

Protein hit no.	Accession no.	Protein description	Protein scor
149	CH10_MOUSE	10 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspe1 PE=1 SV=2	57
43	1433B_MOUSE	14-3-3 protein beta/alpha OS=Mus musculus GN=Ywhab PE=1 SV=3	189
39	1433E_MOUSE	14-3-3 protein epsilon OS=Mus musculus GN=Ywhae PE=1 SV=1	193
52	1433F_MOUSE	14-3-3 protein eta OS=Mus musculus GN=Ywhah PE=1 SV=2	155
40	1433G_MOUSE	14-3-3 protein gamma OS=Mus musculus GN=Ywhag PE=1 SV=2	193
60	1433S_MOUSE	14-3-3 protein sigma OS=Mus musculus GN=Sfn PE=1 SV=2	127
22	1433Z_MOUSE	14-3-3 protein zeta/delta OS=Mus musculus GN=Ywhaz PE=1 SV=1	279
79	RS19_MOUSE	40S ribosomal protein S19 OS=Mus musculus GN=Rps19 PE=1 SV=3	104
152	RLA0_MOUSE	60S acidic ribosomal protein P0 OS=Mus musculus GN=Rplp0 PE=1 SV=3	56
48	GRP78_MOUSE	78 kDa glucose-regulated protein OS=Mus musculus GN=Hspa5 PE=1 SV=3	167
96	AN32A_MOUSE	Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Mus musculus GN=Anp32a PE=1 SV=1	82
6	ACTB_MOUSE	Actin, cytoplasmic 1 OS=Mus musculus GN=Actb PE=1 SV=1	835
108	CAP1_MOUSE	Adenylyl cyclase-associated protein 1 OS=Mus musculus GN=Cap1 PE=1 SV=4	75
147	ARF1_MOUSE	ADP-ribosylation factor 1 OS=Mus musculus GN=Arf1 PE=1 SV=2	57
51	RAGE_MOUSE	Advanced glycosylation end product-specific receptor OS=Mus musculus GN=Ager PE=1 SV=1	157
65	AL1A7_MOUSE	Aldehyde dehydrogenase, cytosolic 1 OS=Mus musculus GN=Aldh1a7 PE=2 SV=1	119
89	ALDH2_MOUSE	Aldehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh2 PE=1 SV=1	90
86	ALDR_MOUSE	Aldose reductase OS=Mus musculus GN=Akr1b1 PE=1 SV=3	92
82	A1AT2_MOUSE	Alpha-1-antitrypsin 1-2 OS=Mus musculus GN=Serpina1b PE=1 SV=2	95
49	A1AT3_MOUSE	Alpha-1-antitrypsin 1-3 OS=Mus musculus GN=Serpina1c PE=1 SV=2	163
83	A1AT4_MOUSE	Alpha-1-antitrypsin 1-4 OS=Mus musculus GN=Serpina1d PE=2 SV=1	95
106	FETUA_MOUSE	Alpha-2-HS-glycoprotein OS=Mus musculus GN=Ahsg PE=1 SV=1	76
15	A2M_MOUSE	Alpha-2-macroglobulin OS=Mus musculus GN=A2m PE=1 SV=3	355

159	ACTN4_MOUSE E	Alpha-actinin-4 OS=Mus musculus GN=Actn4 PE=1 SV=1	52
24	ENOA_MOUSE	Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3	243
121	ACE_MOUSE	Angiotensin-converting enzyme OS=Mus musculus GN=Ace PE=1 SV=3	67
131	ANXA2_MOUSE E	Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	64
37	ANXA5_MOUSE E	Annexin A5 OS=Mus musculus GN=Anxa5 PE=1 SV=1	198
36	APOA1_MOUSE E	Apolipoprotein A-I OS=Mus musculus GN=Apoa1 PE=1 SV=2	202
130	BCAM_MOUSE E	Basal cell adhesion molecule OS=Mus musculus GN=Bcam PE=2 SV=1	64
20	ACTBL_MOUSE E	Beta-actin-like protein 2 OS=Mus musculus GN=Actbl2 PE=1 SV=1	300
104	ENOB_MOUSE	Beta-enolase OS=Mus musculus GN=Eno3 PE=1 SV=3	77
66	CALM_MOUSE	Calmodulin OS=Mus musculus GN=Calm1 PE=1 SV=2	119
132	CALX_MOUSE	Calnexin OS=Mus musculus GN=Canx PE=1 SV=1	63
54	CALR_MOUSE	Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1	146
28	CAH2_MOUSE	Carbonic anhydrase 2 OS=Mus musculus GN=Ca2 PE=1 SV=4	232
163	CAH3_MOUSE	Carbonic anhydrase 3 OS=Mus musculus GN=Ca3 PE=1 SV=3	49
5	CBR2_MOUSE	Carbonyl reductase [NADPH] 2 OS=Mus musculus GN=Cbr2 PE=1 SV=1	1255
44	EST1C_MOUSE E	Carboxylesterase 1C OS=Mus musculus GN=Ces1c PE=1 SV=4	189
33	CES1D_MOUSE E	Carboxylesterase 1D OS=Mus musculus GN=Ces1d PE=1 SV=1	211
129	CATA_MOUSE	Catalase OS=Mus musculus GN=Cat PE=1 SV=4	64
94	CDC42_MOUSE E	Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	83
157	CLIC5_MOUSE	Chloride intracellular channel protein 5 OS=Mus musculus GN=Clic5 PE=1 SV=1	54
110	CISY_MOUSE	Citrate synthase, mitochondrial OS=Mus musculus GN=Cs PE=1 SV=1	73
155	COF1_MOUSE	Cofilin-1 OS=Mus musculus GN=Cfl1 PE=1 SV=3	56
67	CO3_MOUSE	Complement C3 OS=Mus musculus GN=C3 PE=1 SV=3	119
190	COR1C_MOUSE SE	Coronin-1C OS=Mus musculus GN=Coro1c PE=1 SV=2	41
141	KCRM_MOUSE E	Creatine kinase M-type OS=Mus musculus GN=Ckm PE=1 SV=1	58

100	CRIP2_MOUSE E	Cysteine-rich protein 2 OS=Mus musculus GN=Crip2 PE=1 SV=1	79
71	CYC_MOUSE	Cytochrome c, somatic OS=Mus musculus GN=Cyccs PE=1 SV=2	117
27	DPYL2_MOUSE E	Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=1 SV=2	233
93	EHD2_MOUSE	EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1	85
127	EF1A1_MOUSE E	Elongation factor 1-alpha 1 OS=Mus musculus GN=Eef1a1 PE=1 SV=3	65
165	EF1B_MOUSE	Elongation factor 1-beta OS=Mus musculus GN=Eef1b PE=1 SV=5	49
103	ENPL_MOUSE	Endoplasmic reticulum protein OS=Mus musculus GN=Hsp90b1 PE=1 SV=2	78
119	FABPH_MOUSE E	Fatty acid-binding protein, heart OS=Mus musculus GN=Fabp3 PE=1 SV=5	67
35	FLNA_MOUSE	Filamin-A OS=Mus musculus GN=Flna PE=1 SV=5	206
38	ALDOA_MOUSE E	Fructose-bisphosphate aldolase A OS=Mus musculus GN=Aldoa PE=1 SV=2	197
135	GELS_MOUSE	Gelsolin OS=Mus musculus GN=Gsn PE=1 SV=3	61
50	GSTA3_MOUSE E	Glutathione S-transferase A3 OS=Mus musculus GN=Gsta3 PE=1 SV=2	158
140	GSTA4_MOUSE E	Glutathione S-transferase A4 OS=Mus musculus GN=Gsta4 PE=1 SV=3	58
30	GSTM1_MOUSE E	Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2	231
42	GSTM2_MOUSE E	Glutathione S-transferase Mu 2 OS=Mus musculus GN=Gstm2 PE=1 SV=2	190
47	GSTM4_MOUSE E	Glutathione S-transferase Mu 3 OS=Mus musculus GN=Gstm3 PE=1 SV=2	168
73	G3P_MOUSE	Glyceraldehyde-3-phosphate dehydrogenase OS=Mus musculus GN=Gapdh PE=1 SV=2	115
117	GIMA4_MOUSE E	GTPase IMAP family member 4 OS=Mus musculus GN=Gimap4 PE=1 SV=1	69
13	HSP7C_MOUSE E	Heat shock cognate 71 kDa protein OS=Mus musculus GN=Hspa8 PE=1 SV=1	388
31	HS90A_MOUSE E	Heat shock protein HSP 90-alpha OS=Mus musculus GN=Hsp90aa1 PE=1 SV=4	223
26	HS90B_MOUSE E	Heat shock protein HSP 90-beta OS=Mus musculus GN=Hsp90ab1 PE=1 SV=3	237
3	HBA_MOUSE	Hemoglobin subunit alpha OS=Mus musculus GN=Hba PE=1 SV=2	5628
1	HBB1_MOUSE	Hemoglobin subunit beta-1 OS=Mus musculus GN=Hbb-b1 PE=1 SV=2	9233
2	HBB2_MOUSE	Hemoglobin subunit beta-2 OS=Mus musculus GN=Hbb-b2 PE=1 SV=2	6521
173	HEMO_MOUSE E	Hemopexin OS=Mus musculus GN=Hpx PE=1 SV=2	47

113	ROA0_MOUSE	Heterogeneous nuclear ribonucleoprotein A0 OS=Mus musculus GN=Hnrnpa0 PE=1 SV=1	70
114	ROA1_MOUSE	Heterogeneous nuclear ribonucleoprotein A1 OS=Mus musculus GN=Hnrnpa1 PE=1 SV=2	70
115	ROA2_MOUSE	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Mus musculus GN=Hnrnpa2b1 PE=1 SV=2	70
18	INMT_MOUSE	Indolethylamine N-methyltransferase OS=Mus musculus GN=Inmt PE=1 SV=1	328
87	K1C18_MOUSE E	Keratin, type I cytoskeletal 18 OS=Mus musculus GN=Krt18 PE=1 SV=5	90
80	K2C1_MOUSE	Keratin, type II cytoskeletal 1 OS=Mus musculus GN=Krt1 PE=1 SV=4	104
58	K2C1B_MOUSE E	Keratin, type II cytoskeletal 1b OS=Mus musculus GN=Krt77 PE=1 SV=1	136
125	K22O_MOUSE	Keratin, type II cytoskeletal 2 oral OS=Mus musculus GN=Krt76 PE=2 SV=1	66
46	K2C5_MOUSE	Keratin, type II cytoskeletal 5 OS=Mus musculus GN=Krt5 PE=1 SV=1	176
116	K2C8_MOUSE	Keratin, type II cytoskeletal 8 OS=Mus musculus GN=Krt8 PE=1 SV=4	70
74	KNG1_MOUSE	Kininogen-1 OS=Mus musculus GN=Kng1 PE=1 SV=1	114
19	MDHM_MOUSE E	Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE=1 SV=3	301
9	MOES_MOUSE E	Moesin OS=Mus musculus GN=Msn PE=1 SV=3	646
45	MYG_MOUSE	Myoglobin OS=Mus musculus GN=Mb PE=1 SV=3	178
185	MYLK_MOUSE	Myosin light chain kinase, smooth muscle OS=Mus musculus GN=Myk PE=1 SV=3	43
68	MARCS_MOUSE SE	Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Marcks PE=1 SV=2	118
69	HMG2_MOUSE SE	Non-histone chromosomal protein HMG-17 OS=Mus musculus GN=Hmgn2 PE=1 SV=2	117
97	NPM_MOUSE	Nucleophosmin OS=Mus musculus GN=Npm1 PE=1 SV=1	82
122	NDKA_MOUSE	Nucleoside diphosphate kinase A OS=Mus musculus GN=Nme1 PE=1 SV=1	66
105	NDKB_MOUSE	Nucleoside diphosphate kinase B OS=Mus musculus GN=Nme2 PE=1 SV=1	76
75	PTMS_MOUSE	Parathyrosin OS=Mus musculus GN=Ptms PE=1 SV=3	112
78	PRDX1_MOUSE E	Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 PE=1 SV=1	110
29	PRDX2_MOUSE E	Peroxiredoxin-2 OS=Mus musculus GN=Prdx2 PE=1 SV=3	232
10	PRDX6_MOUSE E	Peroxiredoxin-6 OS=Mus musculus GN=Prdx6 PE=1 SV=3	549
91	PEBP1_MOUSE E	Phosphatidylethanolamine-binding protein 1 OS=Mus musculus GN=Pebp1 PE=1 SV=3	85

77	PGAM1_MOUSE E	Phosphoglycerate mutase 1 OS=Mus musculus GN=Pgam1 PE=1 SV=3	110
95	PLST_MOUSE	Plastin-3 OS=Mus musculus GN=Pls3 PE=1 SV=3	82
32	PTRF_MOUSE	Polymerase I and transcript release factor OS=Mus musculus GN=Ptrf PE=1 SV=1	216
133	LMNA_MOUSE	Prelamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2	63
34	PROF1_MOUSE E	Profilin-1 OS=Mus musculus GN=Pfn1 PE=1 SV=2	210
138	SAP_MOUSE	Prosaposin OS=Mus musculus GN=Psap PE=1 SV=2	60
109	PDIA3_MOUSE E	Protein disulfide-isomerase A3 OS=Mus musculus GN=Pdia3 PE=1 SV=2	74
70	PDIA1_MOUSE E	Protein disulfide-isomerase OS=Mus musculus GN=P4hb PE=1 SV=2	117
90	S10A6_MOUSE E	Protein S100-A6 OS=Mus musculus GN=S100a6 PE=1 SV=3	87
99	PTMA_MOUSE	Prothymosin alpha OS=Mus musculus GN=Ptma PE=1 SV=2	80
63	KPYM_MOUSE	Pyruvate kinase PKM OS=Mus musculus GN=Pkm PE=1 SV=4	122
84	RAB1A_MOUSE E	Ras-related protein Rab-1A OS=Mus musculus GN=Rab1A PE=1 SV=3	92
92	RAB1B_MOUSE E	Ras-related protein Rab-1B OS=Mus musculus GN=Rab1b PE=1 SV=1	85
16	AL1A1_MOUSE E	Retinal dehydrogenase 1 OS=Mus musculus GN=Aldh1a1 PE=1 SV=5	341
61	GDIR1_MOUSE E	Rho GDP-dissociation inhibitor 1 OS=Mus musculus GN=Arhgdia PE=1 SV=3	126
14	SBP1_MOUSE	Selenium-binding protein 1 OS=Mus musculus GN=Selenbp1 PE=1 SV=2	373
11	SPA3K_MOUSE E	Serine protease inhibitor A3K OS=Mus musculus GN=Serpina3k PE=1 SV=2	422
8	TRFE_MOUSE	Serotransferrin OS=Mus musculus GN=Tf PE=1 SV=1	759
85	SDPR_MOUSE	Serum deprivation-response protein OS=Mus musculus GN=Sdpr PE=1 SV=3	92
143	SUMO2_MOUSE SE	Small ubiquitin-related modifier 2 OS=Mus musculus GN=Sumo2 PE=1 SV=1	58
55	SPTN1_MOUSE E	Spectrin alpha chain, non-erythrocytic 1 OS=Mus musculus GN=Sptan1 PE=1 SV=4	142
111	SPTB2_MOUSE E	Spectrin beta chain, non-erythrocytic 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2	72
128	GRP75_MOUSE E	Stress-70 protein, mitochondrial OS=Mus musculus GN=Hspa9 PE=1 SV=3	65
12	SODC_MOUSE E	Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2	397
64	SODM_MOUSE E	Superoxide dismutase [Mn], mitochondrial OS=Mus musculus GN=Sod2 PE=1 SV=3	120

17	TLN1_MOUSE	Talin-1 OS=Mus musculus GN=Tln1 PE=1 SV=2	336
107	THIO_MOUSE	Thioredoxin OS=Mus musculus GN=Txn PE=1 SV=3	75
153	TYB4_MOUSE	Thymosin beta-4 OS=Mus musculus GN=Tmsb4x PE=1 SV=1	56
81	TALDO_MOUSE E	Transaldolase OS=Mus musculus GN=Taldo1 PE=1 SV=2	100
41	TAGL_MOUSE	Transgelin OS=Mus musculus GN=Tagln PE=1 SV=3	191
21	TAGL2_MOUSE E	Transgelin-2 OS=Mus musculus GN=Tagln2 PE=1 SV=4	293
123	TERA_MOUSE	Transitional endoplasmic reticulum ATPase OS=Mus musculus GN=Vcp PE=1 SV=4	66
76	TKT_MOUSE	Transketolase OS=Mus musculus GN=Tkt PE=1 SV=1	112
59	TPIS_MOUSE	Triosephosphate isomerase OS=Mus musculus GN=Tpi1 PE=1 SV=4	127
101	TPM1_MOUSE	Tropomyosin alpha-1 chain OS=Mus musculus GN=Tpm1 PE=1 SV=1	79
57	TPM3_MOUSE	Tropomyosin alpha-3 chain OS=Mus musculus GN=Tpm3 PE=1 SV=3	136
53	TPM4_MOUSE	Tropomyosin alpha-4 chain OS=Mus musculus GN=Tpm4 PE=2 SV=3	147
148	TBA1A_MOUSE E	Tubulin alpha-1A chain OS=Mus musculus GN=Tuba1a PE=1 SV=1	57
62	TPPP3_MOUSE E	Tubulin polymerization-promoting protein family member 3 OS=Mus musculus GN=Tppp3 PE=1 SV=1	123
25	RS27A_MOUSE E	Ubiquitin-40S ribosomal protein S27a OS=Mus musculus GN=Rps27a PE=1 SV=2	238
23	UTER_MOUSE	Uteroglobin OS=Mus musculus GN=Scgb1a1 PE=1 SV=1	269
56	VIME_MOUSE	Vimentin OS=Mus musculus GN=Vim PE=1 SV=3	139
7	VINC_MOUSE	Vinculin OS=Mus musculus GN=Vcl PE=1 SV=4	796

MW [kDa]	Sequence covered %	Protein PI	115/114 ratio (znop)	N	SD
12685	19.6	7.93	0.792	2	1.003
31187	22.4	4.77	1.07	6	1.101
32031	18.8	4.63	0.962	6	1.039
31358	16.3	4.81	1.065	7	[1.065]
31017	20.2	4.8	1.042	8	[1.055]
30519	8.5	4.72	1.091	5	1.081
30918	30.2	4.73	1.045	7	1.099
18381	13.8	10.41	1.058	2	1.029
37215	3.5	5.91	1.016	2	1.003
81404	9.9	5.07	1.027	8	1.059
31540	10.9	3.99	0.979	3	1.055
44868	36.3	5.29	1.425	20	1.063
57717	6.8	7.16	0.908	3	1.043
22459	14.4	6.32	1.248	2	1.129
45224	10.2	5.78	0.943	6	[1.027]
60397	8.6	7.6	1.019	4	1.113
61383	5.6	7.53	1.123	4	1.13
39876	10.8	6.71	0.885	2	1.058
49974	8.2	5.32	1.078	3	1.191
49535	10.7	5.25	1.072	5	1.192
49710	8.2	5.24	1.078	3	1.191
40107	14.2	6.04	1.012	3	1.088
180541	8.5	6.24	0.879	12	[1.081]

113493	2.9	5.25	0.881	2	1.096
52431	18.9	6.37	0.799	10	1.08
160446	2.1	6.1	0.905	3	1.063
43781	5	7.55	1.231	2	1.53
38946	26.6	4.83	0.999	4	1.035
33479	16.3	5.51	1.095	3	1.068
71153	5.1	5.85	1.022	2	1.054
44991	15.7	5.3	1.462	11	1.084
52747	8.1	6.73	0.594	4	1.077
18124	18.8	4.09	0.673	4	1.233
76060	1.7	4.5	1.071	2	1.031
54443	12	4.33	0.936	6	[1.095]
31989	38.1	6.49	0.863	5	1.139
32460	4.6	6.89	0.375	2	1.533
28339	70.5	9.1	1.128	29	1.062
65138	10.3	4.97	0.868	6	1.077
67022	14	6.17	0.882	9	1.059
64743	4	7.72	0.967	3	1.091
24115	9.9	6.15	1.046	2	1.007
31578	7.2	5.64	1.42	2	1.105
55968	6.9	8.72	0.891	2	1.005
22478	8.4	8.22	1.008	2	1.139
204900	3.2	6.29	1.033	4	1.085
58971	4.2	6.65	0.939	2	1.098
48101	4.7	6.58	0.578	2	1.017



26670	9.1	8.94	1.497	3	[1.091]
14428	28.6	9.61	0.708	4	1.093
67316	15	5.95	1.06	4	[1.052]
67038	3.9	6.08	1.214	3	1.244
57275	3.2	9.1	1.101	2	1.008
28275	6.2	4.53	0.944	2	1.055
103744	2.9	4.74	1.205	2	1.217
16827	11.3	6.11	0.893	2	1.052
306546	3.2	5.68	1.011	6	1.094
43590	17.6	8.31	0.84	6	1.071
92262	2.9	5.83	0.834	2	1.082
28416	27.1	8.76	1.23	5	1.108
28285	5.9	6.77	1.282	2	1.186
28783	40.4	7.71	1.096	8	1.084
29008	29.4	6.9	0.899	6	1.088
28319	22.5	7.63	0.964	6	[1.099]
39908	6.6	8.44	1.208	2	1.116
27172	7.8	6.54	1.263	2	1.099
78937	21.5	5.37	0.93	12	1.055
96729	9.3	4.93	0.939	8	1.092
94457	9.4	4.97	1.055	9	1.077
16851	67.6	7.96	1.207	162	[1.084]
17651	84.4	7.12	1.4	222	[1.054]
17834	95.2	7.85	1.389	156	[1.060]
55053	4.3	7.92	0.897	2	1.091

33388	2.3	9.35	0.83	2	1.06
37005	2.2	9.27	0.83	2	1.06
40308	2	8.97	0.83	2	1.06
32541	25	6	0.89	11	1.062
50968	3.8	5.22	0.404	2	1.902
71023	3.6	8.39	0.137	3	1.411
65547	4.2	7.74	0.149	2	1.848
68408	2.7	8.68	0.36	3	2.461
66092	6	7.59	0.368	7	[1.830]
59431	5.1	5.7	0.709	3	1.058
79982	4.2	6.05	0.889	3	1.058
39703	34	8.93	0.777	11	1.071
77184	25.5	6.22	0.916	25	[1.046]
20131	28.6	7.07	0.717	5	1.101
236379	1.1	5.92	0.919	2	1.007
33148	16.8	4.34	1.02	4	1.11
12731	13.3	10	1.099	3	1.068
37577	9.9	4.62	0.927	2	1.019
19162	23	6.84	0.901	3	1.029
19317	25.7	6.97	0.934	3	1.021
13441	22.8	4.17	0.976	3	1.066
25228	19.6	8.26	1.029	3	1.03
23920	31.8	5.2	0.935	9	1.043
27685	63.8	5.71	1.058	15	1.047
22829	11.2	5.19	0.93	2	1.179

31644	4.3	6.67	0.976	4	1.055
78460	4.9	5.42	0.93	2	1.022
49691	18.4	5.43	0.891	3	1.124
80187	3.3	6.54	0.988	2	1.088
16526	37.9	8.46	0.935	9	1.129
68519	3.6	5.07	0.98	2	1.077
64504	3.8	5.88	0.952	3	1.194
64694	7.5	4.77	0.965	4	1.048
11675	16.9	5.3	1.01	2	1.055
13544	23.4	3.7	0.97	2	1.041
63744	8.3	7.18	0.832	5	1.042
25441	10.7	5.93	0.94	3	1.086
24761	14.4	5.55	0.987	3	1.006
60847	22.4	7.92	0.996	10	1.107
26321	18.6	5.12	0.935	4	1.075
56831	35	5.87	0.966	12	1.054
51311	29.4	5.05	0.927	11	1.041
86923	30.7	6.94	0.877	24	1.039
52113	11	5.15	0.973	4	1.095
12207	23.2	5.32	0.879	2	1.235
313742	3.1	5.2	0.879	7	1.027
300825	1.1	5.4	1.024	2	1.025
81427	3.2	5.81	0.836	3	1.121
17512	42.9	6.02	1.029	8	[1.049]
27221	6.8	8.8	0.864	4	1.026

295033	5.1	5.84	0.966	12	1.033
13817	27.6	4.8	0.95	3	1.858
7117	26	5.02	1.108	2	1.014
42257	11	6.57	0.967	4	1.116
25201	36.8	8.85	0.995	8	1.077
24537	46.2	8.39	1.006	8	1.05
96734	1.7	5.14	1.098	2	1.105
74624	4.2	7.23	0.866	3	1.013
35900	13	5.56	0.797	3	1.045
38471	12	4.69	0.843	3	1.087
38640	16.5	4.68	0.843	3	1.087
32289	20.6	4.65	0.935	5	1.079
53538	3.8	4.94	1.531	2	1.081
23127	17	9.18	0.928	4	1.083
21962	25.6	9.68	0.982	6	1.057
11705	25	6.15	0.952	6	[1.220]
57015	10.9	5.06	0.741	5	1.096
128056	25.2	5.77	0.951	24	1.023