

Protein	Accession Number	Molecular Weight	Mann Whitney Test (Bonferroni Corrected)	h WT run 1	Pool WT run 2	hool WT run 3	WT 1	WT 2	WT 3	Δ43 1	Δ43 2	Δ43 3	A57G 1	A57G 2	A57G 3
Elongation factor 2 OS=Mus musculus GN=Eef2 PE=1 SV=2	EF2_MOUSE	95 kDa	< 0.0001	27	28	30	28	27	30	30	28	27	30	28	27
Complement C3 OS=Mus musculus GN=C3 PE=1 SV=3	CO3_MOUSE	186 kDa	< 0.0001	32	30	35	30	32	35	35	30	32	35	30	32
Aldehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh2 PE=1 SV=1	ALDH2_MOUSE	57 kDa	< 0.0001	12	12	12	12	12	12	12	12	12	12	12	12
Heat shock protein HSP 90-beta OS=Mus musculus GN=Hsp90ab1 PE=1 SV=3	HS90B_MOUSE	83 kDa	< 0.0001	27	27	26	27	27	26	26	27	27	26	27	27
Leucine-rich PPR motif-containing protein, mitochondrial OS=Mus musculus GN=LrpprlPPRC_MOUSE		157 kDa	< 0.0001	21	25	22	25	21	22	22	25	21	22	25	21
NAD(P) transhydrogenase, mitochondrial OS=Mus musculus GN=Nnt PE=1 SV=2	NNTM_MOUSE	114 kDa	< 0.0001	50	67	66	67	50	66	66	67	50	66	67	50
Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial OS=Mus musculus GN=AL4A1_MOUSE		62 kDa	< 0.0001	12	14	19	14	12	19	19	14	12	19	14	12
Stress-induced-phosphoprotein 1 OS=Mus musculus GN=Stip1 PE=1 SV=1	STIP1_MOUSE	63 kDa	< 0.0001	13	20	19	20	13	19	19	20	13	19	20	13
Cluster of Troponin T, cardiac muscle OS=Mus musculus GN=Tnnt2 PE=4 SV=1 (K3W4R3W4R6_MOUSE [3])		35 kDa	0.00018	43	37	39	37	43	39	39	37	43	39	37	43
Mitochondrial 2-oxoglutarate/malate carrier protein OS=Mus musculus GN=Slc25a11 M2OM_MOUSE		34 kDa	0.0017	11	10	13	10	11	13	10	11	13	10	11	13
Alpha-1B-glycoprotein OS=Mus musculus GN=A1bg PE=1 SV=1	A1BG_MOUSE	57 kDa	0.0018	8	17	8	17	8	8	8	17	8	8	17	8
Elongation factor 1-gamma OS=Mus musculus GN=Eef1g PE=1 SV=3	EF1G_MOUSE	50 kDa	0.0025	11	9	8	9	11	8	8	9	11	8	9	11
Cluster of Myosin-6 OS=Mus musculus GN=Mylh6 PE=1 SV=2 (MYH6_MOUSE [10])	MYH6_MOUSE	224 kDa	0.0039	667	562	569	562	667	569	569	562	667	569	562	667
Succinyl-CoA ligase [GDP-forming] subunit beta, mitochondrial OS=Mus musculus GN=SUCB2_MOUSE		47 kDa	0.0068	18	19	18	19	18	18	19	18	18	18	19	18
Cluster of Inorganic pyrophosphatase 2, mitochondrial OS=Mus musculus GN=Ppa2 PED3Z636_MOUSE [2]		38 kDa	0.0069	9	13	15	13	9	15	15	13	9	15	13	9
Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondrial OS=Mus CHCH3_MOUSE		26 kDa	0.0084	14	12	15	12	14	15	15	12	14	15	12	14
Cluster of Myosin regulatory light chain 2, ventricular/cardiac muscle isoform OS=MusMLRV_MOUSE		19 kDa	0.011	58	53	52	53	58	52	52	53	58	52	53	58
Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3	CLH1_MOUSE	192 kDa	0.017	21	20	18	20	21	18	18	20	21	18	20	21
Histidine rich calcium binding protein, isoform CRA_a OS=Mus musculus GN=Hrc PE=4G5E8I6_MOUSE		85 kDa	0.017	16	61	49	61	16	49	49	61	16	49	61	16
Sarcalumenin OS=Mus musculus GN=Srl PE=1 SV=1	SRCA_MOUSE	99 kDa	0.02	34	27	33	27	34	33	27	34	33	27	34	33
Cytochrome c oxidase subunit 5B, mitochondrial OS=Mus musculus GN=Cox5b PE=1 SV=1 S COX5B_MOUSE		14 kDa	0.02	31	40	44	40	31	44	44	40	31	44	40	31
Serpin H1 OS=Mus musculus GN=Serpinh1 PE=1 SV=3	SERPH_MOUSE	47 kDa	0.021	9	13	10	13	9	10	10	13	9	10	13	9
Calcium-binding mitochondrial carrier protein Aralar1 OS=Mus musculus GN=Slc25a12CMC1_MOUSE		75 kDa	0.024	11	11	13	11	11	13	13	11	11	13	11	11
Short-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=AaCAD5_MOUSE		45 kDa	0.031	21	28	31	28	21	31	31	28	21	31	28	21
Heterogeneous nuclear ribonucleoprotein U OS=Mus musculus GN=Hnrnpu PE=4 SV=2 G3XA10_MOUSE (+1)		87 kDa	0.038	7	8	7	8	7	7	7	8	7	7	8	7
Cluster of long-chain-fatty-acid-CoA ligase 1 OS=Mus musculus GN=Acs1 PE=1 SV=2 ACSL1_MOUSE [4]		78 kDa	0.039	41	28	35	28	41	35	35	28	41	35	28	41
Citrate lyase subunit beta-like protein, mitochondrial OS=Mus musculus GN=Clybl PE=CLYBL_MOUSE		38 kDa	0.042	9	10	11	10	9	11	11	10	9	11	10	9
Cytochrome b-c1 complex subunit 7 OS=Mus musculus GN=Uqcrb PE=3 SV=1	Q9CQB4_MOUSE (+1)	14 kDa	0.046	10	16	14	16	10	14	14	16	10	14	16	10
Isovaleryl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Ivd PE=1 SV=1	IVD_MOUSE	46 kDa	0.048	23	30	30	30	23	30	30	23	30	30	23	30
NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial OS=Mus musculus GN=msu9QXP3_MOUSE (+1)		20 kDa	0.048	11	18	16	18	11	16	16	18	11	16	18	11
Protein Ogdh OS=Mus musculus GN=Ogdh1 PE=2 SV=1	E9Q7L0_MOUSE	117 kDa	0.052	3	11	11	11	3	11	11	3	11	3	11	3
Cluster of Creatine kinase S-type, mitochondrial OS=Mus musculus GN=Ckmt2 PE=1 SV=1 SKCRS_MOUSE		47 kDa	0.055	45	44	43	44	45	43	43	44	45	43	44	45
Nidogen-1 OS=Mus musculus GN=Nid1 PE=1 SV=2	NID1_MOUSE	137 kDa	0.056	11	11	10	11	11	10	10	11	11	10	11	11
Cluster of Aspartate aminotransferase, cytoplasmic OS=Mus musculus GN=Got1 PE=1 SV=1 AATC_MOUSE [2]		46 kDa	0.06	31	30	35	30	31	35	35	30	31	35	30	31
Cluster of 6-phosphofruktokinase, muscle type OS=Mus musculus GN=PfkM PE=1 SV=3K6PF_MOUSE		85 kDa	0.062	18	26	23	26	18	23	23	26	18	23	26	18
Cytoplasmic dynein 1 heavy chain 1 OS=Mus musculus GN=Dync1h1 PE=1 SV=2	DYHC1_MOUSE	532 kDa	0.067	18	26	24	26	18	24	24	26	18	24	26	18
Isocitrate dehydrogenase [NADP], mitochondrial OS=Mus musculus GN=ldh2 PE=1 SV=4DHP_MOUSE		51 kDa	0.068	48	49	50	49	48	50	49	48	50	49	48	50
Transgelin OS=Mus musculus GN=Tagln PE=1 SV=3	TAGL_MOUSE	23 kDa	0.08	6	11	9	11	6	9	9	11	6	9	11	6
Cluster of Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 (K1CK1C10_HUMAN [11])		60 kDa	0.09	20	13	11	13	20	11	11	13	20	11	13	20
Annexin A5 OS=Mus musculus GN=Anxa5 PE=1 SV=1	ANXA5_MOUSE	36 kDa	0.11	14	13	14	13	14	14	14	13	14	14	13	14
Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Mus musculus GN=Ppif PE=1 SV=1 SPPIF_MOUSE		22 kDa	0.12	8	6	8	6	8	8	8	6	8	8	6	8
Fumarate hydratase, mitochondrial OS=Mus musculus GN=Fh PE=1 SV=3	FUMH_MOUSE (+1)	54 kDa	0.13	26	34	37	34	26	37	37	34	26	37	34	26
L-lactate dehydrogenase B chain OS=Mus musculus GN=Ldhb PE=1 SV=2	LDHB_MOUSE	37 kDa	0.15	35	34	32	34	35	32	32	34	35	32	34	35
40S ribosomal protein S11 OS=Mus musculus GN=Rps11 PE=2 SV=3	RS11_MOUSE	18 kDa	0.15	4	5	7	5	4	7	7	5	4	7	5	4
60S ribosomal protein L7 OS=Mus musculus GN=Rpl7 PE=2 SV=2	RL7_MOUSE	31 kDa	0.17	7	9	10	9	7	10	10	9	7	10	9	7
Dehydrogenase/reductase SDR family member 4 OS=Mus musculus GN=Dhrs4 PE=2 SV=2 S DHRSA_MOUSE		30 kDa	0.18	7	11	12	11	7	12	12	11	7	12	11	7
Profilin-1 OS=Mus musculus GN=Pfn1 PE=1 SV=2	PROF1_MOUSE	15 kDa	0.18	7	8	10	8	7	10	10	8	7	10	8	7
BAG family molecular chaperone regulator 3 OS=Mus musculus GN=Bag3 PE=1 SV=2	BAG3_MOUSE	62 kDa	0.19	17	28	29	28	17	29	29	28	17	29	28	17
Ryanodine receptor 2 OS=Mus musculus GN=Ryr2 PE=1 SV=1	RYR2_MOUSE	565 kDa	0.2	26	28	41	28	26	41	41	28	26	41	28	26
Lamin-B1 OS=Mus musculus GN=Lmnb1 PE=1 SV=3	LMNB1_MOUSE	67 kDa	0.22	10	15	17	15	10	17	17	15	10	17	15	10
Histone H4 OS=Mus musculus GN=Hist1h4a PE=1 SV=2	H4_MOUSE	11 kDa	0.25	17	14	15	14	17	15	15	14	17	15	14	17
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitochondrial OS=NDUA9_MOUSE		43 kDa	0.29	19	20	20	20	19	20	20	20	19	20	20	19
Voltage-dependent anion-selective channel protein 1 OS=Mus musculus GN=Vdac1 PEVdac1_MOUSE		32 kDa	0.33	22	19	20	19	22	20	19	22	20	19	22	20
SET and MYND domain-containing protein 1 OS=Mus musculus GN=Smyd1 PE=2 SV=1 G5E8R7_MOUSE (+1)		53 kDa	0.37	11	14	12	14	11	12	12	14	11	12	14	11
ATP synthase subunit alpha, mitochondrial OS=Mus musculus GN=Atp5a1 PE=1 SV=1	ATPA_MOUSE	60 kDa	0.38	82	74	73	74	82	73	73	74	82	73	74	82
Cluster of cAMP-dependent protein kinase type I-alpha regulatory subunit OS=Mus m KAPO_MOUSE		43 kDa	0.53	10	8	7	8	10	7	7	8	10	7	8	10
NADP-dependent malic enzyme, mitochondrial OS=Mus musculus GN=Me3 PE=1 SV=2MAON_MOUSE		67 kDa	0.55	5	9	7	9	5	7	7	9	5	7	9	5
Carnitine O-acetyltransferase OS=Mus musculus GN=Crat PE=1 SV=3	CACP_MOUSE (+1)	71 kDa	0.62	23	29	27	29	23	27	27	29	23	27	29	23
ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b PE=1 SV=2	ATPB_MOUSE	56 kDa	0.68	72	71	66	71	72	66	66	71	72	66	71	72
60S ribosomal protein L34 OS=Mus musculus GN=Rpl34 PE=3 SV=2	RL34_MOUSE	13 kDa	0.71	5	5	4	5	5	4	4	5	5	4	5	5
Hsc70-interacting protein OS=Mus musculus GN=St13 PE=2 SV=1	F10A1_MOUSE	42 kDa	0.72	7	10	9	10	7	9	9	10	7	9	10	7
Protein Ahnak OS=Mus musculus GN=Ahnak PE=2 SV=1	E9Q616_MOUSE	604 kDa	0.96	35	49	46	49	35	46	46	49	35	46	49	35
Cluster of isoform 2 of Titin OS=Mus musculus GN=Ttn (sp) [A2ASS6-2] TITIN_MOUSE	sp [A2ASS6-2] TITIN_MOUSE?		1	997	1169	1237	1169	997	1237	1237	1169	997	1237	1169	997
Cluster of Actin, alpha cardiac muscle 1 OS=Mus musculus GN=Actc1 PE=1 SV=1 (ACTCACTC_MOUSE [5])		42 kDa	1	87	90	88	90	87	88	88	90	87	88	90	87

Cluster of Serum albumin OS=Mus musculus GN=Alb PE=1 SV=3 (ALBU_MOUSE)	ALBU_MOUSE [2]	69 kDa	1	130	132	123	132	130	123	123	132	130	123	132	130
Cluster of Tropomyosin 1, alpha, isoform CRA_c OS=Mus musculus GN=Tpm1 PE=2 SV Q8BP43_MOUSE [10]	Q8BP43_MOUSE [10]	33 kDa	1	137	118	115	118	137	115	115	118	137	115	118	137
Cluster of Beta-globin OS=Mus musculus GN=Hbb-b1 PE=3 SV=1 (ABDUK4_MOUSE)	ABDUK4_MOUSE [4]	16 kDa	1	38	43	44	43	38	44	44	43	38	44	43	38
Cluster of Myoglobin OS=Mus musculus GN=Mb PE=1 SV=3 (MYG_MOUSE)	MYG_MOUSE	17 kDa	1	42	47	44	47	42	44	44	47	42	44	47	42
Cluster of Hemoglobin subunit alpha OS=Mus musculus GN=Hba PE=1 SV=2 (HBA_MCHBA_MOUSE [3])	HBA_MCHBA_MOUSE [3]	15 kDa	1	20	26	26	26	20	26	26	26	20	26	26	20
Aconitate hydratase, mitochondrial OS=Mus musculus GN=Aco2 PE=1 SV=1	ACON_MOUSE	85 kDa	1	84	91	92	91	84	92	92	91	84	92	91	84
Myosin-binding protein C, cardiac-type OS=Mus musculus GN=Mybpc3 PE=2 SV=1	E9Q9T8_MOUSE (+1)	141 kDa	1	106	128	118	128	106	118	118	128	106	118	128	106
Cluster of Isoform 2 of Sarcolemmal/endoplasmic reticulum calcium ATPase 2 OS=Mu sp O52143-2 AT2A2_MOUSE ?	O52143-2_MOUSE ?	1	99	103	109	103	99	109	109	103	99	109	103	99	103
Cluster of Myosin light chain 3 OS=Mus musculus GN=MyI3 PE=1 SV=4 (MYL3_MOUSE MYL3_MOUSE [3])	MYL3_MOUSE [3]	22 kDa	1	41	39	36	39	41	36	36	39	41	36	39	41
Cluster of Fatty acid-binding protein, heart OS=Mus musculus GN=Fabp3 PE=1 SV=5 (FABPH_MOUSE [3])	FABPH_MOUSE [3]	15 kDa	1	37	41	38	41	37	38	38	41	37	38	41	37
Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE=1 SV=3	MDHM_MOUSE	36 kDa	1	45	59	54	59	45	54	54	59	45	54	59	45
Cluster of Alpha-actinin-2 OS=Mus musculus GN=Actn2 PE=1 SV=2 (ACTN2_MOUSE)	ACTN2_MOUSE [5]	104 kDa	1	83	92	88	92	83	88	88	92	83	88	92	83
Myomesin-1 OS=Mus musculus GN=Myom1 PE=1 SV=2	MYOM1_MOUSE	185 kDa	1	85	85	88	85	85	88	88	85	85	88	85	85
Myomesin 2 OS=Mus musculus GN=Myom2 PE=2 SV=1	Q14BI5_MOUSE	165 kDa	1	77	87	96	87	77	96	96	87	77	96	87	77
Very long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=ACADV_MOUSE	ACADV_MOUSE	71 kDa	1	66	70	73	70	66	73	73	70	66	73	70	66
Trifunctional enzyme subunit alpha, mitochondrial OS=Mus musculus GN=Hadha PE=:ECHA_MOUSE	ECHA_MOUSE	83 kDa	1	55	50	44	50	55	44	44	50	55	44	50	55
Cluster of ADP/ATP translocase 1 OS=Mus musculus GN=Slc25a4 PE=1 SV=4 (ADT1_MOUSE ADT1_MOUSE [2])	ADT1_MOUSE [2]	33 kDa	1	40	35	40	35	40	40	40	35	40	40	35	40
Cluster of Creatine kinase M-type OS=Mus musculus GN=Ckm PE=1 SV=1 (KCRM_MOUSE KCRM_MOUSE [2])	KCRM_MOUSE [2]	43 kDa	1	49	42	34	42	49	34	34	42	49	34	42	49
Long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Ac ACADL_MOUSE	ACADL_MOUSE	48 kDa	1	50	56	58	56	50	58	58	56	50	58	56	50
Cluster of Glycogen phosphorylase, muscle form OS=Mus musculus GN=Pygm PE=1 SV PYGM_MOUSE [3]	PYGM_MOUSE [3]	97 kDa	1	67	71	73	71	67	73	73	71	67	73	71	67
Cluster of Glyceraldehyde-3-phosphate dehydrogenase OS=Mus musculus GN=Gm50f D3Z029_MOUSE [4]	D3Z029_MOUSE [4]	36 kDa	1	27	33	34	33	27	34	34	33	27	34	33	27
Malate dehydrogenase, cytoplasmic OS=Mus musculus GN=Mdh1 PE=1 SV=3	MDHC_MOUSE	37 kDa	1	33	43	41	43	33	41	41	43	33	41	43	33
Desmin OS=Mus musculus GN=Des PE=1 SV=3	DESM_MOUSE	53 kDa	1	53	53	50	53	53	50	50	53	53	50	53	53
NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial OS=Mus musculus GN=NDU5_MOUSE	NDU5_MOUSE	80 kDa	1	50	57	63	57	50	63	63	57	50	63	57	50
Desmoplakin OS=Mus musculus GN=Dsp PE=2 SV=1	DESP_MOUSE	333 kDa	1	66	78	70	78	66	70	70	78	66	70	78	66
Cluster of Citrate synthase, mitochondrial OS=Mus musculus GN=Cs PE=1 SV=1 (CISY_MOUSE CISY_MOUSE [2])	CISY_MOUSE [2]	52 kDa	1	29	38	37	38	29	37	37	38	29	37	38	29
3-oxoacyl-CoA thiolase, mitochondrial OS=Mus musculus GN=Acaa2 PE=1 SV=3	THIM_MOUSE	42 kDa	1	52	38	44	38	52	44	44	38	52	44	38	52
Cluster of Plectin OS=Mus musculus GN=Plec PE=1 SV=2 (PLEC_MOUSE)	PLEC_MOUSE [2]	534 kDa	1	73	83	90	83	73	90	90	83	73	90	83	73
Cluster of Serotransferrin OS=Mus musculus GN=Tf PE=1 SV=1 (TRFE_MOUSE)	TRFE_MOUSE [4]	77 kDa	1	53	66	65	66	53	65	65	66	53	65	66	53
Electron transfer flavoprotein subunit alpha, mitochondrial OS=Mus musculus GN=Etff ETFA_MOUSE	ETF_A_MOUSE	35 kDa	1	33	30	32	30	33	32	32	30	33	32	30	33
Troponin I, cardiac muscle OS=Mus musculus GN=Tnni3 PE=1 SV=2	TNNI3_MOUSE	24 kDa	1	33	35	35	35	33	35	35	33	35	35	33	33
Cluster of Fructose-bisphosphate aldolase OS=Mus musculus GN=Aldoa PE=2 SV=1 (A1 AGZ144_MOUSE [3])	AGZ144_MOUSE [3]	45 kDa	1	47	44	50	44	47	50	50	44	47	50	44	47
Cluster of Heat shock cognate 71 kDa protein OS=Mus musculus GN=Hspa8 PE=1 SV=:HSP7C_MOUSE [5]	HSP7C_MOUSE [5]	71 kDa	1	40	53	53	53	40	53	53	53	40	53	53	40
Trifunctional enzyme subunit beta, mitochondrial OS=Mus musculus GN=Hadhb PE=1 ECHB_MOUSE	ECHB_MOUSE	51 kDa	1	37	39	38	39	37	38	38	39	37	38	39	37
Cluster of Spectrin alpha chain, non-erythrocytic 1 OS=Mus musculus GN=Sptan1 PE=:E9Q447_MOUSE [2]	E9Q447_MOUSE [2]	285 kDa	1	76	74	71	74	76	71	71	74	76	71	74	76
Cluster of Obscurin OS=Mus musculus GN=Obscn PE=2 (OBSCN_MOUSE)	OBSCN_MOUSE [3]	966 kDa	1	47	81	75	81	47	75	75	81	47	75	81	47
Aspartate aminotransferase, mitochondrial OS=Mus musculus GN=Got2 PE=1 SV=1	AATM_MOUSE	47 kDa	1	50	37	43	37	50	43	43	37	50	43	37	50
Stress-70 protein, mitochondrial OS=Mus musculus GN=Hspa9 PE=1 SV=3	GRP75_MOUSE	73 kDa	1	49	50	51	50	49	51	51	50	49	51	50	49
Cluster of Mitochondrial inner membrane protein OS=Mus musculus GN=Immt PE=2 :E9Q800_MOUSE [3]	E9Q800_MOUSE [3]	76 kDa	1	52	56	61	56	52	61	61	56	52	61	56	52
Cluster of Nebulette OS=Mus musculus GN=Neb1 PE=2 SV=1 (B7ZCI2_MOUSE)	B7ZCI2_MOUSE [3]	116 kDa	1	47	73	68	73	47	68	68	73	47	68	73	47
Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Mus r DHSA_MOUSE	DHSA_MOUSE	73 kDa	1	42	50	54	50	42	54	54	50	42	54	50	42
Medium-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN ACADM_MOUSE	ACADM_MOUSE	46 kDa	1	30	36	34	36	30	34	34	36	30	34	36	30
Cytochrome b-c1 complex subunit 2, mitochondrial OS=Mus musculus GN=Uqcrc2 PE: QCR2_MOUSE	QCR2_MOUSE	48 kDa	1	35	51	55	51	35	55	55	51	35	55	51	35
Cluster of 14-3-3 protein gamma OS=Mus musculus GN=Yvhag PE=1 SV=2 (1433G_MI1433G_MOUSE [9])	1433G_MOUSE [9]	28 kDa	1	56	58	63	58	56	63	63	58	56	63	58	56
Cluster of Beta-enolase OS=Mus musculus GN=Eno3 PE=1 SV=3 (ENOB_MOUSE)	ENOB_MOUSE [3]	47 kDa	1	38	33	28	33	38	28	28	33	38	28	33	38
Cluster of Phosphoglycerate kinase 1 OS=Mus musculus GN=Pgk1 PE=1 SV=4 (PGK1_A_PGK1_MOUSE [2])	PGK1_MOUSE [2]	45 kDa	1	45	35	40	35	45	40	40	35	45	40	35	45
Cluster of Vimentin OS=Mus musculus GN=Vim PE=1 SV=3 (VIME_MOUSE)	VIME_MOUSE [19]	54 kDa	1	55	64	51	64	55	51	51	64	55	51	64	55
Cluster of Isoform M1 of Pyruvate kinase PKM OS=Mus musculus GN=Pkm (sp) P52480-2 KPYM_MOUSE ?	P52480-2_MOUSE ?	1	48	39	38	39	38	48	38	38	39	48	38	39	48
Iso citrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Mus musculus GN=IDH3A_MOUSE	IDH3A_MOUSE	40 kDa	1	34	34	36	34	34	36	36	34	34	36	34	34
Electron transfer flavoprotein subunit beta OS=Mus musculus GN=Etff PE=1 SV=3	ETFB_MOUSE	28 kDa	1	32	28	28	28	32	28	28	28	32	28	28	32
Acetyl-CoA acetyltransferase, mitochondrial OS=Mus musculus GN=Acac1 PE=1 SV=1	THIL_MOUSE	45 kDa	1	38	53	44	53	38	44	44	53	38	44	53	38
60 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspd1 PE=1 SV=1	CH60_MOUSE	61 kDa	1	48	37	37	37	48	37	37	37	48	37	37	48
Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial OS=Mus mu ETFD_MOUSE	ETFD_MOUSE	68 kDa	1	37	44	53	44	37	53	53	44	37	53	44	37
Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mitochondrial CODPA_MOUSE	CODPA_MOUSE	43 kDa	1	37	36	41	36	37	41	41	36	37	41	36	37
Vinculin OS=Mus musculus GN=Vcl PE=1 SV=4	VINC_MOUSE	117 kDa	1	50	59	57	59	50	57	57	59	50	57	59	50
Cluster of Cytochrome c, somatic OS=Mus musculus GN=Cycs PE=1 SV=2 (CYC_MOUSE CYC_MOUSE [2])	CYC_MOUSE [2]	12 kDa	1	17	15	22	15	17	22	22	15	17	22	15	17
ATP synthase subunit gamma OS=Mus musculus GN=Atp5c1 PE=3 SV=1	A2AKU9_MOUSE	33 kDa	1	25	33	29	33	25	29	29	33	25	29	33	25
Cluster of Sodium/potassium-transporting ATPase subunit alpha-1 OS=Mus musculus AT1A1_MOUSE [6]	AT1A1_MOUSE [6]	113 kDa	1	38	44	52	44	38	52	52	44	38	52	44	38
Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial OS=Mus musculus GN=SUCB1_MOUSE	SUCB1_MOUSE	50 kDa	1	35	35	35	35	35	35	35	35	35	35	35	35
Cytochrome b-c1 complex subunit 1, mitochondrial OS=Mus musculus GN=Uqcrc1 PE: QCR1_MOUSE	QCR1_MOUSE	53 kDa	1	36	33	35	33	36	35	35	33	36	35	33	36
Cluster of Histone H2B type 1-H OS=Mus musculus GN=Hist1h2bh PE=1 SV=3 (H2B1H_MOUSE H2B1H_MOUSE [6])	H2B1H_MOUSE [6]	14 kDa	1	13	16	11	16	13	11	11	16	13	11	16	13
Cluster of LIM domain-binding protein 3 OS=Mus musculus GN=Ldb3 PE=1 SV=1 (LDB:LDB3_MOUSE [2])	LDB3_MOUSE [2]	76 kDa	1	30	39	39	39	30	39	39	39	30	39	39	30
Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Mus musculus ODPB_MOUSE	ODPB_MOUSE	39 kDa	1	26	30	33	30	26	33	33	30	26	33	30	26
Troponin C, slow skeletal and cardiac muscles OS=Mus musculus GN=Tnnc1 PE=2 SV=: TNNC1_MOUSE	TNNC1_MOUSE	18 kDa	1	23	17	17	23	17	17	17	23	17	17	23	17
Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Mus ml DHSB_MOUSE	DHSB_MOUSE	32 kDa	1	26	32	29	32	26	29	29	32	26	29	32	26
Cluster of Phosphoglycerate mutase 2 OS=Mus musculus GN=Pgam2 PE=1 SV=3 (PGAI_PGAM2_MOUSE [2])	PGAM2_MOUSE [2]	29 kDa	1	32	41	41	41	32	41	41	41	32	41	41	32

2-oxoglutarate dehydrogenase, mitochondrial OS=Mus musculus GN=Ogdh PE=1 SV=:	ODO1_MOUSE	116 kDa	1	36	41	50	41	36	50	50	41	36	50	41	36
Dihydropyridol dehydrogenase, mitochondrial OS=Mus musculus GN=Dld PE=1 SV=2	DLDH_MOUSE	54 kDa	1	25	32	34	32	25	34	34	32	25	34	32	25
Cytochrome c oxidase subunit 5A, mitochondrial OS=Mus musculus GN=Cox5a PE=1 S	COX5A_MOUSE	16 kDa	1	21	23	22	23	22	23	22	23	21	22	23	21
Enoyl-CoA hydratase, mitochondrial OS=Mus musculus GN=Echs1 PE=1 SV=1	ECHM_MOUSE	31 kDa	1	24	29	26	29	24	26	26	29	24	26	29	24
Cluster of Annexin A6 OS=Mus musculus GN=Anxa6 PE=1 SV=3 (ANXA6_MOUSE)	ANXA6_MOUSE [2]	76 kDa	1	42	35	40	35	42	40	40	35	42	40	35	42
Cluster of Prelamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2 (LMNA_MOUSE)	LMNA_MOUSE [2]	74 kDa	1	34	43	45	43	34	45	45	43	34	45	43	34
MCG10343, isoform CRA_b OS=Mus musculus GN=Slc25a3 PE=3 SV=1	GSE902_MOUSE	40 kDa	1	19	23	27	23	19	27	27	23	19	27	23	19
Cluster of NADH dehydrogenase [ubiquinone] flavoprotein 1, mitochondrial OS=Mus	NDUVU1_MOUSE [2]	51 kDa	1	26	25	20	25	26	20	20	25	26	20	25	26
Apolipoprotein A-I OS=Mus musculus GN=Apoa1 PE=1 SV=2	APOA1_MOUSE	31 kDa	1	33	25	32	25	33	32	32	25	33	32	25	33
Elongation factor Tu OS=Mus musculus GN=Gm9755 PE=3 SV=1	D3YVN7_MOUSE (+1)	50 kDa	1	32	36	37	36	32	37	37	36	32	37	36	32
Triosephosphate isomerase OS=Mus musculus GN=Tpi1 PE=1 SV=4	TPIS_MOUSE	32 kDa	1	17	32	27	32	17	27	27	32	17	27	32	17
ATP synthase-coupling factor 6, mitochondrial OS=Mus musculus GN=Atp5j PE=1 SV=:	ATP5J_MOUSE	12 kDa	1	18	17	16	17	18	16	16	17	18	16	17	18
Spectrin beta chain, non-erythrocytic 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2	SPTB2_MOUSE	274 kDa	1	34	48	44	48	34	44	44	48	34	44	48	34
Cluster of Histone H1.4 OS=Mus musculus GN=Hist1h1e PE=1 SV=2 (H14_MOUSE)	H14_MOUSE [3]	22 kDa	1	21	27	24	27	21	24	24	27	21	24	27	21
Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Mus musculus GN=Hadh	HCDD_MOUSE	34 kDa	1	20	21	22	21	20	22	22	21	20	22	21	20
Nascent polypeptide-associated complex subunit alpha, muscle-specific form OS=Mus	NACAM_MOUSE	220 kDa	1	19	37	46	37	19	46	46	37	19	46	37	19
Cluster of Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2	GSTM1_MOUSE [5]	26 kDa	1	30	41	36	41	30	36	36	41	30	36	41	30
Trypsin OS=Sus scrofa PE=1 SV=1	TRYP_PIG	24 kDa	1	14	13	13	13	14	13	13	13	14	13	13	14
Cluster of Alpha-1-antitrypsin 1-1 OS=Mus musculus GN=Serpina1a PE=1 SV=4 (A1AT1	A1AT1_MOUSE [4]	46 kDa	1	28	30	27	30	28	27	27	30	28	27	30	28
Alpha-crystallin B chain OS=Mus musculus GN=Cryab PE=1 SV=2	CRYAB_MOUSE	20 kDa	1	21	18	17	18	21	17	17	18	21	17	18	21
Isoform Cytoplasmic-peroxisomal of Peroxiredoxin-5, mitochondrial OS=Mus muscul	sp P99029-2 PRDX5_MOUSE?		1	20	13	19	13	20	19	19	13	20	19	13	20
Cluster of Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Mus musculu	COX4I1_MOUSE [2]	20 kDa	1	25	17	19	17	25	19	19	17	25	19	17	25
Iso citrate dehydrogenase 3 (NAD+) beta OS=Mus musculus GN=ldh3b PE=2 SV=1	Q91VA7_MOUSE	42 kDa	1	19	31	29	31	19	29	29	31	19	29	31	19
Dihydropyridylsine-residue acetyltransferase component of pyruvate dehydrogenase	ODP2_MOUSE	68 kDa	1	21	27	26	27	21	26	26	27	21	26	27	21
Cluster of ATP synthase subunit O, mitochondrial OS=Mus musculus GN=Atp5o PE=1	ATPO_MOUSE	23 kDa	1	17	18	17	18	17	17	17	18	17	17	18	17
Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondrial OS=Mus musculus	(SCOT1_MOUSE	56 kDa	1	17	27	27	27	17	27	27	27	17	27	27	17
10 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspe1 PE=1 SV=2	CH10_MOUSE	11 kDa	1	12	15	16	15	12	16	16	15	12	16	15	12
Cluster of L-lactate dehydrogenase OS=Mus musculus GN=Ldha PE=3 SV=1 (GSE8N5_	JGSE8N5_MOUSE [3]	40 kDa	1	21	19	20	19	21	20	20	19	21	20	19	21
Superoxide dismutase [Mn], mitochondrial OS=Mus musculus GN=Sod2 PE=1 SV=3	SODM_MOUSE	25 kDa	1	16	20	19	20	16	19	19	20	16	19	20	16
Cluster of Tubulin beta-4B chain OS=Mus musculus GN=Tubb4b PE=1 SV=1 (TBB4B_M	TBB4B_MOUSE [4]	50 kDa	1	20	32	32	32	20	32	32	32	20	32	32	20
Myozenin-2 OS=Mus musculus GN=Myoz2 PE=1 SV=1	MYOZ2_MOUSE	30 kDa	1	24	28	28	28	24	28	28	28	24	28	28	24
Succinyl-CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial OS=Mus muscul	SUCA_MOUSE	36 kDa	1	24	27	25	27	24	25	25	27	24	25	27	24
ATP synthase subunit b, mitochondrial OS=Mus musculus GN=Atp5f1 PE=1 SV=1	AT5F1_MOUSE	29 kDa	1	23	23	20	23	23	20	20	23	23	20	23	23
Cluster of Carnitine O-palmitoyltransferase 2, mitochondrial OS=Mus musculus GN=C	CPT2_MOUSE	74 kDa	1	21	33	30	33	21	30	30	33	21	30	33	21
Cytochrome c oxidase subunit 6B1 OS=Mus musculus GN=Cox6b1 PE=1 SV=2	CX6B1_MOUSE	10 kDa	1	11	11	8	11	11	8	8	11	11	8	11	11
Cluster of Basement membrane-specific heparan sulfate proteoglycan core protein O	B1B0C7_MOUSE [2]	469 kDa	1	28	51	42	51	28	42	42	51	28	42	51	28
Cluster of Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 PE=1 SV=1 (PRDX1_MOUSE)	PRDX1_MOUSE	22 kDa	1	19	20	19	20	19	19	19	20	19	19	20	19
Cluster of Nucleoside diphosphate kinase OS=Mus musculus GN=Gm20390 PE=2 SV=1	E9PZF0_MOUSE [3]	30 kDa	1	17	26	22	26	17	22	22	26	17	22	26	17
Enoyl-CoA delta isomerase 1, mitochondrial OS=Mus musculus GN=Eci1 PE=2 SV=2	ECI1_MOUSE	32 kDa	1	24	28	26	28	24	26	26	28	24	26	28	24
Cluster of Sorbin and SH3 domain-containing protein 2 OS=Mus musculus GN=Sorbs2	B2RX09_MOUSE [5]	72 kDa	1	23	27	35	27	23	35	35	27	23	35	27	23
Cluster of ATP synthase subunit d, mitochondrial OS=Mus musculus GN=Atp5h PE=1	SATP5H_MOUSE [2]	19 kDa	1	22	18	19	18	22	19	19	18	22	19	18	22
Cluster of Isoform 2 of Adenylate kinase isoenzyme 1 OS=Mus musculus GN=Ak1 (sp	sp Q9R0Y5-2 KAD1_MOUSE?		1	14	19	19	19	14	19	19	19	14	19	19	14
78 kDa glucose-regulated protein OS=Mus musculus GN=Hspa5 PE=1 SV=3	GRP78_MOUSE	72 kDa	1	30	27	27	27	30	27	27	27	30	27	27	30
Carnitine O-palmitoyltransferase 1, muscle isoform OS=Mus musculus GN=Cpt1b PE=:	CPT1B_MOUSE	88 kDa	1	20	23	26	23	20	26	26	23	20	26	23	20
Phosphoglucomutase-1 OS=Mus musculus GN=Pgm1 PE=1 SV=4	PGM1_MOUSE	61 kDa	1	20	27	26	27	20	26	26	27	20	26	27	20
Filamin-C OS=Mus musculus GN=Flnc PE=1 SV=3	FLNC_MOUSE	291 kDa	1	29	36	41	36	29	41	41	36	29	41	36	29
Dihydropyridylsine-residue succinyltransferase component of 2-oxoglutarate dehydr	ODO2_MOUSE	49 kDa	1	13	12	17	12	13	17	17	12	13	17	12	13
Thioredoxin-dependent peroxide reductase, mitochondrial OS=Mus musculus GN=Pr	PrC PRDX3_MOUSE	28 kDa	1	14	16	17	16	14	17	17	16	14	17	16	14
Transitional endoplasmic reticulum ATPase OS=Mus musculus GN=Vcp PE=1 SV=4	TERA_MOUSE	89 kDa	1	23	22	28	22	23	28	28	22	23	28	22	23
NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial OS=Mus mus	NDU52_MOUSE	53 kDa	1	19	20	25	20	19	25	25	20	19	25	20	19
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10, mitochondrial	OS=NDUAA_MOUSE	41 kDa	1	19	20	16	20	19	16	16	20	19	16	20	19
Alpha-2-macroglobulin OS=Mus musculus GN=A2m PE=1 SV=3	A2M_MOUSE (+1)	166 kDa	1	20	27	27	27	20	27	27	27	20	27	27	20
Cluster of Glucose-6-phosphate isomerase OS=Mus musculus GN=Gpi PE=1 SV=4 (G6P	G6PI_MOUSE [2]	63 kDa	1	16	17	23	17	16	23	23	17	16	23	17	16
2,4-dienoyl-CoA reductase, mitochondrial OS=Mus musculus GN=Decr1 PE=1 SV=1	DECR_MOUSE	36 kDa	1	12	13	13	13	12	13	13	13	12	13	13	12
Aldose reductase OS=Mus musculus GN=Akr1b1 PE=1 SV=3	ALDR_MOUSE	36 kDa	1	24	29	26	29	24	26	26	29	24	26	29	24
ES1 protein homolog, mitochondrial OS=Mus musculus GN=D10Jhu81e PE=1 SV=1	ES1_MOUSE	28 kDa	1	15	16	17	16	15	17	17	16	15	17	16	15
Cytochrome b-c1 complex subunit 6, mitochondrial OS=Mus musculus GN=Uqcqh PE=	QCR6_MOUSE	10 kDa	1	16	11	8	11	16	8	8	11	16	8	11	16
Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Mus musculus GN=Uqcrl	CRC1_MOUSE	29 kDa	1	15	14	17	14	15	17	17	14	15	17	14	15
Voltage-dependent anion-selective channel protein 3 OS=Mus musculus GN=Vdac3	P1J3QM3_MOUSE (+1)	31 kDa	1	12	15	17	15	12	17	17	15	12	17	15	12
Polymerase I and transcript release factor OS=Mus musculus GN=Ptrf PE=1 SV=1	PTRF_MOUSE	44 kDa	1	20	21	23	21	20	23	23	21	20	23	21	20
Cluster of Tubulin alpha-4A chain OS=Mus musculus GN=Tuba4a PE=1 SV=1 (TBA4A_A	TBA4A_MOUSE [5]	50 kDa	1	25	20	25	20	25	25	25	20	25	25	20	25
NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Mus musculus	NDUV2_MOUSE	27 kDa	1	14	20	21	20	14	21	21	20	14	21	20	14
Methylmalonate-semialdehyde dehydrogenase [acylating], mitochondrial OS=Mus	mtMMSA_MOUSE	58 kDa	1	20	21	27	21	20	27	27	21	20	27	21	20
Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Mus musculus GN=	Ec ECH1_MOUSE	36 kDa	1	21	20	19	20	21	19	19	20	21	19	20	21
Cysteine and glycine-rich protein 3 OS=Mus musculus GN=Csrp3 PE=1 SV=1	CSR3_MOUSE	21 kDa	1	18	22	18	22	18	18	18	22	18	18	22	18
Dystrophin OS=Mus musculus GN=Dmd PE=1 SV=3	DMD_MOUSE	426 kDa	1	20	22	19	22	20	19	19	22	20	19	22	20

Myosin regulatory light chain 2, atrial isoform OS=Mus musculus GN=Myl7 PE=2 SV=1 MLRA_MOUSE	19 kDa	1	32	18	23	18	23	18	32	23	23	18	32	23	18	32
Cluster of Ubiquitin-40S ribosomal protein S27a OS=Mus musculus GN=Rps27a PE=1 RS27A_MOUSE [2]	18 kDa	1	9	14	11	14	9	11	11	14	9	11	14	9	11	14
Cluster of Acyl-coenzyme A thioesterase 2, mitochondrial OS=Mus musculus GN=Acot ACOT2_MOUSE [3]	50 kDa	1	20	14	14	14	20	14	14	14	14	20	14	14	14	20
Cytochrome c1, heme protein, mitochondrial OS=Mus musculus GN=Cyc1 PE=1 SV=1 CY1_MOUSE	35 kDa	1	12	14	14	14	12	14	14	14	14	12	14	14	14	12
NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Mus musculus NDUS3_MOUSE	30 kDa	1	13	15	14	15	13	14	14	14	15	13	14	15	13	14
NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Mus musculus NDUS6_MOUSE	13 kDa	1	14	13	15	13	14	15	15	13	14	15	13	14	15	13
Cluster of Histone H3.1 OS=Mus musculus GN=Hist1h3a PE=1 SV=2 (H31_MOUSE) H31_MOUSE [4]	15 kDa	1	6	10	9	10	6	9	9	10	6	9	10	6	9	10
Cluster of PDZ and LIM domain protein 5 OS=Mus musculus GN=Pdlim5 PE=1 SV=4 (PDLIM5_MOUSE [3])	63 kDa	1	16	21	17	21	16	17	17	21	16	17	21	16	17	21
Protein NipSnap homolog 2 OS=Mus musculus GN=Gbas PE=2 SV=1	33 kDa	1	15	18	18	18	15	18	18	18	18	15	18	18	18	15
Cluster of Propionyl-CoA carboxylase alpha chain, mitochondrial OS=Mus musculus GIPCCA_MOUSE [2]	80 kDa	1	17	22	20	22	17	20	20	22	17	20	22	17	20	22
Cluster of Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3 (ENOA_MOUSE) ENOA_MOUSE [2]	47 kDa	1	19	14	16	14	19	16	16	14	19	16	14	19	16	14
Acyl carrier protein, mitochondrial OS=Mus musculus GN=Ndufab1 PE=1 SV=1	17 kDa	1	13	0	0	13	0	0	0	13	0	0	13	0	0	13
Elongation factor 1-alpha 2 OS=Mus musculus GN=Eef1a2 PE=1 SV=1	50 kDa	1	9	14	11	14	9	11	11	14	9	11	14	9	11	14
Peptidyl-prolyl cis-trans isomerase A OS=Mus musculus GN=Ppia PE=1 SV=2	18 kDa	1	10	9	10	9	10	10	10	9	10	10	9	10	10	9
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=Mus musculus NDUA7_MOUSE	13 kDa	1	12	14	18	14	12	18	18	14	12	18	14	12	18	14
Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Mus musculus GN=Hnrnpa2b1 ROA2_MOUSE	37 kDa	1	12	20	19	20	12	19	19	20	12	19	20	12	19	20
Ubiquinone biosynthesis protein COQ9, mitochondrial OS=Mus musculus GN=Coo9 PFCOQ9_MOUSE	35 kDa	1	13	21	15	21	13	15	15	21	13	15	21	13	15	21
NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Mus musculus NDUS8_MOUSE	24 kDa	1	10	13	12	13	10	12	12	13	10	12	13	10	12	13
Peroxisome oxidase OS=Mus musculus GN=Prdx6 PE=2 SV=1	22 kDa	1	12	12	17	12	12	17	17	12	12	17	12	12	17	12
Cluster of Talin-1 OS=Mus musculus GN=Tln1 PE=1 SV=2 (TLN1_MOUSE) TLN1_MOUSE	270 kDa	1	18	23	16	23	18	16	16	23	18	16	23	18	16	23
Myosin-9 OS=Mus musculus GN=Myh9 PE=1 SV=4	226 kDa	1	16	14	12	14	16	12	12	14	16	12	14	16	12	14
Cluster of Serine protease inhibitor A3K OS=Mus musculus GN=Serpina3k PE=1 SV=2 (SPA3K_MOUSE [2])	47 kDa	1	16	12	8	12	16	8	8	12	16	8	12	16	8	12
UTP--glucose-1-phosphate uridylyltransferase OS=Mus musculus GN=Ugp2 PE=2 SV=3 UGPA_MOUSE	57 kDa	1	12	15	19	15	12	19	19	15	12	19	15	12	19	15
Apoptosis-inducing factor 1, mitochondrial OS=Mus musculus GN=Aifm1 PE=1 SV=1	67 kDa	1	18	19	22	19	18	22	22	19	18	22	19	18	22	19
Cytochrome c oxidase subunit 6C OS=Mus musculus GN=Cox6c PE=1 SV=3	8 kDa	1	9	10	8	10	9	8	8	10	9	8	10	9	8	10
Four and a half LIM domains protein 2 OS=Mus musculus GN=Fhl2 PE=1 SV=1	32 kDa	1	14	22	21	22	14	21	21	22	14	21	22	14	21	22
Lon protease homolog, mitochondrial OS=Mus musculus GN=Lonp1 PE=1 SV=2	106 kDa	1	16	16	12	16	16	12	12	16	16	12	16	16	12	16
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=Mus musculus NDUA8_MOUSE	20 kDa	1	12	17	18	17	12	18	18	17	12	18	17	12	18	17
Cluster of Sorbin and SH3 domain-containing protein 1 OS=Mus musculus GN=Sorbs1 SRBS1_MOUSE [2]	143 kDa	1	17	20	19	20	17	19	19	20	17	19	20	17	19	20
Calsequestrin-2 OS=Mus musculus GN=Casq2 PE=2 SV=3	48 kDa	1	10	7	8	7	10	8	8	7	10	8	7	10	8	7
Isocitrate dehydrogenase [NAD] subunit gamma 1, mitochondrial OS=Mus musculus CIDHG1_MOUSE	43 kDa	1	16	11	14	11	16	14	14	11	16	14	11	16	14	11
Cluster of Actin-binding LIM protein 1 OS=Mus musculus GN=Ablim1 PE=2 SV=1 (EQ9:EQ9C1_MOUSE [2])	84 kDa	1	13	15	26	15	13	26	26	15	13	26	15	13	26	15
Laminin subunit gamma-1 OS=Mus musculus GN=Lamc1 PE=2 SV=1	177 kDa	1	15	22	19	22	15	19	19	22	15	19	22	15	19	22
Histone H1.5 OS=Mus musculus GN=Hist1h1b PE=1 SV=2	23 kDa	1	5	5	7	5	5	7	7	5	5	7	5	5	7	5
Prohibitin OS=Mus musculus GN=Phb PE=1 SV=1	30 kDa	1	16	15	15	15	16	15	15	16	15	15	16	15	15	16
Cytochrome b-c1 complex subunit 8 OS=Mus musculus GN=Uqcrcq PE=1 SV=3	10 kDa	1	7	12	10	12	7	10	12	7	10	12	7	10	12	7
Heat shock protein beta-1 OS=Mus musculus GN=Hspb1 PE=1 SV=3	23 kDa	1	15	17	11	17	15	11	11	17	15	11	17	15	11	17
Cluster of Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2 (SOD SODC_MOUSE)	16 kDa	1	9	24	21	24	9	21	21	24	9	21	24	9	21	24
Moesin OS=Mus musculus GN=Msn PE=1 SV=3	68 kDa	1	15	17	16	17	15	16	16	17	15	16	17	15	16	17
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=Mus musculus NDUA4_MOUSE	9 kDa	1	10	13	12	13	10	12	12	13	10	12	13	10	12	13
Cluster of 3-hydroxyisobutyryl-CoA hydrolase, mitochondrial OS=Mus musculus GN=HHIBCH_MOUSE [2]	43 kDa	1	16	20	16	20	16	16	16	20	16	16	20	16	16	20
Plakophilin 2 OS=Mus musculus GN=Pkp2 PE=2 SV=1	88 kDa	1	16	17	16	17	16	16	16	17	16	16	17	16	16	17
Cluster of Ribosome-binding protein 1 OS=Mus musculus GN=Rrbp1 PE=2 SV=1 (A2AV A2AVJ7_MOUSE [2])	158 kDa	1	6	15	16	15	6	16	16	15	6	16	15	6	16	15
Protein Trdn OS=Mus musculus GN=Trdn PE=2 SV=1	78 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0	0	4
Protein disulfide-isomerase A3 OS=Mus musculus GN=Pdia3 PE=1 SV=2	57 kDa	1	11	21	24	21	11	24	24	21	11	24	21	11	24	21
Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3	82 kDa	1	9	24	21	24	9	21	21	24	9	21	24	9	21	24
Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1	48 kDa	1	11	16	17	16	11	17	17	16	11	17	16	11	17	16
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 2 OS=Mus musculus NDUA2_MOUSE	11 kDa	1	9	11	13	11	9	13	13	11	9	13	11	9	13	11
Protein DJ-1 OS=Mus musculus GN=Park7 PE=1 SV=1	20 kDa	1	13	18	15	18	13	15	15	18	13	15	18	13	15	18
Pyruvate dehydrogenase protein X component, mitochondrial OS=Mus musculus GN=ODPX_MOUSE	54 kDa	1	10	9	12	9	10	12	12	9	10	12	9	10	12	9
Acetyl-coenzyme A synthetase 2-like, mitochondrial OS=Mus musculus GN=Acss1 PE= ACSS1_MOUSE	75 kDa	1	14	13	19	13	14	19	19	13	14	19	13	14	19	13
Cluster of EH domain-containing protein 4 OS=Mus musculus GN=Ehd4 PE=1 SV=1 (EH EHD4_MOUSE [2])	61 kDa	1	18	9	13	9	18	13	13	9	18	13	9	18	13	9
Hexokinase 1, isoform CRA_f OS=Mus musculus GN=Hk1 PE=3 SV=1	102 kDa	1	15	13	13	13	15	13	13	13	15	13	13	15	13	13
Cluster of Histone H2A.x OS=Mus musculus GN=H2afx PE=1 SV=2 (H2AX_MOUSE) H2AX_MOUSE [10]	15 kDa	1	3	8	6	8	3	6	6	8	3	6	8	3	6	8
Cluster of Ubiquitin-like modifier-activating enzyme 1 OS=Mus musculus GN=Uba1 PE UBA1_MOUSE [2]	118 kDa	1	14	18	19	18	14	19	18	14	19	18	14	19	18	14
Cytochrome c oxidase subunit 2 OS=Mus musculus GN=Mtco2 PE=1 SV=1	26 kDa	1	0	7	9	7	0	9	9	7	0	9	7	0	9	7
EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1	61 kDa	1	13	10	9	10	13	9	9	10	13	9	9	10	13	9
Voltage-dependent anion-selective channel protein 2 (Fragment) OS=Mus musculus G3UX26_MOUSE (+1)	30 kDa	1	12	16	16	16	12	16	16	16	12	16	16	16	12	16
ATP synthase subunit delta, mitochondrial OS=Mus musculus GN=Atp5d PE=1 SV=1	18 kDa	1	8	0	0	0	8	0	0	0	8	0	0	0	8	0
Hemopexin OS=Mus musculus GN=Hpx PE=1 SV=2	51 kDa	1	13	13	14	13	13	14	14	13	13	14	13	13	14	13
Cluster of NADH dehydrogenase [ubiquinone] flavoprotein 3, mitochondrial OS=Mus musculus NDUV3_MOUSE [2]	12 kDa	1	9	9	12	9	9	12	12	9	9	12	9	9	12	9
Cluster of Murinoglobulin-1 OS=Mus musculus GN=Mug1 PE=1 SV=3 (MUG1_MOUSE) MUG1_MOUSE [2]	165 kDa	1	8	8	9	8	8	9	8	8	9	8	8	9	8	8
Tripartite motif-containing protein 72 OS=Mus musculus GN=Trim72 PE=1 SV=1	53 kDa	1	12	11	13	12	11	13	12	11	13	12	11	13	12	11
Cluster of Protein Neb OS=Mus musculus GN=Neb PE=2 SV=1 (A2AQA9_MOUSE) A2AQA9_MOUSE [3]	801 kDa	1	4	0	0	0	4	0	0	0	4	0	0	0	4	0
Cluster of Hexokinase-2 OS=Mus musculus GN=Hk2 PE=1 SV=1 (HXK2_MOUSE) HXK2_MOUSE	103 kDa	1	14	18	15	18	14	15	15	18	14	15	18	14	15	18
Nucleolin OS=Mus musculus GN=Ncl PE=1 SV=2	77 kDa	1	13	11	11	11	13	11	11	11	13	11	11	11	11	13

Serum deprivation-response protein OS=Mus musculus GN=Sdpr PE=1 SV=3	SDPR_MOUSE	47 kDa	1	9	20	18	20	9	18	18	20	9	18	20	9
Cluster of Kininogen-1 OS=Mus musculus GN=Kng1 PE=1 SV=1 (KNG1_MOUSE)	KNG1_MOUSE [2]	73 kDa	1	11	16	14	16	11	14	14	16	11	14	16	11
Prohibitin-2 OS=Mus musculus GN=Phb2 PE=1 SV=1	PHB2_MOUSE	33 kDa	1	13	13	18	13	13	18	18	13	13	18	13	13
60S ribosomal protein L4 OS=Mus musculus GN=Rpl4 PE=1 SV=3	RL4_MOUSE	47 kDa	1	4	6	7	6	4	7	7	6	4	7	6	4
Cluster of Microtubule-associated protein (Fragment) OS=Mus musculus GN=Map4 PE=F7Ck47_MOUSE [2]	MAP4_MOUSE [2]	153 kDa	1	7	12	18	12	7	18	18	12	7	18	12	7
Cluster of Ras-related protein Rab-1B OS=Mus musculus GN=Rab1b PE=1 SV=1 (RAB1B_MOUSE) [7]	RAB1B_MOUSE [7]	22 kDa	1	12	15	7	15	12	7	7	15	12	7	15	12
Endoplasmic reticulum chaperone protein OS=Mus musculus GN=Hsp90b1 PE=1 SV=2	ENP1_MOUSE	92 kDa	1	9	14	17	14	9	17	17	14	9	17	14	9
Heat shock 70 kDa protein 4 OS=Mus musculus GN=Hspa4 PE=2 SV=1	Q3U2G2_MOUSE	94 kDa	1	9	16	14	16	9	14	14	16	9	14	16	9
Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=Tgm2 PE=1 SV=1 (TGM2_MOUSE)	TGM2_MOUSE	77 kDa	1	9	20	17	20	9	17	17	20	9	17	20	9
Cluster of Carboxylesterase 1C OS=Mus musculus GN=Ces1c PE=1 SV=4 (EST1C_MOUSE; EST1C_MOUSE [3])	EST1C_MOUSE [3]	61 kDa	1	12	16	13	16	12	13	13	16	12	13	16	12
NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Mus musculus GN=Ndufc2 PE=1 SV=1 (NDUC2_MOUSE)	NDUC2_MOUSE	14 kDa	1	9	11	12	11	9	12	12	11	9	12	11	9
Cluster of Laminin subunit alpha-2 OS=Mus musculus GN=Lama2 PE=1 SV=1 (LAMA2_MOUSE; LAMA2_MOUSE [2])	LAMA2_MOUSE [2]	343 kDa	1	8	14	13	14	8	13	13	14	8	13	14	8
Cluster of Myosin-11 OS=Mus musculus GN=Myh11 PE=1 SV=1 (MYH11_MOUSE)	MYH11_MOUSE [2]	227 kDa	1	13	13	10	13	10	10	10	13	10	13	10	13
Cluster of Selenium-binding protein 1 OS=Mus musculus GN=Selenbp1 PE=1 SV=2 (SB_SBP1_MOUSE [2])	SB_SBP1_MOUSE [2]	53 kDa	1	13	14	11	14	13	11	11	14	13	11	14	13
Cluster of Myosin light chain 4 (Fragment) OS=Mus musculus GN=Myl4 PE=2 SV=1 (A2A2A6Q8_MOUSE [3])	Myl4_MOUSE [3]	21 kDa	1	18	8	10	8	18	10	10	8	18	10	8	18
Heterogeneous nuclear ribonucleoprotein M OS=Mus musculus GN=Hnmpm PE=2 SV=2 (BNK3_MOUSE; BNK3_MOUSE (+3))	BNK3_MOUSE (+3)	72 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
Phosphatidylethanolamine-binding protein 1 OS=Mus musculus GN=Pebp1 PE=2 SV=1 (D3Z1V4_MOUSE (+1))	Pebp1_MOUSE (+1)	23 kDa	1	5	9	8	9	5	8	8	9	5	8	9	5
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=Mus musculus GN=Ndub6_MOUSE	NDUB6_MOUSE	15 kDa	1	8	7	10	7	8	10	10	7	8	10	7	8
Peroxisomal protein OS=Mus musculus GN=Prdx2 PE=1 SV=3	PRDX2_MOUSE	22 kDa	1	9	7	7	7	9	7	7	7	9	7	7	9
Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	ANXA2_MOUSE	39 kDa	1	13	9	14	9	13	14	14	9	13	14	9	13
Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	LEG1_MOUSE	15 kDa	1	10	15	11	15	10	11	11	15	10	11	15	10
Acyl-CoA dehydrogenase family member 9, mitochondrial OS=Mus musculus GN=Aca1 ACAD9_MOUSE	ACAD9_MOUSE	69 kDa	1	8	13	13	13	8	13	13	13	8	13	13	8
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Mus musculus GN=Ndub10_MOUSE	NDUB10_MOUSE	21 kDa	1	8	7	6	7	8	6	6	7	8	6	7	8
von Willebrand factor A domain-containing protein 8 OS=Mus musculus GN=Vwa8 PE=VWA8_MOUSE	VWA8_MOUSE	213 kDa	1	13	17	18	17	13	18	18	17	13	18	17	13
Cluster of Enoyl-CoA delta isomerase 2, mitochondrial OS=Mus musculus GN=Eci2 PE=ECI2_MOUSE [5]	ECI2_MOUSE [5]	43 kDa	1	7	7	6	7	6	6	6	7	7	6	7	7
Protein disulfide-isomerase OS=Mus musculus GN=P4hb PE=1 SV=2	PDI1_MOUSE	57 kDa	1	14	10	10	10	14	10	10	10	14	10	10	14
Cluster of Ankyrin-2 OS=Mus musculus GN=Ank2 PE=1 SV=2 (ANK2_MOUSE)	ANK2_MOUSE [4]	426 kDa	1	2	5	2	5	2	2	2	5	2	2	5	2
Perilipin-4 OS=Mus musculus GN=Plin4 PE=1 SV=2	PLIN4_MOUSE	139 kDa	1	6	18	16	18	6	16	16	18	6	16	18	6
Tropomodulin-1 OS=Mus musculus GN=Tmod1 PE=2 SV=2	TMOD1_MOUSE	40 kDa	1	12	9	12	9	12	12	12	9	12	12	9	12
Cluster of D-beta-hydroxybutyrate dehydrogenase, mitochondrial OS=Mus musculus GN=BDH_MOUSE [2]	BDH_MOUSE [2]	38 kDa	1	8	10	10	10	8	10	10	10	8	10	10	8
Creatine kinase B-type OS=Mus musculus GN=Ckb PE=1 SV=1	KCRB_MOUSE	43 kDa	1	7	6	8	6	7	8	8	6	7	8	6	7
Chaperone activity of bc1 complex-like, mitochondrial OS=Mus musculus GN=Adck3 PADC3_MOUSE	PADC3_MOUSE	72 kDa	1	8	15	12	15	8	12	12	15	8	12	15	8
Cluster of Protein Tns1 OS=Mus musculus GN=Tns1 PE=2 SV=1 (E9Q0S6_MOUSE)	E9Q0S6_MOUSE	201 kDa	1	11	12	14	12	11	14	14	12	11	14	12	11
Striated muscle-specific serine/threonine-protein kinase OS=Mus musculus GN=Speg E9Q2S5_MOUSE	E9Q2S5_MOUSE	354 kDa	1	8	10	9	10	8	9	9	10	8	9	10	8
60S ribosomal protein L6 OS=Mus musculus GN=Rpl6 PE=1 SV=3	RL6_MOUSE	34 kDa	1	7	14	12	14	7	12	12	14	7	12	14	7
Nesprin-2 OS=Mus musculus GN=Syne2 PE=1 SV=2	SYNE2_MOUSE	783 kDa	1	3	0	0	3	0	0	0	3	0	0	3	0
Cluster of Glutathione S-transferase omega-1 OS=Mus musculus GN=Gsto1 PE=2 SV=2 (GSTO1_MOUSE)	GSTO1_MOUSE	27 kDa	1	7	9	12	9	7	12	12	9	7	12	9	7
Cluster of Calpastatin OS=Mus musculus GN=Cast PE=1 SV=2 (ICAL_MOUSE)	ICAL_MOUSE [3]	85 kDa	1	10	13	19	13	10	19	19	13	10	19	13	10
Non-specific lipid-transfer protein OS=Mus musculus GN=Scp2 PE=1 SV=3	NLTP_MOUSE	59 kDa	1	8	11	8	11	8	8	11	8	8	11	8	11
Calcium-binding mitochondrial carrier protein Aralar2 OS=Mus musculus GN=Slc25a1; CMC2_MOUSE	CMC2_MOUSE	74 kDa	1	13	11	10	11	13	10	10	11	13	10	11	13
Cysteine-rich protein 2 OS=Mus musculus GN=Crip2 PE=1 SV=1	CRIP2_MOUSE	23 kDa	1	0	9	7	9	0	7	7	9	0	7	9	0
Uncharacterized protein OS=Mus musculus GN=Rps3a2 PE=3 SV=1	D3Z6C3_MOUSE	30 kDa	1	7	11	12	11	7	12	12	11	7	12	11	7
Glutamate dehydrogenase 1, mitochondrial OS=Mus musculus GN=Glud1 PE=1 SV=1	DHE1_MOUSE	61 kDa	1	11	16	14	16	11	14	14	16	11	14	16	11
Cluster of Catenin alpha-1 OS=Mus musculus GN=Cttna1 PE=1 SV=1 (CTNA1_MOUSE)	CTNA1_MOUSE	100 kDa	1	8	11	9	11	8	9	9	11	8	9	11	8
Nexilin OS=Mus musculus GN=Nexn PE=1 SV=3	NEXN_MOUSE	72 kDa	1	10	14	12	14	10	12	12	14	10	12	14	10
Cluster of Talin-2 OS=Mus musculus GN=Tln2 PE=4 SV=1 (E9PUM4_MOUSE)	E9PUM4_MOUSE [2]	272 kDa	1	8	10	9	10	8	9	9	10	8	9	10	8
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Mus musculus GN=Ndub13_MOUSE	NDUB13_MOUSE	17 kDa	1	10	7	9	7	10	9	9	7	10	9	7	10
Phosphoglucomutase-like protein 5 OS=Mus musculus GN=Pgm5 PE=1 SV=2	PGM5_MOUSE	62 kDa	1	7	9	11	9	7	11	11	9	7	11	9	7
Cluster of Dynamin-like 120 kDa protein, mitochondrial OS=Mus musculus GN=Opa1 F7Bx01_MOUSE [3]	F7Bx01_MOUSE [3]	113 kDa	1	16	14	17	14	16	17	17	14	16	17	14	16
Heat shock protein HSP 90-alpha OS=Mus musculus GN=Hsp90aa1 PE=1 SV=4	H90A_MOUSE	85 kDa	1	14	10	12	10	14	12	12	10	14	12	10	14
Protein Agl OS=Mus musculus GN=Agl PE=4 SV=1	F8VPN4_MOUSE	174 kDa	1	11	15	10	15	11	10	10	15	11	10	15	11
Plasminogen activator inhibitor 1 RNA-binding protein OS=Mus musculus GN=Serbp1 PAIRB_MOUSE	PAIRB_MOUSE	45 kDa	1	7	11	8	11	7	8	8	11	7	8	11	7
Propionyl-CoA carboxylase beta chain, mitochondrial OS=Mus musculus GN=Pccb PE=PCCB_MOUSE	PCCB_MOUSE	58 kDa	1	11	11	15	11	11	15	15	11	11	15	11	11
Laminin subunit beta-1 OS=Mus musculus GN=Lamb1 PE=2 SV=1	E9QN70_MOUSE	202 kDa	1	9	8	10	8	9	10	10	8	9	10	8	9
Lumican OS=Mus musculus GN=Lum PE=1 SV=2	LUM_MOUSE	38 kDa	1	7	10	15	10	7	15	15	10	7	15	10	7
Filamin, alpha OS=Mus musculus GN=Flna PE=4 SV=1	B7FAU9_MOUSE (+2)	280 kDa	1	10	18	12	18	10	12	12	18	10	12	18	10
Cluster of MCG5732 OS=Mus musculus GN=Gm4978 PE=4 SV=2 (F6VBB8_MOUSE)	F6VBB8_MOUSE [5]	30 kDa	1	3	9	9	9	3	9	9	9	3	9	9	3
Acyl-coenzyme A thioesterase 13 OS=Mus musculus GN=Acot13 PE=1 SV=1	ACOT13_MOUSE	15 kDa	1	6	AC013_MOUSE	7	6	6	7	7	6	6	7	6	6
Cluster of 40S ribosomal protein S4, X isoform OS=Mus musculus GN=Rps4x PE=2 SV=1	R54X_MOUSE [3]	30 kDa	1	5	6	7	6	5	7	7	6	5	7	6	5
Calmodulin OS=Mus musculus GN=Calm1 PE=1 SV=2	CALM_MOUSE (+1)	17 kDa	1	8	8	9	8	8	9	9	8	8	9	8	8
Protein Fga OS=Mus musculus GN=Fga PE=2 SV=1	E9PV24_MOUSE (+1)	87 kDa	1	8	12	12	12	8	12	12	12	8	12	12	8
Protein Col6a3 OS=Mus musculus GN=Col6a3 PE=4 SV=2	E9PWQ3_MOUSE	354 kDa	1	0	4	7	4	0	7	7	4	0	7	4	0
Histone H1.0 OS=Mus musculus GN=H1f0 PE=2 SV=4	H10_MOUSE	21 kDa	1	7	5	7	5	7	7	7	5	7	5	7	5
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 9 OS=Mus musculus GN=Ndub9_MOUSE	NDUB9_MOUSE	22 kDa	1	0	6	8	6	0	8	8	6	0	8	6	0
Heat shock protein beta-6 OS=Mus musculus GN=Hspb6 PE=1 SV=1	HSPB6_MOUSE	18 kDa	1	11	10	12	10	11	12	12	10	11	12	10	11
Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondrial OS=Mus musculus GN=MCCA_MOUSE	MCCA_MOUSE	79 kDa	1	9	11	17	11	9	17	17	11	9	17	11	9

NADH dehydrogenase [ubiquinone] iron-sulfur protein 5 OS=Mus musculus GN=Ndufs NDUS5_MOUSE	13 kDa	1	7	5	7	5	7	7	7	5	7	7	5	7
ATP synthase subunit epsilon, mitochondrial OS=Mus musculus GN=Atp5e PE=2 SV=2 ATP5E_MOUSE	6 kDa	1	4	8	7	8	4	7	7	8	4	7	8	4
Platelet glycoprotein 4 OS=Mus musculus GN=Cd36 PE=1 SV=2 CD36_MOUSE	53 kDa	1	6	5	5	5	6	5	5	5	6	5	5	6
Ig kappa chain C region OS=Mus musculus PE=1 SV=1 IGKC_MOUSE	12 kDa	1	6	9	8	9	6	8	8	9	6	8	9	6
Protein phosphatase 1 regulatory subunit 12B OS=Mus musculus GN=Ppp1r12b PE=1 MYPT2_MOUSE	109 kDa	1	7	10	12	10	7	12	12	10	7	12	10	7
Cluster of Protein Pabpc6 OS=Mus musculus GN=Pabpc6 PE=2 SV=1 (Q9D4E6_MOUSE Q9D4E6_MOUSE [4])	71 kDa	1	4	3	6	3	4	6	6	3	4	6	3	4
Alpha-2-HS-glycoprotein OS=Mus musculus GN=Ahsfg PE=1 SV=1 FETUA_MOUSE	37 kDa	1	8	9	11	9	8	11	11	9	8	11	9	8
3-hydroxyacyl-CoA dehydrogenase type-2 OS=Mus musculus GN=Hsd17b10 PE=2 SV=2 A2AFQ2_MOUSE (+1)	28 kDa	1	5	13	15	13	5	15	15	13	5	15	13	5
Sodium/potassium-transporting ATPase subunit beta-1 OS=Mus musculus GN=Atp1b1AT1B1_MOUSE	35 kDa	1	5	8	8	5	8	8	8	5	8	8	5	8
Cluster of Catalase OS=Mus musculus GN=Cat PE=1 SV=4 (CATA_MOUSE) CATA_MOUSE [2]	60 kDa	1	11	20	10	20	11	10	10	20	11	10	20	11
Cluster of Protein Gm10923 OS=Mus musculus GN=Gm10923 PE=3 SV=1 (D3YU46_M D3YU46_MOUSE [4])	17 kDa	1	8	9	10	9	8	10	10	9	8	10	9	8
High mobility group protein B1 OS=Mus musculus GN=Hmgb1 PE=1 SV=2 HMGB1_MOUSE	25 kDa	1	10	9	16	9	10	16	16	9	10	16	9	10
Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial OS=Mus musculus GN=MCCB_MOUSE	61 kDa	1	9	9	13	9	9	13	13	9	9	13	9	9
Protein NDRG2 OS=Mus musculus GN=Ndrg2 PE=1 SV=1 NDRG2_MOUSE	41 kDa	1	0	5	8	5	0	8	8	5	0	8	5	0
Cluster of NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 5, mitochondrion NDU5_MOUSE [2]	22 kDa	1	8	5	5	8	5	5	5	8	5	5	8	5
ATPase inhibitor, mitochondrial OS=Mus musculus GN=Atpif1 PE=2 SV=1 E9PV44_MOUSE	9 kDa	1	8	10	9	10	8	9	9	10	8	9	10	8
Cluster of Hmg2 protein OS=Mus musculus GN=Hmg2 PE=4 SV=1 (Q5XK38_MOUSE Q5XK38_MOUSE [2])	10 kDa	1	6	0	0	0	6	0	0	0	6	0	0	6
Cluster of Microtubule-associated protein tau OS=Mus musculus GN=Mapt PE=2 SV=1 A2A5Y6_MOUSE [4]	78 kDa	1	7	7	9	7	7	9	9	7	7	9	7	7
Carbonic anhydrase 2 OS=Mus musculus GN=Ca2 PE=1 SV=4 CAH2_MOUSE	29 kDa	1	9	6	10	6	9	10	10	6	9	10	6	9
Protein Golgb1 OS=Mus musculus GN=Golgb1 PE=2 SV=1 E9QAH1_MOUSE	365 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Adenosine kinase OS=Mus musculus GN=Adk PE=1 SV=2 ADK_MOUSE (+1)	40 kDa	1	7	7	9	7	7	9	9	7	7	9	7	7
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm5611 PE=4 SV=2 (E9Q5T E9Q5T3_MOUSE [2])	30 kDa	1	5	4	7	4	5	7	7	4	5	7	4	5
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 OS=Mus musculus NDUA5_MOUSE	13 kDa	1	7	11	7	11	7	7	7	11	7	7	11	7
Cluster of sp K1C9_HUMAN sp K1C9_HUMAN [2]	?	1	6	13	9	13	6	9	9	13	6	9	13	6
Natriuretic peptides A OS=Mus musculus GN=Nppa PE=1 SV=2 ANF_MOUSE	17 kDa	1	6	9	8	9	6	8	8	9	6	8	9	6
Cluster of Antithrombin-III OS=Mus musculus GN=Serpinc1 PE=1 SV=1 (ANT3_MOUSE ANT3_MOUSE)	52 kDa	1	7	9	10	9	7	10	10	9	7	10	9	7
ATP synthase subunit e, mitochondrial OS=Mus musculus GN=Atp5i PE=1 SV=2 ATP5I_MOUSE	8 kDa	1	0	6	5	6	0	5	5	6	0	5	6	0
14 kDa phosphohistidine phosphatase OS=Mus musculus GN=Phpt1 PE=1 SV=1 PPH14_MOUSE	14 kDa	1	9	10	7	10	9	7	7	10	9	7	10	9
Transaldolase OS=Mus musculus GN=Taldo1 PE=1 SV=2 TALDO_MOUSE	37 kDa	1	6	7	12	7	6	12	12	7	6	12	7	6
Cluster of Basigin OS=Mus musculus GN=Bsg PE=1 SV=2 (BASI_MOUSE) BASI_MOUSE [3]	42 kDa	1	5	9	12	9	5	12	12	9	5	12	9	5
Cluster of Kinetin OS=Mus musculus GN=Ktn1 PE=2 SV=1 (F8VQC7_MOUSE) F8VQC7_MOUSE [2]	153 kDa	1	6	4	8	4	6	8	8	4	6	8	4	6
Proteasome activator complex subunit 1 (Fragment) OS=Mus musculus GN=Psm1 PE G3UXZ5_MOUSE (+1)	27 kDa	1	7	9	10	9	7	10	10	9	7	10	9	7
60S ribosomal protein L3 OS=Mus musculus GN=Rpl3 PE=2 SV=3 RL3_MOUSE	46 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Cluster of Heterogeneous nuclear ribonucleoprotein K OS=Mus musculus GN=Hnrnpk B2M1R6_MOUSE [4]	49 kDa	1	10	12	11	12	10	11	11	12	10	11	12	10
Cofilin-1 OS=Mus musculus GN=Cf1 PE=1 SV=3 COF1_MOUSE (+1)	19 kDa	1	9	11	9	11	9	9	9	11	9	9	11	9
DNA polymerase theta OS=Mus musculus GN=Polq PE=2 SV=1 Q80XB7_MOUSE	281 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cluster of Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha 2AAA_MOUSE	65 kDa	1	7	3	8	3	7	8	8	3	7	8	3	7
Cluster of Methylglutaconyl-CoA hydratase, mitochondrial OS=Mus musculus GN=Auh AUHM_MOUSE [2]	33 kDa	1	5	10	8	10	5	8	8	10	5	8	10	5
Cluster of CAP-Gly domain-containing linker protein 1 (Fragment) OS=Mus musculus CF7CB97_MOUSE [5]	136 kDa	1	4	11	4	11	4	4	4	11	4	4	11	4
Cluster of Cathepsin D OS=Mus musculus GN=Ctsd PE=1 SV=1 (CATD_MOUSE) CATD_MOUSE [2]	45 kDa	1	5	9	11	9	5	11	11	9	5	11	9	5
WD repeat-containing protein 1 OS=Mus musculus GN=Wdr1 PE=1 SV=3 WDR1_MOUSE	66 kDa	1	9	9	9	9	9	9	9	9	9	9	9	9
Serine-rich coiled-coil domain-containing protein 1 OS=Mus musculus GN=Ccser1 PE=1 CSSE1_MOUSE	98 kDa	1	3	0	0	3	0	0	0	3	0	0	3	0
Dextrin OS=Mus musculus GN=Dstn PE=1 SV=3 DEST_MOUSE	19 kDa	1	5	5	8	5	5	8	8	5	5	8	5	5
Fibrous sheath-interacting protein 2 OS=Mus musculus GN=Fisp2 PE=1 SV=3 FISP2_MOUSE	785 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
GrpE protein homolog 1, mitochondrial OS=Mus musculus GN=Grpel1 PE=1 SV=1 GRPE1_MOUSE	24 kDa	1	8	9	6	9	8	6	6	9	8	6	9	8
O-acetyl-ADP-ribose deacetylase MACROD1 OS=Mus musculus GN=Macro1 PE=2 SV=1 MACD1_MOUSE	35 kDa	1	5	9	6	9	5	6	6	9	5	6	9	5
Isoleucine-tRNA ligase, mitochondrial OS=Mus musculus GN=lars2 PE=2 SV=1 SYIN_MOUSE	113 kDa	1	4	9	12	9	4	12	12	9	4	12	9	4
Beta-taxilin OS=Mus musculus GN=Txlnb PE=2 SV=2 TXLNB_MOUSE	77 kDa	1	6	10	10	10	6	10	10	10	6	10	10	6
AFG3-like protein 2 OS=Mus musculus GN=Afg3l2 PE=1 SV=1 AFG32_MOUSE	90 kDa	1	6	10	10	10	6	10	10	10	6	10	10	6
Cluster of MCG142036 OS=Mus musculus GN=Acad12 PE=3 SV=1 (D327X0_MOUSE) D327X0_MOUSE [2]	61 kDa	1	6	11	12	11	6	12	12	11	6	12	11	6
Dynamin-1-like protein OS=Mus musculus GN=Dnm1l PE=1 SV=2 DNM1L_MOUSE (+1)	83 kDa	1	0	10	8	10	0	8	8	10	0	8	10	0
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm18025 PE=3 SV=2 (E9Q1 E9Q1N8_MOUSE [5])	29 kDa	1	0	5	8	5	0	8	8	5	0	8	5	0
Protein Myh15 OS=Mus musculus GN=Myh15 PE=4 SV=1 EQQ264_MOUSE	222 kDa	1	3	1	2	1	3	2	2	1	3	2	1	3
Cluster of Gelsolin OS=Mus musculus GN=Gsn PE=1 SV=3 (GELS_MOUSE) GELS_MOUSE	86 kDa	1	6	9	8	9	6	8	8	9	6	8	9	6
Cluster of Glutathione S-transferase A4 OS=Mus musculus GN=Gsta4 PE=1 SV=3 (GST7 GSTA4_MOUSE [3])	26 kDa	1	5	9	7	9	5	7	7	9	5	7	9	5
Hydroxysteroid dehydrogenase-like protein 2 OS=Mus musculus GN=Hsd12 PE=2 SV=1 HSDL2_MOUSE	54 kDa	1	8	13	10	13	8	10	10	13	8	10	13	8
Adenylate kinase 2, mitochondrial OS=Mus musculus GN=Ak2 PE=1 SV=5 KAD2_MOUSE	26 kDa	1	9	6	5	6	9	5	5	6	9	5	6	9
Ubiquitin-conjugating enzyme E2 N OS=Mus musculus GN=Ube2n PE=1 SV=1 UBE2N_MOUSE	17 kDa	1	5	7	7	7	5	7	7	7	5	7	7	5
Lactoylglutathione lyase OS=Mus musculus GN=Glo1 PE=1 SV=3 LGLU_MOUSE	21 kDa	1	6	6	9	6	6	9	9	6	6	9	6	6
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Mus musculus GNDUB7_MOUSE	16 kDa	1	3	5	6	5	3	6	6	5	3	6	5	3
Cluster of Mitogen-activated protein kinase 12 OS=Mus musculus GN=Mapk12 PE=1 SV=1 SMK12_MOUSE [2]	42 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1 A2AL78_MOUSE	26 kDa	1	0	6	6	6	0	6	6	6	0	6	6	0
Band 3 anion transport protein OS=Mus musculus GN=Slc4a1 PE=1 SV=1 B3AT_MOUSE (+1)	103 kDa	1	4	6	7	6	4	7	7	6	4	7	6	4
CDGSH iron-sulfur domain-containing protein 1 OS=Mus musculus GN=Cisd1 PE=1 SV=1 CISD1_MOUSE	12 kDa	1	7	7	0	7	7	0	0	7	7	0	7	7
Band 4.1-like protein 2 OS=Mus musculus GN=Epb41l2 PE=1 SV=2 E41L2_MOUSE	110 kDa	1	2	5	2	5	2	2	2	5	2	2	5	2
Cluster of Protein Cald1 OS=Mus musculus GN=Cald1 PE=2 SV=1 (E9QA15_MOUSE) E9QA15_MOUSE [2]	89 kDa	1	5	8	5	8	5	5	5	8	5	5	8	5

Eukaryotic translation initiation factor 3 subunit A OS=Mus musculus GN=Eif3a PE=1 S EIF3A_MOUSE	162 kDa	1	5	12	8	12	5	8	8	12	5	8	12	5
Cluster of Kinesin-1 heavy chain OS=Mus musculus GN=Kif5b PE=1 SV=3 (KINH_MOUSE KINH_MOUSE)	110 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
Spectrin beta 1 OS=Mus musculus GN=Sptb PE=2 SV=1	268 kDa	1	7	9	8	9	7	8	8	9	7	8	9	7
Cluster of Thioredoxin OS=Mus musculus GN=Txn PE=1 SV=3 (THIO_MOUSE)	12 kDa	1	5	7	5	7	5	5	5	7	5	7	5	7
Cluster of Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Mus m AN32A_MOUSE [3]	29 kDa	1	6	8	11	8	6	11	11	8	6	11	8	6
Branched-chain-amino-acid aminotransferase, mitochondrial OS=Mus musculus GN=EBCAT2_MOUSE	44 kDa	1	7	7	8	7	7	8	8	7	7	8	7	7
Xin actin-binding repeat-containing protein 1 OS=Mus musculus GN=Xirp1 PE=4 SV=1	124 kDa	1	5	11	12	11	5	12	12	11	5	12	11	5
Protein unc-45 homolog B OS=Mus musculus GN=Unc45b PE=1 SV=1	104 kDa	1	4	9	13	9	4	13	13	9	4	13	9	4
Vigilin OS=Mus musculus GN=Hdlbp PE=1 SV=1	142 kDa	1	3	0	3	0	3	3	3	0	3	3	0	3
Isocitrate dehydrogenase [NADP] cytoplasmic OS=Mus musculus GN=ldh1 PE=1 SV=2	47 kDa	1	8	9	6	9	8	6	6	9	8	6	9	8
LETM1 and EF-hand domain-containing protein 1, mitochondrial OS=Mus musculus GI LETM1_MOUSE	83 kDa	1	4	10	9	10	4	9	9	10	4	9	10	4
NADH dehydrogenase [ubiquinone] iron-sulfur protein 7, mitochondrial OS=Mus mus NDUS7_MOUSE	25 kDa	1	4	8	7	8	4	7	7	8	4	7	8	4
Ornithine aminotransferase, mitochondrial OS=Mus musculus GN=Oat PE=1 SV=1	48 kDa	1	4	7	8	7	4	8	8	7	4	8	7	4
Cluster of 40S ribosomal protein S9 OS=Mus musculus GN=Rps9 PE=2 SV=3 (RS9_MOUSE RS9_MOUSE [4])	23 kDa	1	6	9	10	9	6	10	10	9	6	10	9	6
Succinate-semialdehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh5:SSDH_MOUSE	56 kDa	1	6	10	11	10	6	11	11	10	6	11	10	6
Acylphosphatase-2 OS=Mus musculus GN=Acp2 PE=2 SV=2	12 kDa	1	4	6	7	6	4	7	7	6	4	7	6	4
Cluster of Caveolin-3 OS=Mus musculus GN=Cav3 PE=1 SV=1 (CAV3_MOUSE)	17 kDa	1	5	10	9	10	5	9	9	10	5	9	10	5
Dnaj homolog subfamily A member 2 OS=Mus musculus GN=Dnaja2 PE=1 SV=1	46 kDa	1	8	9	10	9	8	10	10	9	8	10	9	8
Myosin-14 OS=Mus musculus GN=Myl14 PE=4 SV=1	229 kDa	1	7	4	6	4	7	6	6	4	7	6	4	7
Amine oxidase [flavin-containing] B OS=Mus musculus GN=Maob PE=1 SV=4	59 kDa	1	5	7	4	7	5	4	4	7	5	4	7	5
Protein Lmo7 OS=Mus musculus GN=Lmo7 PE=2 SV=1	193 kDa	1	4	8	7	8	4	7	7	8	4	7	8	4
Fibrinogen beta chain OS=Mus musculus GN=Fgb PE=2 SV=1	55 kDa	1	7	8	9	8	7	9	9	8	7	9	8	7
Glutathione S-transferase P 1 OS=Mus musculus GN=Gstp1 PE=1 SV=2	24 kDa	1	3	7	7	7	3	7	7	7	3	7	7	3
Puromycin-sensitive aminopeptidase OS=Mus musculus GN=Npepps PE=1 SV=2	103 kDa	1	5	3	4	3	5	4	4	3	5	4	3	5
Regulator of microtubule dynamics protein 1 OS=Mus musculus GN=Rmdn1 PE=2 SV=	35 kDa	1	5	9	8	9	5	8	8	9	5	8	9	5
Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=1 SV=2	62 kDa	1	4	7	7	7	4	7	7	7	4	7	7	4
Cluster of Ceruloplasmin OS=Mus musculus GN=Cp PE=4 SV=1 (G3X8Q5_MOUSE)	124 kDa	1	6	7	7	7	6	7	7	7	6	7	7	6
Glutathione S-transferase kappa 1 OS=Mus musculus GN=Gstk1 PE=1 SV=3	26 kDa	1	5	7	7	7	5	7	7	7	5	7	7	5
Cluster of Sarcolemmal membrane-associated protein OS=Mus musculus GN=Slmap P H7BX64_MOUSE [3]	95 kDa	1	5	9	6	9	5	6	6	9	5	6	9	5
Mitochondrial carnitine/acylcarnitine carrier protein OS=Mus musculus GN=Slc25a20 MCAT_MOUSE	33 kDa	1	7	7	7	7	7	7	7	7	7	7	7	7
Methylmalonyl-CoA mutase, mitochondrial OS=Mus musculus GN=Mut PE=2 SV=2	83 kDa	1	8	0	6	0	8	6	6	0	8	6	0	8
Proteasome subunit alpha type-1 OS=Mus musculus GN=Psm1 PE=1 SV=1	30 kDa	1	3	9	7	9	3	7	7	9	3	7	9	3
Spectrin alpha chain, erythrocytic 1 OS=Mus musculus GN=Spta1 PE=2 SV=3	280 kDa	1	6	8	5	8	6	5	5	8	6	5	8	6
Short/branched chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musc ACDSB_MOUSE	48 kDa	1	0	7	6	7	0	6	6	7	0	6	7	0
Apolipoprotein A-IV OS=Mus musculus GN=Apoa4 PE=2 SV=3	45 kDa	1	5	5	10	5	5	10	10	5	5	10	5	5
Cluster of F-actin-capping protein subunit beta OS=Mus musculus GN=Capzb PE=1 SV:CAPZB_MOUSE [2]	31 kDa	1	7	8	11	8	7	11	11	8	7	11	8	7
Cofilin-2 OS=Mus musculus GN=Cfl2 PE=1 SV=1	19 kDa	1	11	6	4	6	11	4	4	6	11	4	6	11
Phospholipid hydroperoxide glutathione peroxidase, nuclear OS=Mus musculus GN=GPX42_MOUSE (+1)	29 kDa	1	0	8	7	8	0	7	7	8	0	7	8	0
Decorin OS=Mus musculus GN=Dcn PE=2 SV=1	40 kDa	1	8	10	10	10	8	10	10	10	8	10	10	8
Ryanodine receptor 3 OS=Mus musculus GN=Ryr3 PE=1 SV=1	551 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Tropomyosin alpha-4 chain OS=Mus musculus GN=Tpm4 PE=2 SV=3	28 kDa	1	9	5	8	5	9	8	8	5	9	8	5	9
LIM and calponin homology domains-containing protein 1 OS=Mus musculus GN=Lim1 D3YU22_MOUSE (+1)	120 kDa	1	0	4	0	0	4	0	4	0	4	0	0	4
Cluster of Synaptopodin-2 OS=Mus musculus GN=Synpo2 PE=2 SV=1 (E9Q1U2_MOUSE E9Q1U2_MOUSE [2])	136 kDa	1	0	5	6	5	0	6	6	5	0	6	5	0
Ferrochelatase, mitochondrial OS=Mus musculus GN=Fech PE=1 SV=2	47 kDa	1	4	9	7	9	4	7	7	9	4	7	9	4
Histidine triad nucleotide-binding protein 2, mitochondrial OS=Mus musculus GN=Hin HINT2_MOUSE	17 kDa	1	7	6	6	6	7	6	6	6	7	6	6	7
Heterochromatin protein 1-binding protein 3 OS=Mus musculus GN=Hp1bp3 PE=1 SV:HP1B3_MOUSE	61 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Adenylosuccinate synthetase OS=Mus musculus GN=Adssl1 PE=3 SV=1	53 kDa	1	8	7	8	7	8	8	8	7	8	8	7	8
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 OS=Mus musculus GN=NDUB4_MOUSE	15 kDa	1	5	0	6	0	5	6	6	0	5	6	0	5
Cluster of 26S proteasome non-ATPase regulatory subunit 2 OS=Mus musculus GN=Ps PSDM2_MOUSE [2]	100 kDa	1	6	11	8	11	6	8	8	11	6	8	11	6
Ig mu chain C region secreted form OS=Mus musculus GN=Igh-6 PE=1 SV=2	50 kDa	1	0	7	8	7	0	8	8	7	0	8	7	0
Sulfated glycoprotein 1 OS=Mus musculus GN=Psap PE=4 SV=1	61 kDa	1	4	8	10	8	4	10	10	8	4	10	8	4
Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase ODB2_MOUSE	53 kDa	1	9	8	5	8	9	5	5	8	9	5	8	9
Cluster of Elongation factor 1-delta (Fragment) OS=Mus musculus GN=Eef1d PE=2 SV: E9QN08_MOUSE [3]	27 kDa	1	5	7	5	7	5	5	5	7	5	5	7	5
Nicotinamide phosphoribosyltransferase OS=Mus musculus GN=Nampt PE=1 SV=1	55 kDa	1	0	8	8	8	0	8	8	8	0	8	8	0
Cluster of Palmelphin OS=Mus musculus GN=Palmd PE=2 SV=1 (Q3UVT7_MOUSE)	63 kDa	1	3	8	4	8	3	4	4	8	3	4	8	3
T-complex protein 1 subunit epsilon OS=Mus musculus GN=Cct5 PE=1 SV=1	60 kDa	1	6	3	8	3	6	8	8	3	6	8	3	6
Cluster of MCG130675 OS=Mus musculus GN=2410018M08Rik PE=2 SV=1 (B2RPU8_MOUSE B2RPU8_MOUSE [2])	84 kDa	1	0	6	5	6	0	5	5	6	0	5	6	0
Cluster of Nuclease-sensitive element-binding protein 1 (Fragment) OS=Mus musculus: A2BGG7_MOUSE [2]	25 kDa	1	0	10	14	10	0	14	14	10	0	14	10	0
ADP/ATP translocase 4 OS=Mus musculus GN=Slc25a31 PE=2 SV=1	35 kDa	1	1	1	1	1	1	1	1	1	1	1	1	1
Cadherin-13 OS=Mus musculus GN=Cdh13 PE=1 SV=2	78 kDa	1	3	7	6	7	3	6	6	7	3	6	7	3
Calnexin OS=Mus musculus GN=Canx PE=1 SV=1	67 kDa	1	4	8	6	8	4	6	6	8	4	6	8	4
Fibrinogen gamma chain OS=Mus musculus GN=Fgg PE=2 SV=1	49 kDa	1	5	5	8	5	5	8	8	5	5	8	5	5
Fibronectin OS=Mus musculus GN=Fn1 PE=1 SV=4	273 kDa	1	3	11	7	11	3	7	7	11	3	7	11	3
Cluster of cAMP-dependent protein kinase catalytic subunit alpha OS=Mus musculus (KAPCA_MOUSE [2])	41 kDa	1	6	8	9	8	6	9	9	8	6	9	8	6
Cluster of 60S ribosomal protein L24 OS=Mus musculus GN=Rpl24 PE=2 SV=2 (RL24_MOUSE RL24_MOUSE)	18 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
40S ribosomal protein S6 OS=Mus musculus GN=Rps6 PE=1 SV=1	29 kDa	1	3	7	7	7	3	7	7	7	3	7	7	3

Transferrin OS=Mus musculus GN=Ttr PE=1 SV=1	TTHY_MOUSE	16 kDa	1	5	5	7	5	5	7	7	5	5	7	5	5
Cluster of Archvillin OS=Mus musculus GN=Svil PE=2 SV=1 (Q8K4L2_MOUSE)	Q8K4L2_MOUSE	227 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Cullin-associated NEDD8-dissociated protein 2 OS=Mus musculus GN=Cand2 PE=1 SV=CAND2_MOUSE	CAND2_MOUSE	136 kDa	1	0	10	7	10	0	7	7	10	0	7	10	0
Histone-lysine N-methyltransferase, H3 lysine-36 and H4 lysine-20-specific OS=Mus mE9QAE4_MOUSE	E9QAE4_MOUSE	296 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Receptor accessory protein 5, isoform CRA_a OS=Mus musculus GN=Reep5 PE=4 SV=1G3X8R0_MOUSE	G3X8R0_MOUSE	21 kDa	1	6	7	8	7	6	8	8	7	6	8	7	6
Laminin subunit beta-2 OS=Mus musculus GN=Lamb2 PE=2 SV=2	LAMB2_MOUSE	197 kDa	1	6	5	7	5	6	7	7	5	6	7	5	6
NADH dehydrogenase (ubiquinone) 1 beta subcomplex subunit 11, mitochondrial OS=NDUBB_MOUSE	NDUBB_MOUSE	17 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Vitamin D-binding protein OS=Mus musculus GN=Gc PE=1 SV=2	VTDB_MOUSE	54 kDa	1	6	7	5	7	6	5	5	7	6	5	7	6
Isoform 2 of Dynamin-1 OS=Mus musculus GN=Dnm1	sp P39053-2 DYN1_MOUSE	51 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Probable C->U-editing enzyme APOBEC-2 OS=Mus musculus GN=Apobec2 PE=1 SV=1 ABEC2_MOUSE	ABEC2_MOUSE	26 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Acyl-coenzyme A thioesterase 9, mitochondrial OS=Mus musculus GN=Acot ACOT9_MOUSE [2]	ACOT9_MOUSE [2]	51 kDa	1	3	3	5	3	3	5	5	3	3	5	3	3
Cluster of ATPase family AAA domain-containing protein 3 OS=Mus musculus GN=Ata ATAD3_MOUSE	ATAD3_MOUSE	67 kDa	1	4	5	4	5	4	4	4	5	4	4	5	4
Elongation factor 1-alpha 1 OS=Mus musculus GN=Eef1a1 PE=1 SV=3	E1A1_MOUSE	50 kDa	1	2	5	8	5	2	8	8	5	2	8	5	2
Acylpyruvate FAHD1, mitochondrial OS=Mus musculus GN=Fahd1 PE=1 SV=2	FAHD1_MOUSE	25 kDa	1	3	7	4	7	3	4	4	7	3	4	7	3
Ferritin heavy chain OS=Mus musculus GN=Fth1 PE=1 SV=2	FTH1_MOUSE	21 kDa	1	4	8	8	8	4	8	8	8	4	8	8	4
Cluster of Eukaryotic initiation factor 4A-II OS=Mus musculus GN=Eif4a2 PE=2 SV=2 (EIF4A2_MOUSE [2])	EIF4A2_MOUSE [2]	46 kDa	1	7	8	6	8	7	6	6	8	7	6	8	7
Eukaryotic translation initiation factor 4B OS=Mus musculus GN=Eif4b PE=1 SV=1	IF4B_MOUSE	69 kDa	1	0	9	7	9	0	7	7	9	0	7	9	0
Adenylosuccinate lyase OS=Mus musculus GN=Adsl PE=2 SV=2	PUR8_MOUSE	55 kDa	1	4	5	3	5	4	3	3	5	4	3	5	4
Peripherin OS=Mus musculus GN=Prph PE=1 SV=2	PERI_MOUSE (+1)	54 kDa	1	1	0	1	0	1	1	1	0	1	1	0	1
[Protein ADP-ribosylarginine] hydrolase-like protein 1 OS=Mus musculus GN=Adprhl1 ARHL1_MOUSE	ARHL1_MOUSE	40 kDa	1	7	7	7	7	7	7	7	7	7	7	7	7
Actin-related protein 2/3 complex subunit 4 OS=Mus musculus GN=Arpc4 PE=1 SV=3	ARPC4_MOUSE	20 kDa	1	3	3	4	3	3	4	4	3	3	4	3	3
Transcription elongation factor B polypeptide 2 OS=Mus musculus GN=Tceb2 PE=1 SV ELOB_MOUSE	ELOB_MOUSE	13 kDa	1	6	5	5	5	6	5	5	5	5	6	5	6
Serine/arginine-rich-splicing factor 1 OS=Mus musculus GN=Srsf1 PE=2 SV=1	H7BX95_MOUSE (+1)	28 kDa	1	8	4	7	4	8	7	7	4	8	7	4	8
Laminin subunit alpha-4 OS=Mus musculus GN=Lama4 PE=1 SV=2	LAMAA_MOUSE	202 kDa	1	3	5	3	5	3	3	3	5	3	3	5	3
Mitochondrial fission process protein 1 OS=Mus musculus GN=Mtfp1 PE=1 SV=1	MTFP1_MOUSE	18 kDa	1	3	0	3	0	3	3	3	0	3	3	0	3
Prostaglandin reductase 2 OS=Mus musculus GN=Ptgr2 PE=1 SV=2	PTGR2_MOUSE	38 kDa	1	6	8	7	8	6	7	7	8	6	7	8	6
Protein Sf3b2 OS=Mus musculus GN=Sf3b2 PE=2 SV=1	Q3UJB0_MOUSE	98 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
40S ribosomal protein S12 OS=Mus musculus GN=Rps12 PE=2 SV=1	Q6ZWZ6_MOUSE	15 kDa	1	5	4	3	4	5	3	3	4	5	3	4	5
26S protease regulatory subunit 7 OS=Mus musculus GN=Psmc2 PE=2 SV=1	Q8BVQ9_MOUSE	53 kDa	1	3	6	7	6	3	7	7	6	3	7	6	3
60S ribosomal protein L5 OS=Mus musculus GN=Rpl5 PE=1 SV=3	RL5_MOUSE	34 kDa	1	4	8	5	8	4	5	5	8	4	5	8	4
Transcription factor A, mitochondrial OS=Mus musculus GN=Tfam PE=1 SV=2	TFAM_MOUSE	28 kDa	1	6	5	7	5	6	7	7	5	6	7	5	6
Glycogenin-1 OS=Mus musculus GN=Gyg1 PE=2 SV=3	GLYG_MOUSE (+1)	37 kDa	1	3	5	6	5	3	6	6	5	3	6	5	3
Muscle-related coiled-coil protein OS=Mus musculus GN=Murc PE=2 SV=1	MURC_MOUSE	41 kDa	1	7	5	6	5	7	6	6	5	7	6	5	7
Cluster of Myosin light polypeptide 6 OS=Mus musculus GN=Myl6 PE=1 SV=3 (MYL6_MOUSE [2])	MYL6_MOUSE [2]	17 kDa	1	3	7	8	7	3	8	8	7	3	8	7	3
26S protease regulatory subunit 4 OS=Mus musculus GN=Psmc1 PE=1 SV=1	PRSD_MOUSE	49 kDa	1	0	7	3	7	0	3	3	7	0	3	7	0
26S proteasome non-ATPase regulatory subunit 1 OS=Mus musculus GN=Psm1 PE=1 SV=1	PSMD1_MOUSE	106 kDa	1	4	4	4	4	4	4	4	4	4	4	4	4
Cluster of Prothymosin alpha OS=Mus musculus GN=Ptma PE=1 SV=2 (PTMA_MOUSE, PTMA_MOUSE)	PTMA_MOUSE	12 kDa	1	3	6	5	6	3	5	5	6	3	5	6	3
Bifunctional purine biosynthesis protein PURH OS=Mus musculus GN=Atic PE=1 SV=2 PUR9_MOUSE	PUR9_MOUSE	64 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Cluster of Protein kinase, cAMP dependent regulatory, type II alpha OS=Mus musculus: Q8K1M3_MOUSE	Q8K1M3_MOUSE	46 kDa	1	4	6	9	6	4	9	9	6	4	9	6	4
Aspartate--tRNA ligase, cytoplasmic OS=Mus musculus GN=Dars PE=2 SV=2	SYDC_MOUSE	57 kDa	1	5	6	8	6	5	8	8	6	5	8	6	5
Cluster of Elongation factor G, mitochondrial OS=Mus musculus GN=Gfm1 PE=2 SV=1 EFGM_MOUSE	EFGM_MOUSE	84 kDa	1	6	5	0	5	6	0	0	5	6	0	5	6
Cluster of Protein Gm4945 OS=Mus musculus GN=Gm4945 PE=4 SV=1 (D3Z102_MOUSE D3Z102_MOUSE)	D3Z102_MOUSE	16 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Cluster of Protein Et14 OS=Mus musculus GN=Et14 PE=2 SV=1 (E9QAU4_MOUSE)	E9QAU4_MOUSE	218 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Coiled-coil protein associated with myosin II and DISC1 OS=Mus musculus GN=Ccdc14 A2AST1_MOUSE (+1)	A2AST1_MOUSE (+1)	165 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Rab GDP dissociation inhibitor beta OS=Mus musculus GN=Gdi2 PE=1 SV=1 GDIB_MOUSE [2]	GDIB_MOUSE [2]	51 kDa	1	5	5	4	5	5	4	4	5	5	4	5	5
Cluster of Glyoxalase domain-containing protein 4 OS=Mus musculus GN=Glod4 PE=2 GLOD4_MOUSE [2]	GLOD4_MOUSE [2]	33 kDa	1	7	7	8	7	7	8	8	7	7	8	7	7
Eukaryotic translation initiation factor 5A-1 OS=Mus musculus GN=Eif5a PE=1 SV=2	IF5A1_MOUSE	17 kDa	1	6	0	5	0	6	5	5	0	6	5	0	6
Junctophilin-2 OS=Mus musculus GN=Jph2 PE=1 SV=2	JPH2_MOUSE	75 kDa	1	10	5	8	5	10	8	8	5	10	8	5	10
Cluster of Poly(rC)-binding protein 1 OS=Mus musculus GN=Pcbp1 PE=1 SV=1 (PCBP1_MOUSE PCBP1_MOUSE [4])	PCBP1_MOUSE	37 kDa	1	3	6	6	6	3	6	6	6	3	6	6	3
Programmed cell death 6-interacting protein OS=Mus musculus GN=Pdc6ip PE=1 SV=1 PDC6_MOUSE	PDC6_MOUSE	96 kDa	1	5	7	5	7	5	7	7	5	4	7	5	4
Cluster of Ank1 protein OS=Mus musculus GN=Ank1 PE=2 SV=1 (Q0VGY9_MOUSE)	Q0VGY9_MOUSE [3]	207 kDa	1	2	2	3	2	2	3	3	2	2	3	2	2
Sorting and assembly machinery component 50 homolog OS=Mus musculus GN=Samr SAM50_MOUSE	SAM50_MOUSE	52 kDa	1	4	6	6	6	4	6	6	6	4	6	6	4
T-complex protein 1 subunit theta OS=Mus musculus GN=Cct8 PE=1 SV=3	TCPO_MOUSE	60 kDa	1	7	8	7	8	7	7	7	8	7	7	8	7
Cluster of Muscular LMNA-interacting protein OS=Mus musculus GN=Mlip PE=4 SV=1 tr V9GWW6 V9GWW6_MOUSE	V9GWW6_MOUSE	4 kDa	1	4	4	8	4	4	8	8	4	4	8	4	4
Cluster of Isoform 2 of Protein phosphatase 1 regulatory subunit 12A OS=Mus musculus sp Q9DBR7-2 MYPT1_MOUSE	Q9DBR7-2_MOUSE	4 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Mitochondrial pyruvate carrier 2 OS=Mus musculus GN=Mpc2 PE=1 SV=1	MPC2_MOUSE	14 kDa	1	4	4	5	4	4	5	5	4	4	5	4	4
Cluster of Replication factor C subunit 1 OS=Mus musculus GN=Rfc1 PE=2 SV=1 (G3UUG3UWX1_MOUSE [6])	G3UUG3UWX1_MOUSE [6]	126 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cluster of Calumenin OS=Mus musculus GN=Calu PE=1 SV=1 (CALU_MOUSE)	CALU_MOUSE	37 kDa	1	7	5	5	5	7	5	5	5	7	5	5	7
60S acidic ribosomal protein P0 (Fragment) OS=Mus musculus GN=Rplp0 PE=2 SV=1	D3YVM5_MOUSE (+2)	16 kDa	1	0	6	6	6	0	6	6	6	0	6	6	0
Filamin-B OS=Mus musculus GN=Flnb PE=1 SV=3	FLNB_MOUSE	278 kDa	1	2	1	5	1	2	5	5	1	2	5	1	2
Pyruvate carboxylase OS=Mus musculus GN=Pcx PE=3 SV=1	G5E8R3_MOUSE (+1)	130 kDa	1	0	7	4	7	0	4	4	7	0	4	7	0
Unconventional myosin-Ic OS=Mus musculus GN=Myo1c PE=1 SV=2	MYO1C_MOUSE (+3)	122 kDa	1	5	7	0	7	5	0	0	7	5	0	7	5
26S protease regulatory subunit 6A OS=Mus musculus GN=Psmc3 PE=1 SV=2	PRSDA_MOUSE	50 kDa	1	3	5	3	5	3	3	3	5	3	3	5	3
Radixin OS=Mus musculus GN=Rdx PE=1 SV=3	RADI_MOUSE	69 kDa	1	3	4	5	4	3	5	5	4	3	5	4	3
60S ribosomal protein L8 OS=Mus musculus GN=Rpl8 PE=2 SV=2	RL8_MOUSE	28 kDa	1	3	5	4	5	3	4	4	5	3	4	5	3
Bifunctional glutamate/proline--tRNA ligase OS=Mus musculus GN=Eprs PE=1 SV=4	SYEP_MOUSE	170 kDa	1	5	4	3	4	5	3	3	4	5	3	4	5

T-complex protein 1 subunit eta OS=Mus musculus GN=Cct7 PE=1 SV=1	TCPH_MOUSE	60 kDa	1	3	5	8	5	3	8	8	5	3	8	5	3
Cluster of STE20-like serine/threonine-protein kinase OS=Mus musculus GN=Slk PE=1 SV=1	SLK_MOUSE [2]	141 kDa	1	4	0	0	0	0	0	0	0	0	0	0	4
Cluster of Uncharacterized protein OS=Mus musculus GN=Gm10260 PE=3 SV=2 (F6YV F6YV7_MOUSE [2])		18 kDa	1	0	0	5	0	0	5	5	0	5	0	0	0
Cluster of 5-azacytidine induced gene 1 OS=Mus musculus GN=Azi1 PE=2 SV=1 (B1AXI B1AXI9_MOUSE [2])		120 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Eukaryotic translation initiation factor 3 subunit G OS=Mus musculus GN=EIF3G PE=1 SV=1	EIF3G_MOUSE	36 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Rho GDP-dissociation inhibitor 1 OS=Mus musculus GN=Arhgdia PE=1 SV=3	GDIR1_MOUSE	23 kDa	1	7	4	5	4	7	5	5	4	7	5	4	7
60S ribosomal protein L10 (Fragment) OS=Mus musculus GN=Rpl10 PE=4 SV=1	I7HLV2_MOUSE (+1)	23 kDa	1	5	3	3	3	5	3	3	3	5	3	3	5
Lipoprotein lipase OS=Mus musculus GN=Lpl PE=1 SV=3	LPL_MOUSE	53 kDa	1	5	6	7	6	5	7	7	6	5	7	6	5
Myosin-10 OS=Mus musculus GN=Myl10 PE=1 SV=2	MYH10_MOUSE (+2)	229 kDa	1	2	1	1	2	1	1	1	2	1	2	1	2
Nuclear pore complex-associated intranuclear coiled-coil protein TPR OS=Mus musculus GN=Q7M739_MOUSE		267 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
60S ribosomal protein L31 OS=Mus musculus GN=Rpl31 PE=2 SV=1	RL31_MOUSE	14 kDa	1	4	6	5	6	4	5	6	4	5	6	4	4
40S ribosomal protein S3 OS=Mus musculus GN=Rps3 PE=1 SV=1	RS3_MOUSE	27 kDa	1	6	6	6	6	6	6	6	6	6	6	6	6
Adenosylhomocysteinase OS=Mus musculus GN=Ahcy PE=1 SV=3	SAHH_MOUSE	48 kDa	1	7	4	4	4	7	4	4	4	7	4	4	7
T-complex protein 1 subunit alpha OS=Mus musculus GN=Tcp1 PE=1 SV=3	TCPA_MOUSE	60 kDa	1	4	7	7	7	4	7	7	7	4	7	7	4
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 3 OS=Mus musculus GN=NDUB3_MOUSE		12 kDa	1	4	4	5	4	4	5	5	4	4	5	4	4
UPF0317 protein C14orf159 homolog, mitochondrial OS=Mus musculus PE=2 SV=1	CN159_MOUSE (+1)	66 kDa	1	7	6	5	6	7	5	5	6	7	5	6	7
Collagen alpha-2(VI) chain OS=Mus musculus GN=Col6a2 PE=2 SV=3	CO6A2_MOUSE	110 kDa	1	3	5	4	5	3	4	4	5	3	4	5	3
60S ribosomal protein L13a OS=Mus musculus GN=Rpl13a PE=2 SV=1	E9Q5A0_MOUSE	48 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Kinesin-like protein KIF21B OS=Mus musculus GN=Kif21b PE=2 SV=1	F8VQE2_MOUSE (+1)	181 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Histone H1.1 OS=Mus musculus GN=Hist1h1a PE=1 SV=2	H11_MOUSE	22 kDa	1	2	0	1	0	2	1	1	0	2	1	0	2
GTP:AMP phosphotransferase, mitochondrial OS=Mus musculus GN=Ak3 PE=1 SV=3	KAD3_MOUSE	25 kDa	1	8	8	7	8	8	7	7	8	8	7	8	8
Cluster of NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrion OS=Mus musculus GN=NDUB8_MOUSE		22 kDa	1	0	6	7	6	0	7	7	6	0	7	6	0
60S ribosomal protein L13 OS=Mus musculus GN=Rpl13 PE=2 SV=3	RL13_MOUSE	24 kDa	1	5	7	4	7	5	4	4	7	5	4	7	5
T-complex protein 1 subunit beta OS=Mus musculus GN=Cct2 PE=1 SV=4	TCPB_MOUSE	57 kDa	1	3	6	3	6	3	3	3	6	3	3	6	3
Mitochondrial import inner membrane translocase subunit TIM44 OS=Mus musculus GN=TIM44_MOUSE		51 kDa	1	4	4	7	4	4	7	7	4	4	7	4	4
Transketolase OS=Mus musculus GN=Tkt PE=1 SV=1	TKT_MOUSE	68 kDa	1	5	5	4	5	5	4	4	5	4	5	4	5
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 6 OS=Mus musculus GN=A2AP32_MOUSE (+1)		12 kDa	1	0	0	4	0	0	4	0	0	4	0	0	0
Cluster of Mitochondrial pyruvate carrier 1 OS=Mus musculus GN=Mpc1 PE=1 SV=1 (MPC1_MOUSE [3])		12 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cluster of Probable helicase senataxin OS=Mus musculus GN=Setx PE=2 SV=1 (SETX_A.SETX_MOUSE)		298 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Ubiquinone biosynthesis monooxygenase COQ6 OS=Mus musculus GN=Coq6 PE=2 SV=2	COQ6_MOUSE	51 kDa	1	5	6	4	6	5	4	4	6	5	4	6	5
Cluster of D-dopachrome decarboxylase OS=Mus musculus GN=Ddt PE=1 SV=3 (DOPD DOPD_MOUSE)		13 kDa	1	0	4	0	0	4	0	4	0	4	0	4	0
MCG116065 OS=Mus musculus GN=Gm6472 PE=4 SV=1	J3QJZ3_MOUSE (+1)	22 kDa	1	5	7	5	7	5	5	5	7	5	5	7	5
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Mus musculus GN=NDUAC_MOUSE		17 kDa	1	6	7	5	7	6	5	5	7	6	5	7	6
Plasminogen OS=Mus musculus GN=Plg PE=1 SV=3	PLMN_MOUSE	91 kDa	1	6	7	8	7	6	8	8	7	6	8	7	6
Cluster of Serine/threonine-protein phosphatase PP1-alpha catalytic subunit OS=Mus musculus GN=PP1A_MOUSE [4]		38 kDa	1	5	6	5	6	5	5	6	5	5	6	5	5
60S ribosomal protein L27 OS=Mus musculus GN=Rpl27 PE=2 SV=2	RL27_MOUSE	16 kDa	1	3	4	5	4	3	5	5	4	3	5	4	3
40S ribosomal protein S25 OS=Mus musculus GN=Rps25 PE=2 SV=1	RS25_MOUSE	14 kDa	1	3	5	4	5	3	4	4	5	3	4	5	3
Heat shock protein 75 kDa, mitochondrial OS=Mus musculus GN=Trap1 PE=1 SV=1	TRAP1_MOUSE	80 kDa	1	4	4	6	4	4	6	6	4	4	6	4	4
Zinc-binding alcohol dehydrogenase domain-containing protein 2 OS=Mus musculus GN=CZADH2_MOUSE		41 kDa	1	4	9	6	9	4	6	6	9	4	6	9	4
Synaptopodin 2-like protein OS=Mus musculus GN=Synpo2l PE=2 SV=1	B2RQK7_MOUSE	103 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
40S ribosomal protein S5 (Fragment) OS=Mus musculus GN=Rps5 PE=3 SV=1	D3YYM6_MOUSE (+2)	20 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Dystroglycan OS=Mus musculus GN=Dag1 PE=1 SV=4	DAG1_MOUSE	97 kDa	1	4	4	4	4	4	4	4	4	4	4	4	4
Eukaryotic translation initiation factor 4 gamma 1 OS=Mus musculus GN=EIF4g1 PE=2 SV=1	E9PVC5_MOUSE	175 kDa	1	3	7	5	7	3	5	5	7	3	5	7	3
Ras-related protein Rab-11A OS=Mus musculus GN=Rab11a PE=2 SV=1	E9Q3P9_MOUSE (+4)	17 kDa	1	4	6	5	6	4	5	5	6	4	5	6	4
Cluster of Protein Serpinb6a OS=Mus musculus GN=Serpib6a PE=2 SV=1 (F8WIV2_M F8WIV2_MOUSE [3])		45 kDa	1	4	6	5	6	4	5	5	6	4	5	6	4
Four and a half LIM domains protein 1 OS=Mus musculus GN=Fhl1 PE=2 SV=3	FHL1_MOUSE	32 kDa	1	4	6	7	6	4	7	7	6	4	7	6	4
Glutaryl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Gcdh PE=2 SV=2	GCDH_MOUSE	49 kDa	1	3	3	3	3	3	3	3	3	3	3	3	3
Importin-5 OS=Mus musculus GN=Ipo5 PE=1 SV=3	IPO5_MOUSE	124 kDa	1	3	7	3	7	3	3	3	7	3	3	7	3
MAP7 domain-containing protein 1 OS=Mus musculus GN=Map7d1 PE=1 SV=1	MA7D1_MOUSE	93 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Cluster of Myosin regulatory light chain 12B OS=Mus musculus GN=Myl12b PE=1 SV=1 (ML12B_MOUSE [3])		20 kDa	1	4	4	4	4	4	4	4	4	4	4	4	4
Myosin light chain kinase 3 OS=Mus musculus GN=Mylk3 PE=1 SV=1	MYLK3_MOUSE	86 kDa	1	2	3	3	3	2	3	3	3	2	3	3	2
Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform OS=Mus musculus GN=PP2AA_MOUSE		36 kDa	1	7	3	3	3	7	3	3	3	7	3	3	7
60S ribosomal protein L10a OS=Mus musculus GN=Rpl10a PE=1 SV=3	RL10A_MOUSE	25 kDa	1	6	5	0	5	6	0	0	5	6	0	5	6
Cluster of T-complex protein 1 subunit delta OS=Mus musculus GN=Cct4 PE=1 SV=3 (TCPD_MOUSE [2])		58 kDa	1	3	6	8	6	3	8	8	6	3	8	6	3
Cluster of NEDD8-conjugating enzyme Ubc12 OS=Mus musculus GN=Ubc2m PE=2 SV=2	UBC12_MOUSE	21 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Cluster of Estradiol 17-beta-dehydrogenase 8 OS=Mus musculus GN=Hsd17b8 PE=1 SV=1 (DH8_MOUSE [2])		27 kDa	1	0	7	7	7	0	7	7	7	0	7	7	0
Cluster of NSF1 cofactor p47 OS=Mus musculus GN=Nsf1c PE=2 SV=1 (A2AT02_MOL A2AT02_MOUSE [4])		37 kDa	1	5	4	7	4	5	7	7	4	5	7	4	5
Ras suppressor protein 1 OS=Mus musculus GN=Rsu1 PE=4 SV=1	A2AUR7_MOUSE (+2)	30 kDa	1	0	7	7	7	0	7	7	7	0	7	7	0
ATPase family AAA domain-containing protein 1 OS=Mus musculus GN=Atad1 PE=1 SV=1	ATAD1_MOUSE	41 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Clustered mitochondria protein homolog OS=Mus musculus GN=Cluh PE=2 SV=2	CLU_MOUSE	148 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Malonyl-CoA decarboxylase, mitochondrial OS=Mus musculus GN=Mlycd PE=2 SV=1	DCMC_MOUSE	55 kDa	1	4	5	6	5	4	6	6	5	4	6	5	4
Protein Rpl3l OS=Mus musculus GN=Rpl3l PE=2 SV=1	E9PWZ3_MOUSE	47 kDa	1	1	6	3	6	1	3	3	6	1	3	6	1
Cluster of Calcium/calmodulin-dependent protein kinase type II subunit delta OS=Mus musculus GN=E9Q1T1_MOUSE [2]		60 kDa	1	4	4	5	4	4	5	4	4	5	4	4	5
Cardiomyopathy-associated protein 5 OS=Mus musculus GN=Cmya5 PE=4 SV=1	E9QLJ0_MOUSE	406 kDa	1	0	7	5	7	0	5	5	7	0	5	7	0
Ezrin OS=Mus musculus GN=Ezr PE=1 SV=3	EZRI_MOUSE	69 kDa	1	1	4	2	4	1	2	2	4	1	2	4	1
Protein FAM162A OS=Mus musculus GN=Fam162a PE=2 SV=1	F162A_MOUSE	18 kDa	1	0	4	5	4	0	5	5	4	0	5	4	0

Cluster of Hydroxyacyl glutathione hydrolase OS=Mus musculus GN=Hagh PE=3 SV=1	G5E8T9_MOUSE [2]	34 kDa	1	0	6	4	6	0	4	4	6	0	4	6	0
Heterogeneous nuclear ribonucleoprotein L (Fragment) OS=Mus musculus GN=Hnrrp G5E924_MOUSE		67 kDa	1	3	6	4	6	3	4	4	6	3	4	6	3
Bifunctional epoxide hydrolase 2 OS=Mus musculus GN=Ephx2 PE=1 SV=2	HYES_MOUSE	63 kDa	1	3	6	4	6	3	4	4	6	3	4	6	3
Kelch-like protein 41 OS=Mus musculus GN=Klhl41 PE=1 SV=1	KLHL41_MOUSE	68 kDa	1	3	5	3	5	3	3	3	5	3	3	5	3
Mitochondrial peptide methionine sulfoxide reductase OS=Mus musculus GN=Msra P MSRA_MOUSE		26 kDa	1	7	6	5	6	7	5	5	6	7	5	6	7
Metaxin-2 OS=Mus musculus GN=Mtx2 PE=1 SV=1	MTX2_MOUSE	30 kDa	1	0	7	5	7	0	5	5	7	0	5	7	0
Cluster of 2-oxoisovalerate dehydrogenase subunit alpha, mitochondrial OS=Mus musculus GN=ODBA_MOUSE [2]		50 kDa	1	4	0	3	0	4	3	3	0	4	3	0	4
Protein disulfide-isomerase A6 OS=Mus musculus GN=Pdia6 PE=1 SV=3	PDIA6_MOUSE (+1)	48 kDa	1	4	4	4	4	4	4	4	4	4	4	4	4
[Pyruvate dehydrogenase [lipoamide]] kinase isozyme 2, mitochondrial OS=Mus musculus GN=PKK2_MOUSE		46 kDa	1	3	5	7	5	3	7	7	5	3	7	5	3
Heterogeneous nuclear ribonucleoprotein A1 OS=Mus musculus GN=Hnmpa1 PE=2 SV=2	Q5EBP8_MOUSE (+1)	39 kDa	1	4	6	7	6	4	7	7	6	4	7	6	4
40S ribosomal protein S8 OS=Mus musculus GN=Rps8 PE=1 SV=2	RS8_MOUSE	24 kDa	1	4	0	0	0	0	0	0	0	0	0	0	0
Septin-7 OS=Mus musculus GN=Sept7 PE=1 SV=1	SEPT7_MOUSE	51 kDa	1	2	3	2	3	2	2	2	2	3	2	3	2
Sulfide:quinone oxidoreductase, mitochondrial OS=Mus musculus GN=Sqrldl PE=2 SV=2	SQRD_MOUSE	50 kDa	1	0	3	0	3	0	0	0	0	3	0	0	3
Phenylalanine--tRNA ligase beta subunit OS=Mus musculus GN=Farsb PE=2 SV=2	SYFB_MOUSE	66 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of T-complex protein 1 subunit zeta OS=Mus musculus GN=Cct6a PE=1 SV=3	ITCP3_MOUSE	58 kDa	1	6	3	3	3	6	3	3	3	6	3	3	6
Dedicator of cytokinesis protein 7 OS=Mus musculus GN=Dock7 PE=2 SV=1	A2A9M5_MOUSE (+2)	241 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Apolipoprotein E OS=Mus musculus GN=ApoE PE=1 SV=2	APOE_MOUSE	36 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Bola-like protein 3 OS=Mus musculus GN=Bola3 PE=2 SV=1	BOLA3_MOUSE	12 kDa	1	3	3	4	3	3	4	4	3	3	4	3	3
Dihydropteridine reductase OS=Mus musculus GN=Qdpr PE=1 SV=2	DHPR_MOUSE	26 kDa	1	3	3	7	3	3	7	3	3	7	3	3	3
Cluster of Proteasome subunit alpha type-7 OS=Mus musculus GN=Psm7 PE=1 SV=1	PSA7_MOUSE	28 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Ryanodine receptor 1 OS=Mus musculus GN=Ryr1 PE=1 SV=1	RYR1_MOUSE	565 kDa	1	0	0	0	0	0	0	0	0	0	0	0	0
Cluster of Isoform 2 of Clathrin light chain B OS=Mus musculus GN=Cltb (sp) Q6IRU5-2 sp) Q6IRU5-2) CLCB_MOUSE?			1	5	4	6	4	5	6	6	4	5	6	4	5
Cluster of Aldose reductase-related protein 1 OS=Mus musculus GN=Akr1b7 PE=2 SV=2	ALD1_MOUSE [3]	36 kDa	1	0	6	3	6	0	3	3	6	0	3	6	0
Apolipoprotein O OS=Mus musculus GN=ApoO PE=2 SV=1	APOO_MOUSE	24 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Cluster of 60S ribosomal protein L12 OS=Mus musculus GN=Rpl12 PE=1 SV=2	RL12_MOUSE	18 kDa	1	3	0	5	0	3	5	5	0	3	5	0	3
Cluster of 60S ribosomal protein L32 OS=Mus musculus GN=Rpl32 PE=2 SV=2	RL32_MOUSE	16 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cluster of Transforming acidic coiled-coil-containing protein 2 OS=Mus musculus GN=EQQ8T1_MOUSE		305 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
6-phosphogluconolactonase OS=Mus musculus GN=Pgl3 PE=2 SV=1	6PGL_MOUSE	27 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Alpha-actinin-4 OS=Mus musculus GN=Actn4 PE=1 SV=1	ACTN4_MOUSE	105 kDa	1	5	6	4	6	5	4	4	6	5	4	6	5
Cytosol aminopeptidase OS=Mus musculus GN=Lap3 PE=1 SV=3	AMPL_MOUSE (+1)	56 kDa	1	0	5	9	5	0	9	9	5	0	9	5	0
Beta-2-glycoprotein 1 OS=Mus musculus GN=ApoH PE=1 SV=1	APOH_MOUSE	39 kDa	1	3	6	3	3	6	6	3	3	6	3	3	3
Thioredoxin reductase 2, mitochondrial OS=Mus musculus GN=Txnrd2 PE=2 SV=1	D3Z0K8_MOUSE (+3)	53 kDa	1	0	6	7	6	0	7	7	6	0	7	6	0
Cluster of Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Mus musculus GN=GBB1_MOUSE [3]		37 kDa	1	4	5	7	5	4	7	7	5	4	7	5	4
28 kDa heat- and acid-stable phosphoprotein OS=Mus musculus GN=Pdap1 PE=1 SV=1	HAP28_MOUSE	21 kDa	1	4	4	4	4	4	4	4	4	4	4	4	4
Integrin-linked protein kinase OS=Mus musculus GN=Ilk PE=1 SV=2	ILK_MOUSE	51 kDa	1	0	6	5	6	0	5	5	6	0	5	6	0
Lamin-B2 OS=Mus musculus GN=Lmb2 PE=1 SV=2	LMBN2_MOUSE	67 kDa	1	1	4	5	4	1	5	5	4	1	5	4	1
Leiomodin-2 OS=Mus musculus GN=Lmod2 PE=1 SV=1	LMOD2_MOUSE	62 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
Nidogen-2 OS=Mus musculus GN=Nid2 PE=1 SV=2	NID2_MOUSE	154 kDa	1	5	5	7	5	5	7	7	5	5	7	5	5
2-oxoisovalerate dehydrogenase subunit beta, mitochondrial OS=Mus musculus GN=EOD8B_MOUSE		43 kDa	1	3	4	7	4	3	7	7	4	3	7	4	3
Ubiquitin carboxyl-terminal hydrolase OS=Mus musculus GN=Usp5 PE=2 SV=1	Q3U4W8_MOUSE (+1)	93 kDa	1	3	3	6	3	3	6	6	3	3	6	3	3
60S ribosomal protein L14 OS=Mus musculus GN=Rpl14 PE=2 SV=3	RL14_MOUSE	24 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Saccharopine dehydrogenase-like oxidoreductase OS=Mus musculus GN=Scppdh PE=2 SV=1	SCPPDL_MOUSE	47 kDa	1	3	4	4	4	3	4	4	4	3	4	4	3
Cluster of RNA-binding motif protein, X chromosome OS=Mus musculus GN=Rbmx PE=RBMX_MOUSE [2]		42 kDa	1	0	0	3	3	0	3	3	3	0	3	3	0
E3 ubiquitin-protein ligase HUWE1 OS=Mus musculus GN=Huwe1 PE=2 SV=1	A2AFQ0_MOUSE (+1)	483 kDa	1	0	0	4	0	4	4	0	0	4	0	4	0
Epiplakin OS=Mus musculus GN=Eppk1 PE=1 SV=2	EPIPL_MOUSE	725 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Cluster of Protein Gm20459 (Fragment) OS=Mus musculus GN=Gm20459 PE=4 SV=1	F7AA26_MOUSE	126 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Fermitin family homolog 2 OS=Mus musculus GN=Fermt2 PE=1 SV=1	FERMT2_MOUSE	78 kDa	1	0	5	7	5	0	7	7	5	0	7	5	0
Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Mus musculus GN=Fkbp3 PE=1 SV=2	FKBP3_MOUSE	25 kDa	1	7	0	6	0	7	6	6	0	7	6	0	7
Heat shock protein beta-8 OS=Mus musculus GN=Hspb8 PE=1 SV=1	HSPB8_MOUSE	22 kDa	1	3	4	5	4	3	5	5	4	3	5	4	3
Methionine-R-sulfoxide reductase B2, mitochondrial OS=Mus musculus GN=Msrb2 PE=MSRB2_MOUSE		19 kDa	1	4	6	4	6	4	6	6	4	4	6	4	4
Methylthioribulose-1-phosphate dehydratase OS=Mus musculus GN=Apip PE=1 SV=1	MTNB_MOUSE	27 kDa	1	0	6	5	6	0	5	5	6	0	5	6	0
Neutral cholesterol ester hydrolase 1 OS=Mus musculus GN=Nceh1 PE=1 SV=1	NCEH1_MOUSE	46 kDa	1	4	0	5	0	4	5	5	0	4	5	0	4
Cytochrome b-c1 complex subunit 9 OS=Mus musculus GN=Uqcrl0 PE=1 SV=1	QCR9_MOUSE	7 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
60S ribosomal protein L28 OS=Mus musculus GN=Rpl28 PE=1 SV=2	RL28_MOUSE	16 kDa	1	4	0	3	0	4	3	3	0	4	3	0	4
60S acidic ribosomal protein P2 OS=Mus musculus GN=Rplp2 PE=1 SV=3	RLA2_MOUSE	12 kDa	1	5	4	0	4	5	0	0	4	5	0	4	5
Sepiapterin reductase OS=Mus musculus GN=Spr PE=1 SV=1	SPRE_MOUSE	28 kDa	1	6	9	7	9	6	7	7	9	6	7	9	6
Mitochondrial import inner membrane translocase subunit Tim9 OS=Mus musculus GN=TIM9_MOUSE		10 kDa	1	5	5	6	5	5	6	6	5	5	6	5	5
DNA topoisomerase 2-binding protein 1 OS=Mus musculus GN=Topbp1 PE=1 SV=2	TOPB1_MOUSE	169 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Hsp90 co-chaperone Cdc37 OS=Mus musculus GN=Cdc37 PE=2 SV=1	CDC37_MOUSE	45 kDa	1	3	3	4	3	3	4	4	3	3	4	3	3
Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1	CO6A1_MOUSE	108 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Catenin alpha-3 OS=Mus musculus GN=Ctnna3 PE=1 SV=2	CTNA3_MOUSE	100 kDa	1	3	6	5	6	3	5	5	6	3	5	6	3
Dnaj homolog subfamily A member 4 OS=Mus musculus GN=Dnaj4 PE=2 SV=1	DNJA4_MOUSE	45 kDa	1	3	7	5	7	3	5	5	7	3	5	7	3
Ubiquitin carboxyl-terminal hydrolase OS=Mus musculus GN=Usp14 PE=2 SV=1	E9PNY8_MOUSE (+1)	52 kDa	1	0	4	4	4	0	4	4	0	4	4	0	4
Protein phosphatase 1 regulatory subunit 7 OS=Mus musculus GN=Ppp1r7 PE=1 SV=2	PP1R7_MOUSE	41 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Peptidyl-prolyl cis-trans isomerase D OS=Mus musculus GN=Ppid PE=1 SV=3	PPID_MOUSE	41 kDa	1	3	4	2	4	3	2	2	4	3	2	4	3
Protein phosphatase 1 regulatory subunit 3A OS=Mus musculus GN=Ppp1r3a PE=1 SV=1	PPR3A_MOUSE	121 kDa	1	4	5	5	5	4	5	5	5	4	5	5	4

Proteasome subunit alpha type-4 OS=Mus musculus GN=Psm4 PE=1 SV=1	PSA4_MOUSE	29 kDa	1	3	0	4	0	3	4	4	0	3	4	0	3
MCG49183 OS=Mus musculus GN=1700009N14Rik PE=2 SV=1	Q14AA6_MOUSE	24 kDa	1	4	6	4	6	4	4	4	6	4	4	6	4
39S ribosomal protein L12, mitochondrial OS=Mus musculus GN=Mrp12 PE=1 SV=2	RM12_MOUSE	22 kDa	1	3	7	7	7	7	7	7	7	7	7	7	7
40S ribosomal protein S10 OS=Mus musculus GN=Rps10 PE=1 SV=1	RS10_MOUSE	19 kDa	1	5	3	0	3	5	0	0	3	5	0	3	5
Mitochondrial import inner membrane translocase subunit Tim10 OS=Mus musculus (TIM10_MOUSE	TIM10_MOUSE	10 kDa	1	6	4	3	4	6	3	3	4	6	3	4	6
Cluster of Alcohol dehydrogenase class-3 OS=Mus musculus GN=Adh5 PE=1 SV=3 (AD_ADHX_MOUSE	ADHX_MOUSE	40 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Cluster of Uncharacterized protein C6orf203 homolog OS=Mus musculus PE=1 SV=1 (C6orf203_MOUSE	C6orf203_MOUSE	28 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of CLIP-associating protein 1 OS=Mus musculus GN=Clasp1 PE=1 SV=2 (CLAP1_CLAP1_MOUSE [4]	CLAP1_MOUSE	169 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
A-kinase anchor protein 12 OS=Mus musculus GN=Akap12 PE=1 SV=1	AKA12_MOUSE	181 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
ATP synthase subunit g, mitochondrial OS=Mus musculus GN=Atp5l PE=1 SV=1	ATP5L_MOUSE	11 kDa	1	5	0	0	0	5	0	0	0	5	0	0	5
Hexaprenyldihydroxybenzoate methyltransferase, mitochondrial OS=Mus musculus G COQ3_MOUSE	COQ3_MOUSE	41 kDa	1	6	6	4	6	6	4	4	6	6	4	6	6
Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Mus musculus GN=Gnb2;GBLP_MOUSE	GBLP_MOUSE	35 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Mus musculus GN=Hnrnpc PE=:HNRPC_MOUSE	HNRPC_MOUSE	34 kDa	1	6	3	3	3	6	3	3	3	6	3	3	6
Heat shock protein beta-7 OS=Mus musculus GN=Hspb7 PE=1 SV=3	HSPB7_MOUSE	19 kDa	1	3	8	7	8	3	7	7	8	3	7	8	3
Cluster of LIM and SH3 domain protein 1 OS=Mus musculus GN=Lasp1 PE=1 SV=1 (LALASASP1_MOUSE	LASP1_MOUSE	30 kDa	1	4	6	3	6	4	3	3	6	4	3	6	4
Methyl-CpG-binding protein 2 OS=Mus musculus GN=Mecp2 PE=1 SV=1	MECP2_MOUSE	52 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Mitochondrial-processing peptidase subunit alpha OS=Mus musculus GN=Pmpca PE=:MPPA_MOUSE	MPPA_MOUSE	58 kDa	1	4	5	5	5	4	5	5	5	5	4	5	5
Na(+)/H(+) exchange regulatory cofactor NHE-RF2 OS=Mus musculus GN=Slc9a3r2 PE: NHRF2_MOUSE	NHRF2_MOUSE	37 kDa	1	4	5	4	5	4	4	4	5	4	4	5	4
Perilipin-3 OS=Mus musculus GN=Plin3 PE=1 SV=1	PLIN3_MOUSE	47 kDa	1	0	4	5	4	0	5	5	4	0	5	4	0
Polyribonucleotide nucleotidyltransferase 1, mitochondrial OS=Mus musculus GN=Pnp1;PNPT1_MOUSE	PNPT1_MOUSE	86 kDa	1	3	0	4	0	3	4	4	0	3	4	0	3
26S protease regulatory subunit 6B OS=Mus musculus GN=Psmc4 PE=1 SV=2	PRSB6_MOUSE	47 kDa	1	3	7	3	7	3	3	3	7	3	3	7	3
60S ribosomal protein L23 OS=Mus musculus GN=Rpl23 PE=1 SV=1	RL23_MOUSE	15 kDa	1	3	6	6	6	3	6	6	6	6	6	6	6
60S ribosomal protein L35a OS=Mus musculus GN=Rpl35a PE=2 SV=2	RL35A_MOUSE	13 kDa	1	3	5	0	5	3	0	0	5	3	0	5	3
Cluster of Septin-11 OS=Mus musculus GN=Sept11 PE=1 SV=4 (SEP11_MOUSE)	SEP11_MOUSE [6]	50 kDa	1	3	3	2	3	3	2	2	3	3	2	3	3
Sulfite oxidase, mitochondrial OS=Mus musculus GN=Suox PE=1 SV=2	SUOX_MOUSE	61 kDa	1	4	3	4	3	4	4	4	3	4	4	3	4
Lysine-tRNA ligase OS=Mus musculus GN=Kars PE=1 SV=1	SYK_MOUSE	68 kDa	1	3	5	4	5	3	4	4	5	3	4	5	3
Telethonin OS=Mus musculus GN=Tcap PE=2 SV=1	TEL_MOUSE	19 kDa	1	5	7	6	5	7	7	7	6	5	7	6	5
Cluster of Heterogeneous nuclear ribonucleoprotein A3 OS=Mus musculus GN=Hnrnp A2AL12_MOUSE	A2AL12_MOUSE	34 kDa	1	2	5	8	5	8	8	8	5	2	8	5	2
Cluster of Isobutyryl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Acad4;D3YTT4_MOUSE [2]	D3YTT4_MOUSE [2]	45 kDa	1	0	6	8	6	0	8	8	6	0	8	6	0
Cysteine-rich protein 1 OS=Mus musculus GN=Crip1 PE=2 SV=2	CRIP1_MOUSE	9 kDa	1	0	0	4	0	4	4	4	0	4	0	4	0
Cytoplasmic dynein 1 light intermediate chain 1 OS=Mus musculus GN=Dync1li1 PE=1 SV=1 (DC1L1_MOUSE	DC1L1_MOUSE	11 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Protein-L-isoaspartate O-methyltransferase OS=Mus musculus GN=Pcmt1 PE=2 SV=1	EOCV0_MOUSE	30 kDa	1	0	6	6	6	0	6	6	6	0	6	6	0
Glutaredoxin-3 OS=Mus musculus GN=Glxr3 PE=1 SV=1	GLRX3_MOUSE	38 kDa	1	3	6	3	6	3	3	3	6	3	3	6	3
Histidine triad nucleotide-binding protein 1 OS=Mus musculus GN=Hint1 PE=1 SV=3	HINT1_MOUSE	14 kDa	1	3	0	4	0	3	4	4	0	3	4	0	3
Monocarboxylate transporter 1 OS=Mus musculus GN=Slc16a1 PE=1 SV=1	MOT1_MOUSE	53 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Myopalladin OS=Mus musculus GN=Mypn PE=2 SV=2	MYPN_MOUSE	144 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Programmed cell death protein 5 OS=Mus musculus GN=Pcd5 PE=1 SV=3	PDCD5_MOUSE	14 kDa	1	5	3	4	3	5	4	4	3	5	4	3	5
Bisphosphoglycerate mutase OS=Mus musculus GN=Bpgm PE=2 SV=2	PMGE_MOUSE	30 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Rab11 family-interacting protein 5 OS=Mus musculus GN=Rab11fp5 PE=1 SV=2	RFIP5_MOUSE	70 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of 60S ribosomal protein L18a OS=Mus musculus GN=Rpl18a PE=1 SV=1 (RL18_MOUSE)	RL18_MOUSE	21 kDa	1	3	0	5	0	3	5	5	0	3	5	0	3
40S ribosomal protein S13 OS=Mus musculus GN=Rps13 PE=1 SV=2	RS13_MOUSE	17 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
40S ribosomal protein S14 OS=Mus musculus GN=Rps14 PE=2 SV=3	RS14_MOUSE	16 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
40S ribosomal protein S19 OS=Mus musculus GN=Rps19 PE=1 SV=3	RS19_MOUSE	16 kDa	1	3	3	3	3	3	3	3	3	3	3	3	3
Glycine-tRNA ligase OS=Mus musculus GN=Gars PE=1 SV=1	SYG_MOUSE	82 kDa	1	3	4	4	4	3	4	4	4	3	4	4	3
Translationally-controlled tumor protein OS=Mus musculus GN=Tpt1 PE=1 SV=1	TCTP_MOUSE	19 kDa	1	4	3	0	3	4	0	0	3	4	0	3	4
40S ribosomal protein S24 OS=Mus musculus GN=Rps24 PE=1 SV=1	RS24_MOUSE (+2)	15 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Cluster of AP-2 complex subunit beta OS=Mus musculus GN=Ap2b1 PE=1 SV=1 (AP2B_MOUSE)	AP2B_MOUSE [5]	105 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Cluster of MCG19772 OS=Mus musculus GN=Trim55 PE=4 SV=1 (G3X8Y1_MOUSE)	G3X8Y1_MOUSE [3]	60 kDa	1	0	0	6	0	0	6	6	0	0	6	0	0
Cluster of Protein 4.1 OS=Mus musculus GN=Epb41 PE=1 SV=2 (41_MOUSE)	41_MOUSE [3]	96 kDa	1	0	3	5	3	0	5	5	3	0	5	3	0
Cluster of S-phase kinase-associated protein 1 OS=Mus musculus GN=Skep1 PE=1 SV=3 (SKP1_MOUSE)	SKP1_MOUSE [2]	19 kDa	1	5	3	0	3	5	0	0	3	5	0	3	5
Alpha-1-syntrophin OS=Mus musculus GN=Snta1 PE=2 SV=1	AZAKD7_MOUSE	53 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Apolipoprotein O-like OS=Mus musculus GN=Apool PE=2 SV=1	APOOL_MOUSE	29 kDa	1	0	0	6	0	0	6	6	0	0	6	0	0
Bcl-2-like protein 13 OS=Mus musculus GN=Bcl2l13 PE=1 SV=2	B2L13_MOUSE	47 kDa	1	5	5	0	5	5	0	0	5	5	0	5	5
Adenylyl cyclase-associated protein 2 OS=Mus musculus GN=Cap2 PE=1 SV=1	CAP2_MOUSE (+1)	53 kDa	1	3	6	5	6	3	5	5	6	3	5	6	3
Cathepsin B OS=Mus musculus GN=Ctsb PE=1 SV=2	CATB_MOUSE	37 kDa	1	4	5	0	5	4	0	0	5	4	0	5	4
Collagen alpha-1(I) chain OS=Mus musculus GN=Col1a1 PE=1 SV=4	CO1A1_MOUSE	138 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
2-methoxy-6-polypropenyl-1,4-benzoquinol methylase, mitochondrial OS=Mus musculus COO5_MOUSE	COO5_MOUSE	37 kDa	1	7	6	4	6	7	4	4	6	7	4	6	7
CapZ-interacting protein OS=Mus musculus GN=Rcsd1 PE=2 SV=1	CPZIP_MOUSE	44 kDa	1	0	5	6	5	0	6	6	5	0	6	5	0
Dynactin subunit 1 OS=Mus musculus GN=Dctn1 PE=1 SV=3	DCTN1_MOUSE	142 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Probable rRNA-processing protein EBP2 OS=Mus musculus GN=Ebna1bp2 PE=2 SV=1	EBP2_MOUSE	35 kDa	1	4	0	0	0	4	0	0	4	0	0	4	0
Glyoxylate reductase/hydroxypropylate reductase OS=Mus musculus GN=Grhpr PE=1 SV=1 (GRHPR_MOUSE	GRHPR_MOUSE	35 kDa	1	4	4	0	4	4	0	0	4	4	0	4	4
Kynurenine--oxoglutarate transaminase 3 OS=Mus musculus GN=Ccbl2 PE=1 SV=1	KAT3_MOUSE	51 kDa	1	3	0	3	0	3	3	3	0	3	3	0	3
NADP-dependent malic enzyme OS=Mus musculus GN=Me1 PE=1 SV=2	MAOX_MOUSE	64 kDa	1	3	5	5	5	3	5	5	5	3	5	5	3
Endonuclease G, mitochondrial OS=Mus musculus GN=Endog PE=2 SV=1	NUCG_MOUSE	32 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Cluster of Oxidoreductase NAD-binding domain-containing protein 1 OS=Mus musculus OXND1_MOUSE	OXND1_MOUSE	35 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
[Pyruvate dehydrogenase [lipoamide]] kinase isozyme 1, mitochondrial OS=Mus musculus PDK1_MOUSE	PDK1_MOUSE	49 kDa	1	5	0	0	0	5	0	0	0	5	0	0	5

Peptidyl-prolyl cis-trans isomerase B OS=Mus musculus GN=Ppib PE=2 SV=2	PPIB_MOUSE	24 kDa	1	5	5	4	5	5	4	4	5	5	4	5	5
Presequence protease, mitochondrial OS=Mus musculus GN=Pitrm1 PE=2 SV=1	PREP_MOUSE (+1)	117 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
26S proteasome non-ATPase regulatory subunit 3 OS=Mus musculus GN=PsmD3 PE=1	PSMD3_MOUSE	61 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Protein Prob1 (Fragment) OS=Mus musculus GN=Prob1 PE=2 SV=1	Q3UKG2_MOUSE	106 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
60S ribosomal protein L18 OS=Mus musculus GN=Rpl18 PE=2 SV=3	RL18_MOUSE	22 kDa	1	3	5	5	5	3	5	5	5	3	5	5	3
Heterogeneous nuclear ribonucleoprotein A/B OS=Mus musculus GN=Hnrrnpab PE=1	ROAA_MOUSE	31 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
28S ribosomal protein S36, mitochondrial OS=Mus musculus GN=Mrps36 PE=1 SV=1	RT36_MOUSE	11 kDa	1	7	4	4	4	7	4	4	4	7	4	4	7
Transgelin-2 OS=Mus musculus GN=Tagln2 PE=1 SV=4	TAGL2_MOUSE	22 kDa	1	0	5	5	5	0	5	5	5	0	5	5	0
Vesicle-associated membrane protein-associated protein A OS=Mus musculus GN=Va1VAPA_MOUSE	VAPA_MOUSE	28 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Vacuolar protein sorting-associated protein 35 OS=Mus musculus GN=Vps35 PE=1 SV=1	VPS35_MOUSE	92 kDa	1	6	4	4	4	6	4	4	4	6	4	4	6
Zyxin OS=Mus musculus GN=Zyx PE=1 SV=2	ZYX_MOUSE	61 kDa	1	0	6	4	6	0	4	4	6	0	4	6	0
Serine/threonine-protein kinase mTOR OS=Mus musculus GN=Mtor PE=1 SV=2	MTOR_MOUSE	289 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Reticulon-4-interacting protein 1, mitochondrial OS=Mus musculus GN=Rtn4ip1 PE=1	RT41_MOUSE	43 kDa	1	4	3	3	3	4	3	3	3	4	3	3	4
Fatty acid-binding protein, epidermal OS=Mus musculus GN=Fabp5 PE=1 SV=3	FABP5_MOUSE	15 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Cluster of ADP-ribosylation factor 3 OS=Mus musculus GN=Arf3 PE=2 SV=2 (ARF3_MOUSE)	ARF3_MOUSE	21 kDa	1	0	0	4	0	4	0	4	0	0	4	0	0
Cluster of Protein Zim1 OS=Mus musculus GN=Zim1 PE=2 SV=1 (Q8C393_MOUSE)	Q8C393_MOUSE	65 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Alcohol dehydrogenase [NADP(+)] OS=Mus musculus GN=Akr1a1 PE=1 SV=3	AK1A1_MOUSE	37 kDa	1	5	5	0	5	5	0	0	5	5	0	5	5
Annexin A7 OS=Mus musculus GN=Anxa7 PE=2 SV=2	ANXA7_MOUSE	50 kDa	1	0	4	5	4	0	5	5	4	0	5	4	0
Aflatoxin B1 aldehyde reductase member 2 OS=Mus musculus GN=Akr7a2 PE=1 SV=3	ARK72_MOUSE	41 kDa	1	0	7	3	7	0	3	3	7	0	3	7	0
Cellular nucleic acid-binding protein OS=Mus musculus GN=Cnbp PE=2 SV=2	CNBP_MOUSE	20 kDa	1	0	5	6	5	0	6	6	5	0	6	5	0
Cluster of Anamorsin OS=Mus musculus GN=Ciapi1 PE=1 SV=1 (CPIN1_MOUSE)	CPIN1_MOUSE [2]	33 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
Peroxisomal multifunctional enzyme type 2 OS=Mus musculus GN=Hsd17b4 PE=1 SV=1	DHB4_MOUSE	79 kDa	1	0	0	0	3	0	0	3	0	0	3	0	0
Eukaryotic translation initiation factor 3 subunit J-A OS=Mus musculus GN=Eif3j1 PE=1	E13J1_MOUSE (+1)	29 kDa	1	0	3	5	3	0	5	5	3	0	5	3	0
Protein FAM136A OS=Mus musculus GN=Fam136a PE=1 SV=1	F136A_MOUSE	16 kDa	1	4	3	3	3	4	3	3	3	4	3	3	4
Protocadherin Fat 4 OS=Mus musculus GN=Fat4 PE=1 SV=2	FAT4_MOUSE	540 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Acyl-protein thioesterase 1 OS=Mus musculus GN=Lyp1a1 PE=4 SV=1	J3QP56_MOUSE (+1)	24 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
Adenylate kinase isoenzyme 4, mitochondrial OS=Mus musculus GN=Ak4 PE=2 SV=1	KAD4_MOUSE	25 kDa	1	6	5	3	5	6	3	3	5	6	3	5	6
Leukotriene A-4 hydrolase OS=Mus musculus GN=Lta4h PE=1 SV=4	LKHA4_MOUSE	69 kDa	1	3	7	6	7	3	6	6	7	3	6	7	3
Protein Sptbn2 OS=Mus musculus GN=Sptbn2 PE=2 SV=1	Q68FG2_MOUSE	271 kDa	1	0	0	0	0	0	0	0	0	0	0	0	0
Ankyrin OS=Mus musculus GN=Rai14 PE=1 SV=1	RAI14_MOUSE	109 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
60S ribosomal protein L37 OS=Mus musculus GN=Rpl37 PE=2 SV=3	RL37_MOUSE	11 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
28S ribosomal protein S22, mitochondrial OS=Mus musculus GN=Mrps22 PE=2 SV=1	RT22_MOUSE	41 kDa	1	3	3	3	3	3	3	3	3	3	3	3	3
Gamma-synuclein OS=Mus musculus GN=Sncg PE=1 SV=1	SYUG_MOUSE	13 kDa	1	3	0	4	0	3	4	4	0	3	4	0	3
Cluster of Hydroxymethylglutaryl-CoA lyase, mitochondrial OS=Mus musculus GN=Hm1HMGCL_MOUSE	HMGCL_MOUSE	34 kDa	1	4	4	0	4	4	0	4	4	0	4	4	4
Myosin Vb, isoform CRA_a OS=Mus musculus GN=Myo5b PE=4 SV=1	G3X9Y9_MOUSE (+2)	213 kDa	1	1	0	0	0	1	0	0	0	1	0	0	1
Cluster of Tripeptidyl-peptidase 2 OS=Mus musculus GN=Tpp2 PE=2 SV=3 (TPP2_MOUSE)	TPP2_MOUSE	140 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Cluster of Heat shock 70 kDa protein 12B OS=Mus musculus GN=Hspa12b PE=1 SV=1 (HS12B_MOUSE)	HS12B_MOUSE	76 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cluster of Cordon-bleu protein-like 1 OS=Mus musculus GN=Cobl1 PE=4 SV=1 (B1AZ1_B1AZ14_MOUSE [2])	B1AZ1_MOUSE [2]	133 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Proteasomal ubiquitin receptor ADRM1 OS=Mus musculus GN=Adrm1 PE=1 SV=2	ADRM1_MOUSE	42 kDa	1	0	5	6	5	0	6	6	5	0	6	5	0
Afamin OS=Mus musculus GN=Afm PE=1 SV=2	AFAM_MOUSE	69 kDa	1	0	6	4	6	0	4	4	6	0	4	6	0
Isoaspartyl peptidase/L-asparaginase OS=Mus musculus GN=Asrgl1 PE=1 SV=1	ASGL1_MOUSE	34 kDa	1	3	5	5	5	3	5	5	3	5	3	5	3
C-1-tetrahydrofolate synthase, cytoplasmic OS=Mus musculus GN=Mthfd1 PE=1 SV=4	CLTC_MOUSE	101 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
UPF0598 protein C8orf82 homolog OS=Mus musculus PE=2 SV=1	CH082_MOUSE	24 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
Putative ATP-dependent Clp protease proteolytic subunit, mitochondrial OS=Mus musculus GN=CLPP_MOUSE	CLPP_MOUSE	30 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Dynactin subunit 2 OS=Mus musculus GN=Dctn2 PE=1 SV=3	DCTN2_MOUSE	44 kDa	1	0	5	5	5	0	5	5	5	0	5	5	0
Dnal homolog subfamily A member 3, mitochondrial OS=Mus musculus GN=Dnaja3 PIDNJA3_MOUSE	PIDNJA3_MOUSE	52 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Dystrobrevin alpha OS=Mus musculus GN=Dtna PE=1 SV=2	DTNA_MOUSE	84 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
Glycerol-3-phosphate dehydrogenase [NAD(+)], cytoplasmic OS=Mus musculus GN=GjGPDA_MOUSE	GjGPDA_MOUSE	38 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Glutathione reductase, mitochondrial OS=Mus musculus GN=Gsr PE=2 SV=3	GSHR_MOUSE (+1)	54 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
High mobility group protein HMG-1/HMG-1 OS=Mus musculus GN=Hmga1 PE=1 SV=4	HMGA1_MOUSE	12 kDa	1	4	4	5	4	4	5	5	4	4	5	4	4
Protein Dnahc7b OS=Mus musculus GN=Dnahc7b PE=4 SV=1	L7N1Y0_MOUSE	467 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
NAD(P)H dehydrogenase [quinone] 1 OS=Mus musculus GN=Nqo1 PE=1 SV=3	NQO1_MOUSE	31 kDa	1	3	7	4	7	3	4	4	7	3	4	7	3
Prolyl endopeptidase OS=Mus musculus GN=Prep PE=2 SV=1	PPCE_MOUSE	81 kDa	1	0	7	3	7	0	3	3	7	0	3	7	0
Proline synthase co-transcribed bacterial homolog protein OS=Mus musculus GN=Pro PROSC_MOUSE	PROSC_MOUSE	30 kDa	1	0	5	0	5	0	0	0	5	0	0	5	0
Vesicle-associated membrane protein, associated protein B and C OS=Mus musculus GN=Q8BH80_MOUSE (+1)	Q8BH80_MOUSE (+1)	27 kDa	1	4	2	4	2	4	4	4	2	4	4	2	4
Cortactin, isoform CRA_a OS=Mus musculus GN=Cttn PE=2 SV=1	Q921L6_MOUSE (+1)	57 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
60S ribosomal protein L21 OS=Mus musculus GN=Rpl21 PE=2 SV=1	Q9CQM8_MOUSE	19 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
60S ribosomal protein L26 OS=Mus musculus GN=Rpl26 PE=2 SV=1	RL26_MOUSE	17 kDa	1	4	0	7	0	4	7	7	0	4	7	0	4
60S ribosomal protein L27a OS=Mus musculus GN=Rpl27a PE=2 SV=5	RL27A_MOUSE	17 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
40S ribosomal protein S16 OS=Mus musculus GN=Rps16 PE=2 SV=4	RS16_MOUSE	16 kDa	1	4	0	0	0	4	0	0	4	0	4	0	4
Splicing factor, proline- and glutamine-rich OS=Mus musculus GN=Sfpq PE=1 SV=1	SFPQ_MOUSE	75 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Serine/arginine-rich splicing factor 7 OS=Mus musculus GN=Srsf7 PE=1 SV=1	SRSF7_MOUSE (+2)	31 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Ubiquitin-conjugating enzyme E2 L3 OS=Mus musculus GN=Ube2l3 PE=2 SV=1	UB2L3_MOUSE	18 kDa	1	4	5	4	5	4	5	5	4	4	5	4	4
Cluster of 60S ribosomal protein L17 OS=Mus musculus GN=Rpl17 PE=2 SV=1 (Q6ZWZ_Q6ZWZ2_MOUSE)	Q6ZWZ2_MOUSE	21 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Cluster of Collagen alpha-1(XV) chain OS=Mus musculus GN=Col15a1 PE=2 SV=1 (A2A_A2AJY2_MOUSE [3])	A2AJY2_MOUSE [3]	138 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Adiponectin OS=Mus musculus GN=Adipoq PE=1 SV=2	ADIPO_MOUSE	27 kDa	1	0	5	5	5	0	5	5	5	0	5	5	0

Cluster of Plastin-3 OS=Mus musculus GN=Pls3 PE=2 SV=1 (B1AX58_MOUSE)	B1AX58_MOUSE [3]	72 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Valacyclovir hydrolase OS=Mus musculus GN=Bphl PE=2 SV=1	BPHL_MOUSE	33 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cadherin-2 OS=Mus musculus GN=Cdh2 PE=1 SV=2	CADH2_MOUSE (+1)	100 kDa	1	3	6	2	6	3	2	2	6	3	2	6	3
Cullin-associated NEDD8-dissociated protein 1 OS=Mus musculus GN=Cand1 PE=2 SV=2	CAND1_MOUSE	136 kDa	1	0	5	0	5	0	0	0	5	0	0	5	0
Carbonyl reductase [NADPH] 1 OS=Mus musculus GN=Cbr1 PE=1 SV=3	CBR1_MOUSE	31 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Cytochrome c oxidase subunit 7A-related protein, mitochondrial OS=Mus musculus GN=COX7R_MOUSE (+1)	COX7R_MOUSE (+1)	12 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
Catenin beta-1 OS=Mus musculus GN=Ctnnb1 PE=1 SV=1	CTNB1_MOUSE	85 kDa	1	2	3	5	3	2	5	5	3	2	5	3	2
Microtubule-associated protein RP/EB family member 2 (Fragment) OS=Mus musculus GN=D3YK8_MOUSE (+2)	D3YK8_MOUSE (+2)	29 kDa	1	4	3	5	3	4	5	5	3	4	5	3	4
DNA damage-binding protein 1 OS=Mus musculus GN=Ddb1 PE=1 SV=2	DDB1_MOUSE	127 kDa	1	0	4	5	4	0	5	5	4	0	5	4	0
Inter alpha-trypsin inhibitor, heavy chain 4 OS=Mus musculus GN=Ith4 PE=2 SV=1	E9PVD2_MOUSE (+2)	105 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
26S protease regulatory subunit 10B OS=Mus musculus GN=Psmc6 PE=1 SV=1	PRS10_MOUSE	44 kDa	1	4	5	4	5	4	4	4	5	4	4	5	4
60S ribosomal protein L9 OS=Mus musculus GN=Rpl9 PE=2 SV=2	RL9_MOUSE	22 kDa	1	0	4	2	4	0	2	2	4	0	2	4	0
SRA stem-loop-interacting RNA-binding protein, mitochondrial OS=Mus musculus GN=SLIRP_MOUSE	SLIRP_MOUSE	13 kDa	1	4	5	0	5	4	0	0	5	4	0	5	4
Transmembrane protein 109 OS=Mus musculus GN=Tmem109 PE=1 SV=2	TM109_MOUSE	26 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Tubulin polymerization-promoting protein family member 3 OS=Mus musculus GN=TPPP3_MOUSE	TPPP3_MOUSE	19 kDa	1	5	3	0	3	5	0	0	3	5	0	3	5
Cluster of Hepatoma-derived growth factor OS=Mus musculus GN=HdGF PE=1 SV=2 (HHDGF_MOUSE [2])	HHDGF_MOUSE [2]	26 kDa	1	2	4	5	4	2	5	5	4	2	5	4	2
Alpha-2-antiplasmin OS=Mus musculus GN=Serpinf2 PE=1 SV=1	A2AP_MOUSE	55 kDa	1	5	0	3	0	5	3	3	0	5	3	0	5
Apolipoprotein B-100 OS=Mus musculus GN=ApoB PE=1 SV=1	APOB_MOUSE	509 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Tight junction protein ZO-1 OS=Mus musculus GN=Tjp1 PE=2 SV=1	B9EHJ3_MOUSE	189 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Filamin-binding LIM protein 1 OS=Mus musculus GN=Fblm1 PE=1 SV=2	FBL1_MOUSE	41 kDa	1	4	4	0	4	0	0	0	4	4	0	4	4
Inorganic pyrophosphatase OS=Mus musculus GN=Ppa1 PE=1 SV=1	IPYR_MOUSE	33 kDa	1	3	3	3	3	3	3	3	3	3	3	3	3
UMP-CMP kinase OS=Mus musculus GN=Cmpk1 PE=1 SV=1	KCY_MOUSE	22 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Proteasome subunit alpha type-6 OS=Mus musculus GN=PsmA6 PE=1 SV=1	PSA6_MOUSE	27 kDa	1	3	4	3	4	3	3	3	4	3	3	4	3
Protein Tfg OS=Mus musculus GN=Tfg PE=2 SV=1	Q9Z1A1_MOUSE	43 kDa	1	4	3	0	3	4	0	0	3	4	0	3	4
Cluster of Isoform 2 of Dynamin 2 OS=Mus musculus GN=Dnm2 (sp P39054-2 DYN2_sp P39054-2 DYN2_MOUSE)?	P39054-2_MOUSE	27 kDa	1	2	2	2	2	2	2	2	2	2	2	2	2
Calycalin-binding protein OS=Mus musculus GN=Cacybp PE=1 SV=1	CYBP_MOUSE	27 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1 (ITB1_MOUSE)	ITB1_MOUSE	88 kDa	1	0	0	4	0	0	0	0	4	0	0	4	0
Cluster of Ig gamma-2B chain C region OS=Mus musculus GN=Igh-3 PE=1 SV=3 (IGG2B_IGG2B_MOUSE)	IGG2B_MOUSE	44 kDa	1	0	3	6	3	0	6	6	3	0	6	3	0
Cluster of Ribosomal protein L15 OS=Mus musculus GN=Gm10020 PE=3 SV=1 (E9QAZ:E9QAZ2_MOUSE [2])	E9QAZ2_MOUSE [2]	24 kDa	1	3	4	0	4	3	0	0	4	3	0	4	3
6-phosphogluconate dehydrogenase, decarboxylating OS=Mus musculus GN=Pgd PE=6 PGD_MOUSE	PGD_MOUSE	53 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Annexin A11 OS=Mus musculus GN=Anxa11 PE=1 SV=2	ANX11_MOUSE	54 kDa	1	0	4	7	4	0	7	7	4	0	7	4	0
Clusterin OS=Mus musculus GN=Clu PE=1 SV=1	CLUS_MOUSE	52 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Mus musculus GN=Fkbp4 PE=1 SV=5	FKBP4_MOUSE	52 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
FUN14 domain-containing protein 2 OS=Mus musculus GN=Fundc2 PE=2 SV=1	FUND2_MOUSE	17 kDa	1	4	3	0	3	4	0	0	3	4	0	3	4
Fragile X mental retardation syndrome-related protein 1 OS=Mus musculus GN=Fxr1 FXR1_MOUSE	FXR1_MOUSE	76 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Ras GTPase-activating protein-binding protein 2 OS=Mus musculus GN=G3bp2 PE=1 S'G3BP2_MOUSE (+1)	G3BP2_MOUSE (+1)	54 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Mimecan OS=Mus musculus GN=Ogn PE=2 SV=1	MIME_MOUSE	34 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
E3 ubiquitin-protein ligase NEDD4 OS=Mus musculus GN=Nedd4 PE=1 SV=3	NEDD4_MOUSE	103 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Protein kinase C and casein kinase substrate in neurons protein 2 OS=Mus musculus GN=CACN2_MOUSE	CACN2_MOUSE	56 kDa	1	4	0	4	0	4	4	4	0	4	0	4	0
Parathyroid hormone-related protein OS=Mus musculus GN=Pthrh PE=2 SV=3	PTHRH_MOUSE	11 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Acp1 protein OS=Mus musculus GN=Acp1 PE=1 SV=1	Q561M1_MOUSE	18 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Ribonuclease inhibitor OS=Mus musculus GN=Rnh1 PE=1 SV=1	RINI_MOUSE	50 kDa	1	0	7	5	7	0	5	5	7	0	5	7	0
Spartin OS=Mus musculus GN=Spg20 PE=2 SV=1	SPG20_MOUSE	73 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Slc8a1 protein OS=Mus musculus GN=Slc8a1 PE=2 SV=1	Q68F10_MOUSE	107 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Glutathione peroxidase 3 OS=Mus musculus GN=Gpx3 PE=2 SV=2	GPX3_MOUSE	25 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Microtubule-associated protein 6 OS=Mus musculus GN=Map6 PE=1 SV=2	MAP6_MOUSE	96 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Sorting nexin 3 OS=Mus musculus GN=Snx3 PE=2 SV=1	Q78ZM0_MOUSE (+1)	19 kDa	1	3	5	4	5	3	4	4	5	3	4	5	3
26S protease regulatory subunit 8 OS=Mus musculus GN=Psmc5 PE=1 SV=1	PRS8_MOUSE	46 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Cysteine desulfurase, mitochondrial OS=Mus musculus GN=Nfs1 PE=2 SV=2 NFS1_MOUSE	NFS1_MOUSE	50 kDa	1	0	0	5	0	0	5	5	0	0	5	0	0
3-hydroxyisobutyrate dehydrogenase, mitochondrial OS=Mus musculus GN=Hibadh P3HI0H_MOUSE	P3HI0H_MOUSE	35 kDa	1	0	0	5	0	0	5	0	0	5	0	0	5
Growth factor receptor-bound protein 2 OS=Mus musculus GN=Grb2 PE=2 SV=1	B1AT92_MOUSE (+1)	24 kDa	1	4	3	3	3	4	3	3	3	4	3	3	4
Complement factor I OS=Mus musculus GN=Cfi PE=1 SV=3	CFAI_MOUSE	67 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cytoskeleton-associated protein 4 OS=Mus musculus GN=Ckap4 PE=2 SV=2	CKAP4_MOUSE	64 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Desmoglein-2 OS=Mus musculus GN=Dsg2 PE=1 SV=3	DSG2_MOUSE	122 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Epoxide hydrolase 1 OS=Mus musculus GN=Ephx1 PE=2 SV=1	E9PWK1_MOUSE (+1)	51 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Eukaryotic translation initiation factor 3 subunit C OS=Mus musculus GN=Eif3c PE=1 S EIF3C_MOUSE	EIF3C_MOUSE	106 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Protein ETHE1, mitochondrial OS=Mus musculus GN=Ethe1 PE=1 SV=2	ETHE1_MOUSE	28 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Eukaryotic translation initiation factor 2 subunit 1 OS=Mus musculus GN=Eif2s1 PE=1 IF2A_MOUSE	IF2A_MOUSE	36 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Platelet-activating factor acetylhydrolase 1B subunit alpha OS=Mus musculus GN=PafaL1S1_MOUSE	PafaL1S1_MOUSE	47 kDa	1	0	3	5	3	0	5	5	3	0	5	3	0
Palladin OS=Mus musculus GN=Palld PE=1 SV=2	PALLD_MOUSE (+1)	152 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Prostaglandin E synthase 2 OS=Mus musculus GN=Ptges2 PE=1 SV=3	PGES2_MOUSE	43 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cluster of 26S proteasome non-ATPase regulatory subunit 13 OS=Mus musculus GN=PSD13_MOUSE	PSD13_MOUSE	43 kDa	1	4	5	4	5	4	4	4	5	4	4	5	4
Target of Myb protein 1 OS=Mus musculus GN=Tom1 PE=2 SV=1	Q3UDC3_MOUSE	57 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
60S ribosomal protein L36a OS=Mus musculus GN=Rpl36a PE=2 SV=2	RL36A_MOUSE	12 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Small glutamine-rich tetratricopeptide repeat-containing protein alpha OS=Mus musculus GN=SGTA_MOUSE (+1)	SGTA_MOUSE (+1)	34 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Serine/arginine-rich splicing factor 3 OS=Mus musculus GN=Srsf3 PE=1 SV=1	SRSF3_MOUSE	19 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0

Carbonyl reductase family member 4 OS=Mus musculus GN=Cbr4 PE=2 SV=2	CBR4_MOUSE	25 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Epimerase family protein SDR39U1 OS=Mus musculus GN=Sdr39u1 PE=1 SV=1	D39U1_MOUSE (+1)	33 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Cluster of 60S ribosomal protein L35 OS=Mus musculus GN=Rpl35 PE=2 SV=1 (RL35_N RL35_MOUSE		15 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Cluster of Heterogeneous nuclear ribonucleoprotein D0 OS=Mus musculus GN=Hnrnp HNRPD_MOUSE [4]		38 kDa	1	0	7	0	7	0	0	0	7	0	0	7	0
Cluster of SH3 domain-binding glutamic acid-rich protein OS=Mus musculus GN=Sh3b SH3BG_MOUSE		23 kDa	1	0	3	0	3	0	0	0	0	0	0	3	0
Tyrosine--tRNA ligase, cytoplasmic OS=Mus musculus GN=Yars PE=4 SV=1	A2A757_MOUSE (+1)	63 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
Annexin A3 OS=Mus musculus GN=Anxa3 PE=1 SV=4	ANXA3_MOUSE	36 kDa	1	3	2	2	2	3	2	2	2	3	2	2	3
Protein Gm20547 OS=Mus musculus GN=Gm20547 PE=2 SV=1	B8JJNO_MOUSE (+1)	142 kDa	1	5	3	0	3	5	0	0	3	5	0	3	5
Corticosteroid-binding globulin OS=Mus musculus GN=Serpina6 PE=1 SV=1	CBG_MOUSE	45 kDa	1	3	0	4	0	3	4	4	0	3	4	0	3
Chloride intracellular channel protein 4 OS=Mus musculus GN=Clic4 PE=1 SV=3	CLIC4_MOUSE	29 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Glycogen [starch] synthase, muscle OS=Mus musculus GN=Gys1 PE=2 SV=1	D3Z0Q6_MOUSE (+1)	77 kDa	1	3	2	3	2	3	3	3	2	3	3	2	3
Protein Coq10a OS=Mus musculus GN=Coq10a PE=2 SV=1	E9Q3H6_MOUSE	29 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Elongation factor Ts, mitochondrial OS=Mus musculus GN=Tsfm PE=2 SV=1	EFTS_MOUSE	35 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
RNA-binding protein FUS (Fragment) OS=Mus musculus GN=Fus PE=4 SV=1	G3UXT7_MOUSE	14 kDa	1	3	4	3	4	3	3	3	4	3	3	4	3
Valine--tRNA ligase (Fragment) OS=Mus musculus GN=Vars PE=2 SV=1	G3UY93_MOUSE	141 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Inter-alpha trypsin inhibitor, heavy chain 2 OS=Mus musculus GN=Itih2 PE=4 SV=1	G3X977_MOUSE (+1)	106 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Matrin-3 OS=Mus musculus GN=Matr3 PE=1 SV=1	MATR3_MOUSE	95 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Ribosylidihydroxynicotinamide dehydrogenase [quinone] OS=Mus musculus GN=Nqo2 P	NQO2_MOUSE	26 kDa	1	3	3	4	3	3	4	4	3	3	4	3	3
Prefoldin subunit 2 OS=Mus musculus GN=Pfdn2 PE=2 SV=2	PF2D2_MOUSE	17 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Proteasome subunit beta type-6 OS=Mus musculus GN=Psm6 PE=1 SV=3	PSB6_MOUSE	25 kDa	1	0	3	3	3	0	3	3	0	3	0	3	0
Multifunctional protein ADE2 OS=Mus musculus GN=Paics PE=1 SV=4	PUR6_MOUSE	47 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
40S ribosomal protein S21 OS=Mus musculus GN=Rps21 PE=2 SV=1	RS21_MOUSE	9 kDa	1	0	3	6	3	0	6	6	3	0	6	3	0
40S ribosomal protein SA OS=Mus musculus GN=Rpsa PE=1 SV=4	RSSA_MOUSE	33 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
NAD-dependent protein deacetylase sirtuin-5, mitochondrial OS=Mus musculus GN=Sirt5	SIRT5_MOUSE	34 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
T-complex protein 1 subunit gamma OS=Mus musculus GN=Cct3 PE=1 SV=1	TCPG_MOUSE	61 kDa	1	4	4	3	4	4	0	4	3	4	4	3	4
Hematological and neurological expressed 1 protein OS=Mus musculus GN=Hn1 PE=1	HN1_MOUSE	16 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
28S ribosomal protein S5, mitochondrial OS=Mus musculus GN=Mrps5 PE=2 SV=1	RT05_MOUSE	48 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Adrenodoxin, mitochondrial OS=Mus musculus GN=Fdx1 PE=2 SV=1	ADX_MOUSE	20 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Protein 1300017J02Rik OS=Mus musculus GN=1300017J02Rik PE=2 SV=1	D3Y36_MOUSE (+1)	69 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Dynein light chain 2, cytoplasmic OS=Mus musculus GN=Dynll2 PE=1 SV=1 (DYL2_MOUSE [2]		10 kDa	1	0	5	0	5	0	0	0	5	0	0	5	0
Cluster of Mitochondrial import inner membrane translocase subunit Tim8 A OS=Mus musculus	MM8_MOUSE	11 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cytoplasmic dynein 1 intermediate chain 2 OS=Mus musculus GN=Dync1i2 PE=2 SV=1 A2BFF5_MOUSE (+4)		71 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Flavin reductase (NADPH) OS=Mus musculus GN=Blvrb PE=2 SV=3	BLVRB_MOUSE	22 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Adenylyl cyclase-associated protein 1 OS=Mus musculus GN=Cap1 PE=1 SV=4	CAP1_MOUSE	52 kDa	1	0	5	3	5	0	3	3	5	0	3	5	0
Enoyl-CoA hydratase domain-containing protein 2, mitochondrial (Fragment) OS=Mus musculus	F6Z5N5_MOUSE (+1)	13 kDa	1	0	0	4	0	0	4	4	0	4	0	4	0
High mobility group protein B2 OS=Mus musculus GN=Hmgb2 PE=1 SV=3	HMG2_MOUSE	24 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Heat shock protein 105 kDa OS=Mus musculus GN=Hsph1 PE=1 SV=2	HS105_MOUSE (+1)	96 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Importin subunit beta-1 OS=Mus musculus GN=Kpnb1 PE=1 SV=2	IMB1_MOUSE	97 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Cob(II)yrinic acid a,c-diamide adenosyltransferase, mitochondrial OS=Mus musculus G	MMAB_MOUSE	26 kDa	1	0	4	0	0	4	0	4	0	0	4	0	0
Non-POU domain-containing octamer-binding protein OS=Mus musculus GN=Nono P	NONO_MOUSE	55 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Nucleosome assembly protein 1-like 4 OS=Mus musculus GN=Nap1l4 PE=1 SV=1	NP1L4_MOUSE	43 kDa	1	2	4	0	4	2	0	0	4	2	0	4	2
Nuclear migration protein nudC OS=Mus musculus GN=Nudc PE=1 SV=1	NUDC_MOUSE	38 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Obg-like ATPase 1 OS=Mus musculus GN=Ola1 PE=1 SV=1	OLA1_MOUSE	45 kDa	1	0	6	3	6	0	3	3	6	0	3	6	0
Heterogeneous nuclear ribonucleoprotein H OS=Mus musculus GN=Hnrnp1 PE=2 SV	Q8C2Q7_MOUSE	51 kDa	1	4	3	1	3	4	1	1	3	4	1	3	4
40S ribosomal protein S20 OS=Mus musculus GN=Rps20 PE=1 SV=1	RS20_MOUSE	13 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Actin-related protein 2 OS=Mus musculus GN=Actr2 PE=1 SV=1	ARP2_MOUSE	45 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
LIM and senescent cell antigen-like-containing domain protein 1 OS=Mus musculus G	IE9QP62_MOUSE (+1)	44 kDa	1	0	2	0	2	0	0	0	2	0	0	2	0
Cluster of ATP-dependent RNA helicase DDX3X OS=Mus musculus GN=Ddx3x PE=1 SV	DDX3X_MOUSE	73 kDa	1	2	5	0	5	2	0	0	5	2	0	5	2
Cluster of Mitochondrial calcium uniporter regulator 1 OS=Mus musculus GN=Ccdc90	MCUR1_MOUSE	38 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Ras-related protein Rap-1b OS=Mus musculus GN=Rap1b PE=2 SV=2 (RAP1	RAP1B_MOUSE [2]	21 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cluster of Serine/arginine-rich-splicing factor 4 OS=Mus musculus GN=Srsf4 PE=2 SV	A2A837_MOUSE	56 kDa	1	0	0	1	0	0	0	0	0	1	0	0	0
ATP-binding cassette sub-family B member 7, mitochondrial OS=Mus musculus GN=A	ABC7_MOUSE	83 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
CDGSH iron sulfur domain 3 OS=Mus musculus GN=Cisd3 PE=2 SV=1	B2RWE3_MOUSE (+1)	15 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Calpain-1 catalytic subunit OS=Mus musculus GN=Capn1 PE=2 SV=1	CAN1_MOUSE	82 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Coatomer subunit delta OS=Mus musculus GN=Arccn1 PE=2 SV=2	COPD_MOUSE	57 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Endothelial differentiation-related factor 1 OS=Mus musculus GN=Edf1 PE=1 SV=1	EDF1_MOUSE	16 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Protein-cysteine N-palmitoyltransferase HHAT-like protein OS=Mus musculus GN=Hh	HHAT_MOUSE	56 kDa	1	3	0	6	0	3	6	6	0	3	6	0	3
Ig gamma-3 chain C region OS=Mus musculus PE=1 SV=2	IHG3_MOUSE	44 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
LYR motif-containing protein 4 OS=Mus musculus GN=Lyrm4 PE=2 SV=1	LYRM4_MOUSE	11 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
NFU1 iron-sulfur cluster scaffold homolog, mitochondrial OS=Mus musculus GN=Nfu1	NFU1_MOUSE	29 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
1-acyl-sn-glycerol-3-phosphate acyltransferase gamma OS=Mus musculus GN=Agpat3	PLCC_MOUSE	43 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Proteasome subunit alpha type-3 OS=Mus musculus GN=Psm3 PE=1 SV=3	PSA3_MOUSE	28 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
28S ribosomal protein S31, mitochondrial OS=Mus musculus GN=Mrps31 PE=2 SV=1	RT31_MOUSE	44 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
tRNA-splicing ligase RtcB homolog OS=Mus musculus GN=D10Wsu52e PE=2 SV=1	RTCB_MOUSE	55 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Selenoprotein P OS=Mus musculus GN=Sepp1 PE=2 SV=3	SEPP1_MOUSE	43 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
PDZ and LIM domain protein 1 OS=Mus musculus GN=Pdlim1 PE=2 SV=4	PDL1_MOUSE	36 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3

Transmembrane protein 65 OS=Mus musculus GN=Tmem65 PE=2 SV=1	TMM65_MOUSE	25 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Protein S100-A13 OS=Mus musculus GN=S100a13 PE=1 SV=1	S10AD_MOUSE	11 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Cluster of Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2 (CYB5_MOUSE)	CYB5_MOUSE [2]	15 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
Ig gamma-2A chain C region secreted form OS=Mus musculus PE=1 SV=1	GCA8_MOUSE	37 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Arginine-tRNA ligase, cytoplasmic OS=Mus musculus GN=Rars PE=2 SV=2	SYRC_MOUSE	76 kDa	1	0	0	2	0	0	2	2	0	0	2	0	0
Isoform 2 of Hepatoma-derived growth factor-related protein 3 OS=Mus musculus Ghs p Q9JMG7-2 HDGR3_MOI?			1	0	0	2	0	0	0	0	2	0	0	2	0
Cluster of Ribose-phosphate pyrophosphokinase OS=Mus musculus GN=Prps13 PE=2 G3UXL2_MOUSE [2]		35 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Cluster of Charged multivesicular body protein 4b OS=Mus musculus GN=Chmp4b PE=CHM4B_MOUSE		25 kDa	1	0	0	3	0	0	0	0	3	0	0	3	0
Cluster of Oxysterol-binding protein 1 OS=Mus musculus GN=Osbp PE=1 SV=3 (OSBP1_OSBP1_MOUSE)		89 kDa	1	0	0	4	0	4	0	4	0	4	0	4	0
Cytosolic 5'-nucleotidase 3 OS=Mus musculus GN=Nt5c3 PE=1 SV=4	SNT3_MOUSE	37 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Actin-related protein 3 OS=Mus musculus GN=Actr3 PE=1 SV=3	ARP3_MOUSE	47 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Carbonic anhydrase 1 OS=Mus musculus GN=Ca1 PE=2 SV=4	CAH1_MOUSE	28 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	CDC42_MOUSE	21 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Coiled-coil-helix-coiled-coil-helix domain-containing protein 6, mitochondrial OS=Mus CHCH6_MOUSE (+1)		30 kDa	1	4	0	0	0	4	0	0	0	4	0	0	4
MCG148436 OS=Mus musculus GN=1700014D04Rik PE=4 SV=1	J3QMS2_MOUSE	109 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Mitochondrial intermediate peptidase OS=Mus musculus GN=Mipep PE=2 SV=1	MPEP_MOUSE	81 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Methionine-R-sulfoxide reductase B3, mitochondrial OS=Mus musculus GN=MsrB3 PE=MSRB3_MOUSE		27 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Omega-amidase NIT2 OS=Mus musculus GN=Nit2 PE=1 SV=1	NIT2_MOUSE	31 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
26S proteasome non-ATPase regulatory subunit 6 OS=Mus musculus GN=Psm6d PE=1 PSM6D_MOUSE		46 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Tyrosine-protein phosphatase non-receptor type 11 OS=Mus musculus GN=Ptpn11 PE=PTN11_MOUSE (+1)		68 kDa	1	0	0	4	0	0	4	0	0	4	0	0	0
Ribosome maturation protein SBDS OS=Mus musculus GN=Sbds PE=1 SV=4	SBDS_MOUSE	29 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Ubiquitin carboxyl-terminal hydrolase isozyme L3 OS=Mus musculus GN=Uchl3 PE=1 UCHL3_MOUSE		26 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Isoform 2 of Protein CDV3 OS=Mus musculus GN=Cdv3	sp Q4VAA2-2 CDV3_MOUSE?		1	0	0	4	0	0	4	4	0	0	4	0	0
Acyl-CoA synthetase family member 2, mitochondrial OS=Mus musculus GN=Acsf2 PE=ACSF2_MOUSE		68 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Echinoderm microtubule-associated protein-like 1 OS=Mus musculus GN=Em11 PE=2 EMAL1_MOUSE		90 kDa	1	0	0	3	0	0	3	0	0	3	0	0	0
von Willebrand factor A domain-containing protein 5A OS=Mus musculus GN=Vwa5a VMA5A_MOUSE		87 kDa	1	3	3	0	3	3	0	0	3	3	0	3	3
Heterogeneous nuclear ribonucleoprotein Q OS=Mus musculus GN=Syncrip PE=2 SV=2 G3UZJ2_MOUSE		59 kDa	1	0	1	0	1	0	0	0	1	0	0	1	0
Neurofilament heavy polypeptide OS=Mus musculus GN=Nefh PE=1 SV=3	NFH_MOUSE	117 kDa	1	0	0	0	0	1	0	0	0	1	0	0	1
Cluster of Annexin A1 OS=Mus musculus GN=Anxa1 PE=1 SV=2 (ANXA1_MOUSE)	ANXA1_MOUSE	39 kDa	1	0	2	4	2	0	4	4	2	0	4	2	0
Cluster of Serine (Or cysteine) peptidase inhibitor, clade A, member 3N, isoform CRA_ G3X8T9_MOUSE [2]		47 kDa	1	1	0	0	0	1	0	0	0	1	0	0	1
ATP-dependent RNA helicase DDX1 OS=Mus musculus GN=Ddx1 PE=1 SV=1	DDX1_MOUSE	83 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
NADH-cytochrome b5 reductase 3 OS=Mus musculus GN=Cyb5r3 PE=2 SV=1	F2Z456_MOUSE (+1)	35 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Mitochondrial glutamate carrier 1 OS=Mus musculus GN=Slc25a22 PE=1 SV=1	GHC1_MOUSE	35 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Glutaredoxin-related protein 5, mitochondrial OS=Mus musculus GN=Glrx5 PE=2 SV=2 GLRX5_MOUSE		16 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Homeodomain-only protein OS=Mus musculus GN=Hopx PE=1 SV=1	HOP_MOUSE	8 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Probable D-lactate dehydrogenase, mitochondrial OS=Mus musculus GN=Ldhd PE=1 SLDHD_MOUSE		52 kDa	1	0	5	0	5	0	0	0	5	0	0	5	0
Trans-2-enoyl-CoA reductase, mitochondrial OS=Mus musculus GN=Mecr PE=2 SV=2 MECR_MOUSE		40 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Nucleobindin-1 OS=Mus musculus GN=Nucb1 PE=1 SV=2	NUCB1_MOUSE	53 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=M OST48_MOUSE		49 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Protein kinase C and casein kinase II substrate protein 3 OS=Mus musculus GN=Pacsin3 PACN3_MOUSE		49 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Proteasome subunit beta type-7 OS=Mus musculus GN=Psmb7 PE=1 SV=1	PSB7_MOUSE	30 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Serum albumin OS=Bos taurus GN=ALB PE=1 SV=4	ALBU_BOVIN	69 kDa	1	2	2	1	2	2	1	1	2	2	1	2	2
Major vault protein OS=Mus musculus GN=Mvp PE=2 SV=1	E9Q3X0_MOUSE (+1)	97 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
GMP reductase 1 OS=Mus musculus GN=Gmpr PE=2 SV=1	GMPR1_MOUSE	37 kDa	1	0	3	4	3	0	4	4	3	0	4	3	0
LDLR chaperone MESD OS=Mus musculus GN=Mesdc2 PE=1 SV=1	MESD_MOUSE	25 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Delta-sarcoglycan OS=Mus musculus GN=Sgcd PE=1 SV=1	SGCD_MOUSE	32 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Translocase of inner mitochondrial membrane domain-containing protein 1 OS=Mus r TIDC1_MOUSE		32 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Sorting nexin-12 OS=Mus musculus GN=Snx12 PE=2 SV=1	SNX12_MOUSE	19 kDa	1	0	2	2	2	0	2	2	2	0	2	2	0
Heterogeneous nuclear ribonucleoprotein R OS=Mus musculus GN=Hnrnpr PE=2 SV=1 Q8VHM5_MOUSE		71 kDa	1	0	1	0	1	0	0	0	1	0	0	1	0
Cluster of Muscleblind-like 1 (Drosophila), isoform CRA_a OS=Mus musculus GN=Mbn G3X9Q0_MOUSE [6]		41 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Cluster of Histone-binding protein RBBP7 OS=Mus musculus GN=Rbbp7 PE=2 SV=1 (A: A2AFI9_MOUSE [3])		44 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Cluster of Histone H2A.Z OS=Mus musculus GN=H2afz PE=1 SV=2 (H2AZ_MOUSE)	H2AZ_MOUSE	14 kDa	1	1	2	3	2	1	3	3	2	1	3	2	1
Cluster of Dihydropyrimidinase-related protein 3 OS=Mus musculus GN=Dpysl3 PE=2 E9PWE8_MOUSE		74 kDa	1	0	0	2	0	0	2	2	0	0	2	0	0
Coiled-coil domain-containing protein 90B, mitochondrial OS=Mus musculus GN=Ccdc CC90B_MOUSE		30 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Gap junction alpha-1 protein OS=Mus musculus GN=Gja1 PE=1 SV=2	CXA1_MOUSE	43 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Vacuolar protein sorting-associated protein 29 OS=Mus musculus GN=Vps29 PE=2 SV=2 D3Z645_MOUSE (+1)		16 kDa	1	0	0	5	0	0	5	5	0	0	5	0	0
Glucosidase 2 subunit beta OS=Mus musculus GN=Prksh PE=1 SV=1	GLU2B_MOUSE	59 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Mus Q8R5L1_MOUSE		31 kDa	1	5	0	0	0	5	0	0	0	5	0	0	5
UV excision repair protein RAD23 homolog B OS=Mus musculus GN=Rad23b PE=1 SV=1 RD23B_MOUSE		44 kDa	1	3	4	3	4	3	3	3	4	3	3	4	3
40S ribosomal protein S23 OS=Mus musculus GN=Rps23 PE=2 SV=3	RS23_MOUSE	16 kDa	1	0	0	5	0	0	5	5	0	0	5	0	0
Aldehyde dehydrogenase family 8 member A1 OS=Mus musculus GN=Aldh8a1 PE=1 S AL8A1_MOUSE		54 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Basal cell adhesion molecule OS=Mus musculus GN=Bcam PE=2 SV=1	BCAM_MOUSE	68 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Thioredoxin domain-containing protein 5 OS=Mus musculus GN=Txdnc5 PE=2 SV=1	E9PXX7_MOUSE	39 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Prostaglandin synthase OS=Mus musculus GN=Fam213b PE=1 SV=1	PGFS_MOUSE	22 kDa	1	0	4	4	4	0	4	4	4	0	4	4	0
UV excision repair protein RAD23 homolog A (Fragment) OS=Mus musculus GN=Rad2. Q3TN85_MOUSE (+2)		35 kDa	1	0	4	3	4	0	3	3	4	0	3	4	0
Protein Taf15 OS=Mus musculus GN=Taf15 PE=2 SV=1	Q8BQ46_MOUSE	59 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0

Beta-casein OS=Bos taurus GN=CSN2 PE=1 SV=2	CASB_BOVIN	25 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Histone H1t OS=Mus musculus GN=Hist1h1t PE=1 SV=4	H1T_MOUSE (+1)	22 kDa	1	1	0	0	0	1	0	0	0	1	0	0	1
5'-AMP-activated protein kinase catalytic subunit alpha-2 OS=Mus musculus GN=Prka AAPK2_MOUSE	AAPK2_MOUSE	62 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Heat shock protein beta-2 OS=Mus musculus GN=Hspb2 PE=2 SV=2	HSPB2_MOUSE	20 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Marcks PE=1 SV=MARCS_MOUSE	MARCS_MOUSE	30 kDa	1	0	5	4	5	0	4	4	5	0	4	5	0
Microtubule-associated protein RP/EB family member 1 OS=Mus musculus GN=Mapr1 MARE1_MOUSE	MARE1_MOUSE	30 kDa	1	0	2	3	2	0	3	3	2	0	3	2	0
Proteasome subunit beta type-1 OS=Mus musculus GN=Psb1 PE=1 SV=1	PSB1_MOUSE	26 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Solute carrier family 25 member 46 OS=Mus musculus GN=Slc25a46 PE=1 SV=1	S2546_MOUSE	46 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
DnaJ homolog subfamily A member 1 OS=Mus musculus GN=Dnaj1 PE=1 SV=1	DNJA1_MOUSE	45 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Aspartyl aminopeptidase OS=Mus musculus GN=Dnpep PE=2 SV=2	DNPEP_MOUSE	52 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Dual specificity mitogen-activated protein kinase kinase 1 OS=Mus musculus GN=Map2k1 MP2K1_MOUSE	MP2K1_MOUSE	43 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1	A2AL85_MOUSE (+1)	81 kDa	1	0	0	2	0	2	0	2	0	0	2	0	0
Proteasome subunit alpha type-5 OS=Mus musculus GN=Psm5 PE=1 SV=1	PSA5_MOUSE	26 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Microtubule-associated protein RP/EB family member 3 OS=Mus musculus GN=Mapr3 MARE3_MOUSE	MARE3_MOUSE	32 kDa	1	0	1	1	1	0	1	1	1	0	1	1	0
Monoglyceride lipase OS=Mus musculus GN=Mgll PE=2 SV=1	D3YYS6_MOUSE (+3)	37 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Fibroblast growth factor 1 OS=Mus musculus GN=Fgf1 PE=2 SV=1	FGF1_MOUSE	17 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Glycerophosphocholine phosphodiesterase GPCPD1 OS=Mus musculus GN=Gpcpd1 P	GPCPD1_MOUSE	77 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Vesicle-associated membrane protein 3 OS=Mus musculus GN=Vamp3 PE=1 SV=1	VAMP3_MOUSE	11 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Mitochondrial antiviral-signaling protein OS=Mus musculus GN=Mavs PE=1 SV=1	MAVS_MOUSE	53 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Biglycan OS=Mus musculus GN=Bgn PE=2 SV=1	PGS1_MOUSE	42 kDa	1	0	2	0	2	0	0	0	2	0	0	2	0
Nitric oxide synthase, inducible OS=Mus musculus GN=Nos2 PE=1 SV=1	NOS2_MOUSE	131 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
Annexin A4 OS=Mus musculus GN=Anxa4 PE=2 SV=4	ANXA4_MOUSE	36 kDa	1	0	2	0	2	0	0	0	2	0	0	2	0
F-actin-capping protein subunit alpha-1 OS=Mus musculus GN=Capza1 PE=1 SV=4	CAZA1_MOUSE (+1)	33 kDa	1	0	2	0	2	0	0	0	2	0	0	2	0
Cluster of Mitochondrial-processing peptidase subunit beta OS=Mus musculus GN=Pn MPPB_MOUSE	MPPB_MOUSE	55 kDa	1	2	2	2	2	2	2	2	2	2	2	2	2
Protein Gyk OS=Mus musculus GN=Gyk PE=2 SV=1	B1ASZ3_MOUSE (+1)	60 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
Alpha-endosulfine OS=Mus musculus GN=Ensa PE=1 SV=1	ENSA_MOUSE (+1)	13 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
Mitochondrial import inner membrane translocase subunit Tim8 B OS=Mus musculus	TIM8B_MOUSE	9 kDa	1	0	3	3	3	0	3	3	3	0	3	3	0
Uncharacterized protein C19orf43 homolog OS=Mus musculus PE=2 SV=1	CS043_MOUSE	18 kDa	1	0	0	4	0	0	4	4	0	0	4	0	0
rRNA 2'-O-methyltransferase fibrillar OS=Mus musculus GN=Fbl PE=1 SV=2	FBRL_MOUSE	34 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2
Cluster of Agrin OS=Mus musculus GN=Agrn PE=4 SV=1 (MOQWP1_MOUSE)	MOQWP1_MOUSE	217 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
Limb and neural patterns, isoform CRA_c OS=Mus musculus GN=Lnp PE=4 SV=1	A2ASL8_MOUSE (+1)	42 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
6-phosphofructokinase, liver type OS=Mus musculus GN=Pfkfb1 PE=1 SV=4	K6P1_MOUSE	85 kDa	1	0	3	1	3	0	1	1	3	0	1	3	0
ATP-dependent (S)-NAD(P)H-hydrate dehydratase OS=Mus musculus GN=Carkd PE=3	J3QMM7_MOUSE (+4)	35 kDa	1	0	3	0	3	0	0	0	3	0	0	3	0
1,4-alpha-glucan-branching enzyme OS=Mus musculus GN=Gbe1 PE=4 SV=2	F6ZHD8_MOUSE (+1)	80 kDa	1	0	4	0	4	0	0	0	4	0	0	4	0
Protein unc-45 homolog A (Fragment) OS=Mus musculus GN=Unc45a PE=2 SV=1	D3YZN8_MOUSE	19 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
sp K22E_HUMAN	sp K22E_HUMAN	?	1	0	0	0	0	0	0	0	0	0	0	0	0
Uncharacterized protein C1orf170 homolog OS=Mus musculus PE=2 SV=1	CA170_MOUSE	88 kDa	1	0	0	2	0	0	2	2	0	0	2	0	0
F-actin-capping protein subunit alpha-2 OS=Mus musculus GN=Capza2 PE=1 SV=3	CAZA2_MOUSE	33 kDa	1	0	0	2	0	0	2	2	0	0	2	0	0
Crk-like protein OS=Mus musculus GN=Crkl PE=1 SV=2	CRKL_MOUSE	34 kDa	1	0	0	3	0	0	3	3	0	0	3	0	0
DNA excision repair protein ERCC-6-like OS=Mus musculus GN=Erc6l PE=1 SV=1	ERCC6L_MOUSE	139 kDa	1	3	0	0	0	3	0	0	0	3	0	0	3
DNA-binding protein A OS=Mus musculus GN=Csda PE=1 SV=2	DBPA_MOUSE	39 kDa	1	0	0	2	0	0	2	2	0	0	2	0	0
MCG11326, isoform CRA_b OS=Mus musculus GN=Hnrrnp3 PE=4 SV=1	D3YWT1_MOUSE (+1)	35 kDa	1	2	0	0	0	2	0	0	0	2	0	0	2