

Supplementary Table 1. Full list of proteins identified by mass spectrometry from SILAC experiment with WT (light) 3xMBT pulldown as compared to O355N mutant (heavy) pulldown assay.

Protein ID	Gene Symbol	Description	55364.8		1.7		1419987.2		1.7		Sum	Corrected_max_ratio_H	Corrected_max_ratio_L		
			Number of peptides	max_heavy/median	max_light/median	Corrected_max_ratio_H	Corrected_max_ratio_L	max_heavy/sum	max_light/sum						
A2AEV7_A2AEV7_MOUSE	Cdccl20	Coiled-coil domain containing 120 OS-Mus musculus GN-Cdccl20 PE1 SV-1	1	50.5	1.9	3.2	26.9	15.07	0.06	59.1	1.9	3.2	26.9	15.98	0.06
Q8B7V7_STAC3_MOUSE	Stac3	SH3 and cysteine-rich domain-containing protein 3 OS-Mus musculus GN-Stac3 PE2 SV-1	1	39.1	2.5	4.2	24.1	14.67	0.07	59.1	2.5	4.1	24.1	14.33	0.07
Q8B7V7_MORC2_MOUSE	Morc2	Cell surface glycoprotein 2 OS-Mus musculus GN-Morc2 PE2 SV-1	1	31.7	1.9	15.5	9.6	9.62	0.11	31.7	1.9	15.5	9.6	9.62	0.11
Q8B7V7_RL32_MOUSE	Rpl32	60S ribosomal protein L22-like OS-Mus musculus GN-Rpl32 PE1 SV-1	1	45.0	3.8	6.5	14.6	5.53	0.12	45.0	3.8	6.5	14.6	5.53	0.12
Q8B807_MORC4_MOUSE	Morc4	MORC family CW-type zinc finger protein 4 OS-Mus musculus GN-Morc4 PE2 SV-2	1	17.1	1.2	2.1	13.8	8.07	0.12	17.1	1.2	2.1	13.8	8.23	0.12
Q1U197_DENN1B_MOUSE	Denn1b	DENN domain-containing protein 1B OS-Mus musculus GN-Denn1b PE2 SV-2	1	73.3	5.6	9.7	13.0	7.58	0.13	73.3	5.6	9.7	13.0	7.73	0.13
Q8P9M1_SFK_MOUSE	Sfk	Lytic enzyme OS-Mus musculus GN-Sfk PE1 SV-1	2	64.0	5.5	9.5	11.6	6.76	0.15	128.0	11.6	18.6	11.6	6.89	0.15
Q8Q2P7_P210L_MOUSE	Nup210	Nuclear pore membrane glycoprotein 210-like OS-Mus musculus GN-Nup210 PE2 SV-2	1	25.6	2.4	4.1	10.8	6.32	0.16	25.6	2.4	4.0	10.8	6.45	0.16
Q70491_STRAF6_MOUSE	Straf6	Stimulated by retinoic acid gene 6 protein OS-Mus musculus GN-Straf6 PE1 SV-1	1	15.4	1.5	2.5	10.5	6.10	0.16	15.4	1.5	2.5	10.5	6.22	0.16
Q80787_LVY5_MOUSE	Lvy5	Lymphocyte antigen 75 OS-Mus musculus GN-Lvy5 PE1 SV-2	1	25.3	2.8	4.9	8.9	5.39	0.19	25.3	2.8	4.9	8.9	5.29	0.19
Q82582_TAK1L_MOUSE	Tak1l	Serine/threonine protein kinase TAK1 OS-Mus musculus GN-Tak1L PE1 SV-1	1	9.9	1.2	2.0	8.5	4.95	0.20	9.9	1.2	2.0	8.5	5.05	0.20
EP9VDA_EP9VDA_MOUSE	Akap13	Protein Akap13 (Fragment) OS-Mus musculus GN-Akap13 PE1 SV-1	1	15.3	2.0	3.4	7.6	4.45	0.22	15.3	2.0	3.4	7.6	4.54	0.22
Q9Q0X4_TRIP4_MOUSE	Tripp4	Activating signal cointegrator 1 OS-Mus musculus GN-Tripp4 PE1 SV-2	1	48.0	6.5	11.2	7.4	4.30	0.23	48.0	6.5	11.2	7.4	4.38	0.23
Q8B81U_LRW01_MOUSE	LRW01	Leucine-rich repeat WD repeat-containing protein 1 OS-Mus musculus GN-LRW01 PE1 SV-1	2	27.7	4.0	6.8	7.0	4.07	0.25	55.3	7.9	13.0	7.9	4.15	0.24
P11087_C0JAL_MOUSE	C0Jal	Genes alpha-10 chain OS-Mus musculus GN-C0Jal PE1 SV-1	2	70.9	10.4	17.8	6.8	3.99	0.25	141.7	20.7	34.8	6.8	4.07	0.25
Q6ZWV7_RL35_MOUSE	Rpl35	60S ribosomal protein L35 OS-Mus musculus GN-Rpl35 PE2 SV-1	5	570.8	85.2	146.1	6.7	3.91	0.26	2496.4	443.3	745.4	5.6	3.35	0.30
FRV9P0_FRV9P0_MOUSE	Tc37	Protein Tc37 OS-Mus musculus GN-Tc37 PE4 SV-1	5	12.7	2.0	3.4	6.4	3.75	0.27	69.6	17.2	28.9	4.0	2.41	0.27
PE1811_KAPK5_MOUSE	Kapks5	CaM-dependent protein kinase catalytic subunit 2 OS-Mus musculus GN-Kapks5 PE1 SV-2	1	15.1	2.4	4.1	6.4	3.73	0.27	15.1	2.4	4.0	6.4	3.80	0.26
P14148_RL7_MOUSE	Rpl7	60S ribosomal protein L7 OS-Mus musculus GN-Rpl7 PE1 SV-2	35	167.0	26.1	44.8	6.4	3.75	0.27	10436.2	1493.9	70.0	4.15	2.44	0.27
P17981_NARX2_MOUSE	Arx2a	T-cell acute-ADP-ribosyltransferase 1 OS-Mus musculus GN-Arx2a PE1 SV-1	1	17.0	2.7	4.6	6.4	3.71	0.27	17.0	2.7	4.5	6.4	3.78	0.26
Q61823_P0DC4_MOUSE	P0dc4	Progremmed cell death protein 4 OS-Mus musculus GN-P0dc4 PE1 SV-2	4	13.6	2.2	3.7	6.3	3.66	0.27	62.6	9.3	15.7	6.7	3.99	0.25
Q8P9V9_P0Y12_MOUSE	P0y12	FY corticosteroid 12 OS-Mus musculus GN-P0Y12 PE4 SV-1	1	17.0	2.2	3.4	6.2	3.62	0.28	17.0	2.2	3.4	6.2	3.69	0.27
Q8B8W1_Tbl1R_MOUSE	Tbl1R	F-box-like-WD repeat-containing protein Tbl1R1 OS-Mus musculus GN-Tbl1R1 PE1 SV-1	3	14.4	2.3	4.0	6.2	3.60	0.28	44.4	18.4	31.0	2.7	1.60	0.63
Q6ZWR4_Q6ZWR4_MOUSE	Rpl36	60S ribosomal protein L36 OS-Mus musculus GN-Rpl36 PE1 SV-1	10	21.9	3.6	6.2	6.0	3.52	0.28	99.9	19.20	32.2	9.9	2.94	0.34
Q82280_SGT_MOUSE	Sgt	Protein SGT OS-Mus musculus GN-Sgt PE1 SV-1	1	7.9	1.3	2.3	4.8	3.57	0.28	7.9	1.3	2.3	4.8	3.57	0.28
Q8E5U9_Q8E5U9_MOUSE	Sclt5a	Neuronal amino acid transporter ASCT2 OS-Mus musculus GN-Sclt5a PE1 SV-1	2	11.7	2.0	3.4	6.0	3.49	0.29	23.4	3.9	6.6	6.0	3.56	0.28
Q80850_ERR1_MOUSE	Erra	Steroid hormone receptor ERR1 OS-Mus musculus GN-Erra PE1 SV-4	1	8.7	1.5	2.5	5.9	3.44	0.29	8.7	1.5	2.5	5.9	3.51	0.28
Q8CH77_RPGF_MOUSE	Rpgpf	Rap guanine nucleotide exchange factor 2 OS-Mus musculus GN-Rpgpf PE1 SV-2	1	11.0	1.9	3.2	5.9	3.43	0.29	11.0	1.9	3.1	5.9	3.50	0.29
Q70RE1_PA2A_MOUSE	Pa2a	Phosphatidylinositol 4-kinase type 2B OS-Mus musculus GN-Pa2a PE1 SV-1	1	9.8	1.7	3.0	5.9	3.41	0.29	9.8	1.7	3.0	5.9	3.57	0.29
Q9QAY4_TOM40_MOUSE	Tom40	Mitochondrial import receptor subunit TOM40 homolog OS-Mus musculus GN-Tom40 MAJ SV-1	1	7.5	1.3	2.2	5.8	3.37	0.30	7.5	1.3	2.2	5.8	3.44	0.29
Q90CK7_AP3L1_MOUSE	Ap3l1	AP-3 complex subunit sigma 1 OS-Mus musculus GN-AP3L1 PE1 SV-2	3	49.3	7.9	13.9	5.7	3.34	0.30	265.5	53.3	8.9	5.0	2.97	0.34
Q79192_MRC2_MOUSE	Mrc2	Mitochondrial carrier MRC2 OS-Mus musculus GN-Mrc2 PE1 SV-1	3	9.7	1.6	2.7	5.9	3.33	0.30	29.1	3.6	6.2	5.9	3.39	0.31
Q82CM0_RL15_MOUSE	Rpl15	60S ribosomal protein L15 OS-Mus musculus GN-Rpl15 PE2 SV-4	23	123.4	21.9	37.5	6.6	3.29	0.30	3531.2	623.4	1048.3	5.7	3.37	0.30
Q61194_P3C2A_MOUSE	P3c2a	Phosphatidylinositol 4-phosphate 3-kinase C domain-containing subunit alpha OS-Mus musculus	1	7.2	1.3	2.2	5.5	3.21	0.31	7.2	1.3	2.2	5.5	3.27	0.31
Q61275_RL23A_MOUSE	Rpl23a	60S ribosomal protein L23A OS-Mus musculus GN-Rpl23a PE1 SV-1	21	131.5	24.2	41.2	5.4	3.17	0.32	632.9	122.34	205.7	5.2	3.08	0.32
Q82QW1_P134C_MOUSE	P134c	Protein FAM134C OS-Mus musculus GN-P134C PE1 SV-1	1	15.1	2.4	4.1	6.4	3.73	0.32	15.1	2.4	4.0	6.4	3.80	0.31
Q80R72_ZNF768_MOUSE	Znf768	Zinc finger protein 768 OS-Mus musculus GN-ZNF768 PE2 SV-1	1	7.2	1.3	2.3	5.4	3.14	0.32	7.2	1.3	2.3	5.4	3.20	0.31
Q54988_SUK_MOUSE	Suk	IR2C-like serine/threonine protein kinase OS-Mus musculus GN-Suk PE1 SV-2	1	6.8	1.3	2.2	5.4	3.14	0.32	6.8	1.3	2.1	5.4	3.20	0.31
Q8400C_PAN3_MOUSE	Pan3	PAB-dependent poly(A)-specific ribonuclease subunit PAN3 OS-Mus musculus GN-Pan3 PE2 SV-3	1	6.0	1.1	1.9	5.4	3.12	0.32	6.0	1.1	1.9	5.4	3.19	0.31
Q82QD0_E9BP8_MOUSE	E9bp8	Protein E9BP8 OS-Mus musculus GN-E9BP8 PE1 SV-1	1	14.9	2.8	4.8	5.3	3.12	0.32	13.7	1.9	3.1	5.6	3.17	0.32
Q80B23_RL37_MOUSE	Rpl37	60S ribosomal protein L37 OS-Mus musculus GN-Rpl37 PE2 SV-3	1	256.9	48.3	82.8	5.3	3.10	0.32	256.9	48.3	81.2	5.3	3.16	0.32
Q90KW0_AR6P1_MOUSE	Ar6p1	AR6-ribosylation factor-like protein 6-interacting protein 1 OS-Mus musculus GN-Ar6p1 PE1 SV-1	2	21.6	4.1	7.0	5.3	3.10	0.32	43.1	8.1	13.7	5.3	3.16	0.32
Q82C93_RL14_MOUSE	Rpl14	60S ribosomal protein L14 OS-Mus musculus GN-Rpl14 PE1 SV-1	1	141.7	26.4	45.9	6.2	3.09	0.32	616.2	122.73	206.2	6.2	3.69	0.32
Q90R91_REEP3_MOUSE	Reep3	Receptor expression-enhancing protein 3 OS-Mus musculus GN-Reep3 PE1 SV-1	1	7.6	1.4	2.5	5.3	3.08	0.32	7.6	1.4	2.4	5.3	3.14	0.32
Q64449_RMR2_MOUSE	Rmr2	C-type mannase receptor 2 OS-Mus musculus GN-Rmr2 PE1 SV-3	1	55.2	10.5	17.9	5.3	3.08	0.32	55.2	10.5	17.6	5.3	3.14	0.32
9P7788_XRN1_MOUSE	Xrn1	5'-3' exonuclease OS-Mus musculus GN-Xrn1 PE1 SV-1	3	13.4	2.6	4.4	5.2	3.04	0.33	40.6	8.2	13.8	5.0	2.95	0.34
Q80R02_WNLC_MOUSE	Wnlc	Protein wnlc homolog OS-Mus musculus GN-Wnlc PE2 SV-1	1	12.9	2.5	4.3	5.9	3.03	0.33	10.1	1.9	3.1	5.9	3.18	0.31
Q8B8V9_CHT18_MOUSE	Chit18	Chromosome transmission fidelity protein 18 homolog OS-Mus musculus GN-Chit18 PE2 SV-2	2	14.2	2.8	4.7	5.2	3.01	0.33	28.5	5.5	9.3	5.2	3.07	0.33
Q80F6E_FAM5A_MOUSE	Fam5a	Protein FAM5A OS-Mus musculus GN-Fam5a PE1 SV-2	1	8.2	1.6	2.7	5.2	3.01	0.33	8.2	1.6	2.7	5.2	3.07	0.33
P47915_RL29_MOUSE	Rpl29	60S ribosomal protein L29 OS-Mus musculus GN-Rpl29 PE2 SV-2	10	114.8	22.4	38.5	5.4	2.99	0.33	2507.1	469.0	788.5	5.3	3.18	0.31
P10201_VAKH1_MOUSE	Vakh1	Vakp1 OS-Mus musculus GN-Vakh1 PE2 SV-4	3	17.7	3.5	5.7	5.1	2.98	0.34	53.6	11.8	19.8	5.1	3.04	0.33
P20357_MAP2_MOUSE	Map2	Microtubule-associated protein 2 OS-Mus musculus GN-Map2 PE1 SV-2	1	24.8	4.9	8.4	5.1	2.96	0.34	24.8	4.9	8.2	5.1	3.02	0.33
Q80B79_ZAM1_MOUSE	Zam1	Junctional adhesion molecule 4 OS-Mus musculus GN-Zam1 PE1 SV-2	2	16.2	3.2	5.5	5.0	2.92	0.34	32.3	6.4	10.8	5.0	2.98	0.34
Q82111_RHB_MOUSE	Rhb	CTP-binding protein RHB OS-Mus musculus GN-Rhb PE1 SV-1	1	23.6	4.7	8.1	6.2	2.92	0.34	23.6	4.7	8.1	6.2	2.98	0.34
P84059_RL19_MOUSE	Rpl19	60S ribosomal protein L19 OS-Mus musculus GN-Rpl19 PE4 SV-1	21	57.0	11.5	19.6	5.0	2.90	0.34	5900.9	1047.8	1761.8	5.6	3.35	0.30
Q81040_K8951_MOUSE	Kia08951	Uncharacterized protein KIA08951-like OS-Mus musculus GN-Kia08951 PE2 SV-1	1	17.2	3.5	5.9	5.0	2.89	0.35	17.2	3.5	5.8	5.0	2.95	0.34
Q80730_GEM7B_MOUSE	Gem7b	Uncharacterized protein OS-Mus musculus GN-Gem7B PE1 SV-1	5	112.4	22.7	38.9	5.0	2.89	0.35	853.9	250.5	421.1	3.4	2.03	0.49
Q9N902_E4L11_MOUSE	E4l11	Band 4, like protein 1 OS-Mus musculus GN-E4L11 PE1 SV-1	1	15.1	2.8	4.9	5.0	2.89	0.35	15.1	2.8	4.9	5.0	2.96	0.34
Q82911_RL32_MOUSE	Rpl32	60S ribosomal protein L32 OS-Mus musculus GN-Rpl32 PE1 SV-2	10	59.0	12.1	20.8	4.9	2.84	0.35	2299.7	444.5	747.5	5.2	3.08	0.33
Q80X95_RAGA_MOUSE	Raga	Ras-related GTP-binding protein A OS-Mus musculus GN-Raga PE2 SV-1	1	7.6	1.6	2.7	4.8	2.82	0.35	7.6	1.6	2.7	4.8	2.87	0.35
Q51500_SNE1_MOUSE	Sne1	Synaptotagmin 1 OS-Mus musculus GN-Sng1 PE1 SV-1	1	14.9	2.8	4.8	5.3	2.81	0.36	14.9	2.8	4.8	5.3	2.86	0.35
Q82W08_RL10_MOUSE	Rpl10	60S ribosomal protein L10 OS-Mus musculus GN-Rpl10 PE1 SV-3	20	57.2	11.9	20.4	4.8	2.80	0.36	1945.8	529.9	891.0	3.7	2.18	0.46
Q90C49_NCBP2_MOUSE	Ncbp2	Nuclear cap-binding protein subunit 2 OS-Mus musculus GN-Ncbp2 PE1 SV-1	1	147.2	30.6	52.5	4.8	2.80	0.36	147.2	30.6	51.5	4.8	2.86	0.35
Q60775_EF1_MOUSE	Ef1	ETS-related transcription factor EF1 OS-Mus musculus GN-Ef1 PE2 SV-1	1	7.1	1.5	2.5	4.8	2.80	0.36	7.1	1.5	2.5	4.8	2.86	0.35
P12759_RL3_MOUSE	Rpl3	60S ribosomal protein L3 OS-Mus musculus GN-Rpl3 PE1 SV-1	47	108.0	20.3	38.3	5.3	2.79	0.36	645.8	144.58	243.8	5.3	3.07	0.34
Q1U332_ZF178_MOUSE	Z														

O35125_LRCC2_MOUSE	Lrrc23	Leucine-rich repeat-containing protein 23 OS=Mus musculus GN=Lrrc23 PE=2 SV=1	1	10.4	2.6	4.4	4.0	2.36	0.42	10.4	2.6	4.3	4.0	2.40	0.42
E9B8R4_E9B8R4_MOUSE	Ard1b3	Protein Ard1b3 OS=Mus musculus GN=Ard1b3 PE=1 SV=1	6	8.5	2.1	3.6	4.0	2.35	0.43	50.4	14.5	24.4	3.5	2.06	0.49
P53032_RL10A_MOUSE	Rpl10a	60S ribosomal protein L10 OS=Mus musculus GN=Rpl10a PE=1 SV=3	6	16.65	36.4	4.4	4.0	2.35	0.43	5088.3	1304.1	2159.3	2.8	1.69	0.43
Q8B20D_KPCCD_MOUSE	Prlkd2	Serine/threonine protein kinase D2 OS=Mus musculus GN=Prlkd2 PE=1 SV=1	1	14.1	3.5	6.0	4.0	2.33	0.43	14.1	3.5	5.9	4.0	2.38	0.42
Q9Q614_Q9Q614_MOUSE	Chd3	Protein Chd3 OS=Mus musculus GN=Chd3 PE=1 SV=3	9	11.8	3.0	5.0	4.0	2.33	0.43	128.8	43.0	72.4	3.0	1.78	0.56
Q92218_SUCB2_MOUSE	Chd3	Chromatin ligase (GDP-forming) mitochondrial OS=Mus musculus GN=Sucb2 PE=1 SV=1	3	7.9	2.0	3.4	4.0	2.31	0.43	7.9	2.0	3.4	4.0	2.36	0.43
Q8R78_TF3CS_MOUSE	Gtf3s5	General transcription factor 3C polypeptide3 OS=Mus musculus GN=Gtf3s5 PE=2 SV=2	6	16.1	4.1	7.0	4.0	2.31	0.43	148.1	24.3	40.9	3.9	2.30	0.44
Q9JH55_MTLCL_MOUSE	Mtcl1	Mitochondrial subunit 1 OS=Mus musculus GN=Mtcl1 PE=1 SV=1	7	12.0	3.0	5.2	4.0	2.31	0.43	7.5	2.21	37.2	3.4	2.02	0.49
Q9U3T1_4F2_MOUSE	Elf1em1	Eukaryotic translation initiation factor 4E transporter OS=Mus musculus GN=Elf1em1 PE=1 SV=1	1	7.8	2.0	3.4	4.0	2.31	0.43	7.8	2.0	3.3	4.0	2.35	0.42
Q9VW72_THA1_MOUSE	Eif1	50S ribosome-related transmembrane protein 1 OS=Mus musculus GN=Eif1 PE=1 SV=1	2	11.2	3.3	5.2	4.0	2.31	0.43	11.2	3.3	5.2	4.0	2.35	0.49
Q9R4QD_WDR33_MOUSE	Wdr33	Core RNA 3' end processing protein WDR33 OS=Mus musculus GN=Wdr33 PE=1 SV=1	7	13.8	3.5	6.0	3.9	2.30	0.43	107.7	36.8	61.9	2.9	1.74	0.57
Q9R9L4_CSN1_MOUSE	Gps1	CCP1 signalosome complex subunit 1 OS=Mus musculus GN=Gps1 PE=1 SV=1	6	11.6	2.9	5.0	3.9	2.30	0.43	175.1	58.4	31.0	4.1	2.42	0.41
Q85C96_PFC1_MOUSE	Pfc1	Enhancer of polycomb target 1 OS=Mus musculus GN=Pfc1 PE=2 SV=1	5	7.2	1.9	3.1	3.9	2.30	0.43	5.9	1.8	3.9	3.8	2.08	0.48
Q8R4T9_PFC1_MOUSE	Sic14a2	Urea transporter 2 OS=Mus musculus GN=Sic14a2 PE=2 SV=2	1	9.5	2.4	4.9	2.9	2.29	0.44	9.5	2.4	4.1	3.9	2.34	0.43
PE2281_RS11_MOUSE	Rps11	40S ribosomal protein S11 OS=Mus musculus GN=Rps11 PE=1 SV=3	7	24.5	6.2	10.7	3.9	2.29	0.44	660.4	236.5	397.6	2.8	1.66	0.60
Q9D101_MBP1_MOUSE	Mbp1	MARCKS-12-binding inhibitory protein 1 OS=Mus musculus GN=Mbp1 PE=1 SV=1	2	14.3	3.6	6.3	3.9	2.29	0.44	28.6	7.3	12.3	3.9	2.34	0.43
Q955U6_BRO4_MOUSE	Bro4	Bromodomain-containing protein 4 OS=Mus musculus GN=Bro4 PE=1 SV=2	16	12.5	3.2	5.5	3.9	2.28	0.44	245.5	78.0	121.1	3.1	1.87	0.53
Q8R4D1_SCAF8_MOUSE	Scaf8	Protein SCAF8 OS=Mus musculus GN=Scaf8 PE=1 SV=1	12	14.3	3.6	6.2	3.9	2.28	0.44	275.8	85.9	144.5	3.2	1.91	0.52
Q8R4K9_RB1M2_MOUSE	Rbm12	Ribosomal protein L2 OS=Mus musculus GN=Rbm12 PE=1 SV=1	2	10.6	2.7	4.7	3.9	2.28	0.44	21.3	5.4	9.1	3.9	2.32	0.43
Q8VDG3_PARN_MOUSE	Parn	Poly(A)-specific ribonuclease PARN OS=Mus musculus GN=Parn PE=1 SV=1	3	10.7	2.7	4.7	3.9	2.27	0.44	28.0	8.2	15.7	3.4	2.04	0.49
PE1255_RL3E_MOUSE	Rpl3e	60S ribosomal protein L24 OS=Mus musculus GN=Rpl3e PE=1 SV=1	2	20.5	4.1	123.6	3.9	2.27	0.44	8359.1	2298.7	3865.3	3.6	2.16	0.46
Q9G17_FAM21_MOUSE	Fam21	WASH complex subunit FAM21 OS=Mus musculus GN=Fam21 PE=1 SV=1	22	11.4	2.9	5.0	3.9	2.27	0.44	283.9	76.8	129.2	3.7	2.20	0.46
AG5H24_E6C6_MOUSE	Exc6b	Exonucyst complex component 6 OS=Mus musculus GN=Exc6b PE=2 SV=1	1	11.3	2.9	5.0	3.9	2.27	0.44	11.3	2.9	4.9	3.9	2.31	0.43
PE2918_RL8_MOUSE	Rpl8	60S ribosomal protein L8 OS=Mus musculus GN=Rpl8 PE=1 SV=2	22	12.4	3.2	5.3	3.9	2.27	0.44	75.0	1743.6	2931.7	4.3	2.57	0.39
Q61543_GSLG_MOUSE	Glg1	Golgi apparatus protein 1 OS=Mus musculus GN=Glg1 PE=1 SV=1	5	19.2	4.9	8.5	3.9	2.26	0.44	99.8	26.8	45.1	3.7	2.21	0.43
A2RSX9_A2RSX9_MOUSE	Arfg1	ArfGTPase-1 OS=Mus musculus GN=Arfg1 PE=2 SV=1	1	30.4	7.8	13.4	3.9	2.26	0.44	30.4	7.8	13.2	3.9	2.31	0.43
Q920D3_MED2_MOUSE	Med28	Mediator of RNA polymerase II transcription subunit 28 OS=Mus musculus GN=Med28 PE=2 SV=1	2	9.5	2.5	4.2	3.9	2.26	0.44	19.0	4.9	8.3	3.9	2.32	0.43
Q9R9Y7_CT12_MOUSE	Rpl12	Ribosomal protein L12 OS=Mus musculus GN=Rpl12 PE=2 SV=3	2	8.6	2.2	3.6	3.9	2.26	0.44	8.6	2.2	3.6	3.9	2.30	0.43
Q9R917_R121_MOUSE	Rpl21	60S ribosomal protein L21 OS=Mus musculus GN=Rpl21 PE=2 SV=2	19	51.5	13.3	22.7	3.9	2.26	0.44	2606.8	697.1	1172.2	3.7	2.21	0.43
Q9R919_OV9D_MOUSE	Olfir791	Olfactory receptor 791 OS=Mus musculus GN=Olfir791 PE=1 SV=1	1	20.8	5.4	9.2	3.9	2.26	0.44	20.8	5.4	9.0	3.9	2.30	0.43
Q9R9Y7_Q8R9Y7_MOUSE	Arhg151	Protein Arhg151 OS=Mus musculus GN=Arhg151 PE=1 SV=1	5	13.6	3.5	6.0	3.9	2.26	0.44	17.4	5.1	9.0	3.9	2.30	0.43
Q9G303_VO2C2_MOUSE	Voc2	Voltage-dependent anion-selective channel protein 2 OS=Mus musculus GN=Voc2 PE=1 SV=2	6	12.5	3.6	6.3	3.9	2.26	0.44	29.1	8.7	14.2	3.5	2.07	0.48
Q9E16_Q9E16_MOUSE	Rhnk3	Protein Rhnk3 OS=Mus musculus GN=Rhnk3 PE=1 SV=1	166	29.9	7.7	13.3	3.9	2.25	0.44	1750.0	1718.3	2889.4	3.4	2.62	0.38
Q9Y1K2_RRP19_MOUSE	Rrp19	Ribosomal RNA processing protein 1 homolog B OS=Mus musculus GN=Rrp19 PE=1 SV=2	2	11.1	2.9	4.9	3.9	2.25	0.44	72.2	5.8	9.7	3.9	2.29	0.44
Q9VCR7_AHEH2_MOUSE	Aheh2	Alpha-helical coiled-coil domain-containing protein 2 OS=Mus musculus GN=Aheh2 PE=2 SV=1	2	23.4	6.1	10.5	3.9	2.25	0.44	121.0	46.8	29.4	3.8	2.19	0.48
P39053_DYNI_MOUSE	Dnm1	Dynamin-1 OS=Mus musculus GN=Dnm1 PE=1 SV=2	32	23.0	6.0	10.3	3.8	2.24	0.45	1081.4	584.7	981.1	1.8	1.10	0.91
Q9R9L4_RL11_MOUSE	Rpl11	60S ribosomal protein L11 OS=Mus musculus GN=Rpl11 PE=1 SV=4	15	72.9	19.0	32.6	3.8	2.24	0.45	4355.7	1357.4	2282.4	3.2	1.91	0.52
Q61687_ATRX_MOUSE	Atrx	Transcriptional regulator ATRX OS=Mus musculus GN=Atrx PE=1 SV=3	26	16.2	4.2	7.3	3.8	2.23	0.45	462.3	126.3	119.4	3.7	2.18	0.46
Q8R811_TOX4_MOUSE	Tox4	TOX high mobility group family member 4 OS=Mus musculus GN=Tox4 PE=1 SV=3	9	20.6	5.4	9.2	3.8	2.23	0.45	191.3	56.7	91.4	3.8	2.18	0.46
E9Q4F1_E9Q4F1_MOUSE	Ankrk11	Protein Ankrk11 OS=Mus musculus GN=Ankrk11 PE=4 SV=2	3	13.4	3.5	6.0	3.8	2.23	0.45	45.1	11.8	21.9	3.8	2.26	0.44
QJUEB3_PUF60_MOUSE	Puf60	Poly(A)-binding interacting factor PUF60 OS=Mus musculus GN=Puf60 PE=2 SV=2	25	30.2	7.9	13.5	3.8	2.23	0.45	85.2	23.3	37.5	1.8	1.27	0.44
QJ31A1_SPH2_MOUSE	Spht2	Sphingosine kinase 2 OS=Mus musculus GN=Spht2 PE=1 SV=1	3	6.7	1.9	3.2	3.8	2.23	0.45	6.7	1.9	3.5	3.8	2.23	0.45
Q9ZQL0_RIPK3_MOUSE	Ripk3	Receptor-interacting serine/threonine protein kinase 3 OS=Mus musculus GN=Ripk3 PE=1 SV=2	6	13.7	3.6	6.1	3.8	2.23	0.45	75.2	23.7	39.9	3.2	1.89	0.53
Q9U70_Q9U70_MOUSE	Nuam1	Protein Nuam1 OS=Mus musculus GN=Nuam1 PE=1 SV=1	196	53.2	13.9	23.9	3.8	2.23	0.45	16854.2	4383.6	7370.9	3.8	2.29	0.44
QJ3PH1_PRRC1_MOUSE	Prrc1	Protein PRRC1 OS=Mus musculus GN=Prrc1 PE=2 SV=1	3	17.7	4.6	7.9	3.8	2.22	0.45	45.8	14.8	25.0	3.1	1.84	0.54
PE8372_TB84B_MOUSE	Tb84b	Tubulin beta 4b chain OS=Mus musculus GN=Tb84b PE=1 SV=1	11	20.3	5.2	9.8	3.8	2.22	0.45	26.8	8.0	14.3	3.8	2.15	0.46
QJ343_BAB1_MOUSE	Babam1	BRIC9 and BRCA1-A complex member 1 OS=Mus musculus GN=Babam1 PE=2 SV=1	1	31.1	8.2	14.0	3.8	2.22	0.45	31.1	8.2	13.7	3.8	2.27	0.45
Q8R922_CIR1A_MOUSE	Cir1a	CIR1A OS=Mus musculus GN=Cir1a PE=2 SV=3	3	12.1	3.2	5.5	3.8	2.22	0.45	41.6	11.5	19.4	3.6	2.15	0.47
Q8W0F4_R120B_MOUSE	R120b	Kinase-like protein K120B OS=Mus musculus GN=R120B PE=1 SV=2	3	12.1	3.2	5.5	3.8	2.21	0.45	30.5	8.2	15.0	3.8	2.18	0.46
PE3743_WNK1_MOUSE	Wnk1	Serine/threonine protein kinase WNK1 OS=Mus musculus GN=Wnk1 PE=1 SV=2	2	9.2	2.4	4.1	3.8	2.21	0.45	18.3	4.8	8.1	3.8	2.25	0.44
Q90394_RUFY3_MOUSE	Rufy3	Protein RUFY3 OS=Mus musculus GN=Rufy3 PE=1 SV=1	2	11.8	3.1	5.4	3.8	2.21	0.45	23.7	6.3	10.5	3.8	2.25	0.44
Q9DCC5_T1AP1_MOUSE	T1ap1	Tight junction-associated protein 1 OS=Mus musculus GN=T1ap1 PE=1 SV=1	7	8.8	2.3	4.0	3.8	2.21	0.45	81.3	22.5	37.8	3.6	2.15	0.47
PE1225_RAP2B_MOUSE	Rap2b	Rap guanine nucleotide exchange factor 2 OS=Mus musculus GN=Rap2b PE=1 SV=2	2	11.5	3.1	5.2	3.8	2.21	0.45	2.9	2.0	3.8	3.8	2.15	0.44
Q92115_COG2_MOUSE	Cog2	Conserved oligomeric Golgi complex subunit 2 OS=Mus musculus GN=Cog2 PE=1 SV=2	4	12.2	3.2	5.5	3.8	2.20	0.46	44.1	17.4	29.3	2.5	1.50	0.61
Q9R912_PSD11_MOUSE	Pamd11	26S proteasome non-ATPase regulatory subunit 11 OS=Mus musculus GN=Pamd11 PE=1 SV=1	8	18.7	5.0	8.5	3.8	2.19	0.46	146.0	46.3	77.9	3.2	1.87	0.53
QJ3JH6_AK1_MOUSE	Aks1	AK2-associated protein kinase 1 OS=Mus musculus GN=Aks1 PE=1 SV=2	10	12.4	3.3	5.7	3.8	2.19	0.46	146.7	40.0	67.3	3.7	2.18	0.46
P14094_AT1B1_MOUSE	At1b1	ATP-binding cassette transporter 1 OS=Mus musculus GN=At1b1 PE=1 SV=1	11	11.3	8.4	14.3	3.8	2.19	0.46	11.3	8.4	14.3	3.8	2.19	0.46
Q9P415_NUCL_MOUSE	Ncl	Nucleolin OS=Mus musculus GN=Ncl PE=1 SV=2	27	57.2	15.3	26.2	3.7	2.18	0.46	1818.1	468.0	787.0	3.9	2.31	0.43
Q92211_RBP1_MOUSE	Rbp1	Arp3 complex subunit beta-1 OS=Mus musculus GN=Rbp1 PE=1 SV=2	18	12.5	3.5	5.7	3.7	2.18	0.46	260.4	60.5	101.8	4.3	2.56	0.39
Q85L13_K072_MOUSE	K072	CDK-NOF transcription-inhibiting protein 2 OS=Mus musculus GN=K072 PE=2 SV=2	12	16.9	4.6	8.2	3.7	2.17	0.46	92.1	24.7	52.1	3.7	2.16	0.46
Q8QAK2_ZZZ2_MOUSE	Zzz2	ZZZ2 zinc finger-containing protein 1 OS=Mus musculus GN=Zzz2 PE=1 SV=2	2	14.5	3.9	6.7	3.7	2.17	0.46	78.8	19.8	33.3	3.8	2.27	0.44
Q81E0D_STX5_MOUSE	Stx5	Syntaxin-5 OS=Mus musculus GN=Stx5 PE=1 SV=3	3	16.1	4.3	7.4	3.7	2.17	0.46	49.8	15.5	26.0	3.2	1.91	0.52
Q8Y95E_APA1_MOUSE	Apa1	Actin filament-associated protein 1 OS=Mus musculus GN=Apa1 PE=1 SV=1	5	13.8	3.7	6.4	3.7	2.16	0.46	72.0	24.2	47.7	2.5	1.51	0.66
Q83752_SBR4_MOUSE	Sbr4	3-O-phosphotyrosine phosphatase OS=Mus musculus GN=Sbr4 PE=1 SV=1	15	19.5	5.1	8.6	3.7	2.16	0.46	221.5	61.6	105.5	3.4	2.07	0.48
Q692N7_WOYF_MOUSE	Woyf	Myoferlin OS=Mus musculus GN=Woyf PE=1 SV=2	15	13.5	3.7	6.3	3.7	2.16	0.46	216.2	66.7	112.2	3.2	1.93	0.52
Q93310_PRI2_MOUSE	Prim2	DNA primase large subunit OS=Mus musculus GN=Prim2 PE=1 SV=1	12	18.8	5.1	8.7	3.7	2.16	0.46	239.1	68.7	115.5	3.5	2.07	0.48
Q9J2W1_XTB_MOUSE	Xtb	Synaptobrevin homolog XTB OS=Mus musculus GN=Xtb PE=1 SV=1	3	13.9	3.8	6.5	3.7	2.15	0.46	34.1	12.6	21.2	2.7	1.61	0.62
Q8C9M6_INT4_MOUSE	Int4	Integrator complex subunit 4 OS=Mus musculus GN=Int4 PE=1 SV=1	4	15.0	4.4	7.6	3.7	2.15	0.46	26.8	76.8	26.8	3.8	2.07	0.51
PHF10	PHF10	PHD finger protein 1 OS=Mus musculus GN=PHF10 PE=1 SV=2	1	5.0	3.7	6.0	3.7	2.15	0.46	89.0	25.3	42.6	3.5	2.09	0.48
Q80882_GPB1_MOUSE	Gpbp1	GTP-binding protein 1 OS=Mus musculus GN=Gpbp1 PE=1 SV=4	4	19.1	5.2	8.9	3.7	2.15	0.47	69.1	19.1	32.2	3.6	2.15	0.47
Q92473_TF3_MOUSE	Tf3	Integrin alpha 3 OS=Mus musculus GN=Itga3 PE=1 SV=1	7	14.7	3.9	6.7	3.7	2.15	0.47	14.7	3.8	6.8	3.4	2.02	0.49
QJ3U29_LRFL1_MOUSE	Lrrf1a	Leucine-rich repeat flightless-interacting protein 1 OS=Mus musculus GN=Lrrf1a PE=1 SV=2	1	5.5	1.5	2.6	3.7	2.15	0.47	5.5	1.5	2.5	3.7	2.19	0.46
P41105_RL28_MOUSE	Rpl28	60S ribosomal protein L28 OS=Mus musculus GN=Rpl28 PE=1 SV=2	19	107.2	29.1	49.9	3.7	2.15	0.47	4167.7	848.7	1427.1	4.9	2.92	0.34
Q9CWC3_C2D2_MOUSE	C2d2p2	CD2 antigen cytoplasmic tail-binding protein 2 OS=Mus musculus GN=C2d2p2 PE=1 SV=1	2	10.2	2.8	4.									

QB407_YIPF4_MOUSE	Yipf4	Protein YIPF4 OS-Mus musculus GN-Yipf4 PE-2 SV-1	1	42.7	12.5	21.5	3.4	1.99	0.50	25.7	12.5	21.1	3.4	2.03	0.49
Q5187_CKIB_MOUSE	Ckib	ES S/MuJ protein ligase CB44 OS-Mus musculus GN-Ckib4 PE-1 SV-2	1	10.5	3.1	5.3	3.4	1.98	0.50	10.5	3.1	5.2	3.4	2.02	0.49
Q9N9W6_GREB3_MOUSE	Greb3	Glucocorticoid-inducible protein 3 beta OS-Mus musculus GN-Greb3 PE-1 SV-2	1	8.0	2.3	3.4	3.4	1.98	0.50	8.0	2.3	3.4	3.4	2.02	0.49
Q7QTK7_LMT2_MOUSE	Lmt2	Integrator complex subunit 7 OS-Mus musculus GN-Lmt2 PE-2 SV-2	18	11.7	3.4	5.9	3.4	1.98	0.50	252.9	80.8	135.9	3.1	1.86	0.54
PE2889_RL30_MOUSE	Rpl30	60S ribosomal protein L30 OS-Mus musculus GN-Rpl30 PE-2 SV-2	17	38.7	11.4	19.5	3.4	1.98	0.50	2555.7	772.1	1298.3	3.3	1.97	0.51
PE1514_RL37A_MOUSE	Rpl37a	60S ribosomal protein L37A OS-Mus musculus GN-Rpl37a PE-2 SV-2	7	118.8	34.9	58.4	3.4	1.98	0.50	1440.8	424.0	679.3	3.4	1.87	0.50
P45766_PSN1_MOUSE	Psn1	Presenilin-1 OS-Mus musculus GN-Psn1 PE-1 SV-1	1	6.0	1.8	3.0	3.4	1.98	0.51	6.0	1.8	3.0	3.4	2.02	0.50
QB867_RL24_MOUSE	Rpl24	60S ribosomal protein L24 OS-Mus musculus GN-Rpl24 PE-1 SV-2	15	42.9	12.6	21.7	3.4	1.98	0.51	3018.5	743.5	1250.1	4.1	2.41	0.41
Q55WV4_MED13_MOUSE	Med13	Mediator of RNA polymerase II transcription subunit 13 OS-Mus musculus GN-Med13 PE-2 SV-2	13	10.8	3.2	5.5	3.4	1.98	0.51	170.6	46.2	77.7	3.7	1.98	0.49
Q55WV6_ACKA_MOUSE	Acka	Adrenomedullin-like receptor 1 OS-Mus musculus GN-Acka PE-1 SV-1	1	19.5	5.6	9.8	3.4	1.98	0.51	195.3	48.1	78.9	3.7	1.98	0.49
QBK30_NSL1_MOUSE	Nsl1	Kinetochores-associated protein NSL1 homolog OS-Mus musculus GN-Nsl1 PE-2 SV-1	1	16.6	4.9	8.4	3.4	1.98	0.51	16.6	4.9	8.2	3.4	2.02	0.50
QBK27_INT10_MOUSE	Int10	Integrator complex subunit 10 OS-Mus musculus GN-Int10 PE-1 SV-3	8	14.6	4.3	7.4	3.4	1.97	0.51	133.4	40.9	68.8	3.3	1.94	0.52
QBVB18_BAMP_MOUSE	Bamp	Branched-chain amino acid aminotransferase 1 OS-Mus musculus GN-Bamp PE-1 SV-1	2	5.0	2.6	4.6	3.4	1.97	0.51	5.0	2.6	4.6	3.4	1.97	0.51
D3D209_D3D209_MOUSE	Dtd23	MGC18410, isoform CRA a OS-Mus musculus GN-Dtd23 PE-3 SV-3	22	15.4	4.6	7.8	3.4	1.97	0.51	105.5	183.1	307.8	2.8	1.65	0.65
QB8G55_MYP2T_MOUSE	Ppp1t2b	Protein phosphatase 1 regulatory subunit 12B OS-Mus musculus GN-Ppp1t2b PE-1 SV-2	1	4.7	1.4	2.4	3.4	1.97	0.51	4.7	1.4	2.4	3.4	2.01	0.50
Q9DBK5_CNOT6_MOUSE	Cnot6	BCRA-NOT transcription complex subunit 8 OS-Mus musculus GN-Cnot6 PE-2 SV-1	5	12.8	3.8	6.5	3.4	1.97	0.51	60.0	18.0	30.3	3.3	1.98	0.51
Q9DBV2_VGUL_MOUSE	Vgula	Vaginotransmembrane protein 1 OS-Mus musculus GN-Vgula PE-1 SV-1	84	43.4	12.9	22.1	3.4	1.96	0.51	4186.9	1438.3	2418.5	2.9	1.73	0.58
Q9IKX8_AFP3M1_MOUSE	Afp3m1	AP 3 complex subunit mu-1 OS-Mus musculus GN-Afp3m1 PE-1 SV-1	1	16.1	4.8	8.2	3.4	1.96	0.51	16.1	4.8	8.1	3.4	2.00	0.50
Q9D176_CTF2_MOUSE	Ctf2	Transcriptional repressor CTF2 OS-Mus musculus GN-Ctf2 PE-1 SV-2	6	21.5	6.4	11.0	3.4	1.96	0.51	122.1	41.9	70.5	2.9	1.73	0.58
Q9D177_TNCL_MOUSE	Tncl	CTX component protein 1 OS-Mus musculus GN-Tncl PE-1 SV-1	1	11.6	3.5	6.0	3.4	1.96	0.51	11.6	3.5	5.8	3.4	1.99	0.50
Q9DBE1_LMAN2_MOUSE	Lman2	Neurular integral membrane protein VPS30 OS-Mus musculus GN-Lman2 PE-2 SV-2	8	21.8	6.5	11.2	3.3	1.95	0.51	166.2	62.0	104.2	2.7	1.59	0.63
E9QAL5_E9QAL5_MOUSE	Inf2	Inverted form 2 OS-Mus musculus GN-Inf2 PE-4 SV-1	23	14.2	4.3	7.3	3.3	1.95	0.51	389.0	134.5	226.2	2.9	1.72	0.58
Q9P9M2_WDR6_MOUSE	Wdr6	WD repeat-containing protein 6 OS-Mus musculus GN-Wdr6 PE-2 SV-1	3	7.0	2.1	3.6	3.3	1.95	0.51	17.4	7.3	12.3	2.4	1.42	0.71
Q7D283_WNT2E_MOUSE	Wnt2b	Protein Wnt-2b OS-Mus musculus GN-Wnt2b PE-2 SV-2	1	62.8	18.8	32.2	3.3	1.95	0.51	62.8	18.8	31.6	3.3	1.99	0.50
Q9P1P5_NUDC_MOUSE	Nudc1	NudC domain-containing protein 1 OS-Mus musculus GN-Nudc1 PE-2 SV-2	1	13.3	4.0	6.8	3.3	1.95	0.51	13.3	4.0	6.7	3.3	1.99	0.50
Q9R0X0_MED20_MOUSE	Med20	Mediator of RNA polymerase II transcription subunit 20 OS-Mus musculus GN-Med20 PE-1 SV-2	2	27.4	8.2	14.1	3.3	1.95	0.51	148.8	16.4	27.6	3.3	1.98	0.50
Q3JTM9_Q3JTM9_MOUSE	Ghrf3c	General transcription factor IIIc, polyphosphatase 3 OS-Mus musculus GN-Ghrf3c PE-1 SV-2	9	8.7	2.6	4.5	3.3	1.95	0.51	86.6	39.6	66.6	2.4	1.45	0.69
Q61161_MAK2_MOUSE	Mak2	Tetraspanin-related protein kinase kinase kinase 2 OS-Mus musculus GN-Mak2 PE-1 SV-2	1	2.0	0.7	1.2	3.3	1.95	0.51	2.0	0.7	1.2	3.3	1.99	0.50
Q9N2K8_INT2_MOUSE	Int2	Integrator complex subunit 2 OS-Mus musculus GN-Int2 PE-2 SV-2	6	13.6	4.1	7.0	3.3	1.94	0.51	78.8	24.7	41.6	3.2	1.89	0.53
QB0111_GNL1_MOUSE	Gnl1	Guanine nucleotide-binding protein-like 3 OS-Mus musculus GN-Gnl1 PE-1 SV-2	6	15.8	4.7	8.1	3.3	1.94	0.51	102.3	25.3	42.5	4.1	2.41	0.41
Q90707_COPF_MOUSE	Copf	Coatomer subunit epsilon OS-Mus musculus GN-Copf PE-2 SV-3	1	56.6	17.0	29.1	3.3	1.94	0.51	603.8	167.6	510.2	3.1	1.77	0.79
Q90C29_C2D3_MOUSE	C2d3	COG1 domain-containing protein 43 OS-Mus musculus GN-C2d3 PE-1 SV-1	2	16.7	4.9	8.4	3.3	1.94	0.51	16.7	4.9	8.3	3.3	1.94	0.51
P59429_C2D3_MOUSE	Cgn	Cingulin OS-Mus musculus GN-Cgn PE-1 SV-1	5	9.7	2.9	5.0	3.3	1.94	0.52	9.9	25.7	43.3	2.3	1.38	0.72
Q90708_Q90708_MOUSE	S100A6	Protein S100A6 OS-Mus musculus GN-S100A6 PE-2 SV-1	1	9.5	2.9	4.9	3.3	1.94	0.52	9.5	2.9	4.8	3.3	1.98	0.51
Q90121_WNF23_MOUSE	Wnf23	Protein Wnf23 OS-Mus musculus GN-Wnf23 PE-1 SV-1	1	9.5	2.9	4.9	3.3	1.94	0.52	9.5	2.9	4.8	3.3	1.98	0.51
Q91210_AHRG1_MOUSE	Ahrg1	Rho guanine nucleotide exchange factor 1 OS-Mus musculus GN-Ahrg1 PE-1 SV-2	13	10.1	3.1	5.2	3.3	1.94	0.52	137.4	45.3	76.2	3.0	1.80	0.55
Q919C3_ELF2_MOUSE	Elf2	ETS-related transcription factor ELF-2 OS-Mus musculus GN-Elf2 PE-1 SV-1	3	9.3	2.8	4.8	3.3	1.94	0.52	30.7	13.6	22.9	2.2	1.34	0.75
QB8845_AKA10_MOUSE	Akap10	A kinase anchor protein 10, mitochondrial OS-Mus musculus GN-Akap10 PE-1 SV-3	3	7.2	2.2	3.7	3.3	1.94	0.52	20.6	7.3	12.3	2.8	1.68	0.60
Q90447_TFC3_MOUSE	Tfc3	Autophagy-related protein 3 OS-Mus musculus GN-Tfc3 PE-1 SV-1	3	11.9	3.4	5.9	3.3	1.94	0.52	11.9	3.4	5.9	3.3	1.94	0.52
Q52K18_SRRM1_MOUSE	Srrm1	Serine/arginine repetitive matrix protein 1 OS-Mus musculus GN-Srrm1 PE-1 SV-2	24	35.2	10.6	18.2	3.3	1.93	0.52	1242.5	364.1	612.3	3.4	2.03	0.49
QB8453_C5N6_MOUSE	Copp6	COP9 signalosome complex subunit 6 OS-Mus musculus GN-Copp6 PE-1 SV-1	1	22.1	6.7	11.4	3.3	1.93	0.52	22.1	6.7	11.2	3.3	1.97	0.51
Q319F5_C3HIF5_MOUSE	C3hif5	A3 subunit of vacuolar-ATPase complex OS-Mus musculus GN-C3hif5 PE-1 SV-1	1	9.0	2.7	4.6	3.3	1.93	0.52	9.0	2.7	4.6	3.3	1.97	0.51
QB0203_EBF1_MOUSE	Ebf1	Eukaryotic translation initiation factor 3 subunit 1 OS-Mus musculus GN-Ebf1 PE-1 SV-1	36	37.5	11.3	19.4	3.3	1.93	0.52	1476.4	463.4	779.2	3.2	1.89	0.53
QB9K53_NFKB2_MOUSE	NfkB2	Nuclear factor kappa-B p100 subunit OS-Mus musculus GN-NfkB2 PE-1 SV-1	23	15.1	4.6	7.8	3.3	1.93	0.52	492.5	162.8	273.7	3.0	1.80	0.56
Q9PFC2_HTB6_MOUSE	Htb6	Integrator complex subunit 6 OS-Mus musculus GN-Htb6 PE-2 SV-1	19	15.0	4.6	7.8	3.3	1.93	0.52	294.8	104.2	175.2	2.8	1.68	0.59
P11459_H59B8_MOUSE	H59b8	Heat shock protein H59 beta OS-Mus musculus GN-H59b8 PE-1 SV-1	40	65.4	19.8	34.0	3.3	1.92	0.52	2473.4	1418.8	2424.3	2.4	1.43	0.70
Q919C8_ZB2A_MOUSE	Zb2a	Translational initiation factor eIF-2B subunit alpha OS-Mus musculus GN-Zb2a PE-2 SV-2	8	15.9	4.8	8.3	3.3	1.92	0.52	126.5	54.0	90.8	2.3	1.39	0.72
P59805_R171_MOUSE	Mrp11	28S ribosomal protein S21, mitochondrial OS-Mus musculus GN-Mrp11 PE-2 SV-1	1	9.0	2.7	4.7	3.3	1.92	0.52	9.0	2.7	4.6	3.3	1.96	0.51
P11279_GH40_MOUSE	Gh40	Guanine nucleotide-binding protein G12b subunit alpha OS-Mus musculus GN-Gh40 PE-1 SV-4	44	11.0	3.4	5.8	3.3	1.92	0.52	444.2	127.7	205.5	2.8	1.68	0.49
Q9P470_F1C2A_MOUSE	F1c2a	Autophagy-related protein 2 homolog OS-Mus musculus GN-F1c2a PE-1 SV-1	4	15.5	4.7	8.1	3.3	1.92	0.52	15.5	4.7	8.1	3.3	1.95	0.51
Q925M1_MED1_MOUSE	Med1	Mediator of RNA polymerase II transcription subunit 1 OS-Mus musculus GN-Med1 PE-1 SV-2	5	8.9	2.7	4.6	3.3	1.92	0.52	44.6	18.6	31.2	2.4	1.42	0.70
QB0KVA_CNOT3_MOUSE	Cnot3	BCRA-NOT transcription complex subunit 3 OS-Mus musculus GN-Cnot3 PE-1 SV-3	13	11.2	3.4	5.8	3.3	1.92	0.52	304.6	144.3	242.6	2.1	1.26	0.80
Q31522_PFM6_MOUSE	Pfm6	15S ribosome non-ATPase regulatory subunit 4 OS-Mus musculus GN-Pfm6 PE-1 SV-1	12	15.0	4.3	7.4	3.3	1.92	0.52	15.0	4.3	7.4	3.3	1.98	0.51
P72544_MAP4_MOUSE	Map4	Microtubule-associated protein 4 OS-Mus musculus GN-Map4 PE-1 SV-3	16	15.8	4.8	8.2	3.3	1.92	0.52	277.4	89.8	150.9	3.1	1.84	0.54
Q919D6_VILL_MOUSE	Vill	Villin-like protein OS-Mus musculus GN-Vill PE-1 SV-1	1	10.4	3.2	5.4	3.3	1.91	0.52	10.4	3.2	5.3	3.3	1.95	0.51
Q90XV9_CHOR_MOUSE	Chor	Chromodomain-helicase-DNA-binding protein 8 OS-Mus musculus GN-Chor PE-1 SV-1	4	7.1	2.2	3.7	3.3	1.91	0.52	30.6	9.2	15.4	3.3	1.98	0.50
Q90QK4_TAF8_MOUSE	Taf8	Transcription initiation factor TIIID subunit 8 OS-Mus musculus GN-Taf8 PE-2 SV-1	1	11.9	3.4	5.9	3.3	1.91	0.52	11.9	3.4	5.9	3.3	1.95	0.51
Q90713_CUL7_MOUSE	Cul7	Cullin-7 nuclear speckle-binding protein OS-Mus musculus GN-Cul7 PE-1 SV-2	12	16.4	5.0	8.6	3.3	1.91	0.52	195.3	110.8	186.2	1.8	1.05	0.95
Q90XN9_NBRP1_MOUSE	Nbrp1	Nuclear speckle-binding regulatory protein 1 OS-Mus musculus GN-Nbrp1 PE-1 SV-1	1	4.9	1.5	2.6	3.3	1.90	0.53	4.9	1.5	2.5	3.3	1.94	0.52
Q51465_SMRD2_MOUSE	Srmd2	SWI/SNF-related matrix-associated activator-dependent regulator of chromatin subfamily D member 2 OS-Mus musculus GN-Srmd2 PE-1 SV-1	26	15.8	4.6	8.0	3.3	1.90	0.53	154.5	46.6	79.7	2.7	1.67	0.64
Q90C04_GEM2_MOUSE	Gem2	Genome-associated protein 2 OS-Mus musculus GN-Gem2 PE-2 SV-1	26	17.4	5.3	9.0	3.3	1.90	0.53	17.4	5.3	9.0	3.3	1.94	0.52
Q90J87_DHX58_MOUSE	Dhx58	Probable ATP-dependent RNA helicase DHX58 OS-Mus musculus GN-Dhx58 PE-1 SV-2	36	21.8	6.7	11.2	3.3	1.90	0.53	904.0	262.0	440.6	3.4	2.05	0.49
Q90G62_DHX29_MOUSE	Dhx29	ATP-dependent RNA helicase DHX29 OS-Mus musculus GN-Dhx29 PE-2 SV-1	4	12.7	3.9	6.7	3.3	1.90	0.53	49.0	17.8	30.0	2.8	1.64	0.61
Q90XW5_COP7_MOUSE	Cop7	Protein FAM63 OS-Mus musculus GN-Cop7 PE-1 SV-1	5	18.5	5.2	8.9	3.3	1.90	0.53	18.5	5.2	8.9	3.3	1.97	0.51
Q90C20_C5N7A_MOUSE	C5n7a	COP9 signalosome complex subunit 7 OS-Mus musculus GN-C5n7a PE-1 SV-2	3	18.5	5.7	9.7	3.3	1.90	0.53	66.4	18.2	30.5	3.7	2.18	0.46
QB8V27_CSPF7_MOUSE	Cspf7	Cleavage and polyadenylation specificity factor subunit 7 OS-Mus musculus GN-Cspf7 PE-1 SV-2	12	22.9	7.1	12.1	3.2	1.89	0.53	351.6	103.0	173.1	3.4	2.03	0.49
Q70126_DYF16_MOUSE	Methg1	Protein Methg1 OS-Mus musculus GN-Methg1 PE-2 SV-1	1	3.4	1.1	1.8	3.2	1.89	0.53	3.4	1.1	1.8	3.2	1.99	0.52
Q92546_PARV6_MOUSE	Parv6	Parvancin OS-Mus musculus GN-Parv6 PE-1 SV-1	1	9.0	2.7	4.6	3.2	1.89	0.53	9.0	2.7	4.6	3.2	1.93	0.52
P62156_P5MD7_MOUSE	P5md7	26S proteasome non-ATPase regulatory subunit 7 OS-Mus musculus GN-P5md7 PE-1 SV-2	2	20.1	6.2	10.6	3.2	1.89	0.53	40.2	12.4	20.9	3.2	1.93	0.52
QB8CF9_PRR31_MOUSE	Prr31	UAF16 small nuclear ribonucleoprotein Prr31 OS-Mus musculus GN-Prr31 PE-1 SV-2	29	36.5	11.3	19.3	3.2	1.89	0.53	1244.2	385.6	648.4	3.2	1.92	0.52
QB0019_CFL1_MOUSE	Cfl1	Transcription initiation factor TIIID subunit 1 OS-Mus musculus GN-Cfl1 PE-1 SV-2	1	15.4	4.5	7.8	3.2	1.89	0.53	15.4	4.5	7.8	3.2	1.89	0.53
QB9VNS_FLNC_MOUSE	Flnc	Filamin-C OS-Mus musculus GN-Flnc PE-1 SV-1	78	71.4	22.1	37.8	3.2	1.89	0.53	7679.4	2517.7	4233.5	3.1	1.81	0.55
Q9N1V3_SRL10_MOUSE	U2surp	U2 snRNP-associated SURF motif-containing protein OS-Mus musculus GN-U2surp PE-2 SV-3	41	37.1	11.5	19.7	3.2	1.88	0.53	1830.8	517.2	1273.0	1.7	1.03	0.97
P13132_KAP3_MOUSE	Kap3b	eAMF-associated protein kinase type II beta regulatory subunit OS-Mus musculus GN-Kap3b PE-1 SV-1													

P35979_RL12_MOUSE	Rpl12	60S ribosomal protein L12 OS-Mus musculus GN-Rpl12 PE1 SV-2	13	82.3	26.8	45.9	3.1	1.79	0.56	6069.0	1779.9	2992.9	3.4	2.03	0.49
P18852_XS6A1_MOUSE	Rps6a1	Ribosomal protein S6 kinase alpha-1 OS-Mus musculus GN-Rps6a1 PE1 SV-1	2	8.3	2.7	4.7	3.1	1.79	0.56	16.7	5.4	8.1	3.1	1.83	0.55
QB8AA_SRR6_MOUSE	Srr6	Signal recognition particle SRP6 OS-Mus musculus GN-SRP6 PE2 SV-2	1	2.1	6.1	10.4	4.1	1.79	0.56	849.5	256.3	143.7	3.1	1.87	0.51
P58871_TB182_MOUSE	Trks1bp1	182 kDa tyrosine-1-binding protein OS-Mus musculus GN-Trks1bp1 PE1 SV-2	70	20.1	6.5	11.2	3.1	1.79	0.56	1804.7	615.5	1035.9	2.9	1.74	0.57
PE6762_MC4L1_MOUSE	Morf141	Mortality factor 4-like protein 1 OS-Mus musculus GN-Morf141 PE1 SV-2	5	8.8	2.9	4.9	3.1	1.79	0.56	40.9	14.9	25.1	2.7	1.63	0.61
QB8925_SBNL1_MOUSE	Sbnl1	Protein strobilomyxine OS-Mus musculus GN-Sbnl1 PE1 SV-2	1	5.8	3.2	5.5	3.1	1.79	0.56	12.9	4.6	140.0	2.7	1.87	0.50
QB0811_ERCC2_MOUSE	Erc2	TFIIH basal transcription factor complex helicase XPB subunit OS-Mus musculus GN-Erc2 PE1 SV-2	1	5.3	1.7	3.0	3.1	1.78	0.56	5.3	1.7	2.9	3.1	1.82	0.55
QJ3U11_ZSWM8_MOUSE	Zwim8	Zinc finger RNA domain-containing protein 8 OS-Mus musculus GN-Zwim8 PE1 SV-1	2	10.0	3.3	5.6	3.1	1.78	0.56	20.0	6.6	11.0	3.1	1.82	0.55
Q13001_RAC1_MOUSE	Rac1	Ras-related C3 botulinum toxin substrate 1 OS-Mus musculus GN-Rac1 PE1 SV-1	1	12.2	4.0	6.8	3.1	1.78	0.56	12.2	4.0	6.7	3.1	1.82	0.55
PE3262_R3D3_MOUSE	Rac3	RAS-related protein 3 OS-Mus musculus GN-Rac3 PE1 SV-1	1	20.9	7.8	14.4	3.1	1.78	0.56	28.9	9.8	18.2	3.1	1.81	0.55
QB8ZB_F142_MOUSE	Fam14a2	Protein FAM14A2 OS-Mus musculus GN-Fam14a2 PE2 SV-2	2	9.2	3.0	5.1	3.1	1.78	0.56	18.3	6.0	10.1	3.1	1.81	0.55
QB8ZB_SREK1_MOUSE	Srek1	Splicing regulatory glutamine/tyrosine-rich protein 1 OS-Mus musculus GN-Srek1 PE2 SV-2	6	22.2	7.3	12.5	3.0	1.78	0.56	113.0	37.3	62.8	3.0	1.80	0.56
Q8V9P7_PSF5_MOUSE	Cpsf6	Cleavage and polyadenylation specificity factor subunit 6 OS-Mus musculus GN-Cpsf6 PE2 SV-2	7	21.4	7.4	12.0	3.0	1.78	0.56	105.6	35.0	100.8	3.0	1.79	0.56
QB0929_TAF5_MOUSE	Taf5	Transcription initiation factor TFIID subunit 5 OS-Mus musculus GN-Taf5 PE2 SV-2	6	9.7	3.2	5.5	3.0	1.78	0.56	73.4	18.6	31.2	4.0	2.35	0.43
QB0WU1_LBN2_MOUSE	Lbn2	Ubiquitin-2 OS-Mus musculus GN-Lbn2 PE1 SV-2	3	9.7	3.2	5.5	3.0	1.78	0.56	33.8	10.3	17.3	3.3	1.96	0.51
Q53256_SFR5_MOUSE	Sfr5	Serin/glycine-rich splicing factor 5 OS-Mus musculus GN-Sfr5 PE1 SV-2	6	159.3	52.4	89.9	3.0	1.77	0.56	1012.9	361.7	608.8	3.2	1.67	0.60
Q52926_DCF2_MOUSE	Dcf2	m7GpppRNA hydrolyase OS-Mus musculus GN-Dcf2 PE1 SV-2	2	23.5	7.7	13.3	3.0	1.77	0.57	82.3	25.2	14.0	3.1	1.81	0.55
QB1R17_NBL1_MOUSE	Fam129b	Niban-like protein 1 OS-Mus musculus GN-Fam129b PE2 SV-2	8	14.1	4.6	8.0	3.0	1.77	0.57	104.3	44.6	75.0	2.3	1.39	0.72
QB2H0_RNC_MOUSE	Drosha	Ribonuclease 3 OS-Mus musculus GN-Drosha PE1 SV-1	2	10.4	3.4	5.9	3.0	1.77	0.57	20.8	6.9	11.5	3.0	1.80	0.55
Q525P7_LC7L3_MOUSE	Luc7l3	LUC7-like protein 3 OS-Mus musculus GN-Luc7l3 PE1 SV-1	21	32.7	10.8	18.5	3.0	1.77	0.57	754.4	318.6	535.7	3.4	1.41	0.71
QB1R01_WT02_MOUSE	Mta2	Mitotic-associated protein MTA2 OS-Mus musculus GN-Mta2 PE1 SV-1	30	27.8	9.2	15.7	3.0	1.77	0.57	618.0	187.6	315.4	3.3	1.96	0.51
QB108_BIN3_MOUSE	Bin3	Bridging integrator 3 OS-Mus musculus GN-Bin3 PE2 SV-1	2	10.5	3.5	5.9	3.0	1.77	0.57	21.0	6.9	11.6	3.0	1.80	0.55
Q56925_D0X46_MOUSE	Ddx46	Probable ATP-dependent RNA helicase DDX46 OS-Mus musculus GN-Ddx46 PE1 SV-2	46	34.5	11.4	19.5	3.0	1.77	0.57	2185.7	649.5	1092.2	3.4	2.00	0.50
Q62276_MDD2_MOUSE	Mdd2	Mediator of RNA polymerase II transcription subunit 22 OS-Mus musculus GN-Mdd2 PE2 SV-2	2	23.5	7.7	13.3	3.0	1.77	0.57	155.5	26.0	3.0	1.80	0.56	
QB042_E90A2_MOUSE	Ugg2	Protein Ugg2 OS-Mus musculus GN-Ugg2 PE4 SV-1	2	6.1	2.0	3.5	3.0	1.77	0.57	12.2	4.0	6.8	3.0	1.80	0.56
P46633_R8118_MOUSE	Rab11b	Ras-related protein Rab-11B OS-Mus musculus GN-Rab11b PE1 SV-3	2	41.9	13.8	23.7	3.0	1.77	0.57	83.8	27.7	46.5	3.0	1.80	0.56
QC8CA_VP53_MOUSE	Vp53	Vacuolar protein sorting-associated protein 53 homolog OS-Mus musculus GN-Vp53 PE2 SV-2	6	11.5	3.8	6.5	3.0	1.77	0.57	74.7	21.7	36.6	3.4	2.04	0.49
P15762_C0R1_MOUSE	C0r1	COR1 and OS-Mus musculus GN-C0r1 PE1 SV-2	1	15.9	5.4	9.8	3.0	1.76	0.57	5.2	1.7	5.2	3.0	1.80	0.56
PE3940_ELOC_MOUSE	Tceb1	Transcription elongation factor B polypeptide 1 OS-Mus musculus GN-Tceb1 PE1 SV-1	1	14.5	4.8	8.2	3.0	1.76	0.57	14.5	4.8	8.1	3.0	1.80	0.56
QB0CV_PFM1_MOUSE	Pfm1	Polyamine-nucleotide factor 1 OS-Mus musculus GN-Pfm1 PE1 SV-1	3	14.8	4.9	8.4	3.0	1.76	0.57	14.4	4.8	30.2	2.6	1.57	0.64
QB02G3_PSF4_MOUSE	Cpsf4	Cleavage and polyadenylation specificity factor subunit 4 OS-Mus musculus GN-Cpsf4 PE2 SV-2	6	12.4	4.1	7.1	3.0	1.76	0.57	103.1	36.6	118.2	3.6	1.67	0.63
QB043_MEPCE_MOUSE	Mepce	72kD cytoplasmic capping enzyme OS-Mus musculus GN-Mepce PE1 SV-2	6	8.2	2.7	4.6	3.0	1.76	0.57	8.2	2.7	4.5	3.0	1.80	0.56
QB4V9_NELF6_MOUSE	Nelf6	Negative elongation factor 6 OS-Mus musculus GN-Nelf6 PE1 SV-2	8	17.9	5.9	10.2	3.0	1.76	0.57	176.6	51.6	86.8	3.4	2.03	0.49
FE7N2F_FE7NF_MOUSE	Lmo7	Protein Lmo7 [Fragment] OS-Mus musculus GN-Lmo7 PE1 SV-1	2	17.0	5.6	9.6	3.0	1.76	0.57	34.0	11.2	18.9	3.0	1.80	0.56
PE0222_EF3E_MOUSE	Ef3e	Eukaryotic translation initiation factor 3 OS-Mus musculus GN-Ef3e PE1 SV-1	6	13.0	4.6	7.6	3.0	1.76	0.57	561.5	95.7	161.5	2.7	1.61	0.99
QBK30_D0X52_MOUSE	Ddx52	Probable ATP-dependent RNA helicase DDX52 OS-Mus musculus GN-Ddx52 PE2 SV-2	22	19.9	6.6	11.3	3.0	1.76	0.57	518.0	189.1	318.0	2.7	1.63	0.61
QB217_FAMN1_MOUSE	Fmn1	Formin-like protein 3 OS-Mus musculus GN-Fmn1 PE1 SV-2	9	12.1	4.0	6.9	3.0	1.76	0.57	112.3	35.2	59.1	3.2	1.90	0.53
QJ3U01_QJ3U01_MOUSE	Srd11	Protein Srd11 OS-Mus musculus GN-Srd11 PE1 SV-1	6	39.4	13.1	22.4	3.0	1.76	0.57	357.3	300.4	505.1	1.2	0.71	1.41
QB1792_LAS1_MOUSE	Las1	LAM and SRS domain protein 1 OS-Mus musculus GN-Las1 PE1 SV-1	1	46.8	16.3	28.6	3.0	1.76	0.57	128.5	43.7	107.5	3.0	1.77	0.56
Q61666_HIRA_MOUSE	Hira	Protein HIRA OS-Mus musculus GN-Hira PE1 SV-3	3	10.5	3.5	6.0	3.0	1.76	0.57	34.9	8.2	13.8	4.3	2.54	0.39
PR7360_ETV6_MOUSE	Etv6	Transcription factor ETV6 OS-Mus musculus GN-Etv6 PE1 SV-1	5	21.5	7.1	12.3	3.0	1.76	0.57	111.1	37.3	62.8	3.0	1.77	0.57
QC2P7_LBP2_MOUSE	Lbp2	DNA replication licensing factor 2 OS-Mus musculus GN-Lbp2 PE1 SV-1	2	12.3	4.1	7.0	3.0	1.76	0.57	41.1	13.7	26.9	3.0	1.79	0.56
Q52126_DYST_MOUSE	Dyst	Dystonin OS-Mus musculus GN-Dyst PE1 SV-2	4	9.0	3.0	5.1	3.0	1.76	0.57	33.1	19.0	31.9	1.7	1.04	0.96
Tubgcp2	Tubgcp2	WASH-tubulin complex component 2 OS-Mus musculus GN-Tubgcp2 PE2 SV-2	11	13.6	4.5	7.7	3.0	1.76	0.57	158.9	47.2	79.3	3.4	2.00	0.50
Q52C77_STRUM_MOUSE	Klaad196	YAMA complex subunit strumpelin OS-Mus musculus GN-Klaad196 PE2 SV-2	13	12.0	4.0	6.9	3.0	1.76	0.57	202.6	67.0	112.6	3.0	1.80	0.56
PE3553_C02B_MOUSE	C02b	C02B and OS-Mus musculus GN-C02b PE1 SV-1	4	52.3	17.4	29.8	3.0	1.76	0.57	44.7	16.6	166.1	2.5	1.48	0.67
P13120_AMP1_MOUSE	Amp1	Aminocoyl tRNA synthetase complex-interacting multifunctional protein 1 OS-Mus musculus GN-Amp1	1	14.4	4.8	8.2	3.0	1.76	0.57	14.4	4.8	8.1	3.0	1.79	0.56
EQ908E_EQ908H_MOUSE	Plekha5	Protein Plekha5 OS-Mus musculus GN-Plekha5 PE1 SV-1	4	9.4	3.1	5.3	3.0	1.76	0.57	40.4	16.6	28.0	2.4	1.44	0.69
Q64512_TFN13_MOUSE	Tfn13	Tyrosine protein phosphatase non-receptor type 13 OS-Mus musculus GN-Tfn13 PE1 SV-2	4	7.5	2.4	4.3	3.0	1.75	0.57	29.9	9.9	16.8	3.0	1.60	0.56
Q5555A_TRPCC4_MOUSE	Trpcc4	Transcription regulator 4 OS-Mus musculus GN-Trpcc4 PE1 SV-1	1	5.4	1.8	3.1	3.0	1.75	0.57	5.4	1.8	3.0	3.0	1.79	0.56
Q6PDN1_MYK_MOUSE	Myk	Myosin light chain, smooth muscle OS-Mus musculus GN-Myk PE1 SV-3	2	10.4	3.5	5.9	3.0	1.75	0.57	20.9	6.9	11.6	3.0	1.79	0.56
Q9C0A_RING_MOUSE	Ring	E4 ubiquitin-protein ligase RING2 OS-Mus musculus GN-Ring PE1 SV-1	4	22.6	7.5	12.9	3.0	1.75	0.57	88.0	29.3	49.2	3.0	1.79	0.56
PE2841_R24_MOUSE	R24	40S ribosomal protein R24 OS-Mus musculus GN-R24 PE1 SV-1	1	73.7	24.4	41.4	3.0	1.75	0.57	187.2	61.6	187.2	3.4	1.87	0.48
Q9C9Y_KAD1_MOUSE	Ak1	Adenylyl kinase isoenzyme 1 OS-Mus musculus GN-Ak1 PE1 SV-1	1	11.1	3.7	6.3	3.0	1.75	0.57	11.1	3.7	6.2	3.0	1.79	0.56
FE1012_FE1AL_MOUSE	Etf1a1	Elongation factor 1-alpha 1 OS-Mus musculus GN-Etf1a1 PE1 SV-1	46	78.6	26.2	44.2	3.0	1.75	0.57	8351.1	3324.0	5589.2	2.5	1.49	0.67
OT4046_VAMP8_MOUSE	Vamp8	Vesicle-associated membrane protein 8 OS-Mus musculus GN-Vamp8 PE1 SV-1	3	45.7	15.2	26.1	3.0	1.75	0.57	129.9	46.4	78.0	2.8	1.67	0.63
QB9752_GNA2_MOUSE	Gna2	GTP-binding protein GNA2 OS-Mus musculus GN-Gna2 PE1 SV-1	4	44.1	14.7	25.6	3.0	1.75	0.57	106.6	36.7	61.9	2.9	1.69	0.61
Q61292_LAMB2_MOUSE	Lamb2	Laminin subunit beta-2 OS-Mus musculus GN-Lamb2 PE2 SV-2	4	10.6	3.5	6.1	3.0	1.75	0.57	40.8	14.9	25.0	2.7	1.63	0.61
Q9DAW9_CNN3_MOUSE	Cnn3	Calponin 3 OS-Mus musculus GN-Cnn3 PE1 SV-1	14	16.7	5.6	9.6	3.0	1.75	0.57	430.2	134.9	226.8	3.2	1.90	0.53
PE3307_P7F6_MOUSE	Ppp1cc	Serin/threonine-protein phosphatase P7F6 gamma catalytic subunit OS-Mus musculus GN-Ppp1cc	1	38.7	11.6	20.4	3.0	1.74	0.57	14.9	5.2	25.1	2.6	1.51	0.65
QB8G10_NELF6_MOUSE	Nelf6	Negative elongation factor 6 OS-Mus musculus GN-Nelf6 PE2 SV-2	9	9.8	3.3	5.6	3.0	1.74	0.57	119.7	43.5	73.2	2.8	1.64	0.61
Q9J914_PSM06_MOUSE	Psm06	26S proteasome non-ATPase regulatory subunit 6 OS-Mus musculus GN-Psm06 PE1 SV-1	3	14.7	4.9	8.4	3.0	1.74	0.57	19.6	17.3	29.0	5.3	3.16	0.32
Q9P977_TIR3_MOUSE	Trim3	E4 ubiquitin-protein ligase TRIM3 OS-Mus musculus GN-Trim3 PE1 SV-2	4	9.1	3.1	5.2	3.0	1.74	0.57	43.7	16.6	27.9	2.6	1.57	0.64
QB0X11_TK2_MOUSE	Tk2	Eukaryotic translation initiation factor 2 OS-Mus musculus GN-Tk2 PE1 SV-1	1	18.8	6.3	10.9	3.0	1.74	0.57	12.2	4.0	12.2	3.0	1.79	0.56
PE5724_PCBP4_MOUSE	Pcbp4	Poly(C)binding protein 4 OS-Mus musculus GN-Pcbp4 PE1 SV-2	2	9.5	3.2	5.5	3.0	1.74	0.57	19.3	6.4	10.7	3.0	1.78	0.56
Q60CE_ASF1A_MOUSE	Asf1a	Histone chaperone ASF1A OS-Mus musculus GN-Asf1a PE1 SV-1	1	11.2	3.8	6.4	3.0	1.74	0.57	11.2	3.8	6.3	3.0	1.78	0.56
QB97F_MED16_MOUSE	Med16	Mediator of RNA polymerase II transcription subunit 16 OS-Mus musculus GN-Med16 PE2 SV-2	6	10.9	3.7	6.3	3.0	1.74	0.57	64.2	26.0	43.7	2.5	1.47	0.68
Q54931_WAPR1_MOUSE	Wapr1														

Q3UMG5_LRCH2_MOUSE	Lrch2	Leucine-rich repeat and calponin homology domain-containing protein 2 OS=Mus musculus GN	3	13.0	4.9	8.4	2.7	1.56	0.64	31.8	12.8	21.5	2.5	1.48	0.63
OB0305_OB11_MOUSE	Obp11	Oxytetracycline-resistance-related protein 1 OS=Mus musculus GN=Obp11 PE=1 SV=2	2	17.6	6.6	11.3	2.7	1.56	0.64	35.1	13.2	22.2	2.7	1.59	0.63
Q9H9S_P68B_MOUSE	Prp8	Transcription factor P68 OS=Mus musculus GN=P68 PE=1 SV=1	4	17.0	6.2	10.8	2.7	1.56	0.64	17.0	25.0	29.0	2.7	1.62	0.63
Q96R7_E0Q67_MOUSE	Utrn	Protein Utrn OS=Mus musculus GN=Utrn PE=1 SV=1	54	22.1	8.3	14.2	2.7	1.56	0.64	1150.3	462.3	777.3	2.5	1.48	0.68
P30415_NTR_MOUSE	Ntr	Nucleo-1 recognition protein OS=Mus musculus GN=Ntr1 PE=1 SV=4	2	14.0	5.2	9.0	2.7	1.55	0.64	27.9	10.5	17.6	2.7	1.58	0.63
Q01231_TFRP2_MOUSE	Tfrp2	Polymorphic tract containing protein OS=Mus musculus GN=Tfrp2 PE=1 SV=2	2	15.5	9.6	6.9	2.7	1.55	0.64	120.5	67.3	101.6	2.7	1.63	0.61
Q90270_ARP10_MOUSE	Actr10	Actin-related protein 10 OS=Mus musculus GN=Actr10 PE=1 SV=2	2	14.2	3.2	5.6	2.7	1.55	0.64	17.3	6.5	10.9	2.7	1.58	0.63
Q9P580_RRP12_MOUSE	Rrp12	RRP12-like protein OS=Mus musculus GN=RRP12 PE=1 SV=1	7	10.8	4.1	6.9	2.7	1.55	0.65	79.8	29.3	49.2	2.7	1.62	0.62
Q61545_EVS_MOUSE	Evs1	RNA-binding protein EVS OS=Mus musculus GN=Evs1 PE=1 SV=2	3	25.1	9.4	16.2	2.7	1.55	0.65	78.0	27.4	46.0	2.8	1.69	0.59
Q8B914_TRM2A_MOUSE	Trm2a	2'-O-methyltransferase TRM2A OS=Mus musculus GN=Trm2a PE=2 SV=1	11	11.2	4.2	7.2	2.7	1.55	0.65	4.2	7.2	11.2	2.7	1.63	0.61
Q24448_IF4G2_MOUSE	If4g2	Eukaryotic translation initiation factor 4 gamma 2 OS=Mus musculus GN=If4g2 PE=1 SV=2	10	11.5	4.3	7.4	2.7	1.55	0.65	239.7	144.7	243.3	1.7	1.99	1.02
Q04690_SMR3_MOUSE	Smr3	SWI/SNF-related matrix-associated actin-activated regulator of chromatin subfamily 3 contain	1	8.5	3.2	5.5	2.7	1.55	0.65	8.5	3.2	5.4	2.7	1.58	0.63
Q51Y62_IP29_MOUSE	Ip29	Interleukin-1 receptor type 2 OS=Mus musculus GN=Ip29 PE=1 SV=3	3	8.8	3.8	4.8	2.7	1.55	0.65	6.0	4.8	15.2	2.6	1.52	0.64
Q9R170_SNP47_MOUSE	Snp47	Synaptonemal-associated protein SNP47 OS=Mus musculus GN=Snp47 PE=1 SV=1	3	8.2	3.1	5.3	2.7	1.55	0.65	21.4	8.4	14.2	2.5	1.51	0.66
Q9R181_TRIM3_MOUSE	Trim3	Tripartite motif-containing protein 3 OS=Mus musculus GN=Trim3 PE=1 SV=1	5	15.7	5.9	10.7	2.7	1.55	0.65	77.0	29.1	49.0	2.6	1.57	0.64
PE1069_GABP2_MOUSE	Gabp2	GAB-binding protein subunit beta 2 OS=Mus musculus GN=Gabp2 PE=2 SV=2	3	9.1	3.4	5.9	2.7	1.55	0.65	24.5	15.7	26.3	1.6	1.93	1.07
Q8V84_NOC3L_MOUSE	Noc3l	Nucleolar complex protein 3 homolog OS=Mus musculus GN=Noc3l PE=2 SV=2	2	29.5	11.1	19.1	2.6	1.55	0.65	1481.5	510.9	859.9	1.9	1.72	0.58
P70271_PDL4_MOUSE	Pdlm4	PDZ and LIM domain protein 4 OS=Mus musculus GN=Pdlm4 PE=1 SV=3	11	31.8	12.0	20.6	2.7	1.55	0.65	379.0	150.4	253.0	2.5	1.50	0.67
Q61407_GTA_MOUSE	Gta4	Golgi to ER traffic protein 4 homolog OS=Mus musculus GN=Gta4 PE=2 SV=2	2	16.3	6.1	10.5	2.7	1.55	0.65	32.6	12.3	20.7	2.7	1.58	0.63
Q80296_GRO295_MOUSE	H16	H16 histone family, member X OS=Mus musculus GN=H16 PE=1 SV=1	12	74.6	28.1	48.2	2.7	1.55	0.65	1481.5	510.9	859.9	1.9	1.72	0.58
Q29193_CDR3_MOUSE	Cdr3	Cytosine-dependent kinase 3 OS=Mus musculus GN=CDK3 PE=1 SV=1	1	12.8	4.8	8.2	2.7	1.55	0.65	61.9	22.5	37.8	2.8	1.64	0.61
Q9J94_EIF3M_MOUSE	Eif3m	Eukaryotic translation initiation factor 3 subunit M OS=Mus musculus GN=Eif3m PE=2 SV=1	6	21.2	8.0	13.7	2.7	1.55	0.65	195.8	51.5	86.5	3.8	2.26	0.44
Q90806_MED1_MOUSE	Med11	Mediator of RNA polymerase II transcription subunit 11 OS=Mus musculus GN=Med11 PE=1 SV=3	2	23.9	9.0	15.4	2.7	1.55	0.65	47.7	18.0	30.3	2.7	1.58	0.63
PE1620_S6141_MOUSE	Sec14	Sec14 nuclear protein subunit alpha isoform 1 OS=Mus musculus GN=Sec14 PE=1 SV=2	2	29.5	11.1	19.1	2.6	1.55	0.65	21.3	59.0	22.3	2.6	1.58	0.63
Q5C800_VPS25_MOUSE	Vps25	Vacuolar protein-sorting-associated protein 25 OS=Mus musculus GN=Vps25 PE=2 SV=1	1	8.7	3.3	5.6	2.6	1.55	0.65	8.7	3.3	5.5	2.6	1.58	0.63
Q8R5C3_ACT1_MOUSE	Actr1b	Beta-actinin OS=Mus musculus GN=Actr1b PE=1 SV=1	13	24.9	9.4	16.1	2.6	1.54	0.65	568.2	275.2	462.7	2.1	1.23	0.81
Q61087_LAMB3_MOUSE	Lamb3	Laminin subunit beta-3 OS=Mus musculus GN=Lamb3 PE=2 SV=2	1	4.5	1.7	2.9	2.6	1.54	0.65	4.5	1.7	2.8	2.6	1.57	0.64
Q9P246_RPL35_MOUSE	Rpl35	Putative RNA-binding protein 15B OS=Mus musculus GN=Rpl35 PE=1 SV=2	9	19.5	5.8	10.0	2.6	1.54	0.65	11.8	129.4	11.8	2.6	1.57	0.64
Q8BK12_TRMR6_MOUSE	Trmr6	Trinucleotide repeat-containing gene 6B protein OS=Mus musculus GN=Trmr6 PE=1 SV=1	3	8.9	3.4	5.8	2.6	1.54	0.65	27.2	11.2	18.8	2.4	1.45	0.69
Q904H1_EXOC2_MOUSE	Exoc2	Exocyst complex component 2 OS=Mus musculus GN=Exoc2 PE=1 SV=1	8	15.6	5.9	10.6	2.6	1.54	0.65	122.8	44.6	74.7	2.8	1.64	0.61
A2JAV2_VIR_MOUSE	Vmr1	Protein vir1 homolog OS=Mus musculus GN=Vmr1 PE=1 SV=1	7	10.1	3.7	6.0	2.6	1.54	0.65	10.1	3.7	50.3	2.4	1.41	0.71
P43247_MAB2_MOUSE	Mab2	DNA mismatch repair protein MAB2 OS=Mus musculus GN=Mab2 PE=1 SV=1	31	22.0	8.3	14.3	2.6	1.54	0.65	60.6	610.6	1026.8	1.1	0.66	1.22
Q53600_SCAMP3_MOUSE	Scamp3	History carrier-associated membrane protein 3 OS=Mus musculus GN=Scamp3 PE=1 SV=3	5	35.5	13.4	23.1	2.6	1.54	0.65	155.1	50.6	85.2	3.1	1.82	0.55
Wd228_HDAC2_MOUSE	Hdac2	Histone deacetylase 2 OS=Mus musculus GN=Hdac2 PE=1 SV=1	7	17.8	6.7	11.6	2.6	1.54	0.65	121.6	43.0	72.3	2.8	1.68	0.59
Q8B207_ANC2_MOUSE	Ank2	Anchorin-2 OS=Mus musculus GN=Ank2 PE=2 SV=1	1	26.9	10.2	18.5	2.6	1.54	0.65	26.9	10.2	18.5	2.6	1.57	0.64
E9PYH6_ESP9H_MOUSE	Setd1a	Protein Setd1a OS=Mus musculus GN=Setd1a PE=4 SV=1	16	11.4	4.3	7.4	2.6	1.54	0.65	238.9	72.8	122.5	3.3	1.95	0.51
PE1750_ARF4_MOUSE	Arf4	ADP-ribosylation factor 4 OS=Mus musculus GN=Arf4 PE=1 SV=2	3	16.0	6.1	10.4	2.6	1.54	0.65	44.8	20.8	35.0	2.1	1.28	0.78
Q4F66_PRC3A_MOUSE	Prp38a	Pre-rRNA-splicing factor 38A OS=Mus musculus GN=Prp38a PE=1 SV=1	6	20.4	7.8	13.3	2.6	1.54	0.65	106.7	66.5	111.8	1.6	1.95	1.05
Q4K44_C22A2_MOUSE	Rps24	28S ribosomal protein L24, isoform 2 OS=Mus musculus GN=C22A2 PE=1 SV=1	23	29.8	10.5	19.2	2.6	1.53	0.65	110.2	65.8	102.2	2.6	1.57	0.64
R97351_R3A_MOUSE	Rps3a	40S ribosomal protein S3a OS=Mus musculus GN=Rps3a PE=1 SV=1	22	85.2	32.4	55.5	2.6	1.54	0.65	3146.4	1214.7	2042.5	2.6	1.54	0.65
Q5190_H4L1_MOUSE	H4-L1	H-1 class I histone core particle antigen, K-D alpha chain OS=Mus musculus GN=H4-L1 PE=1 SV=1	1	10.4	4.0	6.8	2.6	1.54	0.65	10.4	4.0	6.7	2.6	1.57	0.64
Q07417_ACD2_MOUSE	Acd2	Actin-specific actin capping protein OS=Mus musculus GN=Acd2 PE=1 SV=1	2	8.8	3.5	6.2	2.6	1.54	0.65	8.8	3.5	6.2	2.6	1.57	0.64
Q58A65_JP4_MOUSE	Spa9	C-hu amino-terminal kinase-interacting protein 4 OS=Mus musculus GN=Spa9 PE=1 SV=2	1	5.6	2.1	3.6	2.6	1.53	0.65	5.6	2.1	3.6	2.6	1.56	0.64
PGC51_CTR8_MOUSE	Chtr8	Chromosome translocation fidelity protein 8 homolog OS=Mus musculus GN=Chtr8 PE=2 SV=1	2	18.8	7.1	12.3	2.6	1.53	0.65	37.6	14.3	24.0	2.6	1.56	0.64
PE8373_TBAC1_MOUSE	Tubac1	Tubulin alpha 1C chain OS=Mus musculus GN=TBAC1 PE=1 SV=1	6	54.3	20.7	35.4	2.6	1.53	0.65	545.5	245.5	412.7	2.2	1.32	0.76
P42206_SFPT2_MOUSE	Sfpt2	Serine phosphorylation factor 2 OS=Mus musculus GN=Sfpt2 PE=1 SV=1	2	19.4	7.4	12.7	2.6	1.53	0.65	16.4	10.5	16.4	2.6	1.57	0.64
Q62450_WAPL_MOUSE	Wapl	Wings apart-like protein homolog OS=Mus musculus GN=Wapl PE=1 SV=2	11	11.8	4.5	7.7	2.6	1.53	0.65	149.1	52.0	87.4	2.9	1.70	0.59
VP372_VAPB_MOUSE	Vapb	Vacuolar protein sorting-associated protein 72 homolog OS=Mus musculus GN=Vapb PE=2 SV=2	3	28.1	10.7	18.4	2.6	1.53	0.65	123.3	50.7	85.2	2.4	1.45	0.69
Q80B75_WAPR2_MOUSE	Wapr2	WAPR2 OS=Mus musculus GN=Wapr2 PE=1 SV=1	2	7.6	2.9	5.0	2.6	1.53	0.65	7.6	2.9	5.0	2.6	1.56	0.64
Q8C820_CHERP_MOUSE	Cherp	Calcium homeostasis endoplasmic reticulum protein OS=Mus musculus GN=Cherp PE=1 SV=1	16	29.0	11.0	18.9	2.6	1.53	0.65	769.6	231.2	388.8	3.3	1.98	0.51
Q8BC18_FERM2_MOUSE	Ferm2	Fermitin family homolog OS=Mus musculus GN=Ferm2 PE=1 SV=1	9	20.2	7.7	13.2	2.6	1.53	0.65	164.4	58.2	97.8	2.8	1.68	0.59
P14575_SRP54_MOUSE	Srp54	Signal recognition particle 54 kDa protein OS=Mus musculus GN=Srp54 PE=1 SV=2	22	26.1	9.9	17.0	2.6	1.53	0.65	80.3	397.2	667.9	2.1	1.28	0.78
Q8A273_MMG17_MOUSE	Mmg17	Membrane magnesium transporter 17 OS=Mus musculus GN=Mmg17 PE=2 SV=1	9	14.9	5.3	9.3	2.6	1.53	0.65	57.5	18.7	31.6	2.6	1.66	0.64
BRU514_B9E15A_MOUSE	Nup205	MGC217156, isoform CRA_b OS=Mus musculus GN=Nup205 PE=2 SV=1	5	8.4	3.2	5.5	2.6	1.53	0.65	45.1	18.6	31.2	2.4	1.45	0.69
P16330_CN3_MOUSE	Cnp	2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Mus musculus GN=Cnp PE=1 SV=3	1	8.0	3.1	5.2	2.6	1.53	0.65	8.0	3.1	5.1	2.6	1.56	0.64
QJ0201_RBM4_MOUSE	Rnpc3	RNA-binding protein 3 OS=Mus musculus GN=Rnpc3 PE=2 SV=2	1	7.2	2.8	4.7	2.6	1.53	0.65	7.2	2.8	4.6	2.6	1.56	0.64
PC2196_PRS8_MOUSE	Prs8	Protein tyrosine phosphatase 8 OS=Mus musculus GN=Prs8 PE=1 SV=1	1	8.0	3.0	4.9	2.6	1.53	0.65	28.7	10.7	14.7	2.6	1.57	0.64
Q62P2V_INO80_MOUSE	Ino80	DNA helicase INO80 OS=Mus musculus GN=INO80 PE=2 SV=2	1	4.9	1.9	3.2	2.6	1.53	0.65	4.9	1.9	3.1	2.6	1.56	0.64
OT7M81_UHRF2_MOUSE	Uhrf2	E1 ubiquitin-protein ligase UHRF2 OS=Mus musculus GN=UHRF2 PE=1 SV=1	3	5.3	2.0	3.5	2.6	1.53	0.65	14.9	8.4	14.2	1.8	1.05	0.95
PE0313_TCF2_MOUSE	Ctcf2	C-terminal protein 1 subunit alpha OS=Mus musculus GN=Ctcf2 PE=1 SV=3	33	19.8	7.0	12.3	2.6	1.53	0.65	334.3	129.2	212.2	2.6	1.54	0.65
Q8R019_HD43_MOUSE	Hd43	Transcriptional adaptor 3 OS=Mus musculus GN=HD43 PE=1 SV=1	6	20.3	7.3	12.6	2.6	1.53	0.65	125.2	45.2	75.9	2.8	1.65	0.61
Q51510_TRD25_MOUSE	Trd25	E1 ubiquitin-protein ligase TRIM25 OS=Mus musculus GN=Trm25 PE=1 SV=2	9	12.8	4.9	8.4	2.6	1.52	0.66	120.3	62.6	105.2	1.9	1.14	0.87
P42567_EP51_MOUSE	Eps15	Epidermal growth factor receptor substrate 15 OS=Mus musculus GN=Eps15 PE=1 SV=1	2	6.2	2.4	4.1	2.6	1.52	0.66	12.4	4.7	8.0	2.6	1.55	0.64
Q52241_HMRP_MOUSE	Hmrp	Progerin-related nuclear protein OS=Mus musculus GN=HMRP PE=1 SV=3	27	10.9	4.0	6.8	2.6	1.52	0.66	69.9	347.5	584.9	2.0	1.14	0.84
Q8C076_GQC76_MOUSE	Gqc76	Kinesin light chain 1 OS=Mus musculus GN=GQC76 PE=1 SV=1	7	13.4	5.1	8.8	2.6	1.52	0.66	112.2	34.1	57.4	2.4	1.42	0.71
D3YVV2_D3YVW2_MOUSE	Golim4	Golgi integral membrane protein 4 OS=Mus musculus GN=Golim4 PE=1 SV=1	4	16.8	6.4	11.0	2.6	1.52	0.66	39.0	23.2	39.0	2.4	1.44	0.69
Q8V808_AK81_MOUSE	Prkab1	5'-NMP-activated protein kinase subunit beta-1 OS=Mus musculus GN=Prkab1 PE=1 SV=2	4	11.3	4.3	7.4	2.6	1.52	0.66	57.3	21.6	36.3	2.7	1.58	0.63
Q52908_R53_MOUSE	R53	52S ribosomal protein OS=Mus musculus GN=R53 PE=1 SV=1	34	46.4	17.0	32.4	2.6	1.52	0.66	5987.8	2288.4	3864.6	2.6	1.55	0.65
Q7F487_FT87_MOUSE	Pamd4	26S proteasome non-ATPase regulatory subunit 4 (Fragment) OS=Mus musculus GN=Pamd4 PE=1 SV=1	2	7.1	2.7	4.7	2.6	1.52	0.66	14.2	5.4	9.2	2.6	1.55	0.64
Q51644_CTR1_MOUSE	Cent3	Centrioin-3 OS=Mus musculus GN=Cent3 PE=2 SV=1	1	8.7	3.3	5.7	2.6	1.52	0.66	8.7	3.3	5.6	2.6	1.55	0.64
Q60932_VDCL1_MOUSE	Vdcl1	Vesicle-docking and uncoating protein 1 OS=Mus musculus GN=VDCL1 PE=1 SV=3	1	7.8	2.9	4.6	2.6	1.52	0.66	14.7	4.6	13.9	2.6	1.65	0.66
Q920U4_ZO2_MOUSE	Tjp2	Tight junction protein ZO-2 OS=Mus musculus GN=Tjp2 PE=1 SV=2	55	35.3	13.5	23.2	2.6	1.52	0.66	2393.5	931.3	1566.0	2.6	1.53	0.65
Q9JH8_E0D21_MOUSE	Kder1	ER lumen protein-retaining receptor 1 OS=Mus musculus GN=Kder1 PE=2 SV=1	2	32.8	12.6	21.6	2.6	1.52	0.66	65.6	25.2	42.3	2.6	1.55	0.65
G5E8V_G5E8V_MOUSE	Arf1p1	MGC18094 OS=Mus musculus GN=Arf1p1 PE=4 SV=1	11	26.3	1										

O9R1C7_PR40A_MOUSE	Pripf40a	PremRNA-processing factor 40 homolog A OS-Mus musculus GN:Pripf40a PE1-SV1	39	35.2	14.0	24.0	2.5	1.47	0.68	2513.8	1019.7	1714.5	2.5	1.47	0.68
O9D4H2_GCLC1_MOUSE	Gccl1	GRIIP and coiled-coil domain-containing protein 1 OS-Mus musculus GN:Gccl1 PE1-SV2	3	6.8	2.7	4.7	2.5	1.47	0.68	219.9	9.2	15.4	2.4	1.42	0.70
OT1L4_TLR2_MOUSE	Tlr2	Toll-like receptor 2 OS-Mus musculus GN:Tlr2 PE1-SV3	6	10.8	7.5	10.8	2.5	1.46	0.68	486.6	150.3	44.7	2.6	1.63	0.61
O90877_O9BQ7_MOUSE	Trim12c	Protein Trim12c OS-Mus musculus GN:Trim12c PE1-SV1	10	18.9	7.5	12.9	2.5	1.46	0.68	237.6	122.1	205.3	1.9	1.16	0.86
O95639_UBPS_MOUSE	Ubp5	Ubiquitin carboxyl-terminal hydrolase 5 OS-Mus musculus GN:Ubp5 PE1-SV1	11	12.2	4.8	8.3	2.5	1.46	0.68	135.8	61.8	104.0	2.2	1.31	0.77
Q172Z1_E5T72_MOUSE	E5T72	Extended synaptotagmin-like protein 1 OS-Mus musculus GN:E5T72 PE1-SV1	14	11.6	4.6	7.9	2.5	1.46	0.68	154.2	79.4	154.8	1.6	1.58	0.67
O99371_TMED9_MOUSE	Tmed9	Transmembrane emp24 domain-containing protein 9 OS-Mus musculus GN:Tmed9 PE1-SV2	11	23.2	9.2	15.8	2.5	1.46	0.68	221.2	86.1	144.8	2.6	1.53	0.65
O8Q114_INT9_MOUSE	Int9	Integrator complex subunit 9 OS-Mus musculus GN:Int9 PE2-SV1	3	9.8	3.9	6.7	2.5	1.46	0.68	28.4	16.6	19.5	2.4	1.46	0.69
O8C181_SF1P1_MOUSE	Sf1p1	Septin 11 OS-Mus musculus GN:Sf1p1 PE1-SV1	7	13.9	5.5	9.5	2.5	1.46	0.68	117.0	43.6	73.3	2.7	1.60	0.69
OT191A_EF3D_MOUSE	Ef3d	RNA-dependent protein kinase (RDK) subunit D OS-Mus musculus GN:Ef3d PE1-SV1	7	13.9	5.5	9.5	2.5	1.46	0.68	117.0	43.6	73.3	2.7	1.60	0.69
O91W69_EFN3_MOUSE	Efn3	Epin-3 OS-Mus musculus GN:Efn3 PE2-SV1	3	4.3	1.7	2.9	2.5	1.46	0.68	11.7	6.6	11.1	1.8	1.06	0.94
O8B85A_O8B85A_MOUSE	Znmp6	Protein Znm6 OS-Mus musculus GN:Znmp6 PE2-SV1	1	4.8	1.9	3.3	2.5	1.46	0.69	4.8	1.9	3.2	2.5	1.49	0.67
O91W16_UBP3_MOUSE	Ubp3	Ubiquitin carboxyl-terminal hydrolase 3 OS-Mus musculus GN:Ubp3 PE2-SV1	12	10.7	7.2	12.5	2.5	1.46	0.69	144.8	55.4	144.4	2.5	2.49	0.67
O9C503_CACTN_MOUSE	Cactn	Catenin OS-Mus musculus GN:Cactn PE1-SV2	4	8.2	3.3	5.6	2.5	1.46	0.69	42.1	13.5	22.7	3.1	1.85	0.54
OT0743_OTUB1_MOUSE	Oub1	Ubiquitin thioesterase OTUB1 OS-Mus musculus GN:Oub1 PE1-SV2	3	6.6	2.6	4.5	2.5	1.46	0.69	27.0	14.9	25.1	1.8	1.08	0.93
O9C2U3_SKL21_MOUSE	Skl21	Superkiller viral integrase activity 2-like 2 OS-Mus musculus GN:Skl21 PE1-SV1	28	20.3	8.1	13.9	2.5	1.46	0.69	707.9	297.4	500.1	2.6	1.54	0.65
P5115D_NAB7A_MOUSE	Nab7a	Ras-related protein Rab-7a OS-Mus musculus GN:Nab7a PE1-SV1	20	15.6	6.2	10.7	2.5	1.46	0.69	691.4	249.7	429.8	2.8	1.65	0.61
O91K7C_AP4M1_MOUSE	Ap4m1	AP-4 complex subunit mu-1 OS-Mus musculus GN:Ap4m1 PE2-SV1	1	8.0	3.2	5.4	2.5	1.46	0.69	8.0	3.2	5.4	2.5	1.49	0.67
O9QZD8_LARP1_MOUSE	Larp1	Lary-retroviral protein 1 OS-Mus musculus GN:Larp1 PE1-SV3	28	15.7	6.3	10.7	2.5	1.46	0.69	570.1	231.1	388.6	2.5	1.47	0.68
O8D19E_RNKS5_MOUSE	Rnks5	Nuclear pore complex protein Nup155 OS-Mus musculus GN:Nup155 PE1-SV1	3	9.4	3.8	6.4	2.5	1.46	0.69	26.4	9.3	15.6	2.8	1.69	0.59
O8C0C0_C92B_MOUSE	C92b	Cytosine-rich protein 3 OS-Mus musculus GN:C92b PE1-SV2	24	5.6	2.7	3.8	2.5	1.46	0.69	42.2	14.5	25.5	2.5	1.49	0.67
O9P9B0_EP9B0_MOUSE	Ahnak2	Protein Ahnak2 (Fragment) OS-Mus musculus GN:Ahnak2 PE1-SV4	12	17.2	6.9	11.1	2.5	1.46	0.69	446.6	150.8	253.6	3.0	1.76	0.57
OT6K16_RA3A_MOUSE	C3baap	DNA-directed RNA polymerase subunit RP34 OS-Mus musculus GN:C3baap PE1-SV2	4	10.3	4.1	7.1	2.5	1.46	0.69	42.7	18.3	30.8	2.3	1.38	0.72
O9P99E_NU155_MOUSE	Nu155	Nuclear pore complex protein Nup155 OS-Mus musculus GN:Nu155 PE1-SV1	12	14.0	5.6	9.6	2.5	1.46	0.69	176.5	109.9	128.6	2.3	1.39	0.72
P62835_RAP1A_MOUSE	Rap1a	Ras-related protein Rap-1A OS-Mus musculus GN:Rap1a PE2-SV1	8	13.3	5.3	9.2	2.5	1.46	0.69	182.3	81.2	136.5	2.2	1.34	0.75
P70698_PRRG1_MOUSE	Ctsp1	CYP synthase 1 OS-Mus musculus GN:Ctsp1 PE1-SV2	23	22.8	9.2	15.7	2.5	1.45	0.69	623.2	287.7	483.7	2.2	1.29	0.78
O9D743_DUS3_MOUSE	Dusp3	Dual specificity tyrosine phosphatase 3 OS-Mus musculus GN:Dusp3 PE1-SV1	1	7.3	2.9	5.0	2.5	1.45	0.69	7.3	2.9	4.9	2.5	1.48	0.64
Q10000_SCR1L_MOUSE	Scrl1	Scavenger receptor class member 1 OS-Mus musculus GN:Scrl1 PE1-SV1	2	5.2	2.7	3.7	2.5	1.45	0.69	7.4	3.2	12.0	2.6	1.47	0.68
P13174_KIF4_MOUSE	Kif4	Chromosome-associated kinesin KIF4 OS-Mus musculus GN:Kif4 PE1-SV3	21	10.2	4.1	7.0	2.5	1.45	0.69	262.0	146.0	245.5	1.8	1.07	0.94
O92626_SNAP3_MOUSE	Snap3	SNARE-associated protein Snap3 OS-Mus musculus GN:Snap3 PE1-SV1	1	16.2	6.5	11.2	2.5	1.45	0.69	16.2	6.5	10.9	2.5	1.48	0.68
Q18865_S5C3_MOUSE	S5c3	S5 histone deacetylase corepressor complex component S5C3 OS-Mus musculus GN:S5c3 PE1-SV1	2	10.7	7.4	10.5	2.5	1.45	0.69	8.6	3.6	8.6	2.5	1.58	0.68
O95647_APK4_MOUSE	Pk4a	5'-AMP-activated protein kinase catalytic subunit alpha-1 OS-Mus musculus GN:Pk4a PE1-SV1	15	11.9	4.8	8.2	2.5	1.45	0.69	242.0	117.6	197.7	2.1	1.22	0.82
O8C0D7_ING4_MOUSE	Ing4	Inhibitor of growth factor 4 OS-Mus musculus GN:Ing4 PE1-SV2	1	57.0	22.9	39.2	2.5	1.45	0.69	57.0	22.9	38.5	2.5	1.48	0.68
O91531_SFPT7_MOUSE	Sfpt7	Septin 7 OS-Mus musculus GN:Sfpt7 PE1-SV1	4	7.7	3.1	5.3	2.5	1.45	0.69	30.9	12.5	21.0	2.5	1.47	0.68
O9P997_SFNP3_MOUSE	Sfnp3	Septin-specific protein 3 OS-Mus musculus GN:Sfnp3 PE1-SV1	1	10.5	4.8	8.0	2.5	1.45	0.69	11.1	4.8	12.1	2.5	1.46	0.68
O9Y797_SBRD1_MOUSE	Sbrd1	SL RNA-binding domain-containing protein 1 OS-Mus musculus GN:Sbrd1 PE2-SV2	1	18.5	7.4	12.7	2.5	1.45	0.69	18.5	7.4	12.5	2.5	1.48	0.68
O8D0D8_WASH1_MOUSE	Wash1	WAS protein family homolog 1 OS-Mus musculus GN:Wash1 PE1-SV1	6	12.7	5.1	8.8	2.5	1.45	0.69	83.2	31.4	52.9	2.6	1.57	0.64
O91691_PNN1_MOUSE	Pnn1	Pinn OS-Mus musculus GN:Pnn1 PE1-SV4	49	24.5	9.9	16.9	2.5	1.45	0.69	227.7	103.6	174.0	2.2	1.30	0.77
O91692_C9Z2_MOUSE	C9z2	Calponin 2 OS-Mus musculus GN:C9z2 PE1-SV1	2	5.6	2.0	2.2	2.5	1.44	0.69	40.4	97.3	104.5	2.6	1.49	0.67
P8R173_FBW1T3_MOUSE	Prrc2b	Protein PRRC2b OS-Mus musculus GN:Prrc2b PE1-SV1	5	7.3	2.9	5.0	2.5	1.44	0.69	44.9	15.5	26.1	2.9	1.72	0.58
O9L9L0_ZN281_MOUSE	Zn281	Zinc finger protein 281 OS-Mus musculus GN:Zn281 PE1-SV1	1	10.5	4.2	7.3	2.5	1.44	0.69	10.5	4.2	7.1	2.5	1.47	0.68
O9B0E9_ECSL2_MOUSE	Ecsl2	RNA-dependent protein kinase (RDK) subunit 2 OS-Mus musculus GN:Ecsl2 PE1-SV1	1	6.6	3.2	5.5	2.5	1.44	0.69	5.3	2.6	4.5	2.5	1.47	0.68
O8QV45_EPM9_MOUSE	Epm9p2a1	EPMA2-interacting protein 1 OS-Mus musculus GN:Epm9p2a1 PE2-SV1	1	6.1	2.5	4.2	2.5	1.44	0.69	6.1	2.5	4.2	2.5	1.47	0.68
O91YU8_SF1_MOUSE	Ppan	Suppressor of SW14 1 homolog OS-Mus musculus GN:Ppan PE1-SV2	4	8.0	3.2	5.5	2.5	1.44	0.69	28.7	12.7	21.4	2.2	1.34	0.75
O91DCT_SF02_MOUSE	Sf02	Stromal cell-derived factor 2 OS-Mus musculus GN:Sf02 PE1-SV1	1	5.4	2.2	3.7	2.5	1.44	0.69	5.4	2.2	3.7	2.5	1.47	0.68
Q10UHQ_NDLB_MOUSE	Ndlb	Nucleolar protein 8 OS-Mus musculus GN:Ndlb PE1-SV1	2	5.2	2.1	3.6	2.5	1.44	0.69	4.2	2.1	3.6	2.5	1.47	0.68
O9B1U0_BKFX5_MOUSE	Uhrf1bp1	Protein Uhrf1bp1 OS-Mus musculus GN:Uhrf1bp1 PE1-SV1	2	3.4	1.4	2.4	2.5	1.44	0.69	6.8	2.8	4.6	2.5	1.47	0.68
P51618_RIP1_MOUSE	Rip1	Ribosomal RNA processing protein 1 homolog A OS-Mus musculus GN:Rip1 PE2-SV2	4	10.0	4.1	7.0	2.5	1.44	0.69	40.1	18.7	31.5	2.1	1.27	0.79
Q11009_TMED1_MOUSE	Tmed1	Transmembrane emp24 domain-containing protein 1 OS-Mus musculus GN:Tmed1 PE2-SV1	2	10.1	7.0	4.1	7.0	2.5	1.44	6.9	12.5	11.5	2.5	1.47	0.68
O9CWC4_INT11_MOUSE	Cpfl1	Integrator complex subunit 11 OS-Mus musculus GN:Cpfl1 PE2-SV1	6	15.7	6.4	10.9	2.5	1.44	0.69	44.8	20.8	44.4	2.6	1.29	0.77
O9P949_NGI1_MOUSE	Gtsp4	Nucleolar GTP-binding protein 1 OS-Mus musculus GN:Gtsp4 PE2-SV3	8	11.8	4.8	8.2	2.5	1.44	0.69	128.1	62.7	105.4	2.0	1.22	0.82
O90B05_PELP1_MOUSE	Pelp1	Protein, glutamic acid- and leucine-rich protein 1 OS-Mus musculus GN:Pelp1 PE1-SV2	13	21.5	8.7	15.0	2.5	1.44	0.70	319.8	125.2	210.4	2.6	1.52	0.67
O91C91_WDHD4_MOUSE	Wdhd4	Winged-helix domain-containing protein 4 OS-Mus musculus GN:Wdhd4 PE2-SV1	2	11.4	6.0	11.4	2.5	1.44	0.70	11.4	6.0	11.4	2.5	1.47	0.68
P29341_PAPB1_MOUSE	Papb1	Polymerase II-binding protein 1 OS-Mus musculus GN:Papb1 PE1-SV2	45	77.6	31.5	54.0	2.5	1.44	0.70	4665.9	2174.0	3655.5	2.1	1.28	0.78
O91052_DPML_MOUSE	Dpm1	Dolichol phosphate mannosyltransferase subunit 1 OS-Mus musculus GN:Dpm1 PE2-SV1	1	8.5	3.4	5.9	2.5	1.44	0.70	8.5	3.4	5.8	2.5	1.46	0.68
EP9F4_P7L_MOUSE	Lmo7	Protein Lmo7 OS-Mus musculus GN:Lmo7 PE1-SV1	35	14.1	5.7	9.8	2.5	1.44	0.70	671.0	257.0	432.1	2.6	1.55	0.64
O90L15_ARHV1_MOUSE	Arhv17	Arh guanine nucleotide exchange factor 17 OS-Mus musculus GN:Arhv17 PE2-SV1	9	10.6	4.3	7.0	2.5	1.44	0.70	91.7	45.4	70.3	2.3	1.43	0.81
O924C3_XPOS_MOUSE	Xpos	Exporin 3 OS-Mus musculus GN:Xpos PE2-SV1	2	7.4	3.0	5.1	2.5	1.43	0.70	14.7	6.0	10.1	2.5	1.46	0.68
O90D83_SF2C3_MOUSE	Sf2c3	Kinectin ortholog protein Sf2c3 OS-Mus musculus GN:Sf2c3 PE2-SV1	1	8.7	3.5	6.0	2.5	1.43	0.70	8.7	3.5	5.9	2.5	1.46	0.68
O90B48_NELLS_MOUSE	Nells	Lymphocyte-specific protein 2 OS-Mus musculus GN:Nells PE2-SV2	16	16.2	6.6	11.3	2.5	1.43	0.70	46.0	16.2	26.0	2.6	1.56	0.68
O9C9C0_TOM2_MOUSE	Tom22	Mitochondrial import receptor subunit TOM22 homolog OS-Mus musculus GN:Tom22 PE2-SV1	4	12.1	8.9	2.5	1.43	0.70	51.2	19.3	32.5	2.6	1.58	0.63	
O91KX5_NUDT5_MOUSE	Nudt5	ADP-sugar pyrophosphatase OS-Mus musculus GN:Nudt5 PE1-SV1	1	29.1	11.9	20.3	2.5	1.43	0.70	29.1	11.9	20.0	2.5	1.46	0.69
P70349_HINT1_MOUSE	Hint1	Histidine triad nucleotide-binding protein 1 OS-Mus musculus GN:Hint1 PE1-SV3	1	18.7	7.6	13.1	2.5	1.43	0.70	18.7	7.6	12.9	2.5	1.46	0.69
O91937_NPM_MOUSE	Npm	Nucleophosmin OS-Mus musculus GN:Npm PE1-SV1	23	22.2	9.1	15.1	2.5	1.43	0.70	156.2	60.5	104.5	2.5	1.47	0.68
P51283_RAB12_MOUSE	Rab12	Ras-related protein Rab-12 OS-Mus musculus GN:Rab12 PE1-SV3	1	7.9	3.2	5.6	2.5	1.43	0.70	7.9	3.2	5.4	2.5	1.46	0.69
P19973_LSP1_MOUSE	Lsp1	Ras-related protein 1 OS-Mus musculus GN:Lsp1 PE1-SV2	1	5.5	2.2	3.8	2.5	1.43	0.70	5.5	2.2	3.8	2.4	1.46	0.69
O91910_CTNNL2_MOUSE	Cttnnb2	CTNNB2 N-terminal-like protein OS-Mus musculus GN:Cttnnb2 PE1-SV1	7	8.8	3.6	6.1	2.4	1.43	0.70	66.3	22.1	37.2	3.0	1.78	0.56
O90R60_RTP4_MOUSE	Rtp4														

P25799_NFKB1_MOUSE	Nfkb1	Nuclear factor NF-kappa-B p105 subunit OS-Mus musculus GN=Nfkb1 PE=1 Sv=2	11	17.6	7.4	12.8	2.4	1.38	0.72	166.2	64.1	107.8	2.6	1.54	0.65
Q80629_FL2D_MOUSE	Wtapp	Pren-mRNA splicing regulator WTAP OS-Mus musculus GN=Wtapp PE=2 Sv=3	6	18.0	7.6	13.0	2.4	1.38	0.72	103.8	45.0	75.7	2.3	1.37	0.73
Q14541_BAKB1_MOUSE	Bak1	Bac-related protein Bak1 OS-Mus musculus GN=Bak1 PE=1 Sv=1	6	22.2	9.4	16.0	2.4	1.38	0.72	137.0	58.0	23.0	2.5	1.47	0.68
P70455_STX4_MOUSE	Stx4	Stxintaxin-4 OS-Mus musculus GN=Stx4 PE=1 Sv=1	3	8.2	3.5	5.9	2.4	1.38	0.72	35.6	10.4	17.5	3.4	2.04	0.49
Q05120_RSU1_MOUSE	Rsu1	Ras suppressor protein 1 OS-Mus musculus GN=Rsu1 PE=2 Sv=3	6	31.0	13.1	22.4	2.4	1.38	0.72	184.6	77.8	130.8	2.4	1.41	0.71
Q04065_C17B_MOUSE	C17orf70	Centrosomal protein of 17 kb OS-Mus musculus GN=C17orf70 PE=1 Sv=2	5	10.0	4.2	7.0	2.4	1.38	0.72	42.7	16.2	24.0	2.3	1.47	0.73
Q02308_CP131_MOUSE	Cep131	Centrosomal protein of 131 kDa OS-Mus musculus GN=Cep131 PE=1 Sv=2	5	12.6	5.3	9.2	2.4	1.38	0.73	56.7	23.9	40.1	2.4	1.41	0.71
P24655_PESA_MOUSE	Pik3r1	Phosphatidylinositol 3-kinase regulator subunit alpha OS-Mus musculus GN=Pik3r1 PE=1 Sv=2	1	6.2	2.6	4.5	2.4	1.38	0.73	6.2	2.6	4.4	2.4	1.41	0.71
Q80853_ECHA_MOUSE	Hadha	Trihydroxyl enzyme subunit alpha, mitochondrial alpha OS-Mus musculus GN=Hadha PE=1 Sv=1	7	12.7	5.4	9.2	2.4	1.38	0.73	105.2	72.9	122.6	1.4	0.86	1.17
Q80871_CARE_MOUSE	Careap	Carboxylase alpha OS-Mus musculus GN=Careap PE=1 Sv=1	6	19.0	8.1	14.0	2.4	1.38	0.73	139.8	89.0	151.8	2.4	1.47	0.68
Q05222_THPP_MOUSE	L3h3pdp	Trans-L-3-hydroxyproline dehydratase OS-Mus musculus GN=L3h3pdp PE=1 Sv=1	1	152.9	64.8	111.2	2.4	1.38	0.73	152.9	64.8	109.0	2.4	1.40	0.71
Q05129_TAP1B_MOUSE	Sap1b	Inhibitor deacetylase complex subunit SAP1B OS-Mus musculus GN=Sap1B PE=1 Sv=1	18	31.5	13.4	22.9	2.4	1.37	0.73	879.0	441.8	742.8	2.0	1.18	0.85
P11967_AFS1_MOUSE	Afs1	AF-1 complex subunit alpha 1 OS-Mus musculus GN=Afs1 PE=1 Sv=1	41	10.6	4.5	7.7	2.4	1.37	0.73	9.8	4.0	7.3	2.8	1.40	0.71
Q02061_PLEC_MOUSE	Plec	Plectin OS-Mus musculus GN=Plec PE=1 Sv=3	130	23.1	9.8	16.8	2.4	1.37	0.73	3257.4	1515.0	2547.5	2.2	1.28	0.78
Q80298_PTH2_MOUSE	Pth2	Pectinyl-RNA hydrolase 2, mitochondrial OS-Mus musculus GN=Pth2 PE=2 Sv=1	3	4.2	1.8	3.0	2.4	1.37	0.73	36.5	14.3	24.1	2.5	1.51	0.66
Q05071_MTNA_MOUSE	Mtna	Methyltransferase L-phosphate isomerase OS-Mus musculus GN=Mtna PE=2 Sv=1	2	8.3	3.5	6.0	2.4	1.37	0.73	16.5	7.0	11.8	2.4	1.40	0.71
Q10301_C1TM_MOUSE	C1orf101	Monofunctional C1-epsilon hydroxylase synthase, mitochondrial OS-Mus musculus GN=C1orf101 PE=1 Sv=1	2	33.8	7.0	22.0	2.4	1.37	0.73	167.3	67.0	112.7	2.5	1.48	0.67
B05160_B9E1B0_MOUSE	Pdz8	PDZ domain containing 8 OS-Mus musculus GN=Pdz8 PE=1 Sv=1	1	11.3	4.8	8.3	2.4	1.37	0.73	11.3	4.8	8.1	2.4	1.40	0.72
Q17700_S2PDH_MOUSE	Sup20h	Transcription factor SPT20 homolog OS-Mus musculus GN=Sup20h PE=1 Sv=1	1	13.7	5.8	10.0	2.4	1.37	0.73	13.7	5.8	9.8	2.4	1.40	0.72
P58773_TPM1_MOUSE	Tpm1	Tropomyosin alpha 1 chain OS-Mus musculus GN=Tpm1 PE=1 Sv=1	21	70.1	28.9	51.2	2.3	1.37	0.73	1951.9	1168.8	1967.1	1.7	0.99	1.01
Q10702_ERK6_MOUSE	Erk6	Interferon-gamma-inducible double-stranded RNA-dependent protein kinase activator A OS-Mus musculus	3	20.4	8.9	14.9	2.3	1.37	0.73	70.0	29.0	42.8	2.4	1.43	0.70
Q01926_BAG2_MOUSE	Bag2	BAG family molecular chaperone regulator 2 OS-Mus musculus GN=Bag2 PE=1 Sv=1	8	22.0	9.4	16.1	2.3	1.37	0.73	180.5	86.4	145.3	2.1	1.24	0.80
Q01905_P48A_MOUSE	P48a	EL SUAD1 protein ligase P48A OS-Mus musculus GN=P48a PE=1 Sv=2	3	7.1	3.0	5.2	2.3	1.37	0.73	38.6	14.4	24.3	2.7	1.59	0.63
Q80156_VTGA_MOUSE	Vtga	Vesicle transport through interaction with 1 SNAREs homolog 1A OS-Mus musculus GN=Vtga11a PE=1 Sv=1	2	13.8	6.0	10.2	2.3	1.37	0.73	3.0	7.1	3.0	2.1	1.40	0.71
Q06941_DDX51_MOUSE	Ddx51	ATP-dependent RNA helicase DDX51 OS-Mus musculus GN=Ddx51 PE=2 Sv=1	8	10.0	4.3	7.3	2.3	1.37	0.73	80.8	38.2	64.3	2.1	1.26	0.80
P21121_MCM6_MOUSE	Mcm6	DNA replication licensing factor MCM6 OS-Mus musculus GN=Mcm6 PE=1 Sv=1	9	11.4	4.9	8.3	2.3	1.37	0.73	133.4	65.0	109.3	2.1	1.22	0.82
P24455_CAPP_MOUSE	Capp	Macrograph-capping protein OS-Mus musculus GN=Capp PE=1 Sv=2	6	13.5	5.8	9.9	2.3	1.37	0.73	173.3	70.1	117.9	2.5	1.47	0.68
Q05822_NKMC_MOUSE	Nkmc	KIF-8 essential regulator OS-Mus musculus GN=Nkmc PE=1 Sv=2	2	13.1	5.6	9.7	2.3	1.37	0.73	17.4	84.3	144.9	2.6	1.47	0.68
Q05800_AKAP8_MOUSE	Akap8	A-kinase anchor protein 8 OS-Mus musculus GN=Akap8 PE=1 Sv=1	9	39.7	17.0	29.1	2.3	1.37	0.73	201.0	147.8	248.6	2.0	1.21	0.83
Q05924_TBC1D1_MOUSE	Tbc1d1	TBC1 domain family member 1 OS-Mus musculus GN=Tbc1d1 PE=1 Sv=1	3	7.6	3.3	5.6	2.3	1.36	0.73	24.4	9.1	15.3	2.7	1.59	0.63
Q05921_SARD1_MOUSE	Sard1	SPFH domain family member 1 OS-Mus musculus GN=Sard1 PE=1 Sv=1	5	12.0	5.1	8.8	2.3	1.36	0.73	49.4	14.0	24.8	2.8	1.59	0.59
Q05974_MED24_MOUSE	Med24	Mediator of RNA polymerase II transcription subunit 24 OS-Mus musculus GN=Med24 PE=1 Sv=1	8	10.5	4.5	7.7	2.3	1.36	0.73	49.9	39.9	67.0	2.3	1.36	0.73
Q61584_FR1_MOUSE	Fxr1	Fragile X mental retardation syndrome-related protein 1 OS-Mus musculus GN=Fxr1 PE=1 Sv=2	22	23.9	10.2	17.5	2.3	1.36	0.73	1024.7	439.2	738.5	2.3	1.39	0.72
Q05387_TM6M_MOUSE	Tmm6	Mitochondrial import inner membrane translocase subunit TM6M OS-Mus musculus GN=Tmm6	3	9.9	4.2	7.3	2.3	1.36	0.73	29.6	11.9	20.1	2.5	1.48	0.68
Q02382_SPTB_MOUSE	Sptb	Structural elongation factor SPTB OS-Mus musculus GN=Sptb PE=1 Sv=2	2	15.5	7.0	12.0	2.3	1.36	0.73	89.2	40.7	68.2	2.3	1.45	0.67
F06559_F06559_MOUSE	Chd8	Chromodomain-helicase-DNA-binding protein 8 (Fragment) OS-Mus musculus GN=Chd8 PE=4 Sv=1	1	10.9	4.7	8.0	2.3	1.36	0.73	10.9	4.7	7.9	2.3	1.39	0.72
Q80851_NUP54_MOUSE	Nup54	Nuclear pore complex protein Nup54 OS-Mus musculus GN=Nup54 PE=1 Sv=1	6	9.0	3.8	6.6	2.3	1.36	0.73	59.1	28.2	47.5	2.1	1.24	0.80
P14348_LAMP1_MOUSE	Lamp1	Lysosome-associated membrane glycoprotein 1 OS-Mus musculus GN=Lamp1 PE=1 Sv=2	3	29.7	12.7	21.8	2.3	1.36	0.73	76.7	32.6	54.7	2.4	1.40	0.71
P07393_HNDS_MOUSE	Hnds	YADP-activating protein OS-Mus musculus GN=Hnds PE=1 Sv=1	2	6.8	2.9	5.0	2.3	1.36	0.73	7.3	5.8	7.8	2.3	1.40	0.71
P14084_ARF5_MOUSE	Arf5	ADP-ribosylation factor 5 OS-Mus musculus GN=Arf5 PE=1 Sv=1	3	11.8	5.7	8.3	2.3	1.36	0.73	53.3	20.8	34.9	2.6	1.52	0.66
Q35841_ARF5_MOUSE	Arf5	Apoptosis inhibitor 5 OS-Mus musculus GN=Arf5 PE=1 Sv=2	15	16.6	7.1	12.2	2.3	1.36	0.73	283.0	121.8	204.7	2.3	1.38	0.72
P17449_APM1_MOUSE	Apm1	Apoptosis inhibitor 1 OS-Mus musculus GN=Apm1 PE=1 Sv=1	1	6.4	2.8	4.9	2.3	1.36	0.73	9.6	4.0	7.0	2.3	1.47	0.68
Q35845_NOM1_MOUSE	Nom1	Nucleolar MFG domain-containing protein 1 OS-Mus musculus GN=Nom1 PE=2 Sv=2	3	10.2	4.4	7.5	2.3	1.36	0.73	27.8	11.4	19.2	2.4	1.45	0.69
P14206_RPSA_MOUSE	Rpsa	40S ribosomal protein SA OS-Mus musculus GN=Rpsa PE=1 Sv=4	19	17.8	7.6	13.1	2.3	1.36	0.73	1078.8	624.2	1049.6	1.7	1.03	0.97
Q91KPE_CHRC1_MOUSE	Chrc1	Chromatin accessibility complex protein 1 OS-Mus musculus GN=Chrc1 PE=1 Sv=1	1	13.0	5.6	9.6	2.3	1.36	0.73	13.0	5.6	9.4	2.3	1.39	0.72
Q91075_HIF1_MOUSE	Hif1	Hypoxia-inducible transcription factor 1 OS-Mus musculus GN=Hif1 PE=1 Sv=2	2	18.1	7.7	13.2	2.3	1.36	0.74	121.8	57.6	101.2	2.5	1.47	0.68
Q91961_PP1R1_MOUSE	Pp1r1	Pren-mRNA processing factor 19 OS-Mus musculus GN=Pp1r1 PE=1 Sv=1	14	58.2	25.0	42.8	2.3	1.36	0.74	1286.8	690.2	1160.9	1.9	1.01	0.90
F62888_F62888_MOUSE	Wiz	Protein WIZ (Fragment) OS-Mus musculus GN=Wiz PE=1 Sv=1	13	25.0	10.7	18.4	2.3	1.36	0.74	306.5	173.8	292.2	1.8	1.05	0.95
Q90961_R602_MOUSE	Erc1	ERC1/R602-interacting/CST1 family member 1 OS-Mus musculus GN=Erc1 PE=1 Sv=1	13	8.7	3.8	6.5	2.3	1.36	0.74	155.8	132.7	204.7	2.4	1.40	0.71
Q05855_RPR8_MOUSE	Rpr8	Ribosomal RNA-processing protein 8 OS-Mus musculus GN=Rpr8 PE=1 Sv=1	8	11.2	4.8	8.3	2.3	1.36	0.74	9.6	16.2	23.3	2.9	1.48	0.72
Q60101_DCL1_MOUSE	Dync1l1	Cytoplasmic dynein 1 light intermediate chain 2 OS-Mus musculus GN=Dync1l1 PE=1 Sv=2	7	22.1	9.5	16.3	2.3	1.36	0.74	158.0	50.3	84.5	3.1	1.87	0.54
P26309_TN1_MOUSE	Tnfr1	Tumor necrosis factor receptor type 1 OS-Mus musculus GN=Tnfr1 PE=1 Sv=1	117	85.0	36.5	62.0	2.3	1.36	0.74	1424.21	588.17	899.0	2.4	1.44	0.69
Q05121_ZC11_MOUSE	Zc11	Zinc finger CCH domain-containing protein 11 OS-Mus musculus GN=Zc11 PE=2 Sv=1	2	14.7	6.3	11.0	2.3	1.36	0.74	183.3	83.3	143.3	2.3	1.40	0.71
Q05083_LRCD4_MOUSE	Lrrcd4	Leucine-rich repeat-containing protein 40 OS-Mus musculus GN=Lrrcd4 PE=2 Sv=1	14	16.3	7.0	12.6	2.3	1.36	0.74	263.2	115.4	194.0	2.3	1.36	0.74
Q07113_BRD2_MOUSE	Brd2	Bromodomain-containing protein 2 OS-Mus musculus GN=Brd2 PE=1 Sv=1	8	10.4	4.5	7.8	2.3	1.36	0.74	128.2	53.2	89.4	2.4	1.43	0.70
Q60436_C1PB_MOUSE	C1pb	Cathepsin-like peptidase B1 protein homolog OS-Mus musculus GN=C1pb PE=1 Sv=1	1	4.4	1.9	3.3	2.3	1.36	0.74	4.4	1.9	3.2	2.3	1.48	0.72
Q05804_WDR2_MOUSE	Wdr2	WD repeat-containing protein 42 OS-Mus musculus GN=Wdr2 PE=1 Sv=1	4	14.6	6.5	11.3	2.3	1.36	0.74	21.6	8.5	14.5	2.3	1.48	0.67
Q61768_KIF5_MOUSE	Kif5b	Kinesin-1 heavy chain OS-Mus musculus GN=Kif5b PE=1 Sv=2	6	13.5	5.8	9.9	2.3	1.36	0.74	74.1	35.3	59.3	2.2	1.38	0.72
Q31704_QST1M_MOUSE	Padi6	Nucleic acid diester isomerase A6 OS-Mus musculus GN=Padi6 PE=2 Sv=1	19	44.2	19.0	32.6	2.3	1.35	0.74	160.5	74.94	126.0	1.9	1.11	0.90
Q05922_Q9912_MOUSE	Nr2f1	GTP-binding protein NR2F1 OS-Mus musculus GN=NR2F1 PE=1 Sv=1	2	23.8	10.4	17.6	2.3	1.35	0.74	205.5	104.0	181.2	2.3	1.45	0.68
F06543_F06543_MOUSE	Ncwr2	Nuclear receptor corepressor 2 OS-Mus musculus GN=Ncwr2 PE=1 Sv=1	20	13.6	5.9	10.3	2.3	1.35	0.74	300.2	161.2	271.1	1.9	1.11	0.90
G58873_G58873_MOUSE	Srp19	Signal recognition particle 19 kDa protein OS-Mus musculus GN=Srp19 PE=4 Sv=1	3	30.0	12.9	22.2	2.3	1.35	0.74	71.8	30.8	51.9	2.3	1.38	0.72
A24243_CuI4B_MOUSE	CuI4b	Cullin-4B OS-Mus musculus GN=CuI4B PE=1 Sv=1	2	13.6	5.9	10.1	2.3	1.35	0.74	27.2	11.7	19.3	2.3	1.38	0.73
Q05342_HMA2_MOUSE	Hma2	Nucleolar subunit alpha OS-Mus musculus GN=Hma2 PE=1 Sv=1	3	9.5	4.2	7.3	2.3	1.35	0.74	30.3	24.0	35.6	2.3	1.39	0.72
Q60FV4_NOP58_MOUSE	Nop58	Cullin protein 58 OS-Mus musculus GN=Nop58 PE=1 Sv=1	31	22.2	9.6	16.5	2.3	1.35	0.74	777.2	366.0	615.4	2.1	1.26	0.79
A2AGH6_MED12_MOUSE	Med12	Mediator of RNA polymerase II transcription subunit 12 OS-Mus musculus GN=Med12 PE=1 Sv=2	5	8.8	3.8	6.6	2.3	1.35	0.74	55.0	23.5	39.5	2.3	1.39	0.72

O31314_RPA1_MOUSE	Poir1a	DNA-directed RNA polymerase subunit RPA1 OS-Mus musculus GN-Poir1a PE1 SV-2	2	6.7	3.0	5.1	2.2	1.31	0.76	13.4	6.0	10.1	2.2	1.33	0.75
Q6A044_LAR4B_MOUSE	Lar4pb	Larval protein 4B OS-Mus musculus GN-Lar4pb PE1 SV-2	6	10.9	4.9	8.3	2.2	1.31	0.76	73.3	31.9	53.7	2.3	1.37	0.73
P62310_SNA3_MOUSE	Sna3	Small nuclear ribonucleoprotein S1 OS-Mus musculus GN-Sna3 PE1 SV-1	4	10.4	21.4	36.8	2.2	1.31	0.76	231.8	390.8	1.8	1.36	0.74	
P70347_TANK_MOUSE	Tank	TRAF family member-associated NF-kappa-B activator OS-Mus musculus GN-Tank PE1 SV-1	1	4.9	2.2	3.8	2.2	1.31	0.76	4.9	2.2	3.7	2.2	1.33	0.75
Q8H442_ARHG2_MOUSE	Arhg2f1	Rho guanine nucleotide exchange factor 12 OS-Mus musculus GN-Arhg2f1 PE1 SV-2	1	7.1	3.2	5.4	2.2	1.31	0.77	7.1	3.2	5.3	2.2	1.33	0.75
Q85B29_CAN2_MOUSE	Can2	Catalytic subunit 2 OS-Mus musculus GN-Can2 PE1 SV-2	4	6.5	2.9	4.6	2.2	1.31	0.77	5.8	2.9	4.7	2.2	1.33	0.75
Q91V81_RBM2_MOUSE	Rbm2a	RNA-binding protein 42 OS-Mus musculus GN-Rbm2a PE1 SV-1	2	9.8	4.4	7.5	2.2	1.31	0.77	19.6	8.7	14.7	2.2	1.33	0.75
O54774_AP3D1_MOUSE	Ap3d1	Ap3 complex subunit delta-1 OS-Mus musculus GN-AP3d1 PE1 SV-1	10	13.1	5.9	10.1	2.2	1.31	0.77	174.6	116.6	196.1	1.5	1.89	1.12
R0V011_RUVB1_MOUSE	Ruvb1	RuvB-like 1 OS-Mus musculus GN-Ruvb1 PE1 SV-1	37	37.2	16.6	28.5	2.2	1.31	0.77	367.47	156.42	2630.2	2.3	1.38	0.73
Q8R191_SNA3_MOUSE	Sna3	Small nuclear ribonucleoprotein S1 OS-Mus musculus GN-Sna3 PE1 SV-1	4	10.4	21.4	36.8	2.2	1.31	0.77	231.8	390.8	1.8	1.36	0.74	
P21617_MGMT_MOUSE	Mgmt	Methylated DNA-protein-cysteine methyltransferase OS-Mus musculus GN-Mgmt PE1 SV-1	1	4.2	1.9	3.2	2.2	1.31	0.77	4.2	1.9	3.1	2.2	1.33	0.75
Q8R465_BBX_MOUSE	Bbx	HMG box transcription factor BBX OS-Mus musculus GN-Bbx PE1 SV-2	1	11.3	5.0	8.7	2.2	1.30	0.77	11.3	5.0	8.5	2.2	1.33	0.75
Q8R141_EHD2_MOUSE	Ehd2	ERL1/Ermen protein-retaining receptor OS-Mus musculus GN-Ehd2 PE1 SV-2	1	24.9	11.1	19.9	2.2	1.30	0.77	24.9	11.1	19.9	2.2	1.33	0.75
Q8K284_TFC1_MOUSE	Gtf1c1	General transcription factor 3c, polypoid type 1 OS-Mus musculus GN-Gtf1c1 PE1 SV-2	2	9.6	7.3	11.1	2.2	1.30	0.77	19.1	8.5	14.4	2.2	1.33	0.75
Q8B771_RF02X_MOUSE	Rbf2ox	RNA-binding protein fox-1 homolog 2 OS-Mus musculus GN-Rbf2ox PE1 SV-2	14	17.6	7.9	13.5	2.2	1.30	0.77	470.5	275.6	463.4	1.7	1.02	0.98
QJ3U71_EYTL1_MOUSE	Eytl1	Extended synaptotagmin-1 OS-Mus musculus GN-Eytl1 PE1 SV-2	28	16.4	7.4	12.6	2.2	1.30	0.77	598.3	258.3	434.3	2.3	1.38	0.73
P84076_ARF1_MOUSE	Arf1	ADP-ribosylation factor 1 OS-Mus musculus GN-Arf1 PE1 SV-2	8	20.6	9.2	15.5	2.2	1.30	0.77	142.2	291.6	242.5	2.0	1.30	0.83
Q5D707_KIAA1551_MOUSE	Kiaa1551	Uncharacterized protein KIAA1551 OS-Mus musculus GN-Kiaa1551 PE-2 SV-2	1	3.1	1.4	2.4	2.2	1.30	0.77	3.1	1.4	2.4	2.2	1.33	0.75
Q8C648_SMC2_MOUSE	Smc2	Structural maintenance of chromosomes protein 2 OS-Mus musculus GN-Smc2 PE1 SV-2	52	29.1	13.1	22.4	2.2	1.30	0.77	1550.4	863.2	1451.5	1.8	1.07	0.94
Q8V86_NSD2_MOUSE	Wnt5c1	Histone H3-lysine methyltransferase NSD2 OS-Mus musculus GN-Wnt5c1 PE1 SV-2	8	12.4	5.5	9.5	2.2	1.30	0.77	107.4	54.7	92.0	2.0	1.17	0.86
F7BRW6_FTRPW6_MOUSE	Sec5a	Protein Sec5 (Fragment) OS-Mus musculus GN-Sec5a PE1 SV-1	3	7.1	3.2	5.3	2.2	1.30	0.77	30.3	20.4	34.3	1.6	0.94	0.96
Q99M9_RNPS1_MOUSE	Rnps1	RNA-binding protein with serine-rich domain 1 OS-Mus musculus GN-Rnps1 PE1 SV-1	11	28.9	12.9	22.2	2.2	1.30	0.77	1048.5	463.9	780.1	2.3	1.34	0.74
Q99BWS_PSD12_MOUSE	Pamd12	26S proteasome non-ATPase regulatory subunit 12 OS-Mus musculus GN-Pamd12 PE1 SV-4	4	18.7	8.4	14.4	2.2	1.30	0.77	73.7	32.7	54.9	2.3	1.34	0.75
P21126_U844_MOUSE	Ubr1a	Ubiquitin-like protein 4A OS-Mus musculus GN-Ubr1a PE1 SV-1	3	14.6	6.5	11.2	2.2	1.30	0.77	39.3	21.1	35.4	1.9	1.11	0.80
Q7T771_ELP1_MOUSE	Htkap	Oligomer complex protein 1 OS-Mus musculus GN-Htkap PE1 SV-2	26	19.3	8.7	14.8	2.2	1.30	0.77	628.4	240.4	404.2	2.6	1.55	0.64
Q9K303_OC3_MOUSE	Orc3	Orient recognition complex subunit 3 OS-Mus musculus GN-Orc3 PE1 SV-1	1	11.8	5.3	9.1	2.2	1.30	0.77	11.8	5.3	8.9	2.2	1.32	0.75
Q60212_SCAM1_MOUSE	Scamp1	Secretory carrier-associated membrane protein 1 OS-Mus musculus GN-Scamp1 PE1 SV-1	2	24.9	11.2	19.2	2.2	1.30	0.77	49.8	22.7	37.6	2.2	1.32	0.76
Q99H0_HNR12_MOUSE	Hnrp12	Transcription initiator factor TIE2 OS-Mus musculus GN-Hnrp12 PE1 SV-1	9	10.7	4.8	8.3	2.2	1.30	0.77	179.8	98.9	146.8	1.7	1.03	0.81
Q6A044_F12A2_MOUSE	Fam120a	Constitutive-associated of PPAR-gamma-like protein 1 OS-Mus musculus GN-FAM120A PE1 SV-1	24	28.0	12.6	21.6	2.2	1.30	0.77	608.7	288.9	485.8	2.1	1.25	0.80
Z3H18	Znfx18	Zinc finger CCHH domain-containing protein 18 OS-Mus musculus GN-Znfx18 PE1 SV-1	31	34.6	15.6	26.7	2.2	1.30	0.77	3110.9	580.8	976.7	5.4	1.39	0.71
Q8R224_WAT12_MOUSE	Wat12	Wnt12-interfering protein 12 OS-Mus musculus GN-Wat12 PE1 SV-1	2	5.8	2.6	4.5	2.2	1.30	0.77	11.7	5.2	6.7	2.2	1.32	0.75
Q8R0V2_CCL1_MOUSE	Cacul1	CCKC-associated and calinin domain-containing protein 1 OS-Mus musculus GN-Cacul1 PE1 SV-2	3	3.4	7.6	5.9	2.2	1.30	0.77	7.6	3.4	5.8	2.2	1.32	0.76
Q99104_MYO5A_MOUSE	Myo5a	Unconventional myosin-Va OS-Mus musculus GN-Myo5a PE1 SV-2	10	9.2	4.2	7.1	2.2	1.29	0.77	93.6	41.6	69.9	2.3	1.34	0.75
Q91Y71_RBM5_MOUSE	Rbm5	RNA-binding protein 5 OS-Mus musculus GN-Rbm5 PE1 SV-1	10	25.0	11.3	19.3	2.2	1.29	0.77	877.8	149.1	250.6	5.9	3.30	0.29
Q8B752_ARHG2_MOUSE	Arhg2	Rho guanine nucleotide exchange factor 2 OS-Mus musculus GN-Arhg2 PE1 SV-1	4	10.7	4.8	8.3	2.2	1.29	0.77	10.7	4.8	8.3	2.2	1.34	0.74
Q8B677_RBGPR_MOUSE	Rab3gap2	Rab3 GTPase-activating protein non-catalytic subunit OS-Mus musculus GN-Rab3gap2 PE1 SV-1	3	10.8	4.9	8.3	2.2	1.29	0.77	40.6	18.4	31.0	2.2	1.31	0.76
P62915_TF2B_MOUSE	Gtf2b	Transcription initiation factor IIIB OS-Mus musculus GN-Gtf2b PE1 SV-1	6	9.8	4.4	7.6	2.2	1.29	0.77	85.6	30.6	51.5	2.8	1.66	0.60
Q8R371_BRD8_MOUSE	Brd8	Bromodomain-containing protein 8 OS-Mus musculus GN-Brd8 PE1 SV-2	7	7.3	3.3	5.7	2.2	1.29	0.77	63.9	24.1	40.5	2.7	1.58	0.63
Q9C001_KAB3_MOUSE	Kab3	Structural maintenance of chromosomes protein 3 OS-Mus musculus GN-Kab3 PE1 SV-1	11	11.7	5.4	9.3	2.2	1.29	0.77	384.5	117.3	193.6	2.0	1.34	0.77
Q8C647_SMC4_MOUSE	Smc4	Structural maintenance of chromosomes protein 4 OS-Mus musculus GN-Smc4 PE1 SV-1	44	28.0	12.6	21.7	2.2	1.29	0.77	1467.3	670.1	1126.8	2.2	1.30	0.77
Q62419_SHG3_MOUSE	Shg3p1	Endophilin-III OS-Mus musculus GN-Shg3p1 PE1 SV-1	13	43.3	19.6	33.5	2.2	1.29	0.77	674.4	689.5	1159.0	1.0	0.58	1.72
Q32328_TCL1_MOUSE	Tcl1a	Endophilin-associated regulatory protein 1 OS-Mus musculus GN-Tcl1a PE1 SV-1	1	11.7	5.4	9.3	2.2	1.29	0.77	11.7	5.4	9.3	2.2	1.32	0.77
P97461_R55_MOUSE	Rps5	40S ribosomal protein S5 OS-Mus musculus GN-Rps5 PE1 SV-3	9	20.1	9.1	15.5	2.2	1.29	0.77	294.5	115.6	194.3	2.5	1.52	0.66
P53564_CUX1_MOUSE	Cux1	Hobox protein cux-1 OS-Mus musculus GN-Cux1 PE1 SV-1	16	11.0	5.0	8.5	2.2	1.29	0.77	182.2	98.3	165.3	1.9	1.10	0.91
E11089_UBR2N_MOUSE	Ubr2n	Ubiquitin-conjugating enzyme E2 N OS-Mus musculus GN-Ubr2n PE1 SV-1	4	49.1	22.2	38.0	2.2	1.29	0.77	190.5	81.2	136.5	2.3	1.40	0.72
Q8R0U9_TAF1_MOUSE	Taf11	Transcription initiator factor TIE1 OS-Mus musculus GN-Taf11 PE1 SV-2	2	11.7	5.4	9.3	2.2	1.29	0.77	11.7	5.4	9.3	2.2	1.32	0.77
P62702_R54X_MOUSE	Rps4x	40S ribosomal protein S4, X isoform OS-Mus musculus GN-Rps4x PE1 SV-2	39	55.1	24.9	42.7	2.2	1.29	0.77	5364.5	1969.0	3310.7	2.7	1.62	0.62
Q91V11_CKAP2_MOUSE	Ckap2	Cytoskeleton-associated protein 2 OS-Mus musculus GN-CKAP2 PE1 SV-1	5	8.4	3.8	6.5	2.2	1.29	0.78	42.0	24.1	40.5	1.7	1.03	0.97
Q8C0C2_SFPA_MOUSE	Farsa	Phenylalanine-tyrosine transferase OS-Mus musculus GN-Farsa PE1 SV-1	6	14.8	6.7	11.2	2.2	1.29	0.78	14.8	6.7	11.2	2.2	1.31	0.76
Q9D0F1_LMAN1_MOUSE	Lman1	Protein ERGIC-53 OS-Mus musculus GN-Lman1 PE1 SV-1	4	21.3	8.6	16.5	2.2	1.29	0.78	59.7	36.2	60.9	1.6	0.98	1.02
Q8RFD1_CHL1_MOUSE	Ctlc1	Clathrin heavy chain 1 OS-Mus musculus GN-Ctlc1 PE1 SV-3	66	26.7	12.1	20.7	2.2	1.29	0.78	2484.6	1068.9	2.3	1.38	0.72	0.72
Q91U75_PHAX_MOUSE	Phax	Phosphorylated actin-binding RNA export protein OS-Mus musculus GN-Phax PE1 SV-1	1	12.4	5.6	9.6	2.2	1.29	0.78	12.4	5.6	9.4	2.2	1.31	0.76
Q91B42_WPRF_MOUSE	Wprf	SMAD3 p55 subfamily member 1 OS-Mus musculus GN-Wprf PE1 SV-1	1	16.4	7.4	12.7	2.2	1.29	0.78	16.4	7.4	12.5	2.2	1.31	0.76
P23607_ZFA_MOUSE	Zfa	Zinc finger autoantigen protein OS-Mus musculus GN-Zfa PE1 SV-2	7	16.4	7.4	12.7	2.2	1.29	0.78	16.4	7.4	12.5	2.2	1.31	0.76
Q62D3X_DDX3_MOUSE	Ddx3x	ATP-dependent RNA helicase DDX3X OS-Mus musculus GN-DDX3X PE1 SV-3	64	56.0	25.4	43.3	2.2	1.28	0.78	6263.2	2703.2	4545.4	2.3	1.38	0.73
Q8R66E_EOX1_MOUSE	Eox1	Early endosome antigen 1 OS-Mus musculus GN-Eox1 PE1 SV-2	25	15.8	7.2	12.3	2.2	1.28	0.78	492.5	198.0	332.9	2.5	1.48	0.68
P98111_PRR1_MOUSE	Prr1	Endoplasmic reticulum protein 1 OS-Mus musculus GN-Prr1 PE1 SV-2	6	16.7	7.6	12.7	2.2	1.28	0.78	16.7	7.6	12.7	2.2	1.34	0.80
Q82B23_STAT5_MOUSE	Stat5a	Signal transducer and activator of transcription 5A OS-Mus musculus GN-Stat5a PE1 SV-1	4	5.6	2.6	4.4	2.2	1.28	0.78	24.3	10.5	17.7	2.3	1.37	0.73
P48804_NOP58_MOUSE	Nop58	Nucleolar protein 9 OS-Mus musculus GN-Nop58 PE1 SV-1	6	14.0	6.4	10.9	2.2	1.28	0.78	92.9	36.6	61.5	2.5	1.51	0.66
Q9V0M6_HNR11_MOUSE	Hnrp11	HnRNP-associated nuclear ribonucleoprotein 11 OS-Mus musculus GN-Hnrp11 PE1 SV-1	18	31.4	14.6	25.8	2.2	1.28	0.78	288.8	118.1	181.8	2.6	1.61	0.63
Q53864_CNS1_MOUSE	Cns1	Alpha-1 syntrophin OS-Mus musculus GN-Cns1 PE1 SV-1	7	15.5	7.1	12.1	2.2	1.28	0.78	119.1	71.0	119.4	1.7	1.00	1.00
Q61234_SNTAL_MOUSE	Sntal	COP1-1 syntrophin OS-Mus musculus GN-Sntal PE1 SV-1	2	10.5	4.8	8.2	2.2	1.28	0.78	21.0	9.6	16.1	2.2	1.30	0.77
Q9C211_UHP1_MOUSE	Uhp11	Protein UHP-1 small nuclear RNA-associated protein 11 OS-Mus musculus GN-Uhp11 PE1 SV-2	1	11.6	5.9	15.3	2.2	1.28	0.78	19.5	8.9	15.0	2.2	1.30	0.77
Q8R8Y6_WTF2_MOUSE	Wtf2	Transcription termination factor 2, mitotic OS-Mus musculus GN-Wtf2 PE1 SV-1	1	19.5	8.9	15.8	2.2	1.28	0.78	119.8	51.8	81.8	2.2	1.30	0.77
Q91XK7_Y1FA_MOUSE	Y1fa	Protein Y1FA OS-Mus musculus GN-Y1fa PE1 SV-1	2	12.6	5.8	9.9	2.2	1.28	0.78	25.2	11.5	19.4	2.2	1.30	0.77
P11440_CDK1_MOUSE	Cdk1	Cyclin-dependent kinase 1 OS-Mus musculus GN-Cdk1 PE1 SV-3	32	31.0	14.2	24.3	2.2	1.28	0.78	1513.3	626.6	1053.6	2.4	1.44	0.70
Q62072_CAND2_MOUSE	Cand2	Cullin-associated NEDD8-associated protein 2 OS-Mus musculus GN-Cand2 PE1 SV-2	7	8.7	4.0	6.9	2.2	1.27	0.78	59.5	23.5	39.6	2.5	1.50	0.67
Q9SK91_NOD1_MOUSE	Nod1	Protein kinase of 40S OS-Mus musculus GN-Nod1 PE1 SV-1	2	6.4	2.9	5.1	2.2	1.27	0.78	6.4	2.9	5.0	2.2	1.32	0.77
P51881_ADT2_MOUSE	Sic25a5	ADP-RT translocase 2 OS-Mus musculus GN-Sic25a5 PE1 SV-3	3	44.5	20.4	34.3	2.2	1.27	0.78	441.2	70.5	118.6	2.0	1.19	0.84
Q99N10_ANR1_MOUSE	Ankr17	Ankyrin repeat domain-containing protein 17 OS-Mus musculus GN-Ankr17 PE1 SV-2	26	13.1	6.0	10.3	2.2	1.27	0.78	138.8	161.6	271.7	2.7	1.61	0.62
Q8R66E_BCL7L_MOUSE	Bcl7l	BCL7L lymphoma 4 protein OS-Mus musculus GN-Bcl7l PE1 SV-1	9	10.7	4.8	8.3	2.2	1.27	0.78	110.4	41.1	69.1	2.7	1.60	0.63
Q53874_SAT1_MOUSE	Sic14a	Neutral amino acid transporter 4 OS-Mus musculus GN-Sic14a PE1 SV-1	1	7.8	3.6	6.1	2.2	1.27	0.79	7.8	3.6	6.0	2.2	1.30	0.77
QJ1D17_FOJ1_MOUSE	Pdim7	PDZ and LIM domain protein 7 OS-Mus musculus GN-Pdim7 PE1 SV-2	20	25.2	11.5	19.8	2.2	1.27	0.79	641.6	326.8	549.5	2.0	1.17	0.86
Q60716_SAMH1_MOUSE	Samh1	Deoxyribose triphosphate triphosphohydrolyase SAMH1 OS-Mus musculus GN-Samh1 PE1 SV-1	16	26.0	11.9	20.4	2.2								

P38647_GRP75_MOUSE	Hspa9p	Stress-70 protein, mitochondrial OS-Mus musculus GN-Hspa9 PE1 S-V1-3	11	17.2	8.1	14.0	2.1	1.23	0.81	175.1	109.9	184.7	1.6	0.95	1.05
OSB06_SRCR1_MOUSE	Rorc1	Serine/arginine-related protein 5 OS-Mus musculus GN-Rorc1 PE1 S-V1-1	3	15.3	7.2	12.4	2.1	1.23	0.81	53.6	29.6	49.7	1.8	1.08	0.99
OSK46_RBP5_MOUSE	Rbp5	Protein ribosomal protein L5 OS-Mus musculus GN-Rbp5 PE1 S-V1-2	1	11.5	10.2	11.6	2.1	1.23	0.81	23.2	23.2	23.2	2.0	1.14	0.66
OSB8N_FLNA_MOUSE	Flna	Filamin-B OS-Mus musculus GN-Flna PE1 S-V1-5	80	66.2	31.4	53.8	2.1	1.23	0.81	7416.9	3051.6	5131.3	2.4	1.45	0.69
OSU57_GZD1_MOUSE	Gzrd1n1	General transcription factor IIH repeat domain-containing protein 1 OS-Mus musculus GN-Gzrd1	2	10.0	4.7	8.1	2.1	1.23	0.81	19.9	9.4	15.9	2.9	1.26	0.80
OSB810_U551_MOUSE	U551	Small nuclear ribonucleoprotein A2 OS-Mus musculus GN-U551 PE1 S-V1-2	57	43.4	20.0	35.0	2.1	1.23	0.81	3226.2	1786.6	3021.8	1.6	1.06	0.84
OSB87K_HAU5_MOUSE	Hau5	HAU5 agmatin-like complex subunit 1 OS-Mus musculus GN-Hau5 PE1 S-V1-2	1	7.2	3.4	5.8	2.1	1.23	0.81	7.2	3.4	5.7	2.1	1.25	0.80
OSQ92Z_HECD1_MOUSE	Hectd1	E ubiquitin-protein ligase HECTD1 OS-Mus musculus GN-Hectd1 PE1 S-V1-2	4	7.2	3.4	5.9	2.1	1.23	0.81	33.7	19.6	32.9	1.7	1.03	0.97
OS4933_NEDD4_MOUSE	Nedd4	E ubiquitin-protein ligase NEDD4 OS-Mus musculus GN-Nedd4 PE1 S-V1-3	12	11.0	5.2	8.9	2.1	1.23	0.81	132.2	76.0	127.8	1.7	1.03	0.97
P5291_NAA1_MOUSE	Naa1	Importin subunit alpha 1 OS-Mus musculus GN-Naa1 PE1 S-V1-2	19	14.3	6.8	11.6	2.1	1.23	0.81	183.1	101.0	181.1	2.0	1.14	0.84
OS61735_CD47_MOUSE	Cd47	Leukocyte surface antigen CD47 OS-Mus musculus GN-CD47 PE1 S-V1-2	2	22.0	10.5	17.9	2.1	1.23	0.81	44.0	20.9	35.2	2.1	1.25	0.80
OS0824_FPI1_MOUSE	Fpi1	Protein Rfx19-3 end-processing factor FPI1 OS-Mus musculus GN-Fpi1 PE1 S-V1-1	20	29.3	13.9	23.9	2.1	1.23	0.81	880.5	389.8	655.4	2.3	1.34	0.74
OS5222_Utk_MOUSE	Utk	ATP-dependent RNA helicase OS-Mus musculus GN-Utk PE1 S-V1-2	21	18.7	15.2	16.2	2.1	1.23	0.81	454.2	245.4	364.5	2.1	1.25	0.80
OS09W6_SNL1_MOUSE	Snl1	Sorting nexin-1 OS-Mus musculus GN-Snl1 PE1 S-V1-1	2	10.1	4.8	7.8	2.1	1.23	0.82	8.5	48.0	80.7	1.9	1.12	0.85
OS08734_BAK_MOUSE	Bak1	Bcl-2 homologous antagonist/killer OS-Mus musculus GN-Bak1 PE1 S-V1-3	2	8.3	3.9	6.8	2.1	1.23	0.82	16.6	7.9	13.3	2.1	1.25	0.80
PE2844_RP51_MOUSE	Rp51	40S ribosomal protein S15 OS-Mus musculus GN-Rp51 PE2 S-V1-2	3	5.4	2.6	4.4	2.1	1.22	0.82	51.7	17.0	28.7	3.0	1.81	0.55
OSR202_S55_MOUSE	S55	Synaptonemal complex protein S55 OS-Mus musculus GN-S55 PE2 S-V1-1	1	9.6	4.6	7.9	2.1	1.22	0.82	71.9	48.5	120.9	0.7	0.40	2.49
OS6418_LBP7_MOUSE	Lbp7	Ubiquitin carboxyl-terminal hydrolase 7 OS-Mus musculus GN-Lbp7 PE1 S-V1-1	12	12.4	5.9	10.1	2.1	1.22	0.82	168.5	77.9	131.1	2.2	1.29	0.78
OS9L48_NMD3_MOUSE	Nmd3	60S ribosomal export protein NMD3 OS-Mus musculus GN-Nmd3 PE2 S-V1-1	2	7.8	3.7	6.4	2.1	1.22	0.82	15.5	7.4	12.5	2.1	1.25	0.80
OS5XAC_O5SK3_MOUSE	Vertf1	Protein Vertf1 OS-Mus musculus GN-Vertf1 PE1 S-V1-1	1	17.6	8.4	14.4	2.1	1.22	0.82	17.6	8.4	14.2	2.1	1.25	0.80
OS9RUB_HCS_MOUSE	Hcs	Hepatocyte growth factor-regulated tyrosine kinase substrate OS-Mus musculus GN-Hcs PE1 S-V1-1	3	9.0	4.8	8.5	2.1	1.22	0.82	12.1	20.8	22.2	2.9	0.77	1.29
OSCK96_NIP7_MOUSE	Nip7	60S ribosome subunit biogenesis protein NIP7 homolog OS-Mus musculus GN-Nip7 PE2 S-V1-1	3	8.5	4.1	7.0	2.1	1.22	0.82	34.8	26.5	44.5	1.3	0.78	1.28
OS0106_HOAC1_MOUSE	Hdac1	Histone deacetylase 1 OS-Mus musculus GN-Hdac1 PE1 S-V1-1	23	21.7	10.4	17.8	2.1	1.22	0.82	918.9	345.0	580.0	2.7	1.58	0.63
OS0205_I7G6_MOUSE	I7g6	Synaptonemal complex protein I7G6 OS-Mus musculus GN-I7g6 PE2 S-V1-1	1	9.6	4.6	7.9	2.1	1.22	0.82	71.9	48.5	120.9	0.7	0.40	2.49
OSQ633_LONM_MOUSE	Lonm1	Long protease homolog, mitochondrial OS-Mus musculus GN-Lonm1 PE1 S-V1-2	11	10.1	4.8	8.2	2.1	1.22	0.82	157.4	63.8	107.3	2.5	1.47	0.68
OS2104_MME6_MOUSE	Mme6	Mediator of RNA polymerase II transcription subunit 6 OS-Mus musculus GN-Mme6 PE1 S-V1-2	1	23.9	11.4	19.6	2.1	1.22	0.82	23.9	11.4	19.2	2.1	1.24	0.80
OS08M4_RL1_MOUSE	Rpl71	60S ribosomal protein L7-like 1 OS-Mus musculus GN-Rpl71 PE2 S-V1-1	2	10.2	4.9	8.3	2.1	1.22	0.82	20.4	9.7	16.4	2.1	1.24	0.80
R14471_EIART1_MOUSE	Eiart1	EIF4E OS-Mus musculus GN-Eiart1 PE1 S-V1-1	1	5.1	2.5	4.1	2.1	1.22	0.82	7.5	2.5	4.1	2.1	1.24	0.80
OS21M4_GCG2_MOUSE	Gcg2	Golgi subfamily A member 2 OS-Mus musculus GN-Gcg2 PE2 S-V1-2	3	25.5	12.2	20.9	2.1	1.22	0.82	74.3	50.7	84.5	1.5	0.88	1.14
OS0207_ZN68_MOUSE	Znf67	Zinc finger protein 687 OS-Mus musculus GN-Znf67 PE1 S-V1-1	2	17.0	8.1	13.9	2.1	1.22	0.82	143.8	70.5	118.6	2.0	1.21	0.83
OS1093_D0V1_MOUSE	D0v1	ATP-dependent RNA helicase D0V1 OS-Mus musculus GN-D0V1 PE1 S-V1-1	29	25.0	12.2	20.9	2.1	1.22	0.82	562.7	245.4	412.7	2.4	1.42	0.70
OS8349_CDC16_MOUSE	Cdc16	Cdk division cyclin protein 16 homolog OS-Mus musculus GN-Cdc16 PE2 S-V1-1	2	5.4	2.6	4.4	2.1	1.22	0.82	10.8	5.2	8.7	2.1	1.21	0.81
OS6411_TOP2B_MOUSE	Top2b	DNA topoisomerase 2-beta OS-Mus musculus GN-Top2b PE1 S-V1-2	20	12.2	5.8	10.0	2.1	1.22	0.82	346.0	219.5	369.1	1.6	0.94	1.07
OS0423_TRF5_MOUSE	Trf5	Transcription factor p65 OS-Mus musculus GN-Trf5 PE1 S-V1-1	1	6.7	3.2	5.5	2.1	1.22	0.82	6.7	3.2	5.4	2.1	1.24	0.81
OS4190_DTR1L_MOUSE	Dtr1l	E3 ubiquitin-protein ligase DTR1L OS-Mus musculus GN-Dtr1l PE2 S-V1-1	2	15.8	5.4	9.2	2.1	1.22	0.82	358.1	186.9	358.1	1.9	1.19	0.84
EQ04M5_HELZ2_MOUSE	Helz2	Helicase with zinc finger domain 2 OS-Mus musculus GN-Helz2 PE1 S-V1-1	1	9.6	4.6	7.9	2.1	1.22	0.82	9.6	4.6	7.7	2.1	1.24	0.81
OS1918_FUBP1_MOUSE	Fubp1	Far upstream element-binding protein 1 OS-Mus musculus GN-Fubp1 PE1 S-V1-1	2	19.4	9.3	16.0	2.1	1.22	0.82	38.8	18.6	31.3	2.1	1.24	0.81
OS9R07_ABP8L_MOUSE	Abp8l	A kinase anchor protein 8-like OS-Mus musculus GN-Abp8l PE1 S-V1-1	4	13.7	6.6	11.3	2.1	1.22	0.82	49.4	25.6	43.1	1.9	1.15	0.87
OS1257_LURCH1_MOUSE	Lurch1	Nucleic acid binding protein LURCH1 OS-Mus musculus GN-Lurch1 PE1 S-V1-1	2	19.9	14.8	25.4	2.1	1.22	0.82	29.6	29.6	29.6	2.0	1.19	0.84
OS8909_NU13_MOUSE	Nup133	Nucleolar pore complex protein Nup133 OS-Mus musculus GN-Nup133 PE1 S-V1-2	5	13.8	6.6	11.4	2.1	1.21	0.82	87.3	33.3	55.9	2.6	1.56	0.64
OS9237_KCC2_MOUSE	Kcc2	Calcium/calmodulin-dependent protein kinase type II subunit gamma OS-Mus musculus GN-Kcc2	4	9.0	4.3	7.4	2.1	1.21	0.82	47.0	18.6	31.2	2.5	1.50	0.66
OS1335_VGAC_MOUSE	Vgac	Calcium-associated protein VGAC OS-Mus musculus GN-Vgac PE1 S-V1-1	1	40.6	18.5	32.7	2.1	1.21	0.82	176.8	40.6	176.8	2.0	1.19	0.84
P12979_MYOG_MOUSE	Myog	Myogenin OS-Mus musculus GN-Myog PE1 S-V1-2	1	32.8	25.1	27.1	2.1	1.21	0.82	32.8	15.8	26.5	2.1	1.24	0.81
OS77M4_MB1A_MOUSE	Myb1a	Myb-binding protein 1A OS-Mus musculus GN-Myb1a PE1 S-V1-2	83	32.2	15.5	26.6	2.1	1.21	0.83	4263.7	1980.2	3329.7	2.2	1.28	0.78
OS1065_CAPRIN1_MOUSE	Caprin1	Caprin-1 OS-Mus musculus GN-Caprin1 PE1 S-V1-1	17	29.2	14.1	24.1	2.1	1.21	0.83	1009.1	388.9	653.9	2.6	1.54	0.65
OS10H4_E2B2_MOUSE	E2b2	Protein E2B transcription factor 2B subunit OS-Mus musculus GN-E2B2 PE1 S-V1-1	2	15.2	7.4	12.7	2.1	1.21	0.83	61.3	14.8	24.9	2.1	1.23	0.81
OS10H6_TAD2A_MOUSE	Tad2a	Transcriptional adapter 2-alpha OS-Mus musculus GN-Tad2a PE1 S-V1-1	1	7.1	3.4	5.9	2.1	1.21	0.83	7.1	3.4	5.7	2.1	1.23	0.81
PE2309_RBP2_MOUSE	Rbp2	Small nuclear ribonucleoprotein G OS-Mus musculus GN-Rbp2 PE1 S-V1-1	3	19.0	9.1	15.7	2.1	1.21	0.83	61.4	26.4	44.4	2.3	1.23	0.81
OS9P97_V07_MOUSE	V07	Esp1p-7 OS-Mus musculus GN-V07 PE1 S-V1-1	10	12.4	5.3	9.2	2.1	1.21	0.83	67.0	40.0	67.0	2.0	1.23	0.72
OS9P92_O5XS_MOUSE	O5xs	Serine/threonine protein kinase O5XS OS-Mus musculus GN-O5XS PE1 S-V1-1	8	14.0	6.8	11.6	2.1	1.21	0.83	112.0	55.8	92.1	2.0	1.22	0.82
OS9VEE_LMCD1_MOUSE	Lmcd1	LIM and cysteine-rich domain protein 1 OS-Mus musculus GN-Lmcd1 PE1 S-V1-1	2	5.8	2.8	4.8	2.1	1.21	0.83	11.5	5.6	9.3	2.1	1.23	0.81
OS9216_NELF1_MOUSE	Nelf1	Negative elongation factor D OS-Mus musculus GN-Nelf1 PE2 S-V1-2	7	11.6	5.6	9.6	2.1	1.21	0.83	113.2	65.4	110.1	1.7	1.03	0.97
OS9151_PNA3_MOUSE	Pna3	PRAL family protein 3 OS-Mus musculus GN-Pna3 PE1 S-V1-2	4	10.3	4.8	8.2	2.1	1.21	0.83	27.2	13.1	23.2	2.1	1.23	0.81
OS9QZ7_A5UN_MOUSE	A5un	Protein asunder homolog OS-Mus musculus GN-A5un PE1 S-V1-2	6	18.3	8.8	15.1	2.1	1.21	0.83	115.5	44.3	74.5	2.6	1.55	0.64
OS88E6_IF74_MOUSE	If74	Intraflagellar transport protein 74 homolog OS-Mus musculus GN-If74 PE1 S-V1-1	2	8.0	3.9	6.6	2.1	1.21	0.83	16.0	7.7	13.0	2.1	1.23	0.81
OS9VE9_S5F4_MOUSE	S5f4	Serine/threonine kinase S5F4 OS-Mus musculus GN-S5F4 PE1 S-V1-1	8	12.2	5.8	10.1	2.1	1.21	0.83	287.0	107.9	185.8	2.7	1.58	0.63
OS9R48_SNS_MOUSE	Sns	Sorting nexin-4 OS-Mus musculus GN-Sns PE1 S-V1-1	2	10.8	4.9	8.6	2.1	1.21	0.83	18.0	9.2	18.0	2.0	1.23	0.81
OS91Y8_SN4_MOUSE	Sns4	Sorting nexin-4 OS-Mus musculus GN-Sns4 PE2 S-V1-2	6	13.9	6.7	11.5	2.1	1.21	0.83	93.1	43.5	73.3	2.1	1.27	0.79
OS91X6_MMTA2_MOUSE	Mmta2	Multiple myeloma tumor-associated protein 2 homolog OS-Mus musculus GN-Mmta2 PE2 S-V1-2	2	10.8	4.7	7.7	2.1	1.21	0.83	29.3	14.2	23.8	2.1	1.23	0.81
OS9P71_P9E1_MOUSE	P9e1	DNA poly(ADP-ribose) polymerase 1 OS-Mus musculus GN-P9e1 PE1 S-V1-1	2	20.8	10.0	17.3	2.1	1.21	0.83	24.1	12.1	20.1	2.1	1.23	0.81
OS8X22_CND1_MOUSE	Cnd2	Condensin complex subunit 1 OS-Mus musculus GN-Cnd2 PE1 S-V1-2	34	18.7	9.0	15.5	2.1	1.21	0.83	783.7	358.2	602.3	2.2	1.30	0.77
OS8127_A5PM_MOUSE	A5pm	Abnormal spindle-like microcephaly-associated protein homolog OS-Mus musculus GN-A5pm	2	8.4	4.1	7.1	2.1	1.21	0.83	16.8	8.1	13.7	2.1	1.23	0.81
OS10R3_S1MP_MOUSE	S1mp	Sarcolemmal membrane-associated protein OS-Mus musculus GN-S1mp PE1 S-V1-2	2	7.1	3.4	5.9	2.1	1.21	0.83	28.4	15.9	26.7	1.8	1.06	0.94
OS8V7E_PC10_MOUSE	Pc10	Cell receptor-associated protein 10 OS-Mus musculus GN-Pc10 PE1 S-V1-1	2	19.9	6.2	11.0	2.1	1.21	0.83	60.2	16.0	31.2	2.1	1.24	0.75
P29037_TBP_MOUSE	Tbp	TATA-box-binding protein OS-Mus musculus GN-Tbp PE1 S-V1-1	2	8.9	4.3	7.4	2.1	1.21	0.83	17.9	8.6	14.5	2.1	1.23	0.81
OS9V8V_NVI_MOUSE	Nvi	Nucleolar valosin-containing protein-like OS-Mus musculus GN-Nvi PE1 S-V1-1	16	14.6	7.1	12.1	2.1	1.20	0.83	297.1	167.6	281.9	1.8	1.05	0.95
OS9Y9K_GAK_MOUSE	Gak	Cyclin G-associated kinase OS-Mus musculus GN-Gak PE1 S-V1-2	9	9.6	4.8	8.2	2.1	1.20	0.83						

Q90206_TOLIP_MOUSE	Tollip	Toll-interacting protein OS=Mus musculus GN= Tollip PE=1 SV=1	4	10.8	5.4	9.3	2.0	1.16	0.86	53.6	23.4	39.3	2.3	1.36	0.73
Q874_CDC7C_MOUSE	Apof9b	Factor VIIIB OS=Mus musculus GN=Apof9b PE=2 SV=1	5	79.9	40.3	69.1	2.0	1.16	0.86	342.7	183.8	309.1	1.9	1.11	0.90
Q889F1_FB1_MOUSE	Raf1	Raf1 OS=Mus musculus GN=Raf1 PE=2 SV=1	6	7.1	3.6	7.2	1.6	1.16	0.86	7.2	3.6	7.2	1.6	1.18	0.94
Q05558_FB2_MOUSE	Fra1	Factor VIIIb intron 22 protein OS=Mus musculus GN=Fra1 PE=2 SV=1	1	6.2	3.1	5.4	2.0	1.16	0.87	6.2	3.1	5.3	2.0	1.18	0.85
Q987D1_TB83_MOUSE	Tubb3	Tubulin beta 3 chain OS=Mus musculus GN=Tubb3 PE=1 SV=1	25	27.7	14.0	24.0	2.0	1.16	0.87	183.7	108.51	182.6	1.8	1.09	0.92
Q987D1_S1NG_MOUSE	Sorting nexin 6 OS=Mus musculus GN=Sorting nexin 6 PE=1 SV=1	3	18.1	9.4	13.2	2.0	1.16	0.87	18.1	9.4	13.2	2.0	1.18	0.92	
Q139F4_FRMD8_MOUSE	Frm8	FERM domain-containing protein 8 OS=Mus musculus GN=Frm8 PE=1 SV=2	1	7.6	3.8	6.0	2.0	1.16	0.87	7.6	3.8	6.5	2.0	1.18	0.85
Q139F4_DCOCS_MOUSE	Exc5	Excyst component 5 OS=Mus musculus GN=Exc5 PE=1 SV=2	2	4.6	2.3	4.0	2.0	1.15	0.87	9.1	4.6	7.8	2.0	1.18	0.85
Q5V5QD_KAT7_MOUSE	Kat7	Histone acetyltransferase KAT7 OS=Mus musculus GN=Kat7 PE=1 SV=1	14	12.0	6.1	10.4	2.0	1.15	0.87	221.6	111.5	187.5	2.0	1.18	0.85
Q5V5QD_M45_MOUSE	Nr4a1	Nuclear receptor coactivator 5 OS=Mus musculus GN=Nr4a1 PE=1 SV=1	25	26.0	13.2	20.5	2.0	1.15	0.87	514.4	260.0	462.0	1.8	1.04	0.92
Q6R18B_TBORA_MOUSE	Tor3a	Tor1A-IP3 OS=Mus musculus GN=Tor3a PE=1 SV=2	1	4.8	2.4	4.1	2.0	1.15	0.87	4.8	2.4	4.0	2.0	1.18	0.85
Q6R18B_KIF15_MOUSE	Kif15	Kinesin-like protein KIF15 OS=Mus musculus GN=Kif15 PE=1 SV=1	3	4.5	2.3	3.9	2.0	1.15	0.87	13.5	13.2	22.2	1.0	0.61	0.65
Q5DB83_U2AF1_MOUSE	U2af1	Splicing factor U2AF 1 OS=Mus musculus GN=U2af1 PE=1 SV=4	9	20.1	10.3	20.5	2.0	1.15	0.87	1116.2	594.1	999.0	1.9	1.12	0.90
Q8BK11_BAI2_MOUSE	Bai2	Brain-specific angiogenesis inhibitor 1-associated protein 2 OS=Mus musculus GN=Bai2 PE=1 SV=1	3	5.0	2.5	4.3	2.0	1.15	0.87	4.8	2.4	12.3	2.0	1.58	0.63
Q80X90_FLNB_MOUSE	Flnb	Filamin-B OS=Mus musculus GN=Flnb PE=1 SV=3	48	32.0	16.2	27.8	2.0	1.15	0.87	1611.3	822.4	1382.9	2.0	1.17	0.86
A2B2E1_LAS1L_MOUSE	Las1l	Ribosomal biogenesis protein LAS1L OS=Mus musculus GN=Las1l PE=1 SV=1	18	14.7	7.5	12.8	2.0	1.15	0.87	311.9	143.7	241.7	2.2	1.29	0.77
P97310_MCM2_MOUSE	Mcm2	DNA replication licensing factor MCM2 OS=Mus musculus GN=Mcm2 PE=1 SV=3	33	12.1	6.1	10.5	2.0	1.15	0.87	559.4	276.6	465.0	1.8	1.10	0.91
Q900F4_NKAP_MOUSE	Nkap	NR-kappa-B activating protein OS=Mus musculus GN=Nkap PE=2 SV=1	4	17.6	8.9	15.5	2.0	1.15	0.87	71.4	33.6	56.6	2.1	1.26	0.79
Q922N8_NCLG6_MOUSE	Act16a	Actin-like protein 6 OS=Mus musculus GN=Act16a PE=1 SV=2	14	28.2	14.3	24.5	2.0	1.15	0.87	889.1	319.6	537.3	2.8	1.65	0.60
Q9SL55_PWP1_MOUSE	Pwp1	Proteolytic tripeptidase protein 1 homolog OS=Mus musculus GN=Pwp1 PE=1 SV=1	1	10.1	5.1	8.8	2.0	1.15	0.87	10.1	5.1	8.6	2.0	1.17	0.85
Q9C2E1_MED18_MOUSE	Med18	Mediator of RNA polymerase II transcription subunit 18 OS=Mus musculus GN=Med18 PE=2 SV=1	6	16.2	8.2	14.1	2.0	1.15	0.87	8.2	4.1	13.8	2.0	1.17	0.85
Q90787_PP1L2_MOUSE	Pp1l2	Peptidyl-prolyl cis-trans isomerase like 2 OS=Mus musculus GN=Pp1l2 PE=1 SV=2	14	9.2	4.6	8.0	2.0	1.15	0.87	131.2	68.1	114.5	1.9	1.15	0.87
Q9C0C0_I5T1_MOUSE	Ist1	IST1 homolog OS=Mus musculus GN=Ist1 PE=2 SV=1	4	21.1	10.7	18.3	2.0	1.15	0.87	163.4	47.0	79.0	3.5	2.07	0.48
Q9L91F_KTCC8_MOUSE	Ktcc8	RNA-splicing ligase Rtt108 homolog OS=Mus musculus GN=Ktcc8 PE=2 SV=1	12	17.9	9.1	15.6	2.0	1.15	0.87	259.0	117.0	230.3	1.9	1.12	0.89
Q43276_H13_MOUSE	Hist13b1b	Histone H13.5 OS=Mus musculus GN=Hist13b1b PE=1 SV=2	13	332.2	168.7	289.2	2.0	1.15	0.87	8623.8	4396.8	7393.1	2.0	1.17	0.86
Q9C0C0_UQCQ2_MOUSE	Uqcq2	Ubiquitin-cytochrome c reductase complex assembly factor 2 OS=Mus musculus GN=Uqcq2 PE=1 SV=1	1	7.1	3.6	6.2	2.0	1.15	0.87	7.1	3.6	6.1	2.0	1.17	0.85
Q9J028_FL1_MOUSE	Fl1	Protein flightless-1 homolog OS=Mus musculus GN=Fl1 PE=1 SV=1	18	12.4	6.3	10.8	2.0	1.15	0.87	247.7	124.7	209.7	2.0	1.18	0.85
Q50522_PGRC1_MOUSE	Pgrc1	Protein phosphatase 1 regulatory subunit 1 OS=Mus musculus GN=Pgrc1 PE=1 SV=1	5	16.1	8.1	12.5	2.0	1.15	0.87	100.0	39.4	66.0	2.1	1.18	0.91
Q90878_ING5_MOUSE	Ing5	Inhibitor of growth protein 5 OS=Mus musculus GN=Ing5 PE=1 SV=1	3	13.1	6.7	11.4	2.0	1.15	0.87	102.6	57.6	96.9	1.8	1.06	0.94
Q9L720_MKCKC_MOUSE	Ctck22bpb	Kinase/threonine protein kinase MKCK beta OS=Mus musculus GN=Ctck22bpb PE=1 SV=2	11	11.0	5.6	9.6	2.0	1.15	0.87	157.7	59.0	99.2	2.7	1.59	0.63
Q139F4_ESRBL2_MOUSE	Ctcfp1	CTCF-associated protein 1 OS=Mus musculus GN=Ctcfp1 PE=1 SV=2	24	16.1	8.1	13.2	2.0	1.15	0.87	303.4	113.7	211.2	2.7	1.59	0.63
Q9L720_C42_MOUSE	Abc2	ATP-binding cassette sub-family 4 member 2 OS=Mus musculus GN=Abc2 PE=2 SV=1	6	13.6	6.9	11.8	2.0	1.15	0.87	65.9	40.0	62.7	1.6	0.98	0.20
Q9C09U_EF1B_MOUSE	Ef1b	Eukaryotic translation initiation factor 1b OS=Mus musculus GN=Ef1b PE=2 SV=2	1	5.0	2.5	4.3	2.0	1.15	0.87	5.0	2.5	4.3	2.0	1.17	0.86
Q61249_IGBP1_MOUSE	Igbp1	Immunoglobulin-binding protein 1 OS=Mus musculus GN=Igbp1 PE=1 SV=1	7	8.4	4.3	7.4	2.0	1.14	0.87	58.9	27.8	46.7	2.1	1.26	0.79
Q50702_E9P02_MOUSE	E9p02	Tropomyosin alpha 2 OS=Mus musculus GN=E9p02 PE=1 SV=1	1	10.1	5.1	9.2	2.0	1.14	0.87	10.1	5.1	9.2	2.0	1.17	0.86
D32087_D32087_MOUSE	Katna2	Kat5 protein 660 ATPase-containing subunit A like 2 OS=Mus musculus GN=Katna2 PE=1 SV=1	1	5.2	2.7	4.6	2.0	1.14	0.87	5.2	2.7	4.5	2.0	1.17	0.86
P56873_SAA2_MOUSE	Saa2	Kyoto syndrome/erythroderma autoantigen 1 homolog OS=Mus musculus GN=Saa2 PE=2 SV=1	2	16.4	8.4	14.3	2.0	1.14	0.88	32.8	16.7	28.1	2.0	1.17	0.86
Q90926_DDA1_MOUSE	Dda1	DDA1 and DDB1-associated protein 1 OS=Mus musculus GN=Dda1 PE=2 SV=1	1	5.8	3.0	5.1	2.0	1.14	0.88	5.8	3.0	5.0	2.0	1.17	0.86
Q9C09U_S9P2_MOUSE	S9p2	Serine/threonine phosphatase 9 OS=Mus musculus GN=S9p2 PE=1 SV=4	1	10.1	5.1	16.9	2.0	1.14	0.88	112.8	48.6	82.8	1.8	1.10	0.92
Q8C0E2_VP26B_MOUSE	Vp26b	Vacuolar protein sorting-associated protein 26B OS=Mus musculus GN=Vp26b PE=1 SV=1	2	12.1	6.2	10.6	2.0	1.14	0.88	24.2	12.4	20.8	2.0	1.16	0.86
P28659_CELF1_MOUSE	Celf1	CUGBP1 Elav-like family member 1 OS=Mus musculus GN=Celf1 PE=1 SV=2	5	12.2	6.3	10.7	2.0	1.14	0.88	54.7	28.9	48.7	1.9	1.12	0.89
Q5S0Q0_CTC1_MOUSE	Ctcf	Cis-acting transcription factor 1 OS=Mus musculus GN=Ctcf PE=1 SV=1	3	6.1	3.1	4.6	2.0	1.14	0.88	6.2	3.1	4.6	2.0	1.18	0.86
P43133_C072_MOUSE	Nr2f2	COUP transcription factor 2 OS=Mus musculus GN=Nr2f2 PE=1 SV=1	8	23.5	12.0	20.6	2.0	1.14	0.88	210.4	96.9	163.0	2.2	1.29	0.77
A2A641_GPAT8_MOUSE	Gpat8	Gpat8 domain-containing protein 8 OS=Mus musculus GN=Gpat8 PE=1 SV=1	12	14.6	7.5	12.8	2.0	1.14	0.88	217.6	105.3	177.1	2.1	1.23	0.81
Q87127_TB3_MOUSE	Tb3	Transducin beta-like protein 3 OS=Mus musculus GN=Tb3 PE=1 SV=1	5	10.2	5.2	9.0	2.0	1.14	0.88	51.6	26.8	45.0	1.9	1.15	0.87
Q18B80_SF130_MOUSE	Sf130	Protein phosphatase 1 regulatory subunit 130 OS=Mus musculus GN=Sf130 PE=1 SV=2	1	10.6	5.4	9.3	2.0	1.14	0.88	10.6	5.4	9.3	2.0	1.15	0.87
Q59555_P9B10_MOUSE	Pim1b10	Protein kinase subunit beta type-10 OS=Mus musculus GN=Pim1b10 PE=1 SV=1	2	12.1	6.2	10.7	1.9	1.14	0.88	24.2	12.4	20.9	1.9	1.16	0.86
Q91W05_TAF5L_MOUSE	Taf5l	TAF5-like RNA polymerase II p300/CBP-associated factor-associated factor 65 kDa subunit 5 OS=	5	7.2	3.7	6.7	1.9	1.14	0.88	36.8	15.9	26.7	1.3	0.78	0.73
Q91W05_PHLB2_MOUSE	Phlb2	Phlebotomus homology 1 domain-containing protein 2 OS=Mus musculus GN=Phlb2 PE=2 SV=2	39	22.0	11.3	18.1	2.0	1.14	0.88	916.1	531.8	864.0	1.9	1.14	0.90
P27441_KRCC3_MOUSE	Krcc3	K-rnc repair cross-complementing protein 3 OS=Mus musculus GN=Krc3 PE=2 SV=1	8	17.2	8.8	14.9	1.9	1.14	0.88	181.1	89.9	151.2	1.6	0.95	0.26
Q61624_ZNF48_MOUSE	Znf48	Zinc finger protein 148 OS=Mus musculus GN=Znf48 PE=1 SV=2	11	9.3	4.8	8.1	1.9	1.14	0.88	110.0	59.9	100.7	1.8	1.09	0.92
Q40336_VP26A_MOUSE	Vp26a	Vacuolar protein sorting-associated protein 26A OS=Mus musculus GN=Vp26a PE=1 SV=1	11	17.3	8.9	15.2	1.9	1.14	0.88	231.7	123.8	208.2	1.9	1.11	0.90
P65113_L13F_MOUSE	L13f	L13 protein eta OS=Mus musculus GN=L13f PE=2 SV=1	5	3.3	1.6	2.8	1.9	1.14	0.88	3.8	1.6	2.6	1.9	1.14	0.89
Q90717_MYO1C_MOUSE	Myo1c	Unconventional myosin-1c OS=Mus musculus GN=Myo1c PE=1 SV=1	49	36.5	18.8	32.2	1.9	1.13	0.88	2390.8	1266.8	2130.1	1.9	1.12	0.89
P48962_AD01_MOUSE	Sic25a4	ADP-ATP translocase 1 OS=Mus musculus GN=Sic25a4 PE=1 SV=4	10	37.5	18.5	31.8	1.9	1.13	0.88	423.3	252.0	423.8	1.7	1.00	1.00
Q92211_PRRF3_MOUSE	Prrf3	UAF1/Small nuclear ribonucleoprotein Prrf3 OS=Mus musculus GN=Prrf3 PE=2 SV=1	10	17.5	9.0	15.5	1.9	1.13	0.88	205.9	126.3	212.3	1.6	0.97	1.00
P14424_AKAP4_MOUSE	Akap4	Kinase A OS=Mus musculus GN=Akap4 PE=1 SV=1	10	9.1	4.6	8.1	1.9	1.13	0.88	12.7	7.1	12.7	1.9	1.17	0.91
Q6Y85_FEE2_MOUSE	Fee2	Fasculation and elongation protein nete 2 OS=Mus musculus GN=Fee2 PE=2 SV=1	2	10.0	5.1	8.8	1.9	1.13	0.88	19.9	10.3	17.3	1.9	1.15	0.87
Q91248_NEFL_MOUSE	Nefl	Negative elongation factor 1 OS=Mus musculus GN=Nefl PE=1 SV=1	3	7.6	3.9	6.7	1.9	1.13	0.88	27.6	20.5	34.4	1.3	0.80	1.25
Q6C746_C13A_MOUSE	C13a	Cell cycle 13A protein OS=Mus musculus GN=C13a PE=2 SV=1	9	18.1	9.1	16.8	1.9	1.13	0.88	208.0	134.0	208.0	1.9	1.12	0.89
Q91W14_WPR4_MOUSE	Wpr4	WD repeat domain phosphoinositide-interacting protein 4 OS=Mus musculus GN=Wpr4 PE=1 SV=1	1	12.9	6.6	11.4	1.9	1.13	0.89	12.9	6.6	11.2	1.9	1.15	0.87
Q8084A_KAN12_MOUSE	Kan12	KAT5 regulatory NLS complex subunit 2 OS=Mus musculus GN=Kan12 PE=2 SV=1	2	8.4	4.3	7.4	1.9	1.13	0.89	16.7	8.6	14.5	1.9	1.15	0.87
P61368_DY1_MOUSE	Dyn1l1	Dynamin light chain 1, cytoplasmic OS=Mus musculus GN=Dyn1l1 PE=1 SV=1	2	52.3	27.0	46.4	1.9	1.13	0.89	104.6	54.1	90.9	1.9	1.15	0.87
Q909D4_CCR2_MOUSE	Ccr2	Chemokine receptor type 2 OS=Mus musculus GN=Ccr2 PE=1 SV=2	7	8.2	4.2	7.3	1.9	1.13	0.89	26.7	14.2	25.5	1.9	1.12	0.88
Q6NS07_LTV1_MOUSE	Ltv1	Protein LTV1 homolog OS=Mus musculus GN=LTV1 PE=2 SV=2	2	9.3	4.8	8.3	1.9	1.13	0.89	18.7	9.7	16.3	1.9	1.15	0.87
Q63318_TB1F_MOUSE	Trimb28	Transcription interactor protein 1-beta OS=Mus musculus GN=Trimb28 PE=1 SV=3	58	34.7	18.0	30.7	1.9	1.13	0.89	4303.7	2096.3	3524.9	2.1	1.22	0.82
Q81248_NTM14_MOUSE	Ntm14	N-terminal Xaa-Pro-Lys-N-methyltransferase 1 OS=Mus musculus GN=Ntm14 PE=2 SV=3	2	7.1	3.7	6.3	1.9	1.13	0.89	14.2	7.4	12.4	1.9	1.15	0.87
Q88712_CTRP1_MOUSE	Ctrp1	Histidine triad domain-containing protein 1 OS=Mus musculus GN=Ctrp1 PE=1 SV=2	3	10.6	5.4	9.3	1.9	1.14	0.88	26.7	14.0	24.2	1.9	1.12	0.88
Q92456_SMC6_MOUSE	Smc6	Structural maintenance of chromosomes protein 6 OS=Mus musculus GN=Smc6 PE=2 SV=1	2	4.7	2.4	4.2	1.9	1.13	0.89	4.9	4.9	8.2	1.9	1.15	0.87
Q6P458_INT1_MOUSE	Ints1	Integrator complex subunit 1 OS=Mus musculus GN=Ints1 PE=2 SV=2	8	5.8	3.0	5.1	1.9	1.13	0.89	49.9	39.2	65.9	1.3	0.76	1.32
P51660_DNR4_MOUSE	Dnr4	Perinatal multifunctional protein 4 OS=Mus musculus GN=Dnr4 PE=1 SV=1	3	10.6	5.4	9.3	1.9	1.12	0.89	10.6	5.4	9.3	1.9	1.10	0.89
Q91M93_ABP4_MOUSE	Abp4	ADP-ribosylation factor-like protein 6-interacting protein 4 OS=Mus musculus GN=Abp4 PE=1 SV=1	9	11.0	6.0	10.3	1.9	1.12	0.89	8.0	4.2	7.0	1.9	1.15	0.87
Q91M93_SAF2_MOUSE	Saf2	Scaffold attachment factor 2B OS=Mus musculus GN=Saf2 PE=2 SV=2	9	11.6	6.0	10.3	1.9	1.12	0.89	131.7	71.2	119.7	1.8	1.10	0.91
Q90111_N															

L0N7N1_KIF14_MOUSE	KIF14	Kinesin-like protein KIF14 OS=Musculus GN:KIF14 PE=1 SV=1	2	7.3	3.9	6.7	1.9	1.08	0.92	145.5	7.8	13.2	1.9	1.10	0.91
P58854_GCF3_MOUSE	Tubgcp3	Gamma-tubulin complex component 3 OS=Musculus GN:Tubgcp3 PE=2 SV=2	3	9.1	4.9	8.4	1.9	1.08	0.92	32.8	32.8	20.7	2.7	1.58	0.83
Q5CYV6_IL12_MOUSE	IL12	Interleukin enhancer-binding factor 2 OS=Musculus GN:IL12 PE=1 SV=1	3	29.8	16.1	27.6	16.1	1.08	0.92	1564.9	469.4	759.9	3.4	2.02	1.69
B0K013_TMF1_MOUSE	Tmf1	TATA-element modulatory factor OS=Musculus GN:Tmf1 PE=1 SV=1	6	5.4	2.9	5.0	1.9	1.08	0.92	32.6	22.0	37.0	1.5	0.88	1.14
Q5QWV8_ASAP1_MOUSE	Asap1	Arf-GAP with SH3 domain, ANK repeat and PH1 domain-containing protein 1 OS=Musculus GN:Asap1 PE=1 SV=1	4	8.8	4.7	8.1	1.9	1.08	0.92	40.0	19.3	32.4	2.1	1.24	0.81
Q52219_TGFL1_MOUSE	Tgfl1	Transforming growth factor-beta type 1 OS=Musculus GN:Tgfl1 PE=1 SV=1	1	11.6	6.3	10.7	6.3	1.08	0.92	11.6	6.3	10.7	6.3	1.08	0.92
Q1UQU0_BRD9_MOUSE	Brd9	Bromodomain-containing protein 9 OS=Musculus GN:Brd9 PE=1 SV=1	3	5.8	3.1	5.3	1.9	1.08	0.92	25.6	12.2	20.6	2.1	1.25	0.80
Q9J956_DERL1_MOUSE	Derl1	Derlin-1 OS=Musculus GN:Derl1 PE=1 SV=1	3	7.4	4.0	6.8	1.9	1.08	0.92	90.9	52.6	88.4	1.9	1.12	0.89
Q9J846_STRM_MOUSE	Strm4	Striatin-4 OS=Musculus GN:Strm4 PE=1 SV=1	2	11.3	6.1	10.4	1.9	1.08	0.92	22.6	12.2	20.5	1.9	1.10	0.91
Q10821_AKCC1_MOUSE	Akcc1	Actin-binding protein 1 OS=Musculus GN:Akcc1 PE=2 SV=1	3	8.5	4.6	8.0	1.9	1.08	0.92	12.9	6.8	11.9	2.1	1.21	0.82
Q10813_CNK7_MOUSE	Cnkp7	COP1 signalosome complex subunit 7 OS=Musculus GN:Cnkp7 PE=1 SV=1	2	12.3	6.6	11.4	1.8	1.08	0.93	24.6	13.3	22.4	1.8	1.10	0.91
Q9CZH7_MKXK1_MOUSE	Mkxk1	Motif-remodeling associated protein 1 OS=Musculus GN:Mkxk1 PE=1 SV=1	1	8.8	4.8	8.2	1.8	1.08	0.93	8.8	4.8	8.0	1.8	1.10	0.91
Q505F1_MKCL1_MOUSE	Mkcl1	Mucin receptor subunit 2 group 1 OS=Musculus GN:Mkcl1 PE=1 SV=1	4	8.8	4.6	8.0	1.8	1.08	0.93	43.9	22.7	43.9	2.7	1.45	1.08
Q51177_PROX1_MOUSE	Prox2	Proxemin-2 OS=Musculus GN:Prox2 PE=1 SV=1	1	7.4	4.0	6.9	1.8	1.08	0.93	7.4	4.0	6.7	1.8	1.10	0.91
P15279_RAB6A_MOUSE	Rab6a	Ras-related protein Rab-6A OS=Musculus GN:Rab6a PE=1 SV=4	13	46.7	25.3	43.3	1.8	1.08	0.93	782.5	381.1	640.8	2.1	1.22	0.82
Q02059_UBA1_MOUSE	Uba1	Ubiquitin-like modifier-activating enzyme 1 OS=Musculus GN:Uba1 PE=1 SV=1	6	8.2	4.5	7.7	1.8	1.08	0.93	52.5	26.4	44.4	2.0	1.18	0.85
Q52186_SRD_MOUSE	Srd4	Translocin-associated protein subunit delta2 OS=Musculus GN:Srd4 PE=2 SV=1	2	15.4	8.2	14.3	1.8	1.08	0.95	15.4	8.2	14.3	1.8	1.10	0.91
Q8E18_SMG8_MOUSE	Smg8	Protein SMG8 OS=Musculus GN:Smg8 PE=2 SV=1	4	7.4	4.0	6.9	1.8	1.07	0.93	22.7	14.5	24.3	1.9	1.14	0.88
Q91011_MPH6_MOUSE	Mphosph6	M-phase phosphoprotein 6 OS=Musculus GN:Mphosph6 PE=1 SV=1	1	4.7	2.5	4.3	1.8	1.07	0.93	4.7	2.5	4.3	1.8	1.10	0.91
P59586_RANBP9_MOUSE	Ranbp9	Ran-binding protein 9 OS=Musculus GN:Ranbp9 PE=1 SV=1	3	10.2	5.6	9.5	1.8	1.07	0.93	26.3	14.0	23.5	1.9	1.12	0.89
Q53850_NUPK2_MOUSE	Nupk2	Nuclear pore glycoprotein k2 OS=Musculus GN:Nupk2 PE=1 SV=2	2	8.8	4.8	8.2	1.8	1.07	0.93	8.8	4.8	8.2	1.8	1.09	0.91
Q88874_CCNK_MOUSE	Ccnk	Cyclin-K OS=Musculus GN:Ccnk PE=1 SV=3	3	6.4	3.5	5.9	1.8	1.07	0.93	19.8	9.9	16.6	2.0	1.19	0.84
Q8BMM0_EMV_MOUSE	Emv1	Protein EMV1 OS=Musculus GN:Emv1 PE=1 SV=2	7	9.0	4.9	8.4	1.8	1.07	0.93	60.2	30.5	51.2	2.0	1.18	0.85
Q529P6_ITIK_MOUSE	Itik	Inhibitor of bruton tyrosine kinase OS=Musculus GN:Itik PE=1 SV=3	1	4.8	2.6	4.5	1.8	1.07	0.93	4.8	2.6	4.4	1.8	1.09	0.92
P63024_VAMP3_MOUSE	Vamp3	Vesicle-associated membrane protein 3 OS=Musculus GN:Vamp3 PE=1 SV=1	2	20.4	11.1	19.1	1.8	1.07	0.93	40.8	22.2	37.4	1.8	1.09	0.92
P97434_MPRIP_MOUSE	Mprip	Myosin phosphatase Rho-interacting protein OS=Musculus GN:Mprip PE=1 SV=2	37	19.4	10.6	18.1	1.8	1.07	0.93	1068.7	441.2	741.9	2.4	1.44	0.69
Q5Q299_QM_MOUSE	Qm	Protein quaking OS=Musculus GN:Qm PE=1 SV=1	9	16.3	8.9	15.6	1.8	1.07	0.93	243.8	241.3	405.7	1.0	0.60	1.66
Q10361_CK2_MOUSE	Ck2	Cyclin-dependent kinase 2 OS=Musculus GN:Ck2 PE=1 SV=1	2	18.8	10.2	17.4	1.8	1.07	0.94	295.0	154.2	329.6	1.8	1.08	0.93
Q17NCA_LC12_MOUSE	Luc12	Putative RNA-binding protein Luc1-like OS=Musculus GN:Luc12 PE=1 SV=1	25	27.4	14.9	25.2	1.8	1.07	0.94	2799.8	1239.1	2083.5	2.3	1.34	0.72
Q9C6B8_EP308_MOUSE	Nup153	Protein Nup153 OS=Musculus GN:Nup153 PE=1 SV=1	8	8.2	4.5	7.7	1.8	1.07	0.94	79.2	35.7	60.0	2.2	1.32	0.76
Q9VSL1_FABR8_MOUSE	Fambr8	Protein FAMBR8 OS=Musculus GN:Fambr8 PE=1 SV=1	1	6.7	3.8	6.7	1.8	1.07	0.94	188.6	80.1	134.2	2.4	1.40	0.71
Q9VSL1_F9S3_MOUSE	F9s3	Vamp/Vps39-like protein OS=Musculus GN:F9S3 PE=2 SV=1	1	6.9	3.8	6.5	1.7	1.07	0.94	3.8	2.0	6.4	1.8	1.09	0.92
Q88E6_CP20A_MOUSE	Cyp20a1	Cytochrome P450 20A1 OS=Musculus GN:Cyp20a1 PE=2 SV=1	3	5.8	3.2	5.4	1.8	1.07	0.94	17.2	9.5	15.9	1.8	1.08	0.93
Q8B5G_MAK16_MOUSE	Mak16	Protein MAK16 homolog OS=Musculus GN:MAK16 PE=1 SV=1	5	14.4	7.9	13.5	1.8	1.07	0.94	58.5	31.2	52.4	1.9	1.12	0.90
Q8B84_F8W8_MOUSE	F8w8	F-box-Wip1 repeat-containing protein 8 OS=Musculus GN:F8W8 PE=1 SV=2	4	10.8	6.0	10.3	1.8	1.07	0.94	13.2	6.8	11.2	1.8	1.08	0.93
Q8CF24_RCOR1_MOUSE	Rcor1	REST corepressor 1 OS=Musculus GN:Rcor1 PE=1 SV=1	6	7.3	4.0	6.8	1.8	1.07	0.94	63.8	37.1	62.3	1.7	1.02	0.98
Q8C704_RBM4_MOUSE	Rbm4	RNA-binding protein 4 OS=Musculus GN:Rbm4 PE=1 SV=1	2	22.5	12.3	21.1	1.8	1.06	0.94	45.0	24.6	41.4	1.8	1.09	0.92
Q70493_SNA12_MOUSE	Sna12	Sorting nexin-12 OS=Musculus GN:Sna12 PE=1 SV=1	3	30.6	16.7	27.8	1.8	1.06	0.94	172.0	90.8	152.6	1.9	1.13	0.89
Q8B7Y1_HAT1_MOUSE	Hat1	Homeobox protein HAT1 OS=Musculus GN:HAT1 PE=1 SV=1	3	10.9	6.0	10.6	1.8	1.06	0.94	14.0	12.0	18.8	1.8	1.08	0.93
P97367_MEI2_MOUSE	Mei2	Homeobox protein Mei2 OS=Musculus GN:Mei2 PE=1 SV=2	2	8.4	4.6	7.9	1.8	1.06	0.94	17.7	9.2	15.4	1.8	1.08	0.92
Q55VY1_CYTB_MOUSE	Spec1	Cytoskeleton 8 OS=Musculus GN:Spec1 PE=1 SV=2	16	14.4	7.9	13.5	1.8	1.06	0.94	260.1	138.1	232.3	1.9	1.12	0.89
Q81811_MUS1_MOUSE	Mus1	Protein mus1-complex subunit 1 OS=Musculus GN:MUS1 PE=2 SV=2	16	12.4	6.8	12.4	1.8	1.06	0.94	47.5	26.2	47.5	1.8	1.08	0.93
Q9P9V0_PRR8_MOUSE	Prrf8	Pre-mRNA-processing splicing factor 8 OS=Musculus GN:Prrf8 PE=1 SV=2	62	25.5	14.0	24.0	1.8	1.06	0.94	2040.6	1052.5	1769.7	1.9	1.16	0.86
Q9WTU0_PHF2_MOUSE	Phf2	Lysine-specific demethylase PHF2 OS=Musculus GN:Phf2 PE=1 SV=2	4	7.8	4.3	7.3	1.8	1.06	0.94	19.7	10.6	34.6	9.4	5.57	0.18
P81839_RAB18_MOUSE	Rab18	Ras-related protein Rab-18 OS=Musculus GN:Rab18 PE=1 SV=1	5	25.9	14.2	24.4	1.8	1.06	0.94	132.6	64.8	108.9	2.1	1.24	0.81
Q03147_C9K7_MOUSE	C9k7	Cyclin-dependent kinase 7 OS=Musculus GN:C9K7 PE=2 SV=2	6	10.4	5.7	9.8	1.8	1.06	0.94	35.8	19.4	35.8	1.8	1.08	0.93
P20029_F9F8_MOUSE	Hsp65	78 kDa glucose-regulated protein OS=Musculus GN:Hsp65 PE=1 SV=1	48	35.7	19.7	33.7	1.8	1.06	0.94	3005.4	2002.1	3366.5	1.5	0.89	1.12
Q55882_MYADM_MOUSE	Myadm	Myoid-associated differentiation marker OS=Musculus GN:Myadm PE=2 SV=2	4	53.7	29.6	50.8	1.8	1.06	0.95	239.0	126.7	213.1	1.9	1.12	0.89
P10306_TNKS1_MOUSE	Tnks1	Tnfr2 transcription factor TNKS1 OS=Musculus GN:Tnks1 PE=1 SV=1	1	10.4	5.6	9.9	1.8	1.06	0.95	10.4	5.6	9.9	1.8	1.08	0.93
Q51W50_CSDC1_MOUSE	Csdc1	CSDC domain-containing protein E1 OS=Musculus GN:CSDC1 PE=2 SV=1	2	9.7	5.3	9.1	1.8	1.06	0.95	19.3	10.7	17.9	1.8	1.08	0.93
Q5SUH6_GSUNH6_MOUSE	Clim1	Clathrin interactor 1 OS=Musculus GN:Clim1 PE=1 SV=1	9	13.2	7.3	12.5	1.8	1.06	0.95	141.5	79.9	134.4	1.8	1.05	0.95
Q9P810_TDF1_MOUSE	Dntt1p1	Deoxyribonucleic acid transferase terminal-interacting protein 1 OS=Musculus GN:Dntt1p1 PE=1 SV=1	2	11.4	6.3	10.8	1.8	1.06	0.95	22.9	12.6	21.3	1.8	1.08	0.93
Q108G52_LIMD1_MOUSE	Limd1	LIM domain-containing protein 1 OS=Musculus GN:LIMD1 PE=2 SV=1	2	16.3	8.9	15.9	1.8	1.06	0.95	35.5	19.3	35.5	1.8	1.08	0.93
Q9O198_SNF2_MOUSE	Snf2	Pre-mRNA-splicing factor SFY2 OS=Musculus GN:Snf2 PE=1 SV=1	2	14.0	7.8	13.3	1.8	1.05	0.95	28.0	15.5	26.1	1.8	1.08	0.93
Q10JF7_LYPL1_MOUSE	Lypl1a1	Lysophospholipase-like protein 1 OS=Musculus GN:Lypl1a1 PE=1 SV=3	1	3.4	1.9	3.2	1.8	1.05	0.95	3.4	1.9	3.1	1.8	1.08	0.93
Q50665_MPL6_MOUSE	Mpl6	Myosin light polypeptide 6 OS=Musculus GN:Mpl6 PE=1 SV=1	28	82.3	45.5	78.1	1.8	1.05	0.95	11198.2	6076.7	10217.9	1.8	1.10	0.91
Q9C4L1_MED1_MOUSE	Med1	Mediator of RNA polymerase II transcription subunit 1 OS=Musculus GN:Med1 PE=1 SV=1	4	7.8	4.1	7.0	1.8	1.05	0.95	14.7	8.1	14.7	1.8	1.07	0.94
Q8VDC5_MED17_MOUSE	Med17	Mediator of RNA polymerase II transcription subunit 17 OS=Musculus GN:Med17 PE=1 SV=1	3	12.8	7.1	12.2	1.8	1.05	0.95	39.2	19.7	33.2	2.0	1.18	0.85
Q9V9P1_VPS4A_MOUSE	Vps4a	Vacuolar protein sorting associated protein 4A OS=Musculus GN:Vps4a PE=1 SV=1	2	13.5	7.5	12.9	1.8	1.05	0.95	27.1	15.0	25.2	1.8	1.07	0.93
Q90B87_HPTFL1_MOUSE	Hptfl1	HPTFL phosphatase 1 regulatory subunit 12A OS=Musculus GN:HPTFL1 PE=1 SV=1	30	19.6	10.9	24.7	1.8	1.05	0.95	106.5	21.1	488.5	2.1	1.25	0.80
Q8CE14_Q8CE14_MOUSE	Ppp1r12a	Protein Ppp1r12a OS=Musculus GN:Ppp1r12a PE=2 SV=1	4	3.6	2.0	3.6	1.8	1.05	0.95	26.1	15.5	26.1	1.7	1.00	1.00
Q8W0Z2_VAC14_MOUSE	Vac14	Protein VAC14 homolog OS=Musculus GN:Vac14 PE=1 SV=1	3	8.4	4.6	8.0	1.8	1.05	0.95	70.7	24.6	41.4	2.9	1.71	0.59
Q8VDD5_MHY9_MOUSE	Mhy9	Myosin-9 OS=Musculus GN:Mhy9 PE=1 SV=4	187	63.9	35.5	60.8	1.8	1.05	0.95	28243.5	15673.8	26358.5	1.8	1.07	0.93
Q9WT05_AKAP2_MOUSE	Akap2	Kinase anchor protein 2 OS=Musculus GN:Akap2 PE=1 SV=1	6	11.1	6.0	10.5	1.8	1.05	0.95	11.1	6.0	10.5	1.7	1.04	0.96
Q9OCT8_CRI2_MOUSE	Crip2	Cysteine-rich protein 2 OS=Musculus GN:Crip2 PE=1 SV=1	2	9.2	5.1	8.8	1.8	1.05	0.95	18.4	10.2	17.2	1.8	1.07	0.94
Q9V9P1_UTP6_MOUSE	Utp6	U3 small nuclear RNA-associated protein 6 homolog OS=Musculus GN:Utp6 PE=2 SV=1	2	4.6	2.6	4.4	1.8	1.05	0.95	3.3	5.2	4.7	1.8	1.07	0.94
Q9BCC9_TF3A4_MOUSE	Tf3a4	Tripartite motif-containing protein 3A OS=Musculus GN:Tf3a4 PE=2 SV=2	1	6.2	3.5	5.9	1.8	1.05	0.95	6.2	3.5	5.8	1.8	1.07	

Q8010_Q8010_MOUSE	Rhoa	Transforming protein RhoA:OS-Musculus:GN-RhoA:PE1-SV1	2	8.0	4.6	7.9	1.7	1.01	0.99	16.0	9.3	15.6	1.7	1.03	0.97
Q52362_Q52362_MOUSE	Spz2h	DNM3-2-related protein polymerase 1, II, and III subunit:PB3D3:OS-Musculus:GN-Polr2h:PE1-SV1	5	16.2	9.4	16.1	1.7	1.01	0.99	80.6	45.4	76.3	1.8	1.06	0.95
PE2414_PE2414_MOUSE	Phf2	Serine/threonine protein phosphatase:PE1:beta:cat:subunit:OS-Musculus:GN-Phf2:PE1-SV1	2	8.6	4.6	8.6	1.7	1.00	1.00	68.6	38.6	115.3	1.8	1.09	0.91
A04087W04_Q40087W04_1	10A07	Protein Sctap:OS-Musculus:GN-Sctap:PE1-SV1	10	10.6	6.2	10.6	1.7	1.00	1.00	111.5	69.6	117.0	1.6	0.95	1.05
O9JX1_G0R2_MOUSE	Gorazp2	Glycyl assembly-stacking protein 2:OS-Musculus:GN-Gorazp2:PE1-SV1	2	7.5	4.3	7.4	1.7	1.00	1.00	15.0	8.7	14.6	1.7	1.02	0.98
Q52002_Q52002_MOUSE	Cy5c1	Cytochrome c5 type I:beta:Musculus:GN-Cy5c1:PE1-SV1	1	7.4	4.3	7.4	1.7	1.00	1.00	21.2	15.6	24.1	1.7	1.02	0.98
Q901W7_Q901W7_MOUSE	Fkbp11	Peptidyl-prolyl cis-trans isomerase FKBP11:OS-Musculus:GN-Fkbp11:PE1-SV1	2	10.6	6.2	10.6	1.7	1.00	1.00	12.2	12.3	10.7	1.7	1.02	0.98
Q62Q33_Q62Q33_MOUSE	Mlec	Mallein:OS-Musculus:GN-Mlec:PE2-SV2	4	11.1	6.4	11.1	1.7	1.00	1.00	54.5	29.5	49.5	1.8	1.10	0.91
Q52003_Q52003_MOUSE	CommD4	COMM domain-containing protein 4:OS-Musculus:GN-CommD4:PE2-SV1	1	8.1	4.7	8.1	1.7	1.00	1.00	81.2	47	8.0	1.7	1.02	0.98
Q6L103_Q6L103_MOUSE	Trio	Triple function domain protein 1:OS-Musculus:GN-Trio:PE1-SV1	4	8.0	4.7	8.0	1.7	1.00	1.00	17.5	17.9	11.1	1.8	1.01	1.00
Q90100_Q90100_MOUSE	Khrp9	Far upstream element-binding protein 2:OS-Musculus:GN-Khrp9:PE1-SV1	11	16.5	9.6	16.5	1.7	1.00	1.00	226.5	137.8	231.7	1.6	0.98	1.02
Q90000_Q90000_MOUSE	Sra9	Serine/arginine-rich signaling factor 9:OS-Musculus:GN-Sra9:PE1-SV1	18	52.0	30.3	51.9	1.7	1.00	1.00	1185.0	725.2	1219.3	1.6	0.97	1.03
PE1021_Q6486_MOUSE	Rar	Rar-related protein Rar:beta:OS-Musculus:GN-Rar:PE1-SV1	3	8.8	5.1	8.7	1.7	1.00	1.00	13.7	13.7	23.1	1.9	0.99	0.72
P70255_P70255_MOUSE	Nlc	Nuclear factor 1 C-type:OS-Musculus:GN-Nlc:PE1-SV1	14	14.6	8.5	14.6	1.7	1.00	1.00	376.1	258.3	434.3	1.5	0.87	1.15
EQ90N1_EQ90N1_MOUSE	Dhv9	ATP-dependent RNA helicase A:OS-Musculus:GN-Dhv9:PE1-SV1	102	109.8	64.1	109.9	1.7	1.00	1.00	15243.4	8663.5	14567.5	1.8	0.95	0.96
O54984_O54984_MOUSE	Ana1	ATPase Ana1:OS-Musculus:GN-Ana1:PE1-SV2	3	22.8	13.3	22.8	1.7	1.00	1.00	99.3	42.3	71.0	2.3	1.40	0.72
Q10252_Q10252_MOUSE	Hmsa2	Hemesevenase 2:OS-Musculus:GN-Hmsa2:PE2-SV1	3	7.6	4.8	7.6	1.7	1.00	1.00	120.7	120.2	20.2	1.4	1.02	0.70
Q9C7W9_Q9C7W9_MOUSE	Golga1	Golgi subfamily A member 1:OS-Musculus:GN-Golga1:PE1-SV2	4	9.2	5.4	9.2	1.7	1.00	1.00	88.8	34.6	58.2	1.1	0.67	1.50
Q904H9_Q904H9_MOUSE	PHF4	PHD finger protein 14:OS-Musculus:GN-PHF4:PE1-SV1	5	15.3	8.9	15.3	1.7	1.00	1.00	93.8	54.7	92.0	1.7	1.02	0.98
EQ3D12_EQ3D12_MOUSE	Sympo	Synaptotagmin:OS-Musculus:GN-Sympo:PE1-SV1	13	11.7	6.9	11.8	1.7	1.00	1.00	174.7	100.2	168.4	1.7	1.04	0.96
Q99172_Q99172_MOUSE	Gcm1	Glutamine-cysteine ligase regulatory subunit:OS-Musculus:GN-Gcm1:PE2-SV1	1	2.5	2.1	2.5	1.7	1.00	1.00	3.2	2.5	2.4	1.7	1.02	0.98
Q85222_Q85222_MOUSE	Endod1	Endonuclease domain-containing protein 1:OS-Musculus:GN-Endod1:PE1-SV2	1	5.3	3.1	5.3	1.7	1.00	1.00	3.5	3.1	5.2	1.7	1.02	0.98
F80V29_F80V29_MOUSE	Igga3	Protein Igga3:OS-Musculus:GN-Igga3:PE4-SV1	3	10.2	5.9	10.2	1.7	1.00	1.00	35.0	31.0	52.2	1.1	0.67	1.49
Q50155_Q50155_MOUSE	Lncr57	Leucine-rich repeat-containing protein 57:OS-Musculus:GN-Lncr57:PE2-SV1	2	4.7	2.8	4.7	1.7	1.00	1.00	9.4	5.5	9.3	1.7	1.02	0.98
Q9DCA5_Q9DCA5_MOUSE	Brix1	Ribosome biogenesis protein BRX1 homolog:OS-Musculus:GN-Brix1:PE2-SV3	6	19.8	11.6	19.8	1.7	1.00	1.00	123.3	74.4	125.1	1.7	0.99	1.02
Q9Y91_Q9Y91_MOUSE	Hadhb	Tripartite oligomer protein beta, mitochondrial:OS-Musculus:GN-Hadhb:PE1-SV1	6	22.9	13.4	22.9	1.7	1.00	1.00	142.7	85.6	143.9	1.7	0.99	1.01
P41819_P41819_MOUSE	Epb41	Protein 4.1:OS-Musculus:GN-Epb41:PE1-SV2	1	7.4	4.8	7.4	1.7	1.00	1.01	8.2	8.2	8.1	1.7	1.01	0.99
Q9XW82_Q9XW82_MOUSE	G7F	G7F protein-activating protein:OS-Musculus:GN-G7F:PE1-SV2	1	8.2	4.3	8.2	1.7	1.00	1.01	12.0	12.0	12.0	1.7	1.01	0.99
P14873_P14873_MOUSE	Map1b	Microtubule-associated protein 1B:OS-Musculus:GN-Map1b:PE1-SV2	45	14.2	8.3	14.2	1.7	0.99	1.01	778.6	428.6	720.7	1.8	0.98	0.93
Q9D70_Q9D70_MOUSE	NOSIP	Nitric oxide synthase interacting protein:OS-Musculus:GN-NOSIP:PE2-SV1	2	14.2	8.3	14.2	1.7	0.99	1.01	28.3	16.6	28.0	1.7	1.01	0.99
Q9P5D8_Q9P5D8_MOUSE	Shc4	Cytoplasmic maintenance of chromosomes fragile hinge domain-containing protein 1:OS-Musculus:GN-Shc4:PE1-SV1	2	14.3	7.8	14.3	1.7	0.99	1.01	189.4	106.7	179.4	1.8	1.06	0.95
Q9V823_Q9V823_MOUSE	Ctlnp1	Ctlnp1 and subunit transmembrane protein 1 homolog:OS-Musculus:GN-Ctlnp1:PE1-SV2	2	5.7	3.4	5.7	1.7	0.99	1.01	6.7	6.7	11.1	1.7	1.01	0.95
Q9P566_Q9P566_MOUSE	Ubn7	UBX domain-containing protein 7:OS-Musculus:GN-Ubn7:PE1-SV2	2	8.8	5.1	8.8	1.7	0.99	1.01	17.5	10.3	17.3	1.7	1.01	0.99
A21772_A21772_MOUSE	Fubp3	MCG130458:OS-Musculus:GN-Fubp3:PE4-SV1	22	19.7	11.6	19.8	1.7	0.99	1.01	622.8	271.9	457.3	2.3	1.36	0.73
Q52119_Q52119_MOUSE	Exoc4	Exosome complex component EXOC4:OS-Musculus:GN-Exoc4:PE2-SV3	1	22.2	13.1	22.2	1.7	0.99	1.01	23.3	12.3	23.3	1.8	1.11	0.84
Q6P404_Q6P404_MOUSE	Rexo4	RNA exonuclease 4:OS-Musculus:GN-Rexo4:PE2-SV2	4	12.6	7.4	12.7	1.7	0.99	1.01	50.9	30.3	50.9	1.7	1.00	1.00
Q9ER00_Q9ER00_MOUSE	Stx12	STX12:OS-Musculus:GN-Stx12:PE1-SV1	5	15.0	8.8	15.1	1.7	0.99	1.01	111.1	60.0	100.9	1.9	1.10	0.91
Q92208_Q92208_MOUSE	Lncr59	Leucine-rich repeat-containing protein 59:OS-Musculus:GN-Lncr59:PE1-SV1	5	11.2	6.6	11.2	1.7	0.99	1.01	64.0	41.0	70.0	1.5	0.91	1.09
Q91420_Q91420_MOUSE	Setd1	GMP-activated transcription factor 1:OS-Musculus:GN-Setd1:PE1-SV2	2	8.5	5.0	8.6	1.7	0.99	1.01	3.0	2.9	4.1	1.7	1.01	0.99
Q98974_Q98974_MOUSE	Setd1b	Histone-N-lysine methyltransferase SETD1B:OS-Musculus:GN-Setd1b:PE1-SV1	2	8.5	5.0	8.6	1.7	0.99	1.01	17.1	10.1	16.9	1.7	1.01	0.99
P97312_P97312_MOUSE	Carp1	Cysteine and glycine-rich protein 1:OS-Musculus:GN-Carp1:PE1-SV1	2	21.9	12.9	22.2	1.7	0.99	1.01	43.8	25.9	43.5	1.7	1.01	0.99
Q6C7V9_Q6C7V9_MOUSE	Utp15	Utp15:OS-Musculus:GN-Utp15:PE1-SV1	12	11.5	6.5	11.5	1.7	0.99	1.01	49.6	28.5	49.6	1.7	1.01	0.99
EQ924_Q924_MOUSE	Elna1	Protein Elna1:OS-Musculus:GN-Elna1:PE1-SV1	1	7.8	4.6	7.8	1.7	0.99	1.01	7.8	4.6	7.8	1.7	1.01	0.99
PE1022_Q9101_MOUSE	Chp1	Calcineurin B homologous protein 1:OS-Musculus:GN-Chp1:PE1-SV2	5	18.8	11.0	19.0	1.7	0.99	1.01	75.9	52.7	88.7	1.4	0.86	1.17
Q8R0F5_Q8R0F5_MOUSE	Rbm2	RNA-binding motif protein, X-linked 2:OS-Musculus:GN-Rbm2:PE1-SV1	4	31.5	18.6	31.9	1.7	0.99	1.01	127.9	78.9	132.7	1.6	0.96	1.04
Q2V0V2_Q2V0V2_MOUSE	Hm2	Protein motif homolog:OS-Musculus:GN-Hm2:PE1-SV1	12	9.9	6.5	9.9	1.7	0.99	1.01	89.9	131.6	151.2	1.5	0.98	1.14
Q91215_Q91215_MOUSE	Baz2a	Bromodomain adjacent to zinc finger domain protein 2A:OS-Musculus:GN-Baz2a:PE1-SV2	6	7.6	4.5	7.7	1.7	0.99	1.02	48.1	30.4	51.0	1.6	0.94	1.06
Q91217_Q91217_MOUSE	Sic7a5	Large neutral amino acid transporter small subunit 1:OS-Musculus:GN-Sic7a5:PE1-SV2	8	9.0	5.3	9.1	1.7	0.98	1.02	87.6	45.5	76.4	1.9	1.15	0.87
P18855_P18855_MOUSE	Pf55	Protein P55:OS-Musculus:GN-Pf55:PE2-SV1	6	11.2	6.6	11.2	1.7	0.98	1.02	11.6	11.6	11.0	1.7	1.00	1.00
P97452_P97452_MOUSE	Bop1	Ribosome biogenesis protein BOP1:OS-Musculus:GN-Bop1:PE1-SV1	1	5.2	3.1	5.2	1.7	0.98	1.02	3.1	3.1	5.2	1.7	1.00	1.00
Q91901_Q91901_MOUSE	H2-k1	H-2 class I histocompatibility antigen, K-B alpha chain:OS-Musculus:GN-H2-K1:PE1-SV1	10	14.0	8.3	14.3	1.7	0.98	1.02	201.3	67.2	1138.6	0.3	1.18	5.66
EP9W99_EP9W99_MOUSE	Protein Rdt1	OS-Musculus:GN-ProRdt1:PE1-SV1	14	13.7	8.1	14.0	1.7	0.98	1.02	190.7	103.6	174.1	1.8	1.10	0.91
Q9R0B7_Q9R0B7_MOUSE	Ran	Zinc finger protein 34B:OS-Musculus:GN-Ran:PE1-SV1	9	9.8	5.8	10.0	1.7	0.98	1.02	11.6	11.6	11.6	1.7	1.00	1.00
Q9V36_Q9V36_MOUSE	Pan1	Paxillin:OS-Musculus:GN-Pan1:PE1-SV1	1	13.4	7.9	13.6	1.7	0.98	1.02	13.4	7.9	13.4	1.7	1.00	1.00
P55122_P55122_MOUSE	Sic12a2	Solute carrier family 12 member 2:OS-Musculus:GN-Sic12a2:PE1-SV1	1	5.7	3.4	5.8	1.7	0.98	1.02	5.7	3.4	5.6	1.7	1.00	1.00
P49717_P49717_MOUSE	Mcm4	DNA replication licensing factor MCM4:OS-Musculus:GN-Mcm4:PE1-SV1	15	14.9	8.9	15.2	1.7	0.98	1.02	267.4	218.5	367.4	1.2	0.72	3.39
Q9L912_Q9L912_MOUSE	Ubr1	Transcription initiation factor eIF-2B:OS-Musculus:GN-Ubr1:PE2-SV1	3	19.7	11.7	19.7	1.7	0.97	1.03	48.5	28.1	48.5	1.8	0.92	1.04
Q9CKG3_Q9CKG3_MOUSE	Ppil4	Peptidyl-prolyl cis-trans isomerase-like 4:OS-Musculus:GN-Ppil4:PE1-SV1	3	7.2	4.3	7.3	1.7	0.98	1.02	30.1	17.0	28.6	1.8	1.05	0.98
C1e1	Exp2	Exp2:OS-Musculus:GN-C1e1:PE1-SV1	4	10.4	6.2	10.4	1.7	0.98	1.02	43.5	22.7	38.2	1.9	1.14	0.88
P10852_P10852_MOUSE	Utr2	472 cell surface antigen:Utr2:OS-Musculus:GN-Utr2:PE1-SV1	7	20.2	12.4	20.2	1.7	0.98	1.02	183.1	109.6	307.8	1.6	0.92	0.95
PC00A1_PC00A1_MOUSE	Chmp6	Charged multivesicular body protein 6:OS-Musculus:GN-Chmp6:PE2-SV2	1	9.9	5.9	10.1	1.7	0.98	1.02	9.9	5.9	9.9	1.7	1.00	1.00
P84089_P84089_MOUSE	Erh	Enhancer of rudimentary homolog:OS-Musculus:GN-Erh:PE1-SV1	3	36.1	21.5	36.9	1.7	0.98	1.02	377.7	202.8	341.1	1.9	1.11	0.90
EQ9ER2_Q99ER2_MOUSE	Strn3	Strn3:OS-Musculus:GN-Strn3:PE1-SV1	1	9.3	5.6	9.5	1.7	0.98	1.02	3.0	2.6	3.0	1.7	1.00	1.00
Q922V5_Q922V5_MOUSE	Pyx1	Pyx1:OS-Musculus:GN-Pyx1:PE1-SV1	2	11.9	6.8	12.0	1.7	0.98	1.02	13.1	12.1	13.1	1.7	1.00	1.00
G3A303_G3A303_MOUSE	Nemc4e4	MCG1618, isoform CRA_c:OS-Musculus:GN-Nemc4e4:PE4-SV1	2	8.5	5.1	8.7	1.7	0.98	1.02	17.0	10.1	17.0	1.7	1.00	1.00
P45856_P45856_MOUSE	Pcy1a1	Choline phosphatidyltransferase A:OS-Musculus:GN-Pcy1a1:PE1-SV1	3	11.4	6.8	11.6</									

O8K1V1_ADRM1_MOUSE	Adrm1	Proteasome ubiquitin receptor ADRM1 OS-Mus musculus GN-Adrm1 PE1 SV-2	1	5.3	3.8	6.5	1.4	0.83	1.21	5.3	3.8	6.3	1.4	0.84	1.19
O874W6_TOML_MOUSE	Tom1	Target of Myosin 1 OS-Mus musculus GN-Tom1 PE1 SV-1	6	8.1	5.8	9.9	1.4	0.83	1.21	80.2	36.3	63.0	1.7	0.99	1.10
P51455_GPT4_MOUSE	Gpt4	GTPase-activating protein 4 OS-Mus musculus GN-Gpt4 PE2 SV-2	1	6.4	5.7	16.7	1.4	0.82	1.21	165.3	24.8	277.9	1.6	0.87	1.18
O8CQ79_TNND9_MOUSE	Tnnd9	Threonine domain-containing protein 9 OS-Mus musculus GN-Tnnd9 PE2 SV-1	1	18.5	13.1	22.5	1.4	0.82	1.21	18.5	13.1	22.0	1.4	0.84	1.19
O61879_MHYH10_MOUSE	Mhy10	Myosin-10 OS-Mus musculus GN-Mhy10 PE1 SV-2	179	70.7	50.1	85.9	1.4	0.82	1.21	27248.5	18183.8	30675.7	1.5	0.89	1.12
O8B0C0_CPS1A_MOUSE	Cps1a	Capsulin OS-Mus musculus GN-Cps1a PE1 SV-1	1	6.4	4.5	7.7	1.4	0.81	1.21	6.4	4.5	7.7	1.4	0.81	1.19
Q10M00_GKAP1_MOUSE	Gkap1	G kinase-anchoring protein 1 OS-Mus musculus GN-Gkap1 PE1 SV-1	1	3.7	2.7	4.6	1.4	0.82	1.22	3.7	2.7	4.5	1.4	0.84	1.19
P13020_GELS_MOUSE	Gels	Gelsolin OS-Mus musculus GN-Gels PE1 SV-3	28	59.2	42.0	72.1	1.4	0.82	1.22	3036.2	2043.0	3435.2	1.5	0.88	1.13
O62282_SVDC_MOUSE	Dars	Apoptate-RNAI Igase, cytoplasmic OS-Mus musculus GN-Dars PE2 SV-2	3	5.3	3.8	6.5	1.4	0.82	1.22	117.5	11.7	19.7	1.5	0.89	1.13
P70377_LAVL_MOUSE	Gln1	GLAV-like protein 1 OS-Mus musculus GN-Gln1 PE1 SV-2	1	2.9	2.1	3.6	1.4	0.82	1.23	2.9	2.1	3.5	1.4	0.83	1.08
P28656_NP11_MOUSE	Nap11	Nucleosome assembly protein 1-like 1 OS-Mus musculus GN-Nap11 PE1 SV-2	1	14.2	10.1	17.3	1.4	0.82	1.22	14.2	10.1	17.0	1.4	0.84	1.20
O5CYN2_SPC23_MOUSE	Spc23	Spc23 peptidase complex subunit 2 OS-Mus musculus GN-Spc23 PE1 SV-1	1	7.7	5.4	9.3	1.4	0.82	1.22	7.7	5.4	9.2	1.4	0.84	1.20
O8C6Y9_AKOS_MOUSE	Rbp89	Actin regulator of STRIP1 OS-Mus musculus GN-Rbp89 PE1 SV-1	6	6.6	6.3	11.8	1.4	0.82	1.22	66.3	39.1	65.8	1.8	1.05	0.85
O8CQ02_MET16_MOUSE	Met16	Methyltransferase-like protein 16 OS-Mus musculus GN-Met16 PE2 SV-1	1	4.2	3.0	4.4	1.4	0.82	1.22	4.2	3.0	4.1	1.4	0.83	1.20
O900M7_CAF1B_MOUSE	Chaf1b	Chromatin assembly factor 1 subunit B OS-Mus musculus GN-Chaf1b PE2 SV-1	4	8.3	5.9	10.1	1.4	0.82	1.22	4.8	3.8	25.6	4.3	0.81	1.24
O90B53_AFRG3_MOUSE	Afrg3p	ADP-ribosylation factor GTPase-activating protein 3 OS-Mus musculus GN-Afrg3p PE1 SV-1	11	7.8	5.6	9.5	1.4	0.82	1.22	111.9	62.7	105.5	1.9	1.12	0.89
O90L20_O90L20_MOUSE	GstP9	Glutathione S-transferase theta 3 OS-Mus musculus GN-GstP9 PE2 SV-1	1	2.9	2.1	3.6	1.4	0.82	1.23	2.9	2.1	3.5	1.4	0.83	1.20
O8BVE3_VATH_MOUSE	Atpev1h	V-type proton ATPase subunit 1 OS-Mus musculus GN-Atpev1h PE1 SV-1	3	4.4	3.1	5.4	1.4	0.82	1.23	15.7	8.3	13.9	1.9	1.13	0.89
O64339_ISG15_MOUSE	Ig15	Ubiquitin-like protein ISG15 OS-Mus musculus GN-Ig15 PE1 SV-4	14	54.5	39.0	66.9	1.4	0.82	1.23	1029.3	689.8	1159.8	1.5	0.89	1.13
O6P8U0_H11L_MOUSE	Hif1p1	Fibronin A-interacting protein 3-like OS-Mus musculus GN-Hif1p1 PE1 SV-2	6	9.1	6.5	11.2	1.4	0.81	1.23	64.9	48.0	80.7	1.4	0.80	1.24
O8C5U7_BRMK_MOUSE	Rbm24	RNA-binding protein 24 OS-Mus musculus GN-Rbm24 PE1 SV-1	4	15.2	10.9	18.7	1.4	0.81	1.23	45.7	31.6	78.9	1.9	0.67	1.49
O921M1_SF3B3_MOUSE	Sf3b3	Dystroglycan 3 subunit 3 OS-Mus musculus GN-Sf3b3 PE2 SV-1	60	38.7	27.8	47.6	1.4	0.81	1.23	3060.0	2236.7	3760.9	1.4	0.81	1.23
O62166_DAG1_MOUSE	Dag1	Dystroglycan OS-Mus musculus GN-Dag1 PE1 SV-4	3	6.3	4.6	7.8	1.4	0.81	1.23	33.9	12.6	21.1	2.7	0.61	0.62
P97376_FG1_MOUSE	Fg1	Protein FG1 OS-Mus musculus GN-Fg1 PE1 SV-2	1	14.6	10.5	17.9	1.4	0.81	1.23	14.6	10.5	17.6	1.4	0.83	1.21
O90D01_HNRP4_MOUSE	Hnrrp4	Heterogeneous nuclear ribonucleoprotein M OS-Mus musculus GN-Hnrrp4 PE1 SV-3	105	36.4	26.1	44.8	1.4	0.81	1.23	12287.0	8236.6	13849.7	1.5	0.89	1.13
O5U409_THOC6_MOUSE	Thoc6	THO complex subunit 6 homolog OS-Mus musculus GN-Thoc6 PE2 SV-1	3	10.7	7.7	13.2	1.4	0.81	1.23	45.4	22.4	37.7	2.0	1.20	0.83
P62204_CALM_MOUSE	Calm1	Calmodulin OS-Mus musculus GN-Calm1 PE1 SV-2	8	65.6	47.1	80.8	1.4	0.81	1.23	1736.7	1079.8	1815.7	1.6	0.96	1.05
O4NVL9_RRP12_MOUSE	Rrp12	RNA-binding protein 12 OS-Mus musculus GN-Rrp12 PE1 SV-2	4	6.5	4.6	8.1	1.4	0.81	1.23	42.3	30.8	47.9	1.6	0.87	1.16
O8QK78_PAF1_MOUSE	Pafl	RNA polymerase II-associated factor 1 homolog OS-Mus musculus GN-Pafl PE2 SV-1	5	6.7	4.8	8.3	1.4	0.81	1.23	33.9	23.9	40.2	1.4	0.84	1.18
O80203_BUD11_MOUSE	Bud11	Protein BUD11 homolog OS-Mus musculus GN-Bud11 PE4 SV-1	5	13.3	9.6	16.5	1.4	0.81	1.24	129.8	67.1	112.8	1.2	0.74	1.36
O8LDV9_WP110_MOUSE	Wp110	SH2 domain nuclear ribonucleoprotein protein RBP110 OS-Mus musculus GN-Wp110 PE2 SV-1	103	11.9	8.6	14.7	1.4	0.81	1.24	172.3	124.8	172.3	1.2	0.84	1.21
O80239_CETNC_MOUSE	Cetnc	Centrin-3 OS-Mus musculus GN-Cetnc PE4 SV-1	3	4.9	3.6	6.1	1.4	0.81	1.24	15.3	9.0	15.2	1.7	1.01	0.99
O81U22_TXIP_MOUSE	Sic25a1	Tricarbonylate transport protein, mitochondrial OS-Mus musculus GN-Sic25a1 PE2 SV-1	1	6.4	4.6	7.9	1.4	0.81	1.24	6.4	4.6	7.7	1.4	0.82	1.22
O31T83_GPTCA_MOUSE	Gpatac4	Gpatac4 domain-containing protein 4 OS-Mus musculus GN-Gpatac4 PE1 SV-1	1	9.0	6.5	11.2	1.4	0.81	1.24	9.0	6.5	11.0	1.4	0.82	1.22
O90113_SAS10_MOUSE	Sas10	Somebody about silencing protein 10 OS-Mus musculus GN-Sas10 PE2 SV-1	1	14.4	10.4	17.9	1.4	0.81	1.24	14.4	10.4	17.7	1.4	0.82	1.22
O9PM46_O9PM46_MOUSE	Polr2c	DNA-directed RNA polymerase I subunit RP83 OS-Mus musculus GN-Polr2c PE1 SV-1	6	24.1	17.5	30.0	1.4	0.81	1.24	197.4	168.3	283.0	1.2	0.70	1.43
PE2305_RUXE_MOUSE	Ruxe	Small nuclear ribonucleoprotein E OS-Mus musculus GN-Ruxe PE2 SV-1	10	21.7	15.7	26.9	1.4	0.80	1.24	850.3	459.0	771.9	1.9	1.10	0.91
O92277_BAZ1L_MOUSE	Baz1b	Tyrosine-protein kinase BAZ1B OS-Mus musculus GN-Baz1b PE1 SV-2	79	19.8	14.4	24.6	1.4	0.80	1.24	2860.7	1712.5	2879.5	1.7	0.99	1.01
P13864_DNMT1_MOUSE	Dnmt1	DNA methyltransferase 1 OS-Mus musculus GN-Dnmt1 PE1 SV-5	32	26.6	19.3	33.1	1.4	0.80	1.24	1119.9	909.1	1529.8	1.2	0.87	1.12
O9NH12_PARD3_MOUSE	Pard3	Partitioning defective 3 homolog OS-Mus musculus GN-Pard3 PE1 SV-2	2	3.7	2.7	4.6	1.4	0.80	1.24	7.4	5.4	9.1	1.4	0.82	1.22
O9CY57_CHTOP_MOUSE	Chtop	Chromatin target of PRMT1 protein OS-Mus musculus GN-Chtop PE1 SV-2	8	10.5	7.7	13.2	1.4	0.80	1.25	157.4	95.5	160.6	1.6	0.98	1.02
O8B9V3_VATH_MOUSE	GstP9	Glutathione S-transferase theta 3 OS-Mus musculus GN-GstP9 PE2 SV-1	1	2.9	2.1	3.6	1.4	0.81	1.25	2.9	2.1	3.6	1.4	0.83	1.08
O8CQ29_POD21_MOUSE	Pod21	POD domain-containing protein 11 OS-Mus musculus GN-Pod21 PE1 SV-1	2	6.9	5.0	8.7	1.4	0.80	1.25	13.9	10.1	17.0	1.4	0.82	1.22
O8R3C0_MCMBP_MOUSE	Mcmcbp	Mini-chromosome maintenance complex-binding protein OS-Mus musculus GN-Mcmcbp PE2 PE1	1	1.7	1.2	2.1	1.4	0.80	1.25	1.7	1.2	2.1	1.4	0.82	1.23
B2R0C6_B2R0C6_MOUSE	Polr2a	Polychrome A (RNA III) DNA-directed polypeptide II OS-Mus musculus GN-Polr2a PE2 SV-1	1	4.4	3.2	5.5	1.4	0.80	1.25	4.4	3.2	5.4	1.4	0.82	1.23
P11522_OAS1_MOUSE	Oas1	2',5'-oligoadenylate synthetase 1 OS-Mus musculus GN-Oas1 PE2 SV-2	1	4.4	3.2	5.5	1.4	0.80	1.25	4.4	3.2	5.4	1.4	0.82	1.23
O3QK00_SPAT5_MOUSE	Spat5a	Spermatogenesis-associated protein 5 OS-Mus musculus GN-Spat5a PE1 SV-2	6	11.1	8.1	13.9	1.4	0.80	1.25	71.7	43.8	73.7	1.6	0.97	1.03
PE6254_I4337_MOUSE	Whaqa	14-3-3 protein theta OS-Mus musculus GN-Whaqa PE1 SV-1	4	12.8	9.4	16.1	1.4	0.80	1.25	70.3	41.6	69.9	1.7	1.01	0.99
O85859_MNKL_MOUSE	Mnkl	MLN-1 OS-Mus musculus GN-Mnkl PE1 SV-2	5	14.0	9.7	16.6	1.4	0.80	1.25	52.0	3266.9	593.1	1.6	0.66	1.04
P98787_COKL_MOUSE	Low1	Lysyl oxidase homolog 1 OS-Mus musculus GN-Low1 PE2 SV-3	1	12.0	8.8	15.1	1.4	0.80	1.25	8.8	12.0	14.8	1.4	0.81	1.23
O89299_ZNF512_MOUSE	Znf512	Zinc finger protein 512 OS-Mus musculus GN-Znf512 PE2 SV-2	6	12.6	9.2	15.8	1.4	0.80	1.26	69.9	50.9	85.6	1.4	0.82	1.22
P53399_O0B2_MOUSE	Upp1	Liponamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase complex 3 OS-Mus musculus GN-Upp1 PE1 SV-1	3	15.8	11.5	19.8	1.4	0.80	1.26	57.9	41.8	70.3	1.4	0.82	1.21
O85559_MNKL_MOUSE	Mnkl	MLN-1 OS-Mus musculus GN-Mnkl PE1 SV-2	5	14.0	9.7	16.6	1.4	0.80	1.26	52.0	3266.9	593.1	1.6	0.66	1.04
O91YW3_DNaj3_MOUSE	Dnaj3	DNA homolog subfamily 3 member C OS-Mus musculus GN-Dnaj3 PE1 SV-1	3	5.9	4.3	7.4	1.4	0.80	1.26	27.9	22.8	38.3	1.2	0.73	1.37
O90319_PICAL_MOUSE	Picalm	Phosphatidylinositol-binding clathrin assembly protein OS-Mus musculus GN-Picalm PE1 SV-1	18	12.5	9.1	15.7	1.4	0.80	1.26	283.3	255.6	429.8	1.1	0.66	1.52
O5CNC3_SHW1_MOUSE	Shw1	SHW domain-containing protein 4 OS-Mus musculus GN-Shw1 PE1 SV-1	18	25.2	18.5	31.6	1.4	0.79	1.26	493.3	367.4	617.7	1.3	0.80	1.25
O9CY72_TTRC_MOUSE	Ttrc	Transcription factor 1 OS-Mus musculus GN-Ttrc PE1 SV-1	1	9.6	7.1	12.7	1.4	0.79	1.26	9.6	7.1	11.9	1.4	0.81	1.24
O91X01_ENY2_MOUSE	Eny2	Transcription and mRNA export factor ENY2 OS-Mus musculus GN-Eny2 PE2 SV-1	1	14.8	10.9	18.7	1.4	0.79	1.27	14.8	10.9	18.4	1.4	0.80	1.24
O91Y11_NEN1_MOUSE	Nen1	Neilin OS-Mus musculus GN-Nen1 PE2 SV-3	4	13.3	9.8	16.8	1.4	0.79	1.27	15.2	38.2	64.2	1.4	0.86	1.16
O91Y11_AKAP9_MOUSE	Akap9	4-kilobase anchor protein OS-Mus musculus GN-Akap9 PE2 SV-1	2	4.8	3.5	5.8	1.4	0.79	1.27	27.9	17.0	28.5	1.6	0.88	1.01
O7TM72_WDR91_MOUSE	Wdr91	WD repeat-containing protein 91 OS-Mus musculus GN-Wdr91 PE1 SV-1	1	5.7	4.2	7.3	1.4	0.79	1.27	5.7	4.2	7.2	1.4	0.80	1.25
P11352_GPX1_MOUSE	Gpx1	Glutathione peroxidase 1 OS-Mus musculus GN-Gpx1 PE1 SV-2	1	4.2	3.1	5.3	1.4	0.79	1.27	4.2	3.1	5.2	1.3	0.80	1.25
O62399_TP052_MOUSE	Tp052	Tumor protein D52 OS-Mus musculus GN-Tp052 PE2 SV-2	1	11.2	8.3	14.3	1.3	0.79	1.27	11.2	8.3	14.0	1.3	0.80	1.25
O91059_PT02_MOUSE	Pt02	Protein tyrosine phosphatase 2 OS-Mus musculus GN-Pt02 PE2 SV-2	1	11.2	8.3	14.3	1.3	0.79	1.27	11.2	8.3	14.0	1.3	0.80	1.25
O8C156_CN2D_MOUSE	Cn2ph	Condensin complex subunit 2 OS-Mus musculus GN-Cn2ph PE2 SV-1	21	14.1	10.5	17.9	1.3	0.79	1.27	434.7	274.6	461.7	1.6	0.94	1.06
P0103_P0103_MOUSE	P4hb	Protein disulfide-isomerase OS-Mus musculus GN-P4hb PE1 SV-2	29	55.0	40.9	70.1	1.3	0.79	1.27	1796.6	1384.8	2328.5	1.3	0.77	1.30
O7J123_FAM8A_MOUSE	Fam8a8	Protein FAM8A OS-Mus musculus GN-Fam8a8 PE2 SV-1	2	10.7	7.9	13.6	1.3	0.78	1.27	89.7	51.7	86.9	1.7	1.03	0.

QB8D9_IF4B_MOUSE	EIf4b	Eukaryotic translation initiation factor 4B OS=Mus musculus GN=EIf4b PE=1 SV=1	1	4.1	3.3	5.6	1.2	0.72	1.38	4.1	3.3	5.5	1.2	0.74	1.35
QB6JK_FHOP_MOUSE	Fhfp5p	FGFR3 oncogene partner OS=Mus musculus GN=Fhfp5p PE=1 SV=1	1	6.7	5.4	9.3	1.2	0.72	1.38	6.7	5.4	9.1	1.2	0.74	1.35
QD220_HNRH_MOUSE	Hnrh	Heterogeneous nuclear ribonucleoprotein A1 C12 OS=Mus musculus GN=Hnrhnc P=1 PE=2	80.9	65.2	111.8	1.2	0.72	1.38	75.74	558.0	439.6	1.2	0.74	1.35	
Q6131_MD31_MOUSE	Med13	Mediator of RNA polymerase II transcription subunit 13-like OS=Mus musculus GN=Med13 P=1 PE=2	2	16.5	13.3	22.8	1.2	0.72	1.38	33.0	26.6	44.8	1.2	0.74	1.36
P16460_ASSY_MOUSE	Ass1	Argininosuccinate synthase OS=Mus musculus GN=Ass1 PE=1 SV=1	2	4.6	3.7	6.3	1.2	0.72	1.38	9.1	7.4	12.4	1.2	0.74	1.36
QD987_CWC2_MOUSE	Cwc2	Pre-mRNA splicing factor CWC2 OS=Mus musculus GN=Cwc2 P=2 SV=2	2	4.6	3.7	6.3	1.2	0.72	1.38	20.5	16.0	26.0	1.2	0.74	1.33
Q6GCI_F13A_MOUSE	Fam193a	Protein FAM193A OS=Mus musculus GN=Fam193a P=2 SV=2	1	7.5	6.1	10.4	1.2	0.72	1.39	7.5	6.1	10.2	1.2	0.73	1.37
QEGP5_F2D2_MOUSE	Setd2	Histone lysine N-methyltransferase SETD2 OS=Mus musculus GN=Setd2 P=1 SV=2	10	8.4	6.8	11.7	1.2	0.72	1.39	101.5	70.8	119.1	1.4	0.85	1.17
Q61245_CBDA_MOUSE	Ccl1a1	Collagen alpha 1(I) chain OS=Mus musculus GN=Ccl1a1 PE=1 SV=2	2	10.5	8.6	14.7	1.2	0.72	1.40	21.1	17.1	28.8	1.2	0.73	1.37
P12382_P34L_MOUSE	P34l	ATP-dependent RNA helicase P34l OS=Mus musculus GN=P34l PE=1 SV=4	2	5.6	4.6	7.8	1.2	0.72	1.41	11.7	9.1	14.2	1.2	0.73	1.37
QJH93_DHX36_MOUSE	Dhx36	ATP-dependent RNA helicase DHX36 OS=Mus musculus GN=Dhx36 PE=1 SV=2	11	6.9	5.7	9.7	1.2	0.72	1.40	113.5	70.8	119.0	1.6	0.95	1.05
AAU401_AZAU1_MOUSE	Raly	RNA-binding protein Raly (Fragmin) OS=Mus musculus GN=Raly PE=1 SV=1	2	94.2	76.9	131.9	1.2	0.71	1.40	188.3	153.9	258.7	1.2	0.73	1.37
QD2146_DKX7_MOUSE	Dkx7	RNA-binding protein DKX7 OS=Mus musculus GN=Dkx7 P=1 SV=1	1	2.1	1.8	3.0	1.2	0.71	1.40	4.0	3.4	5.8	1.2	0.73	1.37
QD281_VATB2_MOUSE	Vatb2	Y-type protein ATPase subunit B, brain isoform OS=Mus musculus GN=Vatb2 P=1 SV=1	2	3.6	2.9	5.1	1.2	0.71	1.40	7.2	5.9	9.9	1.2	0.73	1.38
Q63916_SDRP_MOUSE	SDrp	Serine deprivation-response protein OS=Mus musculus GN=SDrp PE=1 SV=3	9	25.5	20.9	34.8	1.2	0.71	1.40	229.9	160.1	269.2	1.4	0.85	1.17
Q58K6_AW3_MOUSE	Ank3	Ankyrin-3 OS=Mus musculus GN=Ank3 P=1 SV=1	6	5.6	4.6	7.8	1.2	0.71	1.41	33.9	27.6	46.4	1.2	0.73	1.37
P47754_CAD2_MOUSE	Cad2	F-actin-capping protein subunit alpha2 OS=Mus musculus GN=Cad2a PE=1 SV=3	2	30.7	25.2	43.2	1.2	0.71	1.41	62.1	50.8	84.7	1.2	0.73	1.36
OS4784_DAPK3_MOUSE	Dapk3	Death-associated protein kinase 3 OS=Mus musculus GN=Dapk3 P=1 SV=1	1	11.7	9.6	16.5	1.2	0.71	1.41	11.7	9.6	16.2	1.2	0.72	1.38
Q9910_DPY30_MOUSE	Dpy30	Protein dpy-30 homolog OS=Mus musculus GN=Dpy30 P=1 SV=1	2	14.3	11.8	20.1	1.2	0.71	1.41	28.6	23.5	39.5	1.2	0.72	1.38
Q63871_P9AB_MOUSE	Polr2k	DNA-dependent RNA polymerase, II, and III subunit P9AB C4 OS=Mus musculus GN=Polr2k P=2 SV=2	1	53.6	44.1	75.6	1.2	0.71	1.41	53.6	44.1	74.1	1.2	0.72	1.38
Q5C9D9_GLK3_MOUSE	Glrx3	Glutathione S-transferase K3 OS=Mus musculus GN=Glrx3 PE=1 SV=1	1	6.3	8.4	5.5	1.2	0.71	1.41	6.3	5.2	8.7	1.2	0.72	1.38
Q8VC08_Q8VC08_MOUSE	Cald1	Caldesmon 1 OS=Mus musculus GN=Cald1 PE=1 SV=1	16	18.5	15.2	26.1	1.2	0.71	1.41	376.1	252.3	424.3	1.5	0.89	1.13
Q61246_CD2A_MOUSE	Cdk2na	Cyclin-dependent kinase inhibitor 2A, isoforms 1/2 OS=Mus musculus GN=Cdk2na P=1 SV=2	2	10.0	8.2	14.1	1.2	0.71	1.41	20.0	16.5	27.7	1.2	0.72	1.38
Q13077_Q2U77_MOUSE	Trim21	E3 ubiquitin-protein ligase TRIM21 OS=Mus musculus GN=Trim21 P=2 SV=2	18	49.5	40.7	69.9	1.2	0.71	1.41	113.7	92.1	155.2	1.2	0.72	1.38
A2BD00_A2BD00_MOUSE	Adnp	Actively-dependent neurotrophin/homocysteine protein OS=Mus musculus GN=Adnp PE=1 SV=1	13	8.1	6.6	11.4	1.2	0.71	1.41	119.9	103.3	173.8	1.2	0.69	1.40
P37889_FBLN2_MOUSE	Fbln2	Fibulin-2 OS=Mus musculus GN=Fbln2 P=1 SV=2	17	40.3	33.2	57.0	1.2	0.71	1.41	950.6	904.1	1520.3	1.1	0.63	1.60
P91783_CD4K1_MOUSE	Cox4k1	Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Mus musculus GN=Cox4k1 PE=1 SV=1	3	7.0	5.8	9.9	1.2	0.71	1.42	20.9	17.0	28.5	1.2	0.73	1.36
QD2142_QD2142_MOUSE	OCAD2	Oxidation-reduction 2 OS=Mus musculus GN=OCAD2 P=1 SV=1	3	4.4	3.7	6.2	1.2	0.71	1.42	9.7	8.2	13.1	1.2	0.73	1.36
Q09000_NCOA3_MOUSE	Ncoa3	Nuclear receptor coactivator 3 OS=Mus musculus GN=Ncoa3 P=1 SV=2	2	56.2	46.4	79.8	1.2	0.71	1.42	112.4	92.8	156.1	1.2	0.72	1.39
Q91V1L_BAG3_MOUSE	Bag3	BAG family molecular chaperone regulator 3 OS=Mus musculus GN=Bag3 PE=1 SV=2	1	9.9	8.2	14.1	1.2	0.71	1.42	9.9	8.2	13.8	1.2	0.72	1.39
Q12M90_ACO1_MOUSE	ACO1	Acyloxy-coenzyme A thioesterase 10, mitochondrial OS=Mus musculus GN=ACO1 P=2 SV=1	1	4.6	3.8	6.3	1.2	0.71	1.42	4.6	3.8	6.2	1.2	0.72	1.39
QD2209_VCP1_MOUSE	Vcp1	Spinnomegal complex protein 1 OS=Mus musculus GN=Vcp1 PE=1 SV=2	1	8.6	7.2	11.6	1.2	0.70	1.42	8.5	7.2	11.4	1.2	0.72	1.39
QD2202_KNOP1_MOUSE	Knop1	Lysine-rich nuclear protein 1 OS=Mus musculus GN=Knop1 PE=1 SV=2	2	8.7	7.2	12.4	1.2	0.70	1.42	17.4	14.5	24.3	1.2	0.72	1.39
Q8R120_KNOP1_MOUSE	Znf330	Zinc finger protein 830 OS=Mus musculus GN=Znf330 P=1 SV=1	1	11.3	9.4	16.1	1.2	0.70	1.42	11.3	9.4	15.8	1.2	0.72	1.40
P49871_CAV1_MOUSE	CAV1	Caveolin-1 OS=Mus musculus GN=CAV1 PE=1 SV=1	2	20.4	16.9	29.0	1.2	0.70	1.42	69.8	58.0	95.2	1.2	0.72	1.40
Q5CQAL_TPPCS_MOUSE	Trappc5	Trafficking protein particle complex subunit 5 OS=Mus musculus GN=Trappc5 P=1 SV=1	1	5.2	4.3	7.4	1.2	0.70	1.43	5.2	4.3	7.3	1.2	0.71	1.40
Q6P549_SHP2_MOUSE	Shp2	Phosphatidylinositol 3,4,5-trisphosphate 5-phosphatase 2 OS=Mus musculus GN=Shp2 P=1 SV=1	2	9.8	8.2	14.1	1.2	0.70	1.43	19.7	16.4	27.6	1.2	0.71	1.40
Q5Z0C6_MSH2_MOUSE	Msh2	RNA-binding protein Msh2 homolog 2 OS=Mus musculus GN=Msh2 P=1 SV=1	5	10.9	9.1	15.6	1.2	0.70	1.43	113.3	84.0	141.2	1.3	0.80	1.25
Q70435_VSY2_MOUSE	Vsy2	Vesicle OS=Mus musculus GN=VSY2 P=1 SV=1	1	10.9	9.1	15.6	1.2	0.70	1.44	10.9	9.1	15.6	1.2	0.71	1.40
QD0606_RPN2_MOUSE	Rpn2	Dolichyl-3-phosphoglycerol-2-protein glycosyltransferase subunit 2 OS=Mus musculus GN=	7	16.8	14.1	24.2	1.2	0.70	1.44	145.5	98.2	165.1	1.5	0.88	1.13
Q59V90_LSM2_MOUSE	Lsm2	U5 snRNP-associated Sm-like protein Lsm2 OS=Mus musculus GN=Lsm2 P=3 SV=1	2	6.3	5.3	9.1	1.2	0.70	1.44	12.6	10.6	17.7	1.2	0.71	1.41
Q16016_DKQ4E_MOUSE	Dkq4e	Striatin OS=Mus musculus GN=Dkq4e P=1 SV=1	1	11.4	9.4	16.1	1.2	0.70	1.44	11.4	9.4	16.1	1.2	0.72	1.40
QVDF2_UHRF1_MOUSE	Uhrf1	E3 ubiquitin-protein ligase UHRF1 OS=Mus musculus GN=Uhrf1 P=1 SV=2	6	16.2	13.7	23.4	1.2	0.69	1.44	120.6	79.5	133.7	1.3	0.76	1.31
Q92188_RED_MOUSE	Ik	Protein Red OS=Mus musculus GN=Ik P=1 SV=2	7	20.3	17.1	29.3	1.2	0.69	1.44	167.7	132.5	222.8	1.3	0.75	1.33
Q51259_P9H2_MOUSE	P9h2	Prohibitin-2 OS=Mus musculus GN=P9h2 P=1 SV=1	8	13.6	11.5	19.7	1.2	0.69	1.45	125.2	109.3	183.8	1.1	0.68	1.47
Q5C9W6_I2A_MOUSE	I2a	Eukaryotic translation initiation factor 2 subunit 1 OS=Mus musculus GN=EIF2L3 P=1 SV=3	9	10.7	9.1	15.6	1.2	0.69	1.45	138.7	114.6	192.2	1.4	0.85	1.14
Q5C575_LR42_MOUSE	Lrrc47	Lysine-rich repeat-containing protein 47 OS=Mus musculus GN=Lrrc47 P=2 SV=1	3	6.7	5.6	9.6	1.2	0.69	1.45	20.6	19.2	32.2	1.1	0.64	1.57
Q85700_RAEL1_MOUSE	Rael1	mRNA export factor OS=Mus musculus GN=Rael1 PE=1 SV=1	3	12.9	10.9	18.7	1.2	0.69	1.45	41.4	34.7	58.3	1.2	0.71	1.41
Q90207_NBN_MOUSE	Nbn	Nibrin OS=Mus musculus GN=NBN PE=1 SV=1	2	4.7	4.0	6.8	1.2	0.69	1.45	4.7	4.0	6.7	1.2	0.70	1.43
Q82846_POG2_MOUSE	Pog2	Page transposable element with ZNF domain OS=Mus musculus GN=Pog2 PE=2 SV=2	4	11.1	9.5	16.3	1.2	0.68	1.46	42.7	37.8	63.5	1.1	0.67	1.49
Q9Q56_DREB_MOUSE	Dreb1	Debrin OS=Mus musculus GN=Dreb1 PE=1 SV=4	26	24.1	20.7	35.4	1.2	0.68	1.47	1038.1	842.1	1415.9	1.2	0.73	1.36
P56959_FUS_MOUSE	Fus	Transcription factor FUS OS=Mus musculus GN=FUS PE=2 SV=1	5	13.2	11.3	19.4	1.2	0.68	1.47	99.7	80.0	80.7	1.2	1.23	0.81
Q80UW6_RPB1_MOUSE	Rpb1	RNA polymerase I, and II, and III subunit RPB1 OS=Mus musculus GN=Rpb1a2 P=2 SV=2	2	20.9	17.6	30.0	1.2	0.68	1.47	44.7	38.5	49.4	1.4	0.63	1.21
Q51V55_PQB1_MOUSE	Pqb1	Polyglutamine-binding protein 1 OS=Mus musculus GN=Pqb1 P=1 SV=1	2	5.4	4.6	8.0	1.2	0.68	1.47	10.8	9.3	15.6	1.2	0.69	1.44
Q91406_PRR4_MOUSE	Prr4	UAF1 small nuclear ribonucleoprotein Prr4 OS=Mus musculus GN=Prr4 P=1 SV=1	7	19.3	16.6	28.4	1.2	0.68	1.47	161.4	123.6	207.8	1.3	0.78	1.29
Q9D39_SDF2L_MOUSE	Sdf2l1	Striatal cell-derived factor 2-like protein 1 OS=Mus musculus GN=SDF2L1 PE=1 SV=2	1	13.5	11.6	19.9	1.2	0.68	1.47	13.5	11.6	19.5	1.2	0.69	1.44
Q9D221_NU1L_MOUSE	Nu1l	NU1L nuclear cluster factor OS=Mus musculus GN=NU1L PE=1 SV=1	1	10.5	9.1	15.6	1.2	0.68	1.47	10.5	9.1	15.6	1.2	0.69	1.44
Q99172_APEB3_MOUSE	Apeb3	DNA-C/EBP-binding protein APEB3 OS=Mus musculus GN=Apeb3 P=1 SV=2	2	20.9	18.1	31.0	1.2	0.67	1.48	41.9	36.2	60.0	1.2	0.69	1.46
Q91077_QCR2_MOUSE	Uqc2c	Cytochrome b-c1 complex subunit 2, mitochondrial OS=Mus musculus GN=Uqc2c P=1 SV=1	2	9.4	8.2	14.0	1.2	0.67	1.48	18.8	16.4	27.5	1.2	0.69	1.45
Q5L0L6_GSK3B_MOUSE	Gsk3b	Glycogen synthase kinase-3 beta OS=Mus musculus GN=GSK3B P=1 SV=1	2	9.4	8.2	14.0	1.2	0.67	1.48	43.5	36.8	58.8	1.2	0.70	1.42
Q61023_DNB1_MOUSE	Dnb1	RNA1 homolog subfamily B member 1 OS=Mus musculus GN=Dnb1 P=2 SV=3	3	14.6	12.7	21.7	1.2	0.67	1.49	68.6	68.0	114.4	1.0	0.60	1.67
Q92102_LAP2B_MOUSE	Lap2b	Lamina-associated polypeptide 2, isoforms beta/delta/epsilon/gamma OS=Mus musculus GN=	4	17.0	14.8	25.3	1.2	0.67	1.49	68.6	60.0	100.8	1.1	0.68	1.47
Q92121_IMP3_MOUSE	Imp3	U3 small nuclear ribonucleoprotein subunit IMP3 OS=Mus musculus GN=Imp3 P=2 SV=1	4	7.5	6.5	11.2	1.2	0.67	1.49	73.9	54.9	92.3	1.3	0.80	1.25
Q90B06_VNR1_MOUSE	Vnr1	Vesicle OS=Mus musculus GN=VNR1 P=1 SV=1	2	9.7	8.2	14.0	1.2	0.67	1.49	9.7	8.2	14.0	1.2	0.68	1.47
PE1117_NUCB2_MOUSE	Nucb2	Nucleobindin-2 OS=Mus musculus GN=Nucb2 P=1 SV=1	2	9.3	8.1	13.9	1.2	0.67	1.50	18.6	16.2	27.3	1.1	0.68	1.47
P31570_PRDX1_MOUSE	Prdx1	Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 P=1 SV=1	5	12.4	10.8	18.5	1.1	0.67	1.50	60.9	57.4	96.6	1.1	0.63	1.59
P126851_PGS1_MOUSE	Pgs1	Rhlygan OS=Mus musculus GN=Pgs1 PE=1 SV=1	2	14.2</											

Q90B20_MOUSE	Ikbbp	Inhibitor of nuclear factor kappa-B kinase-interacting protein OS-Mus musculus GN-Ikbbp PE-2	1	7.6	7.8	7.8	1.0	0.57	1.76	7.6	7.8	13.1	1.0	0.58	1.72
Q90B20_MOUSE	Atp5o	ATP synthase subunit O, mitochondrial OS-Mus musculus GN-Atp5o PE1 SV-1	1	5.4	5.6	5.6	1.0	0.57	1.76	5.4	5.6	5.4	1.0	0.58	1.72
GN9593_GN9593_MOUSE	GN9593	GN9593 OS-Mus musculus GN-GN9593 PE1 SV-1	1	24.9	24.9	24.9	1.0	0.57	1.76	118.7	118.7	201.3	1.3	0.75	21.4
OS130_N1_MOUSE	Emg1	Ribosomal RNA small subunit methyltransferase NEP1 OS-Mus musculus GN-Emg1 PE2 SV-1	5	6.1	6.3	6.8	1.0	0.56	1.78	34.2	30.1	50.6	1.1	0.68	1.48
Q92105_C1_MOUSE	Clic1	Chloride intracellular channel protein 1 OS-Mus musculus GN-Clic1 PE1 SV-3	1	3.6	3.8	4.6	1.0	0.56	1.78	3.6	3.8	6.3	1.0	0.57	1.75
Q94012_RALY_MOUSE	Ralyl	RNA-binding protein G1 OS-Mus musculus GN-Ralyl PE2 SV-2	24	31.2	31.2	31.2	1.0	0.56	1.79	316.7	295.2	487.1	1.1	0.63	1.59
Q90801_HNRP1_MOUSE	Hnmp1	Heterogeneous nuclear ribonucleoprotein 1 OS-Mus musculus GN-Hnmp1 PE1 SV-2	19	13.3	13.9	23.8	1.0	0.56	1.79	310.5	296.7	498.1	1.0	0.62	1.61
Q90716_ARPC3_MOUSE	Arpc3	Actin-related protein 2/3 complex subunit 3 OS-Mus musculus GN-Arpc3 PE1 SV-2	9	9.3	9.7	16.7	1.0	0.56	1.80	146.0	124.7	209.6	1.2	0.70	1.44
Q92121_TADBP_MOUSE	Tadbp	TAR DNA-binding protein 4 OS-Mus musculus GN-Tadbp PE1 SV-1	30	31.8	33.5	57.4	0.9	0.55	1.81	412.23	425.72	715.83	1.0	0.58	1.74
Q92122_VIME_MOUSE	Vime	Vimentin OS-Mus musculus GN-Vime PE1 SV-2	3	10.3	10.3	10.3	0.9	0.55	1.81	28.0	28.0	318.0	1.0	0.57	1.87
Q90219_CNPY2_MOUSE	Cnpy2	Protein canopy homolog 2 OS-Mus musculus GN-Cnpy2 PE1 SV-1	2	13.2	13.9	23.9	0.9	0.55	1.82	26.3	27.9	46.8	0.9	0.56	1.78
Q90194_RFA2_MOUSE	Rfa2	Replication protein A 32 kDa subunit OS-Mus musculus GN-Rfa2 PE1 SV-1	4	29.4	31.1	53.4	0.9	0.55	1.82	102.9	131.7	221.4	0.8	0.46	2.15
Q92PML_ARPC5_MOUSE	Arpc5	Actin-related protein 2/3 complex subunit 5 OS-Mus musculus GN-Arpc5 PE2 SV-2	3	38.5	38.5	38.5	0.9	0.55	1.82	38.8	38.8	68.4	0.9	0.52	1.79
Q90W54_FACT1_MOUSE	Fma2e24	CAAM premyr protease 1 homolog OS-Mus musculus GN-Fma2e24 PE1 SV-2	1	5.6	5.9	10.2	0.9	0.55	1.83	11.1	11.9	20.0	0.9	0.56	1.79
Q90C51_T176A_MOUSE	Tmem176a	Transmembrane protein 176A OS-Mus musculus GN-Tmem176a PE2 SV-2	1	2.6	2.8	4.7	0.9	0.55	1.83	2.6	2.8	4.6	0.9	0.56	1.79
Q90852_KMT2B_MOUSE	Kmt2b	Histone-lysine N-methyltransferase 2B OS-Mus musculus GN-Kmt2b PE1 SV-3	1	4.3	4.6	7.8	0.9	0.55	1.83	4.3	4.6	7.7	0.9	0.56	1.80
Q92T72_F038B_MOUSE	Fam208b	Protein FAM208B OS-Mus musculus GN-Fam208b PE2 SV-2	1	7.6	8.3	14.2	0.9	0.55	1.83	6.0	6.6	10.1	0.9	0.56	1.80
Q90883_CHMB_MOUSE	Chmbp4	Charged-methyltransferase body protein 4B OS-Mus musculus GN-Chmbp4 PE1 SV-2	1	7.2	7.7	13.8	0.9	0.55	1.83	7.2	7.7	13.0	0.9	0.56	1.80
P15538_TRD3A_MOUSE	Trm30a	Tripartite motif-containing protein 30A OS-Mus musculus GN-Trm30a PE1 SV-2	1	6.0	6.4	11.0	0.9	0.54	1.84	6.0	6.4	10.7	0.9	0.55	1.80
F24261_HKRC2_MOUSE	Herc6	HEC15-1 protein ligase Herc6 OS-Mus musculus GN-Herc6 PE2 SV-1	1	2.3	2.5	4.3	0.9	0.54	1.84	2.3	2.5	4.2	0.9	0.55	1.81
Q98362_TBMA_MOUSE	Tbma4	Tubulin alpha 4A chain OS-Mus musculus GN-Tbma4 PE1 SV-1	4	14.7	15.8	28.1	0.9	0.54	1.85	57.7	58.7	86.9	0.9	0.68	1.48
GX981X_GX981X_MOUSE	Heat1	HEAT repeat containing 1 OS-Mus musculus GN-Heat1 PE4 SV-1	2	3.7	4.0	6.9	0.9	0.54	1.85	3.7	4.0	6.9	0.9	0.55	1.81
Q88M11_SVIC_MOUSE	Lars	lysine- ϵ -NH ₂ ligase, cytoplasmic OS-Mus musculus GN-Lars PE1 SV-2	2	9.0	9.7	16.7	0.9	0.54	1.86	17.9	19.4	32.7	0.9	0.55	1.83
Q925H1_TRPS1_MOUSE	Trps1	Zinc finger transcription factor TRPS1 OS-Mus musculus GN-Trps1 PE1 SV-1	4	7.6	8.3	14.2	0.9	0.54	1.86	31.3	35.0	52.6	1.1	0.66	1.50
P52480_KPYM_MOUSE	Pkm	Pyruvate kinase PKM OS-Mus musculus GN-Pkm PE1 SV-4	2	8.7	9.5	16.2	0.9	0.54	1.87	17.4	18.9	31.8	0.9	0.55	1.83
Q90824_PBB_MOUSE	Cbfh	Cerebin-binding factor subunit beta OS-Mus musculus GN-Cbfh PE1 SV-1	3	12.1	13.2	22.6	0.9	0.53	1.87	35.2	38.5	64.7	0.9	0.54	1.84
Q92G68_HFSA_MOUSE	Hf25a5	Intron-activator protein 205-A OS-Mus musculus GN-Hf25a5 PE2 SV-1	2	5.5	6.0	10.3	0.9	0.53	1.87	11.0	12.0	20.1	0.9	0.55	1.83
Q91481_VKCL1_MOUSE	Vkcl1	5.5 kDa-binding protein OS-Mus musculus GN-Vkcl1 PE2 SV-2	1	4.8	5.5	9.5	0.9	0.53	1.88	4.8	5.5	9.5	0.9	0.51	1.87
Q91754_VMA21_MOUSE	Vma21	Vacuolar ATPase assembly integral membrane protein Vma21 OS-Mus musculus GN-Vma21 PE1 SV-1	1	5.3	5.8	9.9	0.9	0.53	1.88	5.3	5.8	9.9	0.9	0.54	1.85
Q85854_MEF2_MOUSE	Mef2	Myelin expression factor 2 OS-Mus musculus GN-Mef2 PE1 SV-1	35	26.7	29.6	50.0	0.9	0.53	1.90	1142.0	1053.7	1771.8	1.1	0.64	1.55
P80801_P044_MOUSE	P044	Protein disulfide isomerase A4 OS-Mus musculus GN-P044 PE1 SV-3	3	5.0	5.6	9.8	0.9	0.52	1.91	15.1	17.9	31.5	0.9	0.64	1.51
P60824_CRRP_MOUSE	Crrp	Cdk2-inducible RNA-binding protein OS-Mus musculus GN-Crrp PE1 SV-1	5	16.3	16.9	29.5	0.9	0.52	1.91	84.2	85.0	160.0	0.9	0.53	1.87
Q90K70_RBM2_MOUSE	Rbm2	RNA-binding motif, single-stranded-interacting protein 2 OS-Mus musculus GN-Rbm2 PE2 SV-2	5	14.7	16.4	28.2	0.9	0.52	1.92	176.5	214.1	360.0	0.8	0.49	2.04
Q91XK1_STX6_MOUSE	Stx6	Syntaxin-6 OS-Mus musculus GN-Stx6 PE1 SV-1	1	2.3	2.6	4.5	0.9	0.52	1.93	2.3	2.6	4.4	0.9	0.53	1.90
Q92088_PPH_MOUSE	Pph	Peptidyl prolyl 4-cis-transaminase OS-Mus musculus GN-Pph PE2 SV-1	1	13.4	15.2	26.0	0.9	0.52	1.94	14.4	16.2	29.7	1.2	0.70	1.45
Q91U11_CFA20_MOUSE	Cfap20	Cilia- and flagella-associated protein 20 OS-Mus musculus GN-Cfap20 PE2 SV-1	4	7.4	8.4	14.3	0.9	0.51	1.94	42.8	33.8	56.9	1.3	0.75	1.33
Q1U071_SMU1_MOUSE	Smu1	WD40 repeat-containing protein SMU1 OS-Mus musculus GN-Smu1 PE2 SV-2	7	14.0	15.9	27.2	0.9	0.51	1.94	107.9	125.1	210.4	0.9	0.51	1.95
Q92P5C_SMAUG_MOUSE	Smaug1	Single-strand selective monofunctional uracil DNA glycosylase OS-Mus musculus GN-Smaug1 PE1 SV-1	1	6.0	6.9	11.7	0.9	0.51	1.94	6.0	6.9	11.5	0.9	0.52	1.91
Q91R15_ELP5_MOUSE	Elp5	RNA polymerase complex 5 OS-Mus musculus GN-Elp5 PE1 SV-1	1	4.4	5.0	8.6	0.9	0.51	1.94	4.4	5.0	8.6	0.9	0.52	1.91
Q91K43_PKH3_MOUSE	Plekha3	Pleckstrin homolog domain-containing family G member 3 OS-Mus musculus GN-Plekha3 PE1 SV-1	1	3.6	4.1	7.0	0.9	0.51	1.95	3.6	4.1	6.9	0.9	0.52	1.91
PE7776_P1B_MOUSE	P1B	Prohibitin OS-Mus musculus GN-P1B PE1 SV-1	6	12.6	14.3	24.5	0.9	0.51	1.95	71.1	79.4	133.5	0.9	0.53	1.88
Q92P52_SHS7_MOUSE	Shs7	SHS7 domain-containing protein OS-Mus musculus GN-Shs7 PE2 SV-2	1	10.7	11.6	20.6	0.9	0.51	1.95	10.7	11.6	20.6	0.9	0.53	1.88
Q92PFA2_QBPFA_MOUSE	Cita	Clathrin light chain A OS-Mus musculus GN-Cita PE1 SV-1	3	7.7	8.9	15.2	0.9	0.51	1.95	12.5	14.8	26.5	1.6	0.71	1.01
Q90B30_SVIC_MOUSE	Iars	Isoleucine-tRNA ligase, cytoplasmic OS-Mus musculus GN-Iars PE2 SV-2	1	3.3	3.8	6.5	0.9	0.51	1.98	3.3	3.8	6.4	0.9	0.52	1.94
Q95142_SFN1_MOUSE	Sfn1	Sfn1-like OS-Mus musculus GN-Sfn1 PE1 SV-2	3	12.7	14.7	25.2	0.9	0.50	1.98	95.9	73.3	123.2	1.3	0.78	1.28
P20552_C02_MOUSE	Zcchc9	Chromatin protein homolog 9 OS-Mus musculus GN-Zcchc9 PE2 SV-2	1	4.8	5.5	9.5	0.9	0.50	1.98	4.8	5.5	9.3	0.9	0.51	1.95
Q90813_ZC03_MOUSE	Zcchc9	Zinc finger CCHC domain-containing protein 9 OS-Mus musculus GN-Zcchc9 PE2 SV-2	3	6.3	7.3	12.5	0.9	0.50	1.99	22.4	20.0	33.6	1.1	0.67	1.50
Q60739_BAG1_MOUSE	Bag1	BAG family molecular chaperone regulator 1 OS-Mus musculus GN-Bag1 PE1 SV-3	2	4.0	4.6	7.9	0.9	0.50	1.99	7.9	9.2	15.5	0.9	0.51	1.95
Q91866_MARE_MOUSE	Mare1	Microtubule-associated protein RFB family member 1 OS-Mus musculus GN-Mare1 PE1 SV-1	1	11.6	13.5	23.3	0.8	0.50	1.99	36.8	33.0	54.6	0.8	0.66	1.51
Q92305_WBP11_MOUSE	Wbp11	WD domain-binding protein 11 OS-Mus musculus GN-Wbp11 PE1 SV-1	1	12.6	13.9	24.6	0.9	0.50	2.01	7.4	8.3	12.4	0.9	0.51	1.97
P57784_R02A_MOUSE	Srnpa1	U2 small nuclear ribonucleoprotein A' OS-Mus musculus GN-Srnpa1 PE1 SV-2	17	11.5	12.6	23.0	0.8	0.49	2.02	84.2	503.8	355.0	0.8	0.48	2.18
Q91009_MP95_MOUSE	Wdr77	Methylcrotonyl coenzyme S OS-Mus musculus GN-Wdr77 PE1 SV-1	3	10.0	11.8	20.3	0.8	0.49	2.03	34.6	34.5	58.0	1.0	0.60	1.67
P10633_THD_MOUSE	Thd	Thymine DNA methyltransferase OS-Mus musculus GN-Thd PE1 SV-3	1	14.0	16.3	28.5	0.8	0.49	2.03	146.7	166.7	284.8	0.8	0.49	2.03
P48678_LHNA_MOUSE	Lhna	Prelamin-A/C OS-Mus musculus GN-Lhna PE1 SV-2	35	33.6	33.9	68.2	0.8	0.49	2.03	1466.8	1893.2	3183.4	0.8	0.46	2.17
Q62241_R02_MOUSE	Srnpc	U1 small nuclear ribonucleoprotein C OS-Mus musculus GN-Srnpc PE2 SV-1	1	15.1	18.0	30.9	0.8	0.49	2.04	15.1	18.0	30.3	0.8	0.50	2.00
Q81951_Q81951_MOUSE	C1qbp	Complement component 1 Q subcomponent-binding protein, mitochondrial OS-Mus musculus GN-C1qbp PE1 SV-1	2	4.2	5.1	8.7	0.8	0.49	2.05	8.5	10.2	17.1	0.8	0.50	2.01
P21436_P1B_MOUSE	P1B	Protein disulfide isomerase A4 OS-Mus musculus GN-P1B PE1 SV-3	3	4.7	5.6	9.8	0.8	0.49	2.05	16.1	19.7	34.6	0.8	0.50	2.01
Q92QW0_CAF1A_MOUSE	Chaf1a	Chromatin assembly factor 1 subunit A OS-Mus musculus GN-Chaf1a PE1 SV-1	2	8.3	10.0	17.1	0.8	0.49	2.06	16.6	19.9	33.5	0.8	0.50	2.02
Q90779_L3BP_MOUSE	L3bp3	Galactin-3-binding protein OS-Mus musculus GN-L3bp3 PE1 SV-1	3	6.2	7.5	12.8	0.8	0.49	2.06	18.3	20.3	34.2	0.9	0.54	1.87
Q92149_P1B_MOUSE	P1B	Protein disulfide isomerase A4 OS-Mus musculus GN-P1B PE1 SV-3	3	4.8	5.6	9.8	0.8	0.49	2.06	16.1	19.7	34.6	0.8	0.50	2.01
P56480_A7P_MOUSE	A7p	ATP synthase subunit beta, mitochondrial OS-Mus musculus GN-A7p5b PE1 SV-4	18	19.9	23.2	39.9	0.8	0.49	2.06	376.8	442.3	743.6	0.9	0.51	1.97
Q95K54_DK1_MOUSE	Dk1	H/ACA ribonucleoprotein complex subunit 4 OS-Mus musculus GN-Dk1 PE1 SV-4	2	7.6	9.1	15.7	0.8	0.49	2.07	15.2	18.3	30.8	0.8	0.49	2.03
Q92VCE_ARPC2_MOUSE	Arpc2	Actin-related protein 2/3 complex subunit 2 OS-Mus musculus GN-Arpc2 PE1 SV-3	9	11.3	13.9	23.6	0.8	0.48	2.08	186.5	211.3	355.3	0.9	0.52	1.91
Q92C86_D04_MOUSE	D04	Homocysteine methyltransferase OS-Mus musculus GN-D04 PE1 SV-1	1	6.6	8.0	14.0	0.8	0.48	2.08	64.7	74.8	128.8	0.8	0.49	2.03
Q91S6M_SELM_MOUSE	Selm	Selenenoprotein M OS-Mus musculus GN-Selm PE1 SV-3	1	6.4	7.9	13.6	0.8	0.47	2.11	6.4	7.9	13.4	0.8	0.48	2.07
Q81625_C05_MOUSE	Mbd211	Cyclic GMP-AMP synthase OS-Mus musculus GN-Mbd211 PE1 SV-1	2	2.7	3.3	5.7	0.8	0.47	2.13	3.3	6.6	11.1	0.8	0.48	2.09
Q91865_Q91865_MOUSE	Cog1	Conserved oligomeric Golgi complex subunit 3 OS-Mus musculus GN-Cog1 PE4 SV-1	1	7.5	9.3	16.0	0.8	0.47	2.14	7.5	9.3	15.			

G3X9A3_G3X9A3_MOUSE	Zbtb21	Protein zbtb21 OS=Mus musculus GN=Zbtb21 PE=1 SV=1	1	4.9	10.8	18.5	0.5	0.27	3.76	4.9	10.8	18.1	0.5	0.27	3.68
O5R1P1_P5B81_MOUSE	Pamrb3	Probasome subunit beta type-3 OS=Mus musculus GN=Pamrb3 PE=1 SV=1	1	2.8	6.2	10.6	0.5	0.26	3.79	2.8	6.2	10.4	0.5	0.27	3.71
Q5C765_GAR1_MOUSE	Gar1	H/ACA ribonucleoprotein complex subunit 1 OS=Mus musculus GN=Gar1 PE=2 SV=1	3	6.1	14.1	24.3	0.4	0.25	3.95	27.5	52.0	87.4	0.5	0.31	3.17
P28301_LYOX_MOUSE	Lox	Protein-lysine 6-oxidase OS=Mus musculus GN=Lox PE=1 SV=1	4	3.7	8.5	14.5	0.4	0.25	3.97	28.4	50.8	85.4	0.6	0.33	3.00
Q852W6_Q852W6_MOUSE	H2-M10.3	Histocompatibility 2, M region locus 10.3 OS=Mus musculus GN=H2-M10.3 PE=2 SV=1	1	3.4	7.8	13.4	0.4	0.25	3.97	3.4	7.8	13.1	0.4	0.26	3.89
PE2827_RAN_MOUSE	Ran	Binding nuclear protein OS=Mus musculus GN=Ran PE=1 SV=2	4	5.4	12.7	21.8	0.4	0.25	4.07	21.4	40.7	63.5	0.4	0.26	3.90
B2KFR8_B2KFR8_MOUSE	Ptdtd4	CaM-specific 3',5'-cyclic phosphodiesterase 4D OS=Mus musculus GN=Ptdtd4 PE=4 SV=1	1	3.5	8.3	14.2	0.4	0.24	4.09	3.5	8.3	14.0	0.4	0.25	4.01
P42669_PURA_MOUSE	Pura	Transcriptional activator protein Pur-alpha OS=Mus musculus GN=Pura PE=1 SV=1	11	14.7	36.0	61.8	0.4	0.24	4.21	181.7	447.9	753.1	0.4	0.24	4.14
Q811F9_RHG32_MOUSE	Arhgap32	Rho GTPase-activating protein 32 OS=Mus musculus GN=Arhgap32 PE=1 SV=2	1	1.5	3.8	6.5	0.4	0.24	4.25	1.5	3.8	6.4	0.4	0.24	4.17
Q81V94_LVJ1_MOUSE	Lvj1	Lysine-61 OS=Mus musculus GN=Lvj1 PE=1 SV=1	1	5.5	13.6	23.3	0.4	0.22	4.36	5.5	13.6	22.9	0.4	0.24	4.18
P70314_PITX1_MOUSE	Pitx1	Pituitary homeobox 1 OS=Mus musculus GN=Pitx1 PE=2 SV=2	1	13.7	34.7	59.5	0.4	0.23	4.35	13.7	34.7	58.4	0.4	0.23	4.26
Q99020_RODA_MOUSE	Hnmpab	Heterogeneous nuclear ribonucleoprotein A/B OS=Mus musculus GN=Hnmpab PE=1 SV=1	11	30.8	78.5	134.7	0.4	0.23	4.38	452.3	1345.9	2263.2	0.3	0.20	5.00
Q80547_HMRB_MOUSE	Hmrbr	Hydranion-mediated motility receptor OS=Mus musculus GN=Hmrbr PE=1 SV=4	1	3.8	16.8	30.3	0.4	0.23	4.41	3.8	9.8	16.5	0.4	0.23	4.33
P97432_NBR1_MOUSE	Nbr1	Next to BRCA1 gene 1 protein OS=Mus musculus GN=Nbr1 PE=1 SV=1	1	1.9	4.9	8.5	0.4	0.22	4.45	1.9	4.9	8.3	0.4	0.23	4.37
P32020_NLTP_MOUSE	Scp2	Non-specific lipid-transfer protein OS=Mus musculus GN=Scp2 PE=1 SV=3	1	3.6	9.3	15.9	0.4	0.22	4.49	3.6	9.3	15.6	0.4	0.23	4.41
P52927_HMG2_MOUSE	Hmg2a	High mobility group protein HMG1-C OS=Mus musculus GN=Hmg2a PE=1 SV=1	1	2.5	6.5	11.1	0.4	0.22	4.52	2.5	6.5	10.9	0.4	0.23	4.43
Q81V94_Q81V94_MOUSE	Dax1	2',5'-oligoadenylate synthetase protein 1 OS=Mus musculus GN=Dax1 PE=2 SV=1	3	3.3	8.5	15.1	0.4	0.22	4.61	3.3	8.5	15.8	0.4	0.21	4.72
Q62011_PDPN_MOUSE	Pdpn	Podoplanin OS=Mus musculus GN=Pdpn PE=1 SV=2	1	16.3	43.8	75.1	0.4	0.22	4.61	16.3	43.8	73.7	0.4	0.22	4.52
P30051_Tead1_MOUSE	Tead1	Transcriptional enhancer factor 1 OS=Mus musculus GN=Tead1 PE=1 SV=2	1	6.1	17.0	29.1	0.4	0.21	4.74	6.1	17.0	28.6	0.4	0.22	4.65
Q3JTN6_Q3JTN6_MOUSE	T70001014Rik	Protein T70001014Rik OS=Mus musculus GN=T70001014Rik PE=2 SV=1	1	2.9	7.9	13.6	0.4	0.21	4.75	2.9	7.9	13.4	0.4	0.21	4.66
P28714_MOX_MOUSE	Mox	Protein mox OS=Mus musculus GN=Mox PE=1 SV=1	2	2.9	8.3	14.2	0.4	0.21	4.85	2.9	8.3	15.2	0.4	0.21	4.76
UF8A	Ufp8a	MCC3697, isoform CRA_c OS=Mus musculus GN=Ufp8a PE=2 SV=1	1	5.8	16.7	28.6	0.3	0.20	4.93	5.8	16.7	28.1	0.3	0.21	4.84
Q9PWV4_ASAH1_MOUSE	Ash1	ACG ceramidase OS=Mus musculus GN=Ash1 PE=1 SV=1	2	4.7	13.6	23.3	0.3	0.20	4.94	9.4	27.1	45.6	0.3	0.21	4.84
B180C7_B180C7_MOUSE	Hsp22	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Mus musculus GN=	13	4.7	13.4	23.0	0.3	0.20	4.95	63.1	165.3	278.0	0.4	0.23	4.41
P25976_UBF1_MOUSE	Ubf1	Nucleolar transcription factor 1 OS=Mus musculus GN=Ubf1 PE=1 SV=1	1	4.6	13.3	22.8	0.3	0.20	4.99	4.6	13.3	22.3	0.3	0.20	4.89
AP6PW4_WDR76_MOUSE	Wdr76	WD repeat-containing protein 76 OS=Mus musculus GN=Wdr76 PE=2 SV=1	2	4.7	13.9	23.8	0.3	0.20	5.04	4.5	27.8	46.7	0.3	0.20	4.94
Q8B8Q3_HNC2_MOUSE	Hnc2	Zinc finger protein baszumc1-2 OS=Mus musculus GN=Hnc2 PE=2 SV=3	1	2.9	8.7	14.9	0.3	0.20	5.04	2.9	8.7	14.6	0.3	0.20	4.94
Q80666_HNRP2_MOUSE	Hnmp2	Heterogeneous nuclear ribonucleoprotein 2 OS=Mus musculus GN=Hnmp2 PE=1 SV=2	1	10.0	25.4	59.4	0.3	0.20	5.04	100.4	249.4	419.7	0.4	0.21	5.00
Q97802_COE1_MOUSE	Ebf1	Transcription factor COE1 OS=Mus musculus GN=Ebf1 PE=1 SV=1	1	2.2	6.5	11.1	0.3	0.19	5.13	2.2	6.5	10.9	0.3	0.20	5.03
Q54962_BAF_MOUSE	Banf1	Barrier-to-autointegration factor OS=Mus musculus GN=Banf1 PE=1 SV=1	3	6.4	19.2	32.9	0.3	0.19	5.15	49.2	163.9	275.5	0.3	0.18	5.60
G3X9W0_G3X9W0_MOUSE	Hnmp2	Heterogeneous nuclear ribonucleoprotein 2, isoform CRA_a OS=Mus musculus GN=Hnmp2 PE=	1	3.1	9.3	16.0	0.3	0.19	5.23	3.1	9.3	15.7	0.3	0.19	5.13
Q47522_C2NC_MOUSE	Cnc	Cyflin-C OS=Mus musculus GN=Cnc PE=2 SV=4	1	10.3	31.8	54.5	0.3	0.19	5.27	10.3	31.8	53.5	0.3	0.19	5.17
Q8V101_NAA40_MOUSE	Naa40	N-acetyltransferase 40 OS=Mus musculus GN=Naa40 PE=2 SV=1	4	3.1	9.8	16.9	0.3	0.19	5.38	11.8	46.8	78.7	0.3	0.15	6.67
Q54790_MAFG_MOUSE	Mafg	Transcription factor MAFG OS=Mus musculus GN=Magf PE=1 SV=1	2	2.9	9.1	15.6	0.3	0.19	5.39	5.8	18.2	30.6	0.3	0.19	5.29
Q8Y9N6_NMG2_MOUSE	Nmg2	Neurotigin-3 OS=Mus musculus GN=Nmg2 PE=1 SV=2	1	1.0	3.2	5.6	0.3	0.19	5.40	1.0	3.2	5.4	0.3	0.19	5.29
Q99P64_RGS18_MOUSE	Rgs18	Regulator of G-protein signaling 18 OS=Mus musculus GN=Rgs18 PE=2 SV=1	1	3.5	11.0	18.9	0.3	0.18	5.48	3.5	11.0	18.5	0.3	0.19	5.34
Q6P1E1_ZMIZ1_MOUSE	Zmiz1	Zinc finger LIM domain-containing protein 1 OS=Mus musculus GN=Zmiz1 PE=2 SV=1	1	6.0	19.1	32.8	0.3	0.18	5.48	6.0	19.1	32.2	0.3	0.19	5.38
Q8V108_RF1_MOUSE	Rfap1	Replication protein A 70 kDa DNA-binding subunit OS=Mus musculus GN=Rfap1 PE=1 SV=1	15	6.0	19.5	33.4	0.3	0.18	5.57	183.1	310.1	523.5	0.6	0.35	2.85
P35552_HMGL_MOUSE	Cpxar	Cytochrome-dependent cytochrome P450 reductase, mitochondrial OS=Mus musculus GN=Cpxar	2	3.1	10.8	18.8	0.3	0.17	5.64	7.2	21.2	35.4	0.3	0.17	5.72
Q8V108_Schistosoma	Japuncin	Japuncin (Blood fluke)	2	3.1	10.8	18.8	0.3	0.17	6.02	6.2	21.7	36.4	0.3	0.17	5.91
Q92124_Q92124_MOUSE	Parp1	Poly (ADP-ribose) polymerase family, member 1 OS=Mus musculus GN=Parp1 PE=1 SV=1	21	7.8	27.3	46.7	0.3	0.17	6.02	62.7	697.9	1173.5	0.3	0.19	5.22
Q92204_MCP2_MOUSE	Mcp2	Methyl-CpG binding protein 2 OS=Mus musculus GN=Mcp2 PE=1 SV=1	3	4.3	15.9	26.6	0.3	0.16	6.25	23.8	16.9	43.8	0.4	0.23	4.37
P17918_PCNA_MOUSE	Pcna	Proliferating cell nuclear antigen OS=Mus musculus GN=Pcna PE=1 SV=2	6	5.4	20.3	34.8	0.3	0.16	6.44	42.3	162.6	273.4	0.3	0.15	6.46
Q61686_CBX5_MOUSE	Cbx5	Chromatin protein homolog 5 OS=Mus musculus GN=Cbx5 PE=1 SV=1	4	13.8	52.3	89.7	0.3	0.15	6.48	46.9	178.5	300.2	0.3	0.16	6.40
Q04750_TOP1_MOUSE	Top1	DNA topoisomerase 1 OS=Mus musculus GN=Top1 PE=1 SV=2	5	5.6	21.4	36.7	0.3	0.15	6.52	27.8	86.1	144.7	0.3	0.19	5.20
Q3LJL2_FRDC1_MOUSE	Frdc1	Frh2 domain-containing protein 1 OS=Mus musculus GN=Frhc1 PE=2 SV=3	1	2.0	7.7	13.2	0.3	0.15	6.57	2.0	7.7	13.0	0.3	0.16	6.45
PE8433_H31_MOUSE	Hist1h3a	Histone H3.1 OS=Mus musculus GN=H3a1 PE=1 SV=2	20	201.4	815.9	1399.1	0.2	0.14	6.95	11787.6	39478.7	66382.6	0.3	0.18	5.63
EP9PW6_EBPW6_MOUSE	Gcat	2-aminio-3-ketobutyrate coenzyme A ligase, mitochondrial OS=Mus musculus GN=Gcat PE=3 SV=1	1	1.0	4.1	7.1	0.2	0.14	6.97	1.0	4.1	6.9	0.2	0.15	6.84
Q8Y9W1_HMG25_MOUSE	Hmg25	Rho GTPase-activating protein 25 OS=Mus musculus GN=Hmg25 PE=1 SV=2	1	3.6	14.9	25.6	0.2	0.14	7.09	3.6	14.9	25.1	0.2	0.14	6.96
Q84857_C0S6A1_MOUSE	C0s6a1	Collagen alpha-1(XI) chain OS=Mus musculus GN=C0s6a1 PE=2 SV=1	1	2.0	8.2	14.1	0.2	0.14	7.15	2.0	8.2	13.8	0.2	0.14	7.01
P08249_MDHM_MOUSE	Mdh2	Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE=1 SV=3	5	4.5	19.2	33.0	0.2	0.14	7.38	24.8	80.8	135.9	0.3	0.18	5.48
Q9CS86_PARB3_MOUSE	Parb3	Partitioning defective 3 homolog 8 OS=Mus musculus GN=Parb3 PE=1 SV=2	1	6.6	29.6	50.8	0.2	0.13	7.65	6.6	29.6	49.8	0.2	0.13	7.50
PE2805_H4_MOUSE	Hist1h4a	Histone H4 OS=Mus musculus GN=H4a PE=1 SV=2	22	45.5	206.3	353.7	0.2	0.13	7.78	980.4	32164.1	54083.2	0.3	0.18	5.48
B7ZNU1_B7ZNU1_MOUSE	Fn1	Fibronectin OS=Mus musculus GN=Fn1 PE=1 SV=1	1	5.9	26.8	45.9	0.2	0.13	7.84	5.9	26.8	45.0	0.2	0.13	7.68
P23198_CBX3_MOUSE	Cbx3	Chromatin protein homolog 3 OS=Mus musculus GN=Cbx3 PE=1 SV=2	11	8.8	41.8	71.6	0.2	0.12	8.16	157.3	956.9	1608.9	0.2	0.10	10.23
O55112_AIF2_MOUSE	Aif2	AIF family member 2 OS=Mus musculus GN=Aif2 PE=2 SV=2	1	3.3	15.7	27.0	0.2	0.12	8.18	3.3	15.7	26.4	0.2	0.12	8.02
P53441_TBP1_MOUSE	Tbp1	Thymosin beta 4 OS=Mus musculus GN=Tbp1 PE=1 SV=1	1	1.2	5.6	9.6	0.2	0.12	8.20	1.2	5.6	9.5	0.2	0.12	8.04
Q8K3C3_L2IC_MOUSE	Lic	Protein L2IC OS=Mus musculus GN=Lic PE=1 SV=1	1	6.5	31.8	54.5	0.2	0.12	8.41	6.5	31.8	53.5	0.2	0.12	8.24
P83917_CBX1_MOUSE	Cbx1	Chromatin protein homolog 1 OS=Mus musculus GN=Cbx1 PE=1 SV=1	12	5.5	27.1	46.5	0.2	0.12	8.43	42.9	222.9	1223.0	0.3	0.18	5.49
Q80X32_C2D2_MOUSE	Hist1h3	Histone H3.2 OS=Mus musculus GN=H3a2 PE=1 SV=1	2	1.2	6.0	10.4	0.2	0.11	8.76	1.2	6.0	12.1	0.2	0.12	8.59
Q90Z08_H2AF_MOUSE	H2afy	Core histone macro-H2A.1 OS=Mus musculus GN=H2afy PE=1 SV=3	21	8.9	46.7	80.1	0.2	0.11	8.96	339.4	1278.2	2149.2	0.3	0.16	3.33
P27661_H2AX_MOUSE	H2ax	Histone H2AX OS=Mus musculus GN=H2afx PE=1 SV=2	4	19.3	101.6	174.2	0.2	0.11	9.04	76.0	438.9	738.1	0.2	0.10	9.71
Q99M09_NASP_MOUSE	Nasp	Nuclear autoantigenic sperm protein OS=Mus musculus GN=Nasp PE=1 SV=2	1	1.7	9.8	16.8	0.2	0.10	10.04	1.7	9.8	16.5	0.2	0.10	9.85
Q62W99_HIS21C_MOUSE	Hist1h2bc	Histone H2B type 1-C/2/G OS=Mus musculus GN=Hist1h2bc PE=1 SV=2	18	31.9	151.3	238.1	0.2	0.10	10.84	3456.8	19930.2	33512.2	0.2	0.10	9.69
Q64522_H2AZB_MOUSE	Hist2h2ab	Histone H2A type 2-B OS=Mus musculus GN=Hist2h2ab PE=1 SV=3	2	6.1	32.8	65.4	0.2	0.09	10.75	12.2	76.3	128.3	0.2	0.09	10.54
Q6QYQ1_SCNAA_MOUSE	Scn10a	Sodium channel protein type 10 subunit alpha OS=Mus musculus GN=Scn10a PE=1 SV=2	1	2.4	15.7	26.9	0.2	0.09	11.17	2.4	15.7	26.3	0.2	0.09	10.95
P70696_H2B1A_MOUSE	Hist1h2ba	Histone H2B type 1-A OS=Mus musculus GN=Hist1h2ba PE=1 SV=4	10	169.7											