-0.02	0.01
Q0VGB7	Serine/threonine-protein phosphatase 4 regulatory subunit 2 OS=Mus musculus GN=Ppp4r2 PE=1 SV=1 - [PP4R2_MOUSE]	0.31	0.26	-0.02	0.43	0.13	0.05	-0.18	-0.55	0.03	-0.08	-0.08	-0.43	0.01
A2AP93	Mitogen-activated protein kinase kinase 7 OS=Mus musculus GN=Map3k7 PE=2 SV=1 - [A2AP93_MOUSE]	0.23	0.30	0.27	0.38	0.06	0.02	0.07	-0.37	0.02	-0.27	0.01	-0.15	0.01
E9Q1Y3	Apolipoprotein B-100 (Fragment) OS=Mus musculus GN=Apob PE=2 SV=1 - [E9Q1Y3_MOUSE]	0.52	-0.07	0.31	-0.01	-0.01	0.20	-0.17	-0.10	-0.16	-0.34	-0.20	-0.47	0.01
Q69ZA1	Cyclin-dependent kinase 13 OS=Mus musculus GN=Cdk13 PE=1 SV=3 - [CDK13_MOUSE]	0.34	0.66	0.27	0.15	0.19	0.17	-0.23	0.19	-0.01	-0.02	-0.03	-0.21	0.01
A0JNY3	Gephyrin OS=Mus musculus GN=Gphn PE=2 SV=1 - [A0JNY3_MOUSE]	-0.22	-0.18	0.03	-0.06	-0.14	-0.16	0.12	0.06	0.06	-0.06	0.08	-0.02	0.01
A2BDX0	Activity-dependent neuroprotector homeobox protein OS=Mus musculus GN=Adnp PE=3 SV=1 - [A2BDX0_MOUSE]	0.91	0.10	0.42	-0.15	0.05	0.15	-0.23	-0.35	-0.31	-0.57	-0.15	-0.85	0.01
P03911	NADH-ubiquinone oxidoreductase chain 4 OS=Mus musculus GN=Mtnd4 PE=1 SV=1 - [NU4M_MOUSE]	-0.27	-0.25	-0.18	-0.17	-0.02	-0.42	-0.08	-0.01	-0.06	0.14	0.16	-0.02	0.01
Q91X51	Golgi reassembly-stacking protein 1 OS=Mus musculus GN=Gorasp1 PE=2 SV=3 - [GORS1_MOUSE]	0.51	0.50	0.17	0.38	0.47	0.41	0.05	-0.91	-0.02	-0.50	0.07	-0.38	0.01

Q9DAS9	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-12 OS=Mus musculus GN=Gng12 PE=1 SV=3 - [GBG12_MOUSE]	-0.16	-0.03	-0.13	0.02	-0.04	-0.18	0.18	0.04	0.05	0.07	0.08	-0.04	0.01
P40237	CD82 antigen OS=Mus musculus GN=Cd82 PE=1 SV=1 - [CD82_MOUSE]	-0.22	-0.44	-0.04	-0.32	0.02	-0.60	-0.14	0.31	0.18	0.21	0.13	0.22	0.01
Q5XG73-2	Isoform 2 of Acyl-CoA-binding domain-containing protein 5 OS=Mus musculus GN=Acbd5 - [ACBD5_MOUSE]	-0.02	-0.12	0.05	-0.21	-0.13	-0.12	0.16	-0.05	0.02	0.27	0.17	0.15	0.01
P08752	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Mus musculus GN=Gnai2 PE=1 SV=5 - [GNAI2_MOUSE]	-0.14	-0.24	-0.09	-0.12	0.08	-0.32	0.12	0.03	0.06	0.15	0.12	0.14	0.01
Q8BVA5	UPF0554 protein C2orf43 homolog OS=Mus musculus PE=2 SV=1 - [CB043_MOUSE]	-0.11	-0.18	0.14	-0.36	-0.48	-0.42	0.42	0.32	-0.02	0.36	-0.10	0.57	0.01
F6RJV6	LanC-like protein 2 (Fragment) OS=Mus musculus GN=Lanc2 PE=2 SV=1 - [F6RJV6_MOUSE]	-0.02	-0.11	0.01	-0.09	-0.06	-0.07	0.11	0.13	-0.03	0.09	-0.02	0.11	0.01
Q8CFV4	Neuritin OS=Mus musculus GN=Nrn1 PE=1 SV=1 - [NRN1_MOUSE]	-1.07	-0.21	-0.52	-0.08	-0.13	-0.02	0.10	0.24	0.50	0.68	0.24	0.33	0.01
Q3TY65	Islet cell autoantigen 1-like protein OS=Mus musculus GN=Ica11 PE=2 SV=1 - [ICA11_MOUSE]	-0.31	-0.70	0.04	-0.17	-0.14	-0.08	0.12	1.36	0.43	1.54	0.30	0.99	0.01
Q9CU62	Structural maintenance of chromosomes protein 1A OS=Mus musculus GN=Smc1a PE=1 SV=4 - [SMC1A_MOUSE]	0.09	-0.07	0.14	0.06	0.09	0.16	0.02	-0.09	-0.23	-0.01	-0.08	-0.14	0.01
Q8VIG6	TRAF-interacting protein OS=Mus musculus GN=Traip PE=1 SV=2 - [TRAIP_MOUSE]	-0.63	-0.83	-0.30	-0.69	0.10	-0.73	-0.03	0.26	-0.28	-0.01	0.30	0.57	0.01
J3KMN0	Protein Kmt2d OS=Mus musculus GN=Kmt2d PE=4 SV=1 - [J3KMN0_MOUSE]	0.19	0.44	-0.01	0.67	0.12	0.87	-0.76	-0.05	-0.08	-0.05	-0.17	-0.33	0.01
Q8HW98	IgLON family member 5 OS=Mus musculus GN=Iglon5 PE=2 SV=2 - [IGLO5_MOUSE]	-0.06	-0.15	-0.33	0.00	-0.14	-0.24	0.13	0.02	0.28	0.01	0.23	-0.07	0.01
Q8BW96-2	Isoform 2 of Calcium/calmodulin-dependent protein kinase type 1D OS=Mus musculus GN=Camk1d - [KCC1D_MOUSE]	-0.08	0.34	0.23	0.44	0.04	0.26	0.08	-0.27	-0.09	-0.44	-0.21	-0.05	0.01

Q0VGU5	Vitamin K epoxide reductase complex subunit 1-like protein 1 OS=Mus musculus GN=Vkorc111 PE=2 SV=1 - [Q0VGU5_MOUSE]	-0.06	-0.54	0.03	-0.37	-0.18	-0.55	0.11	0.39	0.07	0.24	-0.03	0.07	0.01
Q9DBE9	pre-rRNA processing protein FTSJ3 OS=Mus musculus GN=Ftsj3 PE=1 SV=1 - [SPB1_MOUSE]	0.12	-0.04	0.26	0.22	0.23	0.49	0.04	-0.17	0.06	-0.21	-0.40	-0.08	0.01
Q8R0W6	NEDD4 family-interacting protein 1 OS=Mus musculus GN=Ndfip1 PE=1 SV=1 - [NFIP1_MOUSE]	-0.50	-0.37	-0.37	-0.38	0.04	-0.28	0.05	0.01	-0.14	-0.07	0.14	0.05	0.01
Q8BUU7	N-acetylglucosamine kinase OS=Mus musculus GN=Galk2 PE=2 SV=1 - [Q8BUU7_MOUSE]	0.10	-0.58	0.22	-0.60	-0.09	-0.60	0.37	-0.01	0.73	0.52	0.30	0.45	0.01
Q61142-2	Isoform 2 of Spindlin-1 OS=Mus musculus GN=Spin1 - [SPIN1_MOUSE]	0.21	0.43	0.01	0.03	0.23	0.37	-0.53	-0.35	-0.03	0.03	-0.01	-0.19	0.01
A2AJ26	ATP-binding cassette sub-family A member 2 OS=Mus musculus GN=Abca2 PE=3 SV=1 - [A2AJ26_MOUSE]	-0.27	-0.14	-0.06	-0.13	-0.19	-0.19	0.14	0.18	0.58	0.00	0.75	0.33	0.01
E9PV86	Protein Mctp1 OS=Mus musculus GN=Mctp1 PE=2 SV=1 - [E9PV86_MOUSE]	0.27	0.04	0.21	0.05	0.17	0.26	0.13	-0.16	0.02	-0.01	-0.14	-0.21	0.01
P61264	Syntaxin-1B OS=Mus musculus GN=Stx1b PE=1 SV=1 - [STX1B_MOUSE]	-0.06	-0.16	-0.03	-0.04	0.00	-0.08	0.02	0.01	0.06	-0.01	0.02	0.09	0.01
F8WJ31	Zinc finger and SCAN domain-containing protein 26 OS=Mus musculus GN=Zscan26 PE=2 SV=1 - [F8WJ31_MOUSE]	-0.19	-1.03	-0.43	-1.07	0.14	-1.02	-0.26	0.61	0.23	0.21	0.16	0.89	0.01
Q8K1M6	Dynamin-1-like protein OS=Mus musculus GN=Dnm1l PE=1 SV=2 - [DNM1L_MOUSE]	0.26	0.06	0.47	0.07	-0.10	0.05	-0.02	-0.14	-0.34	-0.45	-0.35	-0.07	0.01
Q9JI11	Serine/threonine-protein kinase 4 OS=Mus musculus GN=Stk4 PE=1 SV=1 - [STK4_MOUSE]	0.15	0.48	0.31	0.02	0.05	0.22	0.13	-0.17	-0.06	-0.04	-0.17	-0.27	0.01
Q3U0J8	TBC1 domain family member 2B OS=Mus musculus GN=Tbc1d2b PE=1 SV=2 - [TBD2B_MOUSE]	0.96	0.77	-0.13	0.25	0.14	0.20	-0.19	-1.26	-0.01	-1.25	-0.12	-0.94	0.01
P53986	Monocarboxylate transporter 1 OS=Mus musculus GN=Sic16a1 PE=1 SV=1 - [MOT1_MOUSE]	-0.04	-0.17	0.07	-0.07	0.00	-0.34	0.09	0.43	0.09	0.28	0.07	0.53	0.01

Q501J7	Phosphatase and actin regulator 4 OS=Mus musculus GN=Phactr4 PE=1 SV=2 - [PHAR4_MOUSE]	-0.37	-0.25	-0.28	-0.42	-0.26	-0.30	-0.23	0.25	0.24	-0.19	0.15	0.01	0.01
Q920M5-3	Isoform C of Coronin-6 OS=Mus musculus GN=Coro6 - [CORO6_MOUSE]	0.02	0.33	0.09	0.22	0.10	0.10	0.16	-0.08	-0.15	-0.12	-0.19	-0.23	0.01
Q9Z2W1	Serine/threonine-protein kinase 25 OS=Mus musculus GN=Stk25 PE=1 SV=2 - [STK25_MOUSE]	-0.55	-0.15	-0.27	-0.09	-0.31	-0.09	0.20	0.32	-0.09	-0.09	0.55	0.15	0.01
Q8CHY6	Transcriptional repressor p66 alpha OS=Mus musculus GN=Gatad2a PE=1 SV=2 - [P66A_MOUSE]	0.21	-0.01	0.10	0.18	0.03	0.52	-0.20	-0.03	-0.23	0.06	-0.28	-0.25	0.01
Q7M6Z0	Reticulon-4 receptor-like 2 OS=Mus musculus GN=Rtn4rl2 PE=2 SV=1 - [R4RL2_MOUSE]	0.14	0.02	-0.03	-0.06	-0.10	-0.14	0.19	-0.01	0.22	0.12	0.20	0.20	0.01
P00158	Cytochrome b OS=Mus musculus GN=Mt-Cyb PE=2 SV=1 - [CYB_MOUSE]	-0.90	-0.44	-0.06	-0.53	-0.62	-0.37	0.34	0.35	-0.15	0.03	-0.41	0.25	0.01
Q922Q4	Pyrroline-5-carboxylate reductase 2 OS=Mus musculus GN=Pycr2 PE=2 SV=1 - [P5CR2_MOUSE]	-0.25	-0.07	-0.19	-0.10	-0.27	-0.01	-0.17	0.35	0.39	0.12	0.15	0.29	0.01
Q9CYG7	Mitochondrial import receptor subunit TOM34 OS=Mus musculus GN=Tomm34 PE=2 SV=1 - [TOM34_MOUSE]	0.52	0.32	0.31	0.51	0.03	0.20	0.07	-0.39	-0.03	-0.01	-0.09	0.18	0.01
Q9CQ20	Mid1-interacting protein 1 OS=Mus musculus GN=Mid1ip1 PE=1 SV=1 - [M1IP1_MOUSE]	0.44	0.41	-0.18	0.24	-0.06	0.12	-0.08	-0.77	-0.23	-0.61	-0.03	-0.74	0.01
Q8BIF0-2	Isoform 2 of CD99 antigen-like protein 2 OS=Mus musculus GN=Cd99l2 - [C99L2_MOUSE]	0.01	-0.13	-0.40	-0.09	-0.27	-0.61	0.02	0.32	0.27	0.13	0.13	-0.07	0.01
Q3UHL1	CaM kinase-like vesicle-associated protein OS=Mus musculus GN=Camkv PE=1 SV=2 - [CAMKV_MOUSE]	0.17	0.08	0.18	0.17	-0.01	0.00	0.05	-0.02	-0.10	-0.13	-0.04	-0.05	0.01
P58044	Isopentenyl-diphosphate Delta-isomerase 1 OS=Mus musculus GN=Idi1 PE=2 SV=1 - [IDI1_MOUSE]	-0.52	-0.16	-0.56	-0.19	-0.03	-0.22	-0.30	0.22	0.40	0.17	0.19	0.16	0.01
Q9DCT1	1,5-anhydro-D-fructose reductase OS=Mus musculus GN=Akr1e2 PE=1 SV=1 - [AKCL2_MOUSE]	0.23	0.22	0.20	0.17	-0.01	0.36	0.06	0.00	-0.04	0.07	-0.01	-0.08	0.01

G3UYK9	Dystrobrevin beta (Fragment) OS=Mus musculus GN=Dtnb PE=4 SV=1 - [G3UYK9_MOUSE]	-0.02	0.37	0.37	0.22	0.47	0.26	0.11	-0.01	-0.08	-0.08	-0.04	0.12	0.01
Q3TIU4	2',5'-phosphodiesterase 12 OS=Mus musculus GN=Pde12 PE=2 SV=2 - [PDE12_MOUSE]	0.24	0.25	0.22	0.16	0.00	0.17	0.18	-0.01	-0.09	-0.11	0.01	-0.16	0.01
Q68FE6	Protein FAM65A OS=Mus musculus GN=Fam65a PE=1 SV=2 - [FA65A_MOUSE]	0.41	0.13	0.42	0.20	0.38	0.12	0.11	0.08	0.01	-0.01	0.11	-0.04	0.01
Q3UTJ2	Sorbin and SH3 domain-containing protein 2 OS=Mus musculus GN=Sorbs2 PE=1 SV=2 - [SRBS2_MOUSE]	-0.10	0.42	0.08	0.10	0.26	0.21	-0.18	-0.24	-0.01	-0.08	-0.05	-0.19	0.01
Q9D9E0-2	Isoform 2 of Solute carrier family 22 member 17 OS=Mus musculus GN=Slc22a17 - [S22AH_MOUSE]	-0.03	-0.30	-0.14	-0.43	0.00	-0.37	0.00	0.29	0.05	0.78	0.12	0.26	0.01
E9Q8S5	Microtubule-associated serine/threonine-protein kinase 3 OS=Mus musculus GN=Mast3 PE=2 SV=1 - [E9Q8S5_MOUSE]	-0.10	-0.24	-0.11	0.09	-0.11	-0.14	0.13	0.01	0.14	0.48	0.24	0.06	0.01
O54946-2	Isoform B of DnaJ homolog subfamily B member 6 OS=Mus musculus GN=Dnajb6 - [DNJB6_MOUSE]	-0.29	-0.11	-0.06	-0.09	-0.02	-0.10	0.00	0.31	-0.02	0.26	0.02	0.28	0.01
A8Y5G5	Protein Zfyve9 OS=Mus musculus GN=Zfyve9 PE=2 SV=1 - [A8Y5G5_MOUSE]	-0.20	-0.08	-0.21	0.05	-0.13	0.07	-0.08	0.25	0.25	0.21	0.11	0.13	0.01
Q9D5V6	Synapse-associated protein 1 OS=Mus musculus GN=Syap1 PE=1 SV=1 - [SYAP1_MOUSE]	-0.07	0.02	0.06	0.08	0.17	0.24	-0.18	-0.15	-0.05	-0.08	-0.12	0.00	0.01
Q99KY4	Cyclin-G-associated kinase OS=Mus musculus GN=Gak PE=1 SV=2 - [GAK_MOUSE]	0.13	0.04	0.28	0.08	0.00	-0.06	0.02	-0.11	-0.19	-0.13	-0.17	-0.09	0.01
Q9Z1W9	STE20/SPS1-related proline-alanine-rich protein kinase OS=Mus musculus GN=Stk39 PE=1 SV=1 - [STK39_MOUSE]	-0.03	-0.09	-0.02	-0.15	0.00	-0.14	0.02	0.05	-0.05	0.17	0.09	0.17	0.01
Q6NZL0	Protein SOGA3 OS=Mus musculus GN=Soga3 PE=2 SV=2 - [SOGA3_MOUSE]	0.07	0.02	0.28	-0.07	0.02	-0.04	-0.09	-0.12	-0.22	-0.14	-0.28	-0.07	0.01
Q5SUC9	Protein SCO1 homolog, mitochondrial OS=Mus musculus GN=Sco1 PE=2 SV=1 - [SCO1_MOUSE]	-0.27	-0.02	0.01	-0.20	-0.16	-0.05	0.03	0.20	0.08	0.06	0.05	-0.02	0.01

Q8BHC4	Dephospho-CoA kinase domain-containing protein OS=Mus musculus GN=DcaKd PE=2 SV=1 - [DCAKD_MOUSE]	1.67	0.25	0.84	0.30	0.92	0.78	0.29	-0.49	0.12	-0.44	0.31	-0.07	0.01
Q9QUI0	Transforming protein RhoA OS=Mus musculus GN=Rhoa PE=1 SV=1 - [RHOA_MOUSE]	-0.53	-0.22	-0.05	-0.02	-0.14	0.01	-0.05	0.24	0.32	0.09	0.17	0.25	0.01
O35114	Lysosome membrane protein 2 OS=Mus musculus GN=Scarb2 PE=1 SV=3 - [SCR2_MOUSE]	-0.16	-0.19	-0.06	-0.04	-0.06	-0.25	-0.09	0.05	0.06	0.14	-0.03	0.01	0.01
P17012	Zinc finger X-chromosomal protein OS=Mus musculus GN=Zfx PE=1 SV=2 - [ZFX_MOUSE]	0.09	0.21	0.02	-0.11	0.17	-0.03	-0.49	-0.42	-0.02	-0.55	-0.02	-0.39	0.01
Q8CH77	Neuron navigator 1 OS=Mus musculus GN=Nav1 PE=1 SV=2 - [NAV1_MOUSE]	-0.01	-0.04	0.14	0.05	0.00	0.03	-0.26	-0.17	-0.06	-0.33	-0.05	-0.07	0.01
Q8K0P3	TLD domain-containing protein 1 OS=Mus musculus GN=Tldc1 PE=2 SV=1 - [TLDC1_MOUSE]	-0.12	-0.08	-0.13	-0.20	-0.19	-0.60	0.34	0.37	0.21	-0.12	0.25	-0.10	0.01
Q9CQ06	39S ribosomal protein L24, mitochondrial OS=Mus musculus GN=Mrl24 PE=2 SV=1 - [RM24_MOUSE]	0.10	0.40	0.08	0.11	-0.02	0.43	0.09	-0.61	-0.13	-0.23	-0.12	-0.24	0.01
Q6P542	ATP-binding cassette sub-family F member 1 OS=Mus musculus GN=Abcf1 PE=1 SV=1 - [ABCF1_MOUSE]	0.03	0.09	0.14	-0.10	0.05	0.14	-0.14	-0.04	-0.12	-0.05	-0.26	-0.02	0.01
Q9DCJ1	Target of rapamycin complex subunit LST8 OS=Mus musculus GN=Mst8 PE=1 SV=1 - [LST8_MOUSE]	0.51	0.13	0.10	0.10	0.08	0.00	-0.21	-0.27	-0.01	-0.04	-0.08	-0.17	0.01
Q9D6W8	Uncharacterized protein C17orf59 homolog OS=Mus musculus PE=1 SV=1 - [CQ059_MOUSE]	0.13	-0.09	0.01	0.26	-0.04	0.22	-0.19	-0.16	-0.07	-0.17	0.01	-0.29	0.01
F8VQK3	Protein Gucy1a2 OS=Mus musculus GN=Gucy1a2 PE=2 SV=1 - [F8VQK3_MOUSE]	-0.04	-0.19	0.03	-0.17	-0.17	-0.17	0.17	0.05	-0.06	0.11	-0.07	0.19	0.01
Q3TIR1-3	Isoform 3 of Trafficking protein particle complex subunit 13 OS=Mus musculus GN=Trappc13 - [TPC13_MOUSE]	0.11	-0.10	-0.06	-0.02	-0.15	0.18	0.09	0.28	0.17	0.26	0.11	0.31	0.01
Q3U5Q7	UMP-CMP kinase 2, mitochondrial OS=Mus musculus GN=Cmpk2 PE=2 SV=2 - [CMPK2_MOUSE]	-0.08	-0.01	0.01	-0.06	-0.15	-0.17	0.19	0.00	0.07	0.00	0.06	0.03	0.01

F8VQF9	Trafficking protein particle complex subunit 10 OS=Mus musculus GN=Trappc10 PE=4 SV=1 - [F8VQF9_MOUSE]	-0.03	-0.16	0.15	-0.04	-0.12	-0.04	0.26	0.10	0.05	0.08	0.14	0.18	0.01
A2AG36	GPI ethanolamine phosphate transferase 3 OS=Mus musculus GN=Pigo PE=2 SV=1 - [A2AG36_MOUSE]	1.08	0.26	0.34	-0.04	0.00	-0.05	-0.52	-0.18	-0.56	-0.22	-0.71	-0.26	0.01
Q9CQA6	Coiled-coil-helix-coiled-coil-helix domain-containing protein 1 OS=Mus musculus GN=Chchd1 PE=2 SV=1 - [CHCH1_MOUSE]	-0.84	-0.11	-0.82	-0.62	-0.09	0.00	-0.23	0.51	1.04	0.03	1.07	0.26	0.01
E9Q6H7	Uncharacterized protein OS=Mus musculus GN=Tmem181b-ps PE=4 SV=1 - [E9Q6H7_MOUSE]	0.85	0.31	-0.06	0.56	-0.10	0.46	0.01	-0.48	-0.07	-0.67	0.00	-0.41	0.01
Q61735-2	Isoform 2 of Leukocyte surface antigen CD47 OS=Mus musculus GN=Cd47 - [CD47_MOUSE]	-0.29	-0.11	-0.20	-0.08	-0.09	-0.22	-0.05	0.02	0.09	-0.17	0.07	0.14	0.01
P46638	Ras-related protein Rab-11B OS=Mus musculus GN=Rab11b PE=1 SV=3 - [RB11B_MOUSE]	0.00	-0.07	-0.08	-0.07	0.00	-0.04	0.23	-0.02	0.03	0.09	0.13	0.08	0.01
Q80U72	Protein scribble homolog OS=Mus musculus GN=Scrib PE=1 SV=2 - [SCRIB_MOUSE]	0.26	0.22	0.14	0.33	-0.01	0.11	0.00	-0.19	-0.04	0.15	-0.08	-0.06	0.01
P59913	Protein-L-isoaspartate O-methyltransferase domain-containing protein 1 OS=Mus musculus GN=Pcmd1 PE=2 SV=1 - [PCMD1_MOUSE]	-0.26	-0.12	0.06	-0.14	-0.05	-0.03	0.17	0.07	-0.02	0.24	0.11	0.02	0.01
Q9JM52-3	Isoform 3 of Misshapen like kinase 1 OS=Mus musculus GN=Mink1 - [MINK1_MOUSE]	0.00	-0.01	0.23	-0.01	0.02	-0.14	-0.14	-0.17	-0.32	-0.03	-0.27	-0.18	0.01
Q61503	5'-nucleotidase OS=Mus musculus GN=Nt5e PE=1 SV=2 - [5NTD_MOUSE]	-0.10	-0.27	-0.24	-0.27	-0.21	-0.21	-0.05	0.37	-0.11	0.39	0.02	0.09	0.01
E9PUA2	Terminal uridylyltransferase 7 OS=Mus musculus GN=Zcchc6 PE=4 SV=1 - [E9PUA2_MOUSE]	-0.26	-0.12	-0.69	0.06	-0.70	-0.08	0.25	0.23	0.72	-0.04	0.35	-0.01	0.01
Q67BT3	Solute carrier family 13 member 5 OS=Mus musculus GN=Slc13a5 PE=2 SV=1 - [S13A5_MOUSE]	-0.97	-0.76	-0.28	-0.92	0.01	-0.81	-0.10	-0.07	0.38	-0.07	-0.13	-0.09	0.01

Q69ZZ6-2	Isoform 2 of Transmembrane and coiled-coil domains protein 1 OS=Mus musculus GN=Tmcc1 - [TMCC1_MOUSE]	0.27	0.24	0.20	0.14	0.03	0.08	-0.01	0.13	-0.19	0.02	-0.21	-0.02	0.01
Q9D9V3-2	Isoform 2 of Ethylmalonyl-CoA decarboxylase OS=Mus musculus GN=Echdc1 - [ECHD1_MOUSE]	0.00	-0.24	-0.06	-0.19	-0.42	-0.33	-0.23	0.17	0.45	0.31	0.11	0.15	0.01
E9PZ00	Sulfated glycoprotein 1 OS=Mus musculus GN=Psap PE=2 SV=1 - [E9PZ00_MOUSE]	-1.25	-0.49	-0.82	-0.28	0.04	-0.18	-0.29	0.19	0.57	0.13	0.55	0.31	0.01
P59110	Sentrin-specific protease 1 OS=Mus musculus GN=Senp1 PE=2 SV=1 - [SENP1_MOUSE]	-0.65	-0.45	-0.27	-0.06	-0.02	-0.09	-0.04	0.53	-0.11	0.52	0.08	0.46	0.01
P97370	Sodium/potassium-transporting ATPase subunit beta-3 OS=Mus musculus GN=Atp1b3 PE=1 SV=1 - [AT1B3_MOUSE]	-0.43	-0.06	-0.04	0.07	-0.10	-0.04	0.16	0.12	0.26	0.13	0.12	0.21	0.01
Q8BJ1	Sodium-dependent neutral amino acid transporter SLC6A17 OS=Mus musculus GN=Slc6a17 PE=1 SV=1 - [S6A17_MOUSE]	0.02	-0.13	0.06	-0.13	-0.07	-0.17	0.07	0.00	0.13	0.11	0.00	0.13	0.01
A2A9P6	Cyclin-dependent kinase 11B OS=Mus musculus GN=Cdk11b PE=2 SV=1 - [A2A9P6_MOUSE]	-0.01	-0.08	0.07	0.19	0.07	0.08	-0.14	-0.16	-0.11	0.02	-0.03	-0.22	0.01
Q920Q6-3	Isoform 3 of RNA-binding protein Musashi homolog 2 OS=Mus musculus GN=Ms12 - [MS12H_MOUSE]	0.34	0.06	0.10	0.09	0.18	0.01	0.11	-0.23	-0.12	-0.12	-0.03	-0.06	0.01
Q80TJ1-4	Isoform 4 of Calcium-dependent secretion activator 1 OS=Mus musculus GN=Cadps - [CAPS1_MOUSE]	0.13	0.90	0.37	0.71	0.25	0.50	0.29	0.17	-0.23	0.00	-0.02	0.00	0.01
P61027	Ras-related protein Rab-10 OS=Mus musculus GN=Rab10 PE=1 SV=1 - [RAB10_MOUSE]	-0.03	-0.35	-0.06	-0.28	-0.10	-0.12	-0.03	0.21	0.00	0.04	-0.04	0.28	0.01
Q0VE82	Copine-7 OS=Mus musculus GN=Cpne7 PE=2 SV=1 - [CPNE7_MOUSE]	0.18	0.31	0.24	-0.24	0.01	0.00	0.06	-0.55	-0.27	-0.27	-0.22	-0.59	0.01
Q9DC48	Pre-mRNA-processing factor 17 OS=Mus musculus GN=Cdc40 PE=1 SV=1 - [PRP17_MOUSE]	0.69	0.22	0.36	-0.28	0.71	0.20	-0.02	-1.09	0.09	-0.74	-0.29	-1.23	0.01
E9Q226	Protein Ccny1 OS=Mus musculus GN=Ccny1 PE=2 SV=1 - [E9Q226_MOUSE]	-0.09	-0.93	-0.01	-0.44	-0.41	-0.29	-0.16	0.13	0.17	0.34	0.08	0.26	0.01

A3KMP2	Tetratricopeptide repeat protein 38 OS=Mus musculus GN=Ttc38 PE=2 SV=2 - [TTC38_MOUSE]	0.02	-0.16	-0.13	0.06	-0.13	-0.19	0.10	0.07	-0.01	0.17	0.36	0.04	0.01
Q61543	Golgi apparatus protein 1 OS=Mus musculus GN=Glg1 PE=1 SV=1 - [GSLG1_MOUSE]	-0.64	-0.14	-0.30	-0.27	-0.14	-0.06	-0.13	-0.03	0.19	0.10	0.12	0.09	0.01
Q9DAR7	m7GpppX diphosphatase OS=Mus musculus GN=Dcps PE=1 SV=1 - [DCPS_MOUSE]	-0.15	-0.12	-0.02	-0.07	-0.06	-0.08	-0.03	-0.01	-0.03	0.00	0.09	-0.03	0.01
G5E890	Transcription factor RFX3 OS=Mus musculus GN=Rfx3 PE=2 SV=1 - [G5E890_MOUSE]	0.17	0.36	0.19	0.03	-0.10	0.35	-0.03	-0.08	-0.02	-0.33	-0.18	-0.03	0.01
Q3TNA1	Xylose kinase OS=Mus musculus GN=Xylb PE=2 SV=1 - [XYLB_MOUSE]	1.11	0.79	0.34	0.67	0.53	0.92	-0.07	0.17	-0.36	0.61	0.53	-0.07	0.01
Q6P1J1	Crmp1 protein OS=Mus musculus GN=Crmp1 PE=2 SV=1 - [Q6P1J1_MOUSE]	-0.03	0.29	0.01	0.15	-0.01	0.21	0.12	-0.16	-0.18	-0.25	-0.15	-0.12	0.01
Q9JI75	Ribosylidihydroxynicotinamide dehydrogenase [quinone] OS=Mus musculus GN=Nqo2 PE=2 SV=3 - [NQO2_MOUSE]	-0.01	-0.11	-0.01	-0.12	-0.13	-0.18	0.20	-0.01	0.05	-0.08	0.21	0.18	0.01
Q3UR70	Transforming growth factor-beta receptor-associated protein 1 OS=Mus musculus GN=Tgfbra1 PE=2 SV=1 - [TGFA1_MOUSE]	0.15	0.24	0.21	0.28	0.11	-0.18	0.17	-0.19	-0.23	-0.50	-0.20	-0.41	0.01
Q9DC22	DDB1- and CUL4-associated factor 6 OS=Mus musculus GN=Dcaf6 PE=1 SV=1 - [DCAF6_MOUSE]	0.87	0.06	0.20	0.24	0.10	0.24	-0.28	-0.33	0.00	0.01	-0.22	-0.14	0.01
Q9CS42	Ribose-phosphate pyrophosphokinase 2 OS=Mus musculus GN=Prps2 PE=1 SV=4 - [PRPS2_MOUSE]	0.39	0.26	0.12	0.20	-0.09	0.28	0.08	-0.66	-0.04	-0.43	-0.04	-0.59	0.01
F6S7W6	Serine/threonine-protein kinase SIK3 (Fragment) OS=Mus musculus GN=SiK3 PE=2 SV=1 - [F6S7W6_MOUSE]	-0.22	0.00	-0.20	-0.16	-0.12	0.01	-0.07	-0.02	0.07	0.09	0.12	0.02	0.01
E9PYK8	Rho guanine nucleotide exchange factor 4 OS=Mus musculus GN=Arhgef4 PE=2 SV=2 - [E9PYK8_MOUSE]	0.77	0.45	-0.07	0.22	0.12	0.12	-0.21	-0.07	-0.54	-0.09	-0.64	0.11	0.01
E9PX57	Kinesin-like protein KIF21B OS=Mus musculus GN=Kif6 PE=2 SV=1 - [E9PX57_MOUSE]	-0.28	-0.14	0.14	-0.13	-0.10	-0.04	-0.28	-0.27	-0.32	-0.45	-0.33	-0.16	0.01

Q6PB89	Adarb2 protein OS=Mus musculus GN=Adarb2 PE=2 SV=1 - [Q6PB89_MOUSE]	-0.05	-0.18	0.32	-0.45	0.01	-0.12	0.07	0.69	0.32	0.74	0.06	0.70	0.01
Q64018	Glycine receptor subunit alpha-1 OS=Mus musculus GN=Glr1 PE=1 SV=2 [GLRA1_MOUSE]	-0.05	-0.21	-0.08	0.03	-0.20	-0.43	0.05	0.11	0.12	-0.10	0.13	0.31	0.01
D3Z698	Lipoma HMGIC fusion partner-like 2 protein (Fragment) OS=Mus musculus GN=Lhpl2 PE=2 SV=1 - [D3Z698_MOUSE]	0.04	-0.22	-0.28	-0.38	-0.13	-0.14	0.49	0.05	0.21	0.29	0.12	-0.19	0.01
Q60963	Platelet-activating factor acetylhydrolase OS=Mus musculus GN=Pla2g7 PE=2 SV=2 - [PAFA_MOUSE]	-0.22	-0.17	-0.12	-0.05	-0.03	0.04	0.07	0.04	0.10	0.06	-0.06	0.14	0.01
J3QQ44	Endophilin-A3 OS=Mus musculus GN=Sh3gl3 PE=4 SV=1 - [J3QQ44_MOUSE]	-2.64	-1.21	-3.12	-0.66	-0.13	0.14	0.21	1.08	0.03	1.06	0.79	1.08	0.01
P32507-2	Isoform Alpha of Poliovirus receptor- related protein 2 OS=Mus musculus GN=Pvrl2 - [PVRL2_MOUSE]	0.55	-1.07	-1.20	-0.82	-1.53	-0.98	0.15	0.21	0.44	0.34	0.28	0.06	0.01
Q810U4-5	Isoform 5 of Neuronal cell adhesion molecule OS=Mus musculus GN=Nrcam - [NRCAM_MOUSE]	-0.04	0.01	0.24	-0.11	0.10	-0.01	0.06	-0.45	-0.29	-0.13	-0.30	-0.34	0.01
Q9WT17-4	Isoform 4 of Unconventional myosin- Ic OS=Mus musculus GN=Myo1c - [MYO1C_MOUSE]	-0.16	-0.22	0.00	-0.26	-0.19	-0.10	0.19	-0.01	-0.08	-0.03	0.08	-0.07	0.01
F6SDD7	DNA topoisomerase (Fragment) OS=Mus musculus GN=Top3b PE=3 SV=1 - [F6SDD7_MOUSE]	0.34	0.21	0.24	-0.08	0.05	0.41	-0.12	-0.35	-0.24	0.09	-0.29	0.11	0.01
B2RQ80	Tnik protein OS=Mus musculus GN=Tnik PE=2 SV=1 - [B2RQ80_MOUSE]	-0.02	-0.07	0.14	0.02	0.09	0.02	-0.13	-0.05	-0.17	-0.05	-0.05	-0.06	0.01
O70589	Peripheral plasma membrane protein CASK OS=Mus musculus GN=Cask PE=1 SV=2 - [CSKP_MOUSE]	-0.77	-0.61	0.00	-0.60	-0.18	-0.47	-0.45	0.28	0.04	-0.05	0.26	0.45	0.01
P38060	Hydroxymethylglutaryl- CoA lyase, mitochondrial OS=Mus musculus GN=Hmgcl PE=1 SV=2 - [HMGCL_MOUSE]	-0.70	-0.29	-0.12	-0.16	-0.18	-0.14	-0.04	0.02	0.07	0.06	0.26	0.01	0.01
B1AUY8	N-alpha- acetyltransferase 10 OS=Mus musculus GN=Naa10 PE=2 SV=1 - [B1AUY8_MOUSE]	-0.02	-0.34	0.06	-0.09	-0.10	-0.09	0.00	0.13	-0.05	0.22	0.32	0.22	0.01

E9Q238	Protein Kank1 OS=Mus musculus GN=Kank1 PE=2 SV=1 - [E9Q238_MOUSE]	0.47	0.49	-0.01	0.43	-0.24	0.19	-0.21	0.15	-0.11	-0.43	-0.48	-0.41	0.01
Q8R464	Cell adhesion molecule 4 OS=Mus musculus GN=Cadm4 PE=1 SV=1 - [CADM4_MOUSE]	-0.15	-0.11	-0.16	-0.01	-0.10	-0.20	-0.09	0.00	0.28	0.06	0.13	0.01	0.01
E9Q0V6	NHS-like protein 2 OS=Mus musculus GN=Nhs12 PE=2 SV=1 -[E9Q0V6_MOUSE]	0.02	0.37	0.04	0.34	0.01	-0.12	-0.05	-0.23	-0.01	-0.43	-0.16	-0.36	0.01
Q80SY6-4	Isoform 4 of Adenosine deaminase-like protein OS=Mus musculus GN=Adal - [ADAL_MOUSE]	-0.30	-0.10	-0.18	-0.08	-0.46	0.36	0.51	0.02	0.66	0.19	0.97	0.20	0.01
Q99K28	ADP-ribosylation factor GTPase-activating protein 2 OS=Mus musculus GN=Arfgap2 PE=1 SV=1 - [ARFG2_MOUSE]	-0.10	0.19	0.10	0.16	0.08	0.27	-0.14	-0.05	-0.07	0.04	-0.09	-0.09	0.01
B7FAU9	Filamin, alpha OS=Mus musculus GN=Flna PE=4 SV=1 - [B7FAU9_MOUSE]	-0.08	0.09	0.31	-0.03	0.02	-0.01	-0.11	-0.09	-0.20	-0.09	-0.21	-0.18	0.01
Q62442-2	Isoform 2 of Vesicle- associated membrane protein 1 OS=Mus musculus GN=Vamp1 - [VAMP1_MOUSE]	0.08	0.05	-0.04	0.03	-0.11	-0.11	0.24	0.06	0.09	0.03	0.13	0.16	0.01
Q61553	Fascin OS=Mus musculus GN=Fscn1 PE=1 SV=4 - [FSCN1_MOUSE]	-0.07	-0.03	-0.01	0.02	-0.08	0.03	0.19	0.13	0.09	-0.01	0.09	0.02	0.01
Q8BL06	Inactive ubiquitin carboxyl-terminal hydrolase 54 OS=Mus musculus GN=Usp54 PE=1 SV=2 - [UBP54_MOUSE]	0.14	0.24	0.11	0.42	0.13	0.41	-0.17	-0.04	-0.09	0.30	-0.13	-0.04	0.01
Q6P5F7	Protein tweety homolog 3 OS=Mus musculus GN=Ttyh3 PE=1 SV=1 - [TTYH3_MOUSE]	-0.15	-0.21	0.05	-0.04	-0.07	-0.30	0.04	0.06	0.21	0.05	-0.05	0.08	0.01
Q9WVF8	Tumor suppressor candidate 2 OS=Mus musculus GN=Tusc2 PE=1 SV=3 - [TUSC2_MOUSE]	-0.39	0.03	-0.33	-0.29	-0.15	0.10	0.18	0.12	0.23	0.07	0.15	-0.04	0.01
Q8VBY2	Calcium/calmodulin- dependent protein kinase kinase 1 OS=Mus musculus GN=Camkk1 PE=1 SV=1 - [KKCC1_MOUSE]	0.18	0.25	0.09	0.18	-0.06	0.12	-0.06	-0.04	-0.09	-0.01	0.09	-0.08	0.01
P51432	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase beta-3 OS=Mus musculus GN=Plcb3 PE=2 SV=2 - [PLCB3_MOUSE]	0.05	0.10	0.05	0.03	-0.01	0.00	0.05	-0.11	-0.14	-0.06	-0.09	-0.22	0.01

G5E876	Hect domain and RLD 3, isoform CRA_a OS=Mus musculus GN=Herc3 PE=4 SV=1 - [G5E876_MOUSE]	-0.34	-0.09	-0.60	-0.46	-0.95	0.00	0.01	-0.03	0.21	0.02	0.13	0.25	0.01
A2RSX9	Arfp1 protein OS=Mus musculus GN=Arfp1 PE=2 SV=1 - [A2RSX9_MOUSE]	0.20	0.10	0.23	0.22	0.16	0.00	-0.04	0.05	-0.17	0.05	-0.23	0.09	0.01
Q99K67	Alpha-aminoadipic semialdehyde synthase, mitochondrial OS=Mus musculus GN=Aass PE=2 SV=1 - [AASS_MOUSE]	0.29	0.15	0.29	0.41	-0.02	0.08	0.28	-0.38	-0.20	-0.41	-0.06	-0.41	0.01
Q8K2C8	Glycerol-3-phosphate acyltransferase 4 OS=Mus musculus GN=Agpat6 PE=2 SV=1 - [GPAT4_MOUSE]	0.37	0.04	0.41	0.11	0.15	0.17	0.25	0.62	0.54	0.42	0.39	0.45	0.01
Q8BIP0	Aspartate-tRNA ligase, mitochondrial OS=Mus musculus GN=Dars2 PE=2 SV=1 - [SYDM_MOUSE]	-0.04	0.05	-0.21	0.00	0.02	-0.06	0.30	0.13	0.27	0.11	0.66	0.11	0.01
Q8BUY8	G-protein coupled receptor-associated sorting protein 2 OS=Mus musculus GN=Gprasp2 PE=2 SV=2 - [GASP2_MOUSE]	0.07	1.00	0.01	1.16	-0.11	0.85	-0.11	-1.75	0.08	-1.71	0.00	-1.38	0.02
Q8BPE4	Transmembrane protein 177 OS=Mus musculus GN=Tmem177 PE=2 SV=1 - [TM177_MOUSE]	0.28	-0.32	0.33	-0.24	0.17	0.01	-0.30	0.00	-0.42	-0.38	-0.73	-0.63	0.02
Q8R361	Rab11 family-interacting protein 5 OS=Mus musculus GN=Rab11fip5 PE=1 SV=2 - [RFIP5_MOUSE]	0.04	-0.06	0.13	0.05	0.08	-0.10	-0.18	-0.12	-0.10	-0.05	-0.20	-0.01	0.02
E9Q447	Spectrin alpha chain, non-erythrocytic 1 OS=Mus musculus GN=Sptan1 PE=2 SV=1 - [E9Q447_MOUSE]	-0.54	-0.25	-0.11	-0.14	0.13	0.21	-0.08	0.39	0.24	0.48	0.31	0.39	0.02
Q62167	ATP-dependent RNA helicase DDX3X OS=Mus musculus GN=Ddx3x PE=1 SV=3 - [DDX3X_MOUSE]	-0.24	-0.20	0.03	-0.08	-0.16	-0.18	0.18	0.24	-0.10	0.17	-0.10	0.06	0.02
O54916	RalBP1-associated Eps domain-containing protein 1 OS=Mus musculus GN=Reps1 PE=1 SV=2 - [REPS1_MOUSE]	0.02	0.02	0.02	0.11	0.16	0.05	-0.06	-0.07	-0.01	0.01	-0.13	0.03	0.02
Q9CZG3	COMM domain-containing protein 8 OS=Mus musculus GN=Comm8 PE=2 SV=1 - [COMD8_MOUSE]	-0.30	-0.16	-0.64	-0.07	-0.29	-0.05	0.07	0.07	-0.07	0.07	0.04	0.18	0.02

H7BXB9	Protein 1700037H04Rik (Fragment) OS=Mus musculus GN=1700037H04Rik PE=2 SV=1 - [H7BXB9_MOUSE]	-0.07	0.34	-0.09	0.03	0.06	0.18	0.07	-0.58	-0.17	-0.29	-0.24	-0.25	0.02
P80315	T-complex protein 1 subunit delta OS=Mus musculus GN=Cct4 PE=1 SV=3 - [TCPD_MOUSE]	0.20	0.51	0.12	0.26	0.05	0.33	0.19	-0.04	-0.03	-0.22	0.04	0.02	0.02
E9QKZ2	Importin-9 OS=Mus musculus GN=Ipo9 PE=2 SV=1 - [E9QKZ2_MOUSE]	0.30	0.02	0.32	0.11	0.16	0.08	0.25	-0.21	-0.22	-0.18	-0.25	-0.05	0.02
Q6P9S0	MTSS1-like protein OS=Mus musculus GN=Mtss1 PE=1 SV=1 - [MTSSL_MOUSE]	-0.15	0.07	0.02	0.03	-0.07	0.06	-0.11	-0.35	-0.07	-0.19	-0.08	-0.25	0.02
E9Q8D3	Protein Zmynd8 OS=Mus musculus GN=Zmynd8 PE=4 SV=2 - [E9Q8D3_MOUSE]	0.31	0.09	0.29	0.41	0.32	0.59	0.09	-0.07	0.25	-0.52	0.08	-0.35	0.02
Q99J83	Autophagy protein 5 OS=Mus musculus GN=Atg5 PE=1 SV=1 - [ATG5_MOUSE]	-0.04	-0.20	-0.70	-0.06	-0.16	-0.58	0.61	-0.20	0.48	0.04	0.79	0.01	0.02
K3W4Q8	Basigin OS=Mus musculus GN=Bsg PE=4 SV=1 - [K3W4Q8_MOUSE]	-0.35	-0.26	-0.08	-0.02	-0.08	-0.31	-0.04	0.05	0.05	-0.01	-0.06	0.13	0.02
Q9CR61	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Mus musculus GN=Ndufb7 PE=1 SV=3 - [NDUB7_MOUSE]	-0.25	-0.08	-0.08	-0.01	-0.03	0.16	0.24	0.31	0.09	0.27	0.04	0.04	0.02
Q3UHQ6-2	Isoform 2 of Protein dopey-2 OS=Mus musculus GN=Dopey2 [DOP2_MOUSE]	-0.31	-0.15	-0.36	-0.08	-0.24	-0.11	-0.04	0.08	0.01	0.23	-0.23	0.05	0.02
Q9CX34	Suppressor of G2 allele of SKP1 homolog OS=Mus musculus GN=Sugt1 PE=2 SV=3 - [SUGT1_MOUSE]	-0.16	-0.06	-0.08	-0.08	0.00	0.00	0.00	0.12	-0.04	0.05	0.06	0.19	0.02
Q9Z160	Conserved oligomeric Golgi complex subunit 1 OS=Mus musculus GN=Cog1 PE=2 SV=3 - [COG1_MOUSE]	0.42	0.40	0.13	-0.02	0.25	-0.06	0.08	-0.13	-0.08	-0.21	-0.02	-0.35	0.02
Q9JI19	Acidic fibroblast growth factor intracellular- binding protein OS=Mus musculus GN=Fibp PE=2 SV=1 - [FIBP_MOUSE]	-0.03	0.24	-0.17	-0.12	-0.20	-0.01	0.00	0.26	0.18	0.16	0.19	0.30	0.02
F6XC25	Coiled-coil and C2 domain-containing protein 1B (Fragment) OS=Mus musculus GN=Cc2d1b PE=4 SV=1 - [F6XC25_MOUSE]	-0.16	0.45	-0.11	0.21	0.09	0.05	-0.27	-0.45	-0.04	-0.13	-0.15	-0.51	0.02

Q8BMD8	Calcium-binding mitochondrial carrier protein SCaMC-1 OS=Mus musculus GN=Slc25a24 PE=2 SV=1 - [SCMC1_MOUSE]	0.14	0.28	0.02	-0.55	-0.26	-0.03	0.66	1.12	0.33	1.17	-0.15	0.74	0.02
Q62092	Neuron-specific protein family member 1 OS=Mus musculus GN=Nsg1 PE=2 SV=3 - [NSG1_MOUSE]	-0.35	-0.38	-0.41	-0.52	-0.16	0.12	-0.31	-0.02	0.20	0.23	0.22	0.26	0.02
Q9D819	Inorganic pyrophosphatase OS=Mus musculus GN=Ppa1 PE=1 SV=1 - [PPYR_MOUSE]	-0.32	-0.28	-0.01	-0.04	-0.19	-0.20	0.24	0.08	-0.08	-0.06	-0.08	0.13	0.02
Q91Z31	Polypyrimidine tract-binding protein 2 OS=Mus musculus GN=Ptbp2 PE=1 SV=2 - [PTBP2_MOUSE]	0.09	0.04	0.21	-0.06	0.07	0.19	0.14	-0.07	-0.19	-0.12	-0.17	-0.19	0.02
Q80YN3	Breast carcinoma-amplified sequence 1 homolog OS=Mus musculus GN=Bcas1 PE=1 SV=3 - [BCAS1_MOUSE]	-0.82	-0.48	-0.45	-0.08	-0.06	-0.37	-0.42	0.32	0.43	0.49	-0.10	0.24	0.02
Q8K2C7-2	Isoform 2 of Protein OS-9 OS=Mus musculus GN=Os9 - [OS9_MOUSE]	0.07	0.84	0.20	0.64	0.09	1.04	-0.21	0.26	-0.17	-0.08	-0.33	-0.09	0.02
P30999	Catenin delta-1 OS=Mus musculus GN=Ctnd1 PE=1 SV=2 - [CTND1_MOUSE]	0.15	-0.01	0.16	-0.08	0.03	-0.11	-0.07	-0.17	-0.12	-0.19	-0.10	-0.17	0.02
P42125	Enoyl-CoA delta isomerase 1, mitochondrial OS=Mus musculus GN=Eci1 PE=2 SV=2 - [EC11_MOUSE]	-0.18	-0.05	-0.39	0.08	-0.37	0.03	0.00	0.22	0.55	0.06	0.40	0.01	0.02
Q6PHS9-4	Isoform 4 of Voltage-dependent calcium channel subunit alpha-2/delta-2 OS=Mus musculus GN=Cacna2d2 - [CA2D2_MOUSE]	-0.30	-0.11	-0.24	-0.02	-0.14	0.03	0.10	-0.05	0.12	0.13	0.12	-0.08	0.02
A2A9X5	5'(3')-deoxyribonucleotidase, cytosolic type OS=Mus musculus GN=Nt5c PE=2 SV=1 - [A2A9X5_MOUSE]	0.60	0.19	-0.13	-0.03	0.20	0.38	-0.28	0.04	-0.12	-0.18	-0.23	-0.25	0.02
Q9R269	Periplakin OS=Mus musculus GN=Ppl PE=1 SV=1 - [PEPL_MOUSE]	0.33	1.77	0.46	-0.35	-0.07	-0.37	-0.21	-2.00	-0.27	-2.28	-0.44	-2.20	0.02
Q78RX3	Small integral membrane protein 12 OS=Mus musculus GN=Smim12 PE=3 SV=1 - [SIM12_MOUSE]	-0.40	-0.09	-0.25	-0.20	-0.16	-0.06	0.10	0.24	-0.07	-0.13	-0.08	0.10	0.02
Q811W2	Cytochrome P450 26B1 OS=Mus musculus GN=Cyp26b1 PE=1 SV=1 - [CP26B_MOUSE]	0.34	1.71	-0.32	0.90	0.47	0.70	0.00	-0.24	-0.11	-0.53	0.02	-1.16	0.02

F8VQD7	Receptor-type tyrosine-protein phosphatase gamma OS=Mus musculus GN=Ptprg PE=2 SV=1 - [F8VQD7_MOUSE]	-0.12	-0.43	-0.02	-0.46	0.08	-0.53	-0.13	0.19	0.13	0.32	0.06	0.11	0.02
P28571-1	Isoform GlyT-1A of Sodium- and chloride-dependent glycine transporter 1 OS=Mus musculus GN=Slc6a9 - [SC6A9_MOUSE]	-0.18	-0.04	-0.04	-0.09	-0.10	-0.20	0.06	-0.08	0.05	0.02	-0.09	0.08	0.02
Q9D7M1	Glucose-induced degradation protein 8 homolog OS=Mus musculus GN=Gid8 PE=2 SV=1 - [GID8_MOUSE]	0.13	0.43	0.00	0.19	0.06	0.63	0.13	-0.34	0.09	-0.38	-0.07	-0.36	0.02
Q0PHV7	Dapper homolog 3 OS=Mus musculus GN=Dact3 PE=1 SV=1 - [DACT3_MOUSE]	0.08	0.30	0.07	0.16	-0.05	0.17	-0.12	-0.24	0.00	-0.22	0.15	-0.26	0.02
E9Q8G3	Protein Srcap OS=Mus musculus GN=Srcap PE=2 SV=1 - [E9Q8G3_MOUSE]	-0.25	-0.42	-0.22	-0.35	-0.26	-0.15	-0.27	0.02	-0.03	0.39	-0.01	0.01	0.02
F6S6C3	Sentrin-specific protease 5 (Fragment) OS=Mus musculus GN=Senp5 PE=2 SV=1 - [F6S6C3_MOUSE]	-0.93	-0.01	-0.32	-0.17	-0.59	0.18	0.51	0.28	0.01	0.68	-0.05	0.23	0.02
Q58A65-2	Isoform 2 of C-Jun-amino-terminal kinase-interacting protein 4 OS=Mus musculus GN=Spag9 - [JIP4_MOUSE]	-0.07	-0.03	0.07	-0.10	0.02	0.06	0.01	-0.28	-0.17	-0.11	-0.14	-0.25	0.02
Q9WUN2	Serine/threonine-protein kinase TBK1 OS=Mus musculus GN=Tbk1 PE=1 SV=1 - [TBK1_MOUSE]	0.08	0.14	0.17	0.01	-0.05	0.10	-0.29	-0.27	0.03	-0.03	-0.01	-0.16	0.02
D3YVQ5	NAD-dependent protein deacetylase sirtuin-3 (Fragment) OS=Mus musculus GN=Sirt3 PE=2 SV=1 - [D3YVQ5_MOUSE]	0.70	0.42	0.38	0.39	0.26	0.60	0.63	0.09	-0.25	-0.14	-0.31	-0.13	0.02
Q9CZH7	Matrix-remodeling-associated protein 7 OS=Mus musculus GN=Mxa7 PE=1 SV=2 - [MXRA7_MOUSE]	-0.90	-0.17	-0.18	-0.42	-0.25	0.02	0.07	-0.07	0.06	0.35	0.00	0.32	0.02
A2CG49	Kalirin OS=Mus musculus GN=Kalrn PE=1 SV=1 - [KALRN_MOUSE]	0.03	0.84	0.03	0.30	-0.14	0.52	-0.38	0.05	-0.38	-0.39	-0.40	0.01	0.02
Q9R0A0	Peroxisomal membrane protein PEX14 OS=Mus musculus GN=Pex14 PE=1 SV=1 - [PEX14_MOUSE]	-0.32	-0.15	-0.09	-0.08	-0.34	-0.27	0.00	0.29	-0.16	0.13	-0.15	0.15	0.02
Q9D787	Peptidyl-prolyl cis-trans isomerase-like 2 OS=Mus musculus GN=Ppil2 PE=2 SV=2 - [PPIL2_MOUSE]	0.33	0.31	0.14	0.58	0.38	0.74	0.01	-0.03	0.29	-0.15	0.17	0.28	0.02

Q8BYM8	Probable cysteine-- tRNA ligase, mitochondrial OS=Mus musculus GN=Cars2 PE=2 SV=2 - [SYCM_MOUSE]	-0.05	-0.07	0.04	-0.10	-0.14	-0.29	0.07	0.37	0.16	0.14	0.04	-0.08	0.02
A2AHC3	Calmodulin-regulated spectrin-associated protein 1 OS=Mus musculus GN=Camsap1 PE=1 SV=1 - [CAMP1_MOUSE]	0.09	-0.15	0.12	0.04	0.06	-0.05	-0.23	-0.17	-0.10	-0.06	-0.10	-0.08	0.02
P51906	Excitatory amino acid transporter 3 OS=Mus musculus GN=Slc1a1 PE=1 SV=2 - [EAA3_MOUSE]	-0.70	-0.13	0.00	-0.18	0.06	-0.67	-0.02	0.15	0.34	0.07	0.26	0.31	0.02
Q8BY89-2	Isoform 2 of Choline transporter-like protein 2 OS=Mus musculus GN=Slc44a2 - [CTL2_MOUSE]	-0.17	-0.24	-0.15	-0.28	-0.18	-0.34	0.13	-0.04	0.50	-0.13	0.41	-0.06	0.02
Q61016	Guanine nucleotide- binding protein G(I)(G(S)G(O) subunit gamma-7 OS=Mus musculus GN=Gng7 PE=2 SV=2 - [GBG7_MOUSE]	-0.46	-0.15	-0.41	0.06	-0.17	-0.25	-0.12	-0.12	0.32	0.01	0.14	0.14	0.02
P60330	Separin OS=Mus musculus GN=Esp1 PE=1 SV=1 - [ESPL1_MOUSE]	0.66	-0.19	0.82	1.17	0.86	0.58	-0.30	0.19	-0.28	0.14	-0.19	0.46	0.02
Q8BV13	COP9 signalosome complex subunit 7b OS=Mus musculus GN=Cops7b PE=1 SV=1 - [CSN7B_MOUSE]	-0.15	-0.22	0.08	-0.21	-0.06	-0.14	0.32	0.34	-0.04	0.26	-0.06	0.03	0.02
P81122	Insulin receptor substrate 2 OS=Mus musculus GN=Irs2 PE=1 SV=2 - [IRS2_MOUSE]	0.19	0.14	-0.02	-0.04	-0.11	0.43	-0.23	-0.16	-0.11	-0.21	-0.22	-0.10	0.02
Q8BIV3	Ran-binding protein 6 OS=Mus musculus GN=Ranbp6 PE=2 SV=3 - [RNBP6_MOUSE]	0.37	0.11	1.06	0.06	0.12	0.01	0.00	-0.54	-0.05	-0.31	-0.10	-0.87	0.02
Q8VDP6	CDP-diacylglycerol-- inositol 3- phosphatidyltransferas e OS=Mus musculus GN=Cdipt PE=1 SV=1 - [CDIPT_MOUSE]	-0.07	-0.30	-0.91	-0.42	-0.88	-0.32	-0.04	-0.09	-0.01	0.11	0.03	-0.11	0.02
Q62178	Semaphorin-4A OS=Mus musculus GN=Sema4a PE=1 SV=2 - [SEM4A_MOUSE]	0.07	0.48	0.12	0.27	0.00	0.16	-0.13	-0.11	0.09	0.03	-0.06	-0.12	0.02
Q5F284	Phospholipid scramblase 3 (Fragment) OS=Mus musculus GN=Plscr3 PE=2 SV=1 - [Q5F284_MOUSE]	0.01	-0.13	-0.17	-0.14	-0.24	-0.18	0.20	-0.01	0.12	-0.11	0.12	-0.09	0.02
P35979	60S ribosomal protein L12 OS=Mus musculus GN=Rpl12 PE=1 SV=2 -[RL12_MOUSE]	0.05	-0.03	-0.01	-0.07	-0.15	-0.03	0.03	0.06	0.22	0.04	0.13	0.01	0.02

Q8BN59	La-related protein 6 OS=Mus musculus GN=Larp6 PE=1 SV=1 - [LARP6_MOUSE]	0.02	0.02	-0.13	0.13	0.00	0.13	-0.14	-0.02	-0.05	-0.08	-0.18	-0.17	0.02
E9QNR6	Protein AI837181 OS=Mus musculus GN=AI837181 PE=2 SV=1 - [E9QNR6_MOUSE]	0.01	-0.11	-0.12	-0.14	-0.16	-0.13	0.21	0.22	-0.15	0.17	0.06	0.01	0.02
P97384	Annexin A11 OS=Mus musculus GN=Anxa11 PE=1 SV=2 - [ANX11_MOUSE]	0.12	-0.15	-0.06	0.02	-0.12	-0.30	0.13	0.08	0.06	0.25	0.12	0.04	0.02
Q922G0	Solute carrier family 25 member 36 OS=Mus musculus GN=Slc25a36 PE=2 SV=1 - [S2536_MOUSE]	0.22	0.22	0.26	-0.08	0.01	-0.03	0.04	-0.59	-0.13	-0.51	-0.11	-0.84	0.02
P61215	Carbonic anhydrase- related protein 10 OS=Mus musculus GN=Ca10 PE=2 SV=1 - [CAH10_MOUSE]	0.21	0.31	0.02	0.42	0.21	0.46	-0.12	0.05	0.08	0.12	0.17	-0.01	0.02
A3KGR9	Minor histocompatibility antigen H13 (Fragment) OS=Mus musculus GN=H13 PE=2 SV=1 - [A3KGR9_MOUSE]	0.36	0.28	0.26	0.69	0.11	0.99	-0.08	0.15	-0.13	0.03	-0.06	0.17	0.02
Q8R1Y2	Uncharacterized protein C16orf45 homolog OS=Mus musculus PE=2 SV=1 - [CP045_MOUSE]	0.06	-0.19	-0.09	0.04	-0.02	-0.21	0.22	0.06	0.12	0.02	0.06	0.06	0.02
Q8R310-2	Isoform 2 of Transmembrane and coiled-coil domains protein 3 OS=Mus musculus GN=Tmcc3 - [TMCC3_MOUSE]	-0.53	-0.09	0.13	0.03	0.05	0.27	-0.15	-0.80	-0.35	-0.80	-0.16	-0.66	0.02
Q9EQ08	Heparan N-sulfatase OS=Mus musculus GN=Sgsh PE=2 SV=1 - [Q9EQ08_MOUSE]	-0.16	0.13	0.19	0.37	0.28	0.07	-0.30	-0.70	0.12	-0.28	0.02	-0.47	0.02
Q9CY45	N(6)-adenine-specific DNA methyltransferase 2 OS=Mus musculus GN=N6amt2 PE=2 SV=1 - [N6MT2_MOUSE]	0.19	0.13	0.08	0.25	-0.10	-0.20	0.99	0.60	0.15	0.32	0.75	0.21	0.02
Q8C0T9-2	Isoform 2 of Adenylate cyclase type 10 OS=Mus musculus GN=Adcy10 - [ADCYA_MOUSE]	-0.25	-0.59	-0.20	-0.45	-0.19	-0.93	0.03	0.45	0.02	0.37	-0.29	-0.32	0.02
Q61749	Translation initiation factor eIF-2B subunit delta OS=Mus musculus GN=EIF2b4 PE=2 SV=2 - [EI2BD_MOUSE]	0.17	0.14	0.31	0.16	0.15	0.08	0.05	0.03	-0.16	-0.33	0.00	0.11	0.02
P52785	Guanylyl cyclase GC-E OS=Mus musculus GN=Gucy2e PE=2 SV=2 - [GUC2E_MOUSE]	0.45	-0.91	-0.18	-0.50	0.10	-0.54	0.50	0.25	0.39	0.42	0.21	0.56	0.02

P52800	Ephrin-B2 OS=Mus musculus GN=Efnb2 PE=1 SV=1 - [EFNB2_MOUSE]	0.18	0.24	0.04	0.34	0.08	-0.05	-0.02	-0.35	0.06	-0.35	-0.10	0.01	0.02
O88630	Golgi SNAP receptor complex member 1 OS=Mus musculus GN=Gosr1 PE=1 SV=2 - [GOSR1_MOUSE]	0.37	1.40	0.08	0.69	0.10	0.25	0.25	-1.15	0.04	-1.29	0.10	-0.94	0.02
F8WGB3	Succinate dehydrogenase cytochrome b560 subunit, mitochondrial OS=Mus musculus GN=Sdhc PE=2 SV=1 - [F8WGB3_MOUSE]	-0.62	-0.33	-0.41	-0.76	-0.17	-0.21	0.05	0.41	-0.11	0.02	-0.41	-0.10	0.02
Q9D164-2	Isoform 2 of FXYD domain-containing ion transport regulator 6 OS=Mus musculus GN=Fxyd6 - [FXYD6_MOUSE]	0.16	-0.43	-0.16	-0.52	-0.01	-0.40	0.04	-0.04	0.17	0.31	0.18	0.13	0.02
A2AM57	Inversin OS=Mus musculus GN=Invs PE=4 SV=1 - [A2AM57_MOUSE]	1.06	-0.10	1.20	0.15	0.30	0.37	0.01	-0.04	-1.03	0.12	-1.33	-0.15	0.02
O09044	Synaptosomal-associated protein 23 OS=Mus musculus GN=Snap23 PE=1 SV=1 - [SNP23_MOUSE]	-0.30	-1.40	-0.25	-0.27	-0.07	-0.20	-0.21	0.95	0.05	0.71	-0.05	0.59	0.02
P56716	Oxygen-regulated protein 1 OS=Mus musculus GN=Rp1 PE=1 SV=2 - [RP1_MOUSE]	0.73	3.02	-0.07	0.49	0.09	0.04	-0.25	-3.35	0.11	-2.65	-0.12	-2.98	0.02
Q8R191	Synaptogyrin-3 OS=Mus musculus GN=Syngr3 PE=1 SV=1 - [SNG3_MOUSE]	0.03	-0.06	-0.06	-0.13	-0.23	-0.11	0.11	0.11	-0.01	0.07	-0.09	0.24	0.02
Q3UID0	SWI/SNF complex subunit SMARCC2 OS=Mus musculus GN=Smarcc2 PE=2 SV=1 - [Q3UID0_MOUSE]	0.08	0.09	0.28	0.09	0.06	0.06	0.06	0.06	-0.22	0.00	-0.10	-0.05	0.02
E9Q705	BolA-like protein 3 OS=Mus musculus GN=Bola3 PE=2 SV=1 - [E9Q705_MOUSE]	-0.47	-0.77	-0.50	-0.55	0.14	-0.31	-0.42	0.37	0.10	0.06	-0.01	0.12	0.02
Q9QY33	Tetraspanin-3 OS=Mus musculus GN=Tspan3 PE=1 SV=1 - [TSN3_MOUSE]	0.51	0.60	-0.07	-0.09	0.07	-0.21	-0.27	-0.79	-0.10	-0.72	0.07	-0.73	0.02
Q7TT50	Serine/threonine-protein kinase MRCK beta OS=Mus musculus GN=Cdc42bbp PE=1 SV=2 - [MRCKB_MOUSE]	0.14	0.15	0.34	0.03	0.00	-0.01	0.15	-0.11	-0.25	-0.20	-0.14	-0.09	0.02
Q8R1N4	NudC domain-containing protein 3 OS=Mus musculus GN=Nudcd3 PE=2 SV=3 - [NUDC3_MOUSE]	0.25	0.27	0.27	0.47	-0.02	0.04	0.15	-0.25	-0.16	-0.10	-0.05	0.10	0.02

Q2WF71	Leucine-rich repeat and fibronectin type III domain-containing protein 1 OS=Mus musculus GN=Lrn1 PE=1 SV=1 - [LRFN1_MOUSE]	0.00	0.67	-0.08	0.29	-0.05	0.67	-0.23	-0.51	-0.05	-0.27	0.06	-0.34	0.02
Q8VHI3	GDP-fucose protein O-fucosyltransferase 2 OS=Mus musculus GN=Poluf2 PE=1 SV=1 - [OFUT2_MOUSE]	0.15	0.22	0.08	0.01	-0.13	0.18	0.06	-0.27	-0.43	-0.06	-0.20	-0.08	0.02
Q9WVT6	Carbonic anhydrase 14 OS=Mus musculus GN=Ca14 PE=1 SV=1 - [CAH14_MOUSE]	-0.02	-0.24	-0.08	-0.43	-0.04	-0.38	0.19	0.05	0.23	-0.08	0.27	-0.17	0.02
O88792	Junctional adhesion molecule A OS=Mus musculus GN=F11r PE=1 SV=2 - [JAM1_MOUSE]	-0.09	-0.34	-0.16	-0.46	0.02	-0.57	0.03	-0.01	0.12	0.10	-0.01	-0.01	0.02
Q8C6M1	Ubiquitin carboxyl-terminal hydrolase 20 OS=Mus musculus GN=Usp20 PE=1 SV=1 - [UBP20_MOUSE]	0.85	-0.02	0.84	0.01	0.67	-0.06	0.01	-0.47	-0.32	-0.27	-0.03	-0.16	0.02
Q920N7	Synaptotagmin-12 OS=Mus musculus GN=Sy12 PE=2 SV=1 - [SYT12_MOUSE]	-0.16	-0.12	0.04	0.00	-0.07	-0.12	0.25	0.05	0.03	0.12	0.03	-0.05	0.02
Q6DI86-2	Isoform 2 of FAST kinase domain-containing protein 1 OS=Mus musculus GN=Fastk1 - [FAKD1_MOUSE]	0.53	1.27	1.06	0.68	0.87	0.80	-0.37	-0.07	-0.09	0.37	1.15	-0.43	0.02
P62849-2	Isoform 2 of 40S ribosomal protein S24 OS=Mus musculus GN=Rps24 - [RS24_MOUSE]	-0.17	0.44	0.13	0.37	-0.06	0.51	0.20	-0.19	-0.20	-0.66	-0.20	-0.31	0.02
E9PVV3	Potassium voltage-gated channel subfamily C member 1 OS=Mus musculus GN=Kcnc1 PE=2 SV=1 - [E9PVV3_MOUSE]	-0.34	-0.13	-0.11	-0.32	-0.07	-0.45	0.17	-0.29	0.20	0.10	0.09	-0.08	0.02
Q8BHK1	Magnesium transporter NIPA1 OS=Mus musculus GN=Nipa1 PE=1 SV=1 - [NIPA1_MOUSE]	-0.64	0.01	-1.56	-0.02	-1.35	-0.07	0.69	0.06	0.89	-0.06	0.30	0.07	0.02
Q9CY34	NEDD8-conjugating enzyme UBE2F OS=Mus musculus GN=Ube2f PE=1 SV=1 - [UB2FA_MOUSE]	-0.44	-0.56	-0.18	-0.42	-0.02	-0.54	-0.07	-0.17	0.03	-0.10	-0.05	-0.11	0.02
Q8C5V0	Spermatogenesis-associated protein 32 OS=Mus musculus GN=Spata32 PE=1 SV=1 - [SPT32_MOUSE]	-0.13	1.61	0.71	1.35	-0.02	1.07	0.09	-3.54	0.05	-2.77	-0.16	-3.82	0.02
A2AFQ9	Gem-associated protein 5 OS=Mus musculus GN=Gemin5 PE=2 SV=1 - [A2AFQ9_MOUSE]	0.08	-0.16	-0.04	0.02	-0.58	-0.15	-0.05	0.15	0.18	0.19	0.44	0.20	0.02

P63073	Eukaryotic translation initiation factor 4E OS=Mus musculus GN=Eif4e PE=1 SV=1 - [IF4E_MOUSE]	-0.45	-0.13	0.01	-0.05	0.00	-0.09	0.22	0.08	-0.05	0.37	0.08	0.09	0.02
G3UWA6	4F2 cell-surface antigen heavy chain OS=Mus musculus GN=Slc3a2 PE=4 SV=1 [G3UWA6_MOUSE]	-0.16	-0.16	-0.02	-0.08	-0.06	-0.24	-0.03	-0.04	0.02	-0.07	0.02	0.01	0.02
A2AKI5	Integrin alpha-V light chain OS=Mus musculus GN=Itgav PE=2 SV=1 - [A2AKI5_MOUSE]	-0.14	-0.20	-0.03	-0.10	-0.15	-0.25	0.05	-0.12	0.16	-0.03	-0.06	-0.03	0.02
A2ATL1	MCG17280 OS=Mus musculus GN=Fam171a1 PE=4 SV=1 - [A2ATL1_MOUSE]	0.21	0.18	-0.02	-0.06	-0.02	0.19	-0.20	-0.10	0.11	-0.16	-0.09	-0.26	0.02
Q9D6Y9	1,4-alpha-glucan-branching enzyme OS=Mus musculus GN=Gbe1 PE=2 SV=1 - [GLGB_MOUSE]	-0.07	0.02	-0.13	-0.08	-0.30	-0.20	0.05	0.03	-0.02	-0.05	0.09	0.01	0.02
Q8BMS4	Hexaprenyldihydroxybenzoate methyltransferase, mitochondrial OS=Mus musculus GN=Coq3 PE=2 SV=1 - [COQ3_MOUSE]	0.16	-0.05	0.15	0.51	-0.04	0.21	-0.04	0.03	-0.19	-0.29	0.03	-0.34	0.02
E9Q1N0	Protein 4932431P20Rik OS=Mus musculus GN=4932431P20Rik PE=2 SV=1 - [E9Q1N0_MOUSE]	0.90	0.11	0.18	-0.05	0.44	0.38	-0.12	-0.50	0.04	0.09	0.01	-0.29	0.02
Q8JZP9	GAS2-like protein 1 OS=Mus musculus GN=Gas2l1 PE=2 SV=1 - [GA2L1_MOUSE]	0.21	0.96	0.30	0.31	0.19	0.11	0.15	-0.45	0.15	-0.24	0.09	-0.30	0.02
Q80VQ0	Aldehyde dehydrogenase family 3 member B1 OS=Mus musculus GN=Aldh3b1 PE=2 SV=1 - [AL3B1_MOUSE]	-0.05	-0.30	-0.07	0.18	-0.06	-0.40	0.11	0.13	0.00	0.44	0.13	0.19	0.02
Q9CQB5	CDGSH iron-sulfur domain-containing protein 2 OS=Mus musculus GN=Cisd2 PE=1 SV=1 - [CISD2_MOUSE]	-0.27	-0.05	-0.09	-0.07	-0.27	0.03	0.16	0.14	-0.03	0.01	-0.06	0.08	0.02
P51150	Ras-related protein Rab-7a OS=Mus musculus GN=Rab7a PE=1 SV=2 - [RAB7A_MOUSE]	-0.06	0.06	-0.07	0.00	-0.03	-0.07	0.03	0.07	0.12	0.00	0.09	0.00	0.02
P70202	Latexin OS=Mus musculus GN=Lxn PE=1 SV=2 - [LXN_MOUSE]	-0.37	-0.26	-0.13	0.00	0.01	-0.21	0.01	0.06	0.06	-0.10	0.12	0.06	0.02
Q8JZM4	Delta and Notch-like epidermal growth factor-related receptor OS=Mus musculus GN=Dner PE=1 SV=1 - [DNER_MOUSE]	0.07	-0.36	-0.12	-0.41	-0.02	-0.62	0.06	0.24	-0.01	-0.06	0.12	0.23	0.02

A2AA86	SUZ domain-containing protein 1 OS=Mus musculus GN=Szrd1 PE=2 SV=1 -[A2AA86_MOUSE]	-0.11	-0.37	-0.43	-0.35	0.16	-0.44	-0.27	0.05	0.37	0.10	0.42	0.06	0.02
Q60967	Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthase 1 OS=Mus musculus GN=Paps1 PE=2 SV=1 [PAPS1_MOUSE]	0.34	0.03	0.21	0.13	0.09	0.19	-0.11	0.08	0.03	0.02	-0.08	0.13	0.02
L7N202	Uncharacterized protein OS=Mus musculus GN=Gm16477 PE=4 SV=1 [L7N202_MOUSE]	0.15	0.31	0.13	0.31	-0.09	0.34	0.18	-0.08	-0.21	-0.01	-0.11	-0.01	0.02
Q921M4-2	Isoform 2 of Golgin subfamily A member 2 OS=Mus musculus GN=Golga2 - [GOGA2_MOUSE]	-0.13	-0.06	-0.05	-0.05	0.06	0.01	-0.24	-0.33	-0.03	-0.34	-0.09	-0.17	0.02
Q3KNY0	Immunoglobulin-like and fibronectin type III domain-containing protein 1 OS=Mus musculus GN=Igf1 PE=1 SV=3 [IGFN1_MOUSE]	0.09	-0.47	0.06	-0.04	0.47	0.09	-0.20	-0.71	-0.24	-0.58	-0.06	-0.72	0.02
Q60972	Histone-binding protein RBBP4 OS=Mus musculus GN=Rbbp4 PE=1 SV=5 [RBBP4_MOUSE]	-0.13	-0.18	-0.22	-0.18	-0.24	-0.21	-0.32	0.16	0.04	0.27	0.18	0.12	0.02
O88572	Low-density lipoprotein receptor-related protein 6 OS=Mus musculus GN=Lrp6 PE=1 SV=1 [LRP6_MOUSE]	0.86	0.57	-0.37	0.39	0.06	0.40	-0.15	-0.58	-0.25	-0.16	-0.34	0.06	0.02
E9QNP0	KxDL motif-containing protein 1 OS=Mus musculus GN=Kxd1 PE=2 SV=1 [E9QNP0_MOUSE]	-1.49	-0.10	-0.58	-0.14	-0.19	-0.08	0.17	0.18	0.58	-0.14	0.69	0.23	0.02
P50586	Tubby protein OS=Mus musculus GN=Tub PE=1 SV=1 [TUB_MOUSE]	-0.10	0.24	-0.17	0.10	0.11	0.12	0.16	-0.97	-0.35	-0.73	-0.29	-1.03	0.02
Q4PJX1-3	Isoform 3 of Protein odr-4 homolog OS=Mus musculus GN=Odr4 - [ODR4_MOUSE]	-0.18	0.20	0.09	0.31	0.05	0.17	0.04	-0.67	-0.48	0.14	-0.48	-0.43	0.02
Q8BLV3	Sodium/hydrogen exchanger 7 OS=Mus musculus GN=Slc9a7 PE=2 SV=1 [SL9A7_MOUSE]	-0.23	-0.17	-0.27	-0.02	-0.07	-0.17	-0.21	0.03	0.11	0.07	-0.01	0.09	0.02
P84075	Neuron-specific calcium-binding protein hippocalcin OS=Mus musculus GN=Hpca PE=1 SV=2 [HPCA_MOUSE]	-0.12	-0.48	-0.20	0.09	0.06	-0.40	-0.11	0.46	0.29	0.04	0.01	0.40	0.02
Q9R062	Glycogenin-1 OS=Mus musculus GN=Gyg1 PE=2 SV=3 [GLYG_MOUSE]	-0.31	-0.17	0.05	-0.03	-0.03	-0.07	0.08	0.08	0.10	0.07	-0.06	0.18	0.02

Q6ZPF3-2	Isoform 2 of T-lymphoma invasion and metastasis-inducing protein 2 OS=Mus musculus GN=Tiam2 - [TIAM2_MOUSE]	0.32	0.27	0.59	-0.08	0.05	0.26	0.11	-0.11	-0.25	0.04	-0.12	-0.06	0.02
A3KG01	Eph receptor B2 OS=Mus musculus GN=Ephb2 PE=2 SV=2 - [A3KG01_MOUSE]	-0.02	-0.60	-0.23	-0.24	-0.09	-0.59	0.04	0.31	-0.04	0.00	-0.02	-0.10	0.02
E9PWP9	Protein 4931408C20Rik OS=Mus musculus GN=4931408C20Rik PE=4 SV=1 - [E9PWP9_MOUSE]	2.93	0.53	0.77	0.60	1.16	0.75	-0.19	-0.06	0.25	-0.14	-0.12	-0.08	0.02
Q5DTU0	Actin filament-associated protein 1-like 2 OS=Mus musculus GN=Afap12 PE=1 SV=2 - [AF1L2_MOUSE]	0.01	0.78	0.11	-0.01	0.20	0.39	0.11	-1.05	-0.56	-0.10	-0.03	-0.34	0.02
P58252	Elongation factor 2 OS=Mus musculus GN=Eef2 PE=1 SV=2 - [EF2_MOUSE]	-0.11	-0.06	0.04	0.00	-0.14	-0.09	0.18	0.10	-0.04	0.01	-0.01	0.12	0.02
O54991	Contactin-associated protein 1 OS=Mus musculus GN=Cntnap1 PE=1 SV=2 - [CNTP1_MOUSE]	-0.08	-0.18	-0.16	-0.11	-0.10	-0.19	0.21	0.01	-0.03	0.00	0.02	-0.15	0.02
Q9D2U5	LSM domain-containing protein 1 OS=Mus musculus GN=Lsm1 PE=2 SV=1 - [LSMD1_MOUSE]	-0.64	-0.30	-0.67	-0.33	-0.65	0.10	-0.18	0.36	-0.16	0.35	-0.34	0.34	0.02
P62869	Transcription elongation factor B polypeptide 2 OS=Mus musculus GN=Tceb2 PE=1 SV=1 - [ELOB_MOUSE]	-0.29	-0.03	-0.16	0.04	0.08	-0.01	0.24	0.16	0.13	-0.01	0.04	0.17	0.02
F6V6T4	Transmembrane emp24 domain-containing protein 2 (Fragment) OS=Mus musculus GN=Tmed2 PE=3 SV=1 - [F6V6T4_MOUSE]	0.08	0.11	0.19	0.03	0.00	0.26	0.20	-0.30	-0.20	-0.33	-0.10	-0.10	0.02
P61460	DEP domain-containing protein 5 OS=Mus musculus GN=Depdc5 PE=1 SV=2 - [DEPD5_MOUSE]	0.71	-0.02	0.11	0.12	0.13	0.06	0.04	-0.20	-0.25	-0.19	0.03	-0.45	0.02
O08966	Solute carrier family 22 member 1 OS=Mus musculus GN=Slc22a1 PE=2 SV=2 - [S22A1_MOUSE]	0.20	-1.03	-0.16	-0.91	-0.01	-1.02	-0.29	0.12	0.40	0.42	0.63	0.16	0.02
O08919	Numb-like protein OS=Mus musculus GN=Numbl PE=1 SV=3 - [NUMBL_MOUSE]	-0.23	-0.09	0.03	0.02	0.01	0.02	0.07	0.03	0.14	0.04	0.28	0.12	0.02

Q91VC9	Growth hormone-inducible transmembrane protein OS=Mus musculus GN=Ghitm PE=2 SV=1 - [GHITM_MOUSE]	0.25	0.81	0.08	0.30	-0.20	0.32	0.31	-0.49	-0.18	-0.65	-0.07	-0.54	0.03
Q60841-3	Isoform 3 of Reelin OS=Mus musculus GN=Reln - [RELN_MOUSE]	-0.09	0.15	-0.19	0.13	-0.21	0.01	0.44	0.86	0.31	2.01	0.35	0.71	0.03
Q7TMC8	L-fucose kinase OS=Mus musculus GN=Fuk PE=2 SV=1 - [Q7TMC8_MOUSE]	-0.34	-0.15	-0.14	0.04	-0.33	0.06	0.08	-0.07	0.20	-0.06	0.15	0.21	0.03
Q8C1B1	Calmodulin-regulated spectrin-associated protein 2 OS=Mus musculus GN=Camsap2 PE=1 SV=3 - [CAMP2_MOUSE]	0.16	0.02	0.20	-0.05	0.06	0.03	-0.10	-0.04	-0.12	-0.09	-0.10	0.09	0.03
B1AXN9	Ribosomal protein S6 kinase alpha-3 OS=Mus musculus GN=Rps6ka3 PE=2 SV=1 - [B1AXN9_MOUSE]	0.19	0.13	0.35	0.34	0.20	-0.07	0.18	0.04	-0.25	0.01	-0.06	-0.15	0.03
P97825	Hematological and neurological expressed 1 protein OS=Mus musculus GN=Hn1 PE=1 SV=3 - [HN1_MOUSE]	-0.07	0.34	-0.40	0.57	-0.03	0.06	-0.76	-0.99	0.06	-0.52	-0.04	-0.89	0.03
Q8BG75	Transmembrane protein 198 OS=Mus musculus GN=Tmem198 PE=2 SV=1 - [TM198_MOUSE]	0.45	0.24	0.21	0.40	-0.04	0.14	-0.17	0.06	0.01	-0.03	0.07	0.09	0.03
Q02780-5	Isoform 5 of Nuclear factor 1 A-type OS=Mus musculus GN=Nfia - [NFIA_MOUSE]	-0.21	-0.20	-0.20	-0.10	-0.15	-0.19	0.13	0.07	0.30	-0.03	0.84	0.06	0.03
Q8C9B9	Death-inducer obliterator 1 OS=Mus musculus GN=Dido1 PE=1 SV=4 - [DIDO1_MOUSE]	0.10	0.00	0.06	0.15	0.09	0.01	-0.28	0.02	-0.02	-0.11	0.01	-0.07	0.03
E9PYG6	Protein Rasa1 OS=Mus musculus GN=Rasa1 PE=2 SV=1 - [E9PYG6_MOUSE]	0.33	0.35	0.10	0.33	-0.02	0.10	-0.08	-0.12	0.01	0.12	0.05	-0.01	0.03
Q62433	Protein NDRG1 OS=Mus musculus GN=Ndr1 PE=1 SV=1 - [NDRG1_MOUSE]	-0.32	-0.11	0.03	-0.01	-0.14	-0.15	0.18	0.18	-0.04	-0.06	-0.02	0.17	0.03
E9Q8T1	Transforming acidic coiled-coil-containing protein 2 OS=Mus musculus GN=Tacc2 PE=2 SV=1 - [E9Q8T1_MOUSE]	-0.02	-1.39	-0.46	-0.05	-0.02	-0.57	-0.17	1.57	-0.21	1.61	0.17	1.28	0.03
Q923T9	Calcium/calmodulin-dependent protein kinase type II subunit gamma OS=Mus musculus GN=Camk2g PE=1 SV=1 - [KCC2G_MOUSE]	-0.09	-0.49	0.07	-0.36	-0.06	-0.59	-0.03	-0.15	0.33	0.36	0.32	-0.07	0.03

P22892	AP-1 complex subunit gamma-1 OS=Mus musculus GN=Ap1g1 PE=1 SV=3 - [AP1G1_MOUSE]	-0.31	-0.11	-0.14	-0.09	-0.21	-0.07	0.25	0.01	-0.03	-0.12	-0.06	0.02	0.03
A2ANP1	2310028H24Rik protein OS=Mus musculus GN=Fam219a PE=2 SV=1 - [A2ANP1_MOUSE]	-0.06	-0.14	-0.15	-0.21	0.14	-0.35	0.29	-0.11	0.23	0.39	-0.09	0.19	0.03
D3YU60	Microsomal glutathione S-transferase 1 OS=Mus musculus GN=Mgst1 PE=2 SV=1 - [D3YU60_MOUSE]	0.16	-0.60	0.04	-0.28	0.30	0.00	0.23	0.50	0.11	0.40	0.26	0.57	0.03
O70405	Serine/threonine-protein kinase ULK1 OS=Mus musculus GN=Ulk1 PE=1 SV=1 - [ULK1_MOUSE]	0.28	0.38	0.01	0.37	-0.07	0.31	-0.23	-0.02	-0.78	0.01	-0.24	0.09	0.03
Q91VU6	DDB1- and CUL4-associated factor 11 OS=Mus musculus GN=Dcaf11 PE=1 SV=1 - [DCA11_MOUSE]	0.16	0.31	0.14	0.22	0.16	0.01	0.15	-0.46	-0.09	-0.52	-0.05	-0.02	0.03
Q32NY4	Metal transporter CNNM3 OS=Mus musculus GN=Cnm3 PE=1 SV=2 - [CNNM3_MOUSE]	-0.25	-0.17	-0.22	0.05	-0.20	-0.55	0.13	-0.26	0.09	0.23	0.02	0.09	0.03
Q6PNC0	DmX-like protein 1 OS=Mus musculus GN=Dmx1 PE=1 SV=1 - [DMXL1_MOUSE]	0.21	0.27	-0.02	0.06	0.10	0.32	0.19	-0.07	-0.19	-0.34	0.04	-0.21	0.03
Q99J09	Methylome protein 50 OS=Mus musculus GN=Wdr77 PE=1 SV=1 - [MEP50_MOUSE]	0.28	-0.22	0.20	-0.21	0.11	-0.28	-0.43	-0.36	-0.17	-0.20	-0.43	-0.37	0.03
Q8BGV0	Probable asparagine--tRNA ligase, mitochondrial OS=Mus musculus GN=Nars2 PE=2 SV=1 - [SYNM_MOUSE]	0.46	0.70	0.24	0.35	-0.09	0.66	0.46	-0.09	-0.07	-0.22	-0.33	0.03	0.03
P62317	Small nuclear ribonucleoprotein Sm D2 OS=Mus musculus GN=Snrpd2 PE=1 SV=1 - [SMD2_MOUSE]	-0.15	-0.07	-0.06	-0.01	-0.07	0.06	-0.04	0.08	0.03	0.02	0.10	0.07	0.03
Q9QYJ0	DnaJ homolog subfamily A member 2 OS=Mus musculus GN=Dnaja2 PE=1 SV=1 - [DNJA2_MOUSE]	-0.21	-0.28	0.05	-0.14	-0.09	-0.08	0.30	0.10	-0.05	0.06	-0.09	0.09	0.03
D3Z5N5	1-phosphatidylinositol 3-phosphate 5-kinase OS=Mus musculus GN=Plkyve PE=2 SV=2 - [D3Z5N5_MOUSE]	0.16	0.16	0.00	0.34	0.10	-0.08	0.00	-0.15	-0.10	-0.07	-0.02	-0.06	0.03
Q80XC6	Protein NRDE2 homolog OS=Mus musculus GN=Nrde2 PE=2 SV=3 - [NRDE2_MOUSE]	-0.49	-0.64	-0.13	-0.36	-0.04	-0.37	-0.20	-0.05	0.15	-0.10	0.05	-0.18	0.03

O54962	Barrier-to-autointegration factor OS=Mus musculus GN=Banf1 PE=1 SV=1 -[BAF_MOUSE]	-0.96	-0.28	-0.74	-0.20	0.07	-0.08	-0.46	0.32	0.41	0.21	0.47	0.21	0.03
Q811S7	Upstream-binding protein 1 OS=Mus musculus GN=Ubp1 PE=1 SV=1 -[UBIP1_MOUSE]	-0.05	0.08	0.38	0.13	0.16	-0.07	0.10	-0.64	-0.26	-0.27	0.02	-0.41	0.03
Q9QYB1	Chloride intracellular channel protein 4 OS=Mus musculus GN=Clic4 PE=1 SV=3 -[CLIC4_MOUSE]	-0.26	-0.09	-0.10	0.00	-0.04	-0.04	-0.10	0.14	0.12	-0.05	0.11	0.18	0.03
Q9DCU2	Plasmalipin OS=Mus musculus GN=Plip PE=2 SV=1 -[PLLIP_MOUSE]	-0.14	-2.06	-0.06	-1.87	0.14	-1.93	-0.12	0.44	0.10	0.65	0.33	0.65	0.03
P0C8B4	Uncharacterized protein C14orf142 homolog OS=Mus musculus PE=2 SV=1 -[CN142_MOUSE]	-1.73	-1.93	-0.72	-0.56	0.03	-1.56	-0.70	-0.31	0.60	0.06	0.59	-0.55	0.03
P70445	Eukaryotic translation initiation factor 4E-binding protein 2 OS=Mus musculus GN=Eif4ebp2 PE=2 SV=1 -[4EBP2_MOUSE]	0.06	0.57	-0.10	0.62	0.07	0.31	-0.32	-1.14	0.26	-1.04	0.10	-1.15	0.03
P23116	Eukaryotic translation initiation factor 3 subunit A OS=Mus musculus GN=Elf3a PE=1 SV=5 -[EIF3A_MOUSE]	0.08	0.04	0.17	-0.03	-0.08	0.13	-0.20	-0.06	-0.07	-0.14	-0.01	0.01	0.03
Q91ZV7	Plexin domain-containing protein 1 OS=Mus musculus GN=Plxdc1 PE=1 SV=2 -[PLDX1_MOUSE]	0.24	0.25	-0.02	0.16	0.45	-0.04	0.15	-0.80	-0.07	-0.79	-0.06	-0.92	0.03
O88839-2	Isoform 2 of Disintegrin and metalloproteinase domain-containing protein 15 OS=Mus musculus GN=Adam15 -[ADA15_MOUSE]	-0.22	-0.31	-0.08	-0.35	0.01	-0.26	-0.05	0.13	-0.16	0.37	-0.10	0.17	0.03
P0C5E4	Phosphatidylinositol phosphatase PTPRQ OS=Mus musculus GN=Ptprq PE=1 SV=1 -[PTPRQ_MOUSE]	0.66	4.06	0.11	1.65	0.00	1.55	0.01	-5.76	0.35	-4.91	0.49	-5.50	0.03
Q9QYF9	Protein NDRG3 OS=Mus musculus GN=Ndr3 PE=1 SV=1 -[NDRG3_MOUSE]	0.03	-0.20	0.01	-0.14	-0.04	-0.29	-0.01	0.12	0.11	-0.02	0.10	0.03	0.03
Q9JJV9	Sodium channel protein type 5 subunit alpha OS=Mus musculus GN=Scn5a PE=1 SV=2 -[SCN5A_MOUSE]	-0.40	0.09	-0.33	-0.34	-0.20	-0.25	-0.11	0.54	-0.23	0.67	0.05	0.51	0.03
F8WI30	Sorting nexin-7 OS=Mus musculus GN=Snx7 PE=2 SV=1 -[F8WI30_MOUSE]	0.42	0.61	0.30	0.07	0.03	0.23	-0.02	-0.16	0.00	0.32	-0.07	-0.28	0.03

Q56A07	Sodium channel subunit beta-2 OS=Mus musculus GN=Scn2b PE=1 SV=1 [SCN2B_MOUSE]	-0.57	-0.23	-0.26	-0.01	0.01	-0.25	-0.24	0.08	0.49	-0.01	0.40	0.12	0.03
Q5SPX8	TBC1 domain family member 10A OS=Mus musculus GN=Tbc1d10a PE=2 SV=1 [Q5SPX8_MOUSE]	0.18	0.36	0.25	0.32	-0.03	0.27	0.26	-0.22	0.12	-0.10	-0.03	-0.07	0.03
Q9D0L7	Armadillo repeat-containing protein 10 OS=Mus musculus GN=Arm10 PE=1 SV=1 [ARM10_MOUSE]	0.06	0.23	0.18	0.02	0.12	0.34	-0.06	0.14	-0.33	0.06	-0.28	-0.01	0.03
Q91Z69	SLIT-ROBO Rho GTPase-activating protein 1 OS=Mus musculus GN=Srgap1 PE=1 SV=2 [SRGP1_MOUSE]	-0.02	0.28	0.19	0.26	-0.04	0.69	-0.12	-0.04	-0.14	-0.06	-0.24	-0.01	0.03
Q3UVX5	Metabotropic glutamate receptor 5 OS=Mus musculus GN=Grm5 PE=2 SV=2 [GRM5_MOUSE]	-0.08	-0.17	0.03	-0.13	-0.04	-0.25	-0.05	0.05	0.02	-0.01	0.01	0.09	0.03
Q61466-2	Isoform 1 of SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily D member 1 OS=Mus musculus GN=Smarcd1 [SMRD1_MOUSE]	0.63	0.20	0.22	0.09	0.12	0.32	0.08	-0.49	0.17	0.13	-0.16	-0.18	0.03
O35375-5	Isoform B0 of Neuropilin-2 OS=Mus musculus GN=Nrp2 [NRP2_MOUSE]	0.96	-0.15	0.26	0.08	-0.10	-0.08	-0.42	-0.22	-0.72	-0.38	-0.32	-0.10	0.03
Q7TMW6	Cytosolic Fe-S cluster assembly factor NARFL OS=Mus musculus GN=Narf1 PE=2 SV=2 [NARFL_MOUSE]	0.16	0.84	0.19	0.42	-0.24	0.43	0.09	-0.57	-0.05	-0.21	0.02	-0.25	0.03
Q6NVG1	Lysophospholipid acyltransferase LPCAT4 OS=Mus musculus GN=Lpcat4 PE=1 SV=1 [LPCT4_MOUSE]	0.27	0.54	0.08	0.12	0.18	0.74	0.11	0.15	-0.33	-0.12	-0.13	0.16	0.03
Q8BG51	Mitochondrial Rho GTPase 1 OS=Mus musculus GN=Rhot1 PE=2 SV=1 [MIRO1_MOUSE]	-0.23	0.01	-0.04	0.09	-0.07	0.19	0.36	0.30	0.05	0.18	0.07	0.16	0.03
Q3TWI9	Transmembrane protein 63B OS=Mus musculus GN=Tmem63b PE=1 SV=1 [TM63B_MOUSE]	-0.15	-0.31	0.05	-0.31	-0.06	-0.20	-0.07	-0.09	0.05	0.15	-0.05	0.22	0.03
Q8BHI4	Kelch repeat and BTB domain-containing protein 3 OS=Mus musculus GN=Kbtbd3 PE=2 SV=1 [KBTB3_MOUSE]	1.67	0.46	3.12	0.32	0.52	0.10	-0.71	-0.01	-0.69	-0.11	-0.60	-0.21	0.03

P70697	Uroporphyrinogen decarboxylase OS=Mus musculus GN=Urod PE=1 SV=2 - [DCUP_MOUSE]	0.00	-0.13	-0.05	-0.14	-0.04	-0.16	0.19	-0.02	0.06	0.15	0.42	-0.06	0.03
E9Q3B5	Uncharacterized protein OS=Mus musculus GN=Gm10774 PE=3 SV=1 [E9Q3B5_MOUSE]	-0.46	-0.16	-0.28	-0.57	-0.33	-0.24	0.13	-0.11	-0.24	0.36	-0.18	-0.21	0.03
E9QKS6	Synaptotagmin-like protein 2 OS=Mus musculus GN=Syt2 PE=2 SV=1 - [E9QKS6_MOUSE]	0.22	-0.43	0.41	-0.13	0.14	-0.42	0.39	0.38	0.26	0.53	0.32	0.46	0.03
Q8CIB6	Transmembrane protein 230 OS=Mus musculus GN=Tmem230 PE=1 SV=1 - [TM230_MOUSE]	0.30	0.14	0.16	0.04	0.17	0.47	-0.02	-0.35	0.32	-0.46	-0.06	-0.26	0.03
Q8K366	Exosc10 protein OS=Mus musculus GN=Exosc10 PE=2 SV=1 - [Q8K366_MOUSE]	0.14	0.78	0.44	0.63	0.42	0.74	0.46	0.11	0.31	-0.01	0.35	-0.09	0.03
Q3TTY5	Keratin, type II cytoskeletal 2 epidermal OS=Mus musculus GN=Krt2 PE=1 SV=1 - [K22E_MOUSE]	0.19	-2.01	-0.23	-2.53	0.15	-2.25	-0.28	0.30	-0.01	0.93	0.47	1.23	0.03
Q8CG03	cGMP-specific 3',5'-cyclic phosphodiesterase OS=Mus musculus GN=Pde5a PE=1 SV=2 - [PDE5A_MOUSE]	0.38	-0.46	-0.07	-0.32	-0.19	-0.42	-0.06	-1.06	-1.03	-0.75	-0.39	-1.06	0.03
Q8BT19	Phosphatidylinositol 4,5-bisphosphate 3-kinase catalytic subunit beta isoform OS=Mus musculus GN=Pk3cb PE=1 SV=2 - [PK3CB_MOUSE]	0.09	0.88	-0.05	0.37	0.10	0.22	0.20	-0.04	-0.33	-0.32	-0.03	-0.60	0.03
Q561M1	Acp1 protein OS=Mus musculus GN=Acp1 PE=2 SV=1 - [Q561M1_MOUSE]	0.44	0.94	0.21	0.78	-0.16	0.03	0.13	-0.18	-0.14	-0.04	-0.15	-0.53	0.03
P60766-1	Isoform 1 of Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 - [CDC42_MOUSE]	-0.95	0.02	-0.31	0.09	-0.12	-0.25	-0.15	0.18	0.48	0.16	0.49	0.14	0.03
Q9CQI7	U2 small nuclear ribonucleoprotein B'' OS=Mus musculus GN=Snrb2 PE=2 SV=1 - [RU2B_MOUSE]	0.38	0.03	-0.26	0.35	0.15	0.23	-0.43	-0.12	-0.06	-0.03	-0.08	-0.19	0.03
Q91Z49-2	Isoform 2 of UAP56-interacting factor OS=Mus musculus GN=Fytd1 - [UIF_MOUSE]	0.21	0.07	-0.14	-0.26	0.12	0.22	0.34	0.37	0.33	0.15	0.41	0.11	0.03
Q80Y17	Lethal(2) giant larvae protein homolog 1 OS=Mus musculus GN=Lgl1 PE=1 SV=1 - [L2GL1_MOUSE]	-0.04	-0.28	0.01	-0.08	0.01	-0.13	0.22	-0.07	0.02	0.27	0.17	-0.02	0.03

A2A7T8	Protein Zmym1 OS=Mus musculus GN=Zmym1 PE=2 SV=1 - [A2A7T8_MOUSE]	0.38	-0.05	0.75	-0.08	0.27	0.03	-0.51	-0.05	-0.24	-0.27	-0.09	0.08	0.03
P97429	Annexin A4 OS=Mus musculus GN=Anxa4 PE=2 SV=4 - [ANXA4_MOUSE]	0.16	-0.22	0.19	-0.19	0.06	-0.19	0.24	0.28	0.07	0.65	0.04	0.72	0.03
P59235	Nucleoporin Nup43 OS=Mus musculus GN=Nup43 PE=2 SV=2 - [NUP43_MOUSE]	-0.17	-0.13	-0.11	-0.36	-0.08	-0.99	0.10	0.10	0.15	0.17	0.26	-0.09	0.03
Q9JJK0	Calcipressin-3 OS=Mus musculus GN=Rcan3 PE=1 SV=1 - [RCAN3_MOUSE]	0.39	0.14	0.16	0.12	0.35	0.01	-0.10	-0.02	0.12	-0.03	0.14	-0.06	0.03
E9Q390	Myoferlin OS=Mus musculus GN=Myof PE=2 SV=2 - [E9Q390_MOUSE]	0.18	0.70	0.15	0.40	0.28	0.14	0.01	0.28	0.04	-0.07	0.10	-0.18	0.03
Q9QYC0-2	Isoform 2 of Alpha- adducin OS=Mus musculus GN=Add1 - [ADDA_MOUSE]	-1.06	-0.42	-0.21	-0.54	-0.03	-0.41	-0.56	0.13	0.06	0.46	0.00	0.27	0.03
Q9EQH2	Endoplasmic reticulum aminopeptidase 1 OS=Mus musculus GN=Erp1 PE=2 SV=2 - [ERAP1_MOUSE]	0.45	-0.21	-0.01	-0.24	0.15	-0.35	0.22	0.26	0.45	0.29	0.48	0.27	0.03
Q9D902	General transcription factor IIE subunit 2 OS=Mus musculus GN=Gtf2e2 PE=2 SV=2 - [T2EB_MOUSE]	0.01	1.40	-0.16	0.33	0.14	0.50	-0.61	-0.55	0.01	-0.33	0.11	-0.48	0.03
Q3URQ7-2	Isoform 2 of Methylenetetrahydrofolate synthase domain- containing protein OS=Mus musculus GN=Mthfsd - [MTHSD_MOUSE]	0.48	0.40	-0.06	0.00	0.02	-0.09	0.11	-0.43	-0.39	0.00	-0.47	-0.97	0.03
P97379	Ras GTPase-activating protein-binding protein 2 OS=Mus musculus GN=G3bp2 PE=1 SV=2 - [G3BP2_MOUSE]	0.00	0.18	-0.08	-0.05	0.01	0.11	-0.21	-0.32	-0.02	-0.30	0.06	-0.18	0.03
F8WHU9	Zinc finger protein ZPR1 (Fragment) OS=Mus musculus GN=Zfp259 PE=4 SV=1 - [F8WHU9_MOUSE]	-0.06	-0.04	-0.23	-0.26	-0.06	-0.03	0.18	0.02	-0.09	0.07	-0.01	0.00	0.03
P31650	Sodium- and chloride- dependent GABA transporter 3 OS=Mus musculus GN=Slc6a11 PE=1 SV=2 - [S6A11_MOUSE]	-0.10	-0.16	-0.16	0.00	-0.09	0.02	-0.10	0.03	0.12	-0.03	0.16	0.11	0.03
Q9JIQ3	Diablo homolog, mitochondrial OS=Mus musculus GN=Diablo PE=1 SV=2 - [DBLOH_MOUSE]	0.00	0.13	0.02	-0.12	-0.13	-0.28	-0.18	-0.23	-0.27	-0.35	-0.19	-0.20	0.03

Q9JIH2	Nuclear pore complex protein Nup50 OS=Mus musculus GN=Nup50 PE=1 SV=3 - [NUP50_MOUSE]	0.02	0.42	-0.01	0.27	0.04	0.07	-0.24	0.02	0.03	-0.18	0.02	-0.16	0.03
G5E881	Mitochondrial translocator assembly and maintenance protein 41 homolog OS=Mus musculus GN=Tamm41 PE=4 SV=1 - [G5E881_MOUSE]	-0.25	-0.11	-0.06	0.01	-0.01	-0.01	0.12	0.04	0.11	-0.04	0.15	-0.02	0.03
Q5F226	Protocadherin Fat 2 OS=Mus musculus GN=Fat2 PE=2 SV=1 - [FAT2_MOUSE]	0.90	0.20	0.48	0.27	0.29	0.26	0.38	-0.04	0.15	0.00	0.03	-0.06	0.03
P0C7N9	Proteasome assembly chaperone 4 OS=Mus musculus GN=Psmg4 PE=1 SV=1 - [PSMG4_MOUSE]	0.02	1.22	0.13	1.10	-0.06	0.13	-0.10	-2.82	0.17	-2.51	-0.12	-2.92	0.03
G3X9S1	Protein FAM13C OS=Mus musculus GN=Fam13c PE=4 SV=1 - [G3X9S1_MOUSE]	0.35	0.72	-0.29	0.87	0.02	0.35	-0.37	-1.63	0.29	-1.78	0.18	-1.54	0.03
J3QNW0	Cytosine-specific methyltransferase OS=Mus musculus GN=Dnmt1 PE=3 SV=1 - [J3QNW0_MOUSE]	-0.05	-0.15	0.17	-0.31	0.35	0.05	-0.84	-0.33	-0.71	-0.11	-0.18	-0.17	0.03
A2AU34	Protein Tmem74b OS=Mus musculus GN=Tmem74b PE=4 SV=1 - [A2AU34_MOUSE]	-1.37	-0.10	-1.85	0.21	-1.72	-0.01	0.14	0.47	-0.03	0.78	0.25	0.28	0.03
Q9DB29	Isoamyl acetate-hydrolyzing esterase 1 homolog OS=Mus musculus GN=lah1 PE=2 SV=1 - [IAH1_MOUSE]	-0.20	-0.32	0.06	-0.35	-0.04	-0.32	0.31	-0.08	-0.06	-0.02	0.03	0.01	0.03
Q8CIE2-2	Isoform 2 of Zinc finger MIZ domain-containing protein 2 OS=Mus musculus GN=Zmiz2 - [ZMIZ2_MOUSE]	0.06	0.39	0.24	0.07	0.14	0.02	-0.19	-0.33	0.21	-0.39	0.10	-0.25	0.03
Q8BHS3	Pre-mRNA-splicing factor RBM22 OS=Mus musculus GN=Rbm22 PE=2 SV=1 - [RBM22_MOUSE]	-0.12	0.41	-0.01	0.49	-0.02	0.73	-0.12	-0.23	-0.13	-0.20	-0.08	-0.22	0.03
Q9CYW4	Haloacid dehalogenase-like hydrolase domain-containing protein 3 OS=Mus musculus GN=Hhd3 PE=2 SV=1 - [HDHD3_MOUSE]	0.29	0.26	0.03	0.01	0.09	-0.13	0.07	-0.03	-0.15	-0.20	-0.14	-0.22	0.03
O88653	Regulator complex protein LAMTOR3 OS=Mus musculus GN=Lamtor3 PE=1 SV=1 - [LTOR3_MOUSE]	-0.06	-0.85	-0.32	-0.72	0.16	-0.94	-0.22	0.06	0.11	0.11	0.16	0.27	0.03

Q3UVK0	Endoplasmic reticulum metalloproteinase 1 OS=Mus musculus GN=Ermp1 PE=1 SV=2 - [ERMP1_MOUSE]	-0.13	0.32	0.17	0.25	0.23	0.13	-0.01	-0.17	0.09	-0.06	0.08	-0.12	0.03
O88196-3	Isoform 3 of E3 ubiquitin-protein ligase TTC3 OS=Mus musculus GN=Ttc3 - [TTC3_MOUSE]	0.38	-0.16	0.10	1.25	-0.02	0.52	-0.12	-0.71	-0.08	-0.66	-0.14	0.02	0.03
Q7TSV4	Phosphoglucomutase-2 OS=Mus musculus GN=Pgm2 PE=1 SV=1 - [PGM2_MOUSE]	-0.03	-0.20	-0.04	-0.15	-0.12	-0.16	0.21	0.18	-0.08	0.19	0.02	-0.14	0.03
Q8C341	SUN domain-containing ossification factor OS=Mus musculus GN=Suco PE=1 SV=3 - [SUCO_MOUSE]	-0.41	-0.29	-0.04	-0.07	-0.17	-0.31	-0.10	1.20	0.23	0.40	-0.07	1.07	0.03
F7BE84	Protein Ankrd50 (Fragment) OS=Mus musculus GN=Ankrd50 PE=2 SV=1 - [F7BE84_MOUSE]	0.23	0.10	0.10	-0.04	0.15	-0.01	0.10	-0.07	-0.03	-0.13	-0.05	-0.05	0.03
F7CK47	Microtubule-associated protein (Fragment) OS=Mus musculus GN=Map4 PE=4 SV=1 - [F7CK47_MOUSE]	0.14	0.39	-0.08	0.17	-0.08	0.42	-0.33	-0.55	0.20	-0.65	0.13	-0.57	0.03
F8VQF3	Polycystic kidney disease and receptor for egg jelly-related protein OS=Mus musculus GN=Pkdrej PE=4 SV=1 - [F8VQF3_MOUSE]	-0.21	4.23	-0.16	-0.20	0.38	-0.56	-0.85	-4.41	-0.13	-3.82	-0.06	-4.09	0.03
Q8R010	Angiotensin-converting enzyme 2 OS=Mus musculus GN=Ace2 PE=1 SV=1 - [ACE2_MOUSE]	-0.01	-0.08	0.16	-0.23	-0.11	-0.77	0.57	0.00	0.39	0.18	0.16	0.04	0.03
P46938-2	Isoform 2 of Yorkie homolog OS=Mus musculus GN=Yap1 - [YAP1_MOUSE]	0.06	0.29	-0.07	0.23	0.34	0.19	-0.24	-0.90	0.10	-0.43	0.18	-0.60	0.03
Q8VHX6-2	Isoform 2 of Filamin-C OS=Mus musculus GN=Flnc - [FLNC_MOUSE]	0.60	0.39	-0.23	0.48	0.10	0.30	0.05	-0.21	-0.35	0.12	-0.18	0.01	0.03
O70274	Protein tyrosine phosphatase type IVA 2 OS=Mus musculus GN=Ptp4a2 PE=1 SV=1 - [TP4A2_MOUSE]	-0.47	-0.10	-0.04	0.00	0.12	-0.01	0.17	0.53	0.10	0.16	0.03	0.17	0.03
P31648	Sodium- and chloride-dependent GABA transporter 1 OS=Mus musculus GN=Slc6a1 PE=1 SV=2 - [SC6A1_MOUSE]	-0.53	-0.44	-0.08	-0.16	-0.09	-0.51	-0.19	-0.05	-0.09	0.02	-0.14	0.23	0.03
Q99N28	Cell adhesion molecule 3 OS=Mus musculus GN=Cadm3 PE=1 SV=1 - [CADM3_MOUSE]	-0.17	-0.01	-0.06	0.01	-0.10	-0.30	0.06	0.05	0.05	-0.05	0.06	0.02	0.03

Q64324	Syntaxin-binding protein 2 OS=Mus musculus GN=Stxbp2 PE=2 SV=1 - [STXB2_MOUSE]	0.71	0.23	0.02	0.21	0.18	0.31	0.17	-0.22	0.11	-0.18	0.09	-0.06	0.03
E9PZ67	Calsequestrin (Fragment) OS=Mus musculus GN=Casq2 PE=2 SV=1 - [E9PZ67_MOUSE]	-0.38	0.49	-0.16	0.24	-0.11	0.19	-0.01	-0.32	-0.49	-0.37	-0.32	-0.54	0.03
O35852	Preprodynorphin (Fragment) OS=Mus musculus GN=Pdyn PE=2 SV=1 - [O35852_MOUSE]	0.31	0.79	-0.18	0.96	0.27	0.76	-0.06	-0.20	0.26	-0.41	0.17	0.03	0.03
P15105	Glutamine synthetase OS=Mus musculus GN=Glul PE=1 SV=6 - [GLNA_MOUSE]	-0.13	-0.04	0.03	0.03	-0.04	-0.03	0.16	0.02	0.00	0.12	0.05	-0.01	0.03
P35762	CD81 antigen OS=Mus musculus GN=Cd81 PE=1 SV=2 - [CD81_MOUSE]	0.01	-0.36	0.00	-0.74	0.12	-0.84	0.02	0.34	0.28	0.17	0.16	0.15	0.03
Q3V4B5	COMM domain-containing protein 6 OS=Mus musculus GN=Comm6 PE=2 SV=1 - [COMD6_MOUSE]	0.24	-0.01	0.24	-0.10	-0.02	-0.42	-0.17	-0.94	-0.04	-0.44	-0.32	-0.71	0.03
Q8R3P0	Aspartoacylase OS=Mus musculus GN=Aspa PE=1 SV=2 - [ACY2_MOUSE]	0.00	-0.21	-0.28	-0.04	-0.20	-0.28	-0.27	0.09	0.19	-0.03	0.25	0.13	0.03
D3Z640	C2 domain-containing protein 3 OS=Mus musculus GN=C2cd3 PE=2 SV=1 - [D3Z640_MOUSE]	0.29	-0.27	-0.69	0.01	0.06	0.04	0.08	0.37	0.30	0.14	0.62	0.41	0.03
F2Z4A3	Protein Fat1 OS=Mus musculus GN=Fat1 PE=2 SV=1 - [F2Z4A3_MOUSE]	0.63	0.56	0.14	0.68	-0.06	0.43	-0.26	0.31	-0.01	0.08	0.02	0.08	0.03
F6RH18	Neurexin-1 OS=Mus musculus GN=Nrxn1 PE=2 SV=2 - [F6RH18_MOUSE]	-0.19	-0.08	-0.14	0.04	0.02	-0.13	-0.10	0.11	0.03	0.01	0.08	0.15	0.03
Q80WG5	Leucine-rich repeat-containing protein 8A OS=Mus musculus GN=Lrrc8a PE=1 SV=1 - [LRC8A_MOUSE]	0.17	-0.25	-0.04	-0.06	-0.05	-0.13	0.21	0.10	0.12	0.11	-0.03	0.12	0.03
P97772	Metabotropic glutamate receptor 1 OS=Mus musculus GN=Grm1 PE=2 SV=2 - [GRM1_MOUSE]	-0.19	-0.19	-0.21	-0.01	-0.21	-0.32	0.10	-0.09	-0.08	0.17	-0.03	-0.19	0.03
Q3UUQ7	GPI inositol-deacylase OS=Mus musculus GN=Pgap1 PE=1 SV=3 - [PGAP1_MOUSE]	-0.06	-0.12	-0.27	-0.14	-0.58	-0.17	0.18	-0.13	0.33	0.10	0.38	-0.27	0.03
Q8BUH8	Sentrin-specific protease 7 OS=Mus musculus GN=Senp7 PE=2 SV=1 - [SENP7_MOUSE]	0.22	-0.15	0.19	-0.16	0.26	0.03	-0.23	-0.17	-0.07	-0.15	-0.09	-0.22	0.03
G3XA00	Glutamate receptor, metabotropic 4 OS=Mus musculus GN=Grm4 PE=3 SV=1 - [G3XA00_MOUSE]	-0.30	-0.10	-0.07	0.16	-0.04	0.02	0.05	0.17	0.19	0.82	0.15	0.29	0.03

Q62261-2	Isoform 2 of Spectrin beta chain, non-erythrocytic 1 OS=Mus musculus GN=Sptbn1 - [SPTB2_MOUSE]	-0.22	0.56	0.41	0.29	-0.02	0.32	-0.05	-0.15	-0.29	-0.24	-0.48	0.25	0.03
P62259	14-3-3 protein epsilon OS=Mus musculus GN=Ywhae PE=1 SV=1 - [1433E_MOUSE]	-0.07	-0.05	-0.18	0.06	0.01	0.00	0.00	0.09	0.05	0.07	0.05	0.13	0.03
E9Q1F5	Protein Myo5c OS=Mus musculus GN=Myo5c PE=2 SV=1 - [E9Q1F5_MOUSE]	0.34	0.89	0.03	0.22	0.31	0.14	-0.31	-1.98	0.21	-1.90	0.18	-1.77	0.03
Q6PAK3	Protein arginine N-methyltransferase 8 OS=Mus musculus GN=Prmt8 PE=2 SV=2 - [ANM8_MOUSE]	-0.16	-0.23	0.07	-0.04	-0.15	-0.20	0.06	0.10	-0.04	0.42	-0.04	0.06	0.03
Q99L12	Chloride channel CLIC-like protein 1 OS=Mus musculus GN=Clicc1 PE=1 SV=1 - [CLCC1_MOUSE]	0.02	0.20	0.29	0.12	0.17	0.26	0.01	-0.10	0.16	0.04	0.15	-0.09	0.03
E9Q2K2	Protein TANC2 (Fragment) OS=Mus musculus GN=Tanc2 PE=2 SV=1 - [E9Q2K2_MOUSE]	-0.03	0.16	0.19	0.20	-0.07	-0.10	-0.17	-0.75	0.12	-0.60	-0.15	-0.86	0.03
F6WM75	Coiled-coil domain-containing protein 167 (Fragment) OS=Mus musculus GN=Ccdc167 PE=2 SV=1 - [F6WM75_MOUSE]	-0.25	-0.15	-0.34	-0.06	-0.47	-0.48	0.14	0.37	0.40	0.62	-0.34	-0.24	0.03
E9Q3H7	Protein Dnah6 OS=Mus musculus GN=Dnah6 PE=2 SV=1 - [E9Q3H7_MOUSE]	0.42	0.67	0.63	0.69	-0.19	0.87	-1.08	-0.02	-0.32	0.30	-0.37	0.47	0.03
Q99NE5-7	Isoform 7 of Regulating synaptic membrane exocytosis protein 1 OS=Mus musculus GN=Rims1 - [RIMS1_MOUSE]	0.12	-0.20	0.33	-0.02	0.09	-0.26	-0.41	-0.10	-0.31	-0.19	-0.27	-0.22	0.03
F8VQ40	Laminin subunit alpha-1 OS=Mus musculus GN=Lama1 PE=4 SV=1 - [F8VQ40_MOUSE]	0.09	0.42	0.10	0.51	0.20	0.54	0.13	0.19	0.00	0.08	-0.05	0.06	0.03
Q8CBM2	Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1 - [Q8CBM2_MOUSE]	-0.34	-0.24	-0.31	-0.19	-0.41	-0.17	0.25	-0.11	-0.31	0.10	-0.17	0.29	0.03
Q8CH09	SURP and G-patch domain-containing protein 2 OS=Mus musculus GN=Supp2 PE=2 SV=2 - [SUGP2_MOUSE]	0.22	0.08	0.24	-0.14	-0.13	0.13	0.09	-0.18	-0.22	-0.16	-0.13	-0.32	0.03
P35991	Tyrosine-protein kinase BTK OS=Mus musculus GN=Btk PE=1 SV=4 - [BTK_MOUSE]	-0.40	-0.59	-0.01	-0.49	-0.22	-0.46	0.50	0.52	-0.43	0.14	-0.33	0.44	0.03

Q810C1	SLIT and NTRK-like protein 1 OS=Mus musculus GN=Slitrk1 PE=2 SV=1 - [SLIK1_MOUSE]	-0.11	0.31	-0.07	0.01	-0.04	0.13	-0.03	-0.76	-0.04	-0.77	-0.10	-0.63	0.03
Q9Z2R6	Protein unc-119 homolog A OS=Mus musculus GN=Unc119 PE=1 SV=1 - [U119A_MOUSE]	-0.19	0.14	0.08	0.20	0.01	0.21	0.01	-0.14	-0.38	0.02	-0.23	-0.10	0.03
H3BJZ7	Protein unc-13 homolog B OS=Mus musculus GN=Unc13a PE=2 SV=1 - [H3BJZ7_MOUSE]	0.19	-0.04	0.27	0.17	0.07	-0.01	0.13	-0.06	-0.15	-0.11	-0.16	0.03	0.03
Q8BUE4	Apoptosis-inducing factor 2 OS=Mus musculus GN=Aifm2 PE=2 SV=1 - [AIFM2_MOUSE]	0.27	0.56	0.59	0.00	0.00	0.26	0.08	-0.12	0.26	-0.36	0.01	-0.26	0.03
Q9JJZ8	Protein flightless-1 homolog OS=Mus musculus GN=Flil PE=1 SV=1 - [FLII_MOUSE]	0.04	-0.17	0.17	-0.06	-0.08	-0.08	-0.08	-0.17	-0.31	-0.20	-0.33	-0.07	0.04
Q61285	ATP-binding cassette sub-family D member 2 OS=Mus musculus GN=Abcd2 PE=1 SV=1 - [ABCD2_MOUSE]	0.95	0.23	0.63	0.10	0.28	0.19	0.30	-0.15	-0.19	0.10	0.02	-0.01	0.04
E9QAB6	Protein Ptar1 OS=Mus musculus GN=Ptar1 PE=4 SV=1 - [E9QAB6_MOUSE]	-0.14	-0.19	-0.07	-0.02	0.17	-0.12	0.23	0.09	0.72	0.29	0.68	-0.11	0.04
Q99K01-2	Isoform 2 of Pyridoxal-dependent decarboxylase domain-containing protein 1 OS=Mus musculus GN=Pdxdc1 - [PDXD1_MOUSE]	0.15	0.00	0.24	-0.08	0.18	-0.01	0.06	-0.13	-0.21	-0.06	-0.25	0.00	0.04
Q8C7K6	Prenylcysteine oxidase-like OS=Mus musculus GN=Pcyox1l PE=2 SV=1 - [PCYXL_MOUSE]	-0.24	-0.02	-0.16	-0.17	-0.21	-0.14	0.05	0.10	-0.08	-0.06	-0.16	-0.06	0.04
Q8BGZ2-2	Isoform 2 of Protein FAM168A OS=Mus musculus GN=Fam168a - [F168A_MOUSE]	0.09	-0.12	-0.01	-0.30	0.30	-0.43	-0.13	0.19	0.55	0.23	0.60	0.33	0.04
P58242	Acid sphingomyelinase-like phosphodiesterase 3b OS=Mus musculus GN=Smpdl3b PE=1 SV=1 - [ASM3B_MOUSE]	0.06	-0.53	-0.03	0.01	-0.19	-0.17	0.05	1.10	0.02	0.96	-0.07	0.74	0.04
B0QZM1	Ubiquitin-conjugating enzyme E2 J2 OS=Mus musculus GN=Ube2j2 PE=2 SV=1 - [B0QZM1_MOUSE]	0.54	-0.30	0.15	0.25	0.11	-0.11	-0.02	-0.42	-0.17	-0.92	0.01	-0.74	0.04
Q08890	Iduronate 2-sulfatase OS=Mus musculus GN=Ids PE=2 SV=3 - [IDS_MOUSE]	0.03	-0.71	-0.15	-0.63	-0.23	-0.85	0.42	-0.10	-0.10	-0.08	-0.05	-0.12	0.04

D3Z030	Leucine-rich repeat-containing protein 16A OS=Mus musculus GN=Lrnc16a PE=2 SV=2 - [D3Z030_MOUSE]	0.23	0.01	0.29	0.16	0.01	-0.24	-0.17	-0.23	-0.15	0.04	-0.28	-0.11	0.04
P58158	Galactosylgalactosylxyloprotein 3-beta-glucuronosyltransferase 3 OS=Mus musculus GN=B3gat3 PE=2 SV=1 - [B3GA3_MOUSE]	0.94	0.17	0.49	-0.10	0.30	0.15	0.13	-0.11	-0.24	0.02	-0.20	-0.12	0.04
F8VPR5	CREB-binding protein OS=Mus musculus GN=Crebbp PE=2 SV=1 - [F8VPR5_MOUSE]	0.39	0.60	0.23	0.22	0.29	0.64	-0.42	-0.69	0.54	0.00	0.10	-0.16	0.04
Q03141-3	Isoform 3 of MAP/microtubule affinity-regulating kinase 3 OS=Mus musculus GN=Mark3 - [MARK3_MOUSE]	0.10	0.05	0.18	0.04	0.06	0.08	0.09	-0.10	-0.06	-0.01	-0.01	-0.29	0.04
Q8BWR8	Rhopilin-2 OS=Mus musculus GN=Rhpn2 PE=1 SV=2 - [RHPN2_MOUSE]	0.46	1.60	-0.13	1.48	-0.25	0.85	0.38	-2.35	0.27	-1.26	0.13	-2.04	0.04
G3X9U9	Fission 1 (Mitochondrial outer membrane) homolog (Yeast), isoform CRA_c OS=Mus musculus GN=Fis1 PE=4 SV=1 - [G3X9U9_MOUSE]	-0.25	0.00	-0.05	-0.06	0.11	0.07	0.17	0.30	0.03	0.06	0.05	0.18	0.04
Q8BGB5	LIM domain-containing protein 2 OS=Mus musculus GN=Limd2 PE=2 SV=1 - [LIMD2_MOUSE]	-0.05	0.90	-0.12	1.14	0.16	1.13	-0.62	0.04	0.06	-0.37	-0.03	-0.06	0.04
A6H5Y3	Methionine synthase OS=Mus musculus GN=Mtr PE=2 SV=1 - [METH_MOUSE]	-0.08	0.50	-0.02	0.61	0.00	0.37	-0.01	-0.29	-0.13	0.11	-0.15	-0.29	0.04
E9QPB2	Suprabasin OS=Mus musculus GN=Sbsn PE=4 SV=1 - [E9QPB2_MOUSE]	0.07	-0.30	-0.46	-0.92	0.04	-0.01	-0.52	0.50	0.36	1.59	0.46	0.99	0.04
Q3TCJ1	BRISC complex subunit Abro1 OS=Mus musculus GN=Fam175b PE=2 SV=1 - [F175B_MOUSE]	0.65	0.14	0.11	0.12	0.25	0.14	0.14	0.03	-0.09	0.04	-0.11	-0.04	0.04
Q8R3L2-4	Isoform 4 of Transcription factor 25 OS=Mus musculus GN=Tcf25 - [TCF25_MOUSE]	0.41	0.17	-0.03	0.27	0.00	0.24	-0.10	0.03	-0.04	0.02	-0.05	0.03	0.04
Q5SNZ0-2	Isoform 2 of Girdin OS=Mus musculus GN=Ccdc88a - [GRDN_MOUSE]	0.02	0.01	0.25	0.07	-0.03	0.14	-0.11	-0.06	-0.27	0.08	-0.31	0.02	0.04
Q3UV17	Keratin, type II cytoskeletal 2 oral OS=Mus musculus GN=Krt76 PE=2 SV=1 - [K22O_MOUSE]	-0.02	-1.06	0.24	-1.33	0.12	-0.81	-0.34	0.31	0.15	0.31	0.57	1.40	0.04

F8WJA7	AF4/FMR2 family member 3 OS=Mus musculus GN=Aff3 PE=2 SV=1 - [F8WJA7_MOUSE]	4.75	0.36	2.01	0.39	2.48	0.48	-0.50	0.29	-0.47	0.04	-0.91	0.12	0.04
P84084	ADP-ribosylation factor 5 OS=Mus musculus GN=Arf5 PE=2 SV=2 - [ARF5_MOUSE]	0.02	-0.28	0.21	-0.12	-0.16	-0.33	0.39	0.40	-0.10	0.24	-0.11	0.49	0.04
Q9DBG5	Perilipin-3 OS=Mus musculus GN=Plin3 PE=1 SV=1 - [PLIN3_MOUSE]	-0.33	0.05	-0.32	-0.09	-0.06	0.08	0.09	-0.06	0.16	0.10	0.17	0.09	0.04
Q9WUL7	ADP-ribosylation factor-like protein 3 OS=Mus musculus GN=Arf3 PE=1 SV=1 - [ARL3_MOUSE]	0.02	-0.07	-0.09	-0.01	-0.23	0.09	0.06	0.14	0.08	0.03	0.10	0.05	0.04
F8VQJ3	Laminin subunit gamma-1 OS=Mus musculus GN=Lamc1 PE=2 SV=1 - [F8VQJ3_MOUSE]	0.11	-0.02	0.51	0.17	0.18	0.13	-0.28	-0.12	-0.56	0.32	-0.42	-0.06	0.04
Q91YL2	E3 ubiquitin-protein ligase RNF126 OS=Mus musculus GN=Rnf126 PE=1 SV=1 - [RN126_MOUSE]	-0.45	-0.08	-0.39	-0.24	-0.05	-0.18	-0.45	0.29	0.30	0.73	0.47	0.04	0.04
D3Z7E5	Glycogen synthase kinase-3 alpha OS=Mus musculus GN=Gsk3a PE=2 SV=1 - [D3Z7E5_MOUSE]	-0.17	-0.01	0.12	0.02	0.00	0.03	0.28	0.38	-0.08	0.16	0.12	0.27	0.04
Q3U0D9	E3 ubiquitin-protein ligase HACE1 OS=Mus musculus GN=Hace1 PE=1 SV=1 - [HACE1_MOUSE]	0.09	-0.52	-0.14	-0.30	-0.45	-0.34	0.21	0.08	-0.03	-0.36	0.09	0.14	0.04
A2A5R8	Double-stranded RNA-binding protein Staufen homolog 1 OS=Mus musculus GN=Stau1 PE=2 SV=1 - [A2A5R8_MOUSE]	0.18	0.00	-0.04	-0.10	0.18	-0.04	-0.09	-0.17	-0.09	-0.35	-0.03	-0.08	0.04
Q80UU9	Membrane-associated progesterone receptor component 2 OS=Mus musculus GN=Pgrmc2 PE=1 SV=2 - [PGR2_MOUSE]	0.04	-0.26	0.02	-0.05	-0.27	-0.17	0.34	0.01	-0.05	-0.02	0.06	0.13	0.04
Q9CYN2	Signal peptidase complex subunit 2 OS=Mus musculus GN=Spcc2 PE=2 SV=1 - [SPCS2_MOUSE]	0.23	0.15	-0.09	0.01	0.03	0.18	-0.02	-1.09	-0.16	-0.42	-0.03	-0.59	0.04
Q91WM1-2	Isoform 2 of Spermatid perinuclear RNA-binding protein OS=Mus musculus GN=Strbp - [STRBP_MOUSE]	0.07	0.23	0.11	-0.02	-0.03	0.11	0.13	-0.06	-0.12	-0.09	-0.09	-0.07	0.04
Q7TMB8	Cytoplasmic FMR1-interacting protein 1 OS=Mus musculus GN=Cyflp1 PE=1 SV=1 - [CYFP1_MOUSE]	0.09	-0.20	0.04	-0.20	-0.19	-0.16	0.22	0.03	-0.02	0.09	-0.02	0.04	0.04

O08582	GTP-binding protein 1 OS=Mus musculus GN=Gtpbp1 PE=1 SV=2 - [GTPB1_MOUSE]	0.15	0.01	0.18	0.15	0.10	0.06	0.17	-0.05	-0.08	-0.07	0.04	-0.03	0.04
Q99NF3	Centrosomal protein of 41 kDa OS=Mus musculus GN=Cep41 PE=1 SV=1 - [CEP41_MOUSE]	-0.08	-0.38	0.23	-0.12	0.04	-0.10	0.06	0.45	0.15	0.13	0.17	0.09	0.04
Q64314	Hematopoietic progenitor cell antigen CD34 OS=Mus musculus GN=Cd34 PE=1 SV=1 - [CD34_MOUSE]	-0.13	-0.12	-0.31	-0.01	0.04	-0.20	-0.14	0.09	0.58	0.05	0.26	0.12	0.04
O54804-2	Isoform 2 of Choline kinase alpha OS=Mus musculus GN=Chka - [CHKA_MOUSE]	0.24	-0.05	0.19	0.15	0.18	0.30	-0.23	-0.95	-0.12	-0.31	-0.11	0.17	0.04
Q8CIG8	Protein arginine N- methyltransferase 5 OS=Mus musculus GN=Prmt5 PE=1 SV=3 -[ANM5_MOUSE]	0.12	-0.02	0.10	0.07	0.01	0.07	-0.07	0.01	-0.21	0.04	-0.09	0.01	0.04
Q9Z2D3	Non-syndromic hearing impairment protein 5 homolog OS=Mus musculus GN=Dfna5 PE=2 SV=1 - [DFNA5_MOUSE]	-0.27	-0.58	-0.13	-0.22	0.01	-0.15	-0.05	-0.04	0.07	0.02	0.00	0.02	0.04
Q6P5H2	Nestin OS=Mus musculus GN=Nes PE=1 SV=1 - [NEST_MOUSE]	-0.09	-1.46	-0.21	-2.27	0.07	-1.34	-0.63	0.37	-0.07	0.72	-0.05	1.96	0.04
O35598	Disintegrin and metalloproteinase domain-containing protein 10 OS=Mus musculus GN=Adam10 PE=1 SV=2 - [ADA10_MOUSE]	0.09	-0.28	-0.02	-0.20	-0.11	-0.42	0.01	0.03	0.10	0.04	0.21	-0.06	0.04
Q920N2	Biotin--protein ligase OS=Mus musculus GN=Hlcs PE=1 SV=1 - [BPL1_MOUSE]	-0.61	-0.32	-0.96	-0.14	-0.03	0.08	0.53	0.30	-0.06	0.12	0.08	-0.15	0.04
P70295	Ancient ubiquitous protein 1 OS=Mus musculus GN=Aup1 PE=2 SV=1 - [AUP1_MOUSE]	0.10	0.06	0.05	-0.03	-0.09	-0.05	0.04	-0.28	-0.07	-0.25	-0.09	-0.15	0.04
Q3UHH1-2	Isoform 2 of Zinc finger SWIM domain- containing protein 8 OS=Mus musculus GN=Zswim8 - [ZSWM8_MOUSE]	0.03	-0.24	0.16	-0.19	0.16	-0.06	0.60	0.40	0.03	0.28	-0.07	0.47	0.04
Q9D115	Methylmalonyl-CoA epimerase, mitochondrial OS=Mus musculus GN=Mcee PE=2 SV=1 - [MCEE_MOUSE]	0.00	0.02	-0.14	0.05	0.06	-0.02	-0.05	-0.17	0.04	-0.29	-0.16	-0.26	0.04
F8WGW3	S1 RNA-binding domain-containing protein 1 OS=Mus musculus GN=Srbd1 PE=2 SV=1 - [F8WGW3_MOUSE]	-0.49	-0.68	-0.05	-0.63	0.26	-0.42	-0.38	0.21	0.28	0.37	0.02	0.20	0.04

Q3TN34	JRAB OS=Mus musculus GN=Mical2 PE=1 SV=1 - [Q3TN34_MOUSE]	0.39	0.07	-0.04	0.57	-0.11	0.05	-0.20	-0.81	-0.15	0.00	-0.15	-1.32	0.04
P46660	Alpha-internexin OS=Mus musculus GN=Ina PE=1 SV=2 - [AINX_MOUSE]	-0.18	0.03	-0.13	-0.04	-0.05	0.03	0.04	-0.05	0.13	0.04	0.13	0.01	0.04
E9Q9C3	Afadin OS=Mus musculus GN=Mlt4 PE=2 SV=1 - [E9Q9C3_MOUSE]	-0.10	0.03	0.21	-0.06	0.03	-0.02	-0.15	-0.16	-0.07	-0.06	-0.15	-0.04	0.04
A2AFI8	RalBP1-associated Eps domain-containing protein 2 OS=Mus musculus GN=Reps2 PE=2 SV=1 - [A2AFI8_MOUSE]	-0.04	0.02	0.01	-0.02	0.10	0.06	-0.12	-0.14	-0.05	-0.01	-0.01	0.00	0.04
Q3TQQ9-3	Isoform 3 of Uncharacterized protein C1orf112 homolog OS=Mus musculus - [CA112_MOUSE]	-0.49	1.16	-0.31	0.11	0.12	0.12	1.25	0.40	1.30	2.02	1.80	-0.12	0.04
Q5DTT2	PH and SEC7 domain-containing protein 1 OS=Mus musculus GN=Psd PE=1 SV=2 - [PSD1_MOUSE]	0.57	-0.16	0.24	0.00	0.13	-0.05	-0.05	-0.30	-0.32	-0.61	-0.14	0.08	0.04
Q62261	Spectrin beta chain, non-erythrocytic 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2 - [SPTB2_MOUSE]	-0.02	0.09	0.31	0.00	0.01	0.07	-0.14	0.03	-0.27	0.02	-0.29	-0.01	0.04
Q3U1F9	Phosphoprotein associated with glycosphingolipid-enriched microdomains 1 OS=Mus musculus GN=Pag1 PE=1 SV=2 - [PHAG1_MOUSE]	-0.09	-0.69	-0.13	-0.54	0.01	-1.09	-0.29	-0.18	0.28	0.60	0.10	0.09	0.04
Q8BYM5	Neuroigin-3 OS=Mus musculus GN=Nlgn3 PE=1 SV=2 - [NLGN3_MOUSE]	0.05	0.03	0.23	0.07	-0.08	0.04	0.06	-0.21	0.04	-0.24	-0.07	-0.27	0.04
Q9QYB5-2	Isoform 1 of Gamma-adducin OS=Mus musculus GN=Add3 - [ADDG_MOUSE]	0.04	-0.71	0.07	-0.34	0.18	-0.40	-0.08	0.23	0.10	0.38	0.30	0.23	0.04
Q8BL48-2	Isoform 2 of RING finger protein unkempt homolog OS=Mus musculus GN=Unk - [UNK_MOUSE]	-0.43	-0.07	-0.15	-0.28	0.17	-0.41	-0.03	0.04	0.39	0.01	0.14	-0.07	0.04
Q99LP6	GrpE protein homolog 1, mitochondrial OS=Mus musculus GN=Grpe1 PE=1 SV=1 - [GRPE1_MOUSE]	-0.25	-0.12	-0.02	-0.12	-0.17	-0.03	0.06	0.36	0.06	0.03	-0.18	0.18	0.04
Q91YJ2	Sorting nexin-4 OS=Mus musculus GN=Snx4 PE=2 SV=1 - [SNX4_MOUSE]	0.47	0.14	0.31	-0.03	-0.03	0.00	0.22	-0.21	-0.25	-0.16	-0.10	-0.20	0.04

Q9CYK1	Tryptophan-tRNA ligase, mitochondrial OS=Mus musculus GN=Wars2 PE=2 SV=2 [SYWM_MOUSE]	0.18	0.28	0.21	0.06	0.07	0.12	0.22	0.10	-0.07	-0.07	-0.09	-0.10	0.04
F2Z4B7	Protein Raph1 OS=Mus musculus GN=Raph1 PE=2 SV=1 [F2Z4B7_MOUSE]	0.39	0.19	-0.10	0.03	-0.09	-0.04	0.07	-0.79	-0.09	-0.20	-0.27	-0.46	0.04
F8WHV4	Collagen alpha-1(XIX) chain OS=Mus musculus GN=Col19a1 PE=2 SV=1 [F8WHV4_MOUSE]	0.17	-0.61	-0.40	-0.35	0.30	-0.92	-0.54	0.52	0.77	0.00	0.76	0.81	0.04
Q9QXX4	Calcium-binding mitochondrial carrier protein Aralar2 OS=Mus musculus GN=Slc25a13 PE=1 SV=1 [CMC2_MOUSE]	-0.10	0.48	-0.19	0.48	-0.33	0.59	0.20	-0.95	-0.33	-0.51	-0.21	-0.46	0.04
E0CYV9	Protein 1110002E22Rik OS=Mus musculus GN=1110002E22Rik PE=4 SV=1 [E0CYV9_MOUSE]	-0.45	0.11	-0.28	0.47	0.11	-0.20	-0.58	-0.42	-0.34	-0.65	0.00	-0.74	0.04
F7BQE4	Rho guanine nucleotide exchange factor 10 (Fragment) OS=Mus musculus GN=Arhgef10 PE=2 SV=1 [F7BQE4_MOUSE]	-0.46	-0.61	-0.09	0.02	-0.04	-0.68	0.11	0.79	-0.18	0.09	-0.24	0.67	0.04
Q920E5	Farnesyl pyrophosphate synthase OS=Mus musculus GN=Fdps PE=2 SV=1 [FPPS_MOUSE]	0.25	0.06	0.30	0.05	-0.03	0.12	0.15	-0.01	-0.38	-0.21	-0.33	0.03	0.04
G3UYG6	PERQ amino acid-rich with GYF domain-containing protein 2 (Fragment) OS=Mus musculus GN=Gigyf2 PE=2 SV=1 [G3UYG6_MOUSE]	0.12	0.10	0.04	-0.02	-0.06	0.08	-0.31	-0.05	-0.06	0.03	-0.14	-0.46	0.04
Q921C5-2	Isoform 2 of Protein bicaudal D homolog 2 OS=Mus musculus GN=Bicd2 [BICD2_MOUSE]	0.13	-0.08	0.07	0.53	0.18	0.08	-0.25	-0.04	0.02	-0.07	0.05	-0.55	0.04
Q8VEH8	Endoplasmic reticulum lectin 1 OS=Mus musculus GN=Erlec1 PE=2 SV=1 [ERLEC_MOUSE]	0.20	-0.15	0.04	-0.06	0.01	-0.04	0.07	0.25	0.08	0.32	0.02	0.42	0.04
Q00558	Factor VIII intron 22 protein OS=Mus musculus GN=F8a1 PE=2 SV=1 [F8I2_MOUSE]	0.00	-0.23	-0.02	0.20	0.05	-0.05	-0.09	-0.51	-0.42	0.02	-0.22	-0.26	0.04
Q8K339	DNA/RNA-binding protein KIN17 OS=Mus musculus GN=Kin PE=2 SV=1 [KIN17_MOUSE]	-0.62	0.95	-0.02	0.01	0.33	-0.03	-0.49	-0.65	0.08	-1.07	-0.15	-1.12	0.04
P62746	Rho-related GTP-binding protein RhoB OS=Mus musculus GN=Rhob PE=1 SV=1 [RHOB_MOUSE]	0.05	-0.21	-0.05	-0.07	-0.05	-0.35	0.24	-0.09	0.29	-0.10	0.24	0.05	0.04

O35600	Retinal-specific ATP-binding cassette transporter OS=Mus musculus GN=Abca4 PE=2 SV=1 - [ABCA4_MOUSE]	-0.01	-0.17	-0.06	-0.78	0.06	-0.37	-0.06	0.27	0.33	0.67	-0.13	0.06	0.04
Q3U319	E3 ubiquitin-protein ligase BRE1B OS=Mus musculus GN=Rnf40 PE=2 SV=2 - [BRE1B_MOUSE]	0.30	-0.21	0.20	0.01	0.11	0.29	-0.11	-0.37	-0.03	-0.11	0.03	-0.09	0.04
P48678	Prelamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2 - [LMNA_MOUSE]	-0.04	0.05	-0.07	0.04	0.19	0.03	-0.30	-0.03	-0.18	-0.02	-0.05	-0.06	0.04
F6QP10	Sn1-specific diacylglycerol lipase beta (Fragment) OS=Mus musculus GN=Daglb PE=4 SV=2 - [F6QP10_MOUSE]	0.01	-0.30	-0.01	-0.18	-0.22	-0.26	0.26	-0.18	0.18	0.03	0.25	-0.13	0.04
Q80W00	Serine/threonine-protein phosphatase 1 regulatory subunit 10 OS=Mus musculus GN=Ppp1r10 PE=1 SV=1 - [PP1RA_MOUSE]	0.16	-0.10	-0.16	-0.19	0.10	-0.16	-0.30	-1.08	-0.07	-0.94	-0.10	-1.00	0.04
E9QP99	Golgin subfamily A member 3 OS=Mus musculus GN=Golga3 PE=2 SV=1 - [E9QP99_MOUSE]	-0.07	0.01	-0.10	0.19	0.15	0.23	-0.42	-0.14	-0.03	-0.04	-0.05	-0.09	0.04
O88962	7-alpha-hydroxycholesterol 4-en-3-one 12-alpha-hydroxylase OS=Mus musculus GN=Cyp8b1 PE=2 SV=1 - [CP8B1_MOUSE]	-0.46	-0.01	-0.42	-0.24	-0.28	0.40	-0.36	0.55	0.24	0.55	0.23	0.51	0.04
Q62421	Endophilin-A3 OS=Mus musculus GN=Sh3gl3 PE=2 SV=1 - [SH3G3_MOUSE]	0.37	0.07	0.11	0.63	-0.01	0.19	0.01	0.07	-0.23	0.06	-0.31	0.07	0.04
Q3TP92	CTD nuclear envelope phosphatase 1 OS=Mus musculus GN=Ctdnep1 PE=2 SV=2 - [CNEP1_MOUSE]	0.35	-0.85	-0.26	-0.76	-0.11	-0.51	-0.16	0.41	0.08	0.50	0.03	0.04	0.04
P97868	E3 ubiquitin-protein ligase RBBP6 OS=Mus musculus GN=Rbbp6 PE=1 SV=5 - [RBBP6_MOUSE]	-0.19	0.90	-0.16	0.60	0.19	0.97	-0.42	-0.07	0.04	-0.46	0.03	-0.23	0.04
Q8JZX5	Leucine rich repeat containing 57 OS=Mus musculus GN=Lrrc57 PE=2 SV=1 - [Q8JZX5_MOUSE]	0.02	-0.22	0.06	-0.01	-0.05	-0.16	0.12	0.24	-0.04	0.04	0.02	0.09	0.04

Q925E7	Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B delta isoform OS=Mus musculus GN=Ppp2r2d PE=2 SV=1 - [ZABD_MOUSE]	0.76	-0.02	0.27	0.08	-0.13	0.11	0.07	-0.14	-0.46	-0.18	-0.37	-0.02	0.04
Q921Q7	Ras and Rab interactor 1 OS=Mus musculus GN=Rin1 PE=1 SV=1 - [RIN1_MOUSE]	0.33	0.07	0.36	0.14	0.07	-0.08	0.06	0.00	-0.15	0.00	-0.25	0.03	0.04
Q9Z1A9	TBC1 domain family member 8 OS=Mus musculus GN=Tbc1d8 PE=2 SV=2 - [TBCD8_MOUSE]	-0.33	-0.34	-0.46	-0.30	0.25	0.04	-0.22	0.34	-0.10	0.20	0.37	0.43	0.04
P23591	GDP-L-fucose synthase OS=Mus musculus GN=Tsta3 PE=2 SV=3 - [FCL_MOUSE]	-0.12	0.21	-0.08	-0.06	-0.15	-0.11	0.02	0.29	0.09	0.17	0.26	-0.06	0.04
O70230-3	Isoform 3 of Zinc finger protein 143 OS=Mus musculus GN=Znf143 - [ZN143_MOUSE]	-0.05	3.52	-0.26	0.24	0.04	0.13	0.00	-5.54	0.00	-4.94	0.05	-5.41	0.04
Q5SU71	E3 ubiquitin/ISG15 ligase TRIM25 OS=Mus musculus GN=Trim25 PE=2 SV=1 - [Q5SU71_MOUSE]	1.39	0.08	0.05	0.05	0.29	-0.05	-0.45	0.01	-0.48	0.03	-0.69	-0.28	0.04
Q8BR63	Protein FAM177A1 OS=Mus musculus GN=Fam177a1 PE=2 SV=1 - [F177A_MOUSE]	-0.14	-0.15	-0.21	0.07	-0.09	-0.08	0.08	0.00	0.18	0.00	-0.01	-0.09	0.04
A2A7B5	Protein Prdm2 OS=Mus musculus GN=Prdm2 PE=2 SV=1 - [A2A7B5_MOUSE]	0.93	0.43	0.02	-0.02	0.37	0.35	-0.26	-0.19	0.21	0.00	0.16	-0.16	0.04
B1AVU6	Protein Gm436 OS=Mus musculus GN=Gm436 PE=4 SV=1 - [B1AVU6_MOUSE]	0.14	-0.13	-0.40	-0.01	-0.29	0.21	0.04	0.46	0.36	-0.19	0.44	0.55	0.04
Q3UMT1	Protein phosphatase 1 regulatory subunit 12C OS=Mus musculus GN=Ppp1r12c PE=1 SV=1 - [PP12C_MOUSE]	-0.37	-0.26	-0.12	-0.28	0.09	-0.21	-0.34	0.04	0.14	0.16	0.25	0.11	0.04
Q8BN51	Thymidine kinase 2, mitochondrial OS=Mus musculus GN=TK2 PE=2 SV=1 - [Q8BN51_MOUSE]	1.13	0.30	-0.35	0.27	0.06	0.23	-0.48	-0.30	-0.71	0.11	-0.28	-0.02	0.04
Q8VE99	Coiled-coil domain-containing protein 115 OS=Mus musculus GN=Cocd115 PE=2 SV=1 - [CC115_MOUSE]	-0.17	-0.67	-0.21	-0.47	-0.08	-0.24	-0.15	-0.09	0.20	-0.14	-0.02	-0.15	0.04
Q91WK1	SPRY domain-containing protein 4 OS=Mus musculus GN=Spryd4 PE=2 SV=1 - [SPRY4_MOUSE]	0.07	-0.06	-0.17	0.07	-0.35	0.01	0.20	0.24	-0.06	0.13	0.07	0.11	0.04

Q4VC33	Macrophage erythroblast attacher OS=Mus musculus GN=Maea PE=1 SV=1 - [MAEA_MOUSE]	0.44	0.58	0.36	0.24	-0.02	0.33	0.50	-0.58	0.08	-0.24	-0.10	-0.78	0.04
Q8BGU2	Cerebellin-2 OS=Mus musculus GN=Cbln2 PE=1 SV=1 - [CBLN2_MOUSE]	0.17	1.63	-0.16	-0.26	0.42	-0.03	-0.65	-2.35	0.10	-1.12	0.22	-2.76	0.04
Q9QXS1-6	Isoform PLEC-0,1C of Plectin OS=Mus musculus GN=Plec - [PLEC_MOUSE]	0.50	0.36	0.25	-0.14	-0.09	-0.10	0.18	-0.46	-0.22	-0.18	-0.13	-0.48	0.04
Q6PIC6	Sodium/potassium-transporting ATPase subunit alpha-3 OS=Mus musculus GN=Atp1a3 PE=1 SV=1 - [AT1A3_MOUSE]	-0.22	-0.05	0.04	0.06	-0.15	-0.13	0.11	0.01	0.09	0.05	-0.02	0.06	0.04
P41731	CD63 antigen OS=Mus musculus GN=Cd63 PE=1 SV=2 - [CD63_MOUSE]	-0.47	-1.07	-0.20	-0.85	-0.08	-0.50	-0.23	-0.28	0.13	-0.04	0.06	-0.31	0.04
Q62443	Neuronal pentraxin-1 OS=Mus musculus GN=Nptx1 PE=2 SV=1 - [NPTX1_MOUSE]	-0.08	0.02	-0.12	0.00	-0.06	0.03	0.06	-0.03	0.12	0.07	0.14	-0.03	0.04
E9QKU6	Serine/threonine-protein phosphatase 4 regulatory subunit 4 OS=Mus musculus GN=Ppp4r4 PE=4 SV=1 - [E9QKU6_MOUSE]	0.09	-0.18	0.22	-0.18	-0.09	-0.17	0.63	-0.01	0.03	0.27	0.19	0.21	0.04
Q9CVD2	Ataxin-3 OS=Mus musculus GN=Atxn3 PE=1 SV=2 - [ATX3_MOUSE]	0.26	0.30	0.11	0.40	0.13	0.14	0.21	-0.06	0.09	0.14	0.10	-0.17	0.04
Q9CW46	Ribonucleoprotein PTB-binding 1 OS=Mus musculus GN=Raver1 PE=1 SV=2 - [RAVR1_MOUSE]	-0.03	0.34	0.18	0.28	0.12	0.23	0.03	-0.04	0.13	-0.45	0.14	-0.36	0.04
P10649	Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2 - [GSTM1_MOUSE]	0.29	0.03	0.22	0.16	0.07	0.15	0.19	-0.07	-0.03	0.05	0.05	-0.02	0.04
P97950	Ras-related protein Rab-33A OS=Mus musculus GN=Rab33a PE=2 SV=1 - [RB33A_MOUSE]	0.69	-0.12	0.20	0.18	-0.07	0.02	0.14	-0.37	-0.22	-0.15	-0.21	-0.27	0.04
P70211-2	Isoform C of Netrin receptor DCC OS=Mus musculus GN=Dcc - [DCC_MOUSE]	0.09	-0.08	-0.07	-0.26	0.00	-0.48	0.10	0.20	0.00	0.03	-0.03	0.27	0.04
Q8R1K1	Ubiquitin-associated domain-containing protein 2 OS=Mus musculus GN=Ubac2 PE=2 SV=1 - [UBAC2_MOUSE]	0.21	0.28	0.25	0.00	-0.12	0.27	0.20	-0.33	-0.21	-0.44	0.12	-0.73	0.04
P58774-2	Isoform 2 of Tropomyosin beta chain OS=Mus musculus GN=Tpm2 - [TPM2_MOUSE]	-0.16	-0.51	-0.37	-0.93	0.28	-0.19	-0.29	0.16	0.68	0.36	0.40	-0.15	0.04

P97765	WW domain-binding protein 2 OS=Mus musculus GN=Wbp2 PE=1 SV=1 - [WBP2_MOUSE]	-0.29	-0.06	-0.24	0.00	-0.09	-0.13	-0.02	0.21	-0.18	0.03	0.00	0.23	0.05
Q9DBT3	Coiled-coil domain-containing protein 97 OS=Mus musculus GN=Ccd97 PE=1 SV=1 - [CCD97_MOUSE]	0.00	0.64	-0.21	0.60	0.19	0.36	-0.32	-0.87	0.23	-0.11	0.14	-0.48	0.05
Q8K207	Uncharacterized protein C1orf21 homolog OS=Mus musculus PE=1 SV=1 - [CA021_MOUSE]	0.03	0.86	-0.13	0.38	0.05	0.12	-0.29	-1.11	0.24	-1.02	0.09	-0.39	0.05
Q8CH72	E3 ubiquitin-protein ligase TRIM32 OS=Mus musculus GN=Trim32 PE=1 SV=2 - [TRIM32_MOUSE]	0.04	0.06	0.25	-0.04	0.01	-0.02	0.14	-0.16	-0.25	-0.09	-0.25	-0.13	0.05
Q61789	Laminin subunit alpha-3 OS=Mus musculus GN=Lama3 PE=1 SV=3 - [LAMA3_MOUSE]	0.18	-0.22	-0.10	-0.08	0.24	-0.06	-0.47	-0.49	-0.14	0.05	-0.26	-0.24	0.05
Q6ZPS6	Ankyrin repeat and IBR domain-containing protein 1 OS=Mus musculus GN=Ankib1 PE=1 SV=2 - [AKIB1_MOUSE]	0.20	0.77	0.11	0.20	0.07	0.52	0.08	-0.13	-0.14	0.26	-0.02	-0.01	0.05
G5E819	Serine/threonine-protein phosphatase OS=Mus musculus GN=Ppp5c PE=3 SV=1 - [G5E819_MOUSE]	-0.22	-0.09	-0.03	0.01	0.04	0.03	-0.05	0.07	0.15	0.09	0.04	0.15	0.05
O35343	Importin subunit alpha-3 OS=Mus musculus GN=Kpna4 PE=2 SV=1 - [IMA3_MOUSE]	-0.92	-0.27	-0.02	-0.08	-0.17	-0.13	-0.30	0.03	0.25	0.19	0.27	0.26	0.05
P47746	Cannabinoid receptor 1 OS=Mus musculus GN=Cnr1 PE=2 SV=1 - [CNR1_MOUSE]	-0.24	-0.21	-0.01	0.04	-0.20	-0.23	0.10	-0.07	0.16	-0.07	0.05	-0.12	0.05
O88736	3-keto-steroid reductase OS=Mus musculus GN=Hsd17b7 PE=2 SV=1 - [DHB7_MOUSE]	1.07	-0.08	0.30	0.01	0.02	-0.01	-0.56	0.03	-1.08	0.11	-0.70	-0.09	0.05
Q8CC86	Nicotinate phosphoribosyltransferase OS=Mus musculus GN=Naprt1 PE=2 SV=1 - [PNCB_MOUSE]	-0.11	-0.11	-0.07	-0.16	0.07	-0.16	-0.09	-0.03	0.09	0.11	0.43	0.14	0.05
Q8VE22	28S ribosomal protein S23, mitochondrial OS=Mus musculus GN=Mps23 PE=2 SV=1 - [RT23_MOUSE]	0.33	0.13	0.15	0.22	-0.03	0.20	0.35	-0.19	-0.34	-0.01	-0.26	-0.19	0.05
F7D291	MTSS1-like protein (Fragment) OS=Mus musculus GN=Mtss1l PE=4 SV=1 - [F7D291_MOUSE]	-0.27	-0.15	-0.07	-0.51	-0.11	0.08	0.32	-0.07	0.25	0.74	0.29	-0.25	0.05

Q60936	Chaperone activity of bc1 complex-like, mitochondrial OS=Mus musculus GN=Adck3 PE=2 SV=2 - [ADCK3_MOUSE]	0.19	0.00	-0.11	0.00	-0.20	-0.24	0.23	0.47	-0.11	0.43	-0.06	0.38	0.05
Q3UTQ8	Cyclin-dependent kinase-like 5 OS=Mus musculus GN=Cdkl5 PE=2 SV=1 - [CDKL5_MOUSE]	-0.05	-0.13	0.10	-0.04	0.09	-0.06	-0.23	-0.06	-0.21	-0.15	-0.05	-0.08	0.05
P00416	Cytochrome c oxidase subunit 3 OS=Mus musculus GN=Mtco3 PE=3 SV=1 - [COX3_MOUSE]	-1.06	-0.67	-0.29	-0.58	-0.45	-0.23	-0.24	0.57	-0.53	0.18	-0.44	0.77	0.05
P29387	Guanine nucleotide-binding protein subunit beta-4 OS=Mus musculus GN=Gnb4 PE=2 SV=4 - [GBB4_MOUSE]	-0.81	0.37	-0.21	-0.34	-0.16	-0.23	-0.03	0.76	0.01	0.03	0.25	0.28	0.05
Q8BVA2	Transmembrane protein 222 OS=Mus musculus GN=Tmem222 PE=2 SV=1 - [TM222_MOUSE]	0.29	-0.03	-0.03	-0.06	-0.05	0.00	0.24	0.07	0.14	0.13	0.71	0.45	0.05
E9PWB1	Pleckstrin homology-like domain family B member 1 OS=Mus musculus GN=Phldb1 PE=2 SV=1 - [E9PWB1_MOUSE]	0.18	0.12	-0.01	0.04	0.01	0.23	-0.11	-0.16	-0.01	0.09	0.09	-0.18	0.05
B8QI35	Liprin-alpha 3 OS=Mus musculus GN=Ppfia3 PE=2 SV=1 - [B8QI35_MOUSE]	0.10	-0.05	0.27	-0.01	0.01	-0.05	0.03	-0.16	-0.18	-0.07	-0.14	-0.05	0.05
Q8CGF6	WD repeat-containing protein 47 OS=Mus musculus GN=Wdr47 PE=1 SV=2 - [WDR47_MOUSE]	-0.42	-0.24	0.03	-0.13	-0.03	-0.11	-0.03	0.05	-0.03	-0.02	0.06	0.06	0.05
Q8BLN5	Lanosterol synthase OS=Mus musculus GN=Lss PE=2 SV=2 - [ERG7_MOUSE]	-0.70	-0.08	-0.26	0.07	-0.38	0.20	0.13	0.36	-0.02	0.20	0.14	0.13	0.05
Q8C788	Sorting nexin 18 OS=Mus musculus GN=Snx18 PE=2 SV=1 - [Q8C788_MOUSE]	-0.14	-0.45	0.10	-0.61	-0.08	-0.20	-0.19	0.49	-0.03	0.20	-0.13	0.55	0.05
Q3U213-2	Isoform 2 of Protein SERAC1 OS=Mus musculus GN=Serac1 - [SRAC1_MOUSE]	-0.24	-0.25	0.01	-0.40	-0.12	-0.19	0.10	-0.13	0.00	0.32	-0.07	-0.16	0.05
Q811Q9-2	Isoform 2 of Choline-phosphate cytidyltransferase B OS=Mus musculus GN=Pcyt1b - [PCY1B_MOUSE]	0.21	0.02	-0.08	-0.07	-0.06	0.05	-0.32	-0.02	-0.12	-0.06	-0.17	-0.10	0.05
Q61627	Glutamate receptor ionotropic, delta-1 OS=Mus musculus GN=Grid1 PE=1 SV=2 - [GRID1_MOUSE]	0.10	-0.08	0.01	-0.15	0.03	0.11	0.01	0.00	0.14	0.21	0.26	0.26	0.05

H3BLA9	Sentrin-specific protease 6 (Fragment) OS=Mus musculus GN=Senp6 PE=2 SV=1 [H3BLA9_MOUSE]	0.21	-0.32	-0.19	-0.28	-0.10	-0.40	-0.01	0.00	0.04	0.08	0.26	-0.05	0.05
Q5SSE9	ATP-binding cassette sub-family A member 13 OS=Mus musculus GN=Abca13 PE=2 SV=1 [ABCAD_MOUSE]	-0.46	0.13	0.04	0.34	0.13	-0.52	-0.52	-1.12	-0.19	-0.32	-0.29	-1.53	0.05
P62267	40S ribosomal protein S23 OS=Mus musculus GN=Rps23 PE=2 SV=3 [RS23_MOUSE]	0.02	0.07	0.40	-0.02	0.05	0.17	0.13	0.02	-0.43	-0.26	-0.41	0.00	0.05
O35495-2	Isoform 2 of Cyclin-dependent kinase 14 OS=Mus musculus GN=Cdk14 [CDK14_MOUSE]	0.01	0.08	0.08	0.06	0.00	0.10	0.04	0.05	-0.09	0.03	-0.12	-0.09	0.05
Q8CDA1	Phosphatidylinositol phosphatase SAC2 OS=Mus musculus GN=Inpp5f PE=1 SV=1 [SAC2_MOUSE]	0.19	0.09	0.16	-0.09	-0.02	0.10	0.14	-0.25	-0.10	-0.04	-0.20	-0.07	0.05
A2AHX7	Bcl-2-like protein 1 (Fragment) OS=Mus musculus GN=Bcl2l1 PE=2 SV=1 [A2AHX7_MOUSE]	0.32	-0.58	0.34	-0.90	0.34	-0.38	0.56	0.19	0.69	0.56	-0.06	0.86	0.05
Q9D358	Low molecular weight phosphotyrosine protein phosphatase OS=Mus musculus GN=Acp1 PE=1 SV=3 [PPAC_MOUSE]	0.03	-0.23	0.09	-0.19	-0.66	-0.13	0.57	-0.04	-0.23	0.48	0.44	0.04	0.05
Q91YW3	DnaJ homolog subfamily C member 3 OS=Mus musculus GN=Dnajc3 PE=1 SV=1 [DNJC3_MOUSE]	0.17	-0.02	0.18	0.05	0.22	0.18	0.07	0.05	-0.06	-0.08	0.10	0.03	0.05
P15532	Nucleoside diphosphate kinase A OS=Mus musculus GN=Nme1 PE=1 SV=1 [NDKA_MOUSE]	0.18	0.06	0.01	0.18	0.21	0.29	0.13	-0.01	0.02	0.00	0.01	0.10	0.05
P55821	Stathmin-2 OS=Mus musculus GN=Stmn2 PE=1 SV=1 [STMN2_MOUSE]	0.15	0.11	-0.16	0.05	0.00	-0.15	-0.11	-0.27	0.08	-0.37	-0.18	-0.30	0.05
Q9WVL5	MORC family CW-type zinc finger protein 1 OS=Mus musculus GN=Morc1 PE=2 SV=1 [MORC1_MOUSE]	-0.85	-0.35	-0.20	-0.08	-0.07	-0.47	-0.54	-0.09	0.33	0.24	0.26	0.27	0.05
Q8VDI9	Alpha-1,2-mannosyltransferase ALG9 OS=Mus musculus GN=Alg9 PE=2 SV=1 [ALG9_MOUSE]	-0.34	0.37	0.40	0.31	-0.10	0.18	0.11	-0.30	-0.36	-0.22	-0.37	-0.05	0.05
Q00915	Retinol-binding protein 1 OS=Mus musculus GN=Rbp1 PE=2 SV=2 [RET1_MOUSE]	-0.05	0.12	0.01	-0.12	0.02	-0.07	0.26	0.08	0.03	0.18	0.18	-0.04	0.05

Q62084	Protein phosphatase 1 regulatory subunit 14B OS=Mus musculus GN=Ppp1r14b PE=1 SV=2 [PP14B_MOUSE]	0.32	0.38	-0.30	0.47	0.41	0.64	-0.47	-0.69	0.45	-0.54	0.37	-0.88	0.05
Q91X96	Guanine nucleotide exchange factor MSS4 OS=Mus musculus GN=Rabif PE=2 SV=1 [MSS4_MOUSE]	-0.61	-0.03	-0.59	-0.20	0.00	0.10	-0.35	0.25	0.35	0.59	0.25	0.02	0.05
Q8BG26	RUN and SH3 domain-containing protein 1 OS=Mus musculus GN=Rusc1 PE=1 SV=2 [RUSC1_MOUSE]	0.24	0.15	0.10	0.08	0.30	-0.18	-0.41	0.03	-0.27	0.05	-0.32	0.07	0.05
Q3TWW8	Protein Srsf6 OS=Mus musculus GN=Srsf6 PE=2 SV=1 [Q3TWW8_MOUSE]	0.10	0.03	-0.14	0.03	0.30	-0.12	-0.24	-0.23	-0.02	-0.54	0.08	-0.40	0.05
O35654	DNA polymerase delta subunit 2 OS=Mus musculus GN=Pold2 PE=2 SV=2 [DPOD2_MOUSE]	0.38	0.00	0.11	0.33	-0.14	0.22	-0.04	-0.59	-0.68	0.36	-0.63	-0.18	0.05
A2RSF9	Protein Serpinb3c OS=Mus musculus GN=Serpinb3c PE=2 SV=1 [A2RSF9_MOUSE]	1.15	-0.02	0.63	-0.30	0.79	0.43	-0.04	-0.44	0.24	0.00	-0.20	-0.27	0.05
P40240	CD9 antigen OS=Mus musculus GN=Cd9 PE=1 SV=2 [CD9_MOUSE]	-0.38	-1.14	-0.16	-1.38	0.09	-1.66	-0.22	-0.09	0.09	0.07	-0.02	0.07	0.05
Q99NF8	Ran-binding protein 17 OS=Mus musculus GN=Ranbp17 PE=2 SV=1 [RBP17_MOUSE]	0.45	0.05	-0.38	-0.10	0.19	-0.10	0.09	0.52	0.47	0.05	0.44	0.52	0.05
Q9CZ44	NSFL1 cofactor p47 OS=Mus musculus GN=Nsf1c PE=1 SV=1 [NSF1C_MOUSE]	0.06	0.14	-0.06	0.19	0.12	0.20	-0.26	0.05	-0.04	0.06	-0.04	0.03	0.05
Q99LH9	SH3 domain-binding protein 5-like OS=Mus musculus GN=Sh3bp5l PE=1 SV=1 [3BP5L_MOUSE]	-0.20	-0.60	-0.20	-0.99	0.10	-0.88	0.01	-0.09	0.20	-0.22	0.09	-0.02	0.05
G3UVV3	MCG1556, isoform CRA_b OS=Mus musculus GN=Zfp407 PE=4 SV=1 [G3UVV3_MOUSE]	-1.22	0.03	-0.22	-0.28	-0.42	-0.21	-0.16	0.39	-0.21	0.41	-0.02	0.13	0.05
P55144-3	Isoform 1 of Tyrosine-protein kinase receptor TYRO3 OS=Mus musculus GN=Tyro3 [TYRO3_MOUSE]	-0.13	-0.19	-0.08	-0.24	0.03	0.04	0.00	0.27	0.06	0.59	-0.14	0.30	0.05
E9PYP8	Beta-crystallin B1 (Fragment) OS=Mus musculus GN=Crybb1 PE=2 SV=1 [E9PYP8_MOUSE]	0.07	-0.93	0.01	-0.25	-0.11	-0.49	0.04	0.16	0.14	-0.18	0.19	0.36	0.05

Q6P6M7	O-phosphoserine tRNA(Sec) selenium transferase OS=Mus musculus GN=Sepsecs PE=1 SV=2 - [SPCS_MOUSE]	0.28	-0.31	-0.12	-0.21	-0.06	-0.31	-0.29	0.37	0.28	0.20	0.34	0.17	0.05
Q3U0K8-2	Isoform 2 of 2- oxoglutarate and iron- dependent oxygenase domain-containing protein 1 OS=Mus musculus GN=Ogfd1 - [OGFD1_MOUSE]	-0.06	-0.42	-0.01	-0.21	-0.09	-0.26	0.23	-0.06	-0.10	0.18	-0.18	0.11	0.05
Q9WTX8	Mitotic spindle assembly checkpoint protein MAD1 OS=Mus musculus GN=Mad11 PE=2 SV=1 - [MD1L1_MOUSE]	0.48	-0.05	0.20	-0.20	0.00	-0.20	-0.15	-0.26	-0.07	-0.44	-0.10	-0.65	0.05
Q5SYL3	UPF0378 protein KIAA0100 OS=Mus musculus GN=Kiaa0100 PE=2 SV=1 - [K0100_MOUSE]	-0.22	-0.02	-0.22	-0.26	0.05	-0.27	0.31	-0.30	0.10	0.03	0.08	0.34	0.05
Q8BGZ1	Hippocalcin-like protein 4 OS=Mus musculus GN=Hpcal4 PE=2 SV=3 - [HPCL4_MOUSE]	-0.02	-0.14	-0.22	-0.04	0.04	-0.12	-0.09	0.12	0.19	-0.07	0.06	0.09	0.05
Q8K3I4	Rab effector MyRIP OS=Mus musculus GN=Myrip PE=1 SV=1 - [MYRIP_MOUSE]	0.93	-0.06	0.24	-0.08	0.23	0.27	-0.09	-0.29	-0.06	-0.28	0.00	-0.06	0.05
O55229	Choline/ethanolamine kinase OS=Mus musculus GN=Chkb PE=1 SV=3 - [CHKB_MOUSE]	0.07	-0.03	0.15	0.03	-0.18	0.14	-0.14	-0.37	-0.07	-0.41	0.04	-0.81	0.05
Q8K3D3-2	Isoform 2 of DNA repair protein SWI5 homolog OS=Mus musculus GN=Swi5 - [SWI5_MOUSE]	-0.22	1.18	-0.07	1.26	0.27	1.58	-0.04	-0.37	0.04	-0.19	-0.04	-0.18	0.05
Q6P8I6	Cytochrome c oxidase assembly protein COX11, mitochondrial OS=Mus musculus GN=Cox11 PE=2 SV=1 -[COX11_MOUSE]	0.56	-0.02	-0.06	0.41	-0.18	0.54	-0.07	1.64	0.80	2.80	0.65	1.74	0.05
D3Z5X4	Uncharacterized protein OS=Mus musculus GN=Gm7964 PE=2 SV=1 - [D3Z5X4_MOUSE]	-0.42	-0.18	-0.20	-0.54	0.06	-0.08	-0.15	0.23	-0.11	0.16	-0.15	0.32	0.05
Q9D2J7	Ankyrin repeat and EF- hand domain- containing protein 1 OS=Mus musculus GN=Ankef1 PE=2 SV=1 - [ANKE1_MOUSE]	-0.50	1.36	-0.26	0.62	0.21	0.35	-0.39	-3.06	0.29	-2.61	0.16	-3.63	0.05
Q61107	Guanylate-binding protein 4 OS=Mus musculus GN=Gbp4 PE=1 SV=1 - [GBP4_MOUSE]	-0.86	-0.25	-0.55	-0.32	0.13	-0.44	-0.45	0.06	0.35	-0.26	0.37	0.16	0.05

Q8R1V4	Transmembrane emp24 domain-containing protein 4 OS=Mus musculus GN=Tmed4 PE=2 SV=1 - [TMED4_MOUSE]	0.00	-0.02	0.09	-0.26	-0.28	0.06	0.31	0.16	-0.03	0.16	0.07	0.03	0.05
P70388-2	Isoform 2 of DNA repair protein RAD50 OS=Mus musculus GN=Rad50 - [RAD50_MOUSE]	3.68	0.72	-0.43	0.27	0.08	0.28	-2.42	-0.32	-0.85	0.01	-0.64	-0.72	0.05
Q9WUD1	STIP1 homology and U box-containing protein 1 OS=Mus musculus GN=Stub1 PE=1 SV=1 - [CHIP_MOUSE]	-0.27	-0.14	-0.03	0.02	-0.01	-0.03	0.01	-0.02	0.10	0.05	0.03	0.05	0.05
Q925I8	OL-protocadherin isoform OS=Mus musculus GN=Pcdh10 PE=2 SV=1 - [Q925I8_MOUSE]	0.12	-0.06	0.03	0.06	-0.11	-0.04	-0.07	-0.26	-0.11	-0.08	-0.01	-0.10	0.05
Q3UYV9	Nuclear cap-binding protein subunit 1 OS=Mus musculus GN=Ncbp1 PE=1 SV=2 - [NCBP1_MOUSE]	-0.01	-0.39	-0.06	-0.29	-0.14	-0.14	0.13	0.13	-0.03	-0.24	0.08	-0.02	0.05
P62315	Small nuclear ribonucleoprotein Sm D1 OS=Mus musculus GN=Snrpd1 PE=1 SV=1 - [SMD1_MOUSE]	0.44	0.16	-0.09	0.24	-0.03	0.33	-0.01	-0.37	0.06	-0.13	0.19	-0.40	0.05
Q8BT60	Copine-3 OS=Mus musculus GN=Cpne3 PE=1 SV=2 - [CPNE3_MOUSE]	-0.13	-0.34	0.09	-0.10	-0.03	-0.20	0.04	0.24	0.08	-0.07	0.06	-0.05	0.05
P68254	14-3-3 protein theta OS=Mus musculus GN=Ywhaq PE=1 SV=1 - [1433T_MOUSE]	-0.49	-0.01	-0.10	0.00	-0.03	0.00	0.09	0.03	0.17	0.06	0.15	0.08	0.05
Q9Z247	Peptidyl-prolyl cis-trans isomerase FKBP9 OS=Mus musculus GN=Fkbp9 PE=1 SV=1 - [FKBP9_MOUSE]	0.16	1.14	-0.24	0.01	0.08	-0.25	-0.06	-0.73	-0.26	-0.90	0.11	-0.96	0.05
Q3U9F6	RING finger protein 166 OS=Mus musculus GN=Rnf166 PE=2 SV=1 - [RN166_MOUSE]	-0.14	-0.92	-0.41	0.05	0.06	-0.13	-0.35	0.61	0.08	0.32	0.07	0.61	0.05
Q9CYH6	Ribosome biogenesis regulatory protein homolog OS=Mus musculus GN=Rrs1 PE=2 SV=1 - [RRS1_MOUSE]	0.10	0.54	-0.04	0.32	0.36	0.39	-0.65	-0.35	0.14	-0.23	0.39	-0.02	0.05
Q9R0P5	Dextrin OS=Mus musculus GN=Dstn PE=1 SV=3 - [DEST_MOUSE]	-0.20	-0.10	-0.11	-0.01	-0.09	0.00	0.01	0.24	-0.06	0.19	-0.12	0.20	0.05
E9PZM4	Protein Chd2 OS=Mus musculus GN=Chd2 PE=2 SV=1 - [E9PZM4_MOUSE]	0.14	0.31	0.21	0.06	0.01	0.03	-0.65	0.07	0.05	-0.14	-0.05	-0.16	0.05
Q80U78-3	Isoform 3 of Pumilio homolog 1 OS=Mus musculus GN=Pum1 - [PUM1_MOUSE]	0.16	-0.16	0.07	0.07	0.13	0.04	-0.25	-0.11	-0.06	-0.01	0.02	-0.09	0.05

Q80T10-3	Isoform 3 of GRAM domain-containing protein 1B OS=Mus musculus GN=Gramd1b - [GRM1B_MOUSE]	0.47	0.37	0.08	0.18	-0.03	-0.23	0.25	-0.46	-0.13	-0.27	-0.15	-0.32	0.05
Q8R2R3	Alpha- and gamma-adaptin-binding protein p34 OS=Mus musculus GN=Aagab PE=2 SV=2 - [AAGAB_MOUSE]	3.57	0.20	0.85	0.59	0.89	0.38	-0.32	-0.03	-0.04	-0.42	-0.13	-0.28	0.05
P70428	Exostosin-2 OS=Mus musculus GN=Ext2 PE=1 SV=2 - [EXT2_MOUSE]	0.51	0.53	-0.05	0.75	0.15	0.62	-0.25	0.43	0.04	0.08	0.12	0.05	0.05
Q9WVA3	Mitotic checkpoint protein BUB3 OS=Mus musculus GN=Bub3 PE=2 SV=2 - [BUB3_MOUSE]	0.04	-0.13	-0.14	-0.32	-0.12	0.02	0.27	0.09	-0.12	0.17	0.04	-0.06	0.05
E9QJS0	28S ribosomal protein S10, mitochondrial OS=Mus musculus GN=Mmps10 PE=3 SV=1 - [E9QJS0_MOUSE]	0.25	0.08	0.24	-0.10	-0.01	0.12	0.15	-0.08	-0.32	-0.27	-0.21	0.06	0.05
Q99MP8-2	Isoform 2 of BRCA1-associated protein OS=Mus musculus GN=Brap - [BRAP_MOUSE]	0.40	-0.34	0.03	-0.44	0.00	0.09	-0.07	0.36	0.16	0.71	0.49	0.28	0.05
E9Q151	Transporter OS=Mus musculus GN=Slc6a8 PE=2 SV=1 - [E9Q151_MOUSE]	0.18	0.70	0.04	0.98	0.03	0.81	-0.09	0.05	0.02	-0.38	0.18	0.29	0.05
P21279	Guanine nucleotide-binding protein G(q) subunit alpha OS=Mus musculus GN=Gnaq PE=1 SV=4 - [GNAQ_MOUSE]	0.12	-0.20	0.08	-0.13	0.00	-0.25	0.26	-0.03	0.10	0.09	0.13	0.06	0.05
Q6GQT9	Nodal modulator 1 OS=Mus musculus GN=Nomo1 PE=1 SV=1 - [NOMO1_MOUSE]	-0.07	-0.12	0.09	-0.10	-0.03	-0.13	0.14	0.02	-0.06	0.03	0.00	0.09	0.05
Q8R205	Zinc finger CCCH domain-containing protein 10 OS=Mus musculus GN=Zc3h10 PE=2 SV=1 - [ZC3HA_MOUSE]	-0.17	0.60	-0.13	0.54	0.25	0.27	-0.29	-0.20	0.15	-0.21	0.05	-0.19	0.05
O35071	Kinesin-like protein KIF1C OS=Mus musculus GN=Kif1c PE=1 SV=2 - [KIF1C_MOUSE]	0.13	-0.10	0.56	-0.05	0.05	-0.24	-0.06	1.14	0.39	1.81	0.25	1.18	0.05
Q4VA53	Sister chromatid cohesion protein PDS5 homolog B OS=Mus musculus GN=Pds5b PE=1 SV=1 - [PDS5B_MOUSE]	0.00	-0.07	0.19	-0.05	0.12	0.18	-0.05	0.08	-0.25	-0.06	-0.27	-0.04	0.05
P18894	D-amino-acid oxidase OS=Mus musculus GN=Dao PE=1 SV=3 - [OXDA_MOUSE]	-0.24	-0.47	0.05	-0.14	-0.10	-0.45	0.14	0.13	-0.24	0.24	-0.23	0.21	0.05

Q3TFP0	Serine/arginine-rich-splicing factor 10 OS=Mus musculus GN=Srsf10 PE=2 SV=1 [Q3TFP0_MOUSE]	-0.18	-0.53	-0.59	-0.50	0.06	-0.56	-0.14	-0.36	0.04	-0.28	0.17	0.08	0.05
Q922P9	Putative oxidoreductase GLYR1 OS=Mus musculus GN=Glyr1 PE=2 SV=1 - [GLYR1_MOUSE]	0.34	-0.11	0.19	-0.06	0.13	0.24	0.16	-0.14	-0.38	-0.13	0.00	-0.12	0.05
A2RT62	F-box/LRR-repeat protein 16 OS=Mus musculus GN=Fbxl16 PE=2 SV=1 - [FXL16_MOUSE]	-0.05	-0.19	0.00	-0.14	-0.10	-0.25	0.20	0.34	-0.19	0.09	-0.12	0.25	0.05
Q6P9N1	Hyccin OS=Mus musculus GN=Fam126a PE=1 SV=2 - [HYCC_MOUSE]	0.84	0.47	-0.28	-0.13	-0.10	0.00	-0.32	-0.83	-0.17	-0.19	-0.05	-0.43	0.05
Q8CI04	Conserved oligomeric Golgi complex subunit 3 OS=Mus musculus GN=Cog3 PE=1 SV=3 - [COG3_MOUSE]	0.30	0.22	0.24	-0.12	-0.05	-0.08	0.04	-0.07	-0.33	-0.07	-0.06	-0.30	0.05
Q9DBF1	Alpha-aminoadipic semialdehyde dehydrogenase OS=Mus musculus GN=Aldh7a1 PE=1 SV=4 [AL7A1_MOUSE]	-0.09	0.01	0.09	-0.06	-0.07	-0.02	0.18	0.00	0.03	0.01	0.03	0.14	0.05
Q61768	Kinesin-1 heavy chain OS=Mus musculus GN=Kif5b PE=1 SV=3 - [KINH_MOUSE]	0.07	-0.05	0.23	-0.01	0.01	0.04	0.04	-0.09	-0.13	-0.01	-0.14	-0.03	0.05
F6ZRK3	Excitatory amino acid transporter 2 (Fragment) OS=Mus musculus GN=Sic1a2 PE=4 SV=1 - [F6ZRK3_MOUSE]	0.37	-1.17	0.18	-0.35	0.23	-0.92	0.09	0.28	0.05	0.60	1.39	0.32	0.05
D3Z6H3	Dynactin 6, isoform CRA_b OS=Mus musculus GN=Dctn6 PE=4 SV=1 - [D3Z6H3_MOUSE]	-0.79	0.03	-0.17	0.10	-0.18	-0.08	-0.03	0.08	0.41	0.03	0.31	0.12	0.05
A2AEB4	Mortality factor 4-like protein 2 (Fragment) OS=Mus musculus GN=Morf4l2 PE=2 SV=1 - [A2AEB4_MOUSE]	0.32	0.69	-0.34	0.13	-0.26	0.28	-0.65	-0.68	0.00	-0.26	0.14	-0.49	0.05
Q4G0F8-2	Isoform 2 of Ubinuclein-1 OS=Mus musculus GN=Ubn1 - [UBN1_MOUSE]	0.43	0.08	0.05	0.17	0.01	0.06	-0.20	0.11	-0.12	0.04	-0.19	0.05	0.05
P48774	Glutathione S-transferase Mu 5 OS=Mus musculus GN=Gstm5 PE=1 SV=1 - [GSTM5_MOUSE]	-0.08	-0.17	-0.05	-0.01	-0.07	-0.07	0.22	0.15	-0.08	0.05	-0.11	0.12	0.05