

Protein FDR Confidence: Combined	Accession	# Unique Peptides	Gene symbol	Description	Abundance Ratio: (ipsi) / (intact)	Abundance Ratio P-Value: (ipsi) / (intact)
High	P01029	6	C4b	Complement C4-B OS=Mus musculus OX=10090 GN=C4b PE=1 SV=3	100,00	0,0000
High	P32261	15	Serpinc1	Antithrombin-III OS=Mus musculus OX=10090 GN=Serpinc1 PE=1 SV=1	12,07	0,0000
High	P01867	3	Igh-3	Ig gamma-2B chain C region OS=Mus musculus OX=10090 GN=Igh-3 PE=1 SV=3	10,82	0,0000
High	P07309	2	Ttr	Transthyretin OS=Mus musculus OX=10090 GN=Ttr PE=1 SV=1	9,62	0,0000
High	P03995	26	Gfap	Glial fibrillary acidic protein OS=Mus musculus OX=10090 GN=Gfap PE=1 SV=4	7,65	0,0000
High	Q00898	6	Serpina1e	Alpha-1-antitrypsin 1-5 OS=Mus musculus OX=10090 GN=Serpina1e PE=1 SV=1	6,96	0,0000
High	Q61703	11	Itih2	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Mus musculus OX=10090 GN=Itih2 PE=1 SV=1	6,92	0,0000
High	P01837	3	Igkc	Immunoglobulin kappa constant OS=Mus musculus OX=10090 GN=Igkc PE=1 SV=2	6,28	0,0000
High	P20918	18	Plg	Plasminogen OS=Mus musculus OX=10090 GN=Plg PE=1 SV=3	6,07	0,0000
High	P07724	43	Alb	Serum albumin OS=Mus musculus OX=10090 GN=Alb PE=1 SV=3	5,80	0,0000
High	P07758	4	Serpina1a	Alpha-1-antitrypsin 1-1 OS=Mus musculus OX=10090 GN=Serpina1a PE=1 SV=4	5,79	0,0000
High	P01864	3		Ig gamma-2A chain C region secreted form OS=Mus musculus OX=10090 PE=1 SV=1	5,72	0,0000
High	P29699	7	Ahsg	Alpha-2-HS-glycoprotein OS=Mus musculus OX=10090 GN=Ahsg PE=1 SV=1	5,29	0,0000
High	P23953	14	Ces1c	Carboxylesterase 1C OS=Mus musculus OX=10090 GN=Ces1c PE=1 SV=4	5,28	0,0000
High	Q9JLF6	8	Tgm1	Protein-glutamine gamma-glutamyltransferase K OS=Mus musculus OX=10090 GN=Tgm1 PE=1 SV=2	5,15	0,0000
High	Q91X72	20	Hpx	Hemopexin OS=Mus musculus OX=10090 GN=Hpx PE=1 SV=2	5,10	0,0000
Medium	Q06770	2	Serpina6	Corticosteroid-binding globulin OS=Mus musculus OX=10090 GN=Serpina6 PE=1 SV=1	5,09	0,0000
High	O08677	11	Kng1	Kininogen-1 OS=Mus musculus OX=10090 GN=Kng1 PE=1 SV=1	4,84	0,0000
High	P21614	16	Gc	Vitamin D-binding protein OS=Mus musculus OX=10090 GN=Gc PE=1 SV=2	4,74	0,0000
High	P22599	4	Serpina1b	Alpha-1-antitrypsin 1-2 OS=Mus musculus OX=10090 GN=Serpina1b PE=1 SV=2	4,59	0,0000
High	Q01339	10	Apoh	Beta-2-glycoprotein 1 OS=Mus musculus OX=10090 GN=Apoh PE=1 SV=1	3,99	0,0000
High	P29788	4	Vtn	Vitronectin OS=Mus musculus OX=10090 GN=Vtn PE=1 SV=2	3,88	0,0000
High	Q9R0H5	1	Krt71	Keratin, type II cytoskeletal 71 OS=Mus musculus OX=10090 GN=Krt71 PE=1 SV=1	3,71	0,0000
High	P01027	43	C3	Complement C3 OS=Mus musculus OX=10090 GN=C3 PE=1 SV=3	3,58	0,0000
High	P07759	12	Serpina3k	Serine protease inhibitor A3K OS=Mus musculus OX=10090 GN=Serpina3k PE=1 SV=2	3,44	0,0000
High	Q60590	3	Orm1	Alpha-1-acid glycoprotein 1 OS=Mus musculus OX=10090 GN=Orm1 PE=1 SV=1	3,43	0,0000
High	Q92111	38	Tf	Serotransferrin OS=Mus musculus OX=10090 GN=Tf PE=1 SV=1	3,41	0,0000
High	P20152	29	Vim	Vimentin OS=Mus musculus OX=10090 GN=Vim PE=1 SV=3	3,21	0,0000
High	Q61704	12	Itih3	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Mus musculus OX=10090 GN=Itih3 PE=1 SV=3	3,11	0,0000
High	P01899	5	H2-D1	H-2 class I histocompatibility antigen, D-B alpha chain OS=Mus musculus OX=10090 GN=H2-D1 PE=1 SV=2	3,09	0,0000
High	P28665	27	Mug1	Murinoglobulin-1 OS=Mus musculus OX=10090 GN=Mug1 PE=1 SV=3	3,02	0,0000
High	Q61702	13	Itih1	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Mus musculus OX=10090 GN=Itih1 PE=1 SV=2	3,00	0,0000
High	Q8VCM7	17	Fgg	Fibrinogen gamma chain OS=Mus musculus OX=10090 GN=Fgg PE=1 SV=1	2,94	0,0000

High	P19221	5	F2	Prothrombin OS=Mus musculus OX=10090 GN=F2 PE=1 SV=1	2,84	0,0000
High	Q9DB25	3	Alg5	Dolichyl-phosphate beta-glucosyltransferase OS=Mus musculus OX=10090 GN=Alg5 PE=1 SV=1	2,69	0,0000
High	P51880	6	Fabp7	Fatty acid-binding protein, brain OS=Mus musculus OX=10090 GN=Fabp7 PE=1 SV=2	2,67	0,0000
High	E9PV24	12	Fga	Fibrinogen alpha chain OS=Mus musculus OX=10090 GN=Fga PE=1 SV=1	2,53	0,0000
High	Q61838	36	Pzp	Pregnancy zone protein OS=Mus musculus OX=10090 GN=Pzp PE=1 SV=3	2,51	0,0000
High	Q8K0E8	16	Fgb	Fibrinogen beta chain OS=Mus musculus OX=10090 GN=Fgb PE=1 SV=1	2,48	0,0000
High	O70200	3	Aif1	Allograft inflammatory factor 1 OS=Mus musculus OX=10090 GN=Aif1 PE=1 SV=1	2,33	0,0000
High	Q61147	21	Cp	Ceruloplasmin OS=Mus musculus OX=10090 GN=Cp PE=1 SV=2	2,29	0,0000
High	Q00623	11	Apoa1	Apolipoprotein A-I OS=Mus musculus OX=10090 GN=Apoa1 PE=1 SV=2	2,29	0,0000
High	Q9CYL5	2	Glipr2	Golgi-associated plant pathogenesis-related protein 1 OS=Mus musculus OX=10090 GN=Glipr2 PE=1 SV=3	2,29	0,0000
High	P05555	5	Itgam	Integrin alpha-M OS=Mus musculus OX=10090 GN=Itgam PE=1 SV=2	2,21	0,0000
High	Q91VS7	1	Mgst1	Microsomal glutathione S-transferase 1 OS=Mus musculus OX=10090 GN=Mgst1 PE=1 SV=3	2,17	0,0000
High	P01869	6	Ighg1	Ig gamma-1 chain C region, membrane-bound form OS=Mus musculus OX=10090 GN=Ighg1 PE=1 SV=2	2,10	0,0000
High	P07356	14	Anxa2	Annexin A2 OS=Mus musculus OX=10090 GN=Anxa2 PE=1 SV=2	2,10	0,0000
High	P19324	11	Serpinh1	Serpin H1 OS=Mus musculus OX=10090 GN=Serpinh1 PE=1 SV=3	2,07	0,0000
High	Q9ESB3	7	Hrg	Histidine-rich glycoprotein OS=Mus musculus OX=10090 GN=Hrg PE=1 SV=2	2,06	0,0011
Medium	Q8BL80	1	Arhgap22	Rho GTPase-activating protein 22 OS=Mus musculus OX=10090 GN=Arhgap22 PE=1 SV=2	2,03	0,0000
High	P14106	6	C1qb	Complement C1q subcomponent subunit B OS=Mus musculus OX=10090 GN=C1qb PE=1 SV=2	2,02	0,0000
High	Q9WVA4	9	Tagln2	Transgelin-2 OS=Mus musculus OX=10090 GN=Tagln2 PE=1 SV=4	2,01	0,0000
High	P16045	5	Lgals1	Galectin-1 OS=Mus musculus OX=10090 GN=Lgals1 PE=1 SV=3	2,01	0,0000
High	Q9JM63	7	Kcnj10	ATP-sensitive inward rectifier potassium channel 10 OS=Mus musculus OX=10090 GN=Kcnj10 PE=1 SV=1	1,99	0,0000
High	P04186	6	Cfb	Complement factor B OS=Mus musculus OX=10090 GN=Cfb PE=1 SV=2	1,99	0,0006
High	Q8VHL0	2	Slc14a1	Urea transporter 1 OS=Mus musculus OX=10090 GN=Slc14a1 PE=1 SV=2	1,97	0,0001
High	P11835	13	Itgb2	Integrin beta-2 OS=Mus musculus OX=10090 GN=Itgb2 PE=1 SV=2	1,91	0,0001
High	P98086	5	C1qa	Complement C1q subcomponent subunit A OS=Mus musculus OX=10090 GN=C1qa PE=1 SV=2	1,88	0,0000
High	P56695	18	Wfs1	Wolframin OS=Mus musculus OX=10090 GN=Wfs1 PE=1 SV=1	1,86	0,0000
Medium	O35566	2	Cd151	CD151 antigen OS=Mus musculus OX=10090 GN=Cd151 PE=1 SV=2	1,85	0,0007
High	P42225	8	Stat1	Signal transducer and activator of transcription 1 OS=Mus musculus OX=10090 GN=Stat1 PE=1 SV=1	1,85	0,0005
High	Q61233	21	Lcp1	Plastin-2 OS=Mus musculus OX=10090 GN=Lcp1 PE=1 SV=4	1,81	0,0003
High	Q00897	4	Serpina1d	Alpha-1-antitrypsin 1-4 OS=Mus musculus OX=10090 GN=Serpina1d PE=1 SV=1	1,81	0,0001
High	Q9JHR7	24	Ide	Insulin-degrading enzyme OS=Mus musculus OX=10090 GN=Ide PE=1 SV=1	1,78	0,0004
High	Q8R2Y2	2	Mcam	Cell surface glycoprotein MUC18 OS=Mus musculus OX=10090 GN=Mcam PE=1 SV=1	1,77	0,0044
Medium	Q9JF0	1	Nap1I5	Nucleosome assembly protein 1-like 5 OS=Mus musculus OX=10090 GN=Nap1I5 PE=1 SV=1	1,76	0,0001
High	Q61599	4	Arhgdib	Rho GDP-dissociation inhibitor 2 OS=Mus musculus OX=10090 GN=Arhgdib PE=1 SV=3	1,76	0,0020
High	Q91VW3	2	Sh3bgrl3	SH3 domain-binding glutamic acid-rich-like protein 3 OS=Mus musculus OX=10090 GN=Sh3bgrl3 PE=1 SV=1	1,76	0,0008
High	Q9Z1Q5	9	Clic1	Chloride intracellular channel protein 1 OS=Mus musculus OX=10090 GN=Clic1 PE=1 SV=3	1,76	0,0003

High	Q02105	5	C1qc	Complement C1q subcomponent subunit C OS=Mus musculus OX=10090 GN=C1qc PE=1 SV=2	1,72	0,0002
High	P06909	16	Cfh	Complement factor H OS=Mus musculus OX=10090 GN=Cfh PE=1 SV=2	1,66	0,0061
High	Q61739	10	Itga6	Integrin alpha-6 OS=Mus musculus OX=10090 GN=Itga6 PE=1 SV=3	1,65	0,0022
High	P62960	5	Ybx1	Y-box-binding protein 1 OS=Mus musculus OX=10090 GN=Ybx1 PE=1 SV=3	1,64	0,0006
High	Q9WVA2	2	Timm8a1	Mitochondrial import inner membrane translocase subunit Tim8 A OS=Mus musculus OX=10090 GN=Timm8a1 PE=1 SV=1	1,63	0,0298
High	P52503	5	Ndufs6	NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Mus musculus OX=10090 GN=Ndufs6 PE=1 SV=2	1,62	0,0023
High	P11276	10	Fn1	Fibronectin OS=Mus musculus OX=10090 GN=Fn1 PE=1 SV=4	1,61	0,0053
High	Q9DCJ9	6	Npl	N-acetylneuraminatase lyase OS=Mus musculus OX=10090 GN=Npl PE=1 SV=1	1,60	0,0092
High	P51910	2	Apod	Apolipoprotein D OS=Mus musculus OX=10090 GN=Apod PE=1 SV=1	1,60	0,0091
High	Q99KR3	3	Lactb2	Endoribonuclease LACTB2 OS=Mus musculus OX=10090 GN=Lactb2 PE=1 SV=1	1,59	0,0093
Medium	O54890	1	Itgb3	Integrin beta-3 OS=Mus musculus OX=10090 GN=Itgb3 PE=1 SV=2	1,59	0,0020
High	P42227	6	Stat3	Signal transducer and activator of transcription 3 OS=Mus musculus OX=10090 GN=Stat3 PE=1 SV=2	1,57	0,0235
High	Q06890	13	Clu	Clusterin OS=Mus musculus OX=10090 GN=Clu PE=1 SV=1	1,56	0,0031
High	O70340	8	Nptx2	Neuronal pentraxin-2 OS=Mus musculus OX=10090 GN=Nptx2 PE=2 SV=1	1,56	0,0021
High	Q8BTM8	35	Flna	Filamin-A OS=Mus musculus OX=10090 GN=Flna PE=1 SV=5	1,56	0,0073
High	Q05816	9	Fabp5	Fatty acid-binding protein 5 OS=Mus musculus OX=10090 GN=Fabp5 PE=1 SV=3	1,56	0,0080
High	Q9WV32	8	Arcp1b	Actin-related protein 2/3 complex subunit 1B OS=Mus musculus OX=10090 GN=Arcp1b PE=1 SV=4	1,55	0,0078
High	Q61FZ6	5	Krt77	Keratin, type II cytoskeletal 1b OS=Mus musculus OX=10090 GN=Krt77 PE=1 SV=1	1,54	0,0020
High	Q91Z38	3	Ttc1	Tetratricopeptide repeat protein 1 OS=Mus musculus OX=10090 GN=Ttc1 PE=1 SV=1	1,54	0,0119
Medium	P20491	1	Fcer1g	High affinity immunoglobulin epsilon receptor subunit gamma OS=Mus musculus OX=10090 GN=Fcer1g PE=1 SV=1	1,54	0,0106
High	Q9DCT8	4	Crip2	Cysteine-rich protein 2 OS=Mus musculus OX=10090 GN=Crip2 PE=1 SV=1	1,54	0,0048
High	A3KMP2	3	Ttc38	Tetratricopeptide repeat protein 38 OS=Mus musculus OX=10090 GN=Ttc38 PE=1 SV=2	1,53	0,0165
High	Q80WW9	1	Ddrgk1	DDRKG domain-containing protein 1 OS=Mus musculus OX=10090 GN=Ddrgk1 PE=1 SV=2	1,53	0,0182
Medium	Q9EST4	2	Psmg2	Proteasome assembly chaperone 2 OS=Mus musculus OX=10090 GN=Psmg2 PE=1 SV=1	1,52	0,0031
High	O89106	3	Fhit	Bis(5'-adenosyl)-triphosphatase OS=Mus musculus OX=10090 GN=Fhit PE=1 SV=3	1,51	0,0213
High	P01872	11	Ighm	Immunoglobulin heavy constant mu OS=Mus musculus OX=10090 GN=Ighm PE=1 SV=2	1,51	0,0066
High	P70689	2	Gjb6	Gap junction beta-6 protein OS=Mus musculus OX=10090 GN=Gjb6 PE=1 SV=1	1,51	0,0154
High	P08228	7	Sod1	Superoxide dismutase [Cu-Zn] OS=Mus musculus OX=10090 GN=Sod1 PE=1 SV=2	1,50	0,0051
High	O55137	5	Acot1	Acyl-coenzyme A thioesterase 1 OS=Mus musculus OX=10090 GN=Acot1 PE=1 SV=1	1,50	0,0111
High	P47867	4	Scg3	Secretogranin-3 OS=Mus musculus OX=10090 GN=Scg3 PE=1 SV=1	1,50	0,0058
High	Q9DC50	6	Crot	Peroxisomal carnitine O-octanoyltransferase OS=Mus musculus OX=10090 GN=Crot PE=1 SV=1	1,49	0,0203
High	P97352	4	S100a13	Protein S100-A13 OS=Mus musculus OX=10090 GN=S100a13 PE=1 SV=1	1,49	0,0055
High	Q62178	9	Sema4a	Semaphorin-4A OS=Mus musculus OX=10090 GN=Sema4a PE=1 SV=2	1,48	0,0073

High	Q9DC11	2	Plxdc2	Plexin domain-containing protein 2 OS=Mus musculus OX=10090 GN=Plxdc2 PE=1 SV=1	1,48	0,0267
High	Q9JL62	9	Gltp	Glycolipid transfer protein OS=Mus musculus OX=10090 GN=Gltp PE=1 SV=3	1,47	0,0112
High	P62313	3	Lsm6	U6 snRNA-associated Sm-like protein LSm6 OS=Mus musculus OX=10090 GN=Lsm6 PE=1 SV=1	1,47	0,0154
High	P97449	16	Anpep	Aminopeptidase N OS=Mus musculus OX=10090 GN=Anpep PE=1 SV=4	1,47	0,0217
High	P21981	13	Tgm2	Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus OX=10090 GN=Tgm2 PE=1 SV=4	1,47	0,0099
High	Q8R332	2	Nup58	Nucleoporin p58/p45 OS=Mus musculus OX=10090 GN=Nup58 PE=1 SV=1	1,47	0,0249
High	O89017	4	Lgmn	Legumain OS=Mus musculus OX=10090 GN=Lgmn PE=1 SV=1	1,47	0,0076
High	Q60829	8	Ppp1r1b	Protein phosphatase 1 regulatory subunit 1B OS=Mus musculus OX=10090 GN=Ppp1r1b PE=1 SV=2	1,46	0,0053
High	Q9WVT6	3	Ca14	Carbonic anhydrase 14 OS=Mus musculus OX=10090 GN=Ca14 PE=1 SV=1	1,46	0,0239
High	Q9ERT9	4	Ppp1r1a	Protein phosphatase 1 regulatory subunit 1A OS=Mus musculus OX=10090 GN=Ppp1r1a PE=1 SV=1	1,45	0,0294
High	Q3UV17	1	Krt76	Keratin, type II cytoskeletal 2 oral OS=Mus musculus OX=10090 GN=Krt76 PE=1 SV=1	1,45	0,0137
High	O35639	17	Anxa3	Annexin A3 OS=Mus musculus OX=10090 GN=Anxa3 PE=1 SV=4	1,44	0,0095
High	O08739	7	Ampd3	AMP deaminase 3 OS=Mus musculus OX=10090 GN=Ampd3 PE=1 SV=2	1,44	0,0309
High	Q8K4M5	3	Commd1	COMM domain-containing protein 1 OS=Mus musculus OX=10090 GN=Commd1 PE=1 SV=2	1,44	0,0212
High	P11438	4	Lamp1	Lysosome-associated membrane glycoprotein 1 OS=Mus musculus OX=10090 GN=Lamp1 PE=1 SV=2	1,44	0,0229
High	Q8VCI5	5	Pex19	Peroxisomal biogenesis factor 19 OS=Mus musculus OX=10090 GN=Pex19 PE=1 SV=1	1,44	0,0460
Medium	Q9R1Q7	1	Plp2	Proteolipid protein 2 OS=Mus musculus OX=10090 GN=Plp2 PE=1 SV=1	1,44	0,0308
High	Q9D708	2	S100a16	Protein S100-A16 OS=Mus musculus OX=10090 GN=S100a16 PE=1 SV=1	1,43	0,0131
High	Q61792	6	Lasp1	LIM and SH3 domain protein 1 OS=Mus musculus OX=10090 GN=Lasp1 PE=1 SV=1	1,43	0,0104
High	Q02819	13	Nucb1	Nucleobindin-1 OS=Mus musculus OX=10090 GN=Nucb1 PE=1 SV=2	1,43	0,0379
High	Q8BHK2	3	Scn3b	Sodium channel subunit beta-3 OS=Mus musculus OX=10090 GN=Scn3b PE=1 SV=1	1,42	0,0348
High	Q8C3X2	2	Ccdc90b	Coiled-coil domain-containing protein 90B, mitochondrial OS=Mus musculus OX=10090 GN=Ccdc90b PE=1 SV=1	1,42	0,0344
High	P57716	12	Ncstn	Nicastrin OS=Mus musculus OX=10090 GN=Ncstn PE=1 SV=3	1,42	0,0200
High	Q91W90	9	Txndc5	Thioredoxin domain-containing protein 5 OS=Mus musculus OX=10090 GN=Txndc5 PE=1 SV=2	1,42	0,0375
High	P63300	3	Selenow	Selenoprotein W OS=Mus musculus OX=10090 GN=Selenow PE=1 SV=3	1,41	0,0425
High	Q9CR20	1	Ier3ip1	Immediate early response 3-interacting protein 1 OS=Mus musculus OX=10090 GN=Ier3ip1 PE=3 SV=1	1,41	0,0382
High	P21460	5	Cst3	Cystatin-C OS=Mus musculus OX=10090 GN=Cst3 PE=1 SV=2	1,41	0,0262
High	P01942	8	Hba	Hemoglobin subunit alpha OS=Mus musculus OX=10090 GN=Hba PE=1 SV=2	1,41	0,0217
High	Q99JF5	4	Mvd	Diphosphomevalonate decarboxylase OS=Mus musculus OX=10090 GN=Mvd PE=1 SV=2	1,41	0,0494
High	Q03517	3	Scg2	Secretogranin-2 OS=Mus musculus OX=10090 GN=Scg2 PE=1 SV=1	1,41	0,0455
High	P17047	5	Lamp2	Lysosome-associated membrane glycoprotein 2 OS=Mus musculus OX=10090 GN=Lamp2 PE=1 SV=2	1,41	0,0271
High	P10605	14	Ctsb	Cathepsin B OS=Mus musculus OX=10090 GN=Ctsb PE=1 SV=2	1,41	0,0264
High	P09055	21	Itgb1	Integrin beta-1 OS=Mus musculus OX=10090 GN=Itgb1 PE=1 SV=1	1,41	0,0142
High	Q00915	9	Rbp1	Retinol-binding protein 1 OS=Mus musculus OX=10090 GN=Rbp1 PE=1 SV=2	1,41	0,0283
High	P13020	22	Gsn	Gelsolin OS=Mus musculus OX=10090 GN=Gsn PE=1 SV=3	1,41	0,0276

High	P18242	13	Ctsd	Cathepsin D OS=Mus musculus OX=10090 GN=Ctsd PE=1 SV=1	1,41	0,0237
High	P16675	7	Ctsa	Lysosomal protective protein OS=Mus musculus OX=10090 GN=Ctsa PE=1 SV=1	1,40	0,0154
High	P17439	6	Gba	Lysosomal acid glucosylceramidase OS=Mus musculus OX=10090 GN=Gba PE=1 SV=1	1,40	0,0472
High	P20060	20	Hexb	Beta-hexosaminidase subunit beta OS=Mus musculus OX=10090 GN=Hexb PE=1 SV=2	1,40	0,0275
High	Q00493	18	Cpe	Carboxypeptidase E OS=Mus musculus OX=10090 GN=Cpe PE=1 SV=2	1,40	0,0188
High	O35114	10	Scarb2	Lysosome membrane protein 2 OS=Mus musculus OX=10090 GN=Scarb2 PE=1 SV=3	1,40	0,0107
High	Q9ER00	12	Stx12	Syntaxin-12 OS=Mus musculus OX=10090 GN=Stx12 PE=1 SV=1	1,39	0,0428
High	Q99J85	15	Nptxr	Neuronal pentraxin receptor OS=Mus musculus OX=10090 GN=Nptxr PE=1 SV=1	1,39	0,0447
High	Q91WE4	1		UPF0729 protein C18orf32 homolog OS=Mus musculus OX=10090 PE=3 SV=1	1,39	0,0476
High	P97371	11	Psme1	Proteasome activator complex subunit 1 OS=Mus musculus OX=10090 GN=Psme1 PE=1 SV=2	1,39	0,0303
High	Q922J6	3	Tspan2	Tetraspanin-2 OS=Mus musculus OX=10090 GN=Tspan2 PE=1 SV=1	1,39	0,0275
High	Q6P5H2	3	Nes	Nestin OS=Mus musculus OX=10090 GN=Nes PE=1 SV=1	1,38	0,0332
High	P04104	4	Krt1	Keratin, type II cytoskeletal 1 OS=Mus musculus OX=10090 GN=Krt1 PE=1 SV=4	1,38	0,0331
High	P56812	6	Pdcd5	Programmed cell death protein 5 OS=Mus musculus OX=10090 GN=Pdcd5 PE=1 SV=3	1,38	0,0383
High	P09103	23	P4hb	Protein disulfide-isomerase OS=Mus musculus OX=10090 GN=P4hb PE=1 SV=2	1,38	0,0390
High	P81122	2	Irs2	Insulin receptor substrate 2 OS=Mus musculus OX=10090 GN=Irs2 PE=1 SV=2	1,37	0,0366
High	Q60766	5	Irgm1	Immunity-related GTPase family M protein 1 OS=Mus musculus OX=10090 GN=Irgm1 PE=1 SV=1	1,37	0,0451
High	P97450	4	Atp5pf	ATP synthase-coupling factor 6, mitochondrial OS=Mus musculus OX=10090 GN=Atp5pf PE=1 SV=1	1,37	0,0284
High	Q6IME9	2	Krt72	Keratin, type II cytoskeletal 72 OS=Mus musculus OX=10090 GN=Krt72 PE=3 SV=1	1,37	0,0424
High	Q9DBS1	11	Tmem43	Transmembrane protein 43 OS=Mus musculus OX=10090 GN=Tmem43 PE=1 SV=1	1,36	0,0471
High	Q6IRU2	11	Tpm4	Tropomyosin alpha-4 chain OS=Mus musculus OX=10090 GN=Tpm4 PE=1 SV=3	1,36	0,0421
High	P70699	18	Gaa	Lysosomal alpha-glucosidase OS=Mus musculus OX=10090 GN=Gaa PE=1 SV=2	1,35	0,0353
High	Q9R099	8	Tbl2	Transducin beta-like protein 2 OS=Mus musculus OX=10090 GN=Tbl2 PE=1 SV=2	1,35	0,0414
High	Q61646	8	Hp	Haptoglobin OS=Mus musculus OX=10090 GN=Hp PE=1 SV=1	1,35	0,0428
High	O35143	2	ATP5IF1	ATPase inhibitor, mitochondrial OS=Mus musculus OX=10090 GN=ATP5IF1 PE=1 SV=2	1,34	0,0346
High	P28667	4	Marcks1	MARCKS-related protein OS=Mus musculus OX=10090 GN=Marcks1 PE=1 SV=2	1,34	0,0344
High	P58771	9	Tpm1	Tropomyosin alpha-1 chain OS=Mus musculus OX=10090 GN=Tpm1 PE=1 SV=1	1,34	0,0429
High	P51855	15	Gss	Glutathione synthetase OS=Mus musculus OX=10090 GN=Gss PE=1 SV=1	1,33	0,0354
High	Q6ZQJ3	9	Mlec	Malectin OS=Mus musculus OX=10090 GN=Mlec PE=1 SV=2	1,33	0,0429
High	Q9D154	12	Serpib1a	Leukocyte elastase inhibitor A OS=Mus musculus OX=10090 GN=Serpib1a PE=1 SV=1	1,32	0,0465
High	Q3URE9	3	Lingo2	Leucine-rich repeat and immunoglobulin-like domain-containing nogo receptor-interacting protein 2 OS=Mus musculus OX=10090 GN=Lingo2 PE=2 SV=1	1,31	0,0482
High	P20444	14	Prkca	Protein kinase C alpha type OS=Mus musculus OX=10090 GN=Prkca PE=1 SV=3	0,78	0,0409
High	Q6NS60	29	Fbxo41	F-box only protein 41 OS=Mus musculus OX=10090 GN=Fbxo41 PE=1 SV=3	0,77	0,0467
High	Q8CJ40	17	Crocc	Rootletin OS=Mus musculus OX=10090 GN=Crocc PE=1 SV=2	0,77	0,0489
High	P35438	24	Grin1	Glutamate receptor ionotropic, NMDA 1 OS=Mus musculus OX=10090 GN=Grin1 PE=1 SV=1	0,77	0,0421

High	Q6PH08	33	Erc2	ERC protein 2 OS=Mus musculus OX=10090 GN=Erc2 PE=1 SV=2	0,77	0,0436
High	Q01097	43	Grin2b	Glutamate receptor ionotropic, NMDA 2B OS=Mus musculus OX=10090 GN=Grin2b PE=1 SV=3	0,76	0,0335
High	Q8BJ42	33	Dlgap2	Disks large-associated protein 2 OS=Mus musculus OX=10090 GN=Dlgap2 PE=1 SV=2	0,76	0,0323
High	Q8C015	21	Pak5	Serine/threonine-protein kinase PAK 5 OS=Mus musculus OX=10090 GN=Pak5 PE=1 SV=1	0,75	0,0397
High	Q4ACU6	53	Shank3	SH3 and multiple ankyrin repeat domains protein 3 OS=Mus musculus OX=10090 GN=Shank3 PE=1 SV=3	0,75	0,0296
High	P48543	4	Kcnj9	G protein-activated inward rectifier potassium channel 3 OS=Mus musculus OX=10090 GN=Kcnj9 PE=1 SV=2	0,75	0,0334
High	Q5DU25	45	Iqsec2	IQ motif and SEC7 domain-containing protein 2 OS=Mus musculus OX=10090 GN=Iqsec2 PE=1 SV=3	0,74	0,0345
High	Q62108	30	Dlg4	Disks large homolog 4 OS=Mus musculus OX=10090 GN=Dlg4 PE=1 SV=1	0,74	0,0171
High	Q8BIZ1	23	Anks1b	Ankyrin repeat and sterile alpha motif domain-containing protein 1B OS=Mus musculus OX=10090 GN=Anks1b PE=1 SV=3	0,73	0,0148
High	Q3V132	1	Slc25a31	ADP/ATP translocase 4 OS=Mus musculus OX=10090 GN=Slc25a31 PE=1 SV=1	0,73	0,0423
High	Q8BXR9	20	Osbpl6	Oxysterol-binding protein-related protein 6 OS=Mus musculus OX=10090 GN=Osbpl6 PE=1 SV=1	0,72	0,0463
High	Q68EF6	25	Begain	Brain-enriched guanylate kinase-associated protein OS=Mus musculus OX=10090 GN=Begain PE=1 SV=2	0,72	0,0102
High	Q61418	7	Cln4	H(+)/Cl(-) exchange transporter 4 OS=Mus musculus OX=10090 GN=Cln4 PE=2 SV=2	0,72	0,0423
High	Q8VHQ9	22	Acot11	Acyl-coenzyme A thioesterase 11 OS=Mus musculus OX=10090 GN=Acot11 PE=1 SV=1	0,72	0,0330
High	Q9Z0V2	13	Kcnd2	Potassium voltage-gated channel subfamily D member 2 OS=Mus musculus OX=10090 GN=Kcnd2 PE=1 SV=1	0,72	0,0448
High	F6SEU4	51	Syngap1	Ras/Rap GTPase-activating protein SynGAP OS=Mus musculus OX=10090 GN=Syngap1 PE=1 SV=2	0,71	0,0104
Medium	Q8BH61	1	F13a1	Coagulation factor XIII A chain OS=Mus musculus OX=10090 GN=F13a1 PE=1 SV=3	0,71	0,0282
High	D3Z7H4	5	Gsg1l	Germ cell-specific gene 1-like protein OS=Mus musculus OX=10090 GN=Gsg1l PE=1 SV=2	0,71	0,0360
High	P01741	1		Ig heavy chain V region OS=Mus musculus OX=10090 PE=1 SV=1	0,71	0,0264
High	Q99KX1	9	Mlf2	Myeloid leukemia factor 2 OS=Mus musculus OX=10090 GN=Mlf2 PE=1 SV=1	0,70	0,0092
High	Q9D7V2	2	Lysmd2	LysM and putative peptidoglycan-binding domain-containing protein 2 OS=Mus musculus OX=10090 GN=Lysmd2 PE=1 SV=2	0,68	0,0156
High	Q3TKT4	6	Smarca4	Transcription activator BRG1 OS=Mus musculus OX=10090 GN=Smarca4 PE=1 SV=1	0,68	0,0183
High	Q9WU79	19	Prodh	Proline dehydrogenase 1, mitochondrial OS=Mus musculus OX=10090 GN=Prodh PE=1 SV=2	0,65	0,0093
High	Q3UXZ6	17	Fam81a	Protein FAM81A OS=Mus musculus OX=10090 GN=Fam81a PE=1 SV=2	0,65	0,0086
High	Q8BGU2	1	Cbln2	Cerebellin-2 OS=Mus musculus OX=10090 GN=Cbln2 PE=1 SV=1	0,64	0,0477
High	A2A8L1	4	Chd5	Chromodomain-helicase-DNA-binding protein 5 OS=Mus musculus OX=10090 GN=Chd5 PE=1 SV=1	0,63	0,0322
High	Q8VHR5	6	Gatad2b	Transcriptional repressor p66-beta OS=Mus musculus OX=10090 GN=Gatad2b PE=1 SV=1	0,63	0,0322
High	P70268	6	Pkn1	Serine/threonine-protein kinase N1 OS=Mus musculus OX=10090 GN=Pkn1 PE=1 SV=3	0,62	0,0264
High	Q9JJ43	4	Rbfox1	RNA binding protein fox-1 homolog 1 OS=Mus musculus OX=10090 GN=Rbfox1 PE=1 SV=3	0,61	0,0020
Medium	Q9CR76	1	Tmem186	Transmembrane protein 186 OS=Mus musculus OX=10090 GN=Tmem186 PE=1 SV=2	0,59	0,0196
High	P62257	2	Ube2h	Ubiquitin-conjugating enzyme E2 H OS=Mus musculus OX=10090 GN=Ube2h PE=1 SV=1	0,56	0,0017
High	Q8BGF9	5	Slc25a44	Solute carrier family 25 member 44 OS=Mus musculus OX=10090 GN=Slc25a44 PE=1 SV=1	0,56	0,0021
High	Q8K0F1	4	Tbc1d23	TBC1 domain family member 23 OS=Mus musculus OX=10090 GN=Tbc1d23 PE=1 SV=1	0,52	0,0009
High	Q9Z0S6	2	Cldn10	Claudin-10 OS=Mus musculus OX=10090 GN=Cldn10 PE=1 SV=2	0,52	0,0005

High	Q8BNN1	3	Spata2l	Spermatogenesis-associated protein 2-like protein OS=Mus musculus OX=10090 GN=Spata2l PE=1 SV=1	0,50	0,0001
High	Q8C4G9	2	Adgra1	Adhesion G protein-coupled receptor A1 OS=Mus musculus OX=10090 GN=Adgra1 PE=2 SV=1	0,49	0,0002
High	B2RWJ3	2	Tmem240	Transmembrane protein 240 OS=Mus musculus OX=10090 GN=Tmem240 PE=1 SV=1	0,46	0,0000
High	O35316	2	Slc6a6	Sodium- and chloride-dependent taurine transporter OS=Mus musculus OX=10090 GN=Slc6a6 PE=1 SV=2	0,41	0,0000
Medium	Q8CIH5	1	Plcg2	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase gamma-2 OS=Mus musculus OX=10090 GN=Plcg2 PE=1 SV=1	0,39	0,0003
High	O35723	2	Dnajb3	DnaJ homolog subfamily B member 3 OS=Mus musculus OX=10090 GN=Dnajb3 PE=2 SV=1	0,34	0,0000
High	P97770	2	Thumpd3	THUMP domain-containing protein 3 OS=Mus musculus OX=10090 GN=Thumpd3 PE=1 SV=1	0,29	0,0000
Medium	Q9D384	1	Snrnp35	U11/U12 small nuclear ribonucleoprotein 35 kDa protein OS=Mus musculus OX=10090 GN=Snrnp35 PE=2 SV=1	0,26	0,0000
High	Q8C0Q9	1	Rapgef5	Rap guanine nucleotide exchange factor 5 OS=Mus musculus OX=10090 GN=Rapgef5 PE=2 SV=2	0,20	0,0000
High	Q60632	1	Nr2f1	COUP transcription factor 1 OS=Mus musculus OX=10090 GN=Nr2f1 PE=2 SV=2	0,10	0,0000
High	Q61738	2	Itga7	Integrin alpha-7 OS=Mus musculus OX=10090 GN=Itga7 PE=1 SV=3	0,05	0,0000
High	Q91WK5	2	Gcsh	Glycine cleavage system H protein, mitochondrial OS=Mus musculus OX=10090 GN=Gcsh PE=1 SV=2	0,01	0,0000
High	P97760	2	Polr2c	DNA-directed RNA polymerase II subunit RPB3 OS=Mus musculus OX=10090 GN=Polr2c PE=1 SV=2	0,01	0,0000
High	O70456	2	Sfn	14-3-3 protein sigma OS=Mus musculus OX=10090 GN=Sfn PE=1 SV=2	0,01	0,0000
Medium	Q9CWR2	1	Smyd3	Histone-lysine N-methyltransferase SMYD3 OS=Mus musculus OX=10090 GN=Smyd3 PE=2 SV=1	0,01	0,0000