24 Hour

Protein hit no.	Accession no.	Protein description	Protein scor	MW [kDa]	Sequence covered %	Protein PI	115/114 ratio (ZnONP20)	N	SD
110	CH10_MOUSE	10 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspe1 PE=1 SV=2	180	12685	33.3	7.93	0.922	5	1.051
54	1433B_MOUSE	14-3-3 protein beta/alpha OS=Mus musculus GN=Ywhab PE=1 SV=3	345	31187	20.3	4.77	0.914	10	1.06
44	1433E_MOUSE	14-3-3 protein epsilon OS=Mus musculus GN=Ywhae PE=1 SV=1	410	32031	29.4	4.63	0.981	12	1.058
118	1433F_MOUSE	14-3-3 protein eta OS=Mus musculus GN=Ywhah PE=1 SV=2	161	31358	15	4.81	0.914	5	1.059
65	1433G_MOUSE	14-3-3 protein gamma OS=Mus musculus GN=Ywhag PE=1 SV=2	300	31017	22.3	4.8	0.899	10	1.043
148	1433S_MOUSE	14-3-3 protein sigma OS=Mus musculus GN=Sfn PE=1 SV=2	124	30519	6.5	4.72	0.906	3	1.079
108	1433T_MOUSE	14-3-3 protein theta OS=Mus musculus GN=Ywhaq PE=1 SV=1	185	30873	12.2	4.69	0.952	4	1.096
20	1433Z_MOUSE	14-3-3 protein zeta/delta OS=Mus musculus GN=Ywhaz PE=1 SV=1	623	30918	40.8	4.73	0.913	14	1.053
236	THIM_MOUSE	3-ketoacyl-CoA thiolase, mitochondrial OS=Mus musculus GN=Acaa2 PE=1 SV=3	71	45774	8.1	8.33	0.812	3	1.106
253	RS18_MOUSE	40S ribosomal protein S18 OS=Mus musculus GN=Rps18 PE=1 SV=3	63	20158	7.2	10.99	0.911	2	1.004
141	RS19_MOUSE	40S ribosomal protein S19 OS=Mus musculus GN=Rps19 PE=1 SV=3	131	18381	13.1	10.41	1.138	3	1.105
124	CH60_MOUSE	60 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspd1 PE=1 SV=1	153	68981	9.2	5.91	1.007	6	1.059
116	RLA0_MOUSE	60S acidic ribosomal protein P0 OS=Mus musculus GN=Rplp0 PE=1 SV=3	164	37215	5.7	5.91	1.012	6	1.043
266	RLA1_MOUSE	60S acidic ribosomal protein P1 OS=Mus musculus GN=Rplp1 PE=2 SV=1	59	12712	14	4.28	0.964	2	1.074
242	RLA2_MOUSE	60S acidic ribosomal protein P2 OS=Mus musculus GN=Rplp2 PE=1 SV=3	67	13229	17.4	4.42	0.932	2	1.026
225	RL13_MOUSE	60S ribosomal protein L13 OS=Mus musculus GN=Rpl13 PE=2 SV=3	77	28083	5.2	11.54	0.978	2	1.081
199	RL23_MOUSE	60S ribosomal protein L23 OS=Mus musculus GN=Rpl23 PE=1 SV=1	85	17254	7.1	10.51	1.14	2	1.035
146	RL28_MOUSE	60S ribosomal protein L28 OS=Mus musculus GN=Rpl28 PE=1 SV=2	127	17931	8.8	12.02	0.95	4	1.07
151	6PGD_MOUSE	6-phosphogluconate dehydrogenase, decarboxylating OS=Mus musculus GN=Pgd PE=1 SV=3	122	58671	5.6	6.81	1.024	5	1.074
55	GRP78_MOUSE	78 kDa glucose-regulated protein OS=Mus musculus GN=Hspa5 PE=1 SV=3	344	81404	11.9	5.07	1.052	12	1.065
134	ACON_MOUSE	Aconitate hydratase, mitochondrial OS=Mus musculus GN=Aco2 PE=1 SV=1	139	93933	7.2	8.08	0.573	5	1.088
11	ACTA_MOUSE	Actin, aortic smooth muscle OS=Mus musculus GN=Acta2 PE=1 SV=1	982	45186	28.4	5.23	1.072	37	1.039
7	ACTB_MOUSE	Actin, cytoplasmic 1 OS=Mus musculus GN=Actb PE=1 SV=1	1578	44868	50.4	5.29	1.095	52	1.042
185	ARP3_MOUSE	Actin-related protein 3 OS=Mus musculus GN=Actr3 PE=1 SV=3	94	51153	7.7	5.61	0.939	3	1.032
277	ACBP_MOUSE	Acyl-CoA-binding protein OS=Mus musculus GN=Dbi PE=1 SV=2	55	12300	25.3	8.78	0.862	2	1.073
156	CAP1_MOUSE	Adenylyl cyclase-associated protein 1 OS=Mus musculus GN=Cap1 PE=1 SV=4	118	57717	12.9	7.16	0.794	3	1.095
69	ARF1_MOUSE	ADP-ribosylation factor 1 OS=Mus musculus GN=Arf1 PE=1 SV=2	286	22459	30.9	6.32	1.023	10	1.079
74	ARF5_MOUSE	ADP-ribosylation factor 5 OS=Mus musculus GN=Arf5 PE=1 SV=2	273	22338	20.6	6.3	1.064	8	1.061
126	RAGE_MOUSE	Advanced glycosylation end product-specific receptor OS=Mus musculus GN=Ager PE=1 SV=1	151	45224	17.9	5.78	0.919	7	1.038

170	AKA12_MOUSE	A-kinase anchor protein 12 OS=Mus musculus GN=Akap12 PE=1 SV=1	103	199347	1.7	4.39	0.975	2	1.073
191	AKAP5_MOUSE	A-kinase anchor protein 5 OS=Mus musculus GN=Akap5 PE=1 SV=2	89	88088	4.3	4.72	0.852	2	1.049
244	AK1A1_MOUSE	Alcohol dehydrogenase [NADP(+)] OS=Mus musculus GN=Akr1a1 PE=1 SV=3	67	39774	8.3	6.9	0.9	3	1.101
223	ADH1_MOUSE	Alcohol dehydrogenase 1 OS=Mus musculus GN=Adh1 PE=1 SV=2	77	44758	4.3	8.44	1.069	2	1.026
34	ALDH2_MOUSE	Aldehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh2 PE=1 SV=1	509	61383	27	7.53	0.93	20	1.049
99	ALDR_MOUSE	Aldose reductase OS=Mus musculus GN=Akr1b1 PE=1 SV=3	205	39876	10.8	6.71	0.94	6	1.024
80	A1AT2_MOUSE	Alpha-1-antitrypsin 1-2 OS=Mus musculus GN=Serpina1b PE=1 SV=2	260	49974	22.3	5.32	1.026	12	1.054
62	A1AT3_MOUSE	Alpha-1-antitrypsin 1-3 OS=Mus musculus GN=Serpina1c PE=1 SV=2	308	49535	22.3	5.25	1.089	12	1.033
88	A1AT4_MOUSE	Alpha-1-antitrypsin 1-4 OS=Mus musculus GN=Serpina1d PE=2 SV=1	233	49710	16.9	5.24	1.031	11	1.039
233	FETUA_MOUSE	Alpha-2-HS-glycoprotein OS=Mus musculus GN=Ahsg PE=1 SV=1	72	40107	2.3	6.04	0.994	2	1.038
37	A2M_MOUSE	Alpha-2-macroglobulin OS=Mus musculus GN=A2m PE=1 SV=3	486	180541	7.8	6.24	0.959	21	1.019
83	ACTN4_MOUSE	Alpha-actinin-4 OS=Mus musculus GN=Actn4 PE=1 SV=1	246	113493	7	5.25	0.928	8	1.069
24	ENOA_MOUSE	Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3	579	52431	30	6.37	0.878	21	1.043
213	PARVA_MOUSE	Alpha-parvin OS=Mus musculus GN=Parva PE=1 SV=1	80	47105	9.4	5.69	0.958	3	1.024
96	ACE_MOUSE	Angiotensin-converting enzyme OS=Mus musculus GN=Ace PE=1 SV=3	208	160446	5.6	6.1	0.938	11	1.055
188	ANXA1_MOUSE	Annexin A1 OS=Mus musculus GN=Anxa1 PE=1 SV=2	92	43263	8.4	6.97	1.668	3	1.525
162	ANXA2_MOUSE	Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	113	43781	8.3	7.55	1.295	4	1.073
66	ANXA3_MOUSE	Annexin A3 OS=Mus musculus GN=Anxa3 PE=1 SV=4	294	40390	26.3	5.5	1.037	10	1.09
41	ANXA5_MOUSE	Annexin A5 OS=Mus musculus GN=Anxa5 PE=1 SV=1	431	38946	29.2	4.83	1.146	15	1.066
176	ANXA6_MOUSE	Annexin A6 OS=Mus musculus GN=Anxa6 PE=1 SV=3	100	84131	4.6	5.34	1.02	3	1.081
86	APOA1_MOUSE	Apolipoprotein A-I OS=Mus musculus GN=Apoa1 PE=1 SV=2	240	33479	21.6	5.51	1.248	10	1.06
284	APOA2_MOUSE	Apolipoprotein A-II OS=Mus musculus GN=Apoa2 PE=1 SV=2	53	12789	9.8	6.56	1.364	2	1.01
157	ATPA_MOUSE	ATP synthase subunit alpha, mitochondrial OS=Mus musculus GN=Atp5a1 PE=1 SV=1	118	64419	8.3	9.22	0.853	3	1.038
164	ATPB_MOUSE	ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b PE=1 SV=2	108	59724	7.6	5.19	0.849	4	1.017
163	ATPD_MOUSE	ATP synthase subunit delta, mitochondrial OS=Mus musculus GN=Atp5d PE=1 SV=1	112	18310	8.3	5.03	0.792	2	1.025
204	ACLY_MOUSE	ATP-citrate synthase OS=Mus musculus GN=Acly PE=1 SV=1	84	131195	2	7.13	0.699	3	[1.063]
189	E41L2_MOUSE	Band 4.1-like protein 2 OS=Mus musculus GN=Epb41l2 PE=1 SV=2	92	121815	2	5.31	0.979	2	1.029
29	ACTBL_MOUSE	Beta-actin-like protein 2 OS=Mus musculus GN=Actbl2 PE=1 SV=1	520	44991	23.7	5.3	1.054	23	1.041
75	ENOB_MOUSE	Beta-enolase OS=Mus musculus GN=Eno3 PE=1 SV=3	273	52747	21.9	6.73	0.862	11	1.045
61	CALM_MOUSE	Calmodulin OS=Mus musculus GN=Calm1 PE=1 SV=2	308	18124	33.6	4.09	1.086	8	1.056
201	CPNS1_MOUSE	Calpain small subunit 1 OS=Mus musculus GN=Capns1 PE=2 SV=1	85	30266	10.4	5.41	0.801	2	1.003
205	CAN2_MOUSE	Calpain-2 catalytic subunit OS=Mus musculus GN=Capn2 PE=2 SV=4	84	86564	1.1	4.86	0.976	2	1.054

230	ICAL_MOUSE	Calpastatin OS=Mus musculus GN=Cast PE=1 SV=2	75	99609	1.6	5.37	0.762	2	1.009
56	CALR_MOUSE	Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1	341	54443	23.6	4.33	0.931	14	1.025
147	CAH1_MOUSE	Carbonic anhydrase 1 OS=Mus musculus GN=Ca1 PE=2 SV=4	127	30521	7.7	6.44	0.986	3	1.061
27	CAH2_MOUSE	Carbonic anhydrase 2 OS=Mus musculus GN=Ca2 PE=1 SV=4	549	31989	44.2	6.49	1.078	18	1.042
93	CAH3_MOUSE	Carbonic anhydrase 3 OS=Mus musculus GN=Ca3 PE=1 SV=3	214	32460	9.2	6.89	0.554	5	1.078
5	CBR2_MOUSE	Carbonyl reductase [NADPH] 2 OS=Mus musculus GN=Cbr2 PE=1 SV=1	2259	28339	62.7	9.1	0.938	60	1.028
64	EST1C_MOUSE	Carboxylesterase 1C OS=Mus musculus GN=Ces1c PE=1 SV=4	303	65138	8.3	4.97	0.998	9	1.053
43	CES1D_MOUSE	Carboxylesterase 1D OS=Mus musculus GN=Ces1d PE=1 SV=1	423	67022	22.8	6.17	0.922	18	1.032
105	CATA_MOUSE	Catalase OS=Mus musculus GN=Cat PE=1 SV=4	189	64743	9.7	7.72	0.856	9	[1.052]
208	CTNB1_MOUSE	Catenin beta-1 OS=Mus musculus GN=Ctnnb1 PE=1 SV=1	83	89813	1.5	5.53	0.86	2	1.113
299	CATD_MOUSE	Cathepsin D OS=Mus musculus GN=Ctsd PE=1 SV=1	49	49616	4.4	6.71	0.941	2	1.066
59	CDC42_MOUSE	Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	322	24115	29.8	6.15	1.04	9	1.038
324	CE164_MOUSE	Centrosomal protein of 164 kDa OS=Mus musculus GN=Cep164 PE=2 SV=2	43	178145	0.8	5.22	0.931	2	1.034
222	CLIC1_MOUSE	Chloride intracellular channel protein 1 OS=Mus musculus GN=Clic1 PE=1 SV=3	78	29866	12.4	5.09	1.092	2	1.03
115	CLIC5_MOUSE	Chloride intracellular channel protein 5 OS=Mus musculus GN=Clic5 PE=1 SV=1	168	31578	12.7	5.64	1.065	4	1.059
178	CISY_MOUSE	Citrate synthase, mitochondrial OS=Mus musculus GN=Cs PE=1 SV=1	99	55968	6.9	8.72	0.634	4	1.304
184	CLH1_MOUSE	Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3	95	206982	1.9	5.48	1.112	3	1.032
60	COF1_MOUSE	Cofilin-1 OS=Mus musculus GN=Cfl1 PE=1 SV=3	320	22478	39.8	8.22	0.932	11	1.061
72	CO3_MOUSE	Complement C3 OS=Mus musculus GN=C3 PE=1 SV=3	279	204900	6.4	6.29	1.079	12	1.051
193	COR1C_MOUSE	Coronin-1C OS=Mus musculus GN=Coro1c PE=1 SV=2	88	58971	4.2	6.65	0.928	4	1.079
219	KCRB_MOUSE	Creatine kinase B-type OS=Mus musculus GN=Ckb PE=1 SV=1	80	45942	9.7	5.4	0.703	4	1.112
63	KCRM_MOUSE	Creatine kinase M-type OS=Mus musculus GN=Ckm PE=1 SV=1	306	48101	20.5	6.58	0.828	12	1.076
290	CSRP1_MOUSE	Cysteine and glycine-rich protein 1 OS=Mus musculus GN=Csrp1 PE=1 SV=3	51	24718	13	8.9	1.156	2	1.032
220	CRIP2_MOUSE	Cysteine-rich protein 2 OS=Mus musculus GN=Crip2 PE=1 SV=1	79	26670	13.9	8.94	1.53	3	1.271
270	CYB5_MOUSE	Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2	57	16817	6.7	4.96	0.944	2	1.019
95	CYC_MOUSE	Cytochrome c, somatic OS=Mus musculus GN=Cycs PE=1 SV=2	210	14428	28.6	9.61	0.729	7	1.054
249	CP2F2_MOUSE	Cytochrome P450 2F2 OS=Mus musculus GN=Cyp2f2 PE=2 SV=1	65	59267	2	7.74	0.892	2	1.055
70	DEST_MOUSE	Destrin OS=Mus musculus GN=Dstn PE=1 SV=3	285	21956	28.5	8.14	0.984	9	1.063
12	DPYL2_MOUSE	Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=1 SV=2	941	67316	47.9	5.95	0.939	29	1.033
265	FMO1_MOUSE	Dimethylaniline monooxygenase [N-oxide-forming] 1 OS=Mus musculus GN=Fmo1 PE=1 SV=1	60	65524	5.3	8.72	1.103	3	1.106
114	FMO2_MOUSE	Dimethylaniline monooxygenase [N-oxide-forming] 2 OS=Mus musculus GN=Fmo2 PE=1 SV=3	170	66635	12.7	8.68	0.962	7	1.061
58	EHD2_MOUSE	EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1	328	67038	20.4	6.08	1.004	15	1.051

94	EHD4_MOUSE	EH domain-containing protein 4 OS=Mus musculus GN=Ehd4 PE=1 SV=1	210	67678	16.8	6.33	1.11	9	1.078
194	ETFA_MOUSE	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Mus musculus GN=Etfa PE=1 SV=2	88	39154	18	8.62	0.694	4	[1.260]
310	ETFB_MOUSE	Electron transfer flavoprotein subunit beta OS=Mus musculus GN=Etfb PE=1 SV=3	45	31681	5.1	8.24	0.776	2	1.021
36	EF1A1_MOUSE	Elongation factor 1-alpha 1 OS=Mus musculus GN=Eef1a1 PE=1 SV=3	494	57275	19.5	9.1	1.083	17	1.027
217	EF1B_MOUSE	Elongation factor 1-beta OS=Mus musculus GN=Eef1b PE=1 SV=5	80	28275	8.9	4.53	1.099	3	1.081
190	EF2_MOUSE	Elongation factor 2 OS=Mus musculus GN=Eef2 PE=1 SV=2	92	105401	5.2	6.41	1.125	4	1.108
235	EVL_MOUSE	Ena/VASP-like protein OS=Mus musculus GN=Evl PE=1 SV=2	71	47717	8.9	8.92	---	1	---
47	ENPL_MOUSE	Endoplasmin OS=Mus musculus GN=Hsp90b1 PE=1 SV=2	379	103744	10.6	4.74	0.937	11	1.037
313	ECHM_MOUSE	Enoyl-CoA hydratase, mitochondrial OS=Mus musculus GN=Echs1 PE=1 SV=1	45	35090	5.2	8.76	0.837	2	1.053
301	IF4B_MOUSE	Eukaryotic translation initiation factor 4B OS=Mus musculus GN=Eif4b PE=1 SV=1	48	74413	2.5	5.47	1.151	2	1.215
260	IF5A1_MOUSE	Eukaryotic translation initiation factor 5A-1 OS=Mus musculus GN=Eif5a PE=1 SV=2	62	19023	10.4	5.08	1.1	2	1.027
192	SODE_MOUSE	Extracellular superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod3 PE=1 SV=1	88	28948	7.6	6.36	0.886	2	1.11
30	EZRI_MOUSE	Ezrin OS=Mus musculus GN=Ezr PE=1 SV=3	519	77525	21.3	5.83	0.924	22	1.047
81	FAS_MOUSE	Fatty acid synthase OS=Mus musculus GN=Fasn PE=1 SV=2	254	289307	3	6.13	0.654	9	1.045
226	FABP4_MOUSE	Fatty acid-binding protein, adipocyte OS=Mus musculus GN=Fabp4 PE=1 SV=3	77	16894	9.1	8.53	0.287	2	1.042
215	FERM2_MOUSE	Fermitin family homolog 2 OS=Mus musculus GN=Fermt2 PE=1 SV=1	80	87237	1.5	6.26	0.966	2	1.004
16	FLNA_MOUSE	Filamin-A OS=Mus musculus GN=Flna PE=1 SV=5	704	306546	9.1	5.68	1.047	25	1.03
123	BLVRB_MOUSE	Flavin reductase (NADPH) OS=Mus musculus GN=Blvrb PE=2 SV=3	154	23716	22.8	6.49	1.041	5	1.032
89	FHL1_MOUSE	Four and a half LIM domains protein 1 OS=Mus musculus GN=Fhl1 PE=1 SV=3	223	38618	11.1	8.76	1.128	5	1.042
21	ALDOA_MOUSE	Fructose-bisphosphate aldolase A OS=Mus musculus GN=Aldoa PE=1 SV=2	616	43590	32.1	8.31	0.928	17	1.03
113	FUMH_MOUSE	Fumarate hydratase, mitochondrial OS=Mus musculus GN=Fh PE=1 SV=3	170	59261	8.1	9.12	0.822	5	1.033
263	LEG1_MOUSE	Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	60	16573	17	5.32	0.852	2	1.033
256	G6PI_MOUSE	Glucose-6-phosphate isomerase OS=Mus musculus GN=Gpi PE=1 SV=4	63	68963	3.2	8.14	0.955	2	1.044
173	GLU2B_MOUSE	Glucosidase 2 subunit beta OS=Mus musculus GN=Prkcsh PE=1 SV=1	102	64149	2.3	4.41	0.907	2	1.103
239	DHE3_MOUSE	Glutamate dehydrogenase 1, mitochondrial OS=Mus musculus GN=Glud1 PE=1 SV=1	70	66618	3.8	8.05	0.891	3	[1.192]
286	GSHR_MOUSE	Glutathione reductase, mitochondrial OS=Mus musculus GN=Gsr PE=1 SV=3	52	59178	1.6	8.19	0.959	2	1.01
179	GSTA3_MOUSE	Glutathione S-transferase A3 OS=Mus musculus GN=Gsta3 PE=1 SV=2	98	28416	13.6	8.76	0.953	4	1.049
275	GSTA4_MOUSE	Glutathione S-transferase A4 OS=Mus musculus GN=Gsta4 PE=1 SV=3	55	28285	6.3	6.77	0.943	2	1.027
49	GSTM1_MOUSE	Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2	372	28783	44.5	7.71	0.959	16	1.046
67	GSTM2_MOUSE	Glutathione S-transferase Mu 2 OS=Mus musculus GN=Gstm2 PE=1 SV=2	290	29008	35.3	6.9	0.955	12	[1.047]
98	GSTM4_MOUSE	Glutathione S-transferase Mu 3 OS=Mus musculus GN=Gstm3 PE=1 SV=2	207	28319	26.6	7.63	0.969	10	[1.062]
102	GSTM7_MOUSE	Glutathione S-transferase Mu 7 OS=Mus musculus GN=Gstm7 PE=1 SV=1	199	28136	20.6	6.34	0.966	9	[1.048]

203	GSTO1_MOUSE	Glutathione S-transferase omega-1 OS=Mus musculus GN=Gsto1 PE=2 SV=2	84	30978	5.8	6.92	1.058	2	1.076
166	GSTP1_MOUSE	Glutathione S-transferase P 1 OS=Mus musculus GN=Gstp1 PE=1 SV=2	106	25605	18.6	7.68	0.933	4	1.111
39	G3P_MOUSE	Glyceraldehyde-3-phosphate dehydrogenase OS=Mus musculus GN=Gapdh PE=1 SV=2	448	39908	25.8	8.44	0.96	12	1.036
317	GON4L_MOUSE	GON-4-like protein OS=Mus musculus GN=Gon4l PE=1 SV=3	44	272140	0.3	4.79	0.99	2	1.024
158	RAN_MOUSE	GTP-binding nuclear protein Ran OS=Mus musculus GN=Ran PE=1 SV=3	116	27284	10.2	7.01	1.142	3	1.077
234	GNAI2_MOUSE	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Mus musculus GN=Gnai2 PE=1 SV=5	71	44958	7.3	5.28	0.859	2	1.007
238	HPT_MOUSE	Haptoglobin OS=Mus musculus GN=Hp PE=1 SV=1	70	42456	8.6	5.88	0.973	3	1.094
48	HS71L_MOUSE	Heat shock 70 kDa protein 1-like OS=Mus musculus GN=Hspa1l PE=2 SV=4	378	78552	11.1	5.91	1.194	12	1.1
294	HSP74_MOUSE	Heat shock 70 kDa protein 4 OS=Mus musculus GN=Hspa4 PE=1 SV=1	50	106678	1.8	5.15	1.055	2	1.038
9	HSP7C_MOUSE	Heat shock cognate 71 kDa protein OS=Mus musculus GN=Hspa8 PE=1 SV=1	1070	78937	33.4	5.37	1.095	39	1.036
150	TRAP1_MOUSE	Heat shock protein 75 kDa, mitochondrial OS=Mus musculus GN=Trap1 PE=1 SV=1	122	87495	2	6.25	1.061	2	1.008
23	HS90A_MOUSE	Heat shock protein HSP 90-alpha OS=Mus musculus GN=Hsp90aa1 PE=1 SV=4	588	96729	16.1	4.93	0.995	20	1.035
14	HS90B_MOUSE	Heat shock protein HSP 90-beta OS=Mus musculus GN=Hsp90ab1 PE=1 SV=3	820	94457	22.1	4.97	0.958	28	1.033
32	HSP72_MOUSE	Heat shock-related 70 kDa protein 2 OS=Mus musculus GN=Hspa2 PE=1 SV=2	516	77466	12.2	5.51	1.108	16	1.057
4	HBA_MOUSE	Hemoglobin subunit alpha OS=Mus musculus GN=Hba PE=1 SV=2	4300	16851	71.8	7.96	1.138	108	1.067
1	HBB1_MOUSE	Hemoglobin subunit beta-1 OS=Mus musculus GN=Hbb-b1 PE=1 SV=2	9684	17651	93.9	7.12	1.178	245	[1.019]
2	HBB2_MOUSE	Hemoglobin subunit beta-2 OS=Mus musculus GN=Hbb-b2 PE=1 SV=2	5647	17834	93.9	7.85	1.193	152	[1.032]
258	HEMO_MOUSE	Hemopexin OS=Mus musculus GN=Hpx PE=1 SV=2	62	55053	5.4	7.92	1.1	3	1.047
186	ROA0_MOUSE	Heterogeneous nuclear ribonucleoprotein A0 OS=Mus musculus GN=Hnrnpa0 PE=1 SV=1	94	33388	2.3	9.35	0.957	3	1.074
145	ROA1_MOUSE	Heterogeneous nuclear ribonucleoprotein A1 OS=Mus musculus GN=Hnrnpa1 PE=1 SV=2	128	37005	7.2	9.27	1.017	5	1.067
149	ROA3_MOUSE	Heterogeneous nuclear ribonucleoprotein A3 OS=Mus musculus GN=Hnrnpa3 PE=1 SV=1	122	42982	4.2	9.1	0.962	4	1.052
195	HNRPF_MOUSE	Heterogeneous nuclear ribonucleoprotein F OS=Mus musculus GN=Hnrnpf PE=1 SV=3	88	48427	3.9	5.31	1.08	2	1.001
136	HNRPK_MOUSE	Heterogeneous nuclear ribonucleoprotein K OS=Mus musculus GN=Hnrnpk PE=1 SV=1	135	54489	7.3	5.39	0.94	4	1.106
138	ROA2_MOUSE	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Mus musculus GN=Hnrnpa2b1 PE=1 SV=2	135	40308	6.8	8.97	1.004	5	1.07
133	HMGB1_MOUSE	High mobility group protein B1 OS=Mus musculus GN=Hmgb1 PE=1 SV=2	141	31357	21.9	5.62	0.907	7	1.052
303	HMGB2_MOUSE	High mobility group protein B2 OS=Mus musculus GN=Hmgb2 PE=1 SV=3	48	30193	10.5	6.88	1.034	2	1.094
155	HCDH_MOUSE	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Mus musculus GN=Hadh PE=1 SV=2	119	39047	7	8.76	0.819	4	1.047
243	GLO2_MOUSE	Hydroxyacylglutathione hydrolase, mitochondrial OS=Mus musculus GN=Hagh PE=1 SV=2	67	37889	3.9	7.66	1.003	2	1.131
300	IGHM_MOUSE	Ig mu chain C region OS=Mus musculus GN=Ighm PE=1 SV=2	49	54383	1.5	6.56	0.964	2	1.043
35	INMT_MOUSE	Indolethylamine N-methyltransferase OS=Mus musculus GN=Inmt PE=1 SV=1	505	32541	37.5	6	0.957	14	1.04
129	ITB1_MOUSE	Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1	147	99008	4	5.68	0.965	4	1.022
268	IRF8_MOUSE	Interferon regulatory factor 8 OS=Mus musculus GN=Irf8 PE=1 SV=1	58	52498	1.9	6.38	0.892	2	1.114

168	IDH3A_MOUSE	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Mus musculus GN=Idh3a PE=1 SV=1	105	43728	3	6.27	0.868	2	1.013
112	IDHP_MOUSE	Isocitrate dehydrogenase [NADP], mitochondrial OS=Mus musculus GN=Idh2 PE=1 SV=3	176	56718	5.5	8.88	0.917	6	1.06
196	PLAK_MOUSE	Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3	88	87102	3.1	5.75	0.963	3	1.101
218	KATL2_MOUSE	Katanin p60 ATPase-containing subunit A-like 2 OS=Mus musculus GN=Katnal2 PE=2 SV=2	80	67489	2.2	7.19	1.075	2	1.19
228	K1C18_MOUSE	Keratin, type I cytoskeletal 18 OS=Mus musculus GN=Krt18 PE=1 SV=5	76	50968	5.9	5.22	1.048	3	1.018
127	K1C19_MOUSE	Keratin, type I cytoskeletal 19 OS=Mus musculus GN=Krt19 PE=1 SV=1	150	47397	13.9	5.28	1.139	6	1.052
262	K22O_MOUSE	Keratin, type II cytoskeletal 2 oral OS=Mus musculus GN=Krt76 PE=2 SV=1	61	68408	2.7	8.68	0.964	2	1.291
246	K2C7_MOUSE	Keratin, type II cytoskeletal 7 OS=Mus musculus GN=Krt7 PE=1 SV=1	66	54713	6.1	5.67	1.11	3	1.207
120	K2C8_MOUSE	Keratin, type II cytoskeletal 8 OS=Mus musculus GN=Krt8 PE=1 SV=4	158	59431	9.8	5.7	1.084	6	1.084
160	KNG1_MOUSE	Kininogen-1 OS=Mus musculus GN=Kng1 PE=1 SV=1	115	79982	4.2	6.05	0.98	4	1.06
287	LGUL_MOUSE	Lactoylglutathione lyase OS=Mus musculus GN=Glo1 PE=1 SV=3	51	23672	9.2	5.24	0.857	2	1.129
282	LAMC1_MOUSE	Laminin subunit gamma-1 OS=Mus musculus GN=Lamc1 PE=1 SV=2	53	193266	0.7	5.08	0.977	2	1.001
289	LKHA4_MOUSE	Leukotriene A-4 hydrolase OS=Mus musculus GN=Lta4h PE=1 SV=4	51	75133	2.9	5.98	1.077	2	1.087
84	LDHA_MOUSE	L-lactate dehydrogenase A chain OS=Mus musculus GN=Ldha PE=1 SV=3	245	40786	28.6	7.62	0.936	12	1.042
111	LDHB_MOUSE	L-lactate dehydrogenase B chain OS=Mus musculus GN=Ldhb PE=1 SV=2	178	40526	12.9	5.7	0.83	6	1.071
119	ACADL_MOUSE	Long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Acadl PE=1 SV=2	158	52378	10.2	8.53	0.715	5	1.063
77	MDHC_MOUSE	Malate dehydrogenase, cytoplasmic OS=Mus musculus GN=Mdh1 PE=1 SV=3	265	41093	20.4	6.16	0.804	11	1.033
53	MDHM_MOUSE	Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE=1 SV=3	352	39703	38.2	8.93	0.737	13	1.086
177	MAP4_MOUSE	Microtubule-associated protein 4 OS=Mus musculus GN=Map4 PE=1 SV=3	99	131074	2.9	4.9	0.932	3	1.175
8	MOES_MOUSE	Moesin OS=Mus musculus GN=Msn PE=1 SV=3	1387	77184	45.4	6.22	0.98	52	1.026
139	MUG1_MOUSE	Murinoglobulin-1 OS=Mus musculus GN=Mug1 PE=1 SV=3	134	180032	2.7	6	1.019	5	1.042
19	MYG_MOUSE	Myoglobin OS=Mus musculus GN=Mb PE=1 SV=3	642	20131	40.3	7.07	0.755	16	1.041
292	MYLK_MOUSE	Myosin light chain kinase, smooth muscle OS=Mus musculus GN=Mylk PE=1 SV=3	50	236379	1.2	5.92	0.927	2	1.05
251	MYL6_MOUSE	Myosin light polypeptide 6 OS=Mus musculus GN=Myl6 PE=1 SV=3	64	18498	13.9	4.56	0.872	2	1.08
28	MARCS_MOUSE	Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Marcks PE=1 SV=2	534	33148	21.7	4.34	0.963	12	1.042
224	DDAH1_MOUSE	N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 OS=Mus musculus GN=Ddah1 PE=1 SV=3	77	34421	3.5	5.64	0.962	2	1.108
227	NHRF2_MOUSE	Na(+)/H(+) exchange regulatory cofactor NHE-RF2 OS=Mus musculus GN=Slc9a3r2 PE=1 SV=2	77	39817	8.3	7.23	1.087	3	1.009
169	YBOX1_MOUSE	Nuclease-sensitive element-binding protein 1 OS=Mus musculus GN=Ybx1 PE=1 SV=3	104	38158	13	9.87	0.951	2	1.183
135	NUCL_MOUSE	Nucleolin OS=Mus musculus GN=Ncl PE=1 SV=2	137	89836	7.6	4.69	1.076	5	1.066
104	NPM_MOUSE	Nucleophosmin OS=Mus musculus GN=Npm1 PE=1 SV=1	195	37577	14.7	4.62	1.084	7	1.047
92	NDKB_MOUSE	Nucleoside diphosphate kinase B OS=Mus musculus GN=Nme2 PE=1 SV=1	215	19317	25.7	6.97	0.901	7	1.035
106	PTMS_MOUSE	Parathymosin OS=Mus musculus GN=Ptms PE=1 SV=3	188	13441	21.8	4.17	0.973	3	1.039

71	PPIA_MOUSE	Peptidyl-prolyl cis-trans isomerase A OS=Mus musculus GN=Ppia PE=1 SV=2	279	20115	38.4	7.74	1.072	11	1.044
152	FKB1A_MOUSE	Peptidyl-prolyl cis-trans isomerase FKBP1A OS=Mus musculus GN=Fkbp1a PE=1 SV=2	121	13258	16.7	7.88	0.897	4	1.017
103	PRDX1_MOUSE	Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 PE=1 SV=1	199	25228	23.6	8.26	0.984	8	1.038
50	PRDX2_MOUSE	Peroxiredoxin-2 OS=Mus musculus GN=Prdx2 PE=1 SV=3	367	23920	31.8	5.2	0.972	13	1.041
182	PRDX4_MOUSE	Peroxiredoxin-4 OS=Mus musculus GN=Prdx4 PE=1 SV=1	95	33667	6.9	6.67	0.94	3	[1.088]
153	PRDX5_MOUSE	Peroxiredoxin-5, mitochondrial OS=Mus musculus GN=Prdx5 PE=1 SV=2	121	24465	18.1	9.1	0.89	4	1.052
15	PRDX6_MOUSE	Peroxiredoxin-6 OS=Mus musculus GN=Prdx6 PE=1 SV=3	804	27685	71.4	5.71	1.04	29	1.03
131	PEBP1_MOUSE	Phosphatidylethanolamine-binding protein 1 OS=Mus musculus GN=Pebp1 PE=1 SV=3	145	22829	11.2	5.19	0.907	3	1.04
128	PGK1_MOUSE	Phosphoglycerate kinase 1 OS=Mus musculus GN=Pgk1 PE=1 SV=4	149	50896	11	8.02	0.962	7	1.053
91	PGAM1_MOUSE	Phosphoglycerate mutase 1 OS=Mus musculus GN=Pgam1 PE=1 SV=3	217	31644	16.5	6.67	0.892	7	1.034
165	PLST_MOUSE	Plastin-3 OS=Mus musculus GN=Pls3 PE=1 SV=3	107	78460	5.2	5.42	0.946	4	1.118
271	CD36_MOUSE	Platelet glycoprotein 4 OS=Mus musculus GN=Cd36 PE=1 SV=2	57	58600	4.4	8.6	0.899	2	1.007
257	PCBP1_MOUSE	Poly(rC)-binding protein 1 OS=Mus musculus GN=Pcbp1 PE=1 SV=1	63	40049	5.3	6.66	0.95	2	1.057
17	PTRF_MOUSE	Polymerase I and transcript release factor OS=Mus musculus GN=Ptrf PE=1 SV=1	690	49691	28.3	5.43	0.794	17	1.04
167	LMNA_MOUSE	Prelamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2	106	80187	3.3	6.54	1.134	2	1.002
25	PROF1_MOUSE	Profilin-1 OS=Mus musculus GN=Pfn1 PE=1 SV=2	569	16526	52.9	8.46	0.921	20	1.044
183	PSA1_MOUSE	Proteasome subunit alpha type-1 OS=Mus musculus GN=Psma1 PE=1 SV=1	95	31631	8.7	6	0.824	3	1.091
175	PSA7_MOUSE	Proteasome subunit alpha type-7 OS=Mus musculus GN=Psma7 PE=1 SV=1	101	31434	8.1	8.59	0.909	3	1.02
200	PSB7_MOUSE	Proteasome subunit beta type-7 OS=Mus musculus GN=Psmb7 PE=1 SV=1	85	33030	6.9	8.14	0.928	3	1.119
87	PDIA3_MOUSE	Protein disulfide-isomerase A3 OS=Mus musculus GN=Pdia3 PE=1 SV=2	237	64504	13.7	5.88	1.008	8	1.034
161	PDIA4_MOUSE	Protein disulfide-isomerase A4 OS=Mus musculus GN=Pdia4 PE=1 SV=3	114	82012	4.9	5.16	0.969	4	1.066
276	PDIA6_MOUSE	Protein disulfide-isomerase A6 OS=Mus musculus GN=Pdia6 PE=1 SV=3	55	53292	5.2	5	1.044	3	1.037
85	PDIA1_MOUSE	Protein disulfide-isomerase OS=Mus musculus GN=P4hb PE=1 SV=2	243	64694	13.4	4.77	1.029	7	1.033
100	PARK7_MOUSE	Protein DJ-1 OS=Mus musculus GN=Park7 PE=1 SV=1	203	22498	37	6.32	0.886	7	1.062
198	S10A6_MOUSE	Protein S100-A6 OS=Mus musculus GN=S100a6 PE=1 SV=3	86	11675	16.9	5.3	1.034	3	1.051
122	S10A9_MOUSE	Protein S100-A9 OS=Mus musculus GN=S100a9 PE=1 SV=3	156	14907	10.6	6.64	2.09	4	1.034
22	PTMA_MOUSE	Prothymosin alpha OS=Mus musculus GN=Ptma PE=1 SV=2	597	13544	23.4	3.7	1.024	16	1.037
42	KPYM_MOUSE	Pyruvate kinase PKM OS=Mus musculus GN=Pkm PE=1 SV=4	428	63744	25.4	7.18	0.873	14	1.035
261	RABL3_MOUSE	Rab-like protein 3 OS=Mus musculus GN=Rabl3 PE=2 SV=1	62	28627	4.7	6.66	0.918	2	1.206
40	RADI_MOUSE	Radixin OS=Mus musculus GN=Rdx PE=1 SV=3	448	77959	15.4	5.91	0.941	19	1.046
159	IQGA1_MOUSE	Ras GTPase-activating-like protein IQGAP1 OS=Mus musculus GN=Iqgap1 PE=1 SV=2	115	208290	2.1	6.07	0.966	5	1.083
181	RAC1_MOUSE	Ras-related C3 botulinum toxin substrate 1 OS=Mus musculus GN=Rac1 PE=1 SV=1	96	24352	13	8.77	1.151	2	1.107

144	RAB10_MOUSE	Ras-related protein Rab-10 OS=Mus musculus GN=Rab10 PE=1 SV=1	129	25881	17	8.59	1.069	4	1.034
209	RB11A_MOUSE	Ras-related protein Rab-11A OS=Mus musculus GN=Rab11a PE=1 SV=3	82	26344	5.1	6.12	0.918	2	1.206
171	RAB15_MOUSE	Ras-related protein Rab-15 OS=Mus musculus GN=Rab15 PE=1 SV=1	103	26839	10.4	5.5	1.073	3	1.046
101	RAB1A_MOUSE	Ras-related protein Rab-1A OS=Mus musculus GN=Rab1A PE=1 SV=3	199	25441	25.9	5.93	0.979	6	1.076
117	RAB1B_MOUSE	Ras-related protein Rab-1B OS=Mus musculus GN=Rab1b PE=1 SV=1	162	24761	21.9	5.55	1.025	5	1.037
172	RAB35_MOUSE	Ras-related protein Rab-35 OS=Mus musculus GN=Rab35 PE=1 SV=1	103	25979	10.9	8.52	1.073	3	1.046
125	RAP1B_MOUSE	Ras-related protein Rap-1b OS=Mus musculus GN=Rap1b PE=2 SV=2	153	23157	14.1	5.65	0.992	3	1.045
267	RRAS2_MOUSE	Ras-related protein R-Ras2 OS=Mus musculus GN=Rras2 PE=1 SV=1	59	25586	5.9	5.74	1.007	2	1.084
280	RTN3_MOUSE	Reticulon-3 OS=Mus musculus GN=Rtn3 PE=1 SV=2	54	114263	1.5	4.83	0.949	2	1.175
31	AL1A1_MOUSE	Retinal dehydrogenase 1 OS=Mus musculus GN=Aldh1a1 PE=1 SV=5	517	60847	20.8	7.92	1.014	17	1.05
78	GDIR1_MOUSE	Rho GDP-dissociation inhibitor 1 OS=Mus musculus GN=Arhgdia PE=1 SV=3	261	26321	18.6	5.12	0.899	10	1.057
38	SBP1_MOUSE	Selenium-binding protein 1 OS=Mus musculus GN=Selenbp1 PE=1 SV=2	448	56831	25.8	5.87	0.974	16	1.022
269	SEPT2_MOUSE	Septin-2 OS=Mus musculus GN=Sept2 PE=1 SV=2	57	45574	2.2	6.1	0.897	2	1.041
264	SEPT7_MOUSE	Septin-7 OS=Mus musculus GN=Sept7 PE=1 SV=1	60	57567	2.3	8.73	0.998	2	1.104
18	SPA3K_MOUSE	Serine protease inhibitor A3K OS=Mus musculus GN=Serpina3k PE=1 SV=2	643	51311	29.4	5.05	1.132	19	1.048
180	2AAA_MOUSE	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Mus musculus GN=Ppp2r1a PE=1 SV=3	98	70680	3.1	5	1.077	3	1.058
13	TRFE_MOUSE	Serotransferrin OS=Mus musculus GN=Tf PE=1 SV=1	890	86923	30.6	6.94	0.938	34	1.04
140	SPB6_MOUSE	Serpin B6 OS=Mus musculus GN=Serpinb6 PE=1 SV=1	132	47602	9	5.53	0.994	4	[1.042]
46	SDPR_MOUSE	Serum deprivation-response protein OS=Mus musculus GN=Sdpr PE=1 SV=3	390	52113	23.7	5.15	0.872	14	1.074
229	DHSO_MOUSE	Sorbitol dehydrogenase OS=Mus musculus GN=Sord PE=1 SV=3	76	42431	2.5	6.56	0.969	2	1.028
304	SORCN_MOUSE	Sorcin OS=Mus musculus GN=Sri PE=1 SV=1	48	22852	5.6	5.32	1.19	2	1.231
51	SPTN1_MOUSE	Spectrin alpha chain, non-erythrocytic 1 OS=Mus musculus GN=Sptan1 PE=1 SV=4	363	313742	5.1	5.2	0.998	15	1.04
207	SPTB2_MOUSE	Spectrin beta chain, non-erythrocytic 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2	83	300825	1.3	5.4	1.029	3	1.02
247	GRP75_MOUSE	Stress-70 protein, mitochondrial OS=Mus musculus GN=Hspa9 PE=1 SV=3	66	81427	4.7	5.81	0.903	3	1.05
214	SUCA_MOUSE	Succinyl-CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial OS=Mus musculus GN=Suclg1 PE=1 SV=4	80	39866	2.6	9.46	0.884	2	1.004
237	SAP_MOUSE	Sulfated glycoprotein 1 OS=Mus musculus GN=Psap PE=1 SV=2	70	68519	1.8	5.07	1.101	2	1.034
45	SODC_MOUSE	Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2	392	17512	39	6.02	0.933	11	1.05
121	SODM_MOUSE	Superoxide dismutase [Mn], mitochondrial OS=Mus musculus GN=Sod2 PE=1 SV=3	156	27221	16.2	8.8	0.885	4	1.044
143	VAT1_MOUSE	Synaptic vesicle membrane protein VAT-1 homolog OS=Mus musculus GN=Vat1 PE=1 SV=3	130	46135	9.9	5.95	0.869	5	1.052
10	TLN1_MOUSE	Talin-1 OS=Mus musculus GN=Tln1 PE=1 SV=2	994	295033	10.8	5.84	1	31	1.032
240	TADBP_MOUSE	TAR DNA-binding protein 43 OS=Mus musculus GN=Tardbp PE=1 SV=1	68	48011	2.9	6.26	0.922	2	1.106
212	TCPA_MOUSE	T-complex protein 1 subunit alpha OS=Mus musculus GN=Tcp1 PE=1 SV=3	80	66687	2.2	5.82	1.061	2	1.185

211	TCPG_MOUSE	T-complex protein 1 subunit gamma OS=Mus musculus GN=Cct3 PE=1 SV=1	81	66239	1.8	6.28	0.995	3	1.041
174	TCPQ_MOUSE	T-complex protein 1 subunit theta OS=Mus musculus GN=Cct8 PE=1 SV=3	102	66318	3.6	5.44	0.942	3	1.07
255	TSN8_MOUSE	Tetraspanin-8 OS=Mus musculus GN=Tspan8 PE=2 SV=1	63	28279	4.3	5.08	0.813	2	1.107
97	THIO_MOUSE	Thioredoxin OS=Mus musculus GN=Txn PE=1 SV=3	207	13817	31.4	4.8	1.064	5	1.047
345	TYB4_MOUSE	Thymosin beta-4 OS=Mus musculus GN=Tmsb4x PE=1 SV=1	39	7117	26	5.02	0.934	2	1.111
221	ZO1_MOUSE	Tight junction protein ZO-1 OS=Mus musculus GN=Tjp1 PE=1 SV=2	78	210652	1.4	6.17	0.932	3	1.096
90	TALDO_MOUSE	Transaldolase OS=Mus musculus GN=Taldo1 PE=1 SV=2	219	42257	16.6	6.57	0.866	6	1.03
154	RHOA_MOUSE	Transforming protein RhoA OS=Mus musculus GN=Rhoa PE=1 SV=1	121	24638	23.8	5.83	1.028	5	1.125
79	TAGL_MOUSE	Transgelin OS=Mus musculus GN=Tagln PE=1 SV=3	260	25201	29.4	8.85	0.988	8	1.036
26	TAGL2_MOUSE	Transgelin-2 OS=Mus musculus GN=Tagln2 PE=1 SV=4	554	24537	50.3	8.39	1.012	19	1.043
82	TERA_MOUSE	Transitional endoplasmic reticulum ATPase OS=Mus musculus GN=Vcp PE=1 SV=4	246	96734	11.8	5.14	1.094	10	1.108
68	TKT_MOUSE	Transketolase OS=Mus musculus GN=Tkt PE=1 SV=1	287	74624	18.3	7.23	0.928	13	1.074
107	TPIS_MOUSE	Triosephosphate isomerase OS=Mus musculus GN=Tpi1 PE=1 SV=4	186	35900	24.1	5.56	0.892	8	1.024
142	TPM1_MOUSE	Tropomyosin alpha-1 chain OS=Mus musculus GN=Tpm1 PE=1 SV=1	131	38471	12.3	4.69	0.956	7	1.085
109	TPM3_MOUSE	Tropomyosin alpha-3 chain OS=Mus musculus GN=Tpm3 PE=1 SV=3	184	38640	16.8	4.68	0.954	9	1.069
52	TPM4_MOUSE	Tropomyosin alpha-4 chain OS=Mus musculus GN=Tpm4 PE=2 SV=3	355	32289	28.2	4.65	0.952	12	1.041
57	TBA1A_MOUSE	Tubulin alpha-1A chain OS=Mus musculus GN=Tuba1a PE=1 SV=1	328	53538	18.2	4.94	1.1	11	1.058
130	TBB4B_MOUSE	Tubulin beta-4B chain OS=Mus musculus GN=Tubb4b PE=1 SV=1	146	52473	15.1	4.79	1.047	6	1.19
132	TBB5_MOUSE	Tubulin beta-5 chain OS=Mus musculus GN=Tubb5 PE=1 SV=1	144	52313	18.5	4.78	1.113	5	1.067
137	TPPP3_MOUSE	Tubulin polymerization-promoting protein family member 3 OS=Mus musculus GN=Tppp3 PE=1 SV=1	135	23127	17	9.18	0.876	4	[1.052]
73	RS27A_MOUSE	Ubiquitin-40S ribosomal protein S27a OS=Mus musculus GN=Rps27a PE=1 SV=2	273	21962	25.6	9.68	1.061	10	1.046
202	UB2L3_MOUSE	Ubiquitin-conjugating enzyme E2 L3 OS=Mus musculus GN=Ube2l3 PE=2 SV=1	84	20726	5.8	8.68	1.046	2	1.068
197	UBE2N_MOUSE	Ubiquitin-conjugating enzyme E2 N OS=Mus musculus GN=Ube2n PE=1 SV=1	88	18614	7.2	6.13	0.972	2	1.043
231	UBA1_MOUSE	Ubiquitin-like modifier-activating enzyme 1 OS=Mus musculus GN=Uba1 PE=1 SV=1	74	126625	1.5	5.43	0.965	3	1.096
76	UTER_MOUSE	Uteroglobin OS=Mus musculus GN=Scgb1a1 PE=1 SV=1	265	11705	19.8	6.15	0.747	7	1.047
33	VIME_MOUSE	Vimentin OS=Mus musculus GN=Vim PE=1 SV=3	509	57015	31.1	5.06	1.155	18	1.068
6	VINC_MOUSE	Vinculin OS=Mus musculus GN=Vcl PE=1 SV=4	1612	128056	37.4	5.77	0.946	59	[1.025]