

Proteins	Accession Number	Molecular Weight	Mann Whitney Test (Bonferroni Corrected)	Average Coefficient of Variance	Δ43 Average	A57G Average	Δ43/A57G	A57G/Δ43
Elongation factor 2 OS=Mus musculus GN=Eef2 PE=1 SV=2	EF2_MOUSE	95 kDa	< 0.0001	20.842738	0.9183	1.06286667	0.863984194	1.157428582
Complement C3 OS=Mus musculus GN=C3 PE=1 SV=3	CO3_MOUSE	186 kDa	< 0.0001	21.553332	0.958566667	1.10473333	0.867690544	1.152484612
Aldehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh2 PEALDH2_MOUSE		57 kDa	< 0.0001	23.078518	0.843066667	1.2276	0.686760074	1.456112605
Heat shock protein HSP 90-beta OS=Mus musculus GN=Hsp90ab1 PE=1 SHS90B_MOUSE		83 kDa	< 0.0001	22.325896	0.931566667	1.0626	0.876686116	1.140659105
Leucine-rich PPR motif-containing protein, mitochondrial OS=Mus musculus PPRC_MOUSE		157 kDa	< 0.0001	21.971369	0.839733333	0.9806	0.856346455	1.167751667
NAD(P) transhydrogenase, mitochondrial OS=Mus musculus GN=Nnt PE=NNTM_MOUSE		114 kDa	< 0.0001	26.87348	0.8075	1.18143333	0.683491804	1.463075335
Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial OS=Mus AL4A1_MOUSE		62 kDa	< 0.0001	21.019696	0.690833333	0.8632	0.800316651	1.249505428
Stress-induced-phosphoprotein 1 OS=Mus musculus GN=Stip1 PE=1 SV=1 STIP1_MOUSE		63 kDa	< 0.0001	19.943316	0.8862	1.0147	0.873361585	1.145001128
Cluster of Troponin T, cardiac muscle OS=Mus musculus GN=Tnnt2 PE=4 K3W4R6_MOUSE [3]		35 kDa	0.00018	26.36347	0.9561	0.84206667	1.13542079	0.880730746
Mitochondrial 2-oxoglutarate/malate carrier protein OS=Mus musculus M2OM_MOUSE		34 kDa	0.0017	22.087899	1.047033333	1.35216667	0.774337483	1.29142657
Alpha-1B-glycoprotein OS=Mus musculus GN=A1bg PE=1 SV=1 A1BG_MOUSE		57 kDa	0.0018	22.157027	1.293566667	0.92943333	1.391779938	0.718504394
Elongation factor 1-gamma OS=Mus musculus GN=Eef1g PE=1 SV=3 EF1G_MOUSE		50 kDa	0.0025	20.053983	0.935866667	1.0707	0.874069923	1.14407323
Cluster of Myosin-6 OS=Mus musculus GN=Myh6 PE=1 SV=2 (MYH6_MOUSE) [10]		224 kDa	0.0039	51.96263	0.974866667	0.88096667	1.106587461	0.903679136
Succinyl-CoA ligase [GDP-forming] subunit beta, mitochondrial OS=Mus musculus SUCB2_MOUSE		47 kDa	0.0068	21.554866	1.041333333	1.18976667	0.875241644	1.142541613
Cluster of Inorganic pyrophosphatase 2, mitochondrial OS=Mus musculus D3Z636_MOUSE [2]		38 kDa	0.0069	21.219736	0.954866667	1.1961	0.798316752	1.252635621
Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondon CHCH3_MOUSE		26 kDa	0.0083	25.52083	1.105033333	0.9405	1.174942407	0.851105547
Cluster of Myosin regulatory light chain 2, ventricular/cardiac muscle iso MLRV_MOUSE		19 kDa	0.011	36.96808	0.857633333	0.79613333	1.077248367	0.928291033
Histidine rich calcium binding protein, isoform CRA_a OS=Mus musculus G5E8J6_MOUSE		85 kDa	0.017	27.835104	0.930766667	1.0272	0.906120197	1.103606346
Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3 CLH1_MOUSE		192 kDa	0.017	21.689476	0.930766667	0.92543333	1.249540756	0.800294024
Sarcalumenin OS=Mus musculus GN=Srl PE=1 SV=1 SRCA_MOUSE		99 kDa	0.02	25.064	1.007733333	1.12463333	0.896055011	1.116002911
Cytochrome c oxidase subunit 5B, mitochondrial OS=Mus musculus GN= COX5B_MOUSE		14 kDa	0.02	25.410923	0.9496	0.85186667	1.11472844	0.897079472
Serpin H1 OS=Mus musculus GN=Serpinh1 PE=1 SV=3 SERPH_MOUSE		47 kDa	0.021	20.842558	0.882433333	1.01086667	0.872947306	1.145544517
Calcium-binding mitochondrial carrier protein Aralar1 OS=Mus musculus CMC1_MOUSE		75 kDa	0.024	20.755787	1.032033333	1.15716667	0.891862307	1.121249314
Short-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus ACADS_MOUSE		45 kDa	0.031	23.044616	0.988366667	1.13993333	0.867039008	1.153350646
Heterogeneous nuclear ribonucleoprotein U OS=Mus musculus GN=Hnrr G3XA10_MOUSE (+1)		87 kDa	0.038	20.39134	0.942833333	1.12273333	0.839766047	1.190807849
Cluster of Long-chain-fatty-acid-CoA ligase 1 OS=Mus musculus GN=Acs ACSL1_MOUSE [4]		78 kDa	0.039	24.848925	1.1165	1.16303333	0.954830759	1.047306018
Citrate lyase subunit beta-like protein, mitochondrial OS=Mus musculus CLYBL_MOUSE		38 kDa	0.042	21.492629	0.9143	1.0197	0.896636266	1.115279449
Cytochrome b-c1 complex subunit 7 OS=Mus musculus GN=Uqcrc PE=3 Q9CQB4_MOUSE (+1)		14 kDa	0.046	22.793165	1.111733333	0.96373333	1.153569452	0.86687455
Isovaleryl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=lvd IVD_MOUSE		46 kDa	0.048	22.403307	0.969633333	1.07926667	0.898418679	1.113066795
NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial E9QPX3_MOUSE (+1)		20 kDa	0.048	28.701907	1.1239	0.94726667	1.186466324	0.842838924
Protein Ogdhl OS=Mus musculus GN=Ogdhl PE=2 SV=1 E9Q7L0_MOUSE		117 kDa	0.052	23.886847	0.976033333	0.68893333	1.416731179	0.70585021
Cluster of Creatine kinase S-type, mitochondrial OS=Mus musculus GN=C KCRS_MOUSE		47 kDa	0.055	26.69867	1.0882	0.99706667	1.091401444	0.91625314
Nidogen-1 OS=Mus musculus GN=Nid1 PE=1 SV=2 NID1_MOUSE		137 kDa	0.056	21.271154	1.133333333	1	1.133333333	0.882352941
Cluster of Aspartate aminotransferase, cytoplasmic OS=Mus musculus G AATC_MOUSE [2]		46 kDa	0.06	21.676573	0.933333333	1.06666667	0.875	1.142857143
Cluster of 6-phosphofructokinase, muscle type OS=Mus musculus GN=Pf K6PF_MOUSE		85 kDa	0.062	22.191806	1.033333333	1.2	0.861111111	1.161290323
Cytoplasmic dynein 1 heavy chain 1 OS=Mus musculus GN=Dync1h1 PE= DYHC1_MOUSE		532 kDa	0.067	20.347059	0.9722	1.03333333	0.935483871	1.068965517
Isocitrate dehydrogenase [NADP], mitochondrial OS=Mus musculus GN= IDHP_MOUSE		51 kDa	0.068	25.900894	0.966666667	1.1	0.878787879	1.137931034
Transgelin OS=Mus musculus GN=Tagln PE=1 SV=3 TAGL_MOUSE		23 kDa	0.08	22.099005	0.9	0.76666667	1.173913043	0.851851852
Cluster of Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PEK1C10_HUMAN [11]		60 kDa	0.09	34.07775	0.8	1.63333333	0.489795918	2.041666667
Annxin A5 OS=Mus musculus GN=Anxa5 PE=1 SV=1 ANXA5_MOUSE		36 kDa	0.11	23.054731	0.8086	0.93333333	0.857142857	1.166666667
Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Mus musculus G PPIF_MOUSE		22 kDa	0.11	25.33626	1.033333333	0.93333333	1.107142857	0.903225806
Fumarate hydratase, mitochondrial OS=Mus musculus GN=Fh PE=1 SV=3 FUMH_MOUSE (+1)		54 kDa	0.13	24.246356	1.1	1.2	0.916666667	1.090909091
L-lactate dehydrogenase B chain OS=Mus musculus GN=Ldhb PE=1 SV=2 LDHB_MOUSE		37 kDa	0.15	22.569285	1.1579	1.1	1.060606061	0.942857143
40S ribosomal protein S11 OS=Mus musculus GN=Rps11 PE=2 SV=3 RS11_MOUSE		18 kDa	0.15	19.645563	0.933333333	1.1	0.848484848	1.178571429
60S ribosomal protein L7 OS=Mus musculus GN=Rpl7 PE=2 SV=2 RL7_MOUSE		31 kDa	0.17	20.453222	1	1.06666667	0.9375	1.066666667
Dehydrogenase/reductase SDR family member 4 OS=Mus musculus GN= DHRS4_MOUSE		30 kDa	0.18	21.563943	1.033333333	1.26666667	0.815789474	1.225806452
Profilin-1 OS=Mus musculus GN=Pfn1 PE=1 SV=2 PROF1_MOUSE		15 kDa	0.18	22.549726	0.766666667	1.03333333	0.741935484	1.347826087
BAG family molecular chaperone regulator 3 OS=Mus musculus GN=Bag: BAG3_MOUSE		62 kDa	0.19	22.774442	0.866666667	0.76666667	1.130434783	0.884615385
Ryanodine receptor 2 OS=Mus musculus GN=Ryr2 PE=1 SV=1 RYR2_MOUSE		565 kDa	0.2	22.876436	1.152133333	1.23333333	0.945945946	1.057142857
Lamin-B1 OS=Mus musculus GN=Lmnb1 PE=1 SV=3 LMNB1_MOUSE		67 kDa	0.22	21.16764	0.966666667	1	0.966666667	1.034482759
Histone H4 OS=Mus musculus GN=Hist1h4a PE=1 SV=2 H4_MOUSE		11 kDa	0.25	23.792167	0.933333333	1.16666667	0.8	1.25

NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitocNDUA9_MOUSE	43 kDa	0.29	21.757936	1	1.13333333	0.882352941	1.13333333
Voltage-dependent anion-selective channel protein 1 OS=Mus musculus VDAC1_MOUSE	32 kDa	0.33	24.582472	1.03333333	1.2	0.861111111	1.161290323
SET and MYND domain-containing protein 1 OS=Mus musculus GN=SmyrG5E8R7_MOUSE (+1)	53 kDa	0.37	21.227764	0.9	1	0.9	1.111111111
ATP synthase subunit alpha, mitochondrial OS=Mus musculus GN=Atp5a ATPA_MOUSE	60 kDa	0.38	27.822062	1	1	1	1
Cluster of cAMP-dependent protein kinase type I-alpha regulatory subunit KAPO_MOUSE	43 kDa	0.53	20.9113	0.83333333	1	0.83333333	1.2
NADP-dependent malic enzyme, mitochondrial OS=Mus musculus GN=MMAON_MOUSE	67 kDa	0.55	20.798239	1.13333333	1	1.13333333	0.882352941
Carnitine O-acetyltransferase OS=Mus musculus GN=Crat PE=1 SV=3 CACP_MOUSE (+1)	71 kDa	0.64	21.854895	1.13333333	1.23333333	0.918918919	1.088235294
ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b ATPB_MOUSE	56 kDa	0.66	28.71563	1.1	1	1.1	0.909090909
60S ribosomal protein L34 OS=Mus musculus GN=Rpl34 PE=3 SV=2 RL34_MOUSE	13 kDa	0.71	19.921245	0.966666667	1.13333333	0.852941176	1.172413793
Hsc70-interacting protein OS=Mus musculus GN=St13 PE=2 SV=1 F10A1_MOUSE	42 kDa	0.72	23.409216	0.83333333	0.9	0.925925926	1.08
Protein Ahnak OS=Mus musculus GN=Ahnak PE=2 SV=1 E9Q616_MOUSE	604 kDa	0.94	20.490229	0.93333333	0.9	1.037037037	0.964285714
Cluster of Isoform 2 of Titin OS=Mus musculus GN=Ttn (sp A2ASS6-2 T11sp A2ASS6-2 TITIN_MOUSE [5	?	1	67.696446	1.03333333	0.96666667	1.068965517	0.935483871
Cluster of Actin, alpha cardiac muscle 1 OS=Mus musculus GN=Actc1 PE=ACTC_MOUSE [5]	42 kDa	1	245.10705	0.966666667	0.96666667	1	1
Cluster of Serum albumin OS=Mus musculus GN=Alb PE=1 SV=3 (ALBU_NALBU_MOUSE [2]	69 kDa	1	27.63054	0.83333333	0.8	1.041666667	0.96
Cluster of Tropomyosin 1, alpha, isoform CRA_c OS=Mus musculus GN=TQ8BP43_MOUSE [10]	33 kDa	1	282.96332	1	0.83333333	1.2	0.83333333
Cluster of Beta-globin OS=Mus musculus GN=Hbb-b1 PE=3 SV=1 (A8DUK A8DUK4_MOUSE [4]	16 kDa	1	55.050486	1.066666667	1.2	0.888888889	1.125
Cluster of Myoglobin OS=Mus musculus GN=Mb PE=1 SV=3 (MYG_MOUSE;MYG_MOUSE	17 kDa	1	34.685573	0.9	0.86666667	1.038461538	0.962962963
Cluster of Hemoglobin subunit alpha OS=Mus musculus GN=Hba PE=1 SV=HBA_MOUSE [3]	15 kDa	1	33.199462	1	1.1	0.909090909	1.1
Aconitate hydratase, mitochondrial OS=Mus musculus GN=Aco2 PE=1 SV=ACON_MOUSE	85 kDa	1	26.961797	1.03333333	1.03333333	1	1
Myosin-binding protein C, cardiac-type OS=Mus musculus GN=Mybpc3 PE9Q9T8_MOUSE (+1)	141 kDa	1	24.626884	1	1	1	1
Cluster of Isoform 2 of Sarcoplasmic/endoplasmic reticulum calcium ATP sp O55143-2 AT2A2_MOUSE [?	1	42.964727	1.2	1.26666667	0.947368421	1.055555556
Cluster of Myosin light chain 3 OS=Mus musculus GN=Myl3 PE=1 SV=4 (NMYL3_MOUSE [3]	22 kDa	1	52.16608	0.9	0.96666667	0.931034483	1.074074074
Cluster of Fatty acid-binding protein, heart OS=Mus musculus GN=Fabp3 FABPH_MOUSE [3]	15 kDa	1	89.24998	0.93333333	0.9	1.037037037	0.964285714
Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE= MDHM_MOUSE	36 kDa	1	26.973465	1.03333333	1.06666667	0.96875	1.032258065
Cluster of Alpha-actinin-2 OS=Mus musculus GN=Actn2 PE=1 SV=2 (ACTNACTN2_MOUSE [5]	104 kDa	1	22.692557	1.03333333	1.1	0.939393939	1.064516129
Myomesin-1 OS=Mus musculus GN=Myom1 PE=1 SV=2 MYOM1_MOUSE	185 kDa	1	22.959904	0.9	0.9	1	1
Myomesin 2 OS=Mus musculus GN=Myom2 PE=2 SV=1 Q14B15_MOUSE	165 kDa	1	24.28174	0.966666667	1	0.966666667	1.034482759
Very long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mu:ACADV_MOUSE	71 kDa	1	24.606413	1.102866667	1.06666667	1.03125	0.96969697
Trifunctional enzyme subunit alpha, mitochondrial OS=Mus musculus GNECHA_MOUSE	83 kDa	1	24.56266	1.114766667	1.06666667	1.03125	0.96969697
Cluster of ADP/ATP translocase 1 OS=Mus musculus GN=Slc25a4 PE=1 SV=ADT1_MOUSE [2]	33 kDa	1	39.76799	1.03333333	1.16666667	0.885714286	1.129032258
Cluster of Creatine kinase M-type OS=Mus musculus GN=Ckm PE=1 SV=1KCRM_MOUSE [2]	43 kDa	1	33.107814	1.066666667	1	1.066666667	0.9375
Long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus mu:ACADL_MOUSE	48 kDa	1	23.451269	1.066666667	1.03333333	1.032258065	0.96875
Cluster of Glycogen phosphorylase, muscle form OS=Mus musculus GN=PYGM_MOUSE [3]	97 kDa	1	61.175907	1.03333333	1.16666667	0.885714286	1.129032258
Cluster of Glyceraldehyde-3-phosphate dehydrogenase OS=Mus musculus D3Z0Z9_MOUSE [4]	36 kDa	1	229.86999	0.966666667	1.03333333	0.935483871	1.068965517
Malate dehydrogenase, cytoplasmic OS=Mus musculus GN=Mdh1 PE=1 MDHC_MOUSE	37 kDa	1	30.740502	1.03333333	1	1.03333333	0.967741935
Desmin OS=Mus musculus GN=Des PE=1 SV=3 DESM_MOUSE	53 kDa	1	23.111612	0.93333333	0.96666667	0.965517241	1.035714286
NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial OS=Mu:NDUS1_MOUSE	80 kDa	1	23.431645	1.1	1.06666667	1.03125	0.96969697
Desmoplakin OS=Mus musculus GN=Dsp PE=2 SV=1 DESP_MOUSE	333 kDa	1	20.91065	1	1	1	1
Cluster of Citrate synthase, mitochondrial OS=Mus musculus GN=Cs PE=:CISY_MOUSE [2]	52 kDa	1	23.998831	1.1	1.13333333	0.970588235	1.03030303
3-ketoacyl-CoA thiolase, mitochondrial OS=Mus musculus GN=Acaa2 PE=THIM_MOUSE	42 kDa	1	23.681417	1.09133333	1.06666667	1.03125	0.96969697
Cluster of Plectin OS=Mus musculus GN=Plec PE=1 SV=2 (PLEC_MOUSE) PLEC_MOUSE [2]	534 kDa	1	20.144194	0.9	0.93333333	0.964285714	1.037037037
Cluster of Serotransferrin OS=Mus musculus GN=Tf PE=1 SV=1 (TRFE_M(TRFE_MOUSE [4]	77 kDa	1	28.733036	0.766666667	0.76666667	1	1
Electron transfer flavoprotein subunit alpha, mitochondrial OS=Mus musculus ETFA_MOUSE	35 kDa	1	24.426578	1.108266667	1.06666667	1.03125	0.96969697
Troponin I, cardiac muscle OS=Mus musculus GN=Tnni3 PE=1 SV=2 TNNI3_MOUSE	24 kDa	1	33.88926	0.766666667	0.76666667	1	1
Cluster of Fructose-bisphosphate aldolase OS=Mus musculus GN=Aldoa IA6ZI44_MOUSE [3]	45 kDa	1	24.518725	0.966666667	0.93333333	1.035714286	0.965517241
Cluster of Heat shock cognate 71 kDa protein OS=Mus musculus GN=Hsp HSP7C_MOUSE [5]	71 kDa	1	25.833014	0.8449	0.96666667	0.896551724	1.115384615
Trifunctional enzyme subunit beta, mitochondrial OS=Mus musculus GN=ECHB_MOUSE	51 kDa	1	22.729386	1	1.03333333	0.967741935	1.03333333
Cluster of Spectrin alpha chain, non-erythrocytic 1 OS=Mus musculus GNE9Q447_MOUSE [2]	285 kDa	1	47.704542	0.83333333	1.03333333	0.806451613	1.24
Cluster of Obscurin OS=Mus musculus GN=Obscn PE=2 SV=2 (OBSCN_M(OBSCN_MOUSE [3]	966 kDa	1	32.373226	1	1	1	1
Aspartate aminotransferase, mitochondrial OS=Mus musculus GN=Got2 AATM_MOUSE	47 kDa	1	24.026474	1.03333333	1	1.03333333	0.967741935
Stress-70 protein, mitochondrial OS=Mus musculus GN=Hspa9 PE=1 SV= GRP75_MOUSE	73 kDa	1	26.49515	0.8759	0.83333333	1.04	0.961538462
Cluster of Mitochondrial inner membrane protein OS=Mus musculus GN E9Q800_MOUSE [3]	76 kDa	1	19.795075	0.9	0.86666667	1.038461538	0.962962963
Cluster of Nebulette OS=Mus musculus GN=Nebi PE=2 SV=1 (B7ZC12_MCB7ZC12_MOUSE [3]	116 kDa	1	22.83046	0.93333333	0.9	1.037037037	0.964285714

Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondria	DHSA_MOUSE	73 kDa	1	23.865125	1.06666667	1	1.06666667	0.9375
Medium-chain specific acyl-CoA dehydrogenase, mitochondrial	OS=Mus ACADM_MOUSE	46 kDa	1	23.873384	0.96666667	1.03333333	0.935483871	1.068965517
Cytochrome b-c1 complex subunit 2, mitochondrial	OS=Mus musculus G QCR2_MOUSE	48 kDa	1	21.78257	1.06666667	1	1.06666667	0.9375
Cluster of 14-3-3 protein gamma	OS=Mus musculus GN=Ywhag PE=1 SV=1433G_MOUSE [9]	28 kDa	1	52.50082	0.96666667	1	0.96666667	1.034482759
Cluster of Beta-enolase	OS=Mus musculus GN=Eno3 PE=1 SV=3 [ENOB_ENOB_MOUSE [3]	47 kDa	1	24.989203	1.1218	1.1	1.03030303	0.970588235
Cluster of Phosphoglycerate kinase 1	OS=Mus musculus GN=Pgk1 PE=1 SPGK1_MOUSE [2]	45 kDa	1	26.37556	0.9	0.9	1	1
Cluster of Vimentin	OS=Mus musculus GN=Vim PE=1 SV=3 [VIME_MOUSVIME_MOUSE [19]	54 kDa	1	42.01593	0.83333333	0.96666667	0.862068966	1.16
Cluster of Isoform M1 of Pyruvate kinase PKM	OS=Mus musculus GN=Pks1p [P52480-2] KPYM_MOUSE [2]	?	1	27.36292	1.06666667	1	1.06666667	0.9375
Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial	OS=Mus m IDH3A_MOUSE	40 kDa	1	23.894511	1.06666667	1.03333333	1.032258065	0.96875
Electron transfer flavoprotein subunit beta	OS=Mus musculus GN=Etfb P ETFB_MOUSE	28 kDa	1	25.35792	1	1	1	1
Acetyl-CoA acetyltransferase, mitochondrial	OS=Mus musculus GN=Acat THIL_MOUSE	45 kDa	1	26.787162	0.93333333	0.86666667	1.076923077	0.928571429
60 kDa heat shock protein, mitochondrial	OS=Mus musculus GN=Hspd1 ICH60_MOUSE	61 kDa	1	32.386965	0.8	0.83333333	0.96	1.041666667
Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondria	ETFDF_MOUSE	68 kDa	1	21.524051	1.03333333	1.06666667	0.96875	1.032258065
Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mi	ODPA_MOUSE	43 kDa	1	22.96659	0.96666667	0.96666667	1	1
Vinculin	OS=Mus musculus GN=Vcl PE=1 SV=4 VINC_MOUSE	117 kDa	1	21.79835	1.06666667	1.06666667	1	1
Cluster of Cytochrome c, somatic	OS=Mus musculus GN=Cycc PE=1 SV=2 CYC_MOUSE [2]	12 kDa	1	41.80578	1.03333333	1.4	0.738095238	1.35483871
ATP synthase subunit gamma	OS=Mus musculus GN=Atp5c1 PE=3 SV=1 A2AKU9_MOUSE	33 kDa	1	23.889059	1.13333333	1.1	1.03030303	0.970588235
Cluster of Sodium/potassium-transporting ATPase subunit alpha-1	OS=Mat1A1_MOUSE [6]	113 kDa	1	23.945752	1.13333333	1.16666667	0.971428571	1.029411765
Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial	OS=Mus r SUCB1_MOUSE	50 kDa	1	24.177477	1.05393333	1.03333333	1	1
Cytochrome b-c1 complex subunit 1, mitochondrial	OS=Mus musculus G QCR1_MOUSE	53 kDa	1	22.792082	0.96666667	0.96666667	1	1
Cluster of Histone H2B type 1-H	OS=Mus musculus GN=Hist1h2b PE=1 H2B1H_MOUSE [6]	14 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of LIM domain-binding protein 3	OS=Mus musculus GN=Ldb3 PE=LDB3_MOUSE [2]	76 kDa	1	31.008857	1	0.93333333	1.071428571	0.93333333
Pyruvate dehydrogenase E1 component subunit beta, mitochondrial	OS=ODPB_MOUSE	39 kDa	1	24.32211	1	0.96666667	1.034482759	0.96666667
Troponin C, slow skeletal and cardiac muscles	OS=Mus musculus GN=Tnni1 TNNC1_MOUSE	18 kDa	1	32.55204	1.23333333	1.1	1.121212121	0.891891892
Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondria	DHSB_MOUSE	32 kDa	1	24.417116	1.06666667	1.03333333	1.032258065	0.96875
Cluster of Phosphoglycerate mutase 2	OS=Mus musculus GN=Pgam2 PE=PGAM2_MOUSE [2]	29 kDa	1	65.760255	0.93333333	1.03333333	0.903225806	1.107142857
2-oxoglutarate dehydrogenase, mitochondrial	OS=Mus musculus GN=Og ODO1_MOUSE	116 kDa	1	24.993779	1.03333333	0.96666667	1.068965517	0.935483871
Dihydropyridine dehydrogenase, mitochondrial	OS=Mus musculus GN=Dld DLDH_MOUSE	54 kDa	1	24.803832	0.93333333	0.9	1.037037037	0.964285714
Cytochrome c oxidase subunit 5A, mitochondrial	OS=Mus musculus GN=COX5A_MOUSE	16 kDa	1	27.938437	0.96666667	0.83333333	1.16	0.862068966
Enoyl-CoA hydratase, mitochondrial	OS=Mus musculus GN=Echs1 PE=1 SECHM_MOUSE	31 kDa	1	23.648663	0.93333333	1.06666667	0.875	1.142857143
Cluster of Annexin A6	OS=Mus musculus GN=Anxa6 PE=1 SV=3 [ANXA6_ANXA6_MOUSE [2]	76 kDa	1	12.771863	1.2	1.25	0.96	1.041666667
Cluster of Prelamin A/C	OS=Mus musculus GN=Lmna PE=1 SV=2 [LMNA_LMNA_MOUSE [2]	74 kDa	1	21.014996	0.9	0.96666667	0.931034483	1.074074074
MCG10343, isoform CRA_b	OS=Mus musculus GN=Slc25a3 PE=3 SV=1 G5E902_MOUSE	40 kDa	1	25.821653	1.06666667	1.16666667	0.914285714	1.09375
Cluster of NADH dehydrogenase [ubiquinone] flavoprotein 1, mitochondria	NDUV1_MOUSE [2]	51 kDa	1	22.568347	1.03333333	1.16666667	0.885714286	1.129032258
Apolipoprotein A-I	OS=Mus musculus GN=Apoa1 PE=1 SV=2 APOA1_MOUSE	31 kDa	1	24.274969	0.9	0.9	1	1
Elongation factor Tu	OS=Mus musculus GN=Gm9755 PE=3 SV=1 D3YVN7_MOUSE (+1)	50 kDa	1	22.029337	1	1	1	1
Triosephosphate isomerase	OS=Mus musculus GN=Tpi1 PE=1 SV=4 TPIS_MOUSE	32 kDa	1	22.664697	0.86666667	0.93333333	0.928571429	1.076923077
ATP synthase-coupling factor 6, mitochondrial	OS=Mus musculus GN=At ATP5J_MOUSE	12 kDa	1	34.429255	0.96666667	0.8	1.208333333	0.827586207
Spectrin beta chain, non-erythrocytic 1	OS=Mus musculus GN=Sptbn1 PESPBT2_MOUSE	274 kDa	1	21.447621	1	1	1	1
Cluster of Histone H1.4	OS=Mus musculus GN=Hist1h1e PE=1 SV=2 [H14_H14_MOUSE [3]	22 kDa	1	22.55826	1	0.93333333	1.071428571	0.93333333
Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial	OS=Mus musculus HCDH_MOUSE	34 kDa	1	25.603878	1.06666667	1.03333333	1.032258065	0.96875
Nascent polypeptide-associated complex subunit alpha, muscle-specific	NACAM_MOUSE	220 kDa	1	21.927986	0.93333333	0.93333333	1	1
Cluster of Glutathione S-transferase Mu 1	OS=Mus musculus GN=Gstm1 GSTM1_MOUSE [5]	26 kDa	1	41.137487	1.06666667	1.1	0.96969697	1.03125
Cluster of Alpha-1-antitrypsin 1-1	OS=Mus musculus GN=Serpina1a PE=1A1AT1_MOUSE [4]	46 kDa	1	78.461015	0.96666667	1.03333333	0.935483871	1.068965517
Alpha-crystallin B chain	OS=Mus musculus GN=Cryab PE=1 SV=2 CRYAB_MOUSE	20 kDa	1	27.167386	0.8	0.8	1	1
Isoform Cytoplasmic+peroxisomal of Peroxiredoxin-5, mitochondrial	OS=sp [P99029-2] PRDX5_MOUSE	?	1	26.87764	1.06666667	1	1.06666667	0.9375
Cluster of Cytochrome c oxidase subunit 4 isoform 1, mitochondria	OS=COX4I_MOUSE [2]	20 kDa	1	23.022999	1	1.06666667	0.9375	1.06666667
Isocitrate dehydrogenase 3 (NAD+) beta	OS=Mus musculus GN=Idh3b PEQ91VA7_MOUSE	42 kDa	1	21.312723	1.13333333	1.13333333	1	1
Dihydropyridine residue acetyltransferase component of pyruvate de	ODP2_MOUSE	68 kDa	1	21.505937	0.9	0.9	1	1
Cluster of ATP synthase subunit O, mitochondrial	OS=Mus musculus GN=ATPO_MOUSE	23 kDa	1	25.889695	0.93333333	0.96666667	0.965517241	1.035714286
Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondria	OS=M SCOT1_MOUSE	56 kDa	1	23.77109	1	1.03333333	0.967741935	1.03333333
10 kDa heat shock protein, mitochondria	OS=Mus musculus GN=Hspe1 ICH10_MOUSE	11 kDa	1	27.067035	0.7375	0.73333333	1	1
Cluster of L-lactate dehydrogenase	OS=Mus musculus GN=Ldha PE=3 SV=G5E8N5_MOUSE [3]	40 kDa	1	20.868783	1.03333333	1.13333333	0.911764706	1.096774194

Superoxide dismutase [Mn], mitochondrial OS=Mus musculus GN=Sod2 ISODM_MOUSE	25 kDa	1	25.42007	1	1	1	1
Cluster of Tubulin beta-4B chain OS=Mus musculus GN=Tubb4b PE=1 SV:TBB4B_MOUSE [4]	50 kDa	1	1126.6539	0.833333333	0.933333333	0.892857143	1.12
Myozenin-2 OS=Mus musculus GN=Myoz2 PE=1 SV=1 MYOZ2_MOUSE	30 kDa	1	24.08687	1.033333333	0.966666667	1.068965517	0.935483871
Succinyl-CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial OS:SUCA_MOUSE	36 kDa	1	23.905633	1.033333333	1.066666667	0.96875	1.032258065
ATP synthase subunit b, mitochondrial OS=Mus musculus GN=Atp5f1 PE:AT5F1_MOUSE	29 kDa	1	25.549453	0.966666667	0.966666667	1	1
Cluster of Carnitine O-palmitoyltransferase 2, mitochondrial OS=Mus muCPT2_MOUSE	74 kDa	1	22.652262	1.033333333	1.1	0.939393939	1.064516129
Cytochrome c oxidase subunit 6B1 OS=Mus musculus GN=Cox6b1 PE=1 CX6B1_MOUSE	10 kDa	1	24.99389	0.966666667	1	0.966666667	1.034482759
Cluster of Basement membrane-specific heparan sulfate proteoglycan ccB1B0C7_MOUSE [2]	469 kDa	1	30.051085	1	0.966666667	1.034482759	0.966666667
Cluster of Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 PE=1 SV=1 (PRD:PRDX1_MOUSE	22 kDa	1	22.737333	1	1.1	0.909090909	1.1
Cluster of Nucleoside diphosphate kinase OS=Mus musculus GN=Gm203 E9PZF0_MOUSE [3]	30 kDa	1	30.421865	0.966666667	0.966666667	1	1
Enoyl-CoA delta isomerase 1, mitochondrial OS=Mus musculus GN=Eci1 ECI1_MOUSE	32 kDa	1	22.536466	1.096	1.1	1	1
Cluster of Sorbin and SH3 domain-containing protein 2 OS=Mus musculus B2RXQ9_MOUSE [5]	72 kDa	1	22.198606	0.766666667	0.733333333	1.045454545	0.956521739
Cluster of ATP synthase subunit d, mitochondrial OS=Mus musculus GN=ATP5H_MOUSE [2]	19 kDa	1	25.14485	0.933333333	0.866666667	1.076923077	0.928571429
Cluster of Isoform 2 of Adenylate kinase isoenzyme 1 OS=Mus musculus sp Q9R0Y5-2 KAD1_MOUSE [2]	?	1	15.728442	0.933333333	0.9	1.037037037	0.964285714
78 kDa glucose-regulated protein OS=Mus musculus GN=Hspa5 PE=1 SV:GRP78_MOUSE	72 kDa	1	22.931659	0.9	0.933333333	0.964285714	1.037037037
Carnitine O-palmitoyltransferase 1, muscle isoform OS=Mus musculus GICPT1B_MOUSE	88 kDa	1	21.052587	1.066666667	1.1	0.96969697	1.03125
Phosphoglucomutase-1 OS=Mus musculus GN=Pgm1 PE=1 SV=4 PGM1_MOUSE	61 kDa	1	20.52368	1.166666667	1.2	0.972222222	1.028571429
Filamin-C OS=Mus musculus GN=Flnc PE=1 SV=3 FLNC_MOUSE	291 kDa	1	21.376863	0.966666667	0.9	1.074074074	0.931034483
Dihydropolyllysine-residue succinyltransferase component of 2-oxoglutarate ODO2_MOUSE	49 kDa	1	23.038653	1	0.966666667	1.034482759	0.966666667
Thioredoxin-dependent peroxide reductase, mitochondrial OS=Mus musculus PRDX3_MOUSE	28 kDa	1	23.396891	1.066666667	1.1	0.96969697	1.03125
Transitional endoplasmic reticulum ATPase OS=Mus musculus GN=Vcp P TERA_MOUSE	89 kDa	1	23.00108	0.996466667	1	1	1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial (NDUS2_MOUSE	53 kDa	1	23.785982	1.133333333	1.166666667	0.971428571	1.029411765
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10, mitochondrial (NDUAA_MOUSE	41 kDa	1	22.33221	1.033333333	1.033333333	1	1
Alpha-2-macroglobulin OS=Mus musculus GN=A2m PE=1 SV=3 A2M_MOUSE (+1)	166 kDa	1	21.242344	1.033333333	1.033333333	1	1
Cluster of Glucose-6-phosphate isomerase OS=Mus musculus GN=Gpi PE G6PI_MOUSE [2]	63 kDa	1	23.293847	1	1.066666667	0.9375	1.066666667
2,4-dienoyl-CoA reductase, mitochondrial OS=Mus musculus GN=Decr1 (DECR_MOUSE	36 kDa	1	23.436627	1.066666667	1.1	0.96969697	1.03125
Aldose reductase OS=Mus musculus GN=Akr1b1 PE=1 SV=3 ALDR_MOUSE	36 kDa	1	21.103968	1	1	1	1
ES1 protein homolog, mitochondrial OS=Mus musculus GN=D10Jhu81e FES1_MOUSE	28 kDa	1	25.15947	0.966666667	1	0.966666667	1.034482759
Cytochrome b-c1 complex subunit 6, mitochondrial OS=Mus musculus G QCR6_MOUSE	10 kDa	1	24.470049	1.133333333	0.966666667	1.172413793	0.852941176
Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Mus musculus UCRI_MOUSE	29 kDa	1	22.218429	1.033333333	1	1.033333333	0.967741935
Voltage-dependent anion-selective channel protein 3 OS=Mus musculus J3QMG3_MOUSE (+1)	31 kDa	1	24.449185	1.066666667	1.166666667	0.914285714	1.09375
Polymerase I and transcript release factor OS=Mus musculus GN=Ptrf PE PTRF_MOUSE	44 kDa	1	21.686059	1.1	1.066666667	1.03125	0.96969697
Cluster of Tubulin alpha-4A chain OS=Mus musculus GN=Tuba4a PE=1 SV:TBA4A_MOUSE [5]	50 kDa	1	69.287926	0.933333333	0.966666667	0.965517241	1.035714286
NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Mus musculus (NDUV2_MOUSE	27 kDa	1	26.41426	1.133333333	1.033333333	1.096774194	0.911764706
Methylmalonate-semialdehyde dehydrogenase [acylating], mitochondrial (MMSA_MOUSE	58 kDa	1	22.055392	1.066666667	1.133333333	0.941176471	1.0625
Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Mus mu ECH1_MOUSE	36 kDa	1	23.647761	1.033333333	1.133333333	0.911764706	1.096774194
Cysteine and glycine-rich protein 3 OS=Mus musculus GN=Csrp3 PE=1 SV:CSR3_MOUSE	21 kDa	1	25.659925	1.033333333	0.966666667	1.068965517	0.935483871
Dystrophin OS=Mus musculus GN=Dmd PE=1 SV=3 DMD_MOUSE	426 kDa	1	21.04594	0.966666667	0.966666667	1	1
Myosin regulatory light chain 2, atrial isoform OS=Mus musculus GN=MyMLRA_MOUSE	19 kDa	1	41.67893	0.8	0.9	0.888888889	1.125
Cluster of Ubiquitin-40S ribosomal protein S27a OS=Mus musculus GN=FRS27A_MOUSE [2]	18 kDa	1	0	1.1	1.1	1	1
Cluster of Acyl-coenzyme A thioesterase 2, mitochondrial OS=Mus musculus ACOT2_MOUSE [3]	50 kDa	1	59.91892	0.8	0.9	0.888888889	1.125
Cytochrome c1, heme protein, mitochondrial OS=Mus musculus GN=Cyc CY1_MOUSE	35 kDa	1	23.00798	1.133333333	1.133333333	1	1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial (NDUS3_MOUSE	30 kDa	1	24.672705	1.033333333	0.966666667	1.068965517	0.935483871
NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial (NDUS6_MOUSE	13 kDa	1	33.005902	0.9	0.766666667	1.173913043	0.851851852
Cluster of Histone H3.1 OS=Mus musculus GN=Hist1h3a PE=1 SV=2 (H31, H31_MOUSE [4]	15 kDa	1	21.202742	0.866666667	0.966666667	0.896551724	1.115384615
Cluster of PDZ and LIM domain protein 5 OS=Mus musculus GN=Pdlim5 (PDLI5_MOUSE [3]	63 kDa	1	30.018938	0.633333333	0.733333333	0.863636364	1.157894737
Protein NipSnap homolog 2 OS=Mus musculus GN=Gbas PE=2 SV=1 NIP52_MOUSE (+1)	33 kDa	1	21.84825	1.1	1.2	0.916666667	1.090909091
Cluster of Propionyl-CoA carboxylase alpha chain, mitochondrial OS=Mus mu:PCCA_MOUSE [2]	80 kDa	1	21.947795	1.033333333	1.033333333	1	1
Cluster of Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3 (ENOA, ENOA_MOUSE [2]	47 kDa	1	23.470576	0.866666667	0.966666667	0.896551724	1.115384615
Acyl carrier protein, mitochondrial OS=Mus musculus GN=Ndufab1 PE=1 ACPM_MOUSE (+2)	17 kDa	1	28.053743	0.9	0.833333333	1.08	0.925925926
Elongation factor 1-alpha 2 OS=Mus musculus GN=Eef1a2 PE=1 SV=1 EF1A2_MOUSE	50 kDa	1	25.993603	1	0.966666667	1.034482759	0.966666667
Peptidyl-prolyl cis-trans isomerase A OS=Mus musculus GN=Ppia PE=1 SV:PIPIA_MOUSE	18 kDa	1	23.900767	0.8513	0.9	0.925925926	1.08
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=MNDUA7_MOUSE	13 kDa	1	24.66094	0.933333333	1	0.933333333	1.071428571

Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Mus musculus GN=ROA2_MOUSE	37 kDa	1	21.48754	0.933333333	1	0.933333333	1.071428571
Ubiquinone biosynthesis protein COQ9, mitochondrial OS=Mus musculus:COQ9_MOUSE	35 kDa	1	21.19581	1.066666667	1	1.066666667	0.9375
NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Mus musculus:NDUS8_MOUSE	24 kDa	1	22.68523	1.066666667	1.033333333	1.032258065	0.96875
Peroxiredoxin-6 OS=Mus musculus GN=Prdx6 PE=2 SV=1 D3Z0Y2_MOUSE (+1)	22 kDa	1	24.262355	0.9	0.833333333	1.08	0.925925926
Cluster of Talin-1 OS=Mus musculus GN=TLn1 PE=1 SV=2 (TLN1_MOUSE) TLN1_MOUSE	270 kDa	1	23.284364	1.066666667	1	1.066666667	0.9375
Myosin-9 OS=Mus musculus GN=Myh9 PE=1 SV=4 MYH9_MOUSE	226 kDa	1	20.120242	1.066666667	1	1.066666667	0.9375
Cluster of Serine protease inhibitor A3K OS=Mus musculus GN=Serpina3 SPA3K_MOUSE [2]	47 kDa	1	24.820502	1.033333333	1.033333333	1	1
UTP--glucose-1-phosphate uridylyltransferase OS=Mus musculus GN=Ug UGPA_MOUSE	57 kDa	1	20.789298	0.933333333	0.833333333	1.12	0.892857143
Apoptosis-inducing factor 1, mitochondrial OS=Mus musculus GN=Aifm1 AIFM1_MOUSE	67 kDa	1	22.026622	1.1	1.066666667	1.03125	0.96969697
Cytochrome c oxidase subunit 6C OS=Mus musculus GN=Cox6c PE=1 SV= COX6C_MOUSE	8 kDa	1	26.715234	0.866666667	1.033333333	0.838709677	1.192307692
Four and a half LIM domains protein 2 OS=Mus musculus GN=Fhl2 PE=1 FHL2_MOUSE	32 kDa	1	25.744793	1.033333333	0.966666667	1.068965517	0.935483871
Lon protease homolog, mitochondrial OS=Mus musculus GN=Lonp1 PE=:LONM_MOUSE	106 kDa	1	21.712473	0.7337	0.766666667	0.956521739	1.045454545
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=MNDUA8_MOUSE	20 kDa	1	22.035451	0.933333333	1.1	0.848484848	1.178571429
Cluster of Sorbin and SH3 domain-containing protein 1 OS=Mus musculus SRBS1_MOUSE [2]	143 kDa	1	61.279005	0.933333333	0.933333333	1	1
Calsequestrin-2 OS=Mus musculus GN=Casq2 PE=2 SV=3 CASQ2_MOUSE (+1)	48 kDa	1	20.76281	1.033333333	1.1	0.939393939	1.064516129
Isocitrate dehydrogenase [NAD] subunit gamma 1, mitochondrial OS=Mt IDHG1_MOUSE	43 kDa	1	22.134587	1.033333333	1.033333333	1	1
Cluster of Actin-binding LIM protein 1 OS=Mus musculus GN=Ablim1 PE=E9Q9C1_MOUSE [2]	84 kDa	1	6.151903	0.9	0.8	1.125	0.888888889
Laminin subunit gamma-1 OS=Mus musculus GN=Lamc1 PE=2 SV=1 F8VQJ3_MOUSE	177 kDa	1	22.924528	1.1	0.966666667	1.137931034	0.878787879
Histone H1.5 OS=Mus musculus GN=Hist1h1b PE=1 SV=2 H15_MOUSE	23 kDa	1	20.209862	1.1	1.1	1	1
Prohibitin OS=Mus musculus GN=Phb PE=1 SV=1 PHB_MOUSE	30 kDa	1	22.333796	0.933333333	0.866666667	1.076923077	0.928571429
Cytochrome b-c1 complex subunit 8 OS=Mus musculus GN=Uqcrc PE=1 QCR8_MOUSE	10 kDa	1	20.729516	1	1.066666667	0.9375	1.066666667
Heat shock protein beta-1 OS=Mus musculus GN=Hspb1 PE=1 SV=3 HSPB1_MOUSE	23 kDa	1	25.343826	0.7064	0.733333333	1	1
Cluster of Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE:SODC_MOUSE	16 kDa	1	23.843075	0.933333333	0.866666667	1.076923077	0.928571429
Moesin OS=Mus musculus GN=Msn PE=1 SV=3 MOES_MOUSE	68 kDa	1	21.800491	0.87933	0.9	0.962962963	1.038461538
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=MNDUA4_MOUSE	9 kDa	1	23.565091	0.933333333	1.033333333	0.903225806	1.107142857
Cluster of 3-hydroxyisobutyryl-CoA hydrolase, mitochondrial OS=Mus musculus HIBCH_MOUSE [2]	43 kDa	1	22.349542	0.866666667	0.833333333	1.04	0.961538462
Plakophilin 2 OS=Mus musculus GN=Pkp2 PE=2 SV=1 Q9CQ73_MOUSE	88 kDa	1	20.409517	1	1	1	1
Cluster of Ribosome-binding protein 1 OS=Mus musculus GN=Rrbp1 PE=A2AVJ7_MOUSE [2]	158 kDa	1	21.087505	0.966666667	0.933333333	1.035714286	0.965517241
Protein Trdn OS=Mus musculus GN=Trdn PE=2 SV=1 E9Q9K5_MOUSE	78 kDa	1	19.730946	1.233333333	1.133333333	1.088235294	0.918918919
Protein disulfide-isomerase A3 OS=Mus musculus GN=Pdia3 PE=1 SV=2 PDIA3_MOUSE	57 kDa	1	21.53763	0.933333333	1	0.933333333	1.071428571
Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3 PLAK_MOUSE	82 kDa	1	23.701087	1	1.066666667	0.9375	1.066666667
Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1 CALR_MOUSE	48 kDa	1	20.603183	0.9	0.933333333	0.964285714	1.037037037
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 2 OS=MNDUA2_MOUSE	11 kDa	1	20.815434	1	1	1	1
Protein DJ-1 OS=Mus musculus GN=Park7 PE=1 SV=1 PARK7_MOUSE	20 kDa	1	22.764306	0.9	0.866666667	1.038461538	0.962962963
Pyruvate dehydrogenase protein X component, mitochondrial OS=Mus musculus ODPX_MOUSE	54 kDa	1	20.695071	0.966666667	0.933333333	1.035714286	0.965517241
Acetyl-coenzyme A synthetase 2-like, mitochondrial OS=Mus musculus GACS2L_MOUSE	75 kDa	1	22.297797	1	1.1	0.909090909	1.1
Cluster of EH domain-containing protein 4 OS=Mus musculus GN=Ehd4 FEHD4_MOUSE [2]	61 kDa	1	73.16867	1.1	1.133333333	0.970588235	1.03030303
Hexokinase 1, isoform CRA_f OS=Mus musculus GN=Hk1 PE=3 SV=1 G3UVV4_MOUSE (+1)	102 kDa	1	19.505818	1.033333333	1.033333333	1	1
Cluster of Histone H2A.x OS=Mus musculus GN=H2afx PE=1 SV=2 (H2AX_H2AX_MOUSE [10])	15 kDa	1	97.976774	0.933333333	1.033333333	0.903225806	1.107142857
Cluster of Ubiquitin-like modifier-activating enzyme 1 OS=Mus musculus UBA1_MOUSE [2]	118 kDa	1	21.83291	0.966666667	1	0.966666667	1.034482759
Cytochrome c oxidase subunit 2 OS=Mus musculus GN=Mtco2 PE=1 SV=:COX2_MOUSE (+1)	26 kDa	1	25.713077	1.1	1.066666667	1.03125	0.96969697
EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1 EHD2_MOUSE	61 kDa	1	21.753658	1	1.133333333	0.882352941	1.133333333
Voltage-dependent anion-selective channel protein 2 (Fragment) OS=Mt G3UX26_MOUSE (+1)	30 kDa	1	23.24543	1.066666667	1.2	0.888888889	1.125
ATP synthase subunit delta, mitochondrial OS=Mus musculus GN=Atp5d ATPD_MOUSE	18 kDa	1	22.777788	1.1	0.866666667	1.269230769	0.787878788
Hemopexin OS=Mus musculus GN=Hpx PE=1 SV=2 HEMO_MOUSE	51 kDa	1	21.900138	1	1.066666667	0.9375	1.066666667
Cluster of NADH dehydrogenase [ubiquinone] flavoprotein 3, mitochondrion OS=Mus musculus NDUV3_MOUSE [2]	12 kDa	1	26.060414	0.966666667	0.8	1.208333333	0.827586207
Cluster of Murinoglobulin-1 OS=Mus musculus GN=Mug1 PE=1 SV=3 (MUG1_MOUSE [2])	165 kDa	1	30.218425	0.833333333	0.933333333	0.892857143	1.12
Tripartite motif-containing protein 72 OS=Mus musculus GN=Trim72 PE=TRI72_MOUSE	53 kDa	1	23.179765	1.066666667	0.933333333	1.142857143	0.875
Cluster of Protein Neb OS=Mus musculus GN=Neb PE=2 SV=1 (A2AQA9_A2AQA9_MOUSE [3])	801 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of Hexokinase-2 OS=Mus musculus GN=Hk2 PE=1 SV=1 (HXK2_M HXK2_MOUSE)	103 kDa	1	20.551997	1	1.033333333	0.967741935	1.033333333
Nucleolin OS=Mus musculus GN=Ncl PE=1 SV=2 NUC1_MOUSE	77 kDa	1	21.239017	0.8041	0.833333333	0.96	1.041666667
Serum deprivation-response protein OS=Mus musculus GN=Sdpr PE=1 SV=SDPR_MOUSE	47 kDa	1	21.971652	1	0.933333333	1.071428571	0.933333333
Cluster of Kininogen-1 OS=Mus musculus GN=Kng1 PE=1 SV=1 (KNG1_M KNG1_MOUSE [2])	73 kDa	1	27.040312	0.933333333	0.9	1.037037037	0.964285714

Prohibitin-2 OS=Mus musculus GN=Phb2 PE=1 SV=1	PHB2_MOUSE	33 kDa	1	20.787968	0.9	0.86666667	1.038461538	0.962962963
60S ribosomal protein L4 OS=Mus musculus GN=Rpl4 PE=1 SV=3	RL4_MOUSE	47 kDa	1	23.343958	0.9	1.03333333	0.870967742	1.148148148
Cluster of Microtubule-associated protein (Fragment) OS=Mus musculus F7CK47_MOUSE [2]		153 kDa	1	24.227646	1.03333333	0.9	1.148148148	0.870967742
Cluster of Ras-related protein Rab-1B OS=Mus musculus GN=Rab1b PE=1RAB1B_MOUSE [7]		22 kDa	1	215.08331	1.03333333	1.1	0.939393939	1.064516129
Endoplasmic reticulum chaperone protein OS=Mus musculus GN=Hsp90b1 PE=1 SV=2	ENPL_MOUSE	92 kDa	1	21.536195	0.96666667	0.9	1.074074074	0.931034483
Heat shock 70 kDa protein 4 OS=Mus musculus GN=Hspa4 PE=2 SV=1	Q3U2G2_MOUSE	94 kDa	1	21.542004	0.85536667	0.93333333	0.928571429	1.076923077
Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=TGM2_MOUSE		77 kDa	1	21.645275	1.06666667	1.1	0.96969697	1.03125
Cluster of Carboxylesterase 1C OS=Mus musculus GN=Ces1c PE=1 SV=4 (EST1C_MOUSE [3])		61 kDa	1	28.846636	1.03333333	1.1	0.939393939	1.064516129
NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Mus musculus GN=NDUC2_MOUSE		14 kDa	1	21.862963	1.03333333	1.1	0.939393939	1.064516129
Cluster of Laminin subunit alpha-2 OS=Mus musculus GN=Lama2 PE=1 SV=1 (S\LAMA2_MOUSE [2])		343 kDa	1	20.730539	0.7	0.93333333	0.75	1.33333333
Cluster of Myosin-11 OS=Mus musculus GN=Myh11 PE=1 SV=1 (MYH11_MOUSE [2])		227 kDa	1	8.001554	1.05	0.95	1.105263158	0.904761905
Cluster of Selenium-binding protein 1 OS=Mus musculus GN=Selenbp1 P SBP1_MOUSE [2]		53 kDa	1	22.65691	1.06666667	1.13333333	0.941176471	1.0625
Cluster of Myosin light chain 4 (Fragment) OS=Mus musculus GN=Myl4 PA2A6Q8_MOUSE [3]		21 kDa	1	0	1	0.25	4	0.25
Heterogeneous nuclear ribonucleoprotein M OS=Mus musculus GN=Hnr B8JK32_MOUSE (+3)		72 kDa	1	19.85611	0.96666667	1	0.96666667	1.034482759
Phosphatidylethanolamine-binding protein 1 OS=Mus musculus GN=Peb D3Z1V4_MOUSE (+1)		23 kDa	1	27.034444	0.8	0.73333333	1.090909091	0.91666667
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=MNDUA6_MOUSE		15 kDa	1	22.54544	1.1	1.03333333	1.064516129	0.939393939
Peroxiredoxin-2 OS=Mus musculus GN=Prdx2 PE=1 SV=3	PRDX2_MOUSE	22 kDa	1	20.521066	1.13333333	1.16666667	0.971428571	1.029411765
Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	ANXA2_MOUSE	39 kDa	1	21.187776	0.96666667	1	0.96666667	1.034482759
Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	LEG1_MOUSE	15 kDa	1	21.617624	0.9	0.96666667	0.931034483	1.074074074
Acyl-CoA dehydrogenase family member 9, mitochondrial OS=Mus musculus ACAD9_MOUSE		69 kDa	1	21.001568	1.03333333	1.03333333	1	1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=MNDUBA_MOUSE		21 kDa	1	24.455515	1.16666667	1.1	1.060606061	0.942857143
von Willebrand factor A domain-containing protein 8 OS=Mus musculus VWA8_MOUSE		213 kDa	1	21.46171	1	1	1	1
Cluster of Enoyl-CoA delta isomerase 2, mitochondrial OS=Mus musculus EC12_MOUSE [5]		43 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Protein disulfide-isomerase OS=Mus musculus GN=P4hb PE=1 SV=2	PDI1_MOUSE	57 kDa	1	21.437772	0.8567	0.83333333	1.04	0.961538462
Cluster of Ankyrin-2 OS=Mus musculus GN=Ank2 PE=1 SV=2 (ANK2_MOUSE [4])		426 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Perilipin-4 OS=Mus musculus GN=Plin4 PE=1 SV=2	PLIN4_MOUSE	139 kDa	1	19.081882	1.1	1.1	1	1
Tropomodulin-1 OS=Mus musculus GN=Tmod1 PE=2 SV=2	TMOD1_MOUSE	40 kDa	1	22.36374	0.93333333	0.93333333	1	1
Cluster of D-beta-hydroxybutyrate dehydrogenase, mitochondrial OS=MDH_MOUSE [2]		38 kDa	1	22.139552	0.73333333	0.66666667	1.1	0.909090909
Creatine kinase B-type OS=Mus musculus GN=Ckb PE=1 SV=1	KCRB_MOUSE	43 kDa	1	25.364146	1.03333333	1.16666667	0.885714286	1.129032258
Chaperone activity of bc1 complex-like, mitochondrial OS=Mus musculus ADCK3_MOUSE		72 kDa	1	22.790584	1.06666667	1.06666667	1	1
Cluster of Protein Tns1 OS=Mus musculus GN=Tns1 PE=2 SV=1 (E9Q0S6_MOUSE [2])		201 kDa	1	22.13987	1	1	1	1
Striated muscle-specific serine/threonine-protein kinase OS=Mus musculus E9Q0Q25_MOUSE		354 kDa	1	22.692575	1	1	1	1
60S ribosomal protein L6 OS=Mus musculus GN=Rpl6 PE=1 SV=3	RL6_MOUSE	34 kDa	1	21.481863	0.96666667	0.96666667	1	1
Nesprin-2 OS=Mus musculus GN=Syne2 PE=1 SV=2	SYNE2_MOUSE	783 kDa	1	9.591509	1	1.3	0.769230769	1.3
Cluster of Glutathione S-transferase omega-1 OS=Mus musculus GN=Gst GSTO1_MOUSE		27 kDa	1	20.586726	0.7771	0.7	1.095238095	0.913043478
Cluster of Calpastatin OS=Mus musculus GN=Cast PE=1 SV=2 (ICAL_MOUSE [3])		85 kDa	1	22.758579	0.9	0.8	1.125	0.888888889
Non-specific lipid-transfer protein OS=Mus musculus GN=Scp2 PE=1 SV=1 (NLTP_MOUSE)		59 kDa	1	20.306158	0.9	0.96666667	0.931034483	1.074074074
Calcium-binding mitochondrial carrier protein Aralar2 OS=Mus musculus CMC2_MOUSE		74 kDa	1	20.084044	1.03333333	1.13333333	0.911764706	1.096774194
Cysteine-rich protein 2 OS=Mus musculus GN=Crip2 PE=1 SV=1	CRIP2_MOUSE	23 kDa	1	50.443006	0.86666667	0.76666667	1.130434783	0.884615385
Uncharacterized protein OS=Mus musculus GN=Rps3a2 PE=3 SV=1	D3Z6C3_MOUSE	30 kDa	1	21.281977	0.93333333	1	0.93333333	1.071428571
Glutamate dehydrogenase 1, mitochondrial OS=Mus musculus GN=Glud DHE3_MOUSE		61 kDa	1	20.491561	1	1.06666667	0.9375	1.06666667
Cluster of Catenin alpha-1 OS=Mus musculus GN=Ctnna1 PE=1 SV=1 (CTCTNA1_MOUSE)		100 kDa	1	20.310375	1.03333333	0.9	1.148148148	0.870967742
Nexilin OS=Mus musculus GN=Nexn PE=1 SV=3	NEXN_MOUSE	72 kDa	1	20.709956	0.93333333	1.03333333	0.903225806	1.107142857
Cluster of Talin-2 OS=Mus musculus GN=Tln2 PE=4 SV=1 (E9PUM4_MOUSE [2])		272 kDa	1	6.36418	0.95	1.05	0.904761905	1.105263158
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=MNDUAD_MOUSE		17 kDa	1	26.50013	1	1	1	1
Phosphoglucomutase-like protein 5 OS=Mus musculus GN=Pgm5 PE=1 SV=1 (S\PGM5_MOUSE)		62 kDa	1	20.74933	1	1.03333333	0.967741935	1.03333333
Cluster of Dynamin-like 120 kDa protein, mitochondrial OS=Mus musculus H7BX01_MOUSE [3]		113 kDa	1	21.181275	0.96666667	1.03333333	0.935483871	1.068965517
Heat shock protein HSP 90-alpha OS=Mus musculus GN=Hsp90aa1 PE=1 SV=1 (HS90A_MOUSE)		85 kDa	1	23.751697	0.78323333	0.9	0.851851852	1.173913043
Protein Agl OS=Mus musculus GN=Agl PE=4 SV=1	F8VFN4_MOUSE	174 kDa	1	21.163423	1.1293	1.13333333	0.970588235	1.03030303
Plasminogen activator inhibitor 1 RNA-binding protein OS=Mus musculus PAIRB_MOUSE		45 kDa	1	19.4952	0.86666667	0.86666667	1	1
Propionyl-CoA carboxylase beta chain, mitochondrial OS=Mus musculus PCCB_MOUSE		58 kDa	1	22.003977	0.96666667	1.03333333	0.935483871	1.068965517
Laminin subunit beta-1 OS=Mus musculus GN=Lamb1 PE=2 SV=1	E9QN70_MOUSE	202 kDa	1	22.540122	1.06666667	1.03333333	1.032258065	0.96875
Lumican OS=Mus musculus GN=Lum PE=1 SV=2	LUM_MOUSE	38 kDa	1	22.302015	0.86666667	0.93333333	0.928571429	1.076923077

Filamin, alpha OS=Mus musculus GN=Flna PE=4 SV=1	B7FAU9_MOUSE (+2)	280 kDa	1	20.803203	1.033333333	0.933333333	1.107142857	0.903225806
Cluster of MCG5732 OS=Mus musculus GN=Gm4978 PE=4 SV=2 (F6VBB8 F6VBB8_MOUSE [5])		30 kDa	1	0	1.2	1.1	1.090909091	0.916666667
Acyl-coenzyme A thioesterase 13 OS=Mus musculus GN=Acot13 PE=1 SV=1 (ACO13_MOUSE)		15 kDa	1	22.309871	1.066666667	1.033333333	1.032258065	0.96875
Cluster of 40S ribosomal protein S4, X isoform OS=Mus musculus GN=RpRS4X_MOUSE [3]		30 kDa	1	0	0.9	1.066666667	0.84375	1.185185185
Calmodulin OS=Mus musculus GN=Calm1 PE=1 SV=2	CALM_MOUSE (+1)	17 kDa	1	24.645248	1	0.933333333	1.071428571	0.933333333
Protein Fga OS=Mus musculus GN=Fga PE=2 SV=1	E9PV24_MOUSE (+1)	87 kDa	1	21.172126	1	1.033333333	0.967741935	1.033333333
Protein Col6a3 OS=Mus musculus GN=Col6a3 PE=4 SV=2	E9PWQ3_MOUSE	354 kDa	1	23.593284	1.366666667	1.066666667	1.28125	0.780487805
Histone H1.0 OS=Mus musculus GN=H1f0 PE=2 SV=4	H10_MOUSE	21 kDa	1	20.203805	1.033333333	1.2	0.861111111	1.161290323
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 9 OS=Mus musculus GN=NDUB9_MOUSE		22 kDa	1	25.329214	1.1	1.133333333	0.970588235	1.03030303
Heat shock protein beta-6 OS=Mus musculus GN=Hspb6 PE=1 SV=1	HSPB6_MOUSE	18 kDa	1	20.452799	0.866666667	0.8	1.083333333	0.923076923
Methylcrotonyl-CoA carboxylase subunit alpha, mitochondrial OS=Mus musculus GN=MCCA_MOUSE		79 kDa	1	21.662836	1.1	1.133333333	0.970588235	1.03030303
NADH dehydrogenase [ubiquinone] iron-sulfur protein 5 OS=Mus musculus GN=NDUS5_MOUSE		13 kDa	1	21.68276	1.033333333	0.9	1.148148148	0.870967742
ATP synthase subunit epsilon, mitochondrial OS=Mus musculus GN=Atp5E_MOUSE		6 kDa	1	27.739382	0.8	1.066666667	0.75	1.333333333
Platelet glycoprotein 4 OS=Mus musculus GN=Cd36 PE=1 SV=2	CD36_MOUSE	53 kDa	1	21.855485	1.133333333	1.233333333	0.918918919	1.088235294
Ig kappa chain C region OS=Mus musculus PE=1 SV=1	IGKC_MOUSE	12 kDa	1	22.266749	0.966666667	0.8	1.208333333	0.827586207
Protein phosphatase 1 regulatory subunit 12B OS=Mus musculus GN=PpMYPT2_MOUSE		109 kDa	1	21.341872	1	0.966666667	1.034482759	0.966666667
Cluster of Protein Pabpc6 OS=Mus musculus GN=Pabpc6 PE=2 SV=1 (Q9IQ9D4E6_MOUSE [4])		71 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Alpha-2-HS-glycoprotein OS=Mus musculus GN=Ahsg PE=1 SV=1	FETUA_MOUSE	37 kDa	1	20.93477	0.9	0.8	1.125	0.888888889
3-hydroxyacyl-CoA dehydrogenase type-2 OS=Mus musculus GN=Hsd17A2AFQ2_MOUSE (+1)		28 kDa	1	29.58014	1.033333333	1.066666667	0.96875	1.032258065
Sodium/potassium-transporting ATPase subunit beta-1 OS=Mus musculus GN=AT1B1_MOUSE		35 kDa	1	21.883778	1.1	1.166666667	0.942857143	1.060606061
Cluster of Catalase OS=Mus musculus GN=Cat PE=1 SV=4 (CATA_MOUSE CATA_MOUSE [2])		60 kDa	1	28.947672	0.933333333	0.9	1.037037037	0.964285714
Cluster of Protein Gm10923 OS=Mus musculus GN=Gm10923 PE=3 SV=1 (D3YU46_MOUSE [4])		17 kDa	1	0	0.5	0.3	1.666666667	0.6
High mobility group protein B1 OS=Mus musculus GN=Hmgb1 PE=1 SV=2 (HMGB1_MOUSE)		25 kDa	1	21.463308	0.9	1	0.9	1.111111111
Methylcrotonyl-CoA carboxylase beta chain, mitochondrial OS=Mus musculus GN=MCCB_MOUSE		61 kDa	1	22.032514	1.033333333	1.133333333	0.911764706	1.096774194
Protein NDRG2 OS=Mus musculus GN=NDRG2 PE=1 SV=1	NDRG2_MOUSE	41 kDa	1	20.763037	1	1	1	1
Cluster of NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 5 OS=Mus musculus GN=NDUB5_MOUSE [2]		22 kDa	1	17.698397	1.1	1.133333333	0.970588235	1.03030303
ATPase inhibitor, mitochondrial OS=Mus musculus GN=Atpif1 PE=2 SV=1 (E9PV44_MOUSE)		9 kDa	1	24.242947	0.833333333	0.633333333	1.315789474	0.76
Cluster of Hmgn2 protein OS=Mus musculus GN=Hmgn2 PE=4 SV=1 (Q5Q5XK38_MOUSE [2])		10 kDa	1	23.643467	1.2	0.7	1.714285714	0.583333333
Cluster of Microtubule-associated protein tau OS=Mus musculus GN=MaA2A5Y6_MOUSE [4]		78 kDa	1	19.519179	0.966666667	0.933333333	1.035714286	0.965517241
Carbonic anhydrase 2 OS=Mus musculus GN=Ca2 PE=1 SV=4	CAH2_MOUSE	29 kDa	1	20.31361	1.033333333	1.066666667	0.96875	1.032258065
Protein Golgb1 OS=Mus musculus GN=Golgb1 PE=2 SV=1	E9QAHI_MOUSE	365 kDa	1	7.870448	1	1.15	0.869565217	1.15
Adenosine kinase OS=Mus musculus GN=Adk PE=1 SV=2	ADK_MOUSE (+1)	40 kDa	1	21.712993	1.033333333	1	1.033333333	0.967741935
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm5611 PE=4 SV=1 (E9Q5T3_MOUSE [2])		30 kDa	1	19.239469	0.9	0.933333333	0.964285714	1.037037037
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 OS=Mus musculus GN=MNDUA5_MOUSE		13 kDa	1	21.889487	1.033333333	1.033333333	1	1
Cluster of sp K1C9_HUMAN	sp K1C9_HUMAN [2]	?	1	26.650304	1.266666667	1.433333333	0.88372093	1.131578947
Natriuretic peptides A OS=Mus musculus GN=Nppa PE=1 SV=2	ANF_MOUSE	17 kDa	1	21.745609	0.7	0.8	0.875	1.142857143
Cluster of Antithrombin-III OS=Mus musculus GN=Serpinc1 PE=1 SV=1 (AANT3_MOUSE)		52 kDa	1	23.352383	0.9	0.933333333	0.964285714	1.037037037
ATP synthase subunit e, mitochondrial OS=Mus musculus GN=Atp5i PE=1 SV=1 (ATP5I_MOUSE)		8 kDa	1	27.29676	1	0.9	1.111111111	0.9
14 kDa phosphohistidine phosphatase OS=Mus musculus GN=Phpt1 PE=1 SV=1 (PHP14_MOUSE)		14 kDa	1	20.79367	1.033333333	1.033333333	1	1
Transaldolase OS=Mus musculus GN=Taldo1 PE=1 SV=2	TALDO_MOUSE	37 kDa	1	19.607647	0.966666667	1.033333333	0.935483871	1.068965517
Cluster of Basigin OS=Mus musculus GN=Bsg PE=1 SV=2 (BASI_MOUSE) BASI_MOUSE [3]		42 kDa	1	0	1	1.15	0.869565217	1.15
Cluster of Kinectin OS=Mus musculus GN=Ktn1 PE=2 SV=1 (F8VQC7_MOUSE) MOI1F8VQC7_MOUSE [2]		153 kDa	1	0	1.1	1	1.1	0.909090909
Proteasome activator complex subunit 1 (Fragment) OS=Mus musculus GN=G3UXZ5_MOUSE (+1)		27 kDa	1	21.807472	1.2	1.133333333	1.058823529	0.944444444
60S ribosomal protein L3 OS=Mus musculus GN=Rpl3 PE=2 SV=3	RL3_MOUSE	46 kDa	1	7.8879684	0.8	0.8	1	1
Cluster of Heterogeneous nuclear ribonucleoprotein K OS=Mus musculus GN=B2M1R6_MOUSE [4]		49 kDa	1	19.67724	0.933333333	1	0.933333333	1.071428571
Cofilin-1 OS=Mus musculus GN=Cfl1 PE=1 SV=3	COF1_MOUSE (+1)	19 kDa	1	21.811543	0.766666667	0.9	0.851851852	1.173913043
DNA polymerase theta OS=Mus musculus GN=Polq PE=2 SV=1	Q80XB7_MOUSE	281 kDa	1	0	2	1.8	1.111111111	0.9
Cluster of Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit 1 OS=Mus musculus GN=PP2A_MOUSE		65 kDa	1	19.637708	1.033333333	1.1	0.939393939	1.064516129
Cluster of Methylglutaconyl-CoA hydratase, mitochondrial OS=Mus musculus GN=AUHM_MOUSE [2]		33 kDa	1	14.166689	1.066666667	1.133333333	0.941176471	1.0625
Cluster of CAP-Gly domain-containing linker protein 1 (Fragment) OS=Mus musculus GN=MtF7CB97_MOUSE [5]		136 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of Cathepsin D OS=Mus musculus GN=Ctsd PE=1 SV=1 (CATD_MOUSE) CATD_MOUSE [2]		45 kDa	1	20.738965	0.8	1	0.8	1.25
WD repeat-containing protein 1 OS=Mus musculus GN=Wdr1 PE=1 SV=3 (WDR1_MOUSE)		66 kDa	1	19.96053	0.966666667	0.933333333	1.035714286	0.965517241
Serine-rich coiled-coil domain-containing protein 1 OS=Mus musculus GN=CCSE1_MOUSE		98 kDa	1	8.25609	0.9	0.8	1.125	0.888888889

Destrin OS=Mus musculus GN=Dstn PE=1 SV=3	DEST_MOUSE	19 kDa	1	21.482916	0.933333333	1.06666667	0.875	1.142857143
Fibrous sheath-interacting protein 2 OS=Mus musculus GN=Fsp2 PE=1 S'FSIP2_MOUSE		785 kDa	1	0	1	1	1	1
GrpE protein homolog 1, mitochondrial OS=Mus musculus GN=Grpel1 PEGRPE1_MOUSE		24 kDa	1	20.165342	0.9	0.96666667	0.931034483	1.074074074
O-acetyl-ADP-ribose deacetylase MACROD1 OS=Mus musculus GN=Macr MACD1_MOUSE		35 kDa	1	22.148654	1	1.23333333	0.810810811	1.233333333
Isoleucine--tRNA ligase, mitochondrial OS=Mus musculus GN=lars2 PE=2 SYIM_MOUSE		113 kDa	1	20.477945	1	1.03333333	0.967741935	1.033333333
Beta-taxilin OS=Mus musculus GN=Txlnb PE=2 SV=2	TXLNB_MOUSE	77 kDa	1	20.961753	0.966666667	1.06666667	0.90625	1.103448276
AFG3-like protein 2 OS=Mus musculus GN=Afg3l2 PE=1 SV=1	AFG32_MOUSE	90 kDa	1	20.113319	0.8884	0.9	1	1
Cluster of MCG142036 OS=Mus musculus GN=Acad12 PE=3 SV=1 (D3Z7XD3Z7X0_MOUSE [2]		61 kDa	1	13.195859	1.133333333	1.1	1.03030303	0.970588235
Dynamin-1-like protein OS=Mus musculus GN=Dnm1l PE=1 SV=2	DNM1L_MOUSE (+1)	83 kDa	1	20.436788	1	1.03333333	0.967741935	1.033333333
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm18025 PE=E9Q1N8_MOUSE [5]		29 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Protein Myh15 OS=Mus musculus GN=Myh15 PE=4 SV=1	E9Q264_MOUSE	222 kDa	1	27.720067	1.333333333	1.23333333	1.081081081	0.925
Cluster of Gelsolin OS=Mus musculus GN=Gsn PE=1 SV=3 (GELS_MOUSE)GELS_MOUSE		86 kDa	1	21.374686	0.833333333	0.93333333	0.892857143	1.12
Cluster of Glutathione S-transferase A4 OS=Mus musculus GN=Gsta4 PE=GSTA4_MOUSE [3]		26 kDa	1	27.96852	0.933333333	1.03333333	0.903225806	1.107142857
Hydroxysteroid dehydrogenase-like protein 2 OS=Mus musculus GN=HscHSDL2_MOUSE		54 kDa	1	21.3858	0.966666667	1	0.966666667	1.034482759
Adenylate kinase 2, mitochondrial OS=Mus musculus GN=AK2 PE=1 SV=5 KAD2_MOUSE		26 kDa	1	20.725907	0.966666667	0.96666667	1	1
Ubiquitin-conjugating enzyme E2 N OS=Mus musculus GN=Ube2n PE=1 UBE2N_MOUSE		17 kDa	1	21.930379	0.9	1	0.9	1.111111111
Lactoylglutathione lyase OS=Mus musculus GN=Glo1 PE=1 SV=3	LGUL_MOUSE	21 kDa	1	21.034554	0.9	0.93333333	0.964285714	1.037037037
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Mus musculus GN=NDUB7_MOUSE		16 kDa	1	30.42457	1.2	1.2	1	1
Cluster of Mitogen-activated protein kinase 12 OS=Mus musculus GN=M MK12_MOUSE [2]		42 kDa	1	8.3054796	1.15	1.1	1.045454545	0.956521739
Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 A2AL78_MOUSE		26 kDa	1	20.37548	1.033333333	0.96666667	1.068965517	0.935483871
Band 3 anion transport protein OS=Mus musculus GN=Slc4a1 PE=1 SV=1 B3AT_MOUSE (+1)		103 kDa	1	20.700495	1.266666667	1.16666667	1.085714286	0.921052632
CDGSH iron-sulfur domain-containing protein 1 OS=Mus musculus GN=C CISD1_MOUSE		12 kDa	1	20.30046	1.1	1.16666667	0.942857143	1.060606061
Band 4.1-like protein 2 OS=Mus musculus GN=Epb4l2 PE=1 SV=2	E41L2_MOUSE	110 kDa	1	39.748502	1.3	1.23333333	1.054054054	0.948717949
Cluster of Protein Cald1 OS=Mus musculus GN=Cald1 PE=2 SV=1 (E9QA1 E9QA15_MOUSE [2]		89 kDa	1	30.69873	0.9	0.73333333	1.227272727	0.814814815
Eukaryotic translation initiation factor 3 subunit A OS=Mus musculus GN=EIF3A_MOUSE		162 kDa	1	23.554555	0.866666667	0.86666667	1	1
Cluster of Kinesin-1 heavy chain OS=Mus musculus GN=Kif5b PE=1 SV=3 KINH_MOUSE		110 kDa	1	29.820034	1.033333333	0.9	1.148148148	0.870967742
Spectrin beta 1 OS=Mus musculus GN=Sptb PE=2 SV=1	Q3UGX2_MOUSE	268 kDa	1	23.465471	1.066666667	1.1	0.96969697	1.03125
Cluster of Thioredoxin OS=Mus musculus GN=Txn PE=1 SV=3 (THIO_MOUSE)THIO_MOUSE		12 kDa	1	25.592077	0.9	1	0.9	1.111111111
Cluster of Acidic leucine-rich nuclear phosphoprotein 32 family member AN32A_MOUSE [3]		29 kDa	1	29.564816	1	0.9	1.111111111	0.9
Branched-chain-amino-acid aminotransferase, mitochondrial OS=Mus musculus GN=BCAT2_MOUSE		44 kDa	1	22.588815	0.866666667	0.9	0.962962963	1.038461538
Xin actin-binding repeat-containing protein 1 OS=Mus musculus GN=Xirp E9QQ93_MOUSE		124 kDa	1	21.231122	0.866666667	0.8	1.083333333	0.923076923
Protein unc-45 homolog B OS=Mus musculus GN=Unc45b PE=1 SV=1	UN45B_MOUSE	104 kDa	1	21.598138	0.9	1.06666667	0.84375	1.185185185
Vigilin OS=Mus musculus GN=Hdlbp PE=1 SV=1	VIGLN_MOUSE	142 kDa	1	14.61967	1.1	1.03333333	1.064516129	0.939393939
Isocitrate dehydrogenase [NADP] cytoplasmic OS=Mus musculus GN=IdhIDHC_MOUSE		47 kDa	1	21.648483	1.2	1.1	1.090909091	0.916666667
LETM1 and EF-hand domain-containing protein 1, mitochondrial OS=Mus musculus GN=LETM1_MOUSE		83 kDa	1	23.618585	1.066666667	1.1	0.96969697	1.03125
NADH dehydrogenase [ubiquinone] iron-sulfur protein 7, mitochondrial OS=Mus musculus GN=NDUS7_MOUSE		25 kDa	1	20.107381	1.033333333	1.1	0.939393939	1.064516129
Ornithine aminotransferase, mitochondrial OS=Mus musculus GN=Oat P OAT_MOUSE		48 kDa	1	22.667679	0.966666667	0.96666667	1	1
Cluster of 40S ribosomal protein S9 OS=Mus musculus GN=Rps9 PE=2 SV RS9_MOUSE [4]		23 kDa	1	20.292385	1.133333333	1.23333333	0.918918919	1.088235294
Succinate-semialdehyde dehydrogenase, mitochondrial OS=Mus musculus GN=SSDH_MOUSE		56 kDa	1	21.072468	1.033333333	1.1	0.939393939	1.064516129
Acylophosphatase-2 OS=Mus musculus GN=Acyp2 PE=2 SV=2	ACYP2_MOUSE	12 kDa	1	23.077793	1.066666667	0.9	1.185185185	0.84375
Cluster of Caveolin-3 OS=Mus musculus GN=Cav3 PE=1 SV=1 (CAV3_MOUSE)CAV3_MOUSE [3]		17 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
DnaJ homolog subfamily A member 2 OS=Mus musculus GN=Dnaja2 PE=DNJA2_MOUSE		46 kDa	1	20.327157	0.933333333	0.96666667	0.965517241	1.035714286
Myosin-14 OS=Mus musculus GN=Myh14 PE=4 SV=1	K3W4R2_MOUSE (+1)	229 kDa	1	23.060788	1.066666667	1.13333333	0.941176471	1.0625
Amine oxidase [flavin-containing] B OS=Mus musculus GN=Maob PE=1 S'AOFB_MOUSE		59 kDa	1	20.96426	1.233333333	1.43333333	0.860465116	1.162162162
Protein Lmo7 OS=Mus musculus GN=Lmo7 PE=2 SV=1	E9PYF4_MOUSE	193 kDa	1	21.196204	1.033333333	1.03333333	1	1
Fibrinogen beta chain OS=Mus musculus GN=Fgb PE=2 SV=1	FIBB_MOUSE	55 kDa	1	20.901106	1	1	1	1
Glutathione S-transferase P 1 OS=Mus musculus GN=Gstp1 PE=1 SV=2	GSTP1_MOUSE	24 kDa	1	22.094472	0.966666667	1.03333333	0.935483871	1.068965517
Puromycin-sensitive aminopeptidase OS=Mus musculus GN=Npepps PE=PSA_MOUSE		103 kDa	1	21.59204	1.066666667	1.2	0.888888889	1.125
Regulator of microtubule dynamics protein 1 OS=Mus musculus GN=Rm1 RMD1_MOUSE		35 kDa	1	21.094505	1	1.16666667	0.857142857	1.166666667
Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=DPYL2_MOUSE		62 kDa	1	22.084574	0.933333333	0.96666667	0.965517241	1.035714286
Cluster of Ceruloplasmin OS=Mus musculus GN=Cp PE=4 SV=1 (G3X8Q5_G3X8Q5_MOUSE [4]		124 kDa	1	44.992045	0.766666667	0.83333333	0.92	1.086956522
Glutathione S-transferase kappa 1 OS=Mus musculus GN=Gstk1 PE=1 SV GSTK1_MOUSE		26 kDa	1	21.262702	1.1	1.2	0.916666667	1.090909091
Cluster of Sarcolemmal membrane-associated protein OS=Mus musculus GN=H7BX64_MOUSE [3]		95 kDa	1	0	0.9	0.95	0.947368421	1.055555556

Mitochondrial carnitine/acylcarnitine carrier protein OS=Mus musculus (MCAT_MOUSE	33 kDa	1	21.43621	1.1	1.2	0.916666667	1.090909091
Methylmalonyl-CoA mutase, mitochondrial OS=Mus musculus GN=Mut f MUTA_MOUSE	83 kDa	1	23.142092	1.066666667	1.133333333	0.941176471	1.0625
Proteasome subunit alpha type-1 OS=Mus musculus GN=Psm1 PE=1 SV PSA1_MOUSE	30 kDa	1	21.214527	0.866666667	0.933333333	0.928571429	1.076923077
Spectrin alpha chain, erythrocytic 1 OS=Mus musculus GN=Spta1 PE=2 SV SPTA1_MOUSE	280 kDa	1	22.224896	1.033333333	0.966666667	1.068965517	0.935483871
Short/branched chain specific acyl-CoA dehydrogenase, mitochondrial O ACDSB_MOUSE	48 kDa	1	20.524439	1.166666667	1.166666667	1	1
Apolipoprotein A-IV OS=Mus musculus GN=Apoa4 PE=2 SV=3 APOA4_MOUSE	45 kDa	1	20.295446	1.033333333	1	1.033333333	0.967741935
Cluster of F-actin-capping protein subunit beta OS=Mus musculus GN=Cc CAPZB_MOUSE [2]	31 kDa	1	0	1.033333333	1	1.033333333	0.967741935
Cofilin-2 OS=Mus musculus GN=Cfl2 PE=1 SV=1 COF2_MOUSE	19 kDa	1	19.80154	0.817533333	0.766666667	1.043478261	0.958333333
Phospholipid hydroperoxide glutathione peroxidase, nuclear OS=Mus musculus GN=GPX42_MOUSE (+1)	29 kDa	1	20.668325	0.966666667	0.933333333	1.035714286	0.965517241
Decorin OS=Mus musculus GN=Dcn PE=2 SV=1 PGS2_MOUSE	40 kDa	1	19.844672	1	1	1	1
Ryanodine receptor 3 OS=Mus musculus GN=Ryr3 PE=1 SV=1 RYR3_MOUSE (+1)	551 kDa	1	0	1	1.1	0.909090909	1.1
Tropomyosin alpha-4 chain OS=Mus musculus GN=Tpm4 PE=2 SV=3 TPM4_MOUSE	28 kDa	1	22.798583	0.833333333	0.7	1.19047619	0.84
LIM and calponin homology domains-containing protein 1 OS=Mus musculus GN=D3YU22_MOUSE (+1)	120 kDa	1	27.701235	1.066666667	1.033333333	1.032258065	0.96875
Cluster of Synaptopodin-2 OS=Mus musculus GN=Synpo2 PE=2 SV=1 (E9)E9Q1U2_MOUSE [2]	136 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Ferrochelatase, mitochondrial OS=Mus musculus GN=Fech PE=1 SV=2 HEMH_MOUSE (+1)	47 kDa	1	21.199597	1	1	1	1
Histidine triad nucleotide-binding protein 2, mitochondrial OS=Mus musculus GN=HINT2_MOUSE	17 kDa	1	20.254165	1.1	1.033333333	1.064516129	0.939393939
Heterochromatin protein 1-binding protein 3 OS=Mus musculus GN=Hp1HP1B3_MOUSE	61 kDa	1	6.35594	0.966666667	1.233333333	0.783783784	1.275862069
Adenylosuccinate synthetase OS=Mus musculus GN=Adss1 PE=3 SV=1 J3QN31_MOUSE (+1)	53 kDa	1	21.082275	1	1.133333333	0.882352941	1.133333333
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 OS=Mus musculus GN=NDUB4_MOUSE	15 kDa	1	22.795871	1	1.1	0.909090909	1.1
Cluster of 26S proteasome non-ATPase regulatory subunit 2 OS=Mus musculus GN=PSMD2_MOUSE [2]	100 kDa	1	14.183229	0.933333333	1	0.933333333	1.071428571
Ig mu chain C region secreted form OS=Mus musculus GN=Igh-6 PE=1 SV IGHM_MOUSE	50 kDa	1	20.744331	0.933333333	0.8	1.166666667	0.857142857
Sulfated glycoprotein 1 OS=Mus musculus GN=Psap PE=4 SV=1 J3QPG5_MOUSE (+3)	61 kDa	1	23.429663	1.1	1.133333333	0.970588235	1.03030303
Lipoamide acyltransferase component of branched-chain alpha-keto acid ODB2_MOUSE	53 kDa	1	21.46635	1.033333333	1.1	0.939393939	1.064516129
Cluster of Elongation factor 1-delta (Fragment) OS=Mus musculus GN=Ee1E9QN08_MOUSE [3]	27 kDa	1	22.249684	0.866666667	0.866666667	1	1
Nicotinamide phosphoribosyltransferase OS=Mus musculus GN=Nampt1 NAMPT_MOUSE	55 kDa	1	24.045411	1.266666667	1.3	0.974358974	1.026315789
Cluster of Palmidolphin OS=Mus musculus GN=Palmd PE=2 SV=1 (Q3UVT Q3UVT7_MOUSE [2]	63 kDa	1	0	0.75	1.05	0.714285714	1.4
T-complex protein 1 subunit epsilon OS=Mus musculus GN=Cct5 PE=1 SV TPCPE_MOUSE	60 kDa	1	19.240971	0.866666667	0.933333333	0.928571429	1.076923077
Cluster of MCG130675 OS=Mus musculus GN=2410018M08Rik PE=2 SV=B2RPU8_MOUSE [2]	84 kDa	1	12.4948	0.8	0.75	1.066666667	0.9375
Cluster of Nuclease-sensitive element-binding protein 1 (Fragment) OS=Mus musculus GN=A2BGG7_MOUSE [2]	25 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
ADP/ATP translocase 4 OS=Mus musculus GN=Slc25a31 PE=2 SV=1 ADT4_MOUSE	35 kDa	1	23.385182	1.133333333	1	1.133333333	0.882352941
Cadherin-13 OS=Mus musculus GN=Cdh13 PE=1 SV=2 CAD13_MOUSE	78 kDa	1	21.390262	1.3	1.166666667	1.114285714	0.897435897
Calnexin OS=Mus musculus GN=Canx PE=1 SV=1 CALX_MOUSE	67 kDa	1	20.065323	1.133333333	1.233333333	0.918918919	1.088235294
Fibrinogen gamma chain OS=Mus musculus GN=Fgg PE=2 SV=1 FIBG_MOUSE	49 kDa	1	20.287552	1.033333333	1.133333333	0.911764706	1.096774194
Fibronectin OS=Mus musculus GN=Fn1 PE=1 SV=4 FINC_MOUSE	273 kDa	1	20.054449	0.933333333	0.966666667	0.965517241	1.035714286
Cluster of cAMP-dependent protein kinase catalytic subunit alpha OS=Mus musculus GN=KAPCA_MOUSE [2]	41 kDa	1	18.989395	1.066666667	1.2	0.888888889	1.125
Cluster of 60S ribosomal protein L24 OS=Mus musculus GN=Rpl24 PE=2 SV RL24_MOUSE	18 kDa	1	22.80761	0.9	1.066666667	0.84375	1.185185185
40S ribosomal protein S6 OS=Mus musculus GN=Rps6 PE=1 SV=1 RS6_MOUSE	29 kDa	1	20.752513	0.933333333	1.1	0.848484848	1.178571429
Transthyretin OS=Mus musculus GN=Ttr PE=1 SV=1 TTHY_MOUSE	16 kDa	1	20.514244	0.966666667	0.833333333	1.16	0.862068966
Cluster of Archvillin OS=Mus musculus GN=Svil PE=2 SV=1 (Q8K4L2_MOUSE Q8K4L2_MOUSE	227 kDa	1	14.063953	0.933333333	0.866666667	1.076923077	0.928571429
Cullin-associated NEDD8-dissociated protein 2 OS=Mus musculus GN=Ca CAND2_MOUSE	136 kDa	1	20.93455	1.033333333	1.066666667	0.96875	1.032258065
Histone-lysine N-methyltransferase, H3 lysine-36 and H4 lysine-20-specific E9QAE4_MOUSE	296 kDa	1	0	0.9	0.8	1.125	0.888888889
Receptor accessory protein 5, isoform CRA_a OS=Mus musculus GN=Ree G3X8R0_MOUSE	21 kDa	1	23.310374	1	1.166666667	0.857142857	1.166666667
Laminin subunit beta-2 OS=Mus musculus GN=Lamb2 PE=2 SV=2 LAMB2_MOUSE	197 kDa	1	20.518704	0.966666667	0.866666667	1.115384615	0.896551724
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 11, mitochondrial NDUBB_MOUSE	17 kDa	1	39.40191	0.9	0.733333333	1.227272727	0.814814815
Vitamin D-binding protein OS=Mus musculus GN=Gc PE=1 SV=2 VTDB_MOUSE	54 kDa	1	20.246282	0.9	0.9	1	1
Probable C->U-editing enzyme APOBEC-2 OS=Mus musculus GN=ApoBec ABEC2_MOUSE	26 kDa	1	22.58346	0.933333333	1.133333333	0.823529412	1.214285714
Cluster of Acyl-coenzyme A thioesterase 9, mitochondrial OS=Mus musculus GN=ACOT9_MOUSE [2]	51 kDa	1	6.115666	1.033333333	0.966666667	1.068965517	0.935483871
Cluster of ATPase family AAA domain-containing protein 3 OS=Mus musculus GN=ATAD3_MOUSE	67 kDa	1	21.057509	0.966666667	0.966666667	1	1
Elongation factor 1-alpha 1 OS=Mus musculus GN=Eef1a1 PE=1 SV=3 EF1A1_MOUSE	50 kDa	1	23.668197	0.733333333	0.733333333	1	1
Acylpyruvate FAHD1, mitochondrial OS=Mus musculus GN=FAhd1 PE=1 SV FAHD1_MOUSE	25 kDa	1	60.405296	1.133333333	1.3	0.871794872	1.147058824
Ferritin heavy chain OS=Mus musculus GN=Fth1 PE=1 SV=2 FRIH_MOUSE	21 kDa	1	19.612518	0.9	0.933333333	0.964285714	1.037037037
Cluster of Eukaryotic initiation factor 4A-II OS=Mus musculus GN=Eif4a2 IF4A2_MOUSE [2]	46 kDa	1	22.808841	1.066666667	1.066666667	1	1
Eukaryotic translation initiation factor 4B OS=Mus musculus GN=Eif4b EIF4B_MOUSE	69 kDa	1	32.508352	0.8	0.766666667	1.043478261	0.958333333

Adenylosuccinate lyase OS=Mus musculus GN=Adsl PE=2 SV=2	PUR8_MOUSE	55 kDa	1	25.604364	0.966666667	1.1	0.878787879	1.137931034
Peripherin OS=Mus musculus GN=Prph PE=1 SV=2	PERI_MOUSE (+1)	54 kDa	1	0	1	1	1	1
[Protein ADP-ribosylarginine] hydrolase-like protein 1 OS=Mus musculus ARHL1_MOUSE	ARHL1_MOUSE	40 kDa	1	21.692301	1.033333333	0.966666667	1.068965517	0.935483871
Actin-related protein 2/3 complex subunit 4 OS=Mus musculus GN=Arpc ARPC4_MOUSE	ARPC4_MOUSE	20 kDa	1	22.25852	0.966666667	1	0.966666667	1.034482759
Transcription elongation factor B polypeptide 2 OS=Mus musculus GN=T ELOB_MOUSE	ELOB_MOUSE	13 kDa	1	20.435576	1	1	1	1
Serine/arginine-rich-splicing factor 1 OS=Mus musculus GN=Srf1 PE=2 S H7BX95_MOUSE (+1)	H7BX95_MOUSE (+1)	28 kDa	1	22.638899	1.033333333	1	1.033333333	0.967741935
Laminin subunit alpha-4 OS=Mus musculus GN=Lama4 PE=1 SV=2	LAMA4_MOUSE	202 kDa	1	21.065247	1	0.966666667	1.034482759	0.966666667
Mitochondrial fission process protein 1 OS=Mus musculus GN=Mtff1 PE=1 MTFP1_MOUSE	MTFP1_MOUSE	18 kDa	1	21.10341	0.966666667	0.966666667	1	1
Prostaglandin reductase 2 OS=Mus musculus GN=Ptgr2 PE=1 SV=2	PTGR2_MOUSE	38 kDa	1	20.81221	0.9	1.066666667	0.84375	1.185185185
Protein Sf3b2 OS=Mus musculus GN=Sf3b2 PE=2 SV=1	Q3UJB0_MOUSE	98 kDa	1	16.187775	0.75	0.85	0.882352941	1.133333333
40S ribosomal protein S12 OS=Mus musculus GN=Rps12 PE=2 SV=1	Q6ZWZ6_MOUSE	15 kDa	1	22.52498	0.833333333	0.933333333	0.892857143	1.12
26S protease regulatory subunit 7 OS=Mus musculus GN=Psmc2 PE=2 SV=1 Q8BVQ9_MOUSE	Q8BVQ9_MOUSE	53 kDa	1	22.642271	0.933333333	0.966666667	0.965517241	1.035714286
60S ribosomal protein L5 OS=Mus musculus GN=Rpl5 PE=1 SV=3	RL5_MOUSE	34 kDa	1	19.330601	0.966666667	1.033333333	0.935483871	1.068965517
Transcription factor A, mitochondrial OS=Mus musculus GN=Tfam PE=1 TFAM_MOUSE	TFAM_MOUSE	28 kDa	1	19.59383	0.966666667	0.933333333	1.035714286	0.965517241
Glycogenin-1 OS=Mus musculus GN=Gyg1 PE=2 SV=3	GLYG_MOUSE (+1)	37 kDa	1	21.226205	0.966666667	0.9	1.074074111	0.931034451
Muscle-related coiled-coil protein OS=Mus musculus GN=Murc PE=2 SV=1 MURC_MOUSE	MURC_MOUSE	41 kDa	1	21.631828	1.066666667	0.933333333	1.142857143	0.875
Cluster of Myosin light polypeptide 6 OS=Mus musculus GN=Myl6 PE=1 MYL6_MOUSE [2]	MYL6_MOUSE [2]	17 kDa	1	23.603867	0.9	0.766666667	1.173913043	0.851851852
26S protease regulatory subunit 4 OS=Mus musculus GN=Psmc1 PE=1 SV=1 SVPRS4_MOUSE	SVPRS4_MOUSE	49 kDa	1	47.199312	1.033333333	1.1	0.939393939	1.064516129
26S proteasome non-ATPase regulatory subunit 1 OS=Mus musculus GN=PSMD1_MOUSE	PSMD1_MOUSE	106 kDa	1	19.74388	0.9	1.033333333	0.870967742	1.148148148
Cluster of Prothymosin alpha OS=Mus musculus GN=Ptma PE=1 SV=2 (P1PTMA_MOUSE	P1PTMA_MOUSE	12 kDa	1	21.409719	1	0.833333333	1.2	0.833333333
Bifunctional purine biosynthesis protein PURH OS=Mus musculus GN=At PUR9_MOUSE	PUR9_MOUSE	64 kDa	1	14.661838	0.9	1	0.9	1.111111111
Cluster of Protein kinase, cAMP dependent regulatory, type II alpha OS=Q8K1M3_MOUSE	Q8K1M3_MOUSE	46 kDa	1	20.898531	1	1.033333333	0.967741935	1.033333333
Aspartate--tRNA ligase, cytoplasmic OS=Mus musculus GN=Dars PE=2 SV=1 SVYDC_MOUSE	SVYDC_MOUSE	57 kDa	1	20.832342	1	1	1	1
Cluster of Elongation factor G, mitochondrial OS=Mus musculus GN=Gfm EFGM_MOUSE	EFGM_MOUSE	84 kDa	1	13.963708	0.933333333	1	0.933333333	1.071428571
Cluster of Protein Gm4945 OS=Mus musculus GN=Gm4945 PE=4 SV=1 (D3Z102_MOUSE	D3Z102_MOUSE	16 kDa	1	19.438279	0.966666667	1.066666667	0.90625	1.103448276
Cluster of Protein Etl4 OS=Mus musculus GN=Etl4 PE=2 SV=1 (E9QAU4_MOUSE	E9QAU4_MOUSE	218 kDa	1	6.695105	1	1	1	1
Coiled-coil protein associated with myosin II and DISC1 OS=Mus musculus A2AST1_MOUSE (+1)	A2AST1_MOUSE (+1)	165 kDa	1	12.835398	1.033333333	1.1	0.939393939	1.064516129
Cluster of Rab GDP dissociation inhibitor beta OS=Mus musculus GN=Gd GDIB_MOUSE [2]	GDIB_MOUSE [2]	51 kDa	1	24.443437	1.066666667	1.066666667	1	1
Cluster of Glyoxalase domain-containing protein 4 OS=Mus musculus GN=GLOD4_MOUSE [2]	GLOD4_MOUSE [2]	33 kDa	1	21.97495	1.133333333	1.233333333	0.918918919	1.088235294
Eukaryotic translation initiation factor 5A-1 OS=Mus musculus GN=Eif5a IF5A1_MOUSE	IF5A1_MOUSE	17 kDa	1	20.03944	0.866666667	0.833333333	1.04	0.961538462
Junctophilin-2 OS=Mus musculus GN=Jph2 PE=1 SV=2	JPH2_MOUSE	75 kDa	1	22.219017	1.1	1	1.1	0.909090909
Cluster of Poly(rC)-binding protein 1 OS=Mus musculus GN=Pcbp1 PE=1 PCBP1_MOUSE [4]	PCBP1_MOUSE [4]	37 kDa	1	35.70203	1.1	1.033333333	1.064516129	0.939393939
Programmed cell death 6-interacting protein OS=Mus musculus GN=Pdc1 PDC6_MOUSE	PDC6_MOUSE	96 kDa	1	22.275603	0.966666667	1	0.966666667	1.034482759
Cluster of Ank1 protein OS=Mus musculus GN=Ank1 PE=2 SV=1 (Q0VGY9_MOUSE [3]	Q0VGY9_MOUSE [3]	207 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Sorting and assembly machinery component 50 homolog OS=Mus musculus SAM50_MOUSE	SAM50_MOUSE	52 kDa	1	47.73003	1.1	1.266666667	0.868421053	1.151515152
T-complex protein 1 subunit theta OS=Mus musculus GN=Cct8 PE=1 SV=1 TCPQ_MOUSE	TCPQ_MOUSE	60 kDa	1	20.282198	0.933333333	1	0.933333333	1.071428571
Cluster of Muscular LMNA-interacting protein OS=Mus musculus GN=Ml tr V9GWW6 V9GWW6_MOUSE	V9GWW6_MOUSE	?	1	22.278617	0.933333333	0.9	1.037037037	0.964285714
Cluster of Isoform 2 of Protein phosphatase 1 regulatory subunit 12A OS=sp Q9DBR7-2 MYPT1_MOUSE	Q9DBR7-2_MOUSE	?	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Mitochondrial pyruvate carrier 2 OS=Mus musculus GN=Mpc2 PE=1 SV=1 MPC2_MOUSE	MPC2_MOUSE	14 kDa	1	22.577049	1.1	1.366666667	0.804878049	1.242424242
Cluster of Replication factor C subunit 1 OS=Mus musculus GN=Rfc1 PE=1 G3UWX1_MOUSE [6]	G3UWX1_MOUSE [6]	126 kDa	1	7.6247565	1.1	1.1	1	1
Cluster of Calumenin OS=Mus musculus GN=Calu PE=1 SV=1 (CALU_MOUSE	CALU_MOUSE	37 kDa	1	20.295848	0.933333333	0.866666667	1.076923077	0.928571429
60S acidic ribosomal protein P0 (Fragment) OS=Mus musculus GN=Rplp0 D3YVM5_MOUSE (+2)	D3YVM5_MOUSE (+2)	16 kDa	1	21.167445	0.933333333	1.033333333	0.903225806	1.107142857
Filamin-B OS=Mus musculus GN=Flnb PE=1 SV=3	FLNB_MOUSE	278 kDa	1	13.790981	1.066666667	1.033333333	1.032258065	0.96875
Pyruvate carboxylase OS=Mus musculus GN=Pcx PE=3 SV=1	G5E8R3_MOUSE (+1)	130 kDa	1	20.360175	1	0.866666667	1.153846154	0.866666667
Unconventional myosin-1c OS=Mus musculus GN=Myo1c PE=1 SV=2	MYO1C_MOUSE (+3)	122 kDa	1	20.641302	1.033333333	1.1	0.939393939	1.064516129
26S protease regulatory subunit 6A OS=Mus musculus GN=Psmc3 PE=1 SPRS6A_MOUSE	SPRS6A_MOUSE	50 kDa	1	23.494495	0.933333333	0.866666667	1.076923077	0.928571429
Radixin OS=Mus musculus GN=Rdx PE=1 SV=3	RADI_MOUSE	69 kDa	1	26.30181	0.933333333	1	0.933333333	1.071428571
60S ribosomal protein L8 OS=Mus musculus GN=Rpl8 PE=2 SV=2	RL8_MOUSE	28 kDa	1	20.572698	0.933333333	1	0.933333333	1.071428571
Bifunctional glutamate/proline--tRNA ligase OS=Mus musculus GN=Eprs SYEP_MOUSE	SYEP_MOUSE	170 kDa	1	19.91574	0.933333333	0.9	1.037037037	0.964285714
T-complex protein 1 subunit eta OS=Mus musculus GN=Cct7 PE=1 SV=1 TCPH_MOUSE	TCPH_MOUSE	60 kDa	1	20.280574	0.966666667	1.033333333	0.935483871	1.068965517
Cluster of STE20-like serine/threonine-protein kinase OS=Mus musculus SLK_MOUSE [2]	SLK_MOUSE [2]	141 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm10260 PE=1 F6YVP7_MOUSE [2]	F6YVP7_MOUSE [2]	18 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of 5-azacytidine induced gene 1 OS=Mus musculus GN=Azi1 PE=2 B1AXI9_MOUSE [2]	B1AXI9_MOUSE [2]	120 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!

Eukaryotic translation initiation factor 3 subunit G OS=Mus musculus GNEIF3G_MOUSE	36 kDa	1	19.377226	0.933333333	0.966666667	0.965517241	1.035714286
Rho GDP-dissociation inhibitor 1 OS=Mus musculus GN=Arhgdia PE=1 SV=1 GDIR1_MOUSE	23 kDa	1	22.530189	0.9	0.966666667	0.931034483	1.074074074
60S ribosomal protein L10 (Fragment) OS=Mus musculus GN=Rpl10 PE=4 I7HLV2_MOUSE (+1)	23 kDa	1	21.37207	0.966666667	1.1	0.878787879	1.137931034
Lipoprotein lipase OS=Mus musculus GN=Lpl PE=1 SV=3 LPL_MOUSE	53 kDa	1	19.726844	1.033333333	1	1.033333333	0.967741935
Myosin-10 OS=Mus musculus GN=Myh10 PE=1 SV=2 MYH10_MOUSE (+2)	229 kDa	1	10.964391	1.033333333	0.866666667	1.192307692	0.838709677
Nuclear pore complex-associated intranuclear coiled-coil protein TPR OS Q7M739_MOUSE	267 kDa	1	16.740105	0.866666667	0.833333333	1.04	0.961538462
60S ribosomal protein L31 OS=Mus musculus GN=Rpl31 PE=2 SV=1 RL31_MOUSE	14 kDa	1	22.355245	0.9	1.033333333	0.870967742	1.148148148
40S ribosomal protein S3 OS=Mus musculus GN=Rps3 PE=1 SV=1 RS3_MOUSE	27 kDa	1	20.922197	0.933333333	1	0.933333333	1.071428571
Adenosylhomocysteinase OS=Mus musculus GN=Ahcy PE=1 SV=3 SAHH_MOUSE	48 kDa	1	19.621514	1	1	1	1
T-complex protein 1 subunit alpha OS=Mus musculus GN=Tcp1 PE=1 SV=1 TCPA_MOUSE	60 kDa	1	22.400063	0.933333333	0.9	1.037037037	0.964285714
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 3 OS=Mt NDUB3_MOUSE	12 kDa	1	26.06619	1	0.933333333	1.071428571	0.933333333
UPF0317 protein C14orf159 homolog, mitochondrial OS=Mus musculus ICN159_MOUSE (+1)	66 kDa	1	23.33662	1.133333333	1.133333333	1	1
Collagen alpha-2(VI) chain OS=Mus musculus GN=Col6a2 PE=2 SV=3 CO6A2_MOUSE	110 kDa	1	20.580691	1.366666667	1.133333333	1.205882353	0.829268293
60S ribosomal protein L13a OS=Mus musculus GN=Rpl13a PE=2 SV=1 E9Q5A0_MOUSE	48 kDa	1	19.168493	0.966666667	1.133333333	0.852941176	1.172413793
Kinesin-like protein KIF21B OS=Mus musculus GN=Kif21b PE=2 SV=1 F8VQE2_MOUSE (+1)	181 kDa	1	8.172655	1.1	1.1	1	1
Histone H1.1 OS=Mus musculus GN=Hist1h1a PE=1 SV=2 H11_MOUSE	22 kDa	1	9.469792	0.9	1	0.9	1.111111111
GTP:AMP phosphotransferase, mitochondrial OS=Mus musculus GN=Ak2KAD3_MOUSE	25 kDa	1	20.30093	0.966666667	0.966666667	1	1
Cluster of NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit NDUB8_MOUSE	22 kDa	1	21.303912	1	1.166666667	0.857142857	1.166666667
60S ribosomal protein L13 OS=Mus musculus GN=Rpl13 PE=2 SV=3 RL13_MOUSE	24 kDa	1	20.44002	0.933333333	1.1	0.848484848	1.178571429
T-complex protein 1 subunit beta OS=Mus musculus GN=Cct2 PE=1 SV=4 TCPB_MOUSE	57 kDa	1	46.331263	1	1.033333333	0.967741935	1.033333333
Mitochondrial import inner membrane translocase subunit TIM44 OS=MTIM44_MOUSE	51 kDa	1	20.449664	1.2	1.1	1.090909091	0.916666667
Transketolase OS=Mus musculus GN=Tkt PE=1 SV=1 TKT_MOUSE	68 kDa	1	21.893649	0.9	1.033333333	0.870967742	1.148148148
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 6 OS=Mt A2AP32_MOUSE (+1)	12 kDa	1	14.414687	1.266666667	1.066666667	1.1875	0.842105263
Cluster of Mitochondrial pyruvate carrier 1 OS=Mus musculus GN=Mpc1 MPC1_MOUSE [3]	12 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of Probable helicase senataxin OS=Mus musculus GN=Setx PE=2 SETX_MOUSE	298 kDa	1	6.5563515	1.3	1.3	1	1
Ubiquinone biosynthesis monooxygenase COQ6 OS=Mus musculus GN=COQ6_MOUSE	51 kDa	1	23.105465	1	1.1	0.909090909	1.1
Cluster of D-dopachrome decarboxylase OS=Mus musculus GN=Ddt PE=1 DOPD_MOUSE	13 kDa	1	25.325394	0.933333333	0.9	1.037037037	0.964285714
MCG116065 OS=Mus musculus GN=Gm6472 PE=4 SV=1 J3QJZ3_MOUSE (+1)	22 kDa	1	19.710933	0.9	0.966666667	0.931034483	1.074074074
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Mt NDUAC_MOUSE	17 kDa	1	21.706855	0.933333333	0.933333333	1	1
Plasminogen OS=Mus musculus GN=Plg PE=1 SV=3 PLMN_MOUSE	91 kDa	1	20.636302	0.933333333	0.933333333	1	1
Cluster of Serine/threonine-protein phosphatase PP1-alpha catalytic subunit PP1A_MOUSE [4]	38 kDa	1	0	0.9	1	0.9	1.111111111
60S ribosomal protein L27 OS=Mus musculus GN=Rpl27 PE=2 SV=2 RL27_MOUSE	16 kDa	1	20.888783	0.513333333	0.966666667	0.531034479	1.883116895
40S ribosomal protein S25 OS=Mus musculus GN=Rps25 PE=2 SV=1 RS25_MOUSE	14 kDa	1	24.065381	1	1.2	0.833333333	1.2
Heat shock protein 75 kDa, mitochondrial OS=Mus musculus GN=Trap1 FTRAP1_MOUSE	80 kDa	1	21.167408	0.966666667	1.066666667	0.90625	1.103448276
Zinc-binding alcohol dehydrogenase domain-containing protein 2 OS=Mt ZADH2_MOUSE	41 kDa	1	22.58135	1	1.1	0.909090909	1.1
Synaptopodin 2-like protein OS=Mus musculus GN=Synpo2l PE=2 SV=1 B2RQK7_MOUSE	103 kDa	1	14.94315	0.9	1	0.9	1.111111111
40S ribosomal protein S5 (Fragment) OS=Mus musculus GN=Rps5 PE=3 SD3YYM6_MOUSE (+2)	20 kDa	1	7.213274	0.95	0.95	1	1
Dystroglycan OS=Mus musculus GN=Dag1 PE=1 SV=4 DAG1_MOUSE	97 kDa	1	21.011098	1.033333333	1.066666667	0.96875	1.032258065
Eukaryotic translation initiation factor 4 gamma 1 OS=Mus musculus GN=E9PVC5_MOUSE	175 kDa	1	22.298212	0.966666667	1	0.966666667	1.034482759
Ras-related protein Rab-11A OS=Mus musculus GN=Rab11a PE=2 SV=1 E9Q3P9_MOUSE (+4)	17 kDa	1	21.746005	1.1	1.133333333	0.970588235	1.03030303
Cluster of Protein Serpinb6a OS=Mus musculus GN=Serpnb6a PE=2 SV=1 F8WIV2_MOUSE [3]	45 kDa	1	20.189424	0.833333333	1.066666667	0.78125	1.28
Four and a half LIM domains protein 1 OS=Mus musculus GN=Fhl1 PE=2 FHL1_MOUSE	32 kDa	1	21.963012	0.866666667	0.966666667	0.896551724	1.115384615
Glutaryl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Gcdl GCDH_MOUSE	49 kDa	1	24.420924	1.033333333	1.166666667	0.885714286	1.129032258
Importin-5 OS=Mus musculus GN=Ipo5 PE=1 SV=3 IPO5_MOUSE	124 kDa	1	25.379056	1	1.166666667	0.857142857	1.166666667
MAP7 domain-containing protein 1 OS=Mus musculus GN=Map7d1 PE=1 IMA7D1_MOUSE	93 kDa	1	18.271294	0.8	0.833333333	0.96	1.041666667
Cluster of Myosin regulatory light chain 12B OS=Mus musculus GN=Myl1 ML12B_MOUSE [3]	20 kDa	1	21.92504	0.833333333	0.7	1.19047619	0.84
Myosin light chain kinase 3 OS=Mus musculus GN=Mylk3 PE=1 SV=1 MYLK3_MOUSE	86 kDa	1	21.65973	0.9	1.033333333	0.870967742	1.148148148
Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform PP2AA_MOUSE	36 kDa	1	25.095513	0.866666667	1.033333333	0.838709677	1.192307692
60S ribosomal protein L10a OS=Mus musculus GN=Rpl10a PE=1 SV=3 RL10A_MOUSE	25 kDa	1	20.442341	1	1.066666667	0.9375	1.066666667
Cluster of T-complex protein 1 subunit delta OS=Mus musculus GN=Cct4 TCPD_MOUSE [2]	58 kDa	1	0	1.05	0.9	1.166666667	0.857142857
Cluster of NEDD8-conjugating enzyme Ubc12 OS=Mus musculus GN=Ubc12 UBC12_MOUSE	21 kDa	1	19.828305	0.966666667	0.933333333	1.035714286	0.965517241
Cluster of Estradiol 17-beta-dehydrogenase 8 OS=Mus musculus GN=HscDH8_MOUSE [2]	27 kDa	1	21.343417	0.933333333	1	0.933333333	1.071428571
Cluster of NSFL1 cofactor p47 OS=Mus musculus GN=Nsf1c PE=2 SV=1 A2AT02_MOUSE [4]	37 kDa	1	0	0.9	1	0.9	1.111111111

Ras suppressor protein 1 OS=Mus musculus GN=Rsu1 PE=4 SV=1	A2AUR7_MOUSE (+2)	30 kDa	1	20.05867	0.96666667	0.96666667	1	1
ATPase family AAA domain-containing protein 1 OS=Mus musculus GN=ATAD1_MOUSE		41 kDa	1	25.589737	1.36666667	1.3	1.051282051	0.951219512
Clustered mitochondria protein homolog OS=Mus musculus GN=Cluh PE CLU_MOUSE		148 kDa	1	20.576704	1.1	1.16666667	0.942857143	1.060606061
Malonyl-CoA decarboxylase, mitochondrial OS=Mus musculus GN=MlyccDCMC_MOUSE		55 kDa	1	20.646286	1.06666667	1.1	0.96969697	1.03125
Protein Rpl31 OS=Mus musculus GN=Rpl31 PE=2 SV=1	E9PWZ3_MOUSE	47 kDa	1	16.227219	1.033333333	1	1.033333333	0.967741935
Cluster of Calcium/calmodulin-dependent protein kinase type II subunit E9Q1T1_MOUSE [2]		60 kDa	1	21.927457	0.9	1	0.9	1.111111111
Cardiomyopathy-associated protein 5 OS=Mus musculus GN=Cmya5 PE=E9QLJ0_MOUSE		406 kDa	1	12.605646	0.933333333	0.9	1.037037037	0.964285714
Ezrin OS=Mus musculus GN=Ezr PE=1 SV=3	EZRI_MOUSE	69 kDa	1	21.269904	1.1	1.16666667	0.942857143	1.060606061
Protein FAM162A OS=Mus musculus GN=Fam162a PE=2 SV=1	F162A_MOUSE	18 kDa	1	27.96448	1.033333333	1.16666667	0.885714286	1.129032258
Cluster of Hydroxyacyl glutathione hydrolase OS=Mus musculus GN=Hag G5E8T9_MOUSE [2]		34 kDa	1	21.000451	0.833333333	0.8	1.041666667	0.96
Heterogeneous nuclear ribonucleoprotein L (Fragment) OS=Mus musculus GN=G5E924_MOUSE		67 kDa	1	19.700597	0.86666667	0.96666667	0.896551724	1.115384615
Bifunctional epoxide hydrolase 2 OS=Mus musculus GN=Ephx2 PE=1 SV= HYES_MOUSE		63 kDa	1	20.851794	1.033333333	1.06666667	0.96875	1.032258065
Kelch-like protein 41 OS=Mus musculus GN=Klh41 PE=1 SV=1	KLH41_MOUSE	68 kDa	1	23.114243	1	1.16666667	0.857142857	1.166666667
Mitochondrial peptide methionine sulfoxide reductase OS=Mus musculus GN=MSRA_MOUSE		26 kDa	1	20.943744	1.06666667	1	1.066666667	0.9375
Metaxin-2 OS=Mus musculus GN=Mtx2 PE=1 SV=1	MTX2_MOUSE	30 kDa	1	22.055636	0.96666667	1.03333333	0.935483871	1.068965517
Cluster of 2-oxoisovalerate dehydrogenase subunit alpha, mitochondrial ODBA_MOUSE [2]		50 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Protein disulfide-isomerase A6 OS=Mus musculus GN=Pdia6 PE=1 SV=3 PDIA6_MOUSE (+1)		48 kDa	1	22.239387	0.933333333	1.13333333	0.823529412	1.214285714
[Pyruvate dehydrogenase (lipoamide)] kinase isozyme 2, mitochondrial CDPK2_MOUSE		46 kDa	1	21.079832	1	1	1	1
Heterogeneous nuclear ribonucleoprotein A1 OS=Mus musculus GN=Hnr Q5EBP8_MOUSE (+1)		39 kDa	1	23.999734	0.8	0.9	0.888888889	1.125
40S ribosomal protein S8 OS=Mus musculus GN=Rps8 PE=1 SV=2	RS8_MOUSE	24 kDa	1	20.622392	0.96666667	1.06666667	0.90625	1.103448276
Septin-7 OS=Mus musculus GN=Sept7 PE=1 SV=1	SEPT7_MOUSE	51 kDa	1	18.973924	0.8	0.9	0.888888889	1.125
Sulfide:quinone oxidoreductase, mitochondrial OS=Mus musculus GN=SrSQRD_MOUSE		50 kDa	1	17.730577	1.15	1.05	1.095238095	0.913043478
Phenylalanine--tRNA ligase beta subunit OS=Mus musculus GN=Farsb PE SYFB_MOUSE		66 kDa	1	7.797211	0.9	1	0.9	1.111111111
Cluster of T-complex protein 1 subunit zeta OS=Mus musculus GN=Cct6aTCPZ_MOUSE		58 kDa	1	22.672342	0.96666667	1	0.966666667	1.034482759
Dedicator of cytokinesis protein 7 OS=Mus musculus GN=Dock7 PE=2 SV A2A9M5_MOUSE (+2)		241 kDa	1	0	1.4	1.7	0.823529412	1.214285714
Apolipoprotein E OS=Mus musculus GN=Apoe PE=1 SV=2	APOE_MOUSE	36 kDa	1	20.109908	0.8	0.8	1	1
BolaA-like protein 3 OS=Mus musculus GN=Bola3 PE=2 SV=1	BOLA3_MOUSE	12 kDa	1	20.363095	0.933333333	0.93333333	1	1
Dihydropteridine reductase OS=Mus musculus GN=Qdpr PE=1 SV=2	DHPR_MOUSE	26 kDa	1	21.434425	0.96666667	1.06666667	0.90625	1.103448276
Cluster of Proteasome subunit alpha type-7 OS=Mus musculus GN=PsmA7_MOUSE		28 kDa	1	20.079666	1	1.03333333	0.967741935	1.033333333
Ryanodine receptor 1 OS=Mus musculus GN=Ryr1 PE=1 SV=1	RYR1_MOUSE	565 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Cluster of Isoform 2 of Clathrin light chain B OS=Mus musculus GN=Cltb [sp] Q6IRU5-2] CLCB_MOUSE [2]		?	1	21.924058	0.95	0.85	1.117647059	0.894736842
Cluster of Aldose reductase-related protein 1 OS=Mus musculus GN=Akr ALD1_MOUSE [3]		36 kDa	1	15.72403	1.1	1.05	1.047619048	0.954545455
Apolipoprotein O OS=Mus musculus GN=Apoo PE=2 SV=1	APOO_MOUSE	24 kDa	1	21.189535	0.96666667	1.06666667	0.90625	1.103448276
Cluster of 60S ribosomal protein L12 OS=Mus musculus GN=Rpl12 PE=1 !RL12_MOUSE		18 kDa	1	22.49839	0.9	1	0.9	1.111111111
Cluster of 60S ribosomal protein L32 OS=Mus musculus GN=Rpl32 PE=2 !RL32_MOUSE		16 kDa	1	20.596597	1.033333333	1.2	0.861111111	1.161290323
Cluster of Transforming acidic coiled-coil-containing protein 2 OS=Mus musculus GN=E9Q8T1_MOUSE		305 kDa	1	13.783537	0.85	0.8	1.0625	0.941176471
6-phosphogluconolactonase OS=Mus musculus GN=PglS PE=2 SV=1	6PGL_MOUSE	27 kDa	1	22.288658	1	1.06666667	0.9375	1.066666667
Alpha-actinin-4 OS=Mus musculus GN=Actn4 PE=1 SV=1	ACTN4_MOUSE	105 kDa	1	20.620131	1	1	1	1
Cytosol aminopeptidase OS=Mus musculus GN=Lap3 PE=1 SV=3	AMPL_MOUSE (+1)	56 kDa	1	21.055734	0.833333333	0.83333333	1	1
Beta-2-glycoprotein 1 OS=Mus musculus GN=Apoh PE=1 SV=1	APOH_MOUSE	39 kDa	1	24.887437	1	0.96666667	1.034482759	0.966666667
Thioredoxin reductase 2, mitochondrial OS=Mus musculus GN=Txnrd2 PID3Z0K8_MOUSE (+3)		53 kDa	1	19.09658	1	0.86666667	1.153846154	0.866666667
Cluster of Guanin nucleotide-binding protein G(I)/G(S)/G(T) subunit bet GBB1_MOUSE [3]		37 kDa	1	19.08933	0.9	0.93333333	0.964285714	1.037037037
28 kDa heat- and acid-stable phosphoprotein OS=Mus musculus GN=PdaHAP28_MOUSE		21 kDa	1	21.498288	0.933333333	0.96666667	0.965517241	1.035714286
Integrin-linked protein kinase OS=Mus musculus GN=Ilk PE=1 SV=2	ILK_MOUSE	51 kDa	1	18.765989	0.933333333	1.06666667	0.875	1.142857143
Lamin-B2 OS=Mus musculus GN=Lmnb2 PE=1 SV=2	LMNB2_MOUSE	67 kDa	1	20.714954	0.96666667	0.96666667	1	1
Leiomodlin-2 OS=Mus musculus GN=Lmod2 PE=1 SV=1	LMOD2_MOUSE	62 kDa	1	20.798174	1	0.93333333	1.071428571	0.933333333
Nidogen-2 OS=Mus musculus GN=Nid2 PE=1 SV=2	NID2_MOUSE	154 kDa	1	19.596416	0.9	0.83333333	1.08	0.925925926
2-oxoisovalerate dehydrogenase subunit beta, mitochondrial OS=Mus musculus GN=ODBB_MOUSE		43 kDa	1	28.30552	1.1	1.3	0.846153846	1.181818182
Ubiquitin carboxyl-terminal hydrolase OS=Mus musculus GN=Usp5 PE=2 Q3U4W8_MOUSE (+1)		93 kDa	1	19.243039	1.033333333	1.03333333	1	1
60S ribosomal protein L14 OS=Mus musculus GN=Rpl14 PE=2 SV=3	RL14_MOUSE	24 kDa	1	20.452224	0.96666667	1.06666667	0.90625	1.103448276
Saccharopine dehydrogenase-like oxidoreductase OS=Mus musculus GN=SCPD_MOUSE		47 kDa	1	20.398413	0.933333333	1	0.933333333	1.071428571
Cluster of RNA-binding motif protein, X chromosome OS=Mus musculus GN=RBMX_MOUSE [2]		42 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
E3 ubiquitin-protein ligase HUWE1 OS=Mus musculus GN=Huwe1 PE=2 SA2AFQ0_MOUSE (+1)		483 kDa	1	8.777281	1.25	1.1	1.136363636	0.88

Epiplakin OS=Mus musculus GN=Eppk1 PE=1 SV=2	EPIPL_MOUSE	725 kDa	1	7.839395	1.1	1	1.1	0.909090909
Cluster of Protein Gm20459 (Fragment) OS=Mus musculus GN=Gm20459 F7AA26_MOUSE		126 kDa	1	23.547421	0.9	0.86666667	1.038461538	0.962962963
Fermitin family homolog 2 OS=Mus musculus GN=Fermt2 PE=1 SV=1	FERM2_MOUSE	78 kDa	1	19.194897	1	1	1	1
Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Mus musculus GN=Fkbp3 FFKBP3_MOUSE		25 kDa	1	22.717509	0.96666667	1.03333333	0.935483871	1.068965517
Heat shock protein beta-8 OS=Mus musculus GN=Hspb8 PE=1 SV=1	HSPB8_MOUSE	22 kDa	1	25.103754	0.7	0.63333333	1.105263158	0.904761905
Methionine-R-sulfoxide reductase B2, mitochondrial OS=Mus musculus (MSRB2_MOUSE		19 kDa	1	22.051401	1	1.16666667	0.857142857	1.16666667
Methylthioribulose-1-phosphate dehydratase OS=Mus musculus GN=ApiMTNB_MOUSE		27 kDa	1	24.600796	0.9	0.9	1	1
Neutral cholesterol ester hydrolase 1 OS=Mus musculus GN=Nceh1 PE=1 NCEH1_MOUSE		46 kDa	1	24.709389	0.9	1.1	0.818181818	1.222222222
Cytochrome b-c1 complex subunit 9 OS=Mus musculus GN=Uqcr10 PE=1 QCR9_MOUSE		7 kDa	1	33.775008	0.93333333	0.9	1.037037037	0.964285714
60S ribosomal protein L28 OS=Mus musculus GN=Rpl28 PE=1 SV=2	RL28_MOUSE	16 kDa	1	20.580837	0.93333333	1.13333333	0.823529412	1.214285714
60S acidic ribosomal protein P2 OS=Mus musculus GN=Rplp2 PE=1 SV=3	RLA2_MOUSE	12 kDa	1	23.697786	0.93333333	0.93333333	1	1
Sepiapterin reductase OS=Mus musculus GN=Spr PE=1 SV=1	SPRE_MOUSE	28 kDa	1	22.759809	0.93333333	0.93333333	1	1
Mitochondrial import inner membrane translocase subunit Tim9 OS=Mu TIM9_MOUSE		10 kDa	1	22.122115	0.9	0.8	1.125	0.888888889
DNA topoisomerase 2-binding protein 1 OS=Mus musculus GN=Topbp1 FTOPB1_MOUSE		169 kDa	1	7.389074	1.3	1.2	1.083333333	0.923076923
Hsp90 co-chaperone Cdc37 OS=Mus musculus GN=Cdc37 PE=2 SV=1	CDC37_MOUSE	45 kDa	1	22.648026	0.96666667	1	0.96666667	1.034482759
Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1	CO6A1_MOUSE	108 kDa	1	20.672217	1.6	1.16666667	1.371428571	0.729166667
Catenin alpha-3 OS=Mus musculus GN=Ctnna3 PE=1 SV=2	CTNA3_MOUSE	100 kDa	1	20.69261	1.1	1.13333333	0.970588235	1.03030303
Dnaj homolog subfamily A member 4 OS=Mus musculus GN=Dnaja4 PE= DNJA4_MOUSE		45 kDa	1	19.615111	0.83333333	0.86666667	0.961538462	1.04
Ubiquitin carboxyl-terminal hydrolase OS=Mus musculus GN=Usp14 PE= E9PYI8_MOUSE (+1)		52 kDa	1	15.679699	0.83333333	0.93333333	0.892857143	1.12
Protein phosphatase 1 regulatory subunit 7 OS=Mus musculus GN=Ppp1 PP1R7_MOUSE		41 kDa	1	22.430208	0.93333333	1	0.93333333	1.071428571
Peptidyl-prolyl cis-trans isomerase D OS=Mus musculus GN=Ppid PE=1 SIPPID_MOUSE		41 kDa	1	20.181602	0.8	0.93333333	0.857142857	1.16666667
Protein phosphatase 1 regulatory subunit 3A OS=Mus musculus GN=Ppp PPR3A_MOUSE		121 kDa	1	22.580025	1.16666667	1.13333333	1.029411765	0.971428571
Proteasome subunit alpha type-4 OS=Mus musculus GN=Psm4 PE=1 SV PSA4_MOUSE		29 kDa	1	19.142711	0.96666667	1.03333333	0.935483871	1.068965517
MCG49183 OS=Mus musculus GN=1700009N14Rik PE=2 SV=1	Q14AA6_MOUSE	24 kDa	1	19.93299	0.96666667	1.06666667	0.90625	1.103448276
39S ribosomal protein L12, mitochondrial OS=Mus musculus GN=Mrpl12 RM12_MOUSE		22 kDa	1	22.030129	0.96666667	1	0.96666667	1.034482759
40S ribosomal protein S10 OS=Mus musculus GN=Rps10 PE=1 SV=1	RS10_MOUSE	19 kDa	1	19.704197	0.86666667	0.93333333	0.928571429	1.076923077
Mitochondrial import inner membrane translocase subunit Tim10 OS=M TIM10_MOUSE		10 kDa	1	21.041359	0.93333333	0.83333333	1.12	0.892857143
Cluster of Alcohol dehydrogenase class-3 OS=Mus musculus GN=Adh5 P1ADHX_MOUSE		40 kDa	1	18.639775	1	1.06666667	0.9375	1.06666667
Cluster of Uncharacterized protein C6orf203 homolog OS=Mus musculus CF203_MOUSE		28 kDa	1	13.368917	1	1	1	1
Cluster of CLIP-associating protein 1 OS=Mus musculus GN=Clasp1 PE=1 CLAP1_MOUSE [4]		169 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
A-kinase anchor protein 12 OS=Mus musculus GN=Akap12 PE=1 SV=1	AKA12_MOUSE	181 kDa	1	13.402957	1.1	0.93333333	1.178571429	0.848484848
ATP synthase subunit g, mitochondrial OS=Mus musculus GN=Atp5l PE=1 ATP5L_MOUSE		11 kDa	1	25.810304	1.03333333	0.96666667	1.068965517	0.935483871
Hexaprenyldihydroxybenzoate methyltransferase, mitochondrial OS=Mu COQ3_MOUSE		41 kDa	1	20.169699	1.06666667	0.96666667	1.103448276	0.90625
Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Mus musculus GBLP_MOUSE		35 kDa	1	13.563631	0.96666667	1	0.96666667	1.034482759
Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Mus musculus GN=HNRPC_MOUSE		34 kDa	1	24.259953	0.93333333	0.93333333	1	1
Heat shock protein beta-7 OS=Mus musculus GN=Hspb7 PE=1 SV=3	HSPB7_MOUSE	19 kDa	1	21.637957	0.93333333	0.83333333	1.12	0.892857143
Cluster of LIM and SH3 domain protein 1 OS=Mus musculus GN=Lasp1 PILASP1_MOUSE		30 kDa	1	19.586591	0.86666667	0.9	0.962962963	1.038461538
Methyl-CpG-binding protein 2 OS=Mus musculus GN=Mecp2 PE=1 SV=1	MECP2_MOUSE	52 kDa	1	20.43711	1.16666667	1.03333333	1.129032258	0.885714286
Mitochondrial-processing peptidase subunit alpha OS=Mus musculus GNMPPA_MOUSE		58 kDa	1	20.922571	0.96666667	1.06666667	0.90625	1.103448276
Na(+)/H(+) exchange regulatory cofactor NHE-RF2 OS=Mus musculus GN=NHRF2_MOUSE		37 kDa	1	25.001833	0.93333333	0.83333333	1.12	0.892857143
Perilipin-3 OS=Mus musculus GN=Plin3 PE=1 SV=1	PLIN3_MOUSE	47 kDa	1	13.871408	0.9	0.96666667	0.931034483	1.074074074
Polyribonucleotide nucleotidyltransferase 1, mitochondrial OS=Mus musculus PNPT1_MOUSE		86 kDa	1	20.113301	1.03333333	1.03333333	1	1
26S protease regulatory subunit 6B OS=Mus musculus GN=Psmc4 PE=1 SPRS6B_MOUSE		47 kDa	1	19.655745	0.9	0.96666667	0.931034483	1.074074074
60S ribosomal protein L23 OS=Mus musculus GN=Rpl23 PE=1 SV=1	RL23_MOUSE	15 kDa	1	21.473497	0.86666667	0.96666667	0.896551724	1.115384615
60S ribosomal protein L35a OS=Mus musculus GN=Rpl35a PE=2 SV=2	RL35A_MOUSE	13 kDa	1	21.747069	0.96666667	1.06666667	0.90625	1.103448276
Cluster of Septin-11 OS=Mus musculus GN=Sept11 PE=1 SV=4 (SEP11_M SEPT11_MOUSE [6])		50 kDa	1	108.40056	0.9	1.25	0.72	1.388888889
Sulfite oxidase, mitochondrial OS=Mus musculus GN=Suox PE=1 SV=2	SUOX_MOUSE	61 kDa	1	20.132692	1.26666667	1.16666667	1.085714286	0.921052632
Lysine--tRNA ligase OS=Mus musculus GN=Kars PE=1 SV=1	SYK_MOUSE	68 kDa	1	23.019235	1.03333333	1.03333333	1	1
Telethonin OS=Mus musculus GN=Tcap PE=2 SV=1	TELT_MOUSE	19 kDa	1	22.621821	0.86666667	0.9	0.962962963	1.038461538
Cluster of Heterogeneous nuclear ribonucleoprotein A3 OS=Mus musculus A2AL12_MOUSE		34 kDa	1	23.40577	0.93333333	0.96666667	0.965517241	1.035714286
Cluster of Isobutyryl-CoA dehydrogenase, mitochondrial OS=Mus musculus D3YTT4_MOUSE [2]		45 kDa	1	0	0.9	1.1	0.818181818	1.222222222
Cysteine-rich protein 1 OS=Mus musculus GN=Crip1 PE=2 SV=2	CRIP1_MOUSE	9 kDa	1	12.790304	0.9	0.9	1	1
Cytoplasmic dynein 1 light intermediate chain 1 OS=Mus musculus GN=LDC1L1_MOUSE		57 kDa	1	20.611545	0.9	0.9	1	1

Protein-L-isoaspartate O-methyltransferase OS=Mus musculus GN=Pcmt E0CYV0_MOUSE	30 kDa	1	20.631564	0.96666667	0.96666667	1	1
Glutaredoxin-3 OS=Mus musculus GN=Glrx3 PE=1 SV=1	38 kDa	1	23.934983	0.93333333	1.06666667	0.875	1.142857143
Histidine triad nucleotide-binding protein 1 OS=Mus musculus GN=Hint1 HINT1_MOUSE	14 kDa	1	20.99829	0.9	0.9	1	1
Monocarboxylate transporter 1 OS=Mus musculus GN=Slc16a1 PE=1 SV=1	53 kDa	1	17.699026	1.16666667	1.2	0.972222222	1.028571429
Myopalladin OS=Mus musculus GN=Mypn PE=2 SV=2	144 kDa	1	13.982381	0.95	0.9	1.055555556	0.947368421
Programmed cell death protein 5 OS=Mus musculus GN=Pcdc5 PE=1 SV=1	14 kDa	1	22.232324	0.8	0.86666667	0.923076923	1.083333333
Bisphosphoglycerate mutase OS=Mus musculus GN=Bpgm PE=2 SV=2	30 kDa	1	20.265861	1.1	1.06666667	1.03125	0.96969697
Rab11 family-interacting protein 5 OS=Mus musculus GN=Rab11fip5 PE=1 SV=1	70 kDa	1	8.03983	0.9	0.8	1.125	0.888888889
Cluster of 60S ribosomal protein L18a OS=Mus musculus GN=Rpl18a PE=1 SV=1	21 kDa	1	19.572401	0.93333333	1.1	0.848484848	1.178571429
40S ribosomal protein S13 OS=Mus musculus GN=Rps13 PE=1 SV=2	17 kDa	1	20.580417	0.93333333	1.2	0.777777778	1.285714286
40S ribosomal protein S14 OS=Mus musculus GN=Rps14 PE=2 SV=3	16 kDa	1	20.842785	0.9	1.13333333	0.794117647	1.259259259
40S ribosomal protein S19 OS=Mus musculus GN=Rps19 PE=1 SV=3	16 kDa	1	21.060506	0.83333333	1	0.833333333	1.2
Glycine--tRNA ligase OS=Mus musculus GN=Gars PE=1 SV=1	82 kDa	1	22.06463	0.73333333	0.73333333	1	1
Translationally-controlled tumor protein OS=Mus musculus GN=Tpt1 PE=1 SV=1	19 kDa	1	21.517941	0.83333333	0.86666667	0.961538462	1.04
40S ribosomal protein S24 OS=Mus musculus GN=Rps24 PE=1 SV=1	15 kDa	1	21.148951	1	1.03333333	0.967741935	1.033333333
Cluster of AP-2 complex subunit beta OS=Mus musculus GN=Ap2b1 PE=1 SV=1	105 kDa	1	6.0669415	0.95	1.05	0.904761905	1.105263158
Cluster of MCG19772 OS=Mus musculus GN=Trim55 PE=4 SV=1	60 kDa	1	32.559413	0.9	0.9	1	1
Cluster of Protein 4.1 OS=Mus musculus GN=Epb41 PE=1 SV=2	96 kDa	1	13.122292	0.95	0.9	1.055555556	0.947368421
Cluster of S-phase kinase-associated protein 1 OS=Mus musculus GN=Skp1 SKP1_MOUSE [2]	19 kDa	1	34.322676	0.9	0.85	1.058823529	0.944444444
Alpha-1-syntrophin OS=Mus musculus GN=Snta1 PE=2 SV=1	53 kDa	1	18.858515	0.96666667	1	0.966666667	1.034482759
Apolipoprotein O-like OS=Mus musculus GN=Apool PE=2 SV=1	29 kDa	1	18.517558	1	1.06666667	0.9375	1.066666667
Bcl-2-like protein 13 OS=Mus musculus GN=Bcl2l13 PE=1 SV=2	47 kDa	1	23.093916	0.96666667	0.96666667	1	1
Adenylyl cyclase-associated protein 2 OS=Mus musculus GN=Cap2 PE=1 SV=1	53 kDa	1	19.970213	0.93333333	1	0.933333333	1.071428571
Cathepsin B OS=Mus musculus GN=Ctsb PE=1 SV=2	37 kDa	1	23.252524	0.8	0.76666667	1.043478261	0.958333333
Collagen alpha-1(I) chain OS=Mus musculus GN=Col1a1 PE=1 SV=4	138 kDa	1	19.584481	1.73333333	1	1.733333333	0.576923077
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial OS=1 COQ5_MOUSE	37 kDa	1	22.51319	1.06666667	1.06666667	1	1
CapZ-interacting protein OS=Mus musculus GN=Rcsd1 PE=2 SV=1	44 kDa	1	14.128383	1.2	1.15	1.043478261	0.958333333
Dynactin subunit 1 OS=Mus musculus GN=Dctn1 PE=1 SV=3	142 kDa	1	19.852781	0.96666667	0.93333333	1.035714286	0.965517241
Probable rRNA-processing protein EBP2 OS=Mus musculus GN=Ebna1bp EBP2_MOUSE	35 kDa	1	9.431601	0.8	1	0.8	1.25
Glyoxylate reductase/hydroxypyruvate reductase OS=Mus musculus GN=GRHPR_MOUSE	35 kDa	1	20.150286	0.83333333	0.86666667	0.961538462	1.04
Kynurenine--oxoglutarate transaminase 3 OS=Mus musculus GN=Ccbl2 PKAT3_MOUSE	51 kDa	1	22.943507	1	0.96666667	1.034482759	0.966666667
NADP-dependent malic enzyme OS=Mus musculus GN=Me1 PE=1 SV=2	64 kDa	1	20.700599	0.96666667	1.03333333	0.935483871	1.068965517
Endonuclease G, mitochondrial OS=Mus musculus GN=Endog PE=2 SV=1	32 kDa	1	25.602296	1	0.93333333	1.071428571	0.933333333
Cluster of Oxidoreductase NAD-binding domain-containing protein 1 OS=OXND1_MOUSE	35 kDa	1	19.868736	0.86666667	0.93333333	0.928571429	1.076923077
[Pyruvate dehydrogenase [lipoamide]] kinase isozyme 1, mitochondrial OS=PKAT1_MOUSE	49 kDa	1	22.547674	1.06666667	1.03333333	1.032258065	0.96875
Peptidyl-prolyl cis-trans isomerase B OS=Mus musculus GN=Ppib PE=2 SV=1	24 kDa	1	20.058267	0.96666667	1.06666667	0.90625	1.103448276
Presequence protease, mitochondrial OS=Mus musculus GN=Pitrm1 PE=1 SV=1	117 kDa	1	20.873213	1	0.96666667	1.034482759	0.966666667
26S proteasome non-ATPase regulatory subunit 3 OS=Mus musculus GN=PSMD3_MOUSE	61 kDa	1	22.767472	1.1	1.13333333	0.970588235	1.03030303
Protein Prob1 (Fragment) OS=Mus musculus GN=Prob1 PE=2 SV=1	106 kDa	1	24.151386	1.03333333	0.86666667	1.192307692	0.838709677
60S ribosomal protein L18 OS=Mus musculus GN=Rpl18 PE=2 SV=3	22 kDa	1	23.327418	0.93333333	1.06666667	0.875	1.142857143
Heterogeneous nuclear ribonucleoprotein A/B OS=Mus musculus GN=Hr ROAA_MOUSE	31 kDa	1	13.051207	0.93333333	0.93333333	1	1
28S ribosomal protein S36, mitochondrial OS=Mus musculus GN=Mrrps3 RT36_MOUSE	11 kDa	1	21.241741	0.96666667	0.86666667	1.115384615	0.896551724
Transgelin-2 OS=Mus musculus GN=Tagln2 PE=1 SV=4	22 kDa	1	22.378351	0.86666667	1	0.866666667	1.153846154
Vesicle-associated membrane protein-associated protein A OS=Mus mus VAPA_MOUSE	28 kDa	1	20.856927	1.03333333	1	1.033333333	0.967741935
Vacuolar protein sorting-associated protein 35 OS=Mus musculus GN=Vps35_MOUSE	92 kDa	1	23.056203	0.9	1.03333333	0.870967742	1.148148148
Zyxin OS=Mus musculus GN=Zyx PE=1 SV=2	61 kDa	1	19.204547	0.83333333	0.76666667	1.086956522	0.92
Serine/threonine-protein kinase mTOR OS=Mus musculus GN=Mtor PE=1 SV=1	289 kDa	1	8.845222	1.1	1.2	0.916666667	1.090909091
Reticulon-4-interacting protein 1, mitochondrial OS=Mus musculus GN=FRT4I1_MOUSE	43 kDa	1	19.890244	0.83333333	0.9	0.925925926	1.08
Fatty acid-binding protein, epidermal OS=Mus musculus GN=Fabp5 PE=1 SV=1	15 kDa	1	20.860764	0.96666667	0.96666667	1	1
Cluster of ADP-ribosylation factor 3 OS=Mus musculus GN=Arf3 PE=2 SV=1	21 kDa	1	22.28918	0.96666667	1.13333333	0.852941176	1.172413793
Cluster of Protein Zim1 OS=Mus musculus GN=Zim1 PE=2 SV=1	65 kDa	1	9.078467	1.6	1.4	1.142857143	0.875
Alcohol dehydrogenase [NADP(+)] OS=Mus musculus GN=Akr1a1 PE=1 SV=1	37 kDa	1	13.325784	0.8	0.9	0.888888889	1.125
Annexin A7 OS=Mus musculus GN=Anxa7 PE=2 SV=2	50 kDa	1	19.596723	0.93333333	0.9	1.037037037	0.964285714

Aflatoxin B1 aldehyde reductase member 2 OS=Mus musculus GN=Akr7zARK72_MOUSE	41 kDa	1	20.213675	1.033333333	1.066666667	0.96875	1.032258065
Cellular nucleic acid-binding protein OS=Mus musculus GN=Cnbp PE=2 S'CNBP_MOUSE	20 kDa	1	12.88657	0.833333333	0.8	1.041666667	0.96
Cluster of Anamorsin OS=Mus musculus GN=Ciapi1 PE=1 SV=1 (CPIN1_ICPIN1_MOUSE [2])	33 kDa	1	0	1.3	1	1.3	0.769230769
Peroxisomal multifunctional enzyme type 2 OS=Mus musculus GN=Hsd1 DHB4_MOUSE	79 kDa	1	20.132968	0.9	0.866666667	1.038461538	0.962962963
Eukaryotic translation initiation factor 3 subunit J-A OS=Mus musculus G EI3JA_MOUSE (+1)	29 kDa	1	12.711407	0.966666667	0.966666667	1	1
Protein FAM136A OS=Mus musculus GN=Fam136a PE=1 SV=1	16 kDa	1	19.216055	0.9	0.8	1.125	0.888888889
Protocadherin Fat 4 OS=Mus musculus GN=Fat4 PE=1 SV=2	540 kDa	1	8.07646	0.9	0.9	1	1
Acyl-protein thioesterase 1 OS=Mus musculus GN=Lyp1a1 PE=4 SV=1	24 kDa	1	19.768785	1.033333333	1	1.033333333	0.967741935
Adenylate kinase isoenzyme 4, mitochondrial OS=Mus musculus GN=Ak4KAD4_MOUSE	25 kDa	1	19.061822	1	1.066666667	0.9375	1.066666667
Leukotriene A-4 hydrolase OS=Mus musculus GN=Lta4h PE=1 SV=4	69 kDa	1	20.320016	0.933333333	1.1	0.848484848	1.178571429
Protein Sptbn2 OS=Mus musculus GN=Sptbn2 PE=2 SV=1	271 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Ankycorbin OS=Mus musculus GN=Rai14 PE=1 SV=1	109 kDa	1	9.875215	1.6	1.3	1.230769231	0.8125
60S ribosomal protein L37 OS=Mus musculus GN=Rpl37 PE=2 SV=3	11 kDa	1	21.650548	0.933333333	1.066666667	0.875	1.142857143
28S ribosomal protein S22, mitochondrial OS=Mus musculus GN=Mrps2;RT22_MOUSE	41 kDa	1	22.052878	0.966666667	0.933333333	1.035714286	0.965517241
Gamma-synuclein OS=Mus musculus GN=Sncg PE=1 SV=1	13 kDa	1	20.362706	0.966666667	0.866666667	1.115384615	0.896551724
Cluster of Hydroxymethylglutaryl-CoA lyase, mitochondrial OS=Mus mus HMGCL_MOUSE	34 kDa	1	18.725693	0.833333333	1	0.833333333	1.2
Myosin Vb, isoform CRA_a OS=Mus musculus GN=Myo5b PE=4 SV=1	213 kDa	1	0	1.1	1.4	0.785714286	1.272727273
Cluster of Tripeptidyl-peptidase 2 OS=Mus musculus GN=Tpp2 PE=2 SV=1	140 kDa	1	22.795229	1.033333333	1	1.033333333	0.967741935
Cluster of Heat shock 70 kDa protein 12B OS=Mus musculus GN=Hspa12 HS12B_MOUSE	76 kDa	1	27.68449	1.033333333	1.133333333	0.911764706	1.096774194
Cluster of Cordon-bleu protein-like 1 OS=Mus musculus GN=Cobl1 PE=4 B1AZ14_MOUSE [2]	133 kDa	1	6.2348075	1	1	1	1
Proteasomal ubiquitin receptor ADRM1 OS=Mus musculus GN=Adrm1 PIADRM1_MOUSE	42 kDa	1	14.206791	0.933333333	0.933333333	1	1
Afamin OS=Mus musculus GN=Afm PE=1 SV=2	69 kDa	1	14.964652	0.95	0.9	1.055555556	0.947368421
Isoaspartyl peptidase/L-asparaginase OS=Mus musculus GN=Asrgl1 PE=1 ASGL1_MOUSE	34 kDa	1	21.772133	0.966666667	1	0.966666667	1.034482759
C-1-tetrahydrofolate synthase, cytoplasmic OS=Mus musculus GN=Mthfc1CTC_MOUSE	101 kDa	1	19.917217	0.9	0.933333333	0.964285714	1.037037037
UPF0598 protein C8orf82 homolog OS=Mus musculus PE=2 SV=1	24 kDa	1	25.05142	1.1	1.1	1	1
Putative ATP-dependent Clp protease proteolytic subunit, mitochondrial CLPP_MOUSE	30 kDa	1	23.712501	1	0.9	1.111111111	0.9
Dynactin subunit 2 OS=Mus musculus GN=Dctn2 PE=1 SV=3	44 kDa	1	19.540466	0.9	0.933333333	0.964285714	1.037037037
DnaJ homolog subfamily A member 3, mitochondrial OS=Mus musculus (DNJA3_MOUSE	52 kDa	1	13.613428	0.966666667	0.9	1.074074074	0.931034483
Dystrobrevin alpha OS=Mus musculus GN=Dtna PE=1 SV=2	84 kDa	1	20.592918	1.1	1.133333333	0.970588235	1.03030303
Glycerol-3-phosphate dehydrogenase [NAD(+)], cytoplasmic OS=Mus muGPDA_MOUSE	38 kDa	1	20.214844	1	0.966666667	1.034482759	0.966666667
Glutathione reductase, mitochondrial OS=Mus musculus GN=Gsr PE=2 S'GSHR_MOUSE (+1)	54 kDa	1	22.370468	0.9	1	0.9	1.111111111
High mobility group protein HMG-I/HMG-Y OS=Mus musculus GN=HmgaHMG1_MOUSE	12 kDa	1	19.789056	1.066666667	1.233333333	0.864864865	1.15625
Protein Dnahc7b OS=Mus musculus GN=Dnahc7b PE=4 SV=1	467 kDa	1	6.6763125	1.1	1.1	1	1
NAD(P)H dehydrogenase [quinone] 1 OS=Mus musculus GN=Nqo1 PE=1	31 kDa	1	20.030507	1.066666667	1	1.066666667	0.9375
Prolyl endopeptidase OS=Mus musculus GN=Prep PE=2 SV=1	81 kDa	1	19.253922	0.933333333	1	0.933333333	1.071428571
Proline synthase co-transcribed bacterial homolog protein OS=Mus mus PROSC_MOUSE	30 kDa	1	19.280142	0.966666667	0.966666667	1	1
Vesicle-associated membrane protein, associated protein B and C OS=M Q8BH80_MOUSE (+1)	27 kDa	1	19.892752	1	0.966666667	1.034482759	0.966666667
Cortactin, isoform CRA_a OS=Mus musculus GN=Cttn PE=2 SV=1	57 kDa	1	19.282375	0.9	0.866666667	1.038461538	0.962962963
60S ribosomal protein L21 OS=Mus musculus GN=Rpl21 PE=2 SV=1	19 kDa	1	19.619861	0.9	1.066666667	0.84375	1.185185185
60S ribosomal protein L26 OS=Mus musculus GN=Rpl26 PE=2 SV=1	17 kDa	1	21.264353	0.833333333	0.933333333	0.892857143	1.12
60S ribosomal protein L27a OS=Mus musculus GN=Rpl27a PE=2 SV=5	17 kDa	1	14.853887	0.9	1.233333333	0.72972973	1.37037037
40S ribosomal protein S16 OS=Mus musculus GN=Rps16 PE=2 SV=4	16 kDa	1	20.263481	0.9	1.066666667	0.84375	1.185185185
Splicing factor, proline- and glutamine-rich OS=Mus musculus GN=Sfpq PSFPQ_MOUSE	75 kDa	1	12.100717	1.033333333	0.966666667	1.068965517	0.935483871
Serine/arginine-rich splicing factor 7 OS=Mus musculus GN=Srsf7 PE=1 S'SRSF7_MOUSE (+2)	31 kDa	1	21.474414	1.033333333	0.866666667	1.192307692	0.838709677
Ubiquitin-conjugating enzyme E2 L3 OS=Mus musculus GN=Ube2l3 PE=2 UB2L3_MOUSE	18 kDa	1	19.088286	0.966666667	0.933333333	1.035714286	0.965517241
Cluster of 60S ribosomal protein L17 OS=Mus musculus GN=Rpl17 PE=2 Q6ZWZ7_MOUSE	21 kDa	1	20.45026	0.866666667	1.166666667	0.742857143	1.346153846
Cluster of Collagen alpha-1(XV) chain OS=Mus musculus GN=Col15a1 PE=A2AJY2_MOUSE [3]	138 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Adiponectin OS=Mus musculus GN=Adipoq PE=1 SV=2	27 kDa	1	14.773202	1.133333333	1.133333333	1	1
Cluster of Plastin-3 OS=Mus musculus GN=Pls3 PE=2 SV=1 (B1AX58_MOIB1AX58_MOUSE [3])	72 kDa	1	33.637613	0.95	0.95	1	1
Valacyclovir hydrolase OS=Mus musculus GN=Bphl PE=2 SV=1	33 kDa	1	22.342654	0.933333333	1	0.933333333	1.071428571
Cadherin-2 OS=Mus musculus GN=Cdh2 PE=1 SV=2	100 kDa	1	24.1404	1.066666667	1	1.066666667	0.9375
Cullin-associated NEDD8-dissociated protein 1 OS=Mus musculus GN=Ca CAND1_MOUSE	136 kDa	1	14.951113	1.066666667	1.133333333	0.941176471	1.0625
Carbonyl reductase [NADPH] 1 OS=Mus musculus GN=Cbr1 PE=1 SV=3	31 kDa	1	20.513207	1.033333333	0.966666667	1.068965517	0.935483871

Cytochrome c oxidase subunit 7A-related protein, mitochondrial OS=Mus COX7R_MOUSE (+1)	12 kDa	1	16.490257	1.1	0.93333333	1.178571429	0.848484848
Catenin beta-1 OS=Mus musculus GN=Ctnnb1 PE=1 SV=1	85 kDa	1	21.237233	1.03333333	1.03333333	1	1
Microtubule-associated protein RP/EB family member 2 (Fragment) OS=ID3YYK8_MOUSE (+2)	29 kDa	1	22.419862	0.86666667	0.86666667	1	1
DNA damage-binding protein 1 OS=Mus musculus GN=Ddb1 PE=1 SV=2	127 kDa	1	15.881756	1	1.13333333	0.882352941	1.133333333
Inter alpha-trypsin inhibitor, heavy chain 4 OS=Mus musculus GN=Itih4 PE9PVD2_MOUSE (+2)	105 kDa	1	8.14969	0.8	0.8	1	1
26S protease regulatory subunit 10B OS=Mus musculus GN=Psmc6 PE=1 PRS10_MOUSE	44 kDa	1	20.165941	0.96666667	1.03333333	0.935483871	1.068965517
60S ribosomal protein L9 OS=Mus musculus GN=Rpl9 PE=2 SV=2	22 kDa	1	24.475066	1	1.06666667	0.9375	1.066666667
SRA stem-loop-interacting RNA-binding protein, mitochondrial OS=Mus SLIRP_MOUSE	13 kDa	1	20.00615	0.86666667	0.9	0.962962963	1.038461538
Transmembrane protein 109 OS=Mus musculus GN=Tmem109 PE=1 SV= TM109_MOUSE	26 kDa	1	18.972135	1.06666667	1.13333333	0.941176471	1.0625
Tubulin polymerization-promoting protein family member 3 OS=Mus muTPPP3_MOUSE	19 kDa	1	21.023977	1.06666667	0.93333333	1.142857143	0.875
Cluster of Hepatoma-derived growth factor OS=Mus musculus GN=Hdgf HDGF_MOUSE [2]	26 kDa	1	12.845804	0.8	0.95	0.842105263	1.1875
Alpha-2-antiplasmin OS=Mus musculus GN=Serpinf2 PE=1 SV=1	55 kDa	1	20.88888	0.93333333	0.96666667	0.965517241	1.035714286
Apolipoprotein B-100 OS=Mus musculus GN=Apob PE=1 SV=1	509 kDa	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Tight junction protein ZO-1 OS=Mus musculus GN=Tjp1 PE=2 SV=1	189 kDa	1	15.30581	0.83333333	0.8	1.041666667	0.96
Filamin-binding LIM protein 1 OS=Mus musculus GN=Fblim1 PE=1 SV=2	41 kDa	1	20.346163	0.96666667	0.9	1.074074074	0.931034483
Inorganic pyrophosphatase OS=Mus musculus GN=Ppa1 PE=1 SV=1	33 kDa	1	19.44908	0.96666667	0.96666667	1	1
UMP-CMP kinase OS=Mus musculus GN=Cmpk1 PE=1 SV=1	22 kDa	1	23.167282	1.03333333	1.1	0.939393939	1.064516129
Proteasome subunit alpha type-6 OS=Mus musculus GN=Psmc6 PE=1 SV=1	27 kDa	1	19.01108	0.93333333	0.96666667	0.965517241	1.035714286
Protein Tfg OS=Mus musculus GN=Tfg PE=2 SV=1	43 kDa	1	20.279233	1	1.03333333	0.967741935	1.033333333
Cluster of Isoform 2 of Dynamin-2 OS=Mus musculus GN=Dnm2 (sp P39 sp P39054-2 DYN2_MOUSE [4	?	1	0	#VALUE!	#VALUE!	#VALUE!	#VALUE!
Calcyclin-binding protein OS=Mus musculus GN=Cacybp PE=1 SV=1	27 kDa	1	20.03453	0.76666667	1	0.766666667	1.304347826
Cluster of Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1 (ITB1_ITB1_MOUSE	88 kDa	1	20.158935	1.13333333	1.03333333	1.096774194	0.911764706
Cluster of Ig gamma-2B chain C region OS=Mus musculus GN=Igh-3 PE=1 IGG2B_MOUSE	44 kDa	1	26.013803	1.13333333	1.16666667	0.971428571	1.029411765
Cluster of Ribosomal protein L15 OS=Mus musculus GN=Gm10020 PE=3 E9QA22_MOUSE [2]	24 kDa	1	19.697095	0.9	1	0.9	1.111111111
6-phosphogluconate dehydrogenase, decarboxylating OS=Mus musculus 6PGD_MOUSE	53 kDa	1	14.397664	1	1.1	0.909090909	1.1
Annexin A11 OS=Mus musculus GN=Anxa11 PE=1 SV=2	54 kDa	1	15.379047	0.95	1	0.95	1.052631579
Clusterin OS=Mus musculus GN=Clu PE=1 SV=1	52 kDa	1	20.539963	0.93333333	0.96666667	0.965517241	1.035714286
Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Mus musculus GN=Fkbp4 FKBP4_MOUSE	52 kDa	1	21.882352	1.06666667	1.23333333	0.864864865	1.15625
FUN14 domain-containing protein 2 OS=Mus musculus GN=Fundc2 PE=2 FUND2_MOUSE	17 kDa	1	20.4234	1.16666667	1.26666667	0.921052632	1.085714286
Fragile X mental retardation syndrome-related protein 1 OS=Mus muscu FXR1_MOUSE	76 kDa	1	19.4449	0.96666667	1.03333333	0.935483871	1.068965517
Ras GTPase-activating protein-binding protein 2 OS=Mus musculus GN=Cg3BP2_MOUSE (+1)	54 kDa	1	6.365376	0.75	0.85	0.882352941	1.133333333
Mimecan OS=Mus musculus GN=Ogn PE=2 SV=1	34 kDa	1	18.601571	0.93333333	1.03333333	0.903225806	1.107142857
E3 ubiquitin-protein ligase NEDD4 OS=Mus musculus GN=Nedd4 PE=1 SV=NEDD4_MOUSE	103 kDa	1	16.120858	1	1.1	0.909090909	1.1
Protein kinase C and casein kinase substrate in neurons protein 2 OS=Mt PACN2_MOUSE	56 kDa	1	19.000797	0.9	0.96666667	0.931034483	1.074074074
Parathyrosin OS=Mus musculus GN=Ptms PE=2 SV=3	11 kDa	1	14.839245	0.83333333	0.86666667	0.961538462	1.04
Acp1 protein OS=Mus musculus GN=Acp1 PE=1 SV=1	18 kDa	1	18.576245	1.2	1.5	0.8	1.25
Ribonuclease inhibitor OS=Mus musculus GN=Rnh1 PE=1 SV=1	50 kDa	1	18.973845	0.93333333	1.1	0.848484848	1.178571429
Spartin OS=Mus musculus GN=Spg20 PE=2 SV=1	73 kDa	1	6.833901	0.56666667	0.66666667	0.85	1.176470588
Slc8a1 protein OS=Mus musculus GN=Slc8a1 PE=2 SV=1	107 kDa	1	13.338606	1.1	1	1.1	0.909090909
Glutathione peroxidase 3 OS=Mus musculus GN=Gpx3 PE=2 SV=2	25 kDa	1	14.786617	0.86666667	1.03333333	0.838709677	1.192307692
Microtubule-associated protein 6 OS=Mus musculus GN=Map6 PE=1 SV= MAP6_MOUSE	96 kDa	1	0	0.9	0.9	1	1
Sorting nexin 3 OS=Mus musculus GN=Snx3 PE=2 SV=1	19 kDa	1	20.551792	0.9	0.93333333	0.964285714	1.037037037
26S protease regulatory subunit 8 OS=Mus musculus GN=Psmc5 PE=1 SV= PRS8_MOUSE	46 kDa	1	12.993085	1	1.06666667	0.9375	1.066666667
Cluster of Cysteine desulfurase, mitochondrial OS=Mus musculus GN=Nf NFS1_MOUSE	50 kDa	1	6.491713	1.06666667	1.06666667	1	1
3-hydroxyisobutyrate dehydrogenase, mitochondrial OS=Mus musculus 3HIDH_MOUSE	35 kDa	1	21.619181	1.1	1.16666667	0.942857143	1.060606061
Growth factor receptor-bound protein 2 OS=Mus musculus GN=Grb2 PE=B1AT92_MOUSE (+1)	24 kDa	1	20.247634	0.9	0.93333333	0.964285714	1.037037037
Complement factor I OS=Mus musculus GN=Cfi PE=1 SV=3	67 kDa	1	13.86267	1.03333333	1	1.033333333	0.967741935
Cytoskeleton-associated protein 4 OS=Mus musculus GN=Ckap4 PE=2 SV= CKAP4_MOUSE	64 kDa	1	20.963126	0.86666667	0.93333333	0.928571429	1.076923077
Desmoglein-2 OS=Mus musculus GN=Dsg2 PE=1 SV=3	122 kDa	1	12.590171	1.06666667	1.06666667	1	1
Epoxide hydrolase 1 OS=Mus musculus GN=Ephx1 PE=2 SV=1	51 kDa	1	21.358167	0.93333333	0.86666667	1.076923077	0.928571429
Eukaryotic translation initiation factor 3 subunit C OS=Mus musculus GN EIF3C_MOUSE	106 kDa	1	6.7104675	1	0.9	1.111111111	0.9
Protein ETHE1, mitochondrial OS=Mus musculus GN=Ethe1 PE=1 SV=2	28 kDa	1	22.189054	1	0.93333333	1.071428571	0.933333333
Eukaryotic translation initiation factor 2 subunit 1 OS=Mus musculus GN IF2A_MOUSE	36 kDa	1	19.372503	0.9	0.9	1	1

Platelet-activating factor acetylhydrolase IB subunit alpha OS=Mus musc LIS1_MOUSE	47 kDa	1	18.949524	1	0.96666667	1.034482759	0.96666667
Palladin OS=Mus musculus GN=Palld PE=1 SV=2	PALLD_MOUSE (+1)	152 kDa	6.361226	0.9	0.83333333	1.08	0.92592592
Prostaglandin E synthase 2 OS=Mus musculus GN=Ptges2 PE=1 SV=3	PGES2_MOUSE	43 kDa	19.298534	1.03333333	1.03333333	1	1
Cluster of 26S proteasome non-ATPase regulatory subunit 13 OS=Mus m PSD13_MOUSE		43 kDa	19.34438	0.96666667	1	0.96666667	1.034482759
Target of Myb protein 1 OS=Mus musculus GN=Tom1 PE=2 SV=1	Q3UDC3_MOUSE	57 kDa	20.43223	0.76666667	0.86666667	0.884615385	1.130434783
60S ribosomal protein L36a OS=Mus musculus GN=Rpl36a PE=2 SV=2	RL36A_MOUSE	12 kDa	20.600867	0.9	1.1	0.818181818	1.22222222
Small glutamine-rich tetratricopeptide repeat-containing protein alpha CSGTA_MOUSE (+1)		34 kDa	14.623615	0.8	0.76666667	1.043478261	0.95833333
Serine/arginine-rich splicing factor 3 OS=Mus musculus GN=Srsf3 PE=1 S' SRSF3_MOUSE		19 kDa	13.05389	0.93333333	0.93333333	1	1
Carbonyl reductase family member 4 OS=Mus musculus GN=Cbr4 PE=2 SCBR4_MOUSE		25 kDa	6.1511613	1	1.1	0.909090909	1.1
Epimerase family protein SDR39U1 OS=Mus musculus GN=Sdr39u1 PE=1 D39U1_MOUSE (+1)		33 kDa	20.439826	0.93333333	1.03333333	0.903225806	1.107142857
Cluster of 60S ribosomal protein L35 OS=Mus musculus GN=Rpl35 PE=2 ; RL35_MOUSE		15 kDa	18.864031	0.9	1.13333333	0.794117647	1.259259259
Cluster of Heterogeneous nuclear ribonucleoprotein D0 OS=Mus muscul HNRPD_MOUSE [4]		38 kDa	652.7471	0.85	1.2	0.708333333	1.411764706
Cluster of SH3 domain-binding glutamic acid-rich protein OS=Mus muscu SH3BG_MOUSE		23 kDa	13.148588	1.03333333	1.03333333	1	1
Tyrosine--tRNA ligase, cytoplasmic OS=Mus musculus GN=Yars PE=4 SV= A2A757_MOUSE (+1)		63 kDa	12.950303	0.73333333	0.83333333	0.88	1.136363636
Annexin A3 OS=Mus musculus GN=Anxa3 PE=1 SV=4	ANXA3_MOUSE	36 kDa	22.18531	1	1.06666667	0.9375	1.06666667
Protein Gm20547 OS=Mus musculus GN=Gm20547 PE=2 SV=1	B8JJN0_MOUSE (+1)	142 kDa	13.875271	0.9	0.93333333	0.964285714	1.037037037
Corticosteroid-binding globulin OS=Mus musculus GN=Serpina6 PE=1 SV CBG_MOUSE		45 kDa	22.933714	0.83333333	0.7	1.19047619	0.84
Chloride intracellular channel protein 4 OS=Mus musculus GN=Clic4 PE=: CLIC4_MOUSE		29 kDa	21.89597	0.86666667	1.03333333	0.838709677	1.192307692
Glycogen [starch] synthase, muscle OS=Mus musculus GN=Gys1 PE=2 SV D3Z0Q6_MOUSE (+1)		77 kDa	22.380893	1.06666667	1.03333333	1.032258065	0.96875
Protein Coq10a OS=Mus musculus GN=Coq10a PE=2 SV=1	E9Q3H6_MOUSE	29 kDa	12.410659	1.13333333	1.16666667	0.971428571	1.029411765
Elongation factor Ts, mitochondrial OS=Mus musculus GN=Tsfm PE=2 SV EFTS_MOUSE		35 kDa	21.897374	1.03333333	1.13333333	0.911764706	1.096774194
RNA-binding protein FUS (Fragment) OS=Mus musculus GN=Fus PE=4 SV G3UXT7_MOUSE		14 kDa	21.779045	0.86666667	0.9	0.962962963	1.038461538
Valine--tRNA ligase (Fragment) OS=Mus musculus GN=Vars PE=2 SV=1	G3UY93_MOUSE	141 kDa	18.587115	0.9	1	0.9	1.111111111
Inter-alpha trypsin inhibitor, heavy chain 2 OS=Mus musculus GN=Itih2 PG3X977_MOUSE (+1)		106 kDa	19.237103	0.93333333	0.96666667	0.965517241	1.035714286
Matrin-3 OS=Mus musculus GN=Matr3 PE=1 SV=1	MATR3_MOUSE	95 kDa	12.629765	0.9	1	0.9	1.111111111
Ribosylidihydroxynicotinamide dehydrogenase [quinone] OS=Mus musculu: NQO2_MOUSE		26 kDa	20.096532	1.16666667	1.2	0.972222222	1.028571429
Prefoldin subunit 2 OS=Mus musculus GN=Pfdn2 PE=2 SV=2	PF2_MOUSE	17 kDa	13.970052	1	1	1	1
Proteasome subunit beta type-6 OS=Mus musculus GN=Psb6 PE=1 SV= PSB6_MOUSE		25 kDa	25.282553	0.96666667	0.93333333	1.035714286	0.965517241
Multifunctional protein ADE2 OS=Mus musculus GN=Paics PE=1 SV=4	PUR6_MOUSE	47 kDa	18.888196	0.93333333	1	0.93333333	1.071428571
40S ribosomal protein S21 OS=Mus musculus GN=Rps21 PE=2 SV=1	RS21_MOUSE	9 kDa	13.698325	0.93333333	1	0.93333333	1.071428571
40S ribosomal protein SA OS=Mus musculus GN=Rpsa PE=1 SV=4	RSSA_MOUSE	33 kDa	23.045151	1.06666667	1.13333333	0.941176471	1.0625
NAD-dependent protein deacetylase sirtuin-5, mitochondrial OS=Mus mus SIR5_MOUSE		34 kDa	20.148633	0.96666667	1	0.96666667	1.034482759
T-complex protein 1 subunit gamma OS=Mus musculus GN=Cct3 PE=1 SV TCGP_MOUSE		61 kDa	22.54952	0.93333333	1.06666667	0.875	1.142857143
Hematological and neurological expressed 1 protein OS=Mus musculus CHN1_MOUSE		16 kDa	15.79018	0.86666667	0.7	1.238095238	0.807692308
28S ribosomal protein S5, mitochondrial OS=Mus musculus GN=Mrrps5 P RT05_MOUSE		48 kDa	29.080287	1.06666667	1.1	0.96969697	1.03125
Adrenodoxin, mitochondrial OS=Mus musculus GN=Fdx1 PE=2 SV=1	ADX_MOUSE	20 kDa	19.335504	1.03333333	0.86666667	1.192307692	0.838709677
Protein 1300017J02Rik OS=Mus musculus GN=1300017J02Rik PE=2 SV=1 D3Y36_MOUSE (+1)		69 kDa	8.220224	1	1	1	1
Cluster of Dynein light chain 2, cytoplasmic OS=Mus musculus GN=Dynll2 DYL2_MOUSE [2]		10 kDa	165.69188	1.05	1.3	0.807692308	1.238095238
Cluster of Mitochondrial import inner membrane translocase subunit TinTIM8A_MOUSE		11 kDa	14.351054	0.96666667	0.73333333	1.318181818	0.75862069
Cytoplasmic dynein 1 intermediate chain 2 OS=Mus musculus GN=Dync1A2BFF5_MOUSE (+4)		71 kDa	20.015031	0.96666667	0.9	1.074074074	0.931034483
Flavin reductase (NADPH) OS=Mus musculus GN=Blvrb PE=2 SV=3	BLVRB_MOUSE	22 kDa	12.3405635	1	1	1	1
Adenylyl cyclase-associated protein 1 OS=Mus musculus GN=Cap1 PE=1 :CAP1_MOUSE		52 kDa	14.57512	1	1.16666667	0.857142857	1.16666667
Enoyl-CoA hydratase domain-containing protein 2, mitochondrial (Fragm F6Z5N5_MOUSE (+1)		13 kDa	13.093527	1.16666667	1.3	0.897435897	1.114285714
High mobility group protein B2 OS=Mus musculus GN=Hmgb2 PE=1 SV=3 HMGB2_MOUSE		24 kDa	12.952177	1	1.06666667	0.9375	1.06666667
Heat shock protein 105 kDa OS=Mus musculus GN=Hsph1 PE=1 SV=2	HS105_MOUSE (+1)	96 kDa	13.001297	0.63333333	0.76666667	0.826086957	1.210526316
Importin subunit beta-1 OS=Mus musculus GN=Kpnb1 PE=1 SV=2	IMB1_MOUSE	97 kDa	12.750782	0.9	1	0.9	1.111111111
Cob(II)rinic acid a,c-diamide adenosyltransferase, mitochondrial OS=Mus MMAB_MOUSE		26 kDa	13.419797	1	0.86666667	1.153846154	0.86666667
Non-POU domain-containing octamer-binding protein OS=Mus musculus: NONO_MOUSE		55 kDa	6.176474	1.05	1.05	1	1
Nucleosome assembly protein 1-like 4 OS=Mus musculus GN=Nap1l4 PE=NP1L4_MOUSE		43 kDa	21.146168	0.93333333	1.03333333	0.903225806	1.107142857
Nuclear migration protein nudC OS=Mus musculus GN=Nudc PE=1 SV=1	NUDC_MOUSE	38 kDa	12.327932	0.83333333	1	0.83333333	1.2
Obg-like ATPase 1 OS=Mus musculus GN=Ola1 PE=1 SV=1	OLA1_MOUSE	45 kDa	12.784268	1.03333333	0.96666667	1.068965517	0.935483871
Heterogeneous nuclear ribonucleoprotein H OS=Mus musculus GN=Hnrr Q8C2Q7_MOUSE		51 kDa	20.153762	0.9	0.96666667	0.931034483	1.074074074
40S ribosomal protein S20 OS=Mus musculus GN=Rps20 PE=1 SV=1	RS20_MOUSE	13 kDa	19.110315	0.96666667	0.96666667	1	1

Actin-related protein 2 OS=Mus musculus GN=Actr2 PE=1 SV=1	ARP2_MOUSE	45 kDa	1	13.074563	0.9	1	0.9	1.111111111
LIM and senescent cell antigen-like-containing domain protein 1 OS=Mu: E9QP62_MOUSE (+1)		44 kDa	1	20.668668	1.1	1.13333333	0.970588235	1.03030303
Cluster of ATP-dependent RNA helicase DDX3X OS=Mus musculus GN=D:DDX3X_MOUSE		73 kDa	1	20.662232	0.966666667	1.06666667	0.90625	1.103448276
Cluster of Mitochondrial calcium uniporter regulator 1 OS=Mus musculus: MCUR1_MOUSE		38 kDa	1	6.2511116	0.9	0.93333333	0.964285714	1.037037037
Cluster of Ras-related protein Rap-1b OS=Mus musculus GN=Rap1b PE=2 RAP1B_MOUSE [2]		21 kDa	1	8.14968	0.85	0.95	0.894736842	1.117647059
Cluster of Serine/arginine-rich-splicing factor 4 OS=Mus musculus GN=Sr A2A837_MOUSE		56 kDa	1	12.12454	0.8	0.85	0.941176471	1.0625
ATP-binding cassette sub-family B member 7, mitochondrial OS=Mus mu ABCB7_MOUSE		83 kDa	1	17.686784	1.05	0.85	1.235294118	0.80952381
CDGSH iron sulfur domain 3 OS=Mus musculus GN=Cisd3 PE=2 SV=1	B2RWE3_MOUSE (+1)	15 kDa	1	19.201045	0.966666667	1.03333333	0.935483871	1.068965517
Calpain-1 catalytic subunit OS=Mus musculus GN=Capn1 PE=2 SV=1	CAN1_MOUSE	82 kDa	1	20.385556	0.966666667	1	0.966666667	1.034482759
Coatomer subunit delta OS=Mus musculus GN=Arcn1 PE=2 SV=2	COPD_MOUSE	57 kDa	1	15.031162	0.75	0.75	1	1
Endothelial differentiation-related factor 1 OS=Mus musculus GN=Edf1 P EDF1_MOUSE		16 kDa	1	6.507688	0.933333333	0.86666667	1.076923077	0.928571429
Protein-cysteine N-palmitoyltransferase HHAT-like protein OS=Mus musci HHATL_MOUSE		56 kDa	1	26.862833	1.233333333	1.23333333	1	1
Ig gamma-3 chain C region OS=Mus musculus PE=1 SV=2	IGHG3_MOUSE	44 kDa	1	21.714237	1.1	1.16666667	0.942857143	1.060606061
LYR motif-containing protein 4 OS=Mus musculus GN=Lym4 PE=2 SV=1	LYRM4_MOUSE	11 kDa	1	19.110554	1.066666667	1.06666667	1	1
NFU1 iron-sulfur cluster scaffold homolog, mitochondrial OS=Mus musci NFU1_MOUSE		29 kDa	1	19.404855	0.966666667	0.83333333	1.16	0.862068966
1-acyl-sn-glycerol-3-phosphate acyltransferase gamma OS=Mus musculus PLCC_MOUSE		43 kDa	1	20.290712	1.1	1.13333333	0.970588235	1.03030303
Proteasome subunit alpha type-3 OS=Mus musculus GN=Psm3 PE=1 SV PSA3_MOUSE		28 kDa	1	12.59461	0.933333333	0.93333333	1	1
28S ribosomal protein S31, mitochondrial OS=Mus musculus GN=Mrps31RT31_MOUSE		44 kDa	1	13.308081	1.066666667	1.06666667	1	1
tRNA-splicing ligase RtcB homolog OS=Mus musculus GN=D10Wsu52e PIRTCB_MOUSE		55 kDa	1	15.091649	0.966666667	0.96666667	1	1
Selenoprotein P OS=Mus musculus GN=Sepp1 PE=2 SV=3	SEPP1_MOUSE	43 kDa	1	33.16094	1.3	0.75	1.733333333	0.576923077
PDZ and LIM domain protein 1 OS=Mus musculus GN=Pdlim1 PE=2 SV=4	PDL1_MOUSE	36 kDa	1	12.832835	0.833333333	0.76666667	1.086956522	0.92
Transmembrane protein 65 OS=Mus musculus GN=Tmem65 PE=2 SV=1	TMM65_MOUSE	25 kDa	1	20.73422	1.066666667	1.13333333	0.941176471	1.0625
Protein S100-A13 OS=Mus musculus GN=S100a13 PE=1 SV=1	S10AD_MOUSE	11 kDa	1	12.679668	0.9	0.86666667	1.038461538	0.962962963
Cluster of Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2 (CYB5E:CYB5_MOUSE [2])		15 kDa	1	21.050896	1.033333333	1.03333333	1	1
Ig gamma-2A chain C region secreted form OS=Mus musculus PE=1 SV=1 GCAB_MOUSE		37 kDa	1	15.098368	1.066666667	0.83333333	1.28	0.78125
Arginine-tRNA ligase, cytoplasmic OS=Mus musculus GN=Rars PE=2 SV= SYRC_MOUSE		76 kDa	1	15.883525	0.95	0.15	6.333333333	0.157894737
Isoform 2 of Hepatoma-derived growth factor-related protein 3 OS=Mus sp Q9JMG7-2 HDGR3_MOUSE		?	1	6.1799414	0.966666667	0.96666667	1	1
Cluster of Ribose-phosphate pyrophosphokinase OS=Mus musculus GN=G3UXL2_MOUSE [2]		35 kDa	1	19.531101	0.766666667	0.9	0.851851852	1.173913043
Cluster of Charged multivesicular body protein 4b OS=Mus musculus GN CHM4B_MOUSE		25 kDa	1	13.183835	1.1	1	1.1	0.909090909
Cluster of Oxysterol-binding protein 1 OS=Mus musculus GN=Osbp PE=1 OSBP1_MOUSE		89 kDa	1	13.223459	0.95	1.1	0.863636364	1.157894737
Cytosolic 5'-nucleotidase 3 OS=Mus musculus GN=Nt5c3 PE=1 SV=4	5NT3_MOUSE	37 kDa	1	20.7719	0.866666667	0.93333333	0.928571429	1.076923077
Actin-related protein 3 OS=Mus musculus GN=Actr3 PE=1 SV=3	ARP3_MOUSE	47 kDa	1	21.758854	0.8	0.83333333	0.96	1.041666667
Carbonic anhydrase 1 OS=Mus musculus GN=Ca1 PE=2 SV=4	CAH1_MOUSE	28 kDa	1	13.645886	1.35	1.15	1.173913043	0.851851852
Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 P CDC42_MOUSE		21 kDa	1	20.504995	1	1.03333333	0.967741935	1.033333333
Coiled-coil-helix-coiled-coil-helix domain-containing protein 6, mitochondon CHCH6_MOUSE (+1)		30 kDa	1	19.882113	0.866666667	0.76666667	1.130434783	0.884615385
MCG148436 OS=Mus musculus GN=1700014D04Rik PE=4 SV=1	J3QMS2_MOUSE	109 kDa	1	6.740225	0.7	0.6	1.166666667	0.857142857
Mitochondrial intermediate peptidase OS=Mus musculus GN=Mipep PE= MIPEP_MOUSE		81 kDa	1	7.379151	1.1	1.15	0.956521739	1.045454545
Methionine-R-sulfoxide reductase B3, mitochondrial OS=Mus musculus (MSRB3_MOUSE		27 kDa	1	13.802345	1	1	1	1
Omega-amidase NIT2 OS=Mus musculus GN=Nit2 PE=1 SV=1	NIT2_MOUSE	31 kDa	1	15.235044	1.2	1.03333333	1.161290323	0.861111111
26S proteasome non-ATPase regulatory subunit 6 OS=Mus musculus GN:PSMD6_MOUSE		46 kDa	1	28.66693	1.166666667	1.13333333	1.029411765	0.971428571
Tyrosine-protein phosphatase non-receptor type 11 OS=Mus musculus CPTN11_MOUSE (+1)		68 kDa	1	15.914597	1.166666667	1.06666667	1.09375	0.914285714
Ribosome maturation protein SBDS OS=Mus musculus GN=5bds PE=1 SV SBDS_MOUSE		29 kDa	1	13.937628	0.933333333	1.03333333	0.903225806	1.107142857
Ubiquitin carboxyl-terminal hydrolase isozyme L3 OS=Mus musculus GN: UCHL3_MOUSE		26 kDa	1	35.251498	0.933333333	0.93333333	1	1
Isoform 2 of Protein CDV3 OS=Mus musculus GN=Cdv3 sp Q4VAA2-2 CDV3_MOUSE		?	1	22.427146	0.833333333	0.7	1.19047619	0.84
Acyl-CoA synthetase family member 2, mitochondrial OS=Mus musculus ACSF2_MOUSE		68 kDa	1	6.544755	0.933333333	0.83333333	1.12	0.892857143
Echinoderm microtubule-associated protein-like 1 OS=Mus musculus GNEMAL1_MOUSE		90 kDa	1	8.173438	0.9	1.06666667	0.84375	1.185185185
von Willebrand factor A domain-containing protein 5A OS=Mus musculus VMA5A_MOUSE		87 kDa	1	12.77978	0.9	1.03333333	0.870967742	1.148148148
Heterogeneous nuclear ribonucleoprotein Q OS=Mus musculus GN=Sync G3UZI2_MOUSE		59 kDa	1	10.1466365	0.733333333	0.93333333	0.785714286	1.272727273
Neurofilament heavy polypeptide OS=Mus musculus GN=Nefh PE=1 SV=: NFH_MOUSE		117 kDa	1	0	1.2	1.3	0.923076923	1.083333333
Cluster of Annexin A1 OS=Mus musculus GN=Anxa1 PE=1 SV=2 (ANXA1_ ANXA1_MOUSE		39 kDa	1	20.375729	1	1	1	1
Cluster of Serine (Or cysteine) peptidase inhibitor, clade A, member 3N, G3X8T9_MOUSE [2]		47 kDa	1	0	0.6	0.6	1	1
ATP-dependent RNA helicase DDX1 OS=Mus musculus GN=Ddx1 PE=1 SV DDX1_MOUSE		83 kDa	1	13.02261	0.9	0.86666667	1.038461538	0.962962963
NADH-cytochrome b5 reductase 3 OS=Mus musculus GN=Cyb5r3 PE=2 S'F2Z456_MOUSE (+1)		35 kDa	1	12.503378	1.033333333	1.03333333	1	1

Mitochondrial glutamate carrier 1 OS=Mus musculus GN=Slc25a22 PE=1 GHC1_MOUSE	35 kDa	1	12.737799	1.1	1.1	1	1
Glutaredoxin-related protein 5, mitochondrial OS=Mus musculus GN=Glr GLRX5_MOUSE	16 kDa	1	18.214183	0.866666667	0.833333333	1.04	0.961538462
Homeodomain-only protein OS=Mus musculus GN=Hopx PE=1 SV=1 HOP_MOUSE	8 kDa	1	22.338365	0.9	0.8	1.125	0.888888889
Probable D-lactate dehydrogenase, mitochondrial OS=Mus musculus GN LDHD_MOUSE	52 kDa	1	16.007045	0.9	1	0.9	1.111111111
Trans-2-enoyl-CoA reductase, mitochondrial OS=Mus musculus GN=Mec MECR_MOUSE	40 kDa	1	22.569466	1.033333333	1.133333333	0.911764706	1.096774194
Nucleobindin-1 OS=Mus musculus GN=Nucb1 PE=1 SV=2 NUCB1_MOUSE	53 kDa	1	22.663568	0.933333333	0.9	1.037037037	0.964285714
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa s OST48_MOUSE	49 kDa	1	6.93763	0.966666667	0.866666667	1.115384615	0.896551724
Protein kinase C and casein kinase II substrate protein 3 OS=Mus musculus GN=PACN3_MOUSE	49 kDa	1	15.184765	1	1	1	1
Proteasome subunit beta type-7 OS=Mus musculus GN=Psb7 PE=1 SV=PSB7_MOUSE	30 kDa	1	14.838177	0.966666667	1	0.966666667	1.034482759
Serum albumin OS=Bos taurus GN=ALB PE=1 SV=4 ALBU_BOVIN	69 kDa	1	23.466004	0.866666667	1.033333333	0.838709677	1.192307692
Major vault protein OS=Mus musculus GN=Mvp PE=2 SV=1 E9Q3X0_MOUSE (+1)	97 kDa	1	13.458279	0.9	0.866666667	1.038461538	0.962962963
GMP reductase 1 OS=Mus musculus GN=Gmpr PE=2 SV=1 GMPR1_MOUSE	37 kDa	1	13.633686	1.033333333	1.1	0.939393939	1.064516129
LDLR chaperone MESD OS=Mus musculus GN=Mesdc2 PE=1 SV=1 MESD_MOUSE	25 kDa	1	12.68479	0.9	0.933333333	0.964285714	1.037037037
Delta-sarcoglycan OS=Mus musculus GN=Sgcd PE=1 SV=1 SGCD_MOUSE	32 kDa	1	13.283147	1.15	1.1	1.045454545	0.956521739
Translocase of inner mitochondrial membrane domain-containing protein TIDC1_MOUSE	32 kDa	1	12.383413	1.166666667	1.1	1.060606061	0.942857143
Sorting nexin-12 OS=Mus musculus GN=Snx12 PE=2 SV=1 SNX12_MOUSE	19 kDa	1	13.479751	0.95	0.9	1.055555556	0.947368421
Heterogeneous nuclear ribonucleoprotein R OS=Mus musculus GN=Hnrr Q8VHM5_MOUSE	71 kDa	1	8.2579695	0.9	0.9	1	1
Cluster of Muscblind-like 1 (Drosophila), isoform CRA_a OS=Mus musculus GN=G3X9Q0_MOUSE [6]	41 kDa	1	0	0.7	1	0.7	1.428571429
Cluster of Histone-binding protein RBBP7 OS=Mus musculus GN=Rbbp7 IA2AFI9_MOUSE [3]	44 kDa	1	6.847463	0.8	0.85	0.941176471	1.0625
Cluster of Histone H2A.Z OS=Mus musculus GN=H2afz PE=1 SV=2 (H2AZ_H2AZ_MOUSE	14 kDa	1	13.498217	1	1.066666667	0.9375	1.066666667
Cluster of Dihydropyrimidinase-related protein 3 OS=Mus musculus GN=E9PWE8_MOUSE	74 kDa	1	7.326679	0.9	0.9	1	1
Coiled-coil domain-containing protein 90B, mitochondrial OS=Mus musculus GN=CC90B_MOUSE	30 kDa	1	12.644841	0.933333333	1	0.933333333	1.071428571
Gap junction alpha-1 protein OS=Mus musculus GN=Gja1 PE=1 SV=2 CXA1_MOUSE	43 kDa	1	20.89954	1.133333333	1.466666667	0.772727273	1.294117647
Vacuolar protein sorting-associated protein 29 OS=Mus musculus GN=Vf D3Z645_MOUSE (+1)	16 kDa	1	6.3425384	1.066666667	1.133333333	0.941176471	1.0625
Glucosidase 2 subunit beta OS=Mus musculus GN=Prkcsh PE=1 SV=1 GLU2B_MOUSE	59 kDa	1	13.613571	0.866666667	0.8	1.083333333	0.923076923
Complement component 1 Q subcomponent-binding protein, mitochondria Q8R5L1_MOUSE	31 kDa	1	29.55775	1.6	1.4	1.142857143	0.875
UV excision repair protein RAD23 homolog B OS=Mus musculus GN=Rad RD23B_MOUSE	44 kDa	1	23.626433	0.9	0.9	1	1
40S ribosomal protein S23 OS=Mus musculus GN=Rps23 PE=2 SV=3 RS23_MOUSE	16 kDa	1	19.126426	0.966666667	1.066666667	0.90625	1.103448276
Aldehyde dehydrogenase family 8 member A1 OS=Mus musculus GN=Al AL8A1_MOUSE	54 kDa	1	6.6674635	1.033333333	1.233333333	0.837837838	1.193548387
Basal cell adhesion molecule OS=Mus musculus GN=Bcam PE=2 SV=1 BCAM_MOUSE	68 kDa	1	12.4283746	1.066666667	1.133333333	0.941176471	1.0625
Thioredoxin domain-containing protein 5 OS=Mus musculus GN=Txndc5 E9PXX7_MOUSE	39 kDa	1	13.339835	1	0.9	1.111111111	0.9
Prostamide/prostaglandin F synthase OS=Mus musculus GN=Fam213b P PGFS_MOUSE	22 kDa	1	13.191435	0.7	0.833333333	0.84	1.19047619
UV excision repair protein RAD23 homolog A (Fragment) OS=Mus musculus GN=Q3TN85_MOUSE (+2)	35 kDa	1	13.322325	1.2	1.15	1.043478261	0.958333333
Protein Taf15 OS=Mus musculus GN=Taf15 PE=2 SV=1 Q8BQ46_MOUSE	59 kDa	1	9.34846	1.15	0.85	1.352941176	0.739130435
Beta-casein OS=Bos taurus GN=CSN2 PE=1 SV=2 CASB_BOVIN	25 kDa	1	29.036477	1.066666667	0.666666667	1.6	0.625
Histone H1t OS=Mus musculus GN=Hist1h1t PE=1 SV=4 H1T_MOUSE (+1)	22 kDa	1	0	1.1	0.9	1.222222222	0.818181818
5'-AMP-activated protein kinase catalytic subunit alpha-2 OS=Mus musculus GN=AAPK2_MOUSE	62 kDa	1	12.972993	1.066666667	1.033333333	1.032258065	0.96875
Heat shock protein beta-2 OS=Mus musculus GN=Hspb2 PE=2 SV=2 HSPB2_MOUSE	20 kDa	1	20.600115	0.9	0.8	1.125	0.888888889
Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Ma MARCS_MOUSE	30 kDa	1	14.273693	1.1	1	1.1	0.909090909
Microtubule-associated protein RP/EB family member 1 OS=Mus musculus GN=MARE1_MOUSE	30 kDa	1	12.230023	0.933333333	0.933333333	1	1
Proteasome subunit beta type-1 OS=Mus musculus GN=Psb1 PE=1 SV=PSB1_MOUSE	26 kDa	1	19.483042	1.033333333	1.033333333	1	1
Solute carrier family 25 member 46 OS=Mus musculus GN=Slc25a46 PE=S2546_MOUSE	46 kDa	1	18.866791	0.966666667	1	0.966666667	1.034482759
Dnaj homolog subfamily A member 1 OS=Mus musculus GN=Dnaja1 PE=DNAJ1_MOUSE	45 kDa	1	12.2515775	0.8	0.95	0.842105263	1.1875
Aspartyl aminopeptidase OS=Mus musculus GN=Dnpep PE=2 SV=2 DNPEP_MOUSE	52 kDa	1	12.876654	1	1.133333333	0.882352941	1.133333333
Dual specificity mitogen-activated protein kinase kinase 1 OS=Mus musculus GN=MP2K1_MOUSE	43 kDa	1	20.728895	0.933333333	0.966666667	0.965517241	1.035714286
Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2A2AL85_MOUSE (+1)	81 kDa	1	7.269213	1	1.1	0.909090909	1.1
Proteasome subunit alpha type-5 OS=Mus musculus GN=Psm5 PE=1 SV=PSA5_MOUSE	26 kDa	1	20.520194	0.833333333	1	0.833333333	1.2
Microtubule-associated protein RP/EB family member 3 OS=Mus musculus GN=MARE3_MOUSE	32 kDa	1	0	0.95	1	0.95	1.052631579
Monoglyceride lipase OS=Mus musculus GN=Mgl1 PE=2 SV=1 D3YYS6_MOUSE (+3)	37 kDa	1	12.48107	1.066666667	1.066666667	1	1
Fibroblast growth factor 1 OS=Mus musculus GN=Fgf1 PE=2 SV=1 FGF1_MOUSE	17 kDa	1	44.51803	1.033333333	0.9	1.148148148	0.870967742
Glycerophosphocholine phosphodiesterase GPCPD1 OS=Mus musculus GN=GPCP1_MOUSE	77 kDa	1	7.084916	1	1	1	1
Vesicle-associated membrane protein 3 OS=Mus musculus GN=Vamp3 P VAMP3_MOUSE	11 kDa	1	6.1215352	1.033333333	1.033333333	1	1
Mitochondrial antiviral-signaling protein OS=Mus musculus GN=Mavs PE=MAVS_MOUSE	53 kDa	1	12.037785	1	1	1	1

Biglycan OS=Mus musculus GN=Bgn PE=2 SV=1	PGS1_MOUSE	42 kDa	1	6.314594	1.1	0.93333333	1.178571429	0.848484848
Nitric oxide synthase, inducible OS=Mus musculus GN=Nos2 PE=1 SV=1	NOS2_MOUSE	131 kDa	1	6.3699104	0.9	0.8	1.125	0.888888889
Annexin A4 OS=Mus musculus GN=Anxa4 PE=2 SV=4	ANXA4_MOUSE	36 kDa	1	7.121446	1	0.9	1.111111111	0.9
F-actin-capping protein subunit alpha-1 OS=Mus musculus GN=Capza1 P CAZA1_MOUSE (+1)	CAZA1_MOUSE (+1)	33 kDa	1	13.759677	1.033333333	0.96666667	1.068965517	0.935483871
Cluster of Mitochondrial-processing peptidase subunit beta OS=Mus mu:MPPB_MOUSE	MPPB_MOUSE	55 kDa	1	19.629665	0.933333333	1.1	0.848484848	1.178571429
Protein Gyk OS=Mus musculus GN=Gyk PE=2 SV=1	B1ASZ3_MOUSE (+1)	60 kDa	1	13.799746	0.9	0.93333333	0.964285714	1.037037037
Alpha-endosulfine OS=Mus musculus GN=Ensa PE=1 SV=1	ENSA_MOUSE (+1)	13 kDa	1	16.527696	1.1	1.05	1.047619048	0.954545455
Mitochondrial import inner membrane translocase subunit Tim8 B OS=MTIM8B_MOUSE	MTIM8B_MOUSE	9 kDa	1	12.960956	0.8	0.95	0.842105263	1.1875
Uncharacterized protein C19orf43 homolog OS=Mus musculus PE=2 SV= CS043_MOUSE	CS043_MOUSE	18 kDa	1	14.598294	1.166666667	1.03333333	1.129032258	0.885714286
rRNA 2'-O-methyltransferase fibrillar OS=Mus musculus GN=Fbl PE=1 SFBRL_MOUSE	SFBRL_MOUSE	34 kDa	1	0	0.933333333	1	0.933333333	1.071428571
Cluster of Agrin OS=Mus musculus GN=Agrn PE=4 SV=1 (M0QWP1_MOUSE M0QWP1_MOUSE)	M0QWP1_MOUSE M0QWP1_MOUSE	217 kDa	1	6.987554	0.95	0.85	1.117647059	0.894736842
Limb and neural patterns, isoform CRA_c OS=Mus musculus GN=Lnp PE=A2ASL8_MOUSE (+1)	A2ASL8_MOUSE (+1)	42 kDa	1	6.77235	0.95	1.1	0.863636364	1.157894737
6-phosphofructokinase, liver type OS=Mus musculus GN=Pfkl PE=1 SV=4 K6PL_MOUSE	K6PL_MOUSE	85 kDa	1	8.306942	0.9	1.15	0.782608696	1.277777778
ATP-dependent (S)-NAD(P)H-hydrate dehydratase OS=Mus musculus GN=J3QMM7_MOUSE (+4)	J3QMM7_MOUSE (+4)	35 kDa	1	13.150911	0.95	0.95	1	1
1,4-alpha-glucan-branching enzyme OS=Mus musculus GN=Gbe1 PE=4 S'F6ZHD8_MOUSE (+1)	S'F6ZHD8_MOUSE (+1)	80 kDa	1	6.1466843	1.15	1.25	0.92	1.086956522
Protein unc-45 homolog A (Fragment) OS=Mus musculus GN=Unc45a PE D3YZN8_MOUSE	D3YZN8_MOUSE	19 kDa	1	6.113683	1.1	1	1.1	0.909090909
sp K22E_HUMAN	sp K22E_HUMAN	?	1	0	0.8	1.1	0.727272727	1.375
Uncharacterized protein C1orf170 homolog OS=Mus musculus PE=2 SV= CA170_MOUSE	CA170_MOUSE	88 kDa	1	7.537425	0.85	0.7	1.214285714	0.823529412
F-actin-capping protein subunit alpha-2 OS=Mus musculus GN=Capza2 P CAZA2_MOUSE	CAZA2_MOUSE	33 kDa	1	0	0.9	0.93333333	0.964285714	1.037037037
Crk-like protein OS=Mus musculus GN=Crkl PE=1 SV=2	CRKL_MOUSE	34 kDa	1	7.626892	1	1.1	0.909090909	1.1
DNA excision repair protein ERCC-6-like OS=Mus musculus GN=Ercc6l PE ERC6L_MOUSE	ERC6L_MOUSE	139 kDa	1	0	1.5	1.6	0.9375	1.066666667
DNA-binding protein A OS=Mus musculus GN=Csda PE=1 SV=2	DBPA_MOUSE	39 kDa	1	0	0.9	0.9	1	1
MCG11326, isoform CRA_b OS=Mus musculus GN=Hnrnp3 PE=4 SV=1 D3YWT1_MOUSE (+1)	D3YWT1_MOUSE (+1)	35 kDa	1	7.199221	1	1	1	1