Supplemental Table

Accession	Score	subcutaneous / light	SD (geo)	# peptides	visceral / light	SD (geo)	# peptides	visceral / subcutaneous	SD (geo)	# peptides	subcutaneous / visceral	SD (geo)	# peptides	Name	signal peptide?
CO3_MOUSE	2356	0.2345	1.5301	40	1.4037	NN	41	3.8962	NN	42	0.2568	NN	42	Complement C3	+
A2M_MOUSE	2021	0.0027	5.4023	22	0.1973	5.208	21	1.0462	NN	33	0.5873	NN	33	Alpha-2-macroglobulin	+
LAMB1_MOUSE	2013	147.8561	NN	35	0.0131	NN	36	0.0465	NN	37	14.0493	3.948	37	Laminin subunit beta 1	+
LAMC1_MOUSE	1988	0.0116	3.3026	36	0.0924	2.401	39	5.3939	1.9632	40	0.1574	NN	40	Laminin subunit gamma-1	+
LAMA2_MOUSE	1409	0.2347	7.3423	25	79.6618	6.516	16	5.8369	5.3994	26	0.079	3.959	26	Laminin subunit alpha-2	+
COEA1_MOUSE	1323	0.1049	5.8287	20	0.0068	3.273	21	0.2932	2.4055	21	3.3802	2.326	22	Collagen alpha-1(XIV) chain	+
FINC_MOUSE	1142	0.0947	NN	24	0.0889	NN	24	3.4279	1.6325	26	0.1466	NN	26	Fibronectin	+
CES3_MOUSE	1132	0	124.26	9	0.2224	26.73	15	7.7748	2.8003	16	0.1269	2.623	15	Carboxylesterase 3	+
HPT_MOUSE	1103	0.6071	NN	30	1.4466	NN	31	2.5003	NN	31	0.3962	NN	31	Haptoglobin	+
CO1A2_MOUSE	925	0.8646	3.0064	14	2.4478	NN	14	1.0944	NN	15	0.37	NN	15	Collagen alpha-2(I) chain	+
TRFE_MOUSE	924	0.1393	NN	17	0.3349	NN	17	1.3117	NN	17	0.3748	1.501	17	Serotransferrin	+
LAMB2_MOUSE	896	2.5529	6.57	13	2.9645	13.3	16	0.5017	NN	17	0.0465	NN	17	Laminin subunit beta-2	+
CFAB_MOUSE	880	1.2984	3.3092	16	0.9025	NN	17	2.2097	1.9195	19	0.4522	1.92	19	Complement factor B	+
CERU_MOUSE	879	0.2978	2.1026	20	0.09174	NN	19	4.2408	1.7977	20	0.2357	1.798	20	Ceruloplasmin	+
SPA3N_MOUSE	865	0.3548	1.941	13	0.0785	NN	14	4.1321	NN	15	0.236	1.397	14	Serine protease inhibitor A3N	+
SPA3K_MOUSE	767	0.0921	NN	13	0.2829	3.77	15	3.1605	1.532	16	0.3164	1.538	16	Serine protease inhibitor A3K	+
ESTN_MOUSE	726	0	117.33	12	410.3341	54.3	12	21.8136	12.037	14	0.0438	6.26	14	Liver carboxylesterase N	+
GELS_MOUSE	725	2.1874	2.7024	12	1.8006	NN	12	1.0615	2.5895	11	1.1021	2.506	12	Gelsolin	+
CO4B_MOUSE	630	4.7321	NN	14	0.1162	NN	14	13.7332	NN	14	0.0729	NN	14	Complement C4-B	+
NID2_MOUSE	618	0.1335	15.389	10	0.0743	17.48	9	3.4691	NN	8	0.288	NN	8	Nidogen-2	+
CATB_MOUSE	596	0.9457	1.5786	11	2.8716	1.5	12	6.2497	NN	13	0.16	NN	13	Cathepsin B	+
ANT3_MOUSE	591	0.9917	5.5633	9	0.0275	14.79	10	0.1622	NN	10	0.4252	1.84	10	Antithrombin-III	+
PGS2_MOUSE	579	0.003	NN	12	0.0878	NN	12	4.737	1.818	12	0.1735	1.827	12	Decorin	+
CO3A1_MOUSE	564	0.0774	NN	13	0.0805	NN	14	2.4722	2.0307	14	0.4055	2.106	13	Collagen alpha-1 (III) chain	+
MRC1_MOUSE	557	0.3938	3.1511	6	0.1811	5.605	7	0.2451	1.9623	7	3.0343	3.176	7	Macrophage mannose receptor 1	+
BCAM_MOUSE	529	2.1488	NN	5	0.3454	9.329	7	0.1956	3.405	6	0.7619	5.058	7	Basal cell adhesion molecule	+
CFAH_MOUSE	496	0.2845	3.5494	4	1.2923	1.629	4	2.8502	2.5445	4	0.1013	3.698	4	Complement factor H	+
MUG2_MOUSE	443	0.8978	10.808	9	0.0303	6.584	8	2.2843	2.6403	9	0.3606	2.441	9	Murinoglobulin-2	+
FIBG_MOUSE	432	23.652	111.11	5	0.0583	9.867	6	0.034	3.3982	5	23.2726	7.762	6	Fibrinogen gamma chain	+
ACE_MOUSE	410	31529	2.2909	8	0.8407	2.853	7	4.8181	2.3835	8	0.2076	2.384	8	Angiotensin-convertine enzyme	+

PTX3_MOUSE	390	67.1404	18.032	4	1.2775	NN	4	18.8319	1.8684	4	0.0531	1.87	4	Pentraxin-related protein PTX3	+
IC1_MOUSE	335	0.3138	1.5059	10	1.9644	NN	10	8.8505	1.6387	11	0.113	1.639	11	Plasma protease C1 inhibitor	+
SPA3M_MOUSE	332	0.0497	NN	7	0.0828	5.602	8	1.9963	2.1728	9	0.501	2.174	9	Serine protease inhibitor A3M	+
SPA3C_MOUSE	331	0.0765	4.283	7	0.6517	2.243	6	4.9315	1.6938	8	0.1036	2.32	8	Serine protease inibitor A3C	+
NGAL_MOUSE	329	0.6059	2.5054	8	2.774	NN	9	3.4368	2.8952	9	0.2909	2.895	9	Neutrophil gelatinase-associated lipocalin	+
CFAD_MOUSE	324	1.3777	1.1318	8	1.3882	1.307	9	1.0363	1.2869	10	0.9651	1.287	10	Complement factor D	+
TSP1_MOUSE	320	0.6749	NN	7	4.6335	1.662	7	11.1698	NN	8	0.0895	NN	8	Thrombospondin-1	+
SODE_MOUSE	273	0.0249	6.6803	7	0.6839	NN	7	32.3922	3.906	7	0.0309	3.908	7	Extracellular superoxide dismutase	+
EST1_MOUSE	273	0		1	1279.303	7248	2	29.1106	7.2684	3	0.0343	7.268	3	Liver carboxylesterase 1	+
VIME_MOUSE	266	19.052	15.548	7	7.874	18.22	7	0.1172	2.9944	7	8.5334	2.995	7	Vimentin	-
SPA3G_MOUSE	264	0.1458	2.0431	5	0.4077	1.614	4	4.1362	1.8379	6	0.2418	1.838	6	Serine protease inhibitor A3G	+
HA11_MOUSE	257	0.064	2.6203	4	0.4928	8.451	4	2.7651	3.6004	4	0.3616	3.602	4	H-2 class I histocompatibility antigen, D-B alpha chain	+
NAR3_MOUSE	257	0.0102	4.8942	5	0.1511	2.59	5	2.2346	2.7769	5	0.4478	2.779	5	Ecto-ADP-ribosyltransferase 3	+
ACTB_MOUSE	252	0.0864	2.2551	3	0.0084	6.181	4	0.0113	12.465	4	2.6692	2.085	4	Actin, cytoplasmic 1	-
ENPP2_MOUSE	248	0.5705	7.9102	4	20.8966	45.79	4	4.1023	3.9229	4	0.2442	3.918	4	Ectonucleotide pyrophosphatase/phosphodiesterase family member 2	+
CATD_MOUSE	241	4.4144	2.6096	5	7.329	1.58	5	15.1463	1.7158	5	0.0661	1.717	5	Cathepsin D	+
DAG1_MOUSE	234	0.0746	12.925	4	0.0326	8.666	3	0.4972	2.7699	4	2.0113	2.77	4	Dystroglycan	+
APOH_MOUSE	234	2.5298	14.217	5	0.0336	8.856	6	0.0429	5.1466	6	14.834	NN	7	Beta-2-glycoprotein 1	+
LDHA_MOUSE	225	6.2293	102.84	5	0.0132	7.267	5	3.738	1.8834	5	94.8116	36.54	6	L-Lactate dehydrogenase A chain	-
1433E_MOUSE	221	0.2642	4.6931	4	0.0752	2.557	3	0.1222	3.3431	4	8.1332	3.365	4	14-3-3 protein epsilon	-
PRDX1_MOUSE	220	0.0557	2.5041	3	0.0009	45.33	2	0.0192	5.6478	3	51.9662	5.666	3	Perioredoxin-1	+
HA18_MOUSE	211	0.096	4.0313	4	17.7184	9.17	4	11.9042	5.2122	4	0.084	5.215	4	H-2 class I histocompatibility antigen, Q8 alpha chain	+
HA17_MOUSE	209	0.0723	15.272	4	12.848	84.12	4	10.7956	5.0913	4	0.0926	5.093	4	H-C class I histocompatibility antigen, Q7 alpha chain	+
PRDX3_MOUSE	205	0.7238	554.81	2	0.2169	18.85	3	12.5873	29.162	4	0.0794	29.18	4	Thioredoxin-dependent peroxide reductase, mitochondrial	-
SPRC_MOUSE	196	0.6078	1.9262	5	0.637	NN	5	1.3967	1.5854	5	0.7159	1.585	5	SPARC	+
CS1B_MOUSE	190	0.1541	5.6117	3	0.025	37.62	3	3.4848	1.8287	3	0.287	1.829	3	Complement C1s-B subcomponent	+
TIMP1_MOUSE	189	0.133	1.0512	2	4.0578	1.085	3	4.7096	11.338	3	0.212	11.32	3	Metalloproteinase inhibitor 1	+
CO4A2_MOUSE	184	0.091	5.164	5	0.0022	14.33	6	0.8723	1.4616	6	1.1459	1.462	6	Collagen alpha-2(IV) chain	+

PGCP_MOUSE	179	0.0341	1.5236	3	0.1248	8.307	3	0.4097	4.2368	3	1.0224	9.465	3	Plasma glutamate carboxypeptidase	+
EGFR_MOUSE	178	0.1552	2.1393	7	0.0291	3.81	7	0.394	NN	7	2.5379	NN	7	Epidermal growth factor receptor	+
CSF1_MOUSE	172	1.8347	1.6213	3	3.7602	1.035	3	2.2038	15.802	3	0.4399	2.846	3	Macrophage colony-stimulating factor 1	+
ECM1_MOUSE	172	1.8373	14.61	5	9.3324	3.051	5	0.1997	9.7599	5	5.0015	9.763	5	Extracellular matrix protein 1	+
ADIPO_MOUSE	171	0.1614		1	0.9812	1.977	2	0.0218	269.86	2	1.1816	4.549	2	Adiponectin	+
SPA3F_MOUSE	165	0.1948	2.4731	3	0.65	1.955	3	3.596	1.6332	4	0.2782	1.622	4	Serine protease inhibitor A3F	+
ACTH_MOUSE	165	1.209	4.5053	3	0.0904	6.231	4	0.0008	22.87	4	3.4122	2.076	4	Actin, gamma-enteric smooth muscle	-
MAOX_MOUSE	161	1.3967		1	0.1691	15.43	2	1.3692	2.8414	2	0.7304	2.841	2	NADP-dependent malic enzyme	-
KV2A7_MOUSE	150	0.0468		1	4.6922		1	39.7792		1	0.0251		1	Ig kappa chain V-II region 26-10	-
CADH5_MOUSE	149	0.0971	2.6599	6	0.1081	NN	6	0.719	NN	6	0.6453	1.948	6	Cadherin-5	+
HA10_MOUSE	149	0.1115	2.5899	2	33.2274	78.69	2	17.1972	22.526	2	0.0581	22.52	2	H-2 class I histocompatability antigen, Q10 alpha chain	+
BPA1_MOUSE	148	0.0004	21.432	12	0.0606	8.403	12	9.2942	4.3497	14	0.0957	6.633	14	Bullous pemphigoid anitgen 1, isoforms 1/2/3/4	-
ANGT_MOUSE	142	2.1243		1	7.5441		1	50.0499		1	0.02		1	Angiotensinogen	+
ITIH2_MOUSE	127	7.2441		1	0.1507		1	0.0169		1	59.1728		1	Inter-alpha-trypsin inhibitor heavy chain H2	+
CATL1_MOUSE	121	0.0077	101.64	2	2.2435	1.381	2	15.3019	15.435	2	0.0654	15.44	2	Cathepsin L1	+
MACF1_MOUSE	120	0.1814	NN	12	0.0355	7.83	13	1.0881	3.3499	13	0.7352	2.833	13	Microtubule-actin cross-linking factor 1	-
CO2_MOUSE	117	0.1838	3.0852	5	0.084	5.556	5	8.581	3.0821	5	0.1166	3.081	5	Complement C2	+
AMRP_MOUSE	114	0.2356	5.7393	4	0.0813	15.69	5	0.7802	12.774	5	0.844	3.519	4	Alpha-2-macroglobulin receptor-associated protein	+
PXDC2_MOUSE	111	0.1589	13.219	2	0.2218	3.478	2	3.1057	12.37	2	0.3197	12.9	2	Plexin domain-containing protein 2	+
SAP_MOUSE	107	3.2836	1.9121	3	2.3312	1.339	3	34.6713	3.8008	4	0.0292	3.785	4	Sulfated glycoprotein 1	+
CD14_MOUSE	107	0.1239		1	5.9176		1	54.9729		1	0.0182		1	Monocyte differentiation antigen CD14	+
CAH3_MOUSE	105	0.0843		1	0.3909	5.124	2	0.0235	9.9987	2	42.5432	9.992	2	Carbonic anhydrase 3	-
CYB5_MOUSE	104	0.082	5.8284	2	0.1961	8.997	2	0.145	50.898	2	7.6463	61.84	2	Cytochrome b5	-
PLXC1_MOUSE	99	1.1268	6.5146	5	0.3903	3.465	5	0.2068	2.2722	5	4.8353	2.272	5	Plexin-C1	+
A2MP_MOUSE	99	0.0201	23.581	3	1.8505	10.53	3	0.3333	46.713	3	0.4291	10.09	3	Alpha-2-macroglobulin-P	+
POSTN_MOUSE	99	3.1756	1.7059	2	0.1401	16.22	2	0.018	18.05	2	4.2549	2.874	2	Periostin	+
DYH3_MOUSE	94	0.076	7.1555	9	1.1202	4.631	10	1.1125	4.1426	10	0.5698	4.153	10	Dynein heavy chain 3, axonemal	-
PLOD1_MOUSE	91	1.0155		1	0.3721	817.9	2	2.3807	1.4291	2	0.4198	1.427	2	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1	+

CO1A1_MOUSE	88	0.3773		1	4.3784		1	13.8932		1	0.0719		1	Collagen alpha-1(I) chain	+
ACTBL_MOUSE	87	218.2882	100.78	2	0.0078	6.04	3	0.003	10.812	3	6.3158	2.942	3	Beta-actin-like protein 2	-
HERC2_MOUSE	86	0.6902	7.5135	7	4.1693	4.218	7	0.7363	3.9473	7	1.3565	3.946	7	Probable E3 ubiquitin-protein ligase HERC2	-
MSH3_MOUSE	85	0.2613	21.788	6	1.5996	7.759	6	2.8729	17.507	6	0.2656	3.625	6	DNA mismatch repair protein Msh3	-
ASPM_MOUSE	82	0.6162	3.8044	8	0.123	4.485	9	2.0761	2.4856	9	0.4821	2.484	9	Abnormal spindle-like microcephaly-associated protein homolog	-
ASAP2_MOUSE	82	0.0002	29.871	4	0.0826	2.924	3	0.7028	1.4637	4	1.4229	1.464	4	Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 2	-
SPTB1_MOUSE	81	0.0737	24.094	9	4.5364	11.6	9	0.3966	6.3907	9	1.7392	7.132	9	Spectrin beta chain, erythrocyte	-
GVIN1_MOUSE	81	3.1184	142.69	6	390.3908	390.4	6	11.8529	75.807	6	0.0694	10.93	6	Interferon-induced very large GTPase 1	-
NAL9C_MOUSE	80	0.94	43.103	3	0.0456	7.369	4	0.0441	16.381	4	22.663	16.35	4		-
DPEP1_MOUSE	79	2.2056	42.514	4	0.0628	9.143	4	0.0357	4.865	4	28.0178	4.864	4	Dipeptidase 1	+
CHD9_MOUSE	78	16.6318	NN	3	3.4233	26.36	3	4.4816	4.5902	5	0.2232	4.589	5	Chromodomain-helicase-DNA-binding protein 9	-
CAD13_MOUSE	78	2.6301	2.9062	2	0.0032	65.78	2	6.5059	3.5891	3	0.01537	3.59	3	Cadherin-13	+
GAS7_MOUSE	77	5.9727	1.1733	2	3.3274	1.129	2	0.2346	NN	3	4.2677	NN	3	Growth arrest specific 7	-
CYC_MOUSE	77	0.0196		1	0.0544		1	0.1362		1	7.3408		1	Cytochrome c, somatic	-
S22A3_MOUSE	71	20.9636	860.78	2	0.0205	2.447	2	0.0241	91.564	2	41.5784	91.69	2	Solute carrier family 22 member 3	-
PAI1_MOUSE	70	0.7036	2.4482	4	4.94	62.59	4	9.7866	7.5482	3	0.1021	7.548	3	Plasminogen activator inhibitor 1	+
ANR53_MOUSE	70	0.1377	18.094	4	0.3512	9.616	3	10.0498	NN	4	0.082	NN	4	Ankyrin repeat domain-containing protein 53	-
MMRN2_MOUSE	70	12.5096	6.0477	2	0.0051		1	4.6797	1224.7	2	0.2137	1225	2	Multimerin-2	+
TTF2_MOUSE	70	0.0769	5.0286	4	0.022	3.893	4	0.1806	2.7587	4	5.5294	2.757	4	Transcription termination factor 2	-
SACS_MOUSE	69	0.0605	5.4009	5	0.0454	5.286	5	0.6014	4.3245	6	1.5689	3.762	6		-
MGAP_MOUSE	69	0.3918	4.6819	8	0.7102	11.54	7	8.7736	2.4715	8	0.1085	NN	8	MAX gene-associated protein	-
ILRL1_MOUSE	68	0.1565		1	3.5857		1	0.4894	35.437	2	2.0441	35.42	2	Interleukin-1 receptor-like 1	+
EVPL_MOUSE	68	0.1793	1.946	5	0.3531	NN	7	3.2461	9.8823	7	0.3189	8.984	7	Envoplakin	-
NAR5_MOUSE	68	0.0015		1	5.3171	20.82	2	0.0518	52.954	2	19.2719	52.96	2	Ecto-ADP-ribosyltransferase 5	+
UBR4_MOUSE	68	0.6076	8.985	4	0.0239	1.433	3	0.3207	47.595	4	0.4693	3.642	4	E3 ubiquitin-protein ligase UBR4	-
GANAB_MOUSE	68	0.1478		1	0.0332		1	0.217		1	4.608		1	Neutral alpha-glucosidase AB	+
NCOR1_MOUSE	68	1.4281	7.223	6	0	NN	7	0.6675	27.994	7	1.4979	28.01	7	Nuclear receptor corepressor 1	-
CC136_MOUSE	67	0.2326	7.6904	3	45.7854	90.66	7	0.169	NN	8	5.1836	NN	8	coiled-coil domain-containing	-

													protein 136	
TRPM7_MOUSE	67	9.5688	NN	6	7.0928	7.806	6	0.0408	3.7778	6	21.7104	2.597	6	Transient receptor potential cation channel subfamily M member 7	-
STRUMP_MOUSE	67	2.1322	1.7139	2	111.9759	5.426	2	16.8345	37.023	2	0.0594	37.05	2	Strumpelin	-
PARD3_MOUSE	67	0.2391	6.643	8	0.0839	4.915	8	0.1408	11.43	8	0.6806	2.786	8	Partitioning defective homolog	-
CD158_MOUSE	66	0.0085	46.142	6	0.166	33.22	6	3.106	7.501	6	0.2552	5.38	6	Coiled-coil domain-containing protein 158	-
ACLY_MOUSE	66	4.3181	12.444	2	0.0022	41.17	4	0.021	45.961	3	8.4078	13.4	4	ATP-citrate synthase	-
ACTN4_MOUSE	66	0.2177	4.877	3	1.0321	9.228	3	1.9221	47.294	3	0.0438	10.03	3	Alpha-actinin-4	-
LRP2_MOUSE	66	0.3556	11.453	12	1.7702	5.511	12	1.3135	2.6414	12	0.4406	3.083	11	Low-density lipoprotein recpetor-related protein 2	+
RPAP1_MOUSE	66	0.3742	3.3283	5	0.7171	23.5	5	3.4288	8.2411	5	0.2915	8.241	5	RNA polymerase II-associated protein 1	-
CAH2_MOUSE	65	0.0839	1.6259	2	0.2907	1.828	2	2.2056	1.3659	2	0.4533	1.367	2	Carbonic anhydrase 2	-
NAL9B_MOUSE	65	0.0155	66.915	3	0.2934	9.635	3	0.7537	7.8936	3	1.3252	7.878	3		-
VINC_MOUSE	65	0.1572	2.1518	3	22.0341	14.25	3	4.645	2.7834	3	0.2153	2.785	3	Vinculin	-
EIF3A_MOUSE	65	0.0715	2.3693	6	0.4094	26.22	7	4.3212	5.0885	7	0.2177	11.56	7	Eukaryotic translation intiation factor 3 subunit A	-
MA7D2_MOUSE	65	0.3336	141.99	5	0.1566	33.83	5	6.7425	NN	5	0.1488	NN	5	MAP7 domain-containing protein 2	-
CO6A1_MOUSE	64	0.064	12.294	2	0.0746	8.676	2	1.9258	1.5233	2	0.5188	1.523	2	Collagen alpha-1(VI) chain	+
BRCA2_MOUSE	64	0	88.789	8	0.3277	5.421	10	3.8641	5.9921	11	0.2566	6.786	10	Breast cancer type 2 susceptibility protein homolog	-
TBCK_MOUSE	64	0.451	5.0835	3	0.656	19.89	3	16.6585	9.2643	4	0.0354	35.91	4		-
MDHM_MOUSE	64	0.4328		1	1.4402		1	12.6742		1	0.079		1	Malate dehydrogenase, mitochondrial	-
LAMP1_MOUSE	64	0.5988		1	1.9068		1	2.5343		1	0.3946		1	Lysosome-associated membrane glycoprotein 1	+
MYO5A_MOUSE	64	0.0991	10.193	4	0.0168	20.25	4	0.0777	4.342	4	2.0474	1.786	4	Myosin-Va	-
ZSWM5_MOUSE	63	0.504	nn	6	0.1894	NN	6	0.1203	1.6787	5	2.7207	2.914	6	Zinc finger SWIM domain-containing protein 5	-
UBP36_MOUSE	62	0.1443	31.736	6	0.0539	8.691	6	0.6201	4.4281	6	1.6127	4.429	6		-
MGT5A_MOUSE	62	0.1529	15.664	2	0.6463	79.39	2	3.4498	1.0897	2	0.29	1.09	2		-
WDR64_MOUSE	62	0.0479	3.0752	3	0.6202	5.376	3	7.6007	3.1374	3	0.1255	3.244	3	WD repeat-containing domain protein 64	-
KIF27_MOUSE	62	0.1525	2.9077	5	1.9513	22.49	6	0.1688	6.6567	6	5.9215	6.655	6		-
SEM7A_MOUSE	62	0.5533		1	1.6165		1	3.2088		1	0.312		1	Semaphorin-7A	+

IFIT2_MOUSE	62	457.5546	7557.6	2	0.0001	110.9	2	0.0997	33.34	2	1.026	33.36	2	Interferon-induced protein with tetratricopeptide repeats 2	-
MINT_MOUSE	62	0.492	9.0921	13	0.0114	NN	13	0.6154	2.3897	13	5.3222	2.854	13	Mxs2-interacting protein	-
ATRX_MOUSE	61	0.9456	5.2326	7	0.1083	4.805	6	0.0796	5.4935	6	12.5475	5.486	6	Transcriptional regulator ATRX	-
MA7D1_MOUSE	61	4.1372	3.0154	3	0.0012	31.26	4	0.2126	3.9806	4	4.7022	3.98	4	MAP7 domain-containing protein 1	-
F91A1	61	15.9334	786.65	2	4.249	18.66	2	0.044	27.682	3	22.7443	27.71	3		-
GLU2B_MOUSE	61	2.8153	1.1815	2	1.5869	3.171	2	0.2904	5.1074	2	3.4447	5.119	2	Glucosidase 2 subunit beta	+