Table S-1a

A list of 8,183 unique protein-specific peptides identified in tumor, sorted by protein name.

**MS/MS spectra were searched against the UniProt Human proteomic database; release 09/07.*
*** The highest Xcorr observed for identified peptide. *** Total count of peptide identifications.*

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
10 kDa heat shock protein, mitochondrial - Homo sapiens (Human)					
P61604	3	6.2295	K.SQGKVLQATVVAVGSGSKGK.G	6	CH10_HUMAN
P61604	1	2.2593	K.FLPLFDR.V	1	CH10_HUMAN
P61604	2	3.8147	R.VLVERSAAETVTK.G	3	CH10_HUMAN
P61604	1	3.074	R.SAAETVTK.G	34	CH10_HUMAN
P61604	2	3.9025	R.KFLPLFDRVLVER.S	3	CH10_HUMAN
P61604	2	3.2619	R.DGDILGKYVD.-	2	CH10_HUMAN
P61604	2	2.9068	R.DGDILGK.Y	5	CH10_HUMAN
P61604	3	3.9242	K.VVLDDKDYFLFRDGDILGK.Y	1	CH10_HUMAN
P61604	2	3.6638	K.VVLDDKDYFLFR.D	2	CH10_HUMAN
P61604	3	4.2717	K.VLQATVVAVGSGSKGKGGEIQPVSVK.V	1	CH10_HUMAN
P61604	2	5.0989	K.VLQATVVAVGSGSKGK.G	3	CH10_HUMAN
P61604	2	5.8123	K.VLQATVVAVGSGSK.G	15	CH10_HUMAN
P61604	2	4.3099	K.VGDKVLLPEYGGTK.V	4	CH10_HUMAN
P61604	1	2.7647	K.GGIMLPEK.S	5	CH10_HUMAN
P61604	2	3.2578	K.FLPLFDRVLVER.S	10	CH10_HUMAN
P61604	2	3.1446	K.GGEIQPVSVK.V	4	CH10_HUMAN
P61604	3	4.4235	K.GGEIQPVSVKVGDKVLLPEYGGTK.V	2	CH10_HUMAN
P61604	1	2.1886	K.GGIM#LPEK.S	1	CH10_HUMAN
P61604	1	2.3902	K.VLLPEYGGTK.V	3	CH10_HUMAN
P61604	3	5.409	K.GGIM#LPEKSQGKVLQATVVAVGSGSK.G	3	CH10_HUMAN
P61604	2	3.5411	K.GGIMLPEKSQGK.V	4	CH10_HUMAN
P61604	3	6.201	K.GGIMLPEKSQGKVLQATVVAVGSGSK.G	2	CH10_HUMAN
P61604	2	4.8678	K.GKGGEIQPVSVK.V	8	CH10_HUMAN
P61604	3	3.7199	K.GKGGEIQPVSVKVGDK.V	1	CH10_HUMAN
P61604	2	6.4404	K.SQGKVLQATVVAVGSGSK.G	9	CH10_HUMAN
P61604	2	3.0761	K.GGIM#LPEKSQGK.V	2	CH10_HUMAN
130 kDa phosphatidylinositol 4,5-biphosphate-dependent ARF1 GTPase-activating protein - Homo					
Q9ULH1	2	3.8327	K.SHPLDLSPNVQSR.D	2	DDEF1_HUMAN
Q9ULH1	2	3.8215	K.GVFPVSFVHILSD.-	1	DDEF1_HUMAN
Q9ULH1	3	4.8537	K.TTNKFEGLSQQSSTSSAK.T	2	DDEF1_HUMAN
Q9ULH1	2	3.5485	K.FEGLSQQSSTSSAK.T	1	DDEF1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9ULH1	2	2.9629	K.GAVPWGNDGGPSSSSK.T	2	DDEF1_HUMAN
Q9ULH1	3	5.5033	R.NAGKGPTGPPSTLPLSTQTSSGSSTLSK.K	1	DDEF1_HUMAN
Q9ULH1	2	3.29	K.GPTGPPSTLPLSTQTSSGSSTLSK.K	1	DDEF1_HUMAN
14 kDa phosphohistidine phosphatase - Homo sapiens (Human)					
Q9NRX4	2	4.1411	K.AKYPDYEVTWANDGY.-	7	PHP14_HUMAN
Q9NRX4	2	6.1032	K.IHVYGYSM#AYGPAQHAISTEK.I	9	PHP14_HUMAN
Q9NRX4	3	4.1347	K.IHVYGYSMAYGPAQHAISTEK.I	3	PHP14_HUMAN
14-3-3 protein beta/alpha - Homo sapiens (Human)					
P31946	4	5.039	R.YLSEVASGDNKQTTVSNSQQAYQEAFEISKK.	1	1433B_HUMAN
14-3-3 protein epsilon - Homo sapiens (Human)					
P62258	3	4.8616	R.YLAEFATGNDRKEAAENSLVAYK.A	2	1433E_HUMAN
P62258	3	4.0153	K.LAEQAERYDEMVESMKK.V	1	1433E_HUMAN
P62258	3	4.9962	K.LAEQAERYDEMVESMK.K	2	1433E_HUMAN
P62258	2	2.8044	K.AAFDDAIAELDTLSEESYK.D	1	1433E_HUMAN
P62258	2	2.999	K.AASDIAMTELPPTHPIR.L	1	1433E_HUMAN
14-3-3 protein gamma - Homo sapiens (Human)					
P61981	2	5.0552	K.NVTELNEPLSNEER.N	3	1433G_HUMAN
14-3-3 protein sigma - Homo sapiens (Human)					
P31947	2	4.0983	K.SNEEGSEEKGPEVR.E	2	1433S_HUMAN
P31947	3	4.5026	K.LAEQAERYEDMAAFMK.G	1	1433S_HUMAN
14-3-3 protein zeta/delta - Homo sapiens (Human)					
P63104	2	4.7685	K.LAEQAERYDDMAACMK.S	2	1433Z_HUMAN
P63104	2	3.9548	R.YLAEVAAGDDKK.G	2	1433Z_HUMAN
182 kDa tankyrase 1-binding protein - Homo sapiens (Human)					
Q9C0C2	2	5.0818	R.KMNMLAGPQPYGGSK.R	2	TB182_HUMAN
Q9C0C2	2	4.4122	R.ESAVGQMGWSGGLSLR.D	2	TB182_HUMAN
Q9C0C2	2	3.2469	R.ESGVGQTDWSGVEAGEFLK.S	2	TB182_HUMAN
Q9C0C2	2	4.7799	R.FAATTVEEILAK.M	5	TB182_HUMAN
Q9C0C2	2	3.446	R.FSEGVLQSPSQDQEK.L	2	TB182_HUMAN
Q9C0C2	2	3.3291	R.GPLAELPSAR.K	1	TB182_HUMAN
Q9C0C2	2	4.2181	R.GSGGLFSPSTAHVPDGALGQR.D	1	TB182_HUMAN
Q9C0C2	2	2.9461	R.KM#NM#LAGPQPYGGSK.R	1	TB182_HUMAN
Q9C0C2	2	3.4882	R.SLPSDLAFNGDLAK.A	2	TB182_HUMAN
Q9C0C2	2	3.6415	R.SYQFGIIGNDR.V	5	TB182_HUMAN
Q9C0C2	2	4.8371	R.WLDDLLASPPPSGGGAR.R	3	TB182_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9C0C2	2	3.8487	R.YSSQDADEQDWEFQKR.D	2	TB182_HUMAN
Q9C0C2	2	3.0428	R.SQEADVQDWEFR.K	2	TB182_HUMAN
Q9C0C2	2	2.8187	R.ESAVGQM#GWSGGLSLR.D	2	TB182_HUMAN
Q9C0C2	2	3.4038	R.GWVGEFSLSVGPQR.E	3	TB182_HUMAN
Q9C0C2	2	3.7875	K.VSAPGVLTAQDR.V	1	TB182_HUMAN
Q9C0C2	3	3.8609	R.ERGVGQADWTPDLGLR.N	1	TB182_HUMAN
Q9C0C2	2	3.9908	K.KIPSVEDSLGEGSR.D	2	TB182_HUMAN
Q9C0C2	2	3.5932	R.VSGAGFSPSSK.M	3	TB182_HUMAN
Q9C0C2	2	2.9072	K.ELGVGQMDWGNNLGLR.D	2	TB182_HUMAN
Q9C0C2	2	3.8792	K.YGQGAGEGSTR.E	7	TB182_HUMAN
Q9C0C2	2	3.3634	R.AAEPQEQEFGK.S	2	TB182_HUMAN
Q9C0C2	2	3.0737	R.DAELQDQEFGK.R	1	TB182_HUMAN
Q9C0C2	2	4.5345	R.DLEVTCDPDSGGSQGLR.G	4	TB182_HUMAN
Q9C0C2	2	2.8165	R.DSLGTYSSR.D	1	TB182_HUMAN
Q9C0C2	2	3.0077	R.DSQGTYSSR.D	6	TB182_HUMAN
Q9C0C2	2	3.3511	R.DVSLGTYGSR.A	4	TB182_HUMAN
Q9C0C2	2	3.3096	R.EMEEELVPTGSEPGDTR.A	1	TB182_HUMAN
Q9C0C2	2	3.9751	R.EAAFSPGQQDWSR.D	2	TB182_HUMAN
Q9C0C2	2	5.8023	K.SKDLAEVGEGGGHSQAR.E	2	TB182_HUMAN
Q9C0C2	2	3.0803	R.EM#EEELVPTGSEPGDTR.A	1	TB182_HUMAN
Q9C0C2	2	3.6967	R.DAELQDQEFGKR.D	4	TB182_HUMAN
1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase beta 1 - Homo sapiens (Human)					
Q9NQ66	2	3.0253	K.YNEIQNDYLR.R	2	PLCB1_HUMAN
26S protease regulatory subunit 7 - Homo sapiens (Human)					
P35998	2	3.5337	K.FVVDLSDQVAPTDIEEGMR.V	1	PRS7_HUMAN
26S proteasome non-ATPase regulatory subunit 2 - Homo sapiens (Human)					
Q13200	3	4.8885	K.APVQPQQSPAAAPGGTDEKPSGK.E	2	PSMD2_HUMAN
26S proteasome non-ATPase regulatory subunit 4 - Homo sapiens (Human)					
P55036	3	4.6507	R.AAAASAAEAGIATTGTEDSDDALLK.M	4	PSMD4_HUMAN
26S proteasome non-ATPase regulatory subunit 9 - Homo sapiens (Human)					
O00233	3	5.5857	R.HNIICLQNDHK.A	1	PSMD9_HUMAN
O00233	2	3.6075	R.SDVDLYQVR.T	5	PSMD9_HUMAN
O00233	3	4.0588	R.RKEEIEAQIK.A	5	PSMD9_HUMAN
O00233	2	4.0098	K.ANYDVLESQK.G	2	PSMD9_HUMAN
O00233	2	4.0943	R.QSGGSSQAGAVTVSDVQELMR.R	3	PSMD9_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O00233	3	4.3134	R.QSGGSSQAGAVTVSDVQELM#R.R	1	PSMD9_HUMAN
O00233	2	4.2292	R.KLGQSESQGPPR.A	4	PSMD9_HUMAN
O00233	2	3.1231	K.QVEEALHQLHAR.D	2	PSMD9_HUMAN
O00233	2	4.3835	K.LGQSESQGPPR.A	4	PSMD9_HUMAN
O00233	2	3.4773	K.GLLGCNIIPLQR.-	4	PSMD9_HUMAN
O00233	2	4.1137	K.GIGM#NEPLVDCEGYPR.S	3	PSMD9_HUMAN
O00233	2	4.015	R.DMAEAHKEAM#SR.K	2	PSMD9_HUMAN
O00233	2	4.3239	K.GIGMNEPLVDCEGYPR.S	4	PSMD9_HUMAN
28 kDa heat- and acid-stable phosphoprotein - Homo sapiens (Human)					
Q13442	1	3.0665	R.M#QSLSLNK.-	8	HAP28_HUMAN
Q13442	3	4.6164	K.RKGVEGLIDIENPNR.V	2	HAP28_HUMAN
Q13442	2	2.9848	R.QYTSPEEIDAQLQAEKQK.A	1	HAP28_HUMAN
Q13442	1	2.9046	K.VTQLDLDGPK.E	3	HAP28_HUMAN
Q13442	3	4.3547	R.EEEEQKEGGDGAAGDPKK.E	4	HAP28_HUMAN
Q13442	3	3.8382	R.EREEIEKQK.A	1	HAP28_HUMAN
Q13442	2	5.3286	R.KGVEGLIDIENPNR.V	9	HAP28_HUMAN
Q13442	3	5.1715	R.QYTSPEEIDAQLQAEK.Q	3	HAP28_HUMAN
Q13442	2	3.1509	K.MHLAGKTEQAK.A	2	HAP28_HUMAN
Q13442	2	3.9878	R.KAKDDATLSGK.R	10	HAP28_HUMAN
Q13442	3	5.3078	K.KVTQLDLDGPKELSR.R	5	HAP28_HUMAN
Q13442	2	3.7963	K.KVTQLDLDGPK.E	6	HAP28_HUMAN
Q13442	3	5.2038	K.KSLDSDESEDEEDDYQQK.R	2	HAP28_HUMAN
Q13442	2	3.2005	K.GVEGLIDIENPNRVAQTTK.K	1	HAP28_HUMAN
Q13442	2	4.829	K.GVEGLIDIENPNR.V	7	HAP28_HUMAN
Q13442	1	2.2097	K.DDATLSGK.R	2	HAP28_HUMAN
Q13442	2	3.9639	K.AKDDATLSGKR.M	3	HAP28_HUMAN
Q13442	2	3.2411	K.AKDDATLSGK.R	9	HAP28_HUMAN
Q13442	2	4.8487	K.SLDSDESEDEEDDYQQK.R	6	HAP28_HUMAN
Q13442	2	3.9825	K.M#HLAGKTEQAK.A	5	HAP28_HUMAN
28S ribosomal protein S31, mitochondrial precursor - Homo sapiens (Human)					
Q92665	2	3.1742	K.ISFSNIISDM#K.V	1	RT31_HUMAN
Q92665	2	3.3302	K.SELLSQLQQHEEESR.A	1	RT31_HUMAN
Q92665	2	4.6086	K.ISFSNIISDMK.V	3	RT31_HUMAN
Q92665	3	4.2414	K.RIEPLSPELVAAASAVADSLPFDKQTTK.S	1	RT31_HUMAN
39S ribosomal protein 54, mitochondrial precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6P161	2	2.8277	K.TLEELDPESR.E	1	RM54_HUMAN
39S ribosomal protein L12, mitochondrial precursor - Homo sapiens (Human)					
P52815	2	3.2881	R.LTEAKPVDKVK.L	4	RM12_HUMAN
P52815	2	3.2401	K.EIKNYIQGINLVQAK.K	2	RM12_HUMAN
P52815	2	4.1911	K.IKAALEAVGGTVVLE.-	1	RM12_HUMAN
P52815	2	4.4916	K.KLVESLPQEIK.A	3	RM12_HUMAN
P52815	2	3.4344	K.LVESLPQEIK.A	2	RM12_HUMAN
P52815	2	4.4337	K.NYIQGINLVQAK.K	4	RM12_HUMAN
P52815	2	3.1783	R.LTEAKPVDK.V	3	RM12_HUMAN
39S ribosomal protein L14, mitochondrial precursor - Homo sapiens (Human)					
Q6P1L8	2	3.1877	K.VGDQILLAIK.G	1	RM14_HUMAN
39S ribosomal protein L34, mitochondrial precursor - Homo sapiens (Human)					
Q9BQ48	2	2.8951	R.GNEYQPSNIK.R	1	RM34_HUMAN
39S ribosomal protein L40, mitochondrial precursor - Homo sapiens (Human)					
Q9NQ50	2	4.0693	K.ATQELIPIEDFITPLK.F	2	RM40_HUMAN
39S ribosomal protein L52, mitochondrial precursor - Homo sapiens (Human)					
Q86TS9	3	4.8027	R.LQQGLAANPSGYGPLTELPDWSYADGRPAP	1	RM52_HUMAN
Q86TS9	3	4.9076	R.LQQGLAANPSGYGPLTELPDWSYADGRPAP	1	RM52_HUMAN
3-hydroxyisobutyrate dehydrogenase, mitochondrial precursor - Homo sapiens (Human)					
P31937	2	3.3806	K.DLGLAQDSATSTK.S	1	3HIDH_HUMAN
P31937	2	3.9306	K.TPVGFIGLGNMGNPM#AK.N	1	3HIDH_HUMAN
P31937	2	4.6762	K.TPVGFIGLGNMGNPMAK.N	1	3HIDH_HUMAN
3-ketoacyl-CoA thiolase, mitochondrial - Homo sapiens (Human)					
P42765	2	3.1017	K.AANDAGYFNDEMAPIEVK.T	2	THIM_HUMAN
4.1G protein - Homo sapiens (Human)					
Q9H1C6	2	3.1119	K.YLMPTIWTVEEK.A	2	Q9H1C6_HUMA
Q9H1C6	2	3.9967	K.SSHETLNIVEEK.K	3	Q9H1C6_HUMA
40S ribosomal protein S11 - Homo sapiens (Human)					
P62280	2	4.2374	R.AYQKQPTIFQNK.K	2	RS11_HUMAN
P62280	3	4.2039	K.RVLLGETGKEK.L	2	RS11_HUMAN
P62280	2	3.1372	K.EAIEGTYIDKK.C	2	RS11_HUMAN
P62280	1	2.439	K.NIGLGFK.T	3	RS11_HUMAN
40S ribosomal protein S13 - Homo sapiens (Human)					
P62277	2	3.5354	K.GLSQSALPYR.R	2	RS13_HUMAN
P62277	2	2.9752	K.GLTPSQIGVILR.D	3	RS13_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P62277	2	3.3002	K.KGLTPSQIGVILR.D	2	RS13_HUMAN
P62277	2	4.3956	K.LTSDDVKEQIYK.L	3	RS13_HUMAN
40S ribosomal protein S14 - Homo sapiens (Human)					
P62263	2	3.1361	K.TPGPGAQSALR.A	2	RS14_HUMAN
P62263	2	3.7213	R.IEDVTPIPSDSTR.R	4	RS14_HUMAN
P62263	2	2.7124	R.IEDVTPIPSDSTRR.K	1	RS14_HUMAN
P62263	3	4.4397	K.ADRDESSPYAAMLAAQDVAQR.C	3	RS14_HUMAN
P62263	2	3.7964	K.IGRIEDVTPIPSDSTR.R	4	RS14_HUMAN
P62263	3	4.5247	R.TKTPGPGAQSALR.A	3	RS14_HUMAN
P62263	2	4.5208	K.IGRIEDVTPIPSDSTRR.K	5	RS14_HUMAN
40S ribosomal protein S15 - Homo sapiens (Human)					
P62841	3	4.7012	K.KEAPPMEKPEVVK.T	3	RS15_HUMAN
40S ribosomal protein S17 - Homo sapiens (Human)					
P08708	2	3.5969	K.LLDFGSLSNLQVTQPTVGM#NFK.T	3	RS17_HUMAN
P08708	3	4.7104	K.LLDFGSLSNLQVTQPTVGMNFK.T	1	RS17_HUMAN
P08708	2	2.7975	K.LQEEERER.R	3	RS17_HUMAN
P08708	2	3.5782	R.LGNDFHTNKR.V	2	RS17_HUMAN
P08708	3	7.3801	R.RDNYVPEVSALDQEIIEVDPDTKEMLK.L	1	RS17_HUMAN
40S ribosomal protein S18 - Homo sapiens (Human)					
P62269	2	3.816	K.RAGELTEDEVER.V	2	RS18_HUMAN
P62269	2	2.8714	R.VITIMQNPR.Q	2	RS18_HUMAN
P62269	2	3.0852	R.KADIDLTKR.A	2	RS18_HUMAN
P62269	2	2.962	R.AGELTEDEVER.V	1	RS18_HUMAN
P62269	3	4.5471	R.AGELTEDEVERVITIMQNPR.Q	1	RS18_HUMAN
40S ribosomal protein S19 - Homo sapiens (Human)					
P39019	1	3.2499	R.GGAGVGSMTK.I	3	RS19_HUMAN
P39019	3	5.736	K.HKELAPYDENWFYTR.A	4	RS19_HUMAN
P39019	2	3.3946	R.VLQALEGLK.M	2	RS19_HUMAN
P39019	2	3.182	R.RVLQALEGLK.M	6	RS19_HUMAN
P39019	2	3.3416	R.IAGQVAAANKKH.-	2	RS19_HUMAN
P39019	2	3.369	R.IAGQVAAANKK.H	3	RS19_HUMAN
P39019	1	3.1372	R.IAGQVAAANK.K	11	RS19_HUMAN
P39019	3	5.1679	R.DLDRIAGQVAAANKK.H	3	RS19_HUMAN
P39019	2	4.869	R.DLDRIAGQVAAANK.K	6	RS19_HUMAN
P39019	2	2.9597	R.ALAAFLKK.S	2	RS19_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P39019	2	2.8849	R.ALAAFLK.K	3	RS19_HUMAN
P39019	3	4.0363	K.LKVPEWVDTVK.L	5	RS19_HUMAN
P39019	1	3.5671	K.DVNQQEFVR.A	8	RS19_HUMAN
P39019	2	3.1808	K.VPEWVDTVK.L	2	RS19_HUMAN
40S ribosomal protein S19-binding protein 1 - Homo sapiens (Human)					
Q86WX3	2	3.4688	R.STVAESVSQQILR.Q	4	S19BP_HUMAN
40S ribosomal protein S2 - Homo sapiens (Human)					
P15880	3	4.2298	K.SPYQEFTDHLVK.T	2	RS2_HUMAN
P15880	1	2.1805	R.TQAPAVATT.-	1	RS2_HUMAN
40S ribosomal protein S20 - Homo sapiens (Human)					
P60866	2	3.7472	R.LIDLHSPSEIVK.Q	4	RS20_HUMAN
P60866	1	2.4006	K.VCADLIR.G	2	RS20_HUMAN
P60866	2	3.3666	K.DTGKTPVEPEVAIHR.I	2	RS20_HUMAN
P60866	2	3.4029	K.RLIDLHSPSEIVK.Q	2	RS20_HUMAN
P60866	3	4.3526	K.TPVEPEVAIHR.I	3	RS20_HUMAN
40S ribosomal protein S21 - Homo sapiens (Human)					
P63220	2	2.9909	R.M#GESDDSILR.L	2	RS21_HUMAN
P63220	2	3.6447	R.MGESDDSILR.L	5	RS21_HUMAN
P63220	2	3.6621	R.RMGESDDSILR.L	2	RS21_HUMAN
40S ribosomal protein S23 - Homo sapiens (Human)					
P62266	2	2.9793	K.GHAVGDIPGVR.F	2	RS23_HUMAN
P62266	2	4.6538	K.VANVSLLALYK.G	5	RS23_HUMAN
P62266	2	3.4091	R.KGHAVGDIPGVR.F	2	RS23_HUMAN
40S ribosomal protein S24 - Homo sapiens (Human)					
P62847	3	4.3922	R.KQMVIDVLHPGK.A	1	RS24_HUMAN
P62847	2	3.3319	K.TTGFGMIYDSLDYAK.K	2	RS24_HUMAN
40S ribosomal protein S25 - Homo sapiens (Human)					
P62851	2	4.3287	R.NTKGGDAPAAGEDA.-	9	RS25_HUMAN
P62851	3	4.5021	R.DKLNNLVLFDKATYDK.L	1	RS25_HUMAN
P62851	2	3.9405	R.DKLNNLVLFDK.A	6	RS25_HUMAN
P62851	2	2.9483	R.AALQELLSK.G	5	RS25_HUMAN
P62851	2	4.6688	K.LNNLVLFDKATYDK.L	3	RS25_HUMAN
P62851	1	2.4474	K.GGDAPAAGEDA.-	2	RS25_HUMAN
P62851	2	3.9868	K.LNNLVLFDK.A	8	RS25_HUMAN
40S ribosomal protein S28 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P62857	2	3.795	R.TGSQGQCTQVR.V	30	RS28_HUMAN
P62857	3	3.9031	K.GPVREGDVLTLLESER.E	2	RS28_HUMAN
P62857	2	3.1235	R.VEFM#DDTSR.S	2	RS28_HUMAN
P62857	1	3.0766	R.VEFMDDTSR.S	6	RS28_HUMAN
P62857	2	3.7871	R.EGDVLTLLESEREAR.R	4	RS28_HUMAN
P62857	3	3.811	K.GPVREGDVLTLLESEREAR.R	1	RS28_HUMAN
P62857	3	4.343	R.NVKGPVREGDVLTLLESER.E	1	RS28_HUMAN
P62857	2	3.5416	R.EGDVLTLLESER.E	1	RS28_HUMAN
40S ribosomal protein S29 - Homo sapiens (Human)					
P62273	2	3.461	K.DIGFIKLD.-	5	RS29_HUMAN
40S ribosomal protein S3 - Homo sapiens (Human)					
P23396	2	3.9876	K.GGKPEPPAMPQPVPTA.-	7	RS3_HUMAN
P23396	2	3.879	K.DEILPTTPISEQK.G	8	RS3_HUMAN
P23396	2	3.5388	K.GGKPEPPAM#PQPVPTA.-	2	RS3_HUMAN
40S ribosomal protein S30 - Homo sapiens (Human)					
P62861	2	2.7041	R.RMQYNR.R	1	RS30_HUMAN
P62861	2	3.416	R.RFVNVVPTFGKK.K	2	RS30_HUMAN
P62861	2	3.933	R.RFVNVVPTFGK.K	10	RS30_HUMAN
P62861	2	2.9714	R.FVNVVPTFGK.K	3	RS30_HUMAN
P62861	2	2.9352	R.FVNVVPTFGKK.K	1	RS30_HUMAN
40S ribosomal protein S3a - Homo sapiens (Human)					
P61247	2	4.1171	K.LMELHGEGSSSGK.A	4	RS3A_HUMAN
P61247	2	2.9479	R.VFEVSLADLQNDEVAFR.K	1	RS3A_HUMAN
P61247	2	3.8729	R.EVQTNDLKEVVNK.L	1	RS3A_HUMAN
P61247	2	3.7253	K.VERADGYEPPVQESV.-	2	RS3A_HUMAN
P61247	2	3.3917	K.FKLITEDVQGK.N	1	RS3A_HUMAN
P61247	2	4.5616	K.ATGDETGAKVER.A	6	RS3A_HUMAN
P61247	2	2.8967	K.APAMFNIR.N	2	RS3A_HUMAN
P61247	2	2.776	K.APAM#FNIR.N	1	RS3A_HUMAN
P61247	3	4.2718	R.VFEVSLADLQNDEVAFRK.F	1	RS3A_HUMAN
P61247	2	3.8477	K.LM#ELHGEGSSSGK.A	2	RS3A_HUMAN
40S ribosomal protein S7 - Homo sapiens (Human)					
P62081	3	4.0439	R.TLTAVHDAILEDLVFPSEIVGK.R	1	RS7_HUMAN
P62081	3	4.8626	R.TLTAVHDAILEDLVFPSEIVGKR.I	2	RS7_HUMAN
P62081	2	3.7921	K.LTGKDVNFEFPEFQL.-	3	RS7_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P62081	2	3.6017	K.DVNFEFPEFQL.-	2	RS7_HUMAN
P62081	3	3.9773	K.AQQNNVEHKVETFSGVYK.K	1	RS7_HUMAN
40S ribosomal protein S9 - Homo sapiens (Human)					
P46781	2	2.9123	R.SPYGGGRPGR.V	2	RS9_HUMAN
P46781	2	3.9264	K.KGQGGAGAGDDEEED.-	10	RS9_HUMAN
45 kDa calcium-binding protein precursor - Homo sapiens (Human)					
Q9BRK5	2	2.8189	K.YSEFFTGSK.L	1	CAB45_HUMAN
4F2 cell-surface antigen heavy chain - Homo sapiens (Human)					
P08195	2	3.848	R.LLTSFLPAQLLR.L	2	4F2_HUMAN
5'-3' exoribonuclease 1 - Homo sapiens (Human)					
Q8IZH2	3	4.8667	K.EAQSSQATPVQTSQPDSSNIVK.V	1	XRN1_HUMAN
5'-3' exoribonuclease 2 - Homo sapiens (Human)					
Q9H0D6	2	2.924	R.GVGAEPLLPWNR.M	1	XRN2_HUMAN
Q9H0D6	2	3.9064	R.NSPGSQVASNPR.Q	4	XRN2_HUMAN
5-azacytidine-induced protein 1 - Homo sapiens (Human)					
Q9UPN4	2	3.57	R.SNSTTQVSQPR.S	2	AZI1_HUMAN
Q9UPN4	3	4.0874	R.RPGSAATTKPIVR.S	1	AZI1_HUMAN
60S acidic ribosomal protein P1 - Homo sapiens (Human)					
P05386	2	4.1974	K.AAGVNVEPFWPGLFAK.A	4	RLA1_HUMAN
60S acidic ribosomal protein P2 - Homo sapiens (Human)					
P05387	2	5.4283	K.VISELNGKNIEDVIAQGIGK.L	2	RLA2_HUMAN
P05387	2	5.3214	-.M#RYVASYLLAALGGNSSPSAK.D	4	RLA2_HUMAN
P05387	2	4.5362	R.YVASYLLAALGGNSSPSAK.D	9	RLA2_HUMAN
P05387	1	2.5898	K.VISELNGK.N	5	RLA2_HUMAN
P05387	2	4.9159	K.NIEDVIAQGIGK.L	9	RLA2_HUMAN
P05387	3	3.9349	K.LASVPAGGAVAVSAAPGSAAPAAGSAPAAAE	1	RLA2_HUMAN
P05387	3	5.1775	K.KILDSVGIEADDDRLNK.V	9	RLA2_HUMAN
P05387	3	5.2299	K.ILDSVGIEADDDRLNKVISELNGK.N	3	RLA2_HUMAN
P05387	3	5.3302	K.ILDSVGIEADDDRLNK.V	8	RLA2_HUMAN
P05387	2	6.5868	-.MRYVASYLLAALGGNSSPSAK.D	3	RLA2_HUMAN
60S ribosomal protein L10-like - Homo sapiens (Human)					
Q96L21	2	4.1762	K.LQNEEHVIEALR.R	1	RL10L_HUMAN
60S ribosomal protein L11 - Homo sapiens (Human)					
P62913	2	3.3533	K.YDGIILPGK.-	1	RL11_HUMAN
P62913	3	4.1802	R.WFQQKYDGIILPGK.-	1	RL11_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
60S ribosomal protein L13 - Homo sapiens (Human)					
P26373	3	4.7021	K.KGDSSAEELKLATQLTGPVMPVR.N	2	RL13_HUMAN
P26373	2	2.9085	R.VITEEEK.N	1	RL13_HUMAN
P26373	2	3.1644	R.GFSLEELR.V	1	RL13_HUMAN
P26373	2	4.9259	R.AKEAAEQDVEKK.K	3	RL13_HUMAN
P26373	3	5.0307	R.AKEAAEQDVEK.K	8	RL13_HUMAN
P26373	2	5.0585	K.STESLQANVQR.L	4	RL13_HUMAN
P26373	2	3.6096	R.VITEEEKNFK.A	6	RL13_HUMAN
P26373	3	5.1744	K.KGDSSAEELKLATQLTGPVM#PVR.N	2	RL13_HUMAN
P26373	2	3.5862	K.KGDSSAEELK.L	11	RL13_HUMAN
P26373	2	3.3468	K.EAAEQDVEKKK.-	2	RL13_HUMAN
P26373	2	3.0777	K.EAAEQDVEKK.K	3	RL13_HUMAN
P26373	1	2.7332	K.EAAEQDVEK.K	2	RL13_HUMAN
P26373	3	4.008	K.ARVITEEEKNFK.A	2	RL13_HUMAN
P26373	2	3.5	K.LATQLTGPVM#PVR.N	3	RL13_HUMAN
P26373	2	3.3552	K.LATQLTGPVMPVR.N	4	RL13_HUMAN
60S ribosomal protein L17 - Homo sapiens (Human)					
P18621	2	3.7031	K.EQIVPKPEEEVAQK.K	4	RL17_HUMAN
P18621	1	2.9724	R.YSLDPENPTK.S	3	RL17_HUMAN
60S ribosomal protein L18 - Homo sapiens (Human)					
Q07020	2	4.3207	K.ILTFDQLALDSPK.G	4	RL18_HUMAN
Q07020	2	2.8749	K.TAVVVGTITDDVR.V	1	RL18_HUMAN
Q07020	2	3.1698	K.APGTPHSHTKPYVR.S	2	RL18_HUMAN
60S ribosomal protein L18a - Homo sapiens (Human)					
Q02543	2	2.9484	K.VEEIAASK.C	1	RL18A_HUMAN
60S ribosomal protein L19 - Homo sapiens (Human)					
P84098	2	2.805	K.RLASSVLR.C	1	RL19_HUMAN
P84098	2	3.263	K.TLSKEEETK.K	5	RL19_HUMAN
P84098	2	3.6307	K.TLSKEEETKK.-	12	RL19_HUMAN
P84098	1	2.2684	R.LASSVLR.C	1	RL19_HUMAN
P84098	3	5.6917	K.KVWLDPNETNEIANANSR.Q	5	RL19_HUMAN
P84098	2	3.8659	K.KLLADQAEAR.R	3	RL19_HUMAN
P84098	3	6.1746	K.KKVWLDPNETNEIANANSR.Q	2	RL19_HUMAN
P84098	2	3.517	K.LLADQAEAR.R	2	RL19_HUMAN
P84098	2	2.8619	K.VWLDPNETNEIANANSR.Q	1	RL19_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
60S ribosomal protein L22 - Homo sapiens (Human)					
P35268	2	4.0305	K.AGNLGGGVVTIER.S	4	RL22_HUMAN
P35268	2	2.8225	K.ITVTSEVPFSK.R	2	RL22_HUMAN
P35268	2	3.8731	R.VVANSKESYELR.Y	1	RL22_HUMAN
60S ribosomal protein L23 - Homo sapiens (Human)					
P62829	2	3.7785	K.GSAITGPVAK.E	4	RL23_HUMAN
P62829	1	2.2764	R.IASNAGSIA.-	3	RL23_HUMAN
60S ribosomal protein L23a - Homo sapiens (Human)					
P62750	2	2.7175	K.FPLTTESAMK.K	1	RL23A_HUMAN
P62750	2	3.8375	R.LAPDYDALDVANKIGII.-	5	RL23A_HUMAN
P62750	2	3.8114	R.LAPDYDALDVANK.I	4	RL23A_HUMAN
P62750	2	2.7094	K.VNTLIRPDGEKK.A	1	RL23A_HUMAN
P62750	1	2.8709	K.LYDIDVAK.V	3	RL23A_HUMAN
P62750	2	3.3299	K.KLYDIDVAK.V	4	RL23A_HUMAN
P62750	2	3.0231	K.KEAPAPPK.A	2	RL23A_HUMAN
P62750	2	4.309	K.KEAPAPPKAEAK.A	10	RL23A_HUMAN
60S ribosomal protein L24 - Homo sapiens (Human)					
P83731	2	5.0274	R.AITGASLADIM#AK.R	4	RL24_HUMAN
P83731	2	4.9368	R.AITGASLADIMAK.R	6	RL24_HUMAN
P83731	2	4.0006	R.AITGASLADIMAKR.N	1	RL24_HUMAN
60S ribosomal protein L26 - Homo sapiens (Human)					
P61254	1	2.3072	K.YKEETIEK.M	1	RL26_HUMAN
P61254	3	3.8526	K.GKYKEETIEKM#QE.-	1	RL26_HUMAN
P61254	3	3.9857	K.GKYKEETIEK.M	4	RL26_HUMAN
P61254	2	3.7247	K.YKEETIEKM#QE.-	1	RL26_HUMAN
60S ribosomal protein L27a - Homo sapiens (Human)					
P46776	2	3.2982	K.LWTLVSEQTR.V	2	RL27A_HUMAN
P46776	2	3.8804	K.TGAAPIIDVVR.S	6	RL27A_HUMAN
P46776	2	2.7369	R.NQSFCPTVNLDK.L	1	RL27A_HUMAN
60S ribosomal protein L29 - Homo sapiens (Human)					
P47914	2	5.1613	K.AQAAAPASVPAQAPK.R	7	RL29_HUMAN
P47914	2	4.4321	K.AQAAAPASVPAQAPKR.T	2	RL29_HUMAN
P47914	2	3.0868	K.DQTKAQAAAPASVPAQAPK.R	1	RL29_HUMAN
P47914	2	2.8052	R.SQRYESLK.G	1	RL29_HUMAN
60S ribosomal protein L3 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P39023	2	3.0106	R.HGSLGFLPR.K	1	RL3_HUMAN
60S ribosomal protein L31 - Homo sapiens (Human)					
P62899	1	2.8453	K.NLQTVNVDEN.-	2	RL31_HUMAN
P62899	2	3.421	R.SAINEVVTR.E	4	RL31_HUMAN
60S ribosomal protein L32 - Homo sapiens (Human)					
P62910	2	2.7779	K.GQILMPNIGYGSNK.K	1	RL32_HUMAN
60S ribosomal protein L34 - Homo sapiens (Human)					
P49207	2	3.1564	R.AFLIEEQK.I	1	RL34_HUMAN
P49207	2	2.7524	R.RLSYNTASNK.T	1	RL34_HUMAN
60S ribosomal protein L35 - Homo sapiens (Human)					
P42766	1	2.1678	K.KKEELLK.Q	1	RL35_HUMAN
P42766	2	3.7014	R.VLTVINQTQK.E	1	RL35_HUMAN
P42766	2	4.4223	R.VLTVINQTQKENLR.K	3	RL35_HUMAN
60S ribosomal protein L35a - Homo sapiens (Human)					
P18077	2	3.0774	K.NNTVTPGGKPNK.T	2	RL35A_HUMAN
60S ribosomal protein L36 - Homo sapiens (Human)					
Q9Y3U8	3	4.0788	R.KREELSNVLAAMR.K	4	RL36_HUMAN
Q9Y3U8	2	4.4971	K.REELSNVLAAMR.K	2	RL36_HUMAN
Q9Y3U8	2	3.5738	R.EELSNVLAAMR.K	2	RL36_HUMAN
60S ribosomal protein L36a - Homo sapiens (Human)					
P83881	1	2.428	K.DSLYAQGK.R	2	RL36A_HUMAN
P83881	2	3.7493	K.GKDSLYAQGK.R	2	RL36A_HUMAN
60S ribosomal protein L36a-like - Homo sapiens (Human)					
Q969Q0	1	2.3439	K.DSLYAQGR.R	2	RL36L_HUMAN
Q969Q0	2	4.0672	K.GKDSLYAQGR.R	5	RL36L_HUMAN
60S ribosomal protein L37a - Homo sapiens (Human)					
P61513	2	2.8343	K.KIEISQHAK.Y	2	RL37A_HUMAN
P61513	2	3.9107	K.TVAGGAWTYNTTSAVTVK.S	2	RL37A_HUMAN
60S ribosomal protein L38 - Homo sapiens (Human)					
P63173	2	4.2019	K.IEEIKDFLLTAR.R	7	RL38_HUMAN
P63173	2	4.1761	R.YLYTLVITDKEK.A	3	RL38_HUMAN
P63173	2	3.2113	K.LKQSLPPGLAVK.E	3	RL38_HUMAN
P63173	2	5.7097	R.KIEEIKDFLLTAR.R	12	RL38_HUMAN
60S ribosomal protein L4 - Homo sapiens (Human)					
P36578	1	5.2782	K.AAAAAAALQAK.S	4	RL4_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P36578	3	4.7738	K.APIRPDIVNFVHTNLR.K	2	RL4_HUMAN
P36578	3	5.1963	R.VDKAAAAAAALQAK.S	3	RL4_HUMAN
60S ribosomal protein L5 - Homo sapiens (Human)					
P46777	2	3.1191	K.RFPGYDSESK.E	1	RL5_HUMAN
P46777	2	2.9751	K.NSVTPDMMEEMYKK.A	1	RL5_HUMAN
P46777	2	3.5234	R.YLMEEDEDAYKK.Q	2	RL5_HUMAN
P46777	2	4.2205	K.HIMGQNVADYMR.Y	3	RL5_HUMAN
P46777	2	3.6949	K.GAVDGGLSIPHSTK.R	2	RL5_HUMAN
P46777	2	3.3185	K.HIM#GQNVADYMR.Y	2	RL5_HUMAN
P46777	2	3.2501	R.YLM#EEDEDAYKK.Q	1	RL5_HUMAN
60S ribosomal protein L8 - Homo sapiens (Human)					
P62917	2	3.4997	K.GIVKDIIHDPGR.G	1	RL8_HUMAN
P62917	2	5.4874	R.ASGNYATVISHNPETK.K	3	RL8_HUMAN
P62917	2	4.3113	R.AVVGVVAGGGR.I	5	RL8_HUMAN
60S ribosomal protein L9 - Homo sapiens (Human)					
P32969	2	4.8703	K.TILSNQTVDIPENVDITLK.G	3	RL9_HUMAN
P32969	1	2.5033	K.GTVQQADE.-	8	RL9_HUMAN
A kinase - Homo sapiens (Human)					
Q5T721	3	5.1998	R.AAGVGWENVLLKEGESASNATETSGPDMTIK	1	Q5T721_HUMAN
Q5T721	2	4.7052	K.KPPQLSEDDIWLK.S	2	Q5T721_HUMAN
Q5T721	2	3.2954	R.AAGVGWENVLLK.E	3	Q5T721_HUMAN
Abhydrolase domain-containing protein 12 - Homo sapiens (Human)					
Q8N2K0	3	4.3517	K.RTEPVALEHER.C	2	ABD12_HUMAN
ABI gene family member 3 - Homo sapiens (Human)					
Q9P2A4	2	2.7748	K.VVTLYPYTSQK.D	1	ABI3_HUMAN
Q9P2A4	3	4.4277	R.IPEPVHLPVVPDGR.L	2	ABI3_HUMAN
Q9P2A4	3	5.3855	R.LSAASSASSLASAGSAEGVGGAPTPK.G	2	ABI3_HUMAN
Abl interactor 1 - Homo sapiens (Human)					
Q8IZP0	3	5.9641	R.KPIDYTVLDDVGHGVK.W	2	ABI1_HUMAN
Q8IZP0	2	2.7427	R.VTGLFPGNYVESIMHYTD.-	1	ABI1_HUMAN
Q8IZP0	2	3.8585	R.HNSTTSSTSSGGYRR.T	2	ABI1_HUMAN
Q8IZP0	2	4.1746	R.HNSTTSSTSSGGYR.R	2	ABI1_HUMAN
Q8IZP0	2	2.8735	R.ALIESYQNLTR.V	1	ABI1_HUMAN
Q8IZP0	3	4.1802	K.TLEPVKPPTVPNDYMTSPAR.L	2	ABI1_HUMAN
Q8IZP0	2	3.1874	R.LGSQHSPGR.T	2	ABI1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Abl interactor 2 - Homo sapiens (Human)					
Q9NYB9	3	5.6067	R.KPIDYTILDDIGHGVK.W	3	ABI2_HUMAN
Q9NYB9	2	2.7706	R.NM#APSQQSPVR.T	1	ABI2_HUMAN
Q9NYB9	2	2.8858	R.NMAPSQQSPVR.T	1	ABI2_HUMAN
Abnormal spindle-like microcephaly-associated protein - Homo sapiens (Human)					
Q8IZT6	1	2.3243	K.EEIAFLK.H	1	ASPM_HUMAN
Absent in melanoma 1 protein - Homo sapiens (Human)					
Q9Y4K1	2	4.0246	K.GTAAESGEEAAR.A	3	AIM1_HUMAN
Q9Y4K1	2	6.4446	R.SFVLPVESTQDVSSQVIPESSEVR.E	4	AIM1_HUMAN
Q9Y4K1	2	4.1225	R.AALDGGVASAASPESKPSPGTK.G	1	AIM1_HUMAN
Q9Y4K1	2	3.5014	K.VTVSEEEILPATR.G	2	AIM1_HUMAN
Q9Y4K1	2	3.5505	K.TLPIQAQSQGSR.T	4	AIM1_HUMAN
Q9Y4K1	2	4.6137	K.SLVLENVTDTAQDIPTTVDTK.D	6	AIM1_HUMAN
Q9Y4K1	2	3.6314	K.SGPQVIPPASEK.T	2	AIM1_HUMAN
Q9Y4K1	2	3.8795	K.LLEKEDSEAADSK.S	3	AIM1_HUMAN
Q9Y4K1	1	2.6071	K.ETAIETK.V	2	AIM1_HUMAN
Q9Y4K1	2	2.8634	K.RASAEQSVLFK.S	2	AIM1_HUMAN
Acidic leucine-rich nuclear phosphoprotein 32 family member A - Homo sapiens (Human)					
P39687	2	4.08	R.KREPEDEGEDDD.-	50	AN32A_HUMAN
P39687	2	3.3231	K.REPEDEGEDDD.-	2	AN32A_HUMAN
Acidic leucine-rich nuclear phosphoprotein 32 family member B - Homo sapiens (Human)					
Q92688	2	2.8082	R.IFGGLDMLAEK.L	2	AN32B_HUMAN
Q92688	2	2.729	K.KLELSENR.I	1	AN32B_HUMAN
Q92688	3	4.2163	R.IFGGLDM#LAEKLPNLTHLNLSGNK.L	1	AN32B_HUMAN
Acidic leucine-rich nuclear phosphoprotein 32 family member C - Homo sapiens (Human)					
O43423	3	3.8504	K.IKDLSTIEPLKQLENLK.S	1	AN32C_HUMAN
Actin filament-associated protein 1-like 1 - Homo sapiens (Human)					
Q8TED9	2	3.1354	R.SPSIVASNQGR.V	2	AF1L1_HUMAN
Actin filament-associated protein 1-like 2 - Homo sapiens (Human)					
Q8N4X5	3	3.9104	K.AVTPASAPDCTPVNSATTLK.N	2	AF1L2_HUMAN
Q8N4X5	2	2.8079	K.NRTEAEVKR.Y	1	AF1L2_HUMAN
Q8N4X5	3	3.773	K.YLSASEYGSSVDGHPEVPETK.D	1	AF1L2_HUMAN
Q8N4X5	2	2.9108	R.YTEEKERLEK.K	1	AF1L2_HUMAN
Actin-bundling protein with BAIAP2 homology - Homo sapiens (Human)					
Q765P7	2	3.3122	K.HGEEVSPAASDLAMVLTR.G	1	Q765P7_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q765P7	3	4.7132	R.VELLRDTEPGPASGGTLGPSGEEAPRPR.M	1	Q765P7_HUMAN
Q765P7	3	4.0343	K.VGSHEQPSGATLQR.R	3	Q765P7_HUMAN
Q765P7	2	2.7976	R.NSNIAQNYR.R	2	Q765P7_HUMAN
Actin-related protein 2/3 complex subunit 1A - Homo sapiens (Human)					
Q92747	2	3.8637	K.TLESSIQGLR.I	3	ARC1A_HUMAN
Actin-related protein 2/3 complex subunit 1B - Homo sapiens (Human)					
O15143	2	5.4343	K.ASSEGGTAAGAGLDSLHK.N	3	ARC1B_HUMAN
O15143	3	6.3371	K.KASSEGGTAAGAGLDSLHK.N	2	ARC1B_HUMAN
O15143	2	4.574	K.NSVSQISVLSGGK.A	3	ARC1B_HUMAN
O15143	2	2.9549	K.SLESALKDLK.I	2	ARC1B_HUMAN
Actin-related protein 2/3 complex subunit 3 - Homo sapiens (Human)					
O15145	2	3.2409	K.LIGNMALLPIR.S	2	ARPC3_HUMAN
O15145	2	3.5553	K.LIGNM#ALLPIR.S	2	ARPC3_HUMAN
Actin-related protein 2/3 complex subunit 5 - Homo sapiens (Human)					
O15511	2	3.6813	K.ALAAGGVGSIVR.V	2	ARPC5_HUMAN
Activated RNA polymerase II transcriptional coactivator p15 - Homo sapiens (Human)					
P53999	2	3.5826	K.ELVSSSSSGSDSDSEVDK.K	1	TCP4_HUMAN
P53999	3	3.7872	K.GISLNPEQWSQLKEQISDIDDAVR.K	1	TCP4_HUMAN
P53999	2	2.7814	K.KQVAPEKPVK.K	1	TCP4_HUMAN
P53999	2	2.8527	K.QSSSSRDDNMFQIGK.M	1	TCP4_HUMAN
P53999	2	2.7975	R.DDNM#FQIGK.M	1	TCP4_HUMAN
P53999	2	2.9967	R.DDNMFQIGK.M	3	TCP4_HUMAN
P53999	3	7.3197	R.KGISLNPEQWSQLKEQISDIDDAVR.K	2	TCP4_HUMAN
Activating signal cointegrator 1 complex subunit 2 - Homo sapiens (Human)					
Q9H1I8	2	4.0673	R.HDSSTAVAGSPR.G	2	ASCC2_HUMAN
Activity-dependent neuroprotector - Homo sapiens (Human)					
Q9H2P0	2	3.4768	K.SVGQGYSVGQSMR.L	1	ADNP_HUMAN
Q9H2P0	2	3.1194	R.IGSLASGNVR.S	2	ADNP_HUMAN
Acyl carrier protein, mitochondrial precursor - Homo sapiens (Human)					
O14561	2	3.1157	K.LSVNSHFMK.D	2	ACPM_HUMAN
O14561	2	2.7053	K.LYDKIDPEK.L	1	ACPM_HUMAN
O14561	3	5.2272	K.LYDKIDPEKLSVNSHFM#K.D	4	ACPM_HUMAN
O14561	3	5.0802	K.LYDKIDPEKLSVNSHFMK.D	5	ACPM_HUMAN
O14561	1	2.2559	R.VLYVLK.L	2	ACPM_HUMAN
Acyl-CoA synthetase medium-chain family member 3 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q53FZ2	2	2.7399	R.KVEFIQELPK.T	1	Q53FZ2_HUMAN
Acylphosphatase-1 - Homo sapiens (Human)					
P07311	2	3.4977	K.LGLVGWVQNTDR.G	2	ACYP1_HUMAN
Acylphosphatase-2 - Homo sapiens (Human)					
P14621	2	2.8391	K.LEYSNFSIRY.-	1	ACYP2_HUMAN
Adaptin ear-binding coat-associated protein 1 - Homo sapiens (Human)					
Q8NC96	2	3.7212	R.GGGLSLLPPPPGGK.V	2	NECP1_HUMAN
Adaptor protein HOFI - Homo sapiens (Human)					
A1X283	2	5.4918	K.SLLDGEGPQAVGGQDVAFSR.S	2	A1X283_HUMAN
A1X283	3	5.7375	K.SQDKSLLDGEGPQAVGGQDVAFSR.S	3	A1X283_HUMAN
A1X283	3	4.4183	K.SKTDLPEEKPDATPQNPFLK.S	1	A1X283_HUMAN
Adenomatous polyposis coli protein - Homo sapiens (Human)					
P25054	3	4.9083	K.VTSHTELTSNQQSANK.T	1	APC_HUMAN
P25054	2	3.3663	R.LQGSSLSSESAR.H	2	APC_HUMAN
P25054	2	2.8621	R.SPTGNTPPVIDSVSEK.A	1	APC_HUMAN
Adenylate cyclase type 9 - Homo sapiens (Human)					
O60503	2	2.9303	K.TSLGSDSSTQAK.D	2	ADCY9_HUMAN
Adenylate kinase isoenzyme 2, mitochondrial - Homo sapiens (Human)					
P54819	3	4.1008	R.LQAYHTQTTPLIEYYR.K	1	KAD2_HUMAN
P54819	2	4.4091	K.LVSDEMVVELIEK.N	2	KAD2_HUMAN
P54819	2	3.0221	R.AVLLGPPGAGK.G	2	KAD2_HUMAN
Adenylosuccinate synthetase isozyme 1 - Homo sapiens (Human)					
Q8N142	2	4.713	R.GRLQQEAAATGSR.V	1	PURA1_HUMAN
Q8N142	2	3.2867	R.LQQEAAATGSR.V	1	PURA1_HUMAN
Adipophilin - Homo sapiens (Human)					
Q99541	2	5.138	K.KVEGFDLVQKPSYYVR.L	4	ADFP_HUMAN
Q99541	2	4.848	K.SELLVEQYLPLTEEELEK.E	2	ADFP_HUMAN
Q99541	2	4.796	K.SVVSGSINTVLGSR.M	4	ADFP_HUMAN
Q99541	2	3.1417	R.MMQLVSSGVENALTK.S	2	ADFP_HUMAN
Q99541	2	4.5007	K.DSVASTITGVMDK.T	8	ADFP_HUMAN
Q99541	2	3.9514	K.DSVASTITGVM#DK.T	3	ADFP_HUMAN
Q99541	2	3.5935	K.DAVTTTVTGAK.D	5	ADFP_HUMAN
Q99541	2	3.1475	K.TKGAVTGSVEK.T	4	ADFP_HUMAN
Q99541	1	2.9607	K.GAVTGSVEK.T	5	ADFP_HUMAN
Adipose most abundant gene transcript 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q15847	3	5.4645	K.QQVEGTAQEAVSAAGAAAQQVVDQATEAGQ	6	APM2_HUMAN
Q15847	1	3.8238	K.TANQASDTFSGIGK.K	8	APM2_HUMAN
Q15847	2	3.7998	K.TANQASDTFSGIGKK.F	1	APM2_HUMAN
Q15847	2	6.4666	K.TTQETIDKTANQASDTFSGIGK.K	22	APM2_HUMAN
Q15847	3	6.5175	K.TTQETIDKTANQASDTFSGIGKK.F	18	APM2_HUMAN
Adipose specific 2 - Homo sapiens (Human)					
Q5TBU2	2	3.4831	K.TTQETIDKTAN.-	5	Q5TBU2_HUMA
ADM precursor [Contains: Adrenomedullin - Homo sapiens (Human)					
P35318	2	4.2788	R.SPEDSSPDAAR.I	4	ADML_HUMAN
ADP-ribose pyrophosphatase, mitochondrial precursor - Homo sapiens (Human)					
Q9BW91	2	3.4396	R.EFGEEALNSLQK.T	2	NUDT9_HUMAN
Q9BW91	2	4.4881	R.WADPQISESNFSPK.F	3	NUDT9_HUMAN
Q9BW91	3	3.9342	R.WGPNHAADPIITR.W	1	NUDT9_HUMAN
Q9BW91	2	3.7012	K.LFSQDHLVIYK.G	1	NUDT9_HUMAN
ADP-ribosylation factor GTPase-activating protein 1 - Homo sapiens (Human)					
Q8N6T3	2	4.724	K.IFDDVSSGVSQLASK.V	3	ARFG1_HUMAN
ADP-ribosylation factor GTPase-activating protein 3 - Homo sapiens (Human)					
Q9NP61	2	3.7783	R.LAYKDLEIQMK.K	2	ARFG3_HUMAN
Q9NP61	3	7.1004	R.SVISHSVTSDMQTIEQESPIMAKPR.K	1	ARFG3_HUMAN
Q9NP61	3	5.1519	R.SVISHSVTSDMQTIEQESPIM#AKPR.K	2	ARFG3_HUMAN
Q9NP61	2	3.1343	R.SSSFSSWDDSSDSYWKK.E	1	ARFG3_HUMAN
Q9NP61	2	2.9065	K.ETSKDTETVLK.T	2	ARFG3_HUMAN
Q9NP61	3	3.9238	K.QAQAADKMKEQEDLAK.V	1	ARFG3_HUMAN
Q9NP61	2	2.8209	K.LANTCFNEIEK.Q	1	ARFG3_HUMAN
Q9NP61	2	3.1644	K.MKEQEDLAK.V	2	ARFG3_HUMAN
Q9NP61	2	2.9993	K.AISSDMYFGR.Q	2	ARFG3_HUMAN
Q9NP61	2	3.853	K.VVSKEESIVSSLR.L	6	ARFG3_HUMAN
ADP-ribosylation factor-like protein 3 - Homo sapiens (Human)					
P36405	2	2.8156	R.ILLLGLDNAGK.T	2	ARL3_HUMAN
ADP-sugar pyrophosphatase - Homo sapiens (Human)					
Q9UKK9	2	2.901	K.EQTADGVAVIPVLQR.T	1	NUDT5_HUMAN
Adrenodoxin, mitochondrial precursor - Homo sapiens (Human)					
P10109	2	3.0831	R.DGETLTTK.G	1	ADX_HUMAN
P10109	1	2.7028	R.VPETVADAR.Q	3	ADX_HUMAN
P10109	2	4.1881	K.SMDNMTVRVPETVADAR.Q	2	ADX_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P10109	2	2.7304	K.SMDNMTVR.V	1	ADX_HUMAN
AF4/FMR2 family member 4 - Homo sapiens (Human)					
Q9UHB7	2	2.806	K.SPAQSDSTTQR.R	1	AFF4_HUMAN
Q9UHB7	2	3.5484	R.EQGTGNSYTDTSGPK.E	1	AFF4_HUMAN
Q9UHB7	2	4.6499	R.GGLKIESETPVDLASSMPSSR.H	1	AFF4_HUMAN
Q9UHB7	2	2.9652	K.IDLNLLTR.I	2	AFF4_HUMAN
Afadin - Homo sapiens (Human)					
P55196	2	3.8192	K.TQVLSPDSLFTAK.F	3	AFAD_HUMAN
P55196	3	4.1623	R.DYEPPSPSPAPGAPPPPPQR.N	4	AFAD_HUMAN
Afamin precursor - Homo sapiens (Human)					
P43652	2	4.7654	K.IAPQLSTEELVSLGEK.M	2	AFAM_HUMAN
Aftiphilin - Homo sapiens (Human)					
Q6ULP2	3	4.0743	K.EEFVPSNHFMPIHEFSENVDSLTSFK.S	1	AFTIN_HUMAN
Q6ULP2	2	4.2883	R.TNMNVVHQNK.Q	3	AFTIN_HUMAN
Q6ULP2	3	5.3487	R.KFTNFQSPNIDPTEENDLDDSLSVK.N	2	AFTIN_HUMAN
Q6ULP2	2	3.1837	K.QGLPTLQQDEFLQSGVQSK.A	1	AFTIN_HUMAN
Q6ULP2	2	3.5653	K.TEEKLDLLTSK.C	2	AFTIN_HUMAN
Q6ULP2	2	3.3968	K.TSDDEVGSPK.E	1	AFTIN_HUMAN
Aggrecan core protein precursor - Homo sapiens (Human)					
P16112	2	3.8252	R.TPCVGDKDSSPGVR.T	2	PGCA_HUMAN
P16112	2	3.0688	R.VNSAYQDK.V	1	PGCA_HUMAN
P16112	2	2.7987	R.AISTRYTLDFDR.A	1	PGCA_HUMAN
P16112	3	3.7293	K.VSLPNYPAIPSDATLEVQSLR.S	1	PGCA_HUMAN
P16112	2	2.9725	K.GTVACGEPPVVEHAR.T	1	PGCA_HUMAN
P16112	2	3.4612	K.FTFQEAANECR.R	4	PGCA_HUMAN
P16112	2	3.0316	K.DRYEINSLVR.Y	2	PGCA_HUMAN
P16112	2	3.2056	R.YQCTEGFVQR.H	4	PGCA_HUMAN
Agrin precursor - Homo sapiens (Human)					
O00468	2	3.15	R.SFLAFPTLR.A	2	AGRIN_HUMAN
O00468	2	2.7985	K.VLGAPVPAFEGR.S	1	AGRIN_HUMAN
O00468	1	2.3823	R.FNAVCLSR.R	2	AGRIN_HUMAN
O00468	3	3.8078	R.RPEMLLRPESCPAR.Q	1	AGRIN_HUMAN
O00468	2	3.2142	R.SADGLTASCLCPATCR.G	1	AGRIN_HUMAN
AHNAK protein - Homo sapiens (Human)					
Q9BVU3	2	3.44	R.SPEPGQTWTR.E	4	Q9BVU3_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BVU3	2	4.0122	R.ELLLPNWQGSGSHGLTIAQR.D	1	Q9BVU3_HUMA
Q9BVU3	2	4.6319	R.DDGVFVQEVTQNSPAAR.T	12	Q9BVU3_HUMA
Q9BVU3	2	3.3952	K.QKEASSQAGAVSVSTPNAGL.-	1	Q9BVU3_HUMA
AHNAK2 protein - Homo sapiens (Human)					
Q5BKX7	2	2.9981	K.ADDESKGSGLGPNEG.-	2	Q5BKX7_HUMA
Q5BKX7	2	3.0144	K.HDLSTEGDSR.G	1	Q5BKX7_HUMA
Q5BKX7	2	2.8335	R.DVDPSLSSATAGGSFQDTEK.A	1	Q5BKX7_HUMA
A-kinase anchor protein 13 - Homo sapiens (Human)					
Q12802	3	4.0443	K.SSPICSTTGDDKLCADSACQQNTVTSSGDLV	1	AKP13_HUMAN
Q12802	2	2.9777	R.IVDAVIEQVK.A	1	AKP13_HUMAN
Q12802	2	3.5684	K.NAASDAEMNHR.S	2	AKP13_HUMAN
Q12802	2	3.4837	K.GADLIEEAASR.I	1	AKP13_HUMAN
Q12802	2	5.6323	K.ALQLSNSPGASSAFLK.A	3	AKP13_HUMAN
Q12802	3	4.4667	R.SREESADAPVDQNSVVIPAAAK.D	2	AKP13_HUMAN
A-kinase anchor protein 4 precursor - Homo sapiens (Human)					
Q5JQC9	3	3.8476	K.SLTSAEKVGEHILKEGLTIWNQK.Q	2	AKAP4_HUMAN
A-kinase anchor protein 8 - Homo sapiens (Human)					
O43823	2	3.5401	R.GGSGGGGEGIQDR.E	2	AKAP8_HUMAN
Alcohol dehydrogenase [NADP+] - Homo sapiens (Human)					
P14550	2	2.8562	R.DAGHPLYPFNDPY.-	1	AK1A1_HUMAN
Alpha-1-acid glycoprotein 1 precursor - Homo sapiens (Human)					
P02763	2	3.3209	K.SDVVYTDWKKDK.C	2	A1AG1_HUMAN
P02763	2	5.3711	R.YVGGQEHFAHLLILR.D	12	A1AG1_HUMAN
P02763	2	3.3009	R.YVGGQEHFAHLLILRDTK.T	2	A1AG1_HUMAN
P02763	3	3.9769	K.TYM#LAFDVNDEKNWGLSVYADKPETTK.E	1	A1AG1_HUMAN
P02763	2	3.3004	K.SDVVYTDWK.K	10	A1AG1_HUMAN
P02763	2	4.4517	K.EQLGEFYEALDCLR.I	10	A1AG1_HUMAN
P02763	3	5.4075	K.TYMLAFDVNDEKNWGLSVYADKPETTK.E	2	A1AG1_HUMAN
P02763	1	3.584	K.SDVVYTDWKK.D	8	A1AG1_HUMAN
Alpha-1-acid glycoprotein 2 precursor - Homo sapiens (Human)					
P19652	1	3.152	R.SDVMYTDWK.K	5	A1AG2_HUMAN
P19652	2	2.7151	R.SDVMYTDWKK.D	1	A1AG2_HUMAN
P19652	2	2.9054	R.SDVM#YTDWKK.D	2	A1AG2_HUMAN
P19652	2	2.9391	R.SDVM#YTDWK.K	1	A1AG2_HUMAN
P19652	2	3.4784	R.EHVAHLLFLR.D	4	A1AG2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P19652	3	4.3797	K.TLMFGSYLDDEKNWGLSFYADKPETTK.E	1	A1AG2_HUMAN
P19652	2	4.4105	R.YEGGREHVAHLLFLR.D	6	A1AG2_HUMAN
Alpha-1-antichymotrypsin precursor - Homo sapiens (Human)					
P01011	2	4.7184	K.ITLLSALVETR.T	4	AACT_HUMAN
P01011	3	5.2817	R.DYNLNDILLQLGIEEAFTSK.A	2	AACT_HUMAN
P01011	2	3.2272	R.NLAVSQVVHK.A	1	AACT_HUMAN
Alpha-1-antitrypsin precursor - Homo sapiens (Human)					
P01009	3	3.7301	K.ELDRDTVFALVNYIFFK.G	1	A1AT_HUMAN
P01009	2	4.468	K.ITPNLAEFAFSLYR.Q	1	A1AT_HUMAN
P01009	2	3.5797	K.KLSSWVLLMK.Y	3	A1AT_HUMAN
Alpha-1B-glycoprotein precursor - Homo sapiens (Human)					
P04217	2	2.785	R.LETPDFQLFK.N	2	A1BG_HUMAN
Alpha-2-antiplasmin precursor - Homo sapiens (Human)					
P08697	2	3.8416	K.LGNQEPGGQTALK.S	3	A2AP_HUMAN
Alpha-2-glycoprotein 1, zinc-binding - Homo sapiens (Human)					
Q5XKQ4	2	3.4938	K.AYLEEECPATLR.K	2	Q5XKQ4_HUMA
Q5XKQ4	2	3.635	K.EIPAWVPFDPAAQITK.Q	1	Q5XKQ4_HUMA
Q5XKQ4	3	6.0105	K.NILDRQDPPSVVVTSHQAPGEK.K	3	Q5XKQ4_HUMA
Q5XKQ4	2	2.8855	K.QKWEAEPVYVQR.A	1	Q5XKQ4_HUMA
Q5XKQ4	3	3.8545	K.YYYDGKDYIEFNKEIPAWVPFDPAAQITK.Q	1	Q5XKQ4_HUMA
Q5XKQ4	2	3.0553	R.AGEVQEPELR.G	1	Q5XKQ4_HUMA
Q5XKQ4	3	4.0206	R.AKAYLEEECPATLRK.Y	1	Q5XKQ4_HUMA
Alpha-2-HS-glycoprotein precursor - Homo sapiens (Human)					
P02765	2	3.8128	K.HTLNQIDEVK.V	4	FETUA_HUMAN
P02765	3	5.3042	R.TVVQPSVGAAAGPVVPPCPGR.I	7	FETUA_HUMAN
P02765	2	3.9976	R.QLKEHAVEGDCDFQLLK.L	5	FETUA_HUMAN
P02765	2	5.861	R.HTFMGVVSLGSPSGEVSHPR.K	9	FETUA_HUMAN
P02765	2	3.2806	K.LDGKFSVVYAK.C	3	FETUA_HUMAN
P02765	1	2.2868	K.FSVVYAK.C	3	FETUA_HUMAN
P02765	2	3.6422	K.EHAVEGDCDFQLLK.L	3	FETUA_HUMAN
P02765	3	6.5038	R.HTFM#GVVSLGSPSGEVSHPR.K	7	FETUA_HUMAN
Alpha-2-macroglobulin precursor - Homo sapiens (Human)					
P01023	2	3.9748	R.IAQWQSFQLEGGLK.Q	1	A2MG_HUMAN
P01023	2	3.4826	K.AIGYLNTGYQR.Q	2	A2MG_HUMAN
P01023	2	3.4432	R.VGFYESDVMGR.G	2	A2MG_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P01023	2	3.5279	R.VGFYESDVM#GR.G	2	A2MG_HUMAN
P01023	3	5.4968	R.TEHPFTVEEFVLPK.F	3	A2MG_HUMAN
P01023	2	3.3019	R.SASNMAIVDVK.M	2	A2MG_HUMAN
P01023	2	3.7238	K.QFSFPLSSEPFQGSYK.V	4	A2MG_HUMAN
P01023	2	3.5646	K.NEDSLVFVQTDK.S	1	A2MG_HUMAN
P01023	2	3.3268	K.LPPNVVEESAR.A	2	A2MG_HUMAN
P01023	2	3.5612	K.HYDGSYSTFGER.Y	1	A2MG_HUMAN
P01023	2	3.9816	K.ALLAYAFALAGNQDK.R	1	A2MG_HUMAN
P01023	3	4.5209	K.DTVIKPLLVEPEGLEK.E	5	A2MG_HUMAN
Alpha-2-macroglobulin receptor-associated protein precursor - Homo sapiens (Human)					
P30533	2	4.8357	K.VHEYNVLLETLSR.T	7	AMRP_HUMAN
P30533	2	3.9472	R.TKELGYTVKK.H	2	AMRP_HUMAN
P30533	3	3.7715	R.TEEIHENVISPSDLSDIK.G	1	AMRP_HUMAN
P30533	3	5.684	R.RVSHQGYSTEAEFEEPR.V	2	AMRP_HUMAN
P30533	2	2.8355	R.NLNVILAK.Y	3	AMRP_HUMAN
P30533	2	4.3115	R.VSHQGYSTEAEFEEPR.V	2	AMRP_HUMAN
P30533	2	3.5531	R.HAESVGDGER.V	19	AMRP_HUMAN
P30533	1	2.9278	K.LNQLWEK.A	3	AMRP_HUMAN
P30533	2	4.7025	K.LKLDGLDEDGEKEAR.L	4	AMRP_HUMAN
P30533	2	4.0836	K.LDGLDEDGEKEAR.L	2	AMRP_HUMAN
P30533	2	2.9767	K.KHLQDLSGR.I	2	AMRP_HUMAN
P30533	1	2.5932	K.HLQDLSGR.I	1	AMRP_HUMAN
P30533	2	4.0784	K.FSGEELDKLWR.E	3	AMRP_HUMAN
P30533	3	4.0719	K.EKVHEYNVLLETLSR.T	2	AMRP_HUMAN
P30533	2	3.4985	R.LAELHADLK.I	2	AMRP_HUMAN
P30533	2	3.6428	K.LRHAESVGDGER.V	2	AMRP_HUMAN
Alpha-crystallin B chain - Homo sapiens (Human)					
P02511	3	5.6838	R.IPADVDPLTITSSLSSDGVLTVNGPR.K	7	CRYAB_HUMAN
P02511	2	3.2138	R.TIPITREEKPAVTAAPK.K	2	CRYAB_HUMAN
P02511	4	4.9023	R.KYRIPADVDPLTITSSLSSDGVLTVNGPR.K	1	CRYAB_HUMAN
P02511	2	3.4263	R.FSVNLDVK.H	6	CRYAB_HUMAN
P02511	1	2.2975	R.EEKPAVTAAPK.K	1	CRYAB_HUMAN
P02511	2	4.3859	R.APSWFDTGLSEMR.L	4	CRYAB_HUMAN
P02511	3	6.2511	K.YRIPADVDPLTITSSLSSDGVLTVNGPR.K	2	CRYAB_HUMAN
P02511	3	5.6321	K.VLGDVIEVHGKHEERQDEHGFISR.E	2	CRYAB_HUMAN
P02511	2	4.8821	K.VLGDVIEVHGKHEER.Q	4	CRYAB_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P02511	2	4.2703	K.VLGDVIEVHGK.H	5	CRYAB_HUMAN
P02511	2	4.6097	K.HFSPEELKVK.V	4	CRYAB_HUMAN
P02511	2	3.4141	K.HEERQDEHGFISR.E	2	CRYAB_HUMAN
P02511	2	2.9098	R.RPFFPFHSPSR.L	1	CRYAB_HUMAN
P02511	2	3.861	R.APSWFDTGLSEM#R.L	5	CRYAB_HUMAN
Alpha-enolase - Homo sapiens (Human)					
P06733	2	2.8779	R.GNPTVEVDLFTSK.G	1	ENOA_HUMAN
P06733	1	2.7037	R.IEEELGSK.A	5	ENOA_HUMAN
P06733	3	3.9343	K.YGKDATNVGDEGGFAPNILENKEGLELLK.T	1	ENOA_HUMAN
P06733	2	2.7421	K.LAMQEFM#ILPVGAANFR.E	1	ENOA_HUMAN
P06733	2	3.8238	K.LAM#QEFMILPVGAANFR.E	2	ENOA_HUMAN
P06733	3	4.3108	R.SGKYDLDFKSPDDPSR.Y	1	ENOA_HUMAN
Alpha-hemoglobin-stabilizing protein - Homo sapiens (Human)					
Q9NZD4	2	3.1162	K.ALQELRQELNTLANPFLAK.Y	1	AHSP_HUMAN
Q9NZD4	2	2.7627	K.SHELPSHPPPSS.-	2	AHSP_HUMAN
Q9NZD4	2	2.7847	R.QELNTLANPFLAK.Y	1	AHSP_HUMAN
Alpha-ketoglutarate-dependent dioxygenase alkB homolog 3 - Homo sapiens (Human)					
Q96Q83	2	2.9382	K.SQAIAQPATTAK.S	1	ALKB3_HUMAN
Alpha-parvin - Homo sapiens (Human)					
Q9NVD7	2	2.9265	K.DDSFLGK.L	1	PARVA_HUMAN
Alpha-protein kinase 2 - Homo sapiens (Human)					
Q86TB3	2	3.1916	K.KAGPETPGEK.K	1	ALPK2_HUMAN
Q86TB3	2	3.047	K.TLQTSTDSVSK.E	1	ALPK2_HUMAN
Q86TB3	3	4.531	R.YKLPTAPEAAENDYPGIQGETR.D	1	ALPK2_HUMAN
Alpha-synuclein - Homo sapiens (Human)					
P37840	2	6.205	K.TVEGAGSIAAATGFVK.K	5	SYUA_HUMAN
P37840	2	2.7514	K.TKQGVAEAAGK.T	1	SYUA_HUMAN
P37840	2	4.6963	K.TVEGAGSIAAATGFVKK.D	5	SYUA_HUMAN
P37840	3	5.2921	K.TKEGVVHGVATVAEK.T	3	SYUA_HUMAN
P37840	2	3.6458	K.AKEGVVAAAEK.T	2	SYUA_HUMAN
P37840	2	3.8864	K.EGVVHGVATVAEK.T	3	SYUA_HUMAN
P37840	3	4.38	K.TKEQVTNVGGAVVTGVTAVAQK.T	3	SYUA_HUMAN
Alpha-taxilin - Homo sapiens (Human)					
P40222	2	2.896	R.DVSEELSR.Q	1	TXLNA_HUMAN
P40222	2	3.2129	R.VQDLSAGGQGSLTDSGPER.R	1	TXLNA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P40222	2	3.8272	R.TAQSGALRDVSEELSR.Q	1	TXLNA_HUMAN
P40222	2	2.7417	R.SLKEEGVQR.A	1	TXLNA_HUMAN
P40222	2	2.9475	R.DKELEGLQVK.I	2	TXLNA_HUMAN
P40222	3	4.1227	K.SSPGQPEAGPEGAQERPSQAAPAVEAEGPG	2	TXLNA_HUMAN
P40222	2	3.7279	K.LRQENMELAER.L	1	TXLNA_HUMAN
P40222	2	2.713	K.ALLEMAEEK.T	1	TXLNA_HUMAN
P40222	2	2.9898	R.SRWESSNK.A	3	TXLNA_HUMAN
AMBP protein precursor [Contains: Alpha-1-microglobulin - Homo sapiens (Human)					
P02760	2	3.3368	R.ETLLQDFR.V	2	AMBP_HUMAN
P02760	2	3.9766	R.TVAACNLPIVR.G	5	AMBP_HUMAN
Aminopeptidase N - Homo sapiens (Human)					
P15144	2	3.1061	R.QYMPWEAALSSLSYFK.L	1	AMPN_HUMAN
Amyloid beta A4 precursor protein-binding family B member 1-interacting protein - Homo sapiens					
Q7Z5R6	2	3.9851	R.SSDTSGSPATPLK.A	3	AB1IP_HUMAN
Q7Z5R6	3	5.7774	K.RQENPGHPGGAGGGEQDFMSDLMK.A	2	AB1IP_HUMAN
Amyloid beta A4 precursor protein-binding family B member 2 - Homo sapiens (Human)					
Q92870	2	2.8026	R.NSPATPPNTLNLR.S	1	APBB2_HUMAN
Q92870	2	2.7612	R.GVLSLIDTLK.Q	1	APBB2_HUMAN
Q92870	2	3.0276	K.SFLNYYADLETSAR.E	1	APBB2_HUMAN
Amyloid beta A4 protein precursor - Homo sapiens (Human)					
P05067	3	5.2179	K.AVIQHFQEKVESLEQEAANER.Q	2	A4_HUMAN
P05067	3	3.9552	K.THPHFVIPYR.C	1	A4_HUMAN
P05067	2	4.1136	R.ISYGNDALM#PSLTETK.T	2	A4_HUMAN
P05067	2	5.2195	R.ISYGNDALMPSLTETK.T	2	A4_HUMAN
Amyotrophic lateral sclerosis 2 chromosomal region candidate gene 4 protein - Homo sapiens (Hu					
Q96Q45	2	3.5329	R.RPSEGNEPSTK.E	2	AL2S4_HUMAN
AN1-type zinc finger protein 6 - Homo sapiens (Human)					
Q6FIF0	3	6.3614	K.AVPETEDVQASVSDTAQQPSEEQSK.S	2	ZFAN6_HUMAN
Anamorsin - Homo sapiens (Human)					
Q6FI81	2	2.8803	K.SREQMSSQPK.S	1	CPIN1_HUMAN
Anaphase-promoting complex subunit CDC26 - Homo sapiens (Human)					
Q8NHZ8	3	5.0574	K.QKEDVEVVGGSDGEGAIGLSSDPK.S	2	CDC26_HUMAN
Q8NHZ8	2	3.4339	K.SREQMINDR.I	4	CDC26_HUMAN
Androgen-induced proliferation inhibitor - Homo sapiens (Human)					
Q9NTI5	2	2.8528	R.KKTPVTEQEEK.L	1	APRIN_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Angiomotin-like protein 2 - Homo sapiens (Human)					
Q9Y2J4	2	2.8668	R.RAELLEQALGNAQGR.A	1	AMOL2_HUMAN
Q9Y2J4	2	2.7205	R.RQDEALR.E	1	AMOL2_HUMAN
Angiopoietin-2 precursor - Homo sapiens (Human)					
O15123	2	3.6009	R.DAPLEYDDSVQR.L	3	ANGP2_HUMAN
ANK repeat and LEM domain-containing protein KIAA0692 - Homo sapiens (Human)					
Q86XL3	2	3.6494	R.RTESEMSAR.I	3	K0692_HUMAN
Ankycorbin - Homo sapiens (Human)					
Q9P0K7	2	3.3075	K.RLESSEAER.K	2	RAI14_HUMAN
Ankyrin repeat and SAM domain-containing protein 1A - Homo sapiens (Human)					
Q92625	3	3.7589	K.ASMQLEETGVHAPGASQPSALDQSK.R	1	ANKS1_HUMAN
Q92625	2	3.0853	R.VGYLTGLPTTNSR.S	1	ANKS1_HUMAN
Ankyrin repeat and SAM domain-containing protein 4B - Homo sapiens (Human)					
Q8N8V4	2	3.1229	K.LPSELLQR.Q	1	ANS4B_HUMAN
Ankyrin repeat and SAM domain-containing protein 6 - Homo sapiens (Human)					
Q68DC2	3	4.1712	K.GHTAESSVSSSSSHR.Q	1	ANKS6_HUMAN
Ankyrin repeat domain-containing protein 47 - Homo sapiens (Human)					
Q6NY19	3	3.7168	K.AAQTESPAEAPSLTQESSPGSMDGDRAVAP	1	ANR47_HUMAN
Q6NY19	2	4.7029	K.FALNQNLPDLGGPR.L	2	ANR47_HUMAN
Q6NY19	2	3.8078	R.EAAEEAAAGAR.A	2	ANR47_HUMAN
Q6NY19	2	2.7639	R.LRELEDQAR.T	1	ANR47_HUMAN
Q6NY19	2	3.0298	R.RAPGPPTSR.R	2	ANR47_HUMAN
Q6NY19	3	3.8922	R.RLELAQTHER.A	4	ANR47_HUMAN
Q6NY19	2	3.2123	R.SSPAPNLAPASPGPAQLQLVR.E	1	ANR47_HUMAN
Ankyrin repeat domain-containing protein 50 - Homo sapiens (Human)					
Q9ULJ7	2	2.7095	K.IMIPSAQQEIGR.S	1	ANR50_HUMAN
Ankyrin repeat domain-containing protein 57 - Homo sapiens (Human)					
Q53LP3	2	5.092	R.IQVTAEPEAPDGPAGPEAR.D	2	ANR57_HUMAN
Ankyrin-1 - Homo sapiens (Human)					
P16157	2	3.2084	R.QQGQEEQVQEAK.N	6	ANK1_HUMAN
P16157	3	5.0855	R.RQQGQEEQVQEAK.N	4	ANK1_HUMAN
P16157	3	4.0196	R.QDDATGAGQDSENEVSLVSGHQR.G	1	ANK1_HUMAN
P16157	2	3.2605	R.DSGEGDTTSLR.L	2	ANK1_HUMAN
P16157	3	8.0517	K.RQDDATGAGQDSENEVSLVSGHQR.G	2	ANK1_HUMAN
P16157	2	3.272	K.HSKDHTSTPNP.-	1	ANK1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P16157	3	4.6544	R.ITHSPTVSQVTER.S	4	ANK1_HUMAN
P16157	2	3.887	R.TAAVLLQNDPNPDVLSK.T	1	ANK1_HUMAN
Annexin A1 - Homo sapiens (Human)					
P04083	2	5.3578	K.GDRSEDFGVNEDLADSDAR.A	3	ANXA1_HUMAN
P04083	2	2.9981	K.GTDVNVFNTILTTR.S	2	ANXA1_HUMAN
P04083	2	2.7629	R.ALYEAGER.R	1	ANXA1_HUMAN
P04083	3	4.0465	R.RKGTDVNVFNTILTTR.S	1	ANXA1_HUMAN
P04083	2	4.1172	R.SEDFGVNEDLADSDAR.A	1	ANXA1_HUMAN
Annexin A2 - Homo sapiens (Human)					
P07355	3	4.8349	K.TDLEKDIISDTSGDFRK.L	1	ANXA2_HUMAN
Annexin A5 - Homo sapiens (Human)					
P08758	2	3.4252	R.GTVTDFPGFDER.A	3	ANXA5_HUMAN
P08758	3	4.4954	K.WGTDEEKFITIFGTR.S	2	ANXA5_HUMAN
P08758	2	3.6604	K.VLTEIIASR.T	4	ANXA5_HUMAN
P08758	2	3.5879	K.NFATSLYSMIK.G	3	ANXA5_HUMAN
P08758	2	3.1335	K.NFATSLYSM#IK.G	2	ANXA5_HUMAN
P08758	2	5.1708	K.GLGTDEESILTLLTSR.S	3	ANXA5_HUMAN
P08758	2	2.7915	K.HALKGAGTNEK.V	2	ANXA5_HUMAN
Antigen KI-67 - Homo sapiens (Human)					
P46013	2	2.7368	K.ESADGLQGETQLLVSR.K	1	KI67_HUMAN
AP-1 complex subunit mu-2 - Homo sapiens (Human)					
Q9Y6Q5	1	2.2171	K.NKSVELEDVK.F	1	AP1M2_HUMAN
AP1 subunit gamma-binding protein 1 - Homo sapiens (Human)					
Q9UMZ2	3	4.1696	R.KLSPFVLSAGSGSPSATSILQK.K	1	SYNG_HUMAN
Q9UMZ2	3	3.8773	K.VTTFVSEDALPETTFPALASFK.D	1	SYNG_HUMAN
Q9UMZ2	2	3.4389	K.LSPFVLSAGSGSPSATSILQK.K	1	SYNG_HUMAN
Q9UMZ2	3	5.3689	R.ELEQTAENKPLGESFAEFR.S	3	SYNG_HUMAN
Q9UMZ2	2	4.0369	R.ETLGQIWALANR.T	3	SYNG_HUMAN
AP2-associated protein kinase 1 - Homo sapiens (Human)					
Q2M2I8	2	3.9259	K.AGQTQPNPGILPIQPALTPR.K	3	AAK1_HUMAN
Q2M2I8	2	4.6192	K.LTAEELLNKDFAK.L	2	AAK1_HUMAN
Q2M2I8	2	2.8103	K.SATTTPSGSPR.T	1	AAK1_HUMAN
Q2M2I8	2	4.9574	K.STQLLQAAAAEASLNK.S	2	AAK1_HUMAN
Q2M2I8	2	5.4892	R.ILSDVTHSAVFGVPASK.S	3	AAK1_HUMAN
Q2M2I8	2	4.5825	R.LTDPIPTTETSIAPR.Q	4	AAK1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
AP-3 complex subunit delta-1 - Homo sapiens (Human)					
O14617	3	4.4879	K.HRPSEADEEELAR.R	1	AP3D1_HUMAN
O14617	3	4.6702	R.ALDIDLDKPLADSEKLPIQK.H	2	AP3D1_HUMAN
AP-3 complex subunit sigma-1 - Homo sapiens (Human)					
Q92572	2	2.7682	K.NMNLPEIPR.N	1	AP3S1_HUMAN
Apolipoprotein A-I precursor - Homo sapiens (Human)					
P02647	2	3.3242	K.VEPLRAELQEGAR.Q	4	APOA1_HUMAN
P02647	3	4.4584	R.VKDLATVYVDVLKDSGR.D	2	APOA1_HUMAN
P02647	2	4.7667	R.VKDLATVYVDVLK.D	7	APOA1_HUMAN
P02647	3	4.3498	R.LAARLEALKENGGAR.L	3	APOA1_HUMAN
P02647	2	4.1289	R.DYVSQFEGSALGK.Q	4	APOA1_HUMAN
P02647	3	4.193	R.AHVDALRTHLAPYSDELRQR.L	1	APOA1_HUMAN
P02647	2	2.8725	R.AELQEGAR.Q	1	APOA1_HUMAN
P02647	2	3.4639	K.WQEEMELYR.Q	2	APOA1_HUMAN
P02647	2	3.3678	K.WQEEM#ELYR.Q	3	APOA1_HUMAN
P02647	2	3.5603	K.DLATVYVDVLK.D	8	APOA1_HUMAN
P02647	3	4.2074	K.QLNLKLLDNWDSVTSTFSK.L	1	APOA1_HUMAN
P02647	4	5.7437	K.LREQLGPVTQEFWDNLEKETEGLR.Q	1	APOA1_HUMAN
P02647	2	3.4792	K.LLDNWDSVTSTFSK.L	2	APOA1_HUMAN
P02647	3	5.3734	K.DSGRDYVSQFEGSALGK.Q	11	APOA1_HUMAN
P02647	3	4.7	K.DLATVYVDVLKDSGRDYVSQFEGSALGK.Q	4	APOA1_HUMAN
P02647	2	2.7523	K.VQPYLDDFQKK.W	1	APOA1_HUMAN
P02647	2	5.1729	K.VSFLSALEEYTK.K	8	APOA1_HUMAN
Apolipoprotein A-II precursor - Homo sapiens (Human)					
P02652	2	4.3009	K.VKSPELQAEAK.S	4	APOA2_HUMAN
P02652	2	3.6595	K.SPELQAEAK.S	5	APOA2_HUMAN
P02652	2	3.4119	K.SKEQLTPLIKK.A	3	APOA2_HUMAN
P02652	3	4.6826	K.AGTELVNFLSYFVELGTQPATQ.-	12	APOA2_HUMAN
P02652	2	3.5988	K.SKEQLTPLIK.K	4	APOA2_HUMAN
Apolipoprotein A-IV precursor - Homo sapiens (Human)					
P06727	3	3.9303	K.KLVPFATELHER.L	1	APOA4_HUMAN
P06727	2	2.8492	R.ISASAEELRQR.L	1	APOA4_HUMAN
P06727	2	2.7584	R.ISASAEELR.Q	1	APOA4_HUMAN
P06727	2	3.7545	R.RVEPYGENFNK.A	2	APOA4_HUMAN
P06727	2	3.9623	K.ALVQQMEQLR.Q	6	APOA4_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P06727	3	3.9041	K.AKIDQNVEELKGR.L	1	APOA4_HUMAN
P06727	3	4.2681	K.SLAELGGHLDQQVEEFRR.R	2	APOA4_HUMAN
P06727	2	3.1313	R.DKVNSFFSTFK.E	2	APOA4_HUMAN
Apolipoprotein B-100 precursor - Homo sapiens (Human)					
P04114	3	5.7359	R.VPSYTLILPSLELPVLHVPR.N	2	APOB_HUMAN
Apolipoprotein B48 receptor - Homo sapiens (Human)					
Q0VD83	3	3.9124	R.EADAGETEEPGAEGAGKGEEVVVVEK.A	1	Q0VD83_HUMA
Q0VD83	2	3.1924	R.ETEDEEAEADR.T	3	Q0VD83_HUMA
Q0VD83	3	4.1913	R.HAGSVKPEASEAFPGAWENR.T	2	Q0VD83_HUMA
Q0VD83	2	4.4229	R.TEEAAESQTAGR.E	7	Q0VD83_HUMA
Apolipoprotein C-II precursor - Homo sapiens (Human)					
P02655	2	2.9032	K.TAAQNLYEK.T	2	APOC2_HUMAN
Apolipoprotein C-III precursor - Homo sapiens (Human)					
P02656	2	3.0207	R.GWVTDGFSSLK.D	2	APOC3_HUMAN
P02656	2	5.8927	K.DALSSVQESQVAQQAR.G	16	APOC3_HUMAN
P02656	2	5.5543	K.TAKDALSSVQESQVAQQAR.G	3	APOC3_HUMAN
Apolipoprotein E precursor - Homo sapiens (Human)					
P02649	2	3.7963	R.AATVGSLAGQPLQER.A	2	APOE_HUMAN
P02649	1	2.1488	R.DADDLQK.R	1	APOE_HUMAN
P02649	2	2.862	R.LAVYQAGAR.E	1	APOE_HUMAN
Apolipoprotein O precursor - Homo sapiens (Human)					
Q9BUR5	2	2.7493	R.SQLEESISQLR.H	1	APOO_HUMAN
Apolipoprotein O-like precursor - Homo sapiens (Human)					
Q6UXV4	2	2.8463	K.HSVPLPTELSSEAK.T	1	APOOL_HUMAN
Apoptosis inhibitor 5 - Homo sapiens (Human)					
Q9BZZ5	2	3.2765	R.ASEDTTSGSPPKK.S	2	API5_HUMAN
Q9BZZ5	2	4.1777	R.ASEDTTSGSPPK.K	3	API5_HUMAN
Q9BZZ5	2	3.313	K.RASEDTTSGSPPK.K	2	API5_HUMAN
Q9BZZ5	3	4.838	K.RASEDTTSGSPPKK.S	6	API5_HUMAN
Apoptosis-associated speck-like protein containing a CARD - Homo sapiens (Human)					
Q9ULZ3	3	5.9553	R.ARDAILDALENLTAEELKK.F	1	ASC_HUMAN
Q9ULZ3	2	4.9201	R.DAILDALENLTAEELKK.F	2	ASC_HUMAN
Apoptosis-inducing factor 3 - Homo sapiens (Human)					
Q96NN9	2	2.7461	K.DLENGQMREVELGWGK.V	1	AIFM3_HUMAN
Apoptosis-related protein 3 precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6UW56	2	2.7912	K.VAFYC*KTTRELM#LHAR.C	1	APR3_HUMAN
Apoptosis-stimulating of p53 protein 2 - Homo sapiens (Human)					
Q13625	2	2.7851	R.QQQQVAEQEK.L	1	ASPP2_HUMAN
Q13625	2	4.5275	K.NFQQAVQSALTK.T	3	ASPP2_HUMAN
Apoptotic chromatin condensation inducer in the nucleus - Homo sapiens (Human)					
Q9UKV3	3	5.7988	R.NLKTEEEEEEEEEEEEDDEEEEGDDEGQK.S	1	ACINU_HUMAN
Q9UKV3	3	7.6193	R.NLKTEEEEEEEEEEEEDDEEEEGDDEGQKS	2	ACINU_HUMAN
Q9UKV3	2	3.2655	R.RLSQPESAEK.H	4	ACINU_HUMAN
Q9UKV3	2	3.5964	R.SAQPLPLKIEELALAK.G	1	ACINU_HUMAN
Q9UKV3	2	3.7244	R.SQEQEVLER.G	3	ACINU_HUMAN
Q9UKV3	2	3.4244	R.TSTSSSSVQAR.R	5	ACINU_HUMAN
Q9UKV3	2	2.8915	R.KISVVSATK.G	1	ACINU_HUMAN
Q9UKV3	2	3.0837	K.TTSPLEEEEREIK.S	2	ACINU_HUMAN
Q9UKV3	1	3.5398	R.WGASTATTQK.K	4	ACINU_HUMAN
Q9UKV3	3	6.6785	K.HSTPHAAFQPNSQIGEEM#SQNSFIK.Q	1	ACINU_HUMAN
Q9UKV3	3	5.6168	K.STLADYSAQKDLEPESDR.S	6	ACINU_HUMAN
Q9UKV3	2	3.5968	K.GALMLENLQK.H	5	ACINU_HUMAN
Q9UKV3	4	5.6488	K.HSTPHAAFQPNSQIGEEMSQNSFIK.Q	3	ACINU_HUMAN
Q9UKV3	2	3.2493	K.KPSISITTESLK.S	2	ACINU_HUMAN
Q9UKV3	2	2.961	K.LLDDLFR.K	2	ACINU_HUMAN
Q9UKV3	2	4.2279	K.LSEGSQPAEEEEDQETPSR.N	8	ACINU_HUMAN
Q9UKV3	2	4.3244	R.LKGALMLENLQK.H	3	ACINU_HUMAN
Q9UKV3	3	3.9447	K.SSSISEEKGDSDDEKPR.K	1	ACINU_HUMAN
Q9UKV3	3	5.1636	K.AESPAEKVPEESVLPLVQK.S	1	ACINU_HUMAN
Q9UKV3	2	2.7092	K.EFKEEGEEIPR.V	1	ACINU_HUMAN
Arachidonate 15-lipoxygenase type II - Homo sapiens (Human)					
O15296	2	2.9726	K.QLNYSLLCLPEDIR.T	1	LX15B_HUMAN
Archaemetzincin-2 - Homo sapiens (Human)					
Q86W34	2	2.8337	K.EWIIKCLAVLQK.-	1	AMZ2_HUMAN
ARF GTPase-activating protein GIT2 - Homo sapiens (Human)					
Q14161	3	4.0797	K.QATTNVYQVQTGSEYTDTSNHSSLK.R	1	GIT2_HUMAN
Arfaptin-1 - Homo sapiens (Human)					
P53367	3	5.3015	K.HSLPSGLGLSETQITSHGFDNTK.E	2	ARFP1_HUMAN
P53367	2	4.4151	K.SGPVILADEIKNPAMEK.L	1	ARFP1_HUMAN
P53367	2	2.737	R.LAQQGSDLIVPAGGQR.T	1	ARFP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Arginine/serine-rich coiled coil protein 1 - Homo sapiens (Human)					
Q96IZ7	3	4.7659	R.VKEIEAIESDSFVQQTFR.S	1	RSRC1_HUMAN
ARHGAP5 protein - Homo sapiens (Human)					
Q05BE8	3	5.3409	R.VPLAHPEDMDPSDNYAEPIDTIFK.Q	2	Q05BE8_HUMA
Q05BE8	2	3.374	K.TIEAGIGKNPR.K	2	Q05BE8_HUMA
Armadillo repeat protein deleted in velo-cardio-facial syndrome - Homo sapiens (Human)					
O00192	2	3.7283	R.FQAEPYGLEDDTR.S	3	ARVC_HUMAN
O00192	2	2.778	R.SLAADDEGGPELEPDYGTATR.R	1	ARVC_HUMAN
Armadillo repeat-containing X-linked protein 1 - Homo sapiens (Human)					
Q9P291	2	3.7995	K.AHSGSHSGGGLEAK.A	4	ARMX1_HUMAN
Ashwin - Homo sapiens (Human)					
Q9BVC5	4	4.7943	R.KLSNSSSSVSPLILSSNLPVNNKTEHNNNDAK	1	ASHWN_HUMAN
Q9BVC5	3	4.0741	R.KRPLIVFDGSSTSTSIK.V	1	ASHWN_HUMAN
Q9BVC5	3	4.9271	R.VNKDSLTDLYVQHAIPLPQR.D	2	ASHWN_HUMAN
Asparaginase-like protein 1 - Homo sapiens (Human)					
Q7L266	2	5.0722	K.GNVAYATSTGGIVNK.M	2	Q7L266_HUMAN
Q7L266	2	3.6327	R.AATVGYGILR.E	7	Q7L266_HUMAN
Aspartate aminotransferase, cytoplasmic - Homo sapiens (Human)					
P17174	2	3.3812	K.NLDYVATSIHEAVTK.I	1	AATC_HUMAN
Aspartate aminotransferase, mitochondrial precursor - Homo sapiens (Human)					
P00505	2	3.082	R.KAEAQIAAK.N	2	AATM_HUMAN
AT hook DNA-binding motif-containing protein 1 - Homo sapiens (Human)					
Q5TGY3	2	3.7895	K.VGTGLLADFLGR.T	1	Q5TGY3_HUMA
Q5TGY3	3	4.6063	R.LSRPPAPPPGDLQYSFFSSPSLANSIR.S	1	Q5TGY3_HUMA
Ataxin-2 - Homo sapiens (Human)					
Q99700	3	6.2635	R.KPGGSGLLASPAAAPSPSSSSVSSSSATAPS	1	ATX2_HUMAN
Ataxin-2-like protein - Homo sapiens (Human)					
Q8WWM7	2	2.8819	K.ISLAPTDVK.E	1	ATX2L_HUMAN
Q8WWM7	2	4.1046	R.GPHHLDNSSPGPGSEAR.G	1	ATX2L_HUMAN
Q8WWM7	3	4.4377	K.GEDKDEGPVAEQVKK.S	1	ATX2L_HUMAN
Q8WWM7	2	2.9629	K.FNEENYGVK.T	2	ATX2L_HUMAN
Q8WWM7	2	3.7036	K.EVDGLLTSEPMGSPVSSK.T	2	ATX2L_HUMAN
Q8WWM7	2	3.7182	K.EKEVDGLLTSEPMGSPVSSK.T	2	ATX2L_HUMAN
Q8WWM7	2	3.6949	K.EFNPTKPLLSVNK.S	3	ATX2L_HUMAN
Q8WWM7	2	2.704	K.STSTPTSPGPR.T	1	ATX2L_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Atherin - Homo sapiens (Human)					
Q6SPF0	2	2.9726	R.RAPPPAVAAR.E	2	SAMD1_HUMAN
AT-hook-containing transcription factor 1 - Homo sapiens (Human)					
Q8WYP5	2	3.5772	R.NLSFNELYPSGTLK.L	1	AHTF1_HUMAN
Q8WYP5	2	3.44	K.SAQQEASADVATPK.M	2	AHTF1_HUMAN
Q8WYP5	3	3.8556	K.SKVPVLDEGLTSVETYTPAIR.A	1	AHTF1_HUMAN
Q8WYP5	2	3.2323	R.GLSQNQQIPQNSVTPR.R	1	AHTF1_HUMAN
Q8WYP5	2	3.4186	R.KINPSEDVGSK.A	2	AHTF1_HUMAN
ATP synthase D chain, mitochondrial - Homo sapiens (Human)					
O75947	2	3.7704	R.IVEYEKEMEK.M	4	ATP5H_HUMAN
O75947	2	3.4584	R.IVEYEKEM#EK.M	2	ATP5H_HUMAN
O75947	2	4.1991	K.YPYWPHQPIENL.-	5	ATP5H_HUMAN
O75947	2	2.9157	K.KYPYWPHQPIENL.-	1	ATP5H_HUMAN
O75947	3	4.7991	K.VPVPEDKYTAQVDAEEKEDVK.S	1	ATP5H_HUMAN
ATP synthase delta chain, mitochondrial precursor - Homo sapiens (Human)					
P30049	2	5.1591	K.AQAELVGTADEATR.A	5	ATPD_HUMAN
P30049	3	3.9954	K.AQAELVGTADEATRAEIQIR.I	4	ATPD_HUMAN
ATP synthase e chain, mitochondrial - Homo sapiens (Human)					
P56385	2	2.7827	R.YNYLKPR.A	1	ATP5I_HUMAN
P56385	1	2.1219	R.ELAEDDSILK.-	1	ATP5I_HUMAN
ATP synthase subunit alpha, mitochondrial precursor - Homo sapiens (Human)					
P25705	2	4.4608	R.VVDALGNAIDGKGPIGSK.T	2	ATPA_HUMAN
P25705	2	2.8568	R.NVQAEEMVEFSSGLK.G	1	ATPA_HUMAN
P25705	2	3.8391	K.TGTAEM#SSILEER.I	3	ATPA_HUMAN
P25705	2	5.2474	R.ILGADTSVDLEETGR.V	2	ATPA_HUMAN
ATP synthase subunit beta, mitochondrial precursor - Homo sapiens (Human)					
P06576	2	2.8935	K.ADKLAEEHSS.-	1	ATPB_HUMAN
P06576	3	4.2714	K.VLDSGAPIKIPVGPETLGR.I	2	ATPB_HUMAN
ATP synthase-coupling factor 6, mitochondrial precursor - Homo sapiens (Human)					
P18859	3	3.9544	K.LKQMFGNADM#NTFPTFK.F	1	ATP5J_HUMAN
P18859	2	5.8939	R.QTSGGPVDASSEYQQELER.E	3	ATP5J_HUMAN
P18859	3	5.3403	K.RQTSGGPVDASSEYQQELERELFK.L	1	ATP5J_HUMAN
P18859	3	4.5872	K.RQTSGGPVDASSEYQQELER.E	2	ATP5J_HUMAN
P18859	2	3.6703	K.QMFGNADMNTFPTFK.F	11	ATP5J_HUMAN
P18859	2	3.3408	K.QM#FGNADMNTFPTFK.F	2	ATP5J_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P18859	3	4.0925	K.LKQMFGNADMNTFPTFK.F	3	ATP5J_HUMAN
P18859	3	4.253	K.LKQM#FGNADMNTFPTFK.F	1	ATP5J_HUMAN
P18859	3	5.1983	K.LKQM#FGNADM#NTFPTFK.F	1	ATP5J_HUMAN
P18859	1	2.3024	K.LFVDKIR.E	2	ATP5J_HUMAN
P18859	2	4.6106	K.FEDPKFEVIEKPQA.-	7	ATP5J_HUMAN
P18859	2	2.8034	K.QM#FGNADM#NTFPTFK.F	1	ATP5J_HUMAN
ATP-binding cassette sub-family F member 3 - Homo sapiens (Human)					
Q9NUQ8	2	2.7127	R.EYEAQQQYR.Q	1	ABCF3_HUMAN
Q9NUQ8	3	6.1564	R.LAEPQSQGNSQVLLDAPIQLSK.I	5	ABCF3_HUMAN
ATP-binding cassette, sub-family F - Homo sapiens (Human)					
Q5STZ8	3	5.3512	K.VLKEKEQQQQQQQQQQK.K	2	Q5STZ8_HUMA
ATP-dependent DNA helicase 2 subunit 1 - Homo sapiens (Human)					
P12956	3	5.2478	K.SGLKKQELLEALTK.H	2	KU70_HUMAN
P12956	3	4.1842	K.KQELLEALTK.H	5	KU70_HUMAN
ATP-dependent DNA helicase Q1 - Homo sapiens (Human)					
P46063	2	4.1069	K.AANMLQQSGSK.N	1	RECQ1_HUMAN
P46063	2	4.1175	K.KAANMLQQSGSK.N	2	RECQ1_HUMAN
ATP-dependent RNA helicase A - Homo sapiens (Human)					
Q08211	2	3.2038	K.AENNSEVGASGYGVPGPTWDR.G	2	DHX9_HUMAN
ATP-dependent RNA helicase DDX19B - Homo sapiens (Human)					
Q9UMR2	3	4.5623	K.TNANAEKTDEEEKEDR.A	4	DD19B_HUMAN
ATP-dependent RNA helicase DDX42 - Homo sapiens (Human)					
Q86XP3	3	5.6802	K.GIPGFGNTGNISGAPVTYPSAGAQGVNNTAS	1	DDX42_HUMAN
Q86XP3	3	4.3563	K.LPQQSHSAFGATSSSSGFGK.S	3	DDX42_HUMAN
Q86XP3	2	2.9243	K.SHFVAASLSNQK.A	1	DDX42_HUMAN
Q86XP3	2	4.4563	K.TDKTADGFAVPEPPK.R	4	DDX42_HUMAN
Q86XP3	3	3.7509	K.TDKTADGFAVPEPPKR.K	1	DDX42_HUMAN
AT-rich interactive domain-containing protein 1A - Homo sapiens (Human)					
O14497	2	3.5179	K.AAAGQESEGPAVGPPQPLGK.E	1	ARI1A_HUMAN
O14497	3	5.8999	R.GGTPGSGAAAAAGSKPPPSSSASASSSSSS	1	ARI1A_HUMAN
AT-rich interactive domain-containing protein 1B - Homo sapiens (Human)					
Q8NFD5	2	4.3983	K.DMGAQYAAASPAWAAAQQR.S	2	ARI1B_HUMAN
Q8NFD5	3	4.0786	K.DSAAGGQADPPGPPLLSKPGDEDDAPPK.M	3	ARI1B_HUMAN
Q8NFD5	2	3.5245	K.DM#GAQYAAASPAWAAAQQR.S	2	ARI1B_HUMAN
Attractin precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O75882	2	4.1074	R.LTGSSGFVTDGPGNYK.Y	2	ATRN_HUMAN
ATXN3 protein - Homo sapiens (Human)					
Q4VBR4	3	4.7839	R.VLEANDGSGMLDEDEEDLQR.A	1	Q4VBR4_HUMA
Autogenous vein graft remodeling associated protein 2 - Homo sapiens (Human)					
Q2HYQ4	2	2.7244	K.KQSQHTSTVNVLQK.F	1	Q2HYQ4_HUMA
Autophagy-related protein 3 - Homo sapiens (Human)					
Q9NT62	2	4.3229	K.ALEVAEYLTPVLK.E	4	ATG3_HUMAN
Axin-1 - Homo sapiens (Human)					
O15169	2	3.1086	K.SASTEVPGASEDAEKNQK.I	1	AXN1_HUMAN
BAG family molecular chaperone regulator 3 - Homo sapiens (Human)					
O95817	2	2.956	R.SSLGSHQLPR.G	4	BAG3_HUMAN
O95817	2	2.854	K.YLMIEEYLTK.E	2	BAG3_HUMAN
O95817	2	3.7073	R.VHTVVDRPQQPMTHR.E	3	BAG3_HUMAN
O95817	2	3.7282	R.SQSPAASDCSSSSSSASLPSSGR.S	1	BAG3_HUMAN
O95817	3	3.7267	R.PAAQPSFHQAQK.T	1	BAG3_HUMAN
O95817	2	4.8734	R.KVQTILEKLEQK.A	5	BAG3_HUMAN
O95817	2	2.8926	R.KVQTILEK.L	1	BAG3_HUMAN
O95817	3	7.1929	R.GMPETTQPDKQCGQVAAAAAAQPPASHGPE	2	BAG3_HUMAN
O95817	2	3.8979	R.FRTEAAAAAPQR.S	5	BAG3_HUMAN
O95817	3	3.7805	K.EVDSKPVSQKPPPPSEK.V	1	BAG3_HUMAN
O95817	2	4.1626	K.VQTILEKLEQK.A	4	BAG3_HUMAN
O95817	2	4.1898	K.ELLALDSVDPEGR.A	6	BAG3_HUMAN
O95817	3	4.3179	K.KNAGNAEDPHTETQQPEATAAATSNPSSMT	1	BAG3_HUMAN
O95817	3	5.1	K.NAGNAEDPHTETQQPEATAAATSNPSSM#TD	1	BAG3_HUMAN
O95817	1	2.4387	K.VEAILEK.V	2	BAG3_HUMAN
O95817	3	4.6328	K.VEAILEKVQGLEQAVDNFEGK.K	1	BAG3_HUMAN
O95817	3	6.4087	K.VEAILEKVQGLEQAVDNFEGKK.T	4	BAG3_HUMAN
O95817	2	5.0659	K.VQGLEQAVDNFEGKK.T	7	BAG3_HUMAN
O95817	3	4.8637	R.VPSEGPKETPSSANGPSR.E	3	BAG3_HUMAN
BAG family molecular chaperone regulator 4 - Homo sapiens (Human)					
O95429	2	4.5469	K.DSSYPYSQSDQSMNR.H	3	BAG4_HUMAN
O95429	2	2.918	R.SSGNSPTPVSR.W	1	BAG4_HUMAN
Band 4.1-like protein 1 - Homo sapiens (Human)					
Q9H4G0	2	3.8655	K.HQASINELKR.T	5	E41L1_HUMAN
Q9H4G0	2	3.5302	R.KKIEPEAVLQTR.V	3	E41L1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9H4G0	2	4.2703	R.VSAMDNTQQVDGSASVGR.E	5	E41L1_HUMAN
Q9H4G0	2	4.3161	R.VSAM#DNTQQVDGSASVGR.E	3	E41L1_HUMAN
Q9H4G0	3	7.0706	R.SLSPIIGKDVLTSTYGATAETLSTSTTTHVTK.T	1	E41L1_HUMAN
Q9H4G0	2	4.2896	K.KIEPEAVLQTR.V	5	E41L1_HUMAN
Q9H4G0	2	2.8268	K.GAPSQDDESGGIEDSPDR.G	1	E41L1_HUMAN
Q9H4G0	2	4.7041	K.GAAAM#IPGPQTVATEIR.S	4	E41L1_HUMAN
Q9H4G0	2	3.9838	R.DLNKGAPSQDDESGGIEDSPDR.G	1	E41L1_HUMAN
Band 4.1-like protein 3 - Homo sapiens (Human)					
Q9Y2J2	2	3.4001	K.VVQETVLVEER.R	1	E41L3_HUMAN
Q9Y2J2	2	4.5506	K.EGSALTEGAKEEGGEEVAK.A	4	E41L3_HUMAN
Q9Y2J2	2	4.8002	R.TFLETSTDTAVTNEWEKR.L	2	E41L3_HUMAN
Q9Y2J2	3	7.2128	R.RVVHASGDASYSAGDSGDAAAQPAFTGIK.G	3	E41L3_HUMAN
Q9Y2J2	3	5.4494	R.VVHASGDASYSAGDSGDAAAQPAFTGIK.G	3	E41L3_HUMAN
Q9Y2J2	2	4.7349	R.KGEEVTPISAIRHEGK.S	1	E41L3_HUMAN
Q9Y2J2	2	3.6864	R.KGEEVTPISAIR.H	2	E41L3_HUMAN
Q9Y2J2	3	3.72	R.IVITGDADIDHDQALAQAIKEAK.E	1	E41L3_HUMAN
Q9Y2J2	2	3.5439	K.VVVHKETEITPEDGED.-	1	E41L3_HUMAN
Q9Y2J2	2	4.0843	K.TESSGIETEPTVHHLPLSTEK.V	1	E41L3_HUMAN
Q9Y2J2	2	3.1333	K.LMDGSEIFSLLESAR.K	1	E41L3_HUMAN
Q9Y2J2	2	3.9476	K.HQTNISELKR.T	3	E41L3_HUMAN
Q9Y2J2	2	6.0492	K.GKEGSALTEGAKEEGGEEVAK.A	5	E41L3_HUMAN
Q9Y2J2	2	4.0768	K.GISQTNLITTVTPEK.K	1	E41L3_HUMAN
Q9Y2J2	2	3.9421	K.AVLEQEETAAASR.E	2	E41L3_HUMAN
Q9Y2J2	3	4.7943	K.TITYESSQVDPGTDLEPGVLMSAQTITSETTS	1	E41L3_HUMAN
Q9Y2J2	2	4.695	K.GISQTNLITTVTPEKK.A	2	E41L3_HUMAN
Barrier-to-autointegration factor - Homo sapiens (Human)					
O75531	2	3.8438	K.AYVVLGQFLVLK.K	2	BAF_HUMAN
O75531	3	5.6158	R.DFVAEPMGEKPVGSLAGIGEVLGK.K	1	BAF_HUMAN
O75531	3	4.4938	R.DFVAEPM#GEKPVGSLAGIGEVLGK.K	1	BAF_HUMAN
O75531	3	6.6133	K.HRDFVAEPMGEKPVGSLAGIGEVLGK.K	3	BAF_HUMAN
Basement membrane-specific heparan sulfate proteoglycan core protein precursor - Homo sapiens					
P98160	1	2.7801	R.VTVTSEGGR.G	2	PGBM_HUMAN
P98160	3	5.0079	R.SPVISIDPPSSTVQQGQDASFK.C	1	PGBM_HUMAN
P98160	3	5.0038	R.LRSPVISIDPPSSTVQQGQDASFK.C	2	PGBM_HUMAN
P98160	2	4.3795	R.LPAVEPTDQAQYLCR.A	2	PGBM_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P98160	2	3.0021	R.LGTVPQFPR.A	1	PGBM_HUMAN
P98160	3	4.2778	R.GM#LEPVQRPDVVLVGAGYR.L	1	PGBM_HUMAN
P98160	2	2.8751	R.GHTPTQPGALNQR.Q	1	PGBM_HUMAN
P98160	2	2.876	R.AQAGANTRPCPS.-	2	PGBM_HUMAN
P98160	2	3.2148	R.AM#DFNGILTIR.N	2	PGBM_HUMAN
P98160	2	3.3497	K.SPAYTLVWTR.L	1	PGBM_HUMAN
P98160	2	3.1641	R.YELGSGLAVLR.S	1	PGBM_HUMAN
P98160	2	2.9884	R.LEGDTLIIPR.V	1	PGBM_HUMAN
Basigin precursor - Homo sapiens (Human)					
P35613	2	4.7616	K.GGVVLKEDALPGQK.T	4	BASI_HUMAN
P35613	2	3.1863	K.SESVPPVTDWAWYK.I	1	BASI_HUMAN
P35613	2	3.4861	R.FFVSSSQGR.S	5	BASI_HUMAN
P35613	3	5.0806	R.RKPEDVLDDDDAGSAPLK.S	1	BASI_HUMAN
P35613	2	4.4375	R.SELHIENLNMEADPGQYR.C	4	BASI_HUMAN
B-cell CLL/lymphoma 7 protein family member A - Homo sapiens (Human)					
Q4VC05	2	4.3369	K.MKLEASQQNSEEM#.-	2	BCL7A_HUMAN
Q4VC05	2	4.1971	K.MKLEASQQNSEEM.-	2	BCL7A_HUMAN
Q4VC05	2	3.2975	K.M#KLEASQQNSEEM.-	1	BCL7A_HUMAN
Q4VC05	4	4.7794	K.VDRQPSGDSGLAAETSAISQDLEGVPPSKK.	1	BCL7A_HUMAN
B-cell lymphoma/leukemia 10 - Homo sapiens (Human)					
O95999	3	4.3396	R.ASTVMYHPEGESSTTPFFSTNSSLNLPVLEV	1	BCL10_HUMAN
B-cell receptor-associated protein 29 - Homo sapiens (Human)					
Q9UHQ4	1	2.3953	K.EYDQLLK.E	1	BAP29_HUMAN
B-cell receptor-associated protein 31 - Homo sapiens (Human)					
P51572	2	3.8029	K.LQAAVDGPMDKKEE.-	2	BAP31_HUMAN
Bcl2 antagonist of cell death - Homo sapiens (Human)					
Q92934	2	3.8061	R.GLGPSPAGDGPSGSGK.H	1	BAD_HUMAN
Bcl-2 related ovarian killer - Homo sapiens (Human)					
Q9UL32	2	2.7427	R.AGLSWSAPERASPAPGGR.L	1	Q9UL32_HUMAN
BCL2/adenovirus E1B 19 kDa protein-interacting protein 3 - Homo sapiens (Human)					
Q12983	2	3.2872	K.NSSQSEEDDIER.R	1	BNIP3_HUMAN
Bcl-2-associated transcription factor 1 - Homo sapiens (Human)					
Q9NYF8	2	2.8185	R.FTDEESR.V	1	BCLF1_HUMAN
Q9NYF8	2	2.9627	R.VFKEENQKGDK.K	1	BCLF1_HUMAN
Q9NYF8	2	3.1056	R.SSFYPDGGDQETAK.T	1	BCLF1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9NYF8	2	3.2761	R.NTEEEGLKYK.S	3	BCLF1_HUMAN
Q9NYF8	2	3.0282	R.KSVLADQGK.S	2	BCLF1_HUMAN
Q9NYF8	3	3.8855	R.FTDEESRVFLLDR.G	1	BCLF1_HUMAN
Q9NYF8	2	2.8521	K.RFTDEESR.V	3	BCLF1_HUMAN
Q9NYF8	2	3.8404	K.LKETGYVVERPSTTK.D	1	BCLF1_HUMAN
Q9NYF8	2	4.5275	K.KAEGEPQEESPLK.S	2	BCLF1_HUMAN
Q9NYF8	2	3.0174	K.FNDSEGDDTEETEDYR.Q	1	BCLF1_HUMAN
Q9NYF8	2	3.5311	K.ETGYVVERPSTTK.D	1	BCLF1_HUMAN
Q9NYF8	1	2.3924	K.SVLADQGK.S	2	BCLF1_HUMAN
Q9NYF8	2	2.7628	K.ETQSPEQVKSEK.L	1	BCLF1_HUMAN
Beta-2-glycoprotein 1 precursor - Homo sapiens (Human)					
P02749	3	4.0776	K.ATFGCHDGYSLDGPEEIECTK.L	1	APOH_HUMAN
P02749	2	3.9359	K.TFYEPGEEITYSCKPGYVSR.G	3	APOH_HUMAN
P02749	1	3.6658	K.TDASDVKPC.-	14	APOH_HUMAN
P02749	2	3.1835	K.ATVVYQGER.V	6	APOH_HUMAN
P02749	2	4.0104	K.KATVVYQGER.V	3	APOH_HUMAN
Beta-adducin - Homo sapiens (Human)					
P35612	3	4.0936	K.KLELDGEKETAPEEPGSPAK.S	1	ADDB_HUMAN
P35612	2	5.6585	K.SAGPQSQLLASVIAEK.S	3	ADDB_HUMAN
P35612	1	2.2952	K.SPLVSPSK.S	2	ADDB_HUMAN
Beta-arrestin-1 - Homo sapiens (Human)					
P49407	1	2.1515	R.QFLMSDK.P	1	ARRB1_HUMAN
Beta-catenin-interacting protein 1 - Homo sapiens (Human)					
Q9NSA3	2	3.6396	R.EGAPGKSPEEMYIQQK.V	1	CNBP1_HUMAN
Q9NSA3	2	4.1064	K.MGSNLTASEEEFLR.T	1	CNBP1_HUMAN
Betaine--homocysteine S-methyltransferase 1 - Homo sapiens (Human)					
Q93088	2	4.3241	K.EATTEQQLKELFEK.Q	2	BHMT1_HUMAN
Q93088	2	2.8561	K.GTAELM#QQK.E	1	BHMT1_HUMAN
Q93088	2	3.2774	K.GTAELMQQK.E	1	BHMT1_HUMAN
Q93088	2	3.2578	K.QGFIDLPEFPFGLEPR.V	2	BHMT1_HUMAN
BIN3 domain-containing protein 1 - Homo sapiens (Human)					
Q7L2J0	2	3.7324	R.ESPGAAATSSSGPQAQQHR.G	2	BN3D1_HUMAN
Q7L2J0	3	4.1021	R.RGGGGTELGPPAPPRPR.N	1	BN3D1_HUMAN
Biogenesis of lysosome-related organelles complex-1 subunit 1 - Homo sapiens (Human)					
P78537	2	3.5052	R.TIATALEYVYK.G	2	BL1S1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Biogenesis of lysosome-related organelles complex-1 subunit 3 - Homo sapiens (Human)					
Q6QNY0	2	4.4885	R.VAGEAAETDSEPEPEPEPTAAPR.D	3	BL1S3_HUMAN
Biotinidase precursor - Homo sapiens (Human)					
P43251	2	2.7646	R.TSIYPFLDFMPSPQVVR.W	1	BTD_HUMAN
Bisphosphoglycerate mutase - Homo sapiens (Human)					
P07738	2	4.6293	R.AVGPHQFLGDQEAIQAAIK.K	4	PMGE_HUMAN
BMP-2-inducible protein kinase - Homo sapiens (Human)					
Q9NSY1	2	4.2319	R.VLQQLQQGDWR.L	2	BMP2K_HUMAN
Q9NSY1	2	5.2646	R.SVADKEAIANFTNQK.N	2	BMP2K_HUMAN
Q9NSY1	2	3.8731	K.SESNEDLFGLVPFDEITGSQQQK.V	1	BMP2K_HUMAN
Q9NSY1	3	3.9698	K.NLPQHRFPAAGLEQEEFDVFTK.A	1	BMP2K_HUMAN
Q9NSY1	2	3.0067	K.KTSVQGQVQK.G	1	BMP2K_HUMAN
Q9NSY1	4	4.8099	K.KVNVQECHAVGPEAHTIPGYPK.S	1	BMP2K_HUMAN
BNIP2 motif-containing molecule at the C-terminal region 1 - Homo sapiens (Human)					
Q58A63	3	3.9633	R.AGPDAVTHDNEWEMLSPQPVQK.N	1	BMCC1_HUMAN
BolA-like protein 1 - Homo sapiens (Human)					
Q9Y3E2	2	3.046	R.FEGLSPLQR.H	1	BOLA1_HUMAN
Q9Y3E2	3	5.0264	R.TKLEEALSPEVLELRNESGGHAVPPGSETHF	2	BOLA1_HUMAN
Brain acid soluble protein 1 - Homo sapiens (Human)					
P80723	2	3.0181	K.GYNVNDEKAK.E	2	BASP_HUMAN
P80723	2	4.5729	K.AEGAATEEEGTPK.E	3	BASP_HUMAN
P80723	2	5.0985	K.KTEAPAAPAAQETK.S	1	BASP_HUMAN
P80723	2	4.2787	K.SDGAPASDSKPGSSEAAPSSK.E	7	BASP_HUMAN
P80723	3	5.7449	K.AQGPAASAEEPKPVEAPAANSDQTVTVKE.-	2	BASP_HUMAN
P80723	3	5.9681	K.AEGAATEEEGTPKESEPQAAAEPAEAK.E	2	BASP_HUMAN
P80723	3	4.5679	K.AEPPKAPEQEQAAPGPAAGGEAPK.A	2	BASP_HUMAN
P80723	2	3.3777	K.ETPAATEAPSSTPK.A	5	BASP_HUMAN
Brain-specific polypeptide PEP-19 - Homo sapiens (Human)					
P48539	2	2.7599	R.AAVAIQSQFR.K	1	PEP19_HUMAN
BRCA2 and CDKN1A-interacting protein - Homo sapiens (Human)					
Q9P287	2	2.7524	K.LLQQLFLK.A	1	BCCIP_HUMAN
Breast carcinoma amplified sequence 2 - Homo sapiens (Human)					
O75934	2	3.3671	R.TIVQLENEIYQIK.Q	2	BCAS2_HUMAN
Bridging integrator 2 - Homo sapiens (Human)					
Q9UBW5	2	4.0155	R.TSLEVSPNPEPPEKPVR.T	7	BIN2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9UBW5	3	5.2264	R.ATASPRPSSGNIPSSPTASGGGSPTSPR.A	2	BIN2_HUMAN
Bromodomain and PHD finger-containing protein 3 - Homo sapiens (Human)					
Q9ULD4	2	3.0641	R.EQDEKTSAVKEELK.Y	1	BRPF3_HUMAN
Bromodomain-containing protein 4 - Homo sapiens (Human)					
O60885	2	4.4262	R.SREDEDALEQAR.R	2	BRD4_HUMAN
O60885	2	2.8959	R.AASVVQPQPLVVVKEEK.I	1	BRD4_HUMAN
Bromodomain-containing protein 8 - Homo sapiens (Human)					
Q9H0E9	2	3.4609	K.LLSTGPTEPWSIR.E	3	BRD8_HUMAN
BTB/POZ domain-containing protein KCTD18 - Homo sapiens (Human)					
Q6PI47	2	3.8169	K.NSASVTVSPASAIQTSAGATANR.F	2	KCD18_HUMAN
BUD13 homolog - Homo sapiens (Human)					
Q9BRD0	2	2.9953	K.ATDSDLSSPR.H	2	BUD13_HUMAN
Q9BRD0	2	3.2003	K.TGLVLTDIQR.E	1	BUD13_HUMAN
Q9BRD0	2	3.6399	R.HGSSDISSPR.R	2	BUD13_HUMAN
Butyrate response factor 1 - Homo sapiens (Human)					
Q07352	2	3.0366	K.QPGGGQVNSSR.Y	1	TISB_HUMAN
C4b-binding protein alpha chain precursor - Homo sapiens (Human)					
P04003	2	4.0149	K.LSLEIEQLELQR.D	1	C4BP_HUMAN
P04003	1	2.1989	K.MALEVYK.L	1	C4BP_HUMAN
P04003	2	2.7717	R.LMQCLPNPEDVK.M	1	C4BP_HUMAN
C4b-binding protein beta chain precursor - Homo sapiens (Human)					
P20851	1	3.1167	K.ALLAFQESK.N	4	C4BB_HUMAN
P20851	2	3.635	K.LIQEAPKPECEK.A	2	C4BB_HUMAN
C7 protein - Homo sapiens (Human)					
A0M8Q6	3	3.7655	R.VTHEGSTVEK.T	2	A0M8Q6_HUMA
Cadherin EGF LAG seven-pass G-type receptor 1 precursor - Homo sapiens (Human)					
Q9NYQ6	2	3.4967	R.TGSAQADGSDSEKP.-	1	CELR1_HUMAN
Cadherin-11 precursor - Homo sapiens (Human)					
P55287	3	4.8952	K.EDIRDNIVSYNDEGGGEEDTQAFDIGTLR.N	1	CAD11_HUMAN
P55287	3	4.5038	R.HTDLDRFFTINPEDGFIK.T	2	CAD11_HUMAN
P55287	3	3.802	R.VHAKDPDAANSPIR.Y	1	CAD11_HUMAN
Cadherin-13 precursor - Homo sapiens (Human)					
P55290	2	3.4957	K.DIQGSLQDIFK.F	3	CAD13_HUMAN
P55290	2	2.7552	R.YEVSSPYFK.V	1	CAD13_HUMAN
P55290	2	3.8718	R.TPHAEDMAELVIVGGK.D	2	CAD13_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P55290	2	4.0999	R.SIVVSPILIPENQR.Q	5	CAD13_HUMAN
P55290	2	3.9592	R.M#TAFDADDPATDNALLR.Y	2	CAD13_HUMAN
P55290	2	4.3333	R.INENTGSVSVTR.T	6	CAD13_HUMAN
P55290	2	3.3306	R.DVGKVVDSDRPER.S	2	CAD13_HUMAN
P55290	2	3.9097	K.RSIVVSPILIPENQR.Q	1	CAD13_HUMAN
P55290	2	4.3427	K.VNSDGGLVALR.N	2	CAD13_HUMAN
Cadherin-16 precursor - Homo sapiens (Human)					
O75309	2	4.32	R.EGAEGQIVLSGDSGK.A	2	CAD16_HUMAN
O75309	2	4.4329	R.AFQVDPTSGSVTLGVLPLR.A	1	CAD16_HUMAN
O75309	2	3.4678	K.ATEGPFAMDPDSGFLLVTR.A	1	CAD16_HUMAN
O75309	3	5.6935	R.LSAEDADAPGSPNSHVVYQLLSPEPEDGVEG	1	CAD16_HUMAN
Cadherin-2 precursor - Homo sapiens (Human)					
P19022	2	2.9489	R.YMQQNIR.Y	1	CADH2_HUMAN
P19022	3	5.6061	K.FLEAGIYEVPIIITDSGNPPK.S	3	CADH2_HUMAN
P19022	4	4.9492	K.IIRQEEGLHAGTMLTTFTAQDPDRYMQQNIR.	1	CADH2_HUMAN
P19022	2	2.8281	K.LSDPANWLK.I	1	CADH2_HUMAN
P19022	2	3.01	R.GPFPQELVR.I	2	CADH2_HUMAN
P19022	1	2.1591	R.ISGGDPTGR.F	1	CADH2_HUMAN
P19022	2	4.0359	R.LNGDFAQLNLK.I	3	CADH2_HUMAN
P19022	2	3.4311	R.M#FVLTVAAENQVPLAK.G	1	CADH2_HUMAN
Cadherin-5 precursor - Homo sapiens (Human)					
P33151	2	3.1732	R.TSDKGQFFR.V	2	CADH5_HUMAN
P33151	3	4.5346	K.KPLIGTVLAMDPDAAR.H	4	CADH5_HUMAN
P33151	2	2.7441	K.ELDSTGTPTGK.E	4	CADH5_HUMAN
P33151	2	3.0493	K.KGDIYNEK.E	2	CADH5_HUMAN
Cadherin-6 precursor - Homo sapiens (Human)					
P55285	2	2.7404	R.DNTDVRDFINQR.L	1	CADH6_HUMAN
P55285	2	4.3121	R.IVVEDVDEPPVFSK.L	5	CADH6_HUMAN
P55285	2	4.0043	K.YILSGDGAGDLFIINENTGDIQATK.R	1	CADH6_HUMAN
P55285	1	2.9402	R.FLYLGPFK.D	5	CADH6_HUMAN
P55285	2	4.0982	R.TGRPVEPESEFIIK.I	2	CADH6_HUMAN
P55285	2	3.5711	R.DIVPEALFLPR.R	2	CADH6_HUMAN
P55285	2	2.8227	R.FLYLGPFKDSATVR.I	1	CADH6_HUMAN
P55285	2	3.9051	K.VEASNPYVEPR.F	4	CADH6_HUMAN
P55285	2	3.6561	K.TPESSPPGTPIGR.I	5	CADH6_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P55285	2	4.3125	K.LHSDQDRGDGSLK.Y	2	CADH6_HUMAN
P55285	2	3.3835	K.LAYILQIR.E	3	CADH6_HUMAN
P55285	2	3.7447	K.LADM#YGGVDSDKDS.-	2	CADH6_HUMAN
P55285	2	4.6542	K.IHDINDNEPIFTK.E	3	CADH6_HUMAN
P55285	2	3.0329	K.VVYSILQGQPYFSVESETGIIK.T	1	CADH6_HUMAN
Cadherin-related tumor suppressor homolog precursor - Homo sapiens (Human)					
Q14517	2	2.9282	R.DM#PAAGSLGSSSR.N	2	FATH_HUMAN
Calbindin - Homo sapiens (Human)					
P05937	2	3.4264	K.LFDSNNDGKLELTEMAR.L	2	CALB1_HUMAN
P05937	3	5.2749	K.NKQDLDINNITTYKK.N	2	CALB1_HUMAN
P05937	2	3.7945	K.TVDDTKLAEYTDLMLK.L	1	CALB1_HUMAN
P05937	2	3.6989	K.ELQNLIQELQQAR.K	1	CALB1_HUMAN
Calcitonin gene-related peptide type 1 receptor precursor - Homo sapiens (Human)					
Q16602	2	3.2538	K.IQFGNSFSNSEALR.S	1	CALRL_HUMAN
Calcium/calmodulin-dependent protein kinase II inhibitor alpha - Homo sapiens (Human)					
Q7Z7J9	2	4.1144	R.LQDTNNFFGAGQNK.R	2	Q7Z7J9_HUMAN
Calcium-regulated heat stable protein 1 - Homo sapiens (Human)					
Q9Y2V2	2	2.7476	R.ASQGPVYK.G	2	CHSP1_HUMAN
Q9Y2V2	2	4.6949	K.LQAVEVVITHLAPGTK.H	1	CHSP1_HUMAN
Q9Y2V2	2	3.2544	K.HETWSGHVISS.-	1	CHSP1_HUMAN
Q9Y2V2	1	2.1802	R.TFSATVR.A	2	CHSP1_HUMAN
Caldesmon - Homo sapiens (Human)					
Q05682	3	4.0982	R.IAYQRNDDDEEEAARER.R	3	CALD1_HUMAN
Q05682	2	2.8327	R.LKEEIERR.R	1	CALD1_HUMAN
Q05682	2	2.7509	R.LKEEIER.R	4	CALD1_HUMAN
Q05682	2	4.6997	R.LEQYTSAIEGTK.S	10	CALD1_HUMAN
Q05682	2	3.4065	R.KVLEEEEQRR.K	2	CALD1_HUMAN
Q05682	2	4.2498	R.KVLEEEEQR.R	8	CALD1_HUMAN
Q05682	1	2.5953	R.KKGFTEVK.S	4	CALD1_HUMAN
Q05682	2	5.7877	R.ASVDTKEAEGAPQVEAGK.R	8	CALD1_HUMAN
Q05682	2	3.5596	R.RKVLEEEEQR.R	1	CALD1_HUMAN
Q05682	2	4.581	R.IAYQRNDDDEEEAAR.E	10	CALD1_HUMAN
Q05682	2	3.8936	R.GETESEEFEKLK.Q	5	CALD1_HUMAN
Q05682	2	3.3038	R.GETESEEFEK.L	2	CALD1_HUMAN
Q05682	3	4.9488	R.ASVDTKEAEGAPQVEAGKR.L	6	CALD1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q05682	2	3.7146	R.M#QNDTAENETTEK.E	1	CALD1_HUMAN
Q05682	2	3.7061	R.IEAEEKAAAQER.E	1	CALD1_HUMAN
Q05682	2	4.3107	R.M#QNDTAENETTEKEEK.S	9	CALD1_HUMAN
Q05682	3	5.0608	R.M#QNDTAENETTEKEEKSESR.Q	4	CALD1_HUMAN
Q05682	2	5.0408	R.MQNDTAENETTEK.E	7	CALD1_HUMAN
Q05682	2	4.8676	R.MQNDTAENETTEKEEK.S	5	CALD1_HUMAN
Q05682	1	3.1494	R.YEIEETETVTK.S	8	CALD1_HUMAN
Q05682	2	4.4105	R.MQNDTAENETTEKEEKSESR.Q	3	CALD1_HUMAN
Q05682	1	2.1882	K.KGFTEVK.S	1	CALD1_HUMAN
Q05682	2	4.0649	R.QERYEIEETETVTK.S	5	CALD1_HUMAN
Q05682	2	3.5347	R.RGETESEEFEKLK.Q	3	CALD1_HUMAN
Q05682	2	2.9206	R.RLKEEIER.R	2	CALD1_HUMAN
Q05682	3	5.2442	R.RM#QNDTAENETTEKEEK.S	3	CALD1_HUMAN
Q05682	2	6.9552	R.RMQNDTAENETTEKEEK.S	4	CALD1_HUMAN
Q05682	2	2.8038	R.RRGETESEEFEK.L	1	CALD1_HUMAN
Q05682	1	3.1948	K.VLEEEEQR.R	7	CALD1_HUMAN
Q05682	2	3.6053	R.RGETESEEFEK.L	8	CALD1_HUMAN
Q05682	2	2.7352	K.MPEDGLSDDK.K	1	CALD1_HUMAN
Q05682	2	2.7245	K.LQEDKPTFK.K	1	CALD1_HUMAN
Q05682	2	3.1496	K.DKEPKEEVK.S	7	CALD1_HUMAN
Q05682	2	3.5712	K.EAEGAPQVEAGK.R	7	CALD1_HUMAN
Q05682	2	3.1432	K.EAEGAPQVEAGKR.L	3	CALD1_HUMAN
Q05682	2	3.2567	K.GSSLKIEER.A	4	CALD1_HUMAN
Q05682	2	3.0342	K.HTENTFSRPGGR.A	1	CALD1_HUMAN
Q05682	2	4.6599	K.IDSRLEQYTSAIEGTK.S	16	CALD1_HUMAN
Q05682	2	2.8711	K.IEERAEFLNK.S	2	CALD1_HUMAN
Q05682	3	3.8924	K.KAQEDKLQTAVLK.K	1	CALD1_HUMAN
Q05682	2	3.2651	K.VEQKIEGK.W	7	CALD1_HUMAN
Q05682	2	3.2542	K.M#PEDGLSDDKKPFK.C	2	CALD1_HUMAN
Q05682	2	3.8014	K.MPEDGLSDDKKPFK.C	2	CALD1_HUMAN
Q05682	3	4.1594	K.NDWRDAEENKK.E	1	CALD1_HUMAN
Q05682	2	3.4931	K.SPAPKPSDLRPGDVSSK.R	2	CALD1_HUMAN
Q05682	2	4.067	K.TTESQEETVVMSLK.N	3	CALD1_HUMAN
Q05682	2	3.1466	K.STHQAAIVSK.I	8	CALD1_HUMAN
Q05682	2	3.1266	K.LKHTENTFSRPGGR.A	2	CALD1_HUMAN
Q05682	2	3.2504	K.SQNGEFM#THK.L	4	CALD1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q05682	2	3.2373	K.PSDLRPGDVSSK.R	2	CALD1_HUMAN
Q05682	2	4.0151	K.SESRQERYEIEETETVTK.S	4	CALD1_HUMAN
Q05682	3	5.4809	K.SAKPTKPAASDLPVPAEGVR.N	22	CALD1_HUMAN
Q05682	2	3.2615	K.REEMRLEAER.I	1	CALD1_HUMAN
Q05682	2	2.7826	K.REEM#RLEAER.I	1	CALD1_HUMAN
Q05682	2	3.0077	K.QQEAALELEELKK.K	2	CALD1_HUMAN
Q05682	2	3.0476	K.QQEAALELEELK.K	1	CALD1_HUMAN
Q05682	2	4.2148	K.QKQQEAALELEELKK.K	7	CALD1_HUMAN
Q05682	2	3.4055	K.QKQQEAALELEELK.K	2	CALD1_HUMAN
Q05682	2	2.7365	K.SQNGEFMTHK.L	1	CALD1_HUMAN
Calmin - Homo sapiens (Human)					
Q96JQ2	2	3.4852	K.FNSDLIDFASTSQAFNK.V	1	CLMN_HUMAN
Q96JQ2	2	3.1418	K.TSDISEPSPESSILSSR.K	1	CLMN_HUMAN
Calmodulin - Homo sapiens (Human)					
P62158	3	3.8719	R.EADIDGDGQVNYEEFVQM#MTAK.-	1	CALM_HUMAN
P62158	3	5.576	R.EADIDGDGQVNYEEFVQMM#TAK.-	2	CALM_HUMAN
P62158	2	4.288	R.EADIDGDGQVNYEEFVQMMTAK.-	41	CALM_HUMAN
Calmodulin-like protein 3 - Homo sapiens (Human)					
P27482	2	3.1442	R.SLGQNPTEAELR.D	2	CALL3_HUMAN
P27482	2	2.8595	K.LSDEEVDEMIR.A	2	CALL3_HUMAN
P27482	2	3.1927	R.AADTDGDGQVNYEEFVR.V	2	CALL3_HUMAN
Calmodulin-like protein 5 - Homo sapiens (Human)					
Q9NZT1	3	6.4726	K.AFSAVDTDGNGTINAQELGAALK.A	4	CALL5_HUMAN
Calmodulin-regulated spectrin-associated protein 1. - Homo sapiens (Human)					
Q5T5Y3	2	3.882	K.ISQQQEQLLMK.S	2	Q5T5Y3_HUMAN
Q5T5Y3	2	3.9734	K.LNETISTLQQAILK.I	1	Q5T5Y3_HUMAN
Q5T5Y3	2	2.7118	K.SIQGEDIPDQR.H	1	Q5T5Y3_HUMAN
Q5T5Y3	3	4.6923	R.TPTDPGLDSALEPSGDPHGK.C	5	Q5T5Y3_HUMAN
Calnexin precursor - Homo sapiens (Human)					
P27824	3	4.7412	R.KIPNPDFFEDLEPFR.M	2	CALX_HUMAN
Calpain-1 catalytic subunit - Homo sapiens (Human)					
P07384	2	2.9606	K.YLGQDYEQLR.V	1	CAN1_HUMAN
Calpastatin - Homo sapiens (Human)					
P20810	2	3.6729	K.AIPVSQQMEGPHLPNK.K	3	ICAL_HUMAN
P20810	3	6.0136	K.EGITGPPADSSKPIGPDDAIDALSSDFTCGSP	1	ICAL_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P20810	2	3.0679	K.AIPVSQQM#EGPHLPNK.K	1	ICAL_HUMAN
Calponin-1 - Homo sapiens (Human)					
P51911	2	3.0425	K.LAQKYDHQR.E	1	CNN1_HUMAN
P51911	2	2.9202	R.IGNNFMDGLK.D	1	CNN1_HUMAN
P51911	2	3.5124	R.GPAYGLSAEVK.N	3	CNN1_HUMAN
P51911	1	3.4509	R.GMTVYGLPR.Q	5	CNN1_HUMAN
P51911	2	3.2996	R.GM#TVYGLPR.Q	2	CNN1_HUMAN
P51911	2	3.4249	R.NIIGLQMGTNK.F	2	CNN1_HUMAN
P51911	2	4.8568	K.LGTDQPLDQATISLQM#GTNK.G	2	CNN1_HUMAN
P51911	2	3.4892	R.NIIGLQM#GTNK.F	3	CNN1_HUMAN
P51911	2	4.271	K.GASQAGMTAPGTKR.Q	4	CNN1_HUMAN
P51911	2	4.4169	K.GASQAGMTAPGTK.R	7	CNN1_HUMAN
P51911	2	3.2283	K.GASQAGM#TAPGTKR.Q	5	CNN1_HUMAN
P51911	2	3.8669	K.GASQAGM#TAPGTK.R	3	CNN1_HUMAN
P51911	2	4.1033	K.FASQQGMTAYGTR.R	8	CNN1_HUMAN
P51911	2	3.735	K.FASQQGM#TAYGTR.R	6	CNN1_HUMAN
P51911	3	5.1869	K.LGTDQPLDQATISLQMGTNK.G	2	CNN1_HUMAN
P51911	2	4.4008	K.INESTQNWHQLENIGNFIK.A	1	CNN1_HUMAN
Calponin-2 - Homo sapiens (Human)					
Q99439	3	3.9143	K.NHILPPMDHSTISLQM#GTNK.C	1	CNN2_HUMAN
Q99439	3	3.7701	K.NHILPPMDHSTISLQMGTNK.C	1	CNN2_HUMAN
Q99439	2	3.6645	R.TWIEGLTGLSIGPDFQK.G	2	CNN2_HUMAN
Calponin-3 - Homo sapiens (Human)					
Q15417	3	5.6028	K.M#QTDKPFDQTTISLQM#GTNK.G	5	CNN3_HUMAN
Q15417	3	6.5828	R.DIYDQKLTLQPVDNSTISLQM#GTNK.V	2	CNN3_HUMAN
Q15417	2	5.2281	K.AGQSVIGLQMGTNK.C	6	CNN3_HUMAN
Q15417	1	2.2462	R.DIYDQK.L	4	CNN3_HUMAN
Q15417	2	3.4124	R.DYQYSDQGIDY.-	1	CNN3_HUMAN
Q15417	3	5.3162	K.MQTDKPFDQTTISLQMGTNK.G	7	CNN3_HUMAN
Q15417	3	5.4805	K.MQTDKPFDQTTISLQM#GTNK.G	5	CNN3_HUMAN
Q15417	3	5.5689	K.M#QTDKPFDQTTISLQMGTNK.G	5	CNN3_HUMAN
Q15417	2	5.4469	K.LTLQPVDNSTISLQM#GTNK.V	13	CNN3_HUMAN
Q15417	2	3.52	K.GMSVYGLGR.Q	10	CNN3_HUMAN
Q15417	2	3.3876	K.GM#SVYGLGR.Q	5	CNN3_HUMAN
Q15417	2	3.0821	K.GASQAGMLAPGTRR.D	3	CNN3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q15417	2	3.2638	K.GASQAGM#LAPGTR.R	2	CNN3_HUMAN
Q15417	2	4.7924	K.AGQSVIGLQM#GTNK.C	4	CNN3_HUMAN
Q15417	2	3.0737	K.LTLQPVDNSTISLQMGTNK.V	1	CNN3_HUMAN
Q15417	2	3.8172	K.GASQAGMLAPGTR.R	3	CNN3_HUMAN
Calreticulin precursor - Homo sapiens (Human)					
P27797	2	4.092	K.IDNSQVESGSLEDDWDFLPPK.K	5	CALR_HUMAN
P27797	2	5.5177	R.FYALSASFEPFSNK.G	7	CALR_HUMAN
P27797	1	2.8118	K.VHVIFNYK.G	3	CALR_HUMAN
P27797	2	3.8153	K.SDFGKFVLSSGK.F	2	CALR_HUMAN
P27797	1	2.2654	K.NVLINK.D	2	CALR_HUMAN
P27797	1	2.5944	K.KVHVIFNYK.G	1	CALR_HUMAN
P27797	3	4.4073	K.IDNSQVESGSLEDDWDFLPPKK.I	11	CALR_HUMAN
P27797	2	4.8963	K.HEQNIDCGGGYVK.L	8	CALR_HUMAN
P27797	2	3.3867	K.GKNVLINK.D	3	CALR_HUMAN
P27797	2	5.9125	K.FYGDEEKDKGLQTSQDAR.F	8	CALR_HUMAN
P27797	2	3.01	K.FYGDEEKDK.G	3	CALR_HUMAN
P27797	1	2.3207	K.FVLSSGK.F	8	CALR_HUMAN
P27797	3	3.9245	K.DPDASKPEDWDER.A	1	CALR_HUMAN
P27797	2	3.444	K.DKGLQTSQDAR.F	2	CALR_HUMAN
P27797	2	4.4458	K.IKDPDASKPEDWDER.A	6	CALR_HUMAN
CALRETICULIN=CALCIUM binding protein - Homo sapiens (Human)					
Q9UDG2	2	4.0545	-.EPAVYFKEQFLDGDGWTSR.W	3	Q9UDG2_HUMA
Calumenin precursor - Homo sapiens (Human)					
O43852	2	2.8291	K.TFDQLTPEESK.E	1	CALU_HUMAN
O43852	2	3.0289	R.WIYEDVER.Q	8	CALU_HUMAN
O43852	2	3.9955	K.TFDQLTPEESKER.L	4	CALU_HUMAN
O43852	3	3.8914	K.VHNDAQSFDYDHDAFLGAEEAK.T	1	CALU_HUMAN
O43852	2	3.5408	R.HLVYESDQNK.D	2	CALU_HUMAN
O43852	3	4.7052	R.HLVYESDQNKDGK.L	5	CALU_HUMAN
O43852	4	5.7109	R.VHHEPQLSDKVHNDAQSFDYDHDAFLGAEE	3	CALU_HUMAN
O43852	3	6.07	K.MDKEETKDWILPSDYDHAEAEAR.H	5	CALU_HUMAN
O43852	3	4.5612	R.VHHEPQLSDK.V	1	CALU_HUMAN
O43852	2	2.9584	K.RWIYEDVER.Q	1	CALU_HUMAN
O43852	4	5.4684	K.MADKDGDLIATKEEFTAFLHPEEYDYMK.D	2	CALU_HUMAN
O43852	2	4.3474	K.MADKDGDLIATK.E	2	CALU_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43852	4	5.6763	K.M#DKEETKDWILPSDYDHAEAEAR.H	3	CALU_HUMAN
O43852	2	3.9758	K.M#ADKDGDLIATK.E	1	CALU_HUMAN
O43852	3	6.9992	K.LTKEEIVDKYDLFVGSQATDFGEALVR.H	1	CALU_HUMAN
O43852	2	3.972	K.IDGDKDGFVTVDELKDWIK.F	1	CALU_HUMAN
O43852	2	3.2703	K.TEREQFVEFR.D	3	CALU_HUMAN
O43852	3	4.5462	K.TEREQFVEFRDK.N	2	CALU_HUMAN
cAMP response element-binding protein - Homo sapiens (Human)					
P16220	2	3.9908	R.TAPTSTIAPGVVM#ASSPALPTQPAEEAAR.K	2	CREB1_HUMAN
P16220	2	3.4504	R.TAPTSTIAPGVVMASSPALPTQPAEEAAR.K	1	CREB1_HUMAN
cAMP response element-binding protein 5 - Homo sapiens (Human)					
Q02930	2	3.0855	R.VVDEDPDER.R	1	CREB5_HUMAN
cAMP-dependent protein kinase inhibitor gamma - Homo sapiens (Human)					
Q9Y2B9	3	6.1853	R.KLAGDMGELALEGAEGQVEGSAPDKEAGNQ	1	IPKG_HUMAN
cAMP-regulated phosphoprotein 19 - Homo sapiens (Human)					
P56211	2	2.7756	R.KPSLVASK.L	2	ARP19_HUMAN
P56211	2	3.5834	R.YPHLGQKPGGSDFLR.K	3	ARP19_HUMAN
CAP-Gly domain-containing linker protein 2 - Homo sapiens (Human)					
Q9UDT6	2	5.2589	R.TSTGSASSSAAVAASSK.E	4	CLIP2_HUMAN
Q9UDT6	2	3.3923	K.ATLNSGPGAQQK.E	2	CLIP2_HUMAN
Q9UDT6	2	4.2967	K.LQEAQEELAGLQR.H	1	CLIP2_HUMAN
Q9UDT6	3	5.0779	K.HDLETAMHVK.E	2	CLIP2_HUMAN
Q9UDT6	3	4.1222	K.MEHQLELGNLQAK.H	1	CLIP2_HUMAN
Q9UDT6	2	3.7574	K.QSSGPSSSPAAAAAPEKPGPK.A	2	CLIP2_HUMAN
Q9UDT6	3	4.4384	K.SKLDSLASDHQK.S	3	CLIP2_HUMAN
Q9UDT6	2	3.2305	K.SLEDLKATLNSGPGAQQK.E	1	CLIP2_HUMAN
Q9UDT6	2	4.2108	K.TGNESGSNLSDSGSVK.R	3	CLIP2_HUMAN
Q9UDT6	2	3.2996	R.DALDQAQQVEK.L	1	CLIP2_HUMAN
Q9UDT6	3	4.3928	R.KTHDASGQLVLISQELLR.K	1	CLIP2_HUMAN
Q9UDT6	4	5.8966	R.RGEIEELQQCLLHSGPPPPDHPDAAEILR.L	2	CLIP2_HUMAN
Q9UDT6	2	3.2006	R.RLEAELETVSR.K	1	CLIP2_HUMAN
Q9UDT6	3	3.7617	K.AQHEQYVAEAEEK.L	1	CLIP2_HUMAN
Q9UDT6	2	3.2752	K.AYQAEVDKLR.A	2	CLIP2_HUMAN
Carbonic anhydrase 1 - Homo sapiens (Human)					
P00915	1	2.9513	K.YSSLAEAASK.A	2	CAH1_HUMAN
P00915	2	3.8797	K.VLDALQAIK.T	6	CAH1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P00915	2	6.3887	K.TSETKHDTSLKPISVSYNPATAK.E	5	CAH1_HUMAN
P00915	2	4.991	K.LYPIANGNNQSPVDIK.T	2	CAH1_HUMAN
P00915	3	5.5628	K.HDTSLKPISVSYNPATAK.E	7	CAH1_HUMAN
P00915	2	4.6203	K.ADGLAVIGVLMK.V	3	CAH1_HUMAN
P00915	3	4.4469	K.YSSLAEAASKADGLAVIGVLMK.V	2	CAH1_HUMAN
Carbonic anhydrase 12 precursor - Homo sapiens (Human)					
O43570	2	2.832	K.WTYFGPDGENSWSK.K	2	CAH12_HUMAN
Carbonic anhydrase 2 - Homo sapiens (Human)					
P00918	3	4.4812	K.AVQQPDGLAVLGIFLK.V	1	CAH2_HUMAN
P00918	2	2.9139	K.VGSAKPGLQK.V	1	CAH2_HUMAN
P00918	3	4.4873	K.YDPSLKPLSVSYDQATSLR.I	2	CAH2_HUMAN
Carbonic anhydrase 9 precursor - Homo sapiens (Human)					
Q16790	2	3.2752	K.GGVSYRPAEVAETGA.-	1	CAH9_HUMAN
Q16790	2	4.6495	R.VIEASFPAGVDSSPR.A	1	CAH9_HUMAN
Q16790	2	2.8425	R.YGGDPPWPR.V	2	CAH9_HUMAN
Carbonyl reductase [NADPH] 1 - Homo sapiens (Human)					
P16152	3	5.5	K.SPEEGAETPVYLALLPPDAEGPHGQFVSEK.	1	CBR1_HUMAN
Carboxylesterase 2 precursor - Homo sapiens (Human)					
O00748	2	3.0233	K.MIPGVVDGVFLPR.H	1	EST2_HUMAN
Carnitine O-palmitoyltransferase I, muscle isoform - Homo sapiens (Human)					
Q92523	1	2.1904	R.KALLDIADLFQVPK.A	1	CPT1B_HUMAN
Casein kinase I isoform epsilon - Homo sapiens (Human)					
P49674	2	3.6805	R.GAPANVSSSDLTGR.Q	3	KC1E_HUMAN
P49674	3	3.9172	R.IPASQTSVPFDHLGK.-	1	KC1E_HUMAN
Caskin-2 - Homo sapiens (Human)					
Q8WXE0	2	5.0275	K.SIGTKEQEGTPSASTK.H	2	CSKI2_HUMAN
Q8WXE0	3	5.9619	K.VPGAGTAPKPVSVACTQLAFSGPK.L	4	CSKI2_HUMAN
Q8WXE0	2	3.4162	R.LSSVSGPSPEPPPLDGSPGPK.E	3	CSKI2_HUMAN
Q8WXE0	3	5.924	R.SHSLSRPGPTEGDAEGEAEGPVGSTLGSYA	1	CSKI2_HUMAN
Q8WXE0	2	2.9355	R.TSPSVTPTPAR.G	1	CSKI2_HUMAN
Q8WXE0	3	3.7515	K.QRPKPAGPPPR.E	1	CSKI2_HUMAN
Catalase - Homo sapiens (Human)					
P04040	2	4.2109	K.GAGAFGYFEVTHDITK.Y	1	CATA_HUMAN
P04040	2	4.0625	R.AFYVNVLNEEQR.K	3	CATA_HUMAN
P04040	3	4.6933	R.AAQKADVLTTGAGNPVGDKLNVITVGPR.G	1	CATA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P04040	2	3.0643	K.LNVITVGPR.G	2	CATA_HUMAN
P04040	2	5.1575	K.ADVLTTGAGNPVGDKLNVITVGPR.G	2	CATA_HUMAN
P04040	3	4.4873	K.NAIHTFVQSGSHLAAR.E	2	CATA_HUMAN
Catenin alpha-1 - Homo sapiens (Human)					
P35221	2	3.8876	K.HVNPVQALSEFK.A	4	CTNA1_HUMAN
P35221	3	4.2536	K.KHVNPVQALSEFK.A	5	CTNA1_HUMAN
Catenin beta-1 - Homo sapiens (Human)					
P35222	3	3.8993	K.AAVSHWQQQSYLDSGIHSGATTTAPSLSGK.	1	CTNB1_HUMAN
Cathepsin B precursor - Homo sapiens (Human)					
P07858	2	2.8314	K.ICEPGYSPTYK.Q	1	CATB_HUMAN
P07858	2	3.8151	K.NGPVEGAFSVYSDFLLYK.S	2	CATB_HUMAN
Cathepsin D precursor - Homo sapiens (Human)					
P07339	2	3.9536	R.DPDAQPGGELM#LGGTDSK.Y	2	CATD_HUMAN
P07339	3	4.7917	R.ISVNNVLPVFDNLMQQK.L	2	CATD_HUMAN
Cathepsin G precursor - Homo sapiens (Human)					
P08311	2	2.7151	R.ENTQQHITAR.R	1	CATG_HUMAN
P08311	2	3.5208	R.RENTQQHITAR.R	3	CATG_HUMAN
Cathepsin L precursor - Homo sapiens (Human)					
P07711	2	3.8077	K.NSWGEEWGM#GGYVK.M	3	CATL_HUMAN
Cation-dependent mannose-6-phosphate receptor precursor - Homo sapiens (Human)					
P20645	2	3.7547	R.HTLADNFNPVSEER.G	1	MPRD_HUMAN
Caveolin-1 - Homo sapiens (Human)					
Q03135	2	4.9137	K.YVDSEGHLYTVPIR.E	5	CAV1_HUMAN
CCDC43 protein - Homo sapiens (Human)					
Q86WV7	2	4.1934	R.NTNVEDVLNAR.K	4	Q86WV7_HUMA
CCR4-NOT transcription complex subunit 2 - Homo sapiens (Human)					
Q9NZN8	2	2.8292	R.GM#SNNTPQLNR.S	2	CNOT2_HUMAN
Q9NZN8	2	3.1481	R.GMSNNTPQLNR.S	2	CNOT2_HUMAN
Q9NZN8	3	4.4517	R.NMMNHSQVGQGIGIPSR.T	2	CNOT2_HUMAN
Q9NZN8	2	3.6151	R.TNSM#SSSGLGSPNR.S	2	CNOT2_HUMAN
Q9NZN8	2	3.6338	R.TNSMSSSGLGSPNR.S	2	CNOT2_HUMAN
CD109 antigen precursor - Homo sapiens (Human)					
Q6YHK3	2	3.3452	K.TLTLPSLPLNSADEIYELR.V	1	CD109_HUMAN
CD2 antigen cytoplasmic tail-binding protein 2 - Homo sapiens (Human)					
O95400	3	4.1997	R.KVTFQGVGDEEDEDEIIVPK.K	1	CD2B2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O95400	2	2.8632	R.KLDPPGGQFYNSK.R	1	CD2B2_HUMAN
O95400	2	3.1672	K.LVDPVAGSGGPGSR.F	2	CD2B2_HUMAN
O95400	2	3.5891	K.VTFQGVGDEEDEDEIIVPK.K	2	CD2B2_HUMAN
CD2-associated protein - Homo sapiens (Human)					
Q9Y5K6	1	3.2427	K.TQSVEITK.T	4	CD2AP_HUMAN
Q9Y5K6	2	3.1666	R.SNLEMEIEK.L	1	CD2AP_HUMAN
Q9Y5K6	2	4.5196	K.ANTTAFLTPLEIK.A	3	CD2AP_HUMAN
Q9Y5K6	2	2.9619	R.ISTYGLPAGGIQPHPQTK.N	2	CD2AP_HUMAN
Q9Y5K6	2	3.3041	R.TLFAYEGTNEDELTFK.E	1	CD2AP_HUMAN
Q9Y5K6	2	3.5318	R.HGNVASLVQR.I	2	CD2AP_HUMAN
Q9Y5K6	2	4.2139	R.GIGFGDIFKEGSVK.L	3	CD2AP_HUMAN
Q9Y5K6	2	2.9748	R.GIGFGDIFK.E	4	CD2AP_HUMAN
Q9Y5K6	2	3.3292	R.ETEFKDDSLPIKR.E	2	CD2AP_HUMAN
Q9Y5K6	2	3.9779	K.YFSLKPEEKDEK.S	2	CD2AP_HUMAN
Q9Y5K6	2	3.1646	K.SVDFDSLTVR.T	1	CD2AP_HUMAN
Q9Y5K6	2	5.721	K.LQEEGWLEGELNGR.R	2	CD2AP_HUMAN
Q9Y5K6	2	3.2512	K.LGLFPSNFVK.E	2	CD2AP_HUMAN
Q9Y5K6	2	4.2026	K.KLQEEGWLEGELNGR.R	1	CD2AP_HUMAN
Q9Y5K6	2	4.342	K.INGEVSSISSKFETEPVSK.L	2	CD2AP_HUMAN
Q9Y5K6	1	2.2828	K.FETEPVSK.L	1	CD2AP_HUMAN
Q9Y5K6	1	2.2239	K.DLEEEK.T	2	CD2AP_HUMAN
Q9Y5K6	2	3.8674	K.APAPKPELIAAEK.K	2	CD2AP_HUMAN
Q9Y5K6	2	3.888	K.AKVETDDVKK.N	3	CD2AP_HUMAN
Q9Y5K6	3	4.1804	K.APAPKPELIAAEKK.Y	4	CD2AP_HUMAN
CD44 antigen precursor - Homo sapiens (Human)					
P16070	3	7.2939	K.SQEMVHLVNKESSETPDQFMTADETR.N	1	CD44_HUMAN
P16070	2	3.0462	K.ESSETPDQFMTADETR.N	1	CD44_HUMAN
P16070	2	3.3969	K.ESSETPDQFM#TADETR.N	2	CD44_HUMAN
P16070	1	2.7331	R.NLQNVDMK.I	3	CD44_HUMAN
CD59 glycoprotein precursor - Homo sapiens (Human)					
P13987	2	3.6944	K.AGLQVYNK.C	11	CD59_HUMAN
CD97 antigen precursor - Homo sapiens (Human)					
P48960	2	4.0846	K.KQAELEEIYESSIR.G	1	CD97_HUMAN
P48960	2	4.34	K.LVDELMEAPGDVEALAPPVR.H	1	CD97_HUMAN
P48960	2	3.857	K.QAELEEIYESSIR.G	2	CD97_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P48960	2	4.0518	K.TSSAEVTIQNVIK.L	2	CD97_HUMAN
CD99 antigen precursor - Homo sapiens (Human)					
P14209	2	3.5951	R.NANAEPAVQR.T	3	CD99_HUMAN
Cdc42 effector protein 1 - Homo sapiens (Human)					
Q00587	2	3.5764	K.LSPVGWVSSSQGK.R	2	BORG5_HUMAN
Q00587	3	4.6946	K.NAISLPQLNQAAYDSLVVGK.L	2	BORG5_HUMAN
Q00587	2	3.2079	R.TPVPSTVQANTFEFADAEEDDEVKV.-	1	BORG5_HUMAN
Cdc42 effector protein 2 - Homo sapiens (Human)					
O14613	3	4.8326	K.FHLLPGTM#VEGPEEDGTFDLPFQFTR.T	2	BORG1_HUMAN
Cdc42 effector protein 4 - Homo sapiens (Human)					
Q9H3Q1	3	4.0458	R.QPDKEFSFMDEEEEDEIRV.-	1	BORG4_HUMAN
Q9H3Q1	2	4.5572	K.AGEPDGESLDEQPSSSSSK.R	14	BORG4_HUMAN
Q9H3Q1	2	3.9884	K.AGEPDGESLDEQPSSSSSKR.S	2	BORG4_HUMAN
Q9H3Q1	2	4.7307	K.ANDGEGGDEEAGTEEAVPR.R	10	BORG4_HUMAN
Q9H3Q1	2	4.9535	K.KANDGEGGDEEAGTEEAVPR.R	2	BORG4_HUMAN
Q9H3Q1	2	3.0085	R.ADLTAEM#ISAPLGDFR.H	1	BORG4_HUMAN
Q9H3Q1	2	2.7788	R.DMLGSLRDSALFVK.N	1	BORG4_HUMAN
Cdc42 GTPase-activating protein - Homo sapiens (Human)					
Q2M1Z3	2	5.5884	R.DPANQSTQGASTAASR.E	3	Q2M1Z3_HUMA
Cdc42-interacting protein 4 - Homo sapiens (Human)					
Q15642	2	3.254	K.EGGEGYVPTSYLR.V	2	CIP4_HUMAN
Q15642	2	2.7302	K.YEAWLAEAESR.V	1	CIP4_HUMAN
Q15642	3	3.989	R.HARPPDPPASAPPDSSSNSASQDTK.E	1	CIP4_HUMAN
Q15642	2	2.9498	R.LQQQLEER.S	1	CIP4_HUMAN
CDNA FLJ10233 fis, clone HEMBB1000266 - Homo sapiens (Human)					
Q9NW82	3	4.6437	R.SGPSEVTGSDASGPDPQLAVTMGFTGFGKK.	1	Q9NW82_HUMA
Q9NW82	2	4.2434	K.NIALDKTDDSNPR.E	2	Q9NW82_HUMA
Q9NW82	3	4.1124	K.TM#FAQVESDDEEAKNEPEWK.K	1	Q9NW82_HUMA
Q9NW82	3	5.0223	R.SGPSEVTGSDASGPDPQLAVTM#GFTGFGK	2	Q9NW82_HUMA
CDNA FLJ13298 fis, clone OVARC1001306, weakly similar to N-MYC PROTO-ONCOGENE PRO					
Q9H8R3	2	3.0681	K.KEADEQLIKETK.T	2	Q9H8R3_HUMA
CDNA FLJ16182 fis, clone BRTHA2002133 - Homo sapiens (Human)					
Q6ZNE3	2	2.7787	R.QEKEIKDIQIGK.S	1	Q6ZNE3_HUMA
CDNA FLJ16607 fis, clone TESTI4010382, highly similar to Rattus norvegicus dynein, cytoplasmic					
Q6ZMX7	2	2.7879	K.TCLEEWTKSAGLEK.F	1	Q6ZMX7_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
CDNA FLJ20036 fis, clone COL00219 - Homo sapiens (Human)					
Q9NXV6	2	2.7088	K.SSSQTSGSLVSK.S	1	Q9NXV6_HUMA
Q9NXV6	2	3.1984	K.SSSQTSTSQLPSK.S	2	Q9NXV6_HUMA
Q9NXV6	2	4.457	K.STSLASVSQLASK.S	1	Q9NXV6_HUMA
Q9NXV6	2	4.304	K.TSSEASVSSSVAK.N	1	Q9NXV6_HUMA
Q9NXV6	2	4.6974	R.SSGISSQNSSTSDGDR.S	4	Q9NXV6_HUMA
CDNA FLJ31051 fis, clone HSYRA2000605, weakly similar to MYOSIN HEAVY CHAIN, CLONE					
Q96ND2	2	3.007	R.VSQDLIETEK.K	2	Q96ND2_HUMA
Q96ND2	2	4.7896	K.TLEGIQYDNSILK.M	2	Q96ND2_HUMA
CDNA FLJ31713 fis, clone NT2RI2006487 - Homo sapiens (Human)					
Q96MY4	2	2.718	R.IQSTSPASVLSSTK.E	1	Q96MY4_HUMA
CDNA FLJ32810 fis, clone TESTI2002729, weakly similar to Homo sapiens oligophrenin 1 - Hom					
Q96M56	2	4.3513	K.SYSGSIQSLTSVGSK.E	4	Q96M56_HUMA
CDNA FLJ35765 fis, clone TESTI2004941 - Homo sapiens (Human)					
Q8NA79	2	2.8535	R.FWAC*LGSSAAKTLTSLSNAELCFPR.Q	1	Q8NA79_HUMA
CDNA FLJ36810 fis, clone ASTRO2001249 - Homo sapiens (Human)					
Q8N9N8	2	2.99	K.EVLGEHIVPSDQQQIVR.V	1	Q8N9N8_HUMA
Q8N9N8	3	5.2074	K.RGDFLIVDPIEEGEKVK.A	1	Q8N9N8_HUMA
Q8N9N8	1	2.5537	R.FLVSMPSKYRK.N	2	Q8N9N8_HUMA
Q8N9N8	3	5.1716	R.TPGNNLHEVETAQGQR.F	4	Q8N9N8_HUMA
CDNA FLJ37649 fis, clone BRHIP2000534, moderately similar to GAMMA-INTERFERON-INDU					
Q8N9E5	2	2.9349	K.TSLSAPPNTSSTENPK.T	1	Q8N9E5_HUMA
CDNA FLJ40925 fis, clone UTERU2006486 - Homo sapiens (Human)					
Q8N7K9	2	3.1736	K.TQTAAETQPSSR.I	2	Q8N7K9_HUMA
CDNA FLJ41046 fis, clone NT2RP7009030 - Homo sapiens (Human)					
Q6ZWI0	2	2.741	K.WVDSGCKGIFSHFSQQR.K	1	Q6ZWI0_HUMA
CDNA FLJ41346 fis, clone BRAWH2005315, moderately similar to Neuronal-STOP protein - Hom					
Q6ZWB8	3	4.5678	K.TQGPVATEPDKDQGSVVPGLLK.G	1	Q6ZWB8_HUMA
CDNA FLJ41802 fis, clone NHNPC2002565 - Homo sapiens (Human)					
Q6ZW13	3	3.8634	K.AEAELPPKLPLQEEEPEDSQSEPSPSAK.Q	1	Q6ZW13_HUMA
CDNA FLJ41861 fis, clone NTONG2008672 - Homo sapiens (Human)					
Q6ZVZ7	3	4.2067	R.SPVQEDRPGPGLGLSTPVPVTEQGTDQIR.T	1	Q6ZVZ7_HUMA
CDNA FLJ42542 fis, clone BRACE3004435 - Homo sapiens (Human)					
Q6ZVI5	2	3.3377	R.SPGSLSPWLLGRR.G	2	Q6ZVI5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
CDNA FLJ44590 fis, clone BLADE2000256, moderately similar to Homo sapiens suppression of tu					
Q6ZTJ7	2	4.125	R.NTGTLGSLEEPAGGASVSAGSR.A	1	Q6ZTJ7_HUMAN
CDNA FLJ45252 fis, clone BRHIP2011199 - Homo sapiens (Human)					
Q6ZSR9	2	3.1368	K.QSIAGSVSITSLSSR.T	1	Q6ZSR9_HUMA
Q6ZSR9	2	4.9756	R.HYSPEDEPSPEAQPIAAYK.I	3	Q6ZSR9_HUMA
CDNA FLJ45505 fis, clone BRTHA2020642, weakly similar to DRR1 protein - Homo sapiens (Hu					
Q6ZSI4	2	3.9902	K.QKLQEEQENAPEFVK.V	3	Q6ZSI4_HUMAN
Q6ZSI4	2	2.8672	R.RTGQEVAQAQES.-	1	Q6ZSI4_HUMAN
Q6ZSI4	3	6.6501	K.LEQLELEKQKLQEEQENAPEFVK.V	2	Q6ZSI4_HUMAN
Q6ZSI4	1	3.205	R.TGQEVAQAQES.-	5	Q6ZSI4_HUMAN
CDNA FLJ46103 fis, clone TESTI2023903, weakly similar to Homo sapiens ubiquilin 1 - Homo sa					
Q6ZRU1	2	3.4814	R.VIYNSSGGFSSNTSANDTLNKVNHTSK.A	1	Q6ZRU1_HUMA
CDNA FLJ46765 fis, clone TRACH3024512 - Homo sapiens (Human)					
Q6ZQZ9	2	2.867	K.LEISRCMGPNCTHQK.S	2	Q6ZQZ9_HUMA
CDNA FLJ46846 fis, clone UTERU3004635, moderately similar to Neuroblast differentiation assoc					
Q6ZQN2	2	3.9082	K.VEGDLKGPEVDIR.D	2	Q6ZQN2_HUMA
Q6ZQN2	2	3.9058	K.M#KGDVDVSLPK.V	3	Q6ZQN2_HUMA
Q6ZQN2	2	4.1111	K.MEGDLKAPEVDIK.G	2	Q6ZQN2_HUMA
Q6ZQN2	2	4.6085	K.MEGDLKAPEVDIKGPK.V	4	Q6ZQN2_HUMA
Q6ZQN2	2	3.9279	K.MKGDVDVSLPK.V	3	Q6ZQN2_HUMA
Q6ZQN2	2	4.5956	K.SPQISM#SDIDLNLK.G	1	Q6ZQN2_HUMA
Q6ZQN2	2	4.4652	K.SPQISMSDIDLNLK.G	1	Q6ZQN2_HUMA
Q6ZQN2	2	2.9502	K.VDIDAPDVSIEGPDAK.L	1	Q6ZQN2_HUMA
Q6ZQN2	2	3.7353	K.VDIDVPDVNIEGPDAK.L	5	Q6ZQN2_HUMA
Q6ZQN2	3	5.2789	K.VESDLKGPEVDIEGPEGK.L	14	Q6ZQN2_HUMA
Q6ZQN2	2	4.8845	K.VGIDTPDIDIHGPEGK.L	8	Q6ZQN2_HUMA
Q6ZQN2	2	2.9173	K.VKGDM#DISLPK.V	2	Q6ZQN2_HUMA
Q6ZQN2	2	4.1215	K.VKGDMDISLPK.V	7	Q6ZQN2_HUMA
Q6ZQN2	3	4.1178	K.M#EGDLKAPEVDIKGPK.V	1	Q6ZQN2_HUMA
Q6ZQN2	3	3.8424	K.SPQISM#SDIDLNLKGPK.I	4	Q6ZQN2_HUMA
Q6ZQN2	2	4.5075	K.VDIDTPDIDIHGPEGK.L	4	Q6ZQN2_HUMA
Q6ZQN2	2	3.6004	K.FSMPGFKGEGPEVDVNLPK.A	1	Q6ZQN2_HUMA
Q6ZQN2	1	3.3349	K.ADIEISGPK.V	5	Q6ZQN2_HUMA
Q6ZQN2	3	4.1653	K.ADLDVSGPKVDIDVPDVNIEGPDAK.L	4	Q6ZQN2_HUMA
Q6ZQN2	2	2.8718	K.APEVDIKGPK.V	2	Q6ZQN2_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6ZQN2	3	3.7552	K.FKMPEINIK.A	2	Q6ZQN2_HUMA
Q6ZQN2	2	4.2188	K.FSM#PGFKGEGPDVDVTLPK.A	6	Q6ZQN2_HUMA
Q6ZQN2	2	3.6342	K.ISMPDIDFNLKGPK.V	2	Q6ZQN2_HUMA
Q6ZQN2	2	3.36	K.FSMPGFKGEGPDVDVTLPK.A	1	Q6ZQN2_HUMA
Q6ZQN2	2	5.2857	K.SPQISMSDIDLNLKGPK.I	6	Q6ZQN2_HUMA
Q6ZQN2	2	3.0345	K.GEGPDVDVTLPK.A	1	Q6ZQN2_HUMA
Q6ZQN2	3	6.3438	K.ADIDVSGPKVDIDTPDIDIHGPEGK.L	2	Q6ZQN2_HUMA
Q6ZQN2	3	4.7311	K.GPQVSGELKGPGVDVNLKGPR.I	1	Q6ZQN2_HUMA
Q6ZQN2	2	5.4945	K.GPSLDIDTPDVNIEGPEGK.L	7	Q6ZQN2_HUMA
Q6ZQN2	2	3.5345	K.IKGDMDISVPK.L	3	Q6ZQN2_HUMA
Q6ZQN2	2	3.4155	K.ISM#PDIDFNLKGPK.V	1	Q6ZQN2_HUMA
Q6ZQN2	2	2.8946	K.ISMPDIDFNLK.G	1	Q6ZQN2_HUMA
Q6ZQN2	2	3.7655	K.FSM#PGFKGEGPEVDVNLPK.A	1	Q6ZQN2_HUMA
CDNA FLJ46889 fis, clone UTERU3017995, highly similar to Homo sapiens likely ortholog of rat					
Q6ZQU3	2	2.8143	K.ASSSILINESEPTTNIQIR.L	1	Q6ZQU3_HUMA
CDNA PSEC0066 fis, clone NT2RP2001087, weakly similar to TETRACYCLINE RESISTANCE P					
Q8NBP5	2	2.751	K.TGTEAEAADSGAVGARR.F	1	Q8NBP5_HUMA
CDNA: FLJ22312 fis, clone HRC05210. - Homo sapiens (Human)					
Q9H6G0	2	4.0736	K.EGESGECAVAESEDR.A	2	Q9H6G0_HUMA
Q9H6G0	3	4.9328	R.TVLEGSTASTSPADHSALPNQSLTVR.E	4	Q9H6G0_HUMA
Q9H6G0	3	4.6627	K.TSDSKEGGEGFTVDTPAK.A	2	Q9H6G0_HUMA
Q9H6G0	2	3.0648	K.AGPATTTSSETR.Q	2	Q9H6G0_HUMA
Q9H6G0	2	3.6206	K.LTGVIVENENITK.E	2	Q9H6G0_HUMA
CDNA: FLJ22427 fis, clone HRC09013 - Homo sapiens (Human)					
Q9H6A9	3	3.9145	R.SFDTVIGAGTPPGLAEPLLVVRPK.D	1	Q9H6A9_HUMA
CDNA: FLJ22461 fis, clone HRC10107 - Homo sapiens (Human)					
Q9H698	2	2.7317	K.M#AEPSSFVCRSTGSLLK.T	1	Q9H698_HUMAN
Q9H698	2	4.3963	R.DVSNLESSGGTENK.A	2	Q9H698_HUMAN
Cell adhesion molecule 1 precursor - Homo sapiens (Human)					
Q9BY67	2	3.0702	K.SDDSVIQLLNPNR.Q	1	CADM1_HUMAN
Cell cycle associated protein 1 - Homo sapiens (Human)					
Q9BV09	2	2.8872	R.DGYQQNFK.R	3	Q9BV09_HUMA
Q9BV09	3	5.3957	K.SSGPPPPSGSSGSEAAAGAGAAAPASQHPA	3	Q9BV09_HUMA
Q9BV09	3	5.5735	K.SSGPPPPSGSSGSEAAAGAGAAAPASQHPA	3	Q9BV09_HUMA
Cell division cycle 2-related protein kinase 7 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9NYV4	2	5.4444	K.DGSGGASGTLQPSSGGGSSNSR.E	10	CD2L7_HUMAN
Q9NYV4	2	3.3223	K.RSNEETDDYGK.A	2	CD2L7_HUMAN
Cell division cycle 5-like protein - Homo sapiens (Human)					
Q99459	3	4.4006	R.SKLVLPAPQISDAELQEVVK.V	1	CDC5L_HUMAN
Q99459	2	4.1327	K.ESDLPSAILQTSGVSEFTK.K	2	CDC5L_HUMAN
Q99459	3	4.8626	K.GGLNTPLHESDFSGVTPQR.Q	3	CDC5L_HUMAN
Q99459	2	3.3984	K.VGQASEIAR.Q	3	CDC5L_HUMAN
Q99459	3	4.1334	R.DKLNINPEDGMADYSDPSYVK.Q	1	CDC5L_HUMAN
Q99459	2	3.9588	R.DNEEETTDDPR.K	3	CDC5L_HUMAN
Q99459	2	2.7785	R.DNEEETTDDPRK.L	1	CDC5L_HUMAN
Q99459	2	3.7514	R.LGLLGLPAPK.N	1	CDC5L_HUMAN
Q99459	2	4.4385	R.QVVQTPNTVLSTPFR.T	1	CDC5L_HUMAN
Cell division cycle and apoptosis regulator protein 1 - Homo sapiens (Human)					
Q8IX12	2	3.1315	R.IQTLPNQNQSQTQPLLK.T	1	CCAR1_HUMAN
Q8IX12	2	3.1798	K.DLSQLQENLK.I	1	CCAR1_HUMAN
Cell division cycle protein 27 homolog - Homo sapiens (Human)					
P30260	2	3.3378	K.SVFSQSGNSR.E	3	CDC27_HUMAN
Cell division cycle-associated protein 4 - Homo sapiens (Human)					
Q9BXL8	2	2.7131	K.C*VGHEEDVEGALAGLK.T	1	CDCA4_HUMAN
Cell division protein kinase 6 - Homo sapiens (Human)					
Q00534	1	2.1127	-.MEKDGLCR.A	1	CDK6_HUMAN
Cell growth-regulating nucleolar protein - Homo sapiens (Human)					
Q9NX58	2	3.4368	K.DAVEQQGEVKK.N	2	LYAR_HUMAN
Q9NX58	3	3.8695	K.LPEHPEGGEPEDDEAPAK.G	1	LYAR_HUMAN
Q9NX58	3	4.5923	K.VKDAVEQQGEVKK.N	2	LYAR_HUMAN
Cell surface glycoprotein MUC18 precursor - Homo sapiens (Human)					
P43121	2	3.3364	R.EVTVPVFYPTEK.V	1	MUC18_HUMAN
P43121	4	4.8582	R.VYKAPEEPNIQVNPLGIPVNSKEPEEVATCVG	1	MUC18_HUMAN
P43121	2	5.1075	R.VRQGQGQSEPGEYEQR.L	4	MUC18_HUMAN
P43121	2	3.3571	R.TQLVNVAIFGPPWM#AFK.E	1	MUC18_HUMAN
P43121	2	3.8509	R.QGQGQSEPGEYEQR.L	4	MUC18_HUMAN
P43121	2	4.6344	R.GATLALTQVTPQDER.I	4	MUC18_HUMAN
P43121	2	2.9861	R.EAEEETTNDNGVLVLEPARK.E	1	MUC18_HUMAN
P43121	2	3.2736	K.VWLEVEPVGMLK.E	4	MUC18_HUMAN
P43121	2	4.1857	K.VWLEVEPVGM#LK.E	4	MUC18_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P43121	2	3.7045	R.GPVLQLHDLKR.E	2	MUC18_HUMAN
Centaurin-delta 3 - Homo sapiens (Human)					
Q8WWN8	3	4.9337	R.TVFGGLSGPATTQRPGLSPALGGPGVSR.S	2	CEND3_HUMAN
Centaurin-gamma 1 - Homo sapiens (Human)					
Q99490	2	3.8156	R.AISAFGPSASINGLVK.D	1	CENG1_HUMAN
Centrin-2 - Homo sapiens (Human)					
P41208	2	4.204	K.MNFGDFLTVM#TQK.M	3	CETN2_HUMAN
P41208	3	4.2831	K.AFKLFDDDETGKISFK.N	1	CETN2_HUMAN
P41208	2	3.0754	R.EAFDLFDADGTGTIDVK.E	1	CETN2_HUMAN
P41208	3	4.699	R.M#SPKPELTEEQKQEIR.E	1	CETN2_HUMAN
P41208	2	4.391	K.MNFGDFLTVMTQK.M	6	CETN2_HUMAN
P41208	2	4.4391	K.M#NFGDFLTVMTQK.M	1	CETN2_HUMAN
P41208	2	3.8807	K.M#ISEIDKEGTGK.M	3	CETN2_HUMAN
P41208	2	3.462	K.M#NFGDFLTVM#TQK.M	1	CETN2_HUMAN
P41208	2	3.9979	K.MISEIDKEGTGK.M	4	CETN2_HUMAN
Centrin-3 - Homo sapiens (Human)					
O15182	2	2.7825	K.DYDREATGK.I	2	CETN3_HUMAN
Centromere protein C 1 - Homo sapiens (Human)					
Q03188	2	3.5916	R.SVQAHEVHQK.I	1	CENPC_HUMAN
Q03188	2	3.5487	R.TISPAESTALFQGR.K	1	CENPC_HUMAN
Centromere protein J - Homo sapiens (Human)					
Q9HC77	2	2.8712	K.QLKEAEGPLPIKAK.P	2	CENPJ_HUMAN
Centromere protein U - Homo sapiens (Human)					
Q71F23	2	2.8956	K.MLKESQM#LTNLK.R	1	CENPU_HUMAN
Centrosomal protein of 170 kDa - Homo sapiens (Human)					
Q5SW79	3	4.1338	R.FGYDTNLFTVVQGEMRVPEEALK.H	1	CE170_HUMAN
Q5SW79	2	2.7058	K.HQDQAVTSSAHHR.G	1	CE170_HUMAN
Centrosome-associated protein 350 - Homo sapiens (Human)					
Q5VT06	2	4.1851	R.LTDSSPSSTSTSNSQR.L	2	CEP35_HUMAN
Centrosome-associated protein CEP250 - Homo sapiens (Human)					
Q9BV73	2	2.9526	K.ATASSPTQQDGR.G	1	CP250_HUMAN
Ceruloplasmin precursor - Homo sapiens (Human)					
P00450	3	4.4193	K.HYYIGIIETTWDYASDHGEK.K	1	CERU_HUMAN
P00450	3	3.9505	K.MYYSAVDPTKDIFTGLIGPMK.I	2	CERU_HUMAN
CGG triplet repeat-binding protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9UFW8	2	3.7587	R.KAEFEEQNVR.K	3	CGBP1_HUMAN
Q9UFW8	2	3.1374	K.VSVIQDFVK.M	2	CGBP1_HUMAN
CGI-151 protein - Homo sapiens (Human)					
Q9Y3E9	2	4.769	R.YAALSDQGLDIK.A	5	Q9Y3E9_HUMA
cGMP-gated cation channel alpha 1 - Homo sapiens (Human)					
P29973	2	2.7449	R.AEFQARIDAIK.Q	1	CNGA1_HUMAN
cGMP-inhibited 3',5'-cyclic phosphodiesterase A - Homo sapiens (Human)					
Q14432	2	3.9324	K.ISAVQFPESADTTAK.Q	2	PDE3A_HUMAN
Q14432	3	4.3325	R.RVSSTWTTTTSATGLPTLEPAPVRR.D	2	PDE3A_HUMAN
Charged multivesicular body protein 1a - Homo sapiens (Human)					
Q9HD42	1	2.4134	K.NMAQVTK.A	1	CHM1A_HUMAN
Q9HD42	1	2.3951	K.VQTAVTMK.G	2	CHM1A_HUMAN
Q9HD42	1	2.2017	R.VDAVASK.V	2	CHM1A_HUMAN
Charged multivesicular body protein 1b - Homo sapiens (Human)					
Q7LBR1	1	2.2475	K.GNMEVAR.I	1	CHM1B_HUMAN
Q7LBR1	2	2.7965	R.VDAVAAR.V	1	CHM1B_HUMAN
Q7LBR1	1	2.2045	K.SMDATLK.T	2	CHM1B_HUMAN
Q7LBR1	2	2.8005	K.HLFNLK.F	1	CHM1B_HUMAN
Q7LBR1	2	2.7566	K.NQAVNFLR.M	1	CHM1B_HUMAN
Charged multivesicular body protein 2a - Homo sapiens (Human)					
O43633	2	2.8938	K.KIIADIKK.M	1	CHM2A_HUMAN
O43633	2	3.0073	R.ANIQAVSLK.I	2	CHM2A_HUMAN
O43633	2	2.9981	R.RKTPEELLR.Q	1	CHM2A_HUMAN
O43633	2	3.183	K.SNNSMAQAMK.G	3	CHM2A_HUMAN
O43633	2	7.1297	K.KAEAAASALADADADLEER.L	5	CHM2A_HUMAN
O43633	2	2.7033	K.IMM#EFER.Q	1	CHM2A_HUMAN
O43633	2	5.7308	K.AEAAASALADADADLEER.L	5	CHM2A_HUMAN
O43633	3	6.94	K.AEAAASALADADADLEERLK.N	5	CHM2A_HUMAN
O43633	3	6.2901	K.KAEAAASALADADADLEERLK.N	6	CHM2A_HUMAN
Charged multivesicular body protein 2b - Homo sapiens (Human)					
Q9UQN3	2	3.2043	K.TLQTM#QNFQK.E	2	CHM2B_HUMAN
Q9UQN3	2	3.4641	K.ATISDEEIER.Q	3	CHM2B_HUMAN
Q9UQN3	2	3.957	K.TLQTMQNFQK.E	8	CHM2B_HUMAN
Q9UQN3	2	3.0627	K.M#AGAMSTTAK.T	1	CHM2B_HUMAN
Q9UQN3	2	3.956	K.TVDDVIKEQNR.E	2	CHM2B_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Charged multivesicular body protein 4b - Homo sapiens (Human)					
Q9H444	2	2.7369	K.IEQELTAAK.K	1	CHM4B_HUMAN
Q9H444	3	4.4901	K.QLAQIDGTLSTIEFQR.E	3	CHM4B_HUMAN
Q9H444	3	4.5378	R.LRDTEEMLSK.K	6	CHM4B_HUMAN
Q9H444	2	4.1112	R.LRDTEEM#LSK.K	3	CHM4B_HUMAN
Q9H444	3	3.8724	R.LRDTEEMLSKK.Q	1	CHM4B_HUMAN
Q9H444	2	3.9603	R.EALENANTNTEVLK.N	2	CHM4B_HUMAN
Q9H444	1	2.5308	K.NMGYAAK.A	8	CHM4B_HUMAN
Q9H444	3	4.081	K.KKEEEDDDMKELENWAGSM#.-	1	CHM4B_HUMAN
Q9H444	1	3.9701	K.KIEQELTAAK.K	7	CHM4B_HUMAN
Q9H444	2	2.8289	K.GGPTPQEAIQR.L	2	CHM4B_HUMAN
Q9H444	3	4.0145	K.KIEQELTAAKK.H	6	CHM4B_HUMAN
Charged multivesicular body protein 5 - Homo sapiens (Human)					
Q9NZZ3	3	7.2902	R.DNLAQQSFNM#EQANYTIQSLK.D	3	CHMP5_HUMAN
Q9NZZ3	1	2.7991	R.LDAELVK.Y	2	CHMP5_HUMAN
Q9NZZ3	3	7.8219	R.DNLAQQSFNMEQANYTIQSLK.D	3	CHMP5_HUMAN
Q9NZZ3	2	3.8334	K.ISRLDAELVK.Y	4	CHMP5_HUMAN
Q9NZZ3	2	4.0659	K.APPPSLTDCIGTVDSR.A	8	CHMP5_HUMAN
Q9NZZ3	2	3.195	K.RMYEQQR.D	7	CHMP5_HUMAN
Charged multivesicular body protein 6 - Homo sapiens (Human)					
Q96FZ7	2	2.9878	R.ILDETQEAVEYQR.Q	1	CHMP6_HUMAN
Chloride channel CLIC-like protein 1 precursor - Homo sapiens (Human)					
Q96S66	3	3.9026	K.AQLKSEAAGSPDQGSTYSPAR.G	1	CLCC1_HUMAN
Q96S66	3	5.5287	K.ESSTESSQSAKPVSGQDTSGNTEGSPAAEK.	1	CLCC1_HUMAN
Chloride intracellular channel protein 1 - Homo sapiens (Human)					
O00299	2	4.1171	K.LAALNPESNTAGLDIFAK.F	1	CLIC1_HUMAN
Chondroitin sulfate proteoglycan 4 precursor - Homo sapiens (Human)					
Q6UVK1	2	3.671	K.SPPSAGYLVMVSR.G	1	CSPG4_HUMAN
Q6UVK1	2	4.032	R.VTGALQFGELQK.Q	2	CSPG4_HUMAN
Q6UVK1	2	2.7293	R.FTQADVDSGR.L	1	CSPG4_HUMAN
Q6UVK1	2	4.122	R.LSDGQGFTQDDIQAGR.V	3	CSPG4_HUMAN
Chromatin modifying protein 4A - Homo sapiens (Human)					
Q14D22	3	4.0535	K.LPSVPSTHLPAGPAPKVDEDEEALKQLAEWV	1	Q14D22_HUMAN
Q14D22	1	3.6689	R.TMELAAQSMK.K	5	Q14D22_HUMAN
Q14D22	3	4.1602	R.FEQQLAQTDGTLSTLEFQREAIENATTNAEVL	1	Q14D22_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q14D22	2	4.4763	R.EAIENATTNAEVLR.T	4	Q14D22_HUMAN
Q14D22	3	4.9987	K.RFEQQLAQTDGTLSTLEFQR.E	4	Q14D22_HUMAN
Q14D22	2	3.0397	K.RAALQALR.R	2	Q14D22_HUMAN
Q14D22	3	4.6483	K.KQEFLEQKIQQELQTAK.K	2	Q14D22_HUMAN
Q14D22	2	2.9087	K.KQEFLEQK.I	2	Q14D22_HUMAN
Q14D22	2	2.7389	K.IQQELQTAKK.Y	1	Q14D22_HUMAN
Q14D22	2	3.2013	K.IQQELQTAK.K	1	Q14D22_HUMAN
Q14D22	2	2.9975	K.EKGPTPEEAIQK.L	2	Q14D22_HUMAN
Q14D22	2	3.1883	R.TMELAAQSMKK.A	2	Q14D22_HUMAN
Q14D22	3	5.9085	K.KRFEQQLAQTDGTLSTLEFQR.E	3	Q14D22_HUMAN
Chromobox protein homolog 1 - Homo sapiens (Human)					
P83916	2	3.7682	K.NSDEADLVPAK.E	4	CBX1_HUMAN
P83916	2	5.0474	K.WKNSDEADLVPAKEANVK.C	1	CBX1_HUMAN
P83916	2	4.6284	K.NSDEADLVPAKEANVK.C	6	CBX1_HUMAN
P83916	3	4.5768	R.GLEPERIIGATDSSGELMFLMK.W	3	CBX1_HUMAN
P83916	3	4.7687	K.KVEEVLEEEEEEYVVEK.V	5	CBX1_HUMAN
P83916	2	2.894	K.GKVEYLLK.W	2	CBX1_HUMAN
P83916	2	3.2955	K.VEEVLEEEEEEYVVEK.V	1	CBX1_HUMAN
Chromobox protein homolog 3 - Homo sapiens (Human)					
Q13185	2	5.2882	K.WKDSDEADLVLAK.E	10	CBX3_HUMAN
Q13185	3	4.5951	R.RVVNGKVEYFLK.W	2	CBX3_HUMAN
Q13185	2	3.7287	R.VVNGKVEYFLK.W	2	CBX3_HUMAN
Q13185	2	3.544	R.LTWHSCPEDEAQ.-	5	CBX3_HUMAN
Q13185	3	4.9268	R.KSLSDSESDDSKSK.K	2	CBX3_HUMAN
Q13185	2	3.8049	R.KSLSDSESDDSK.S	1	CBX3_HUMAN
Q13185	2	4.091	K.VEEAEPEEFVVEK.V	5	CBX3_HUMAN
Q13185	2	4.1556	K.SLSDSESDDSKSK.K	6	CBX3_HUMAN
Q13185	2	3.619	K.SLSDSESDDSK.S	5	CBX3_HUMAN
Q13185	2	3.6324	K.SKKVEEAEPEEFVVEK.V	2	CBX3_HUMAN
Q13185	2	4.2671	K.DSDEADLVLAK.E	4	CBX3_HUMAN
Q13185	2	5.9361	K.KVEEAEPEEFVVEK.V	6	CBX3_HUMAN
Chromobox protein homolog 5 - Homo sapiens (Human)					
P45973	2	3.6464	R.KSNFSNSADDIK.S	1	CBX5_HUMAN
P45973	3	3.9953	R.LTWHAYPEDAENKEK.E	1	CBX5_HUMAN
P45973	2	3.0372	K.REQSNDIAR.G	1	CBX5_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P45973	2	2.9408	K.KREQSNDIAR.G	2	CBX5_HUMAN
P45973	2	3.0355	K.SNFSNSADDIK.S	1	CBX5_HUMAN
Chromodomain Y-like protein - Homo sapiens (Human)					
Q9Y232	2	2.9993	K.NSQLFAASQK.F	2	CDYL1_HUMAN
Q9Y232	4	6.114	R.TAVDGFQSESPEKLDPVEQGQEDTVAPEVA	2	CDYL1_HUMAN
Chromodomain-helicase-DNA-binding protein 2 - Homo sapiens (Human)					
O14647	2	2.878	K.NNPDYNWNVR.K	1	CHD2_HUMAN
Chromodomain-helicase-DNA-binding protein 6 - Homo sapiens (Human)					
Q8TD26	2	3.331	R.SPEESTESTDSQK.R	2	CHD6_HUMAN
Chromodomain-helicase-DNA-binding protein 8 - Homo sapiens (Human)					
Q9HCK8	3	3.8873	K.VLSASEVAALSSPASSAPHSGGK.T	1	CHD8_HUMAN
Chromosome 20 open reading frame 3 - Homo sapiens (Human)					
A2A2F9	2	3.4482	-.VVTDDDGQAPEAK.D	1	A2A2F9_HUMAN
Ciliary dynein heavy chain 8 - Homo sapiens (Human)					
Q96JB1	2	2.7747	R.DEMDEITQGLISVMK.R	1	DYH8_HUMAN
Cingulin - Homo sapiens (Human)					
Q9P2M7	3	4.6791	R.LASSEGFQKPSASLSQLESQNQLLQER.L	1	CING_HUMAN
Cingulin-like protein 1 - Homo sapiens (Human)					
Q0VF96	3	4.0586	R.QLLEQTLKDLEYELEAK.S	2	Q0VF96_HUMAN
Q0VF96	2	4.3672	K.GQQELTQQTNEETAK.Q	1	Q0VF96_HUMAN
Q0VF96	3	4.0455	R.SQHNEKVEENSTLQQR.L	1	Q0VF96_HUMAN
Q0VF96	2	4.856	R.SNTSEQDQAGTEMR.V	4	Q0VF96_HUMAN
Q0VF96	2	4.1063	R.SNTSEQDQAGTEM#R.V	2	Q0VF96_HUMAN
Q0VF96	2	4.411	R.ALENELEAAQGNLSQTTQEQK.Q	3	Q0VF96_HUMAN
Q0VF96	2	3.8472	K.VLETEGSQESTVIR.A	2	Q0VF96_HUMAN
Q0VF96	2	3.7335	K.NQQNIKEERER.M	2	Q0VF96_HUMAN
Q0VF96	2	3.913	K.GALIEELLQAK.Q	4	Q0VF96_HUMAN
Q0VF96	4	5.0791	K.QMEDKVSQLEMELEEERNNSDLLSER.I	1	Q0VF96_HUMAN
Cisplatin resistance-associated overexpressed protein - Homo sapiens (Human)					
O95232	3	4.4874	K.ESDTKNEVNGTSEDIKSEGDTQSN.-	2	CROP_HUMAN
C-jun-amino-terminal kinase-interacting protein 4 - Homo sapiens (Human)					
O60271	3	4.9777	K.LHQLSGSDQLESTAHSR.I	2	JIP4_HUMAN
O60271	2	3.315	R.VESLESQTR.Q	2	JIP4_HUMAN
O60271	2	4.2137	R.EVENLILENTQLLETK.N	1	JIP4_HUMAN
O60271	2	2.7701	R.ETDYPAGEDLSESGQVDK.A	1	JIP4_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O60271	2	3.1632	K.FQELSQPR.S	3	JIP4_HUMAN
O60271	3	5.8404	K.ERPISLGIFPLPAGDGLLTPDAQK.G	5	JIP4_HUMAN
O60271	2	4.4563	K.AFDFLSEETEASLASR.R	1	JIP4_HUMAN
O60271	3	3.7146	K.ATTPASTANSDVATIPTDTPLKEENEGFVK.V	1	JIP4_HUMAN
O60271	2	4.0007	K.HIEVQVAQETR.N	2	JIP4_HUMAN
Clathrin interactor 1 - Homo sapiens (Human)					
Q14677	2	4.4045	K.LGELSDKIGSTIDDTISK.F	4	EPN4_HUMAN
Q14677	2	3.2596	R.SQNTDMVQK.S	2	EPN4_HUMAN
Q14677	2	3.412	K.QDAFANFANFSK.-	2	EPN4_HUMAN
Q14677	2	4.931	K.HIHITQATETTTTR.H	4	EPN4_HUMAN
Q14677	2	4.1788	K.GEFKDEEETVTTK.H	2	EPN4_HUMAN
Q14677	2	3.8101	K.ASPDQNASTHTPQSSVK.T	1	EPN4_HUMAN
Q14677	3	3.8947	R.SPKGEFKDEEETVTTK.H	1	EPN4_HUMAN
Clathrin light chain A - Homo sapiens (Human)					
P09496	1	2.5357	R.LCDFNPK.S	8	CLCA_HUMAN
P09496	2	3.1365	R.LQSEPESIRK.W	2	CLCA_HUMAN
P09496	2	3.1965	R.LQSEPESIR.K	1	CLCA_HUMAN
P09496	2	3.0662	R.LEALDANSRK.Q	2	CLCA_HUMAN
P09496	2	4.6894	K.AIKELEEWYARQDEQLQK.T	3	CLCA_HUMAN
P09496	2	3.3462	R.LEALDANSR.K	7	CLCA_HUMAN
P09496	2	4.0546	R.EEQMERLEALDANSR.K	4	CLCA_HUMAN
P09496	2	3.1986	K.WREEQMERLEALDANSR.K	2	CLCA_HUMAN
P09496	2	4.211	K.ELEEWYARQDEQLQK.T	3	CLCA_HUMAN
P09496	2	3.1494	K.WREEQMER.L	5	CLCA_HUMAN
P09496	2	4.4843	K.AIKELEEWYAR.Q	6	CLCA_HUMAN
P09496	2	3.3991	R.KQEAEWKEK.A	1	CLCA_HUMAN
Clathrin light chain B - Homo sapiens (Human)					
P09497	3	5.3901	R.LQELDAASKVTEQEWR.E	7	CLCB_HUMAN
P09497	2	3.1257	R.LTQEPESIR.K	1	CLCB_HUMAN
P09497	1	2.9208	R.LQELDAASK.V	8	CLCB_HUMAN
P09497	2	3.1658	R.KRLQELDAASK.V	2	CLCB_HUMAN
P09497	2	3.016	K.VTEQEWREK.A	2	CLCB_HUMAN
P09497	2	2.7329	K.VTEQEWR.E	1	CLCB_HUMAN
P09497	3	3.729	K.RLQELDAASKVTEQEWR.E	1	CLCB_HUMAN
P09497	2	3.0719	K.RLQELDAASK.V	3	CLCB_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P09497	2	3.33	K.KDLEEWNQR.Q	2	CLCB_HUMAN
P09497	2	3.1844	K.ESKEETPGTEWEK.V	3	CLCB_HUMAN
P09497	2	3.3731	K.DLEEWNQR.Q	4	CLCB_HUMAN
P09497	3	4.0964	K.AKKDLEEWNQR.Q	3	CLCB_HUMAN
P09497	2	3.0919	R.LTQEPESIRK.W	2	CLCB_HUMAN
P09497	1	3.8528	K.VAQLCDFNPK.S	14	CLCB_HUMAN
Cleavage and polyadenylation specificity factor 6 - Homo sapiens (Human)					
Q16630	2	3.013	K.QFLSQFEM#QSR.K	1	CPSF6_HUMAN
Q16630	3	4.8342	R.AVSDASAGDYGSAIETLVTAISLIK.Q	1	CPSF6_HUMAN
Q16630	2	5.0897	R.TPLSEAEFEEIMNR.N	4	CPSF6_HUMAN
Cleavage stimulation factor 64 kDa subunit - Homo sapiens (Human)					
P33240	2	2.9244	R.GSLPANVPTPR.G	1	CSTF2_HUMAN
Cleavage stimulation factor 64 kDa subunit, tau variant - Homo sapiens (Human)					
Q9H0L4	2	3.3662	R.AMETEVLETR.V	1	CSTFT_HUMAN
Q9H0L4	3	5.3225	R.GPMTGGIQGPGPINIGAGGPPQGPR.Q	1	CSTFT_HUMAN
CLIP-associating protein 1 - Homo sapiens (Human)					
Q7Z460	2	4.8618	K.LLGSGYGGLTGGSSR.G	4	CLAP1_HUMAN
Q7Z460	2	3.0774	R.AQTTNSNSSSSSDVSTHS.-	1	CLAP1_HUMAN
Q7Z460	2	3.6595	R.DGGAASPATEGR.G	4	CLAP1_HUMAN
Q7Z460	2	4.4484	R.QSSGSATNVASTPDNR.G	3	CLAP1_HUMAN
Q7Z460	2	3.302	R.RQSSGSATNVASTPDNR.G	1	CLAP1_HUMAN
Clusterin precursor - Homo sapiens (Human)					
P10909	2	3.4205	K.TLLSNLEEAK.K	2	CLUS_HUMAN
P10909	2	5.8935	R.VTTVASHTSDSDVPSGVTEVVVK.L	4	CLUS_HUMAN
P10909	2	2.827	R.KYNELLK.S	1	CLUS_HUMAN
P10909	2	4.1371	R.KTLLSNLEEAKK.K	6	CLUS_HUMAN
P10909	2	4.1083	R.KTLLSNLEEAK.K	5	CLUS_HUMAN
P10909	2	3.7331	K.TLLSNLEEAKK.K	2	CLUS_HUMAN
P10909	2	3.4479	R.ELDESLQVAER.L	3	CLUS_HUMAN
P10909	2	3.3966	K.LFDSDPITVTVPVEVSR.K	1	CLUS_HUMAN
P10909	1	2.3564	K.FMETVAEK.A	1	CLUS_HUMAN
P10909	2	5.2034	K.YVNKEIQNAVNGVK.Q	2	CLUS_HUMAN
c-Myc-responsive protein Rcl - Homo sapiens (Human)					
O43598	3	5.91	R.FGTVLTEHVAAAELGAR.G	7	RCL_HUMAN
O43598	3	4.4622	R.RFGTVLTEHVAAAELGAR.G	3	RCL_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
CNKSR family member 3 - Homo sapiens (Human)					
Q5SGD5	2	2.8913	R.IPPIIEESSSPPYR.F	1	Q5SGD5_HUMA
Coagulation factor IX precursor - Homo sapiens (Human)					
P00740	2	3.6098	K.FGSGYVSGWGR.V	1	FA9_HUMAN
Coatomer subunit beta' - Homo sapiens (Human)					
P35606	2	3.9623	R.NVMEEGKDFQPSR.S	1	COPB2_HUMAN
Coatomer subunit epsilon - Homo sapiens (Human)					
O14579	2	3.1751	K.ENDFDRLVLQYAPSA.-	1	COPE_HUMAN
Cofilin-1 - Homo sapiens (Human)					
P23528	2	3.726	R.KSSTPEEVKK.R	12	COF1_HUMAN
P23528	2	5.4504	K.EILVGDVGQTVDDPYATFVK.M	3	COF1_HUMAN
P23528	3	4.6163	K.HELQANCYEEVKDR.C	1	COF1_HUMAN
P23528	2	4.7926	K.LGGSAVISLEGKPL.-	53	COF1_HUMAN
P23528	3	5.5788	K.LTGIKHELQANCYEEVKDR.C	3	COF1_HUMAN
P23528	2	3.1054	K.MLPDKDCR.Y	1	COF1_HUMAN
P23528	2	3.6561	K.NIILEEGKEILVGDVGQTVDDPYATFVK.M	2	COF1_HUMAN
P23528	2	3.2404	R.KSSTPEEVK.K	6	COF1_HUMAN
P23528	2	3.197	K.AVLFCLSEDKK.N	2	COF1_HUMAN
Cofilin-2 - Homo sapiens (Human)					
Q9Y281	2	3.4594	K.LGGNVVVSLEGKPL.-	2	COF2_HUMAN
Coiled-coil and C2 domain-containing protein 1A - Homo sapiens (Human)					
Q6P1N0	2	2.9671	K.LANQDEGPEDEEDEVPK.K	1	C2D1A_HUMAN
Q6P1N0	2	3.4788	R.TPQSGSAPTAK.A	1	C2D1A_HUMAN
Q6P1N0	2	2.9776	R.YQVAAAQAK.S	1	C2D1A_HUMAN
Coiled-coil and C2 domain-containing protein 1B - Homo sapiens (Human)					
Q5T0F9	2	3.4575	R.FGAVLEALEK.G	2	C2D1B_HUMAN
Coiled-coil domain-containing protein 102A - Homo sapiens (Human)					
Q96A19	2	3.5339	R.MSEDLEDELGAR.S	1	C102A_HUMAN
Q96A19	3	4.0898	R.SLDEQTEQSENLQVQLEHLQSR.L	2	C102A_HUMAN
Q96A19	2	2.8932	R.LQAENAAEWGR.R	1	C102A_HUMAN
Q96A19	3	4.3013	R.DVGSERPPGSQELELVESLLK.S	1	C102A_HUMAN
Q96A19	2	2.975	K.GSLLTILGSPSPER.M	1	C102A_HUMAN
Q96A19	2	2.9038	K.ELAQAEDELDEAHNQAR.K	1	C102A_HUMAN
Coiled-coil domain-containing protein 104 - Homo sapiens (Human)					
Q96G28	2	3.3009	R.TKQIQNMEQK.G	1	CC104_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96G28	2	3.167	K.NIEMQLQAIR.I	2	CC104_HUMAN
Coiled-coil domain-containing protein 117 - Homo sapiens (Human)					
Q8IWD4	2	3.2131	K.HVAAGTAFPQR.T	1	CC117_HUMAN
Coiled-coil domain-containing protein 12 - Homo sapiens (Human)					
Q8WUD4	3	4.5732	K.VKEQLEAAKPEPVIEEVDLANLAPR.K	2	CCD12_HUMAN
Coiled-coil domain-containing protein 124 - Homo sapiens (Human)					
Q96CT7	2	2.8906	R.LKQENPNMR.L	2	CC124_HUMAN
Q96CT7	2	3.3765	R.SPDNPMNQR.A	2	CC124_HUMAN
Q96CT7	2	3.3932	R.SPDNPM#NQR.A	12	CC124_HUMAN
Q96CT7	2	4.0558	R.VLEEGSVEAR.T	5	CC124_HUMAN
Q96CT7	2	2.9811	R.RLDQLER.K	1	CC124_HUMAN
Q96CT7	2	5.1333	R.AAFTAFEEAQLPR.L	7	CC124_HUMAN
Q96CT7	2	3.8843	K.SHLEVPLEENVNR.R	3	CC124_HUMAN
Q96CT7	2	4.3339	R.LLEEEDSKLKGGK.A	2	CC124_HUMAN
Q96CT7	2	3.3516	R.LLEEEDSKLK.G	4	CC124_HUMAN
Coiled-coil domain-containing protein 128 - Homo sapiens (Human)					
Q6ZMI0	3	3.748	K.KGVVDEQANSAALKEQLK.M	1	CC128_HUMAN
Coiled-coil domain-containing protein 137 - Homo sapiens (Human)					
Q6PK04	2	2.9064	K.NQDEQEIPFR.L	2	CC137_HUMAN
Q6PK04	3	4.0208	R.KGESDGAYIHR.M	2	CC137_HUMAN
Coiled-coil domain-containing protein 16 - Homo sapiens (Human)					
Q96NB3	3	5.2684	R.ENTAEALPEGFFDDPEVDAR.V	1	CCD16_HUMAN
Q96NB3	3	5.884	R.RENTAEALPEGFFDDPEVDAR.V	2	CCD16_HUMAN
Q96NB3	3	4.8813	R.ATPSKPSGLSLLPDYEDEEEEEEEEEGDGER	3	CCD16_HUMAN
Q96NB3	3	3.7012	K.RVINQEELRR.L	1	CCD16_HUMAN
Q96NB3	2	3.7104	K.EASQGSSASSAPQSVKR.K	2	CCD16_HUMAN
Q96NB3	2	4.6593	K.EASQGSSASSAPQSVK.R	5	CCD16_HUMAN
Coiled-coil domain-containing protein 23 - Homo sapiens (Human)					
Q8N300	2	4.2229	R.VMTELEQQQFDEFCK.Q	3	CCD23_HUMAN
Coiled-coil domain-containing protein 28A - Homo sapiens (Human)					
Q8IWP9	3	4.5129	K.GAQSTPIQHSFLTDVSDVQEMER.G	1	CC28A_HUMAN
Q8IWP9	2	2.9667	K.LHLADAQDVPNTSAS.-	2	CC28A_HUMAN
Q8IWP9	2	4.0026	R.GLLSLLNDFHSGK.L	1	CC28A_HUMAN
Coiled-coil domain-containing protein 47 precursor - Homo sapiens (Human)					
Q96A33	2	2.9892	R.RLEEAALR.R	1	CCD47_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Coiled-coil domain-containing protein 50 - Homo sapiens (Human)					
Q8IVM0	3	3.9496	R.KLQEEELLATQVDMR.A	2	CCD50_HUMAN
Q8IVM0	2	5.1467	R.MAHRDQEWYDAEIAR.K	2	CCD50_HUMAN
Q8IVM0	3	6.0654	R.YKDLEQQDCEIAQEIQEK.L	7	CCD50_HUMAN
Q8IVM0	3	4.8127	R.M#AHRDQEWYDAEIAR.K	2	CCD50_HUMAN
Q8IVM0	2	2.8481	R.LVQHDLQVAK.Q	1	CCD50_HUMAN
Q8IVM0	2	4.1771	R.LLQEKELQEEK.K	3	CCD50_HUMAN
Q8IVM0	2	2.8445	R.LLMAEEK.K	1	CCD50_HUMAN
Q8IVM0	2	4.1127	R.KLQEEELLATQVDM#R.A	2	CCD50_HUMAN
Q8IVM0	2	3.5813	R.IQEKKDEDIAR.L	2	CCD50_HUMAN
Q8IVM0	3	6.4327	R.DFAVLEDHTLAHSLQEQEIEHHLASNVQR.N	2	CCD50_HUMAN
Q8IVM0	2	3.9046	R.AYADSYYYEDGGMKPR.V	3	CCD50_HUMAN
Q8IVM0	2	2.9901	R.AYADSYYYEDGGM#KPR.V	1	CCD50_HUMAN
Q8IVM0	2	3.5539	R.AAQVAQDEEIAR.L	2	CCD50_HUMAN
Q8IVM0	1	2.1558	K.LAIEAER.R	1	CCD50_HUMAN
Q8IVM0	2	3.3676	K.QLQEEDLKAQAQLQK.R	2	CCD50_HUMAN
Coiled-coil domain-containing protein 53 - Homo sapiens (Human)					
Q9Y3C0	2	3.0055	K.M#VQVGVPVMAIR.N	2	CCD53_HUMAN
Q9Y3C0	2	3.3698	K.MVQVGVPVMAIR.N	3	CCD53_HUMAN
Coiled-coil domain-containing protein 55 - Homo sapiens (Human)					
Q9H0G5	2	2.8721	R.NNEETVMSAR.D	2	CCD55_HUMAN
Q9H0G5	2	4.2419	R.NQEKPSNSESSLGAK.H	2	CCD55_HUMAN
Q9H0G5	2	3.6308	K.RNNEETVMSAR.D	3	CCD55_HUMAN
Q9H0G5	2	2.8557	K.HRLTEEGQEK.G	1	CCD55_HUMAN
Q9H0G5	2	4.9839	R.HLLNQAVGEEEVPK.C	2	CCD55_HUMAN
Coiled-coil domain-containing protein 56 - Homo sapiens (Human)					
Q9Y2R0	2	3.4321	K.RGEAPFAQR.I	2	CCD56_HUMAN
Coiled-coil domain-containing protein 58 - Homo sapiens (Human)					
Q4VC31	2	3.8804	K.WMQSELNVEEVVNDR.S	2	CCD58_HUMAN
Q4VC31	3	4.1805	R.TIDDRIVHELNTTVPTASFAGK.I	2	CCD58_HUMAN
Coiled-coil domain-containing protein 6 - Homo sapiens (Human)					
Q16204	3	5.3304	R.KLM#QLQHEKGELEQHLEQEQEFQVNK.L	1	CCDC6_HUMAN
Coiled-coil domain-containing protein 7 - Homo sapiens (Human)					
Q96M83	3	3.8248	K.IQEYPQITAQSGRLIEK.R	1	CCDC7_HUMAN
Coiled-coil domain-containing protein 71 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8IV32	1	2.1207	K.GPGAGPR.R	1	CCD71_HUMAN
Coiled-coil domain-containing protein 72 - Homo sapiens (Human)					
Q9Y2S6	3	4.3885	K.AAGKGPLATGGIKK.S	4	CCD72_HUMAN
Q9Y2S6	2	3.0536	K.GPLATGGIKK.S	1	CCD72_HUMAN
Q9Y2S6	1	2.4519	K.GPLATGGIK.K	2	CCD72_HUMAN
Q9Y2S6	2	3.3331	K.EMDEEDKAFKQK.Q	2	CCD72_HUMAN
Q9Y2S6	2	2.721	K.EM#DEEDKAFKQK.Q	1	CCD72_HUMAN
Q9Y2S6	1	2.7472	K.EMDEEDKAFK.Q	2	CCD72_HUMAN
Coiled-coil domain-containing protein 86 - Homo sapiens (Human)					
Q9H6F5	3	4.7019	K.KLNKEELPVIPK.G	1	CCD86_HUMAN
Q9H6F5	3	4.6356	R.ALVEFESNPEETREPGSPPSVQR.A	2	CCD86_HUMAN
Q9H6F5	2	3.5671	R.DTLALLQK.Q	2	CCD86_HUMAN
Q9H6F5	2	3.2828	R.KAEVVQVIR.N	2	CCD86_HUMAN
Coiled-coil domain-containing protein 9 - Homo sapiens (Human)					
Q9Y3X0	2	4.5446	R.SVEKENVAVESEK.N	3	CCDC9_HUMAN
Q9Y3X0	2	2.7669	R.KNEALIR.R	1	CCDC9_HUMAN
Q9Y3X0	2	2.7387	K.EGAASPAPETPQPTSPETSPK.E	1	CCDC9_HUMAN
Q9Y3X0	2	3.9539	K.MNEEMEKIAEYER.N	2	CCDC9_HUMAN
Q9Y3X0	3	5.3366	R.SVEKENVAVESEKNLGPSR.R	3	CCDC9_HUMAN
Q9Y3X0	2	3.4398	R.EGVLEPNPVR.N	3	CCDC9_HUMAN
Coiled-coil domain-containing protein 91 - Homo sapiens (Human)					
Q7Z6B0	2	2.8292	K.RLDQVIR.Q	1	CCD91_HUMAN
Q7Z6B0	2	2.8963	K.VSQEIQK.A	2	CCD91_HUMAN
Q7Z6B0	2	2.7062	R.KISQETVK.A	1	CCD91_HUMAN
Coiled-coil-helix-coiled-coil-helix domain-containing protein 3 - Homo sapiens (Human)					
Q9NX63	2	3.2715	K.AAEEVEAK.F	3	CHCH3_HUMAN
Q9NX63	2	4.1042	R.YSGAYGASVSDEELK.R	1	CHCH3_HUMAN
Coiled-coil-helix-coiled-coil-helix domain-containing protein 6 - Homo sapiens (Human)					
Q9BRQ6	2	3.674	R.SGSSGGQQPSGMK.E	2	CHCH6_HUMAN
Coiled-coil-helix-coiled-coil-helix domain-containing protein 8 - Homo sapiens (Human)					
Q9NYJ1	2	5.0633	R.VKKDDEEEDPLDQLISR.S	3	Q9NYJ1_HUMA
Q9NYJ1	2	3.1484	K.DCMSEQQAR.R	1	Q9NYJ1_HUMA
Coilin - Homo sapiens (Human)					
P38432	3	4.7568	K.LLELTSSYSPDVSDYKEGR.I	2	P80C_HUMAN
Cold-inducible RNA-binding protein - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q14011	2	4.4056	R.SSGGSYRDSYDSYATHNE.-	3	CIRBP_HUMAN
Q14011	2	3.7864	R.DSYDSYATHNE.-	9	CIRBP_HUMAN
Collagen alpha-1(I) chain precursor - Homo sapiens (Human)					
P02452	2	3.1469	R.EGAPGAEGSPGR.D	3	CO1A1_HUMAN
P02452	2	2.7846	R.GVQGPPGPAGPR.G	1	CO1A1_HUMAN
P02452	2	3.5411	R.GQAGVM#GFPGPK.G	8	CO1A1_HUMAN
P02452	2	2.7354	R.GPPGPM#GPPGLAGPPGESGR.E	1	CO1A1_HUMAN
P02452	3	4.3564	R.GPAGPQGPRGDKGETGEQGDR.G	3	CO1A1_HUMAN
P02452	1	3.0082	R.GPAGPQGPR.G	28	CO1A1_HUMAN
P02452	2	2.8516	K.GEAGPQGPR.G	10	CO1A1_HUMAN
P02452	2	5.6542	R.GFSGLDGAKGDAGPAGPK.G	7	CO1A1_HUMAN
P02452	1	2.5014	R.GFSGLDGAK.G	9	CO1A1_HUMAN
P02452	2	4.8302	R.GETGPAGPAGPVGPVGAR.G	3	CO1A1_HUMAN
P02452	2	3.545	K.GADGSPGKDGVR.G	12	CO1A1_HUMAN
P02452	2	3.1931	K.STGGISVPGPMGPSGPR.G	5	CO1A1_HUMAN
P02452	2	5.5649	K.DGEAGAQGPPGPAGPAGER.G	12	CO1A1_HUMAN
P02452	3	3.739	K.GDRGDAGPKGADGSPGK.D	1	CO1A1_HUMAN
P02452	2	4.7982	K.GSPGADGPAGAPGTPGPQGIAGQR.G	1	CO1A1_HUMAN
P02452	2	4.1196	K.GESGPSGPAGPTGAR.G	2	CO1A1_HUMAN
P02452	3	4.2392	K.QGPSGASGERGPPGPM#GPPGLAGPPGES	1	CO1A1_HUMAN
P02452	2	5.9967	K.SGDRGETGPAGPAGPVGPVGAR.G	14	CO1A1_HUMAN
P02452	2	2.9419	K.STGGISVPGPM#GPSGPR.G	3	CO1A1_HUMAN
Collagen alpha-1(III) chain precursor - Homo sapiens (Human)					
P02461	2	4.145	R.GPVGPSGPPGK.D	9	CO3A1_HUMAN
P02461	2	2.7021	R.GLPGPPGSNGNPGPPGPSGSPGK.D	1	CO3A1_HUMAN
Collagen alpha-1(V) chain precursor - Homo sapiens (Human)					
P20908	3	4.4907	R.FGGGGDAGSKGPM#VSAQESQAQAILQQAR	1	CO5A1_HUMAN
P20908	2	2.9511	R.GPAGPM#GLTGR.P	1	CO5A1_HUMAN
P20908	2	3.313	R.GPAGPMGLTGR.P	3	CO5A1_HUMAN
Collagen alpha-1(VI) chain precursor - Homo sapiens (Human)					
P12109	2	3.274	R.GDPGEAGPQGDQGR.E	1	CO6A1_HUMAN
P12109	2	3.6111	K.VFSVAITPDHLEPR.L	4	CO6A1_HUMAN
P12109	2	3.3965	R.DAEEAISQTIDTIVDM#IK.N	1	CO6A1_HUMAN
Collagen alpha-1(XII) chain precursor - Homo sapiens (Human)					
Q99715	3	4.0811	K.MIATDPDDTHAYNVADFESLSR.I	1	COCA1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Collagen alpha-1(XV) chain precursor [Contains: Endostatin - Homo sapiens (Human)					
P39059	3	5.7784	R.IILYYTEPGSHVSQEAAAFSVPVMTHR.W	3	COFA1_HUMAN
P39059	2	3.6721	R.NLVTAFSNM#DDMLQK.A	2	COFA1_HUMAN
P39059	2	4.5253	R.TADTAVTGLASPLSTGK.I	2	COFA1_HUMAN
P39059	2	5.3571	R.GGVLFAITDAFQK.V	5	COFA1_HUMAN
Collagen alpha-1(XVI) chain precursor - Homo sapiens (Human)					
Q07092	2	3.2113	R.DGQQGQTGLR.G	3	COGA1_HUMAN
Q07092	2	4.7322	R.DTQSNELIEINPQSEGK.V	1	COGA1_HUMAN
Collagen alpha-1(XVIII) chain precursor [Contains: Endostatin] - Homo sapiens (Human)					
P39060	3	4.893	R.GTDNEVAALQPPVVQLHDSNPYPR.R	7	COIA1_HUMAN
P39060	3	5.7042	R.ELLREETGAALKPR.L	6	COIA1_HUMAN
Collagen alpha-2(I) chain precursor - Homo sapiens (Human)					
P08123	2	5.7675	R.GEAGAAGPAGPAGPR.G	4	CO1A2_HUMAN
P08123	2	3.1962	R.PGPIGPAGAR.G	1	CO1A2_HUMAN
P08123	2	3.6499	R.GPAGPSGPAGK.D	17	CO1A2_HUMAN
P08123	2	3.8671	R.GPAGPSGPAGKDGR.T	5	CO1A2_HUMAN
P08123	2	3.4978	R.GPRGDQGPVGR.T	1	CO1A2_HUMAN
P08123	2	3.4847	R.GPSGPQGIR.G	13	CO1A2_HUMAN
P08123	1	2.2591	R.GVVGPQGAR.G	1	CO1A2_HUMAN
P08123	2	2.8503	R.GASGPAGVR.G	2	CO1A2_HUMAN
P08123	2	2.8542	R.GPSGPQGIRGDK.G	2	CO1A2_HUMAN
P08123	1	2.2277	R.AGVM#GPPGSR.G	2	CO1A2_HUMAN
P08123	3	4.7361	K.NGDKGHAGLAGAR.G	15	CO1A2_HUMAN
P08123	2	6.8429	K.HGNRGETGPSGPVGPAGAVGPR.G	7	CO1A2_HUMAN
P08123	2	3.83	K.GVGLGPGPMGLMGPR.G	4	CO1A2_HUMAN
P08123	2	3.7929	K.GVGLGPGPM#GLMGPR.G	4	CO1A2_HUMAN
P08123	2	3.3148	K.GVGLGPGPM#GLM#GPR.G	4	CO1A2_HUMAN
P08123	2	2.8097	K.GHAGLAGAR.G	3	CO1A2_HUMAN
P08123	2	4.8346	K.GEIGAVGNAGPAGPAGPR.G	4	CO1A2_HUMAN
P08123	2	4.478	R.GETGPSGPVGPAGAVGPR.G	5	CO1A2_HUMAN
Collagen alpha-2(IV) chain precursor [Contains: Canstatin] - Homo sapiens (Human)					
P08572	2	3.5002	R.SVSIGYLLVK.H	3	CO4A2_HUMAN
P08572	2	3.3353	R.GLPGEVLGAQPGPR.G	3	CO4A2_HUMAN
P08572	2	3.7264	R.GLDGYQGPDGPR.G	6	CO4A2_HUMAN
P08572	2	3.7859	R.GEQGFMGNTGPTGAVGDR.G	2	CO4A2_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P08572	2	3.2479	K.IAVQPGTVGPQGR.R	4	CO4A2_HUMAN
P08572	2	4.5456	R.GEQGFM#GNTGPTGAVGDR.G	4	CO4A2_HUMAN
Collagen alpha-2(V) chain precursor - Homo sapiens (Human)					
P05997	2	3.7822	K.SGLGSQVGLM#PGSVGPVGPR.G	2	CO5A2_HUMAN
Collagen alpha-2(VI) chain precursor - Homo sapiens (Human)					
P12110	2	3.2916	R.LFAVAPNQNLK.E	2	CO6A2_HUMAN
P12110	2	3.205	R.DIASTPHELYR.N	2	CO6A2_HUMAN
P12110	2	3.5703	K.NLQGISSFR.R	5	CO6A2_HUMAN
P12110	3	5.0019	R.YGGLHFSDQVEVFSPPGSDR.A	2	CO6A2_HUMAN
Collagen alpha-3(V) chain precursor - Homo sapiens (Human)					
P25940	3	4.7255	K.AAREDEEGDDSTM#GPDFR.A	3	CO5A3_HUMAN
P25940	2	3.667	K.AAREDEEGDDSTMGPDFR.A	2	CO5A3_HUMAN
P25940	2	3.0625	K.ALGVQGGQAGVPEGPGFCPQR.T	2	CO5A3_HUMAN
P25940	2	2.9516	K.RGPSGHMGR.E	4	CO5A3_HUMAN
P25940	2	4.081	R.EDEEGDDSTM#GPDFR.A	4	CO5A3_HUMAN
Collagen alpha-3(VI) chain precursor - Homo sapiens (Human)					
P12111	2	3.8407	K.LSDAGITPLFLTR.Q	1	CO6A3_HUMAN
P12111	2	4.147	K.SVEDAQDVSLALTQR.G	1	CO6A3_HUMAN
P12111	2	2.9263	K.IASNSATAFR.V	1	CO6A3_HUMAN
P12111	2	3.0234	R.GETGDDGRDGVGSEGR.R	1	CO6A3_HUMAN
Collagen triple helix repeat-containing protein 1 precursor - Homo sapiens (Human)					
Q96CG8	2	2.7081	R.DGFKGEKGECLR.E	1	CTHR1_HUMAN
Complement C1q subcomponent subunit C precursor - Homo sapiens (Human)					
P02747	2	3.1032	K.FQSVFTVTR.Q	1	C1QC_HUMAN
Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alpha chain; C3					
P01024	2	4.4517	R.IPIEDGSGEVVLSR.K	3	CO3_HUMAN
P01024	3	5.6196	R.TKKQELSEAEQATR.T	2	CO3_HUMAN
P01024	1	2.1285	R.SVQLTEK.R	1	CO3_HUMAN
P01024	2	3.765	R.ILLQGTPVAQMTEDAVDAER.L	1	CO3_HUMAN
P01024	3	6.159	R.LESEETMVLEAHDAQGDVPVTVTVHDFPGK.	2	CO3_HUMAN
P01024	2	4.7787	R.VPVAVQGEDTVQSLTQGDGVAK.L	2	CO3_HUMAN
P01024	3	4.2685	R.EGVQKEDIPPADLSDQVPDTESETR.I	3	CO3_HUMAN
P01024	3	4.0319	K.VQLSNDFDEYIMAIEQTIK.S	3	CO3_HUMAN
P01024	3	5.2525	K.VQLSNDFDEYIM#AIEQTIK.S	28	CO3_HUMAN
P01024	2	4.1901	K.SGSDEVQVGQQR.T	4	CO3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P01024	2	4.2141	K.QLYNVEATSYALLALLQLK.D	1	CO3_HUMAN
P01024	2	3.0486	K.KLVLSSEK.T	1	CO3_HUMAN
P01024	1	2.3818	R.QNQELK.V	1	CO3_HUMAN
P01024	2	3.056	R.EPGQDLVVLPLSITTDFIPSFR.L	2	CO3_HUMAN
Complement component 1 Q subcomponent-binding protein, mitochondrial precursor - Homo sapie					
Q07021	2	4.9411	K.M#SGGWELELNGTEAK.L	4	C1QBP_HUMAN
Q07021	2	5.1507	K.MSGGWELELNGTEAK.L	7	C1QBP_HUMAN
Q07021	3	5.4927	K.ALVLDCHYPEDEVGQEDEAESDIFSIR.E	4	C1QBP_HUMAN
Q07021	2	3.3815	K.AFVDFLSDEIKEER.K	2	C1QBP_HUMAN
Complement factor H precursor - Homo sapiens (Human)					
P08603	2	2.8156	K.SCDIPVFMNAR.T	1	CFAH_HUMAN
P08603	2	3.8114	K.SSNLIILEEHLK.N	2	CFAH_HUMAN
P08603	3	5.2904	R.DTSCVNPPTVQNAYIVSR.Q	2	CFAH_HUMAN
P08603	2	3.3292	R.SSQESYAHGTK.L	7	CFAH_HUMAN
P08603	2	2.7915	R.TKNDFTWFK.L	1	CFAH_HUMAN
P08603	2	3.8566	K.EQVQSCGPPPELLNGNVK.E	1	CFAH_HUMAN
Condensin complex subunit 3 - Homo sapiens (Human)					
Q9BPX3	2	4.4529	K.SKLNLAQFLNEDLS.-	2	CND3_HUMAN
Condensin-II complex subunit G2 - Homo sapiens (Human)					
Q86XI2	2	2.7305	K.EDTGKTAFVM#LLR.R	1	CNDG2_HUMAN
Conserved nuclear protein NHN1 - Homo sapiens (Human)					
Q86VM9	2	2.7797	R.DSSTQPPK.S	1	Q86VM9_HUMA
Q86VM9	2	2.8258	K.AVEDAIAR.K	1	Q86VM9_HUMA
COP9 signalosome complex subunit 3 - Homo sapiens (Human)					
Q9UNS2	2	3.2618	K.SMGSQEDDSGNKPSSYS.-	2	CSN3_HUMAN
Q9UNS2	2	3.8375	K.SM#GSQEDDSGNKPSSYS.-	3	CSN3_HUMAN
COP9 signalosome complex subunit 4 - Homo sapiens (Human)					
Q9BT78	2	2.9634	K.LYNNITFEELGALLEIPAAK.A	1	CSN4_HUMAN
Copper chaperone for superoxide dismutase - Homo sapiens (Human)					
O14618	2	3.6314	R.SLIIDEGEDDLGRGGHPLSK.I	1	CCS_HUMAN
O14618	3	4.0438	K.GM#GSGQLQNLGAAVAILGGPGTVQGVVR.F	1	CCS_HUMAN
O14618	2	2.8578	R.FLQLTPER.C	1	CCS_HUMAN
O14618	2	4.6663	R.MEDEQLKVWDVIGR.S	3	CCS_HUMAN
O14618	2	3.0239	R.SAGLFQNPK.Q	1	CCS_HUMAN
Copper transport protein ATOX1 - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O00244	3	4.3084	K.LGGVKYDIDLPNKK.V	3	ATOX1_HUMAN
O00244	2	2.7502	K.YDIDLPNKK.V	2	ATOX1_HUMAN
O00244	2	3.6366	K.TGKTVSYLGLE.-	4	ATOX1_HUMAN
Cordon-bleu protein-like 1 - Homo sapiens (Human)					
Q53SF7	2	3.5811	R.QSSLTFQSSDPEQMR.Q	2	COBL1_HUMAN
Q53SF7	2	3.6638	R.DQTASAPATPLVNK.H	4	COBL1_HUMAN
Q53SF7	3	5.2399	R.ASNAQAKPSSFFLQMQK.R	2	COBL1_HUMAN
Q53SF7	3	4.7547	R.DTGTAPFAPNLEEINNILESK.F	1	COBL1_HUMAN
Q53SF7	2	3.8654	R.LSHSM#SPDAQDGH.-	3	COBL1_HUMAN
Q53SF7	2	4.7735	R.SNTISKPYISNTLPSDAPK.K	2	COBL1_HUMAN
Q53SF7	3	4.0702	R.SRLSHSM#SPDAQDGH.-	1	COBL1_HUMAN
Q53SF7	3	3.9811	R.TEHQVPSSVSSPDDAMVSPLKPAPK.M	1	COBL1_HUMAN
Q53SF7	2	3.7056	R.VSGHYVTSAAAK.S	2	COBL1_HUMAN
Q53SF7	2	5.1797	R.AGSLQLSSMSAGNSSLR.R	3	COBL1_HUMAN
Q53SF7	2	2.9501	R.VTIPSNTISVNGR.S	1	COBL1_HUMAN
Q53SF7	3	3.7255	K.ETAIQTEDSAISESPEEPLPNLKPKPNLR.T	1	COBL1_HUMAN
Q53SF7	2	2.9577	R.AGSLQLSSM#SAGNSSLR.R	1	COBL1_HUMAN
Q53SF7	3	4.0334	K.EAERDMLPSPEQTLSPLSK.M	1	COBL1_HUMAN
Q53SF7	2	3.0041	K.IPPHQSDENSR.V	3	COBL1_HUMAN
Q53SF7	2	3.2285	K.KTEINVEGVAK.N	2	COBL1_HUMAN
Q53SF7	2	2.9121	K.LNQPSAEK.T	2	COBL1_HUMAN
Q53SF7	1	2.1196	K.NSTASYLK.N	1	COBL1_HUMAN
Q53SF7	1	2.2652	K.NYPLYR.Q	2	COBL1_HUMAN
Q53SF7	2	3.605	K.PSSFFLQMQK.R	2	COBL1_HUMAN
Q53SF7	2	3.7445	K.RVSGHYVTSAAAK.S	2	COBL1_HUMAN
Q53SF7	3	3.8346	K.SM#SVDETDKSPCEAGR.V	1	COBL1_HUMAN
Q53SF7	3	3.7001	K.ELTNKEAERDMLPSPEQTLSPLSK.M	1	COBL1_HUMAN
Core histone macro-H2A.1 - Homo sapiens (Human)					
O75367	2	3.726	K.AASADSTTEGTPADGFTVLSTK.S	1	H2AY_HUMAN
Coronin-1A - Homo sapiens (Human)					
P31146	2	3.341	R.AAPEASGTPSSDAVSR.L	1	COR1A_HUMAN
P31146	2	3.163	R.DAGPLLISLK.D	2	COR1A_HUMAN
P31146	2	3.9307	R.KLQATVQELQK.R	2	COR1A_HUMAN
P31146	3	4.0193	R.KSDLFQEDLYPPTAGPDPALTAEEWLGGR.D	1	COR1A_HUMAN
Coronin-1B - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BR76	2	5.6276	R.AGEAGKLEEVMQELR.A	5	COR1B_HUMAN
Q9BR76	3	3.7683	R.KSDLFQDDLYPDTAGPEAALEAEEWVSGR.D	1	COR1B_HUMAN
Coronin-1C - Homo sapiens (Human)					
Q9ULV4	2	4.6324	K.KTTDTASVQNEAK.L	2	COR1C_HUMAN
Q9ULV4	2	3.2934	R.ISKLEQQMAK.I	1	COR1C_HUMAN
Q9ULV4	2	3.0091	R.ISKLEQQM#AK.I	1	COR1C_HUMAN
Cortactin-binding protein 2 - Homo sapiens (Human)					
Q8WZ74	2	3.6144	K.TVASTPSSLPQGNR.V	3	CTTB2_HUMAN
Craniofacial development protein 1 - Homo sapiens (Human)					
Q9UEE9	2	3.9874	K.VFDFAGEEVR.V	4	CFDP1_HUMAN
Q9UEE9	2	3.4243	R.SSGMSSLLGK.I	1	CFDP1_HUMAN
Q9UEE9	2	3.1549	R.KAQSIPAR.K	2	CFDP1_HUMAN
Q9UEE9	4	5.4889	K.SKLDWESFKEEEGIGEELAIHNR.G	1	CFDP1_HUMAN
Q9UEE9	3	5.741	K.KKEDELWASFLNDVGPK.S	5	CFDP1_HUMAN
Q9UEE9	2	2.7864	K.EVDATSKEAK.S	1	CFDP1_HUMAN
Q9UEE9	2	4.8471	K.EKPQANVPSALPSLPAGSGLK.R	3	CFDP1_HUMAN
Q9UEE9	2	3.0791	K.EDEVDGEEQTQK.T	2	CFDP1_HUMAN
Q9UEE9	2	4.0014	K.AEELEKPKETEK.V	2	CFDP1_HUMAN
Q9UEE9	2	2.7283	R.SSGM#SSLLGK.I	1	CFDP1_HUMAN
CREB-regulated transcription coactivator 3 - Homo sapiens (Human)					
Q6UUV7	2	3.0237	R.AFEQLMTDLTLSR.V	1	Q6UUV7_HUMA
Crk-like protein - Homo sapiens (Human)					
P46109	2	2.9956	R.DSSTCPGDYVLSVSENSR.V	1	CRKL_HUMAN
P46109	2	2.7602	R.VGMIPVPYVEK.L	1	CRKL_HUMAN
P46109	2	2.7685	R.SAWYMGPVSR.Q	1	CRKL_HUMAN
P46109	2	4.8791	R.MNINGQWEGEVNGR.K	2	CRKL_HUMAN
P46109	2	3.2083	R.M#NINGQWEGEVNGR.K	1	CRKL_HUMAN
P46109	3	4.1998	R.KGLFPFTHVK.I	2	CRKL_HUMAN
P46109	2	2.8188	R.HGMFLVR.D	2	CRKL_HUMAN
P46109	2	4.8338	K.TALALEVGDIVK.V	3	CRKL_HUMAN
P46109	2	4.2672	K.IHYLDTTTLIEPAPR.Y	2	CRKL_HUMAN
P46109	3	3.7147	K.IGDQEFDHLPALLEFYK.I	1	CRKL_HUMAN
P46109	3	4.1515	R.FKIGDQEFDHLPALLEFYK.I	1	CRKL_HUMAN
CRSP complex subunit 7 - Homo sapiens (Human)					
O95402	3	3.8907	K.ASVLQQLDRVDETPGPPHPK.G	1	CRSP7_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O95402	3	4.2847	K.GSVPSPSPRPQALDATQVPSPLPLAQPSTPP	1	CRSP7_HUMAN
CTD small phosphatase-like protein 2 - Homo sapiens (Human)					
Q3ZTU1	3	4.4707	R.KASQQSNQIQTQR.T	2	Q3ZTU1_HUMA
CTTNBP2 N-terminal-like protein - Homo sapiens (Human)					
Q9P2B4	2	3.874	R.DLSPTLIDNSAAK.Q	7	CT2NL_HUMAN
Q9P2B4	2	3.3189	R.NLANTANPR.G	3	CT2NL_HUMAN
Q9P2B4	3	5.9833	R.FTSQQGPIKPVSPNSSPFGTDYR.N	4	CT2NL_HUMAN
Q9P2B4	2	3.1523	R.GDTSHSPTPGK.V	5	CT2NL_HUMAN
C-type lectin domain family 14 member A precursor - Homo sapiens (Human)					
Q86T13	2	3.4926	R.DRAEGALLAESPLGSSDA.-	1	CLC14_HUMAN
Cubilin precursor - Homo sapiens (Human)					
O60494	2	3.3958	R.FVTDGSVTASGFR.L	2	CUBN_HUMAN
O60494	1	2.1772	R.SSGDSMFIK.L	1	CUBN_HUMAN
O60494	2	3.4843	R.SPFFPNVYPGER.T	4	CUBN_HUMAN
O60494	2	3.9436	R.SPENPMQVSSTGNELAIR.F	2	CUBN_HUMAN
O60494	2	3.3786	R.VGDADGPLMWR.L	2	CUBN_HUMAN
O60494	2	3.1043	K.YDDCEGGSVAR.C	1	CUBN_HUMAN
O60494	2	3.2484	K.SSSTENHGFMAK.F	1	CUBN_HUMAN
O60494	2	2.7678	K.SITNSVWIR.F	1	CUBN_HUMAN
O60494	2	2.9857	K.LRTDEGQQGR.G	2	CUBN_HUMAN
O60494	2	3.3132	R.ITLMFNNLR.L	1	CUBN_HUMAN
O60494	3	4.0843	R.KFQGLQQTVDKK.V	3	CUBN_HUMAN
Cullin-4A - Homo sapiens (Human)					
Q13619	2	3.1022	R.KGSFSALVGR.T	2	CUL4A_HUMAN
Cullin-5 - Homo sapiens (Human)					
Q93034	2	2.8267	K.SNVEDSIVR.K	2	CUL5_HUMAN
Cyclin-dependent kinase 6 inhibitor - Homo sapiens (Human)					
P42773	2	3.1079	R.VVEFLVK.H	2	CDN2C_HUMAN
Cyclin-dependent kinase inhibitor 1B - Homo sapiens (Human)					
P46527	2	4.2299	R.TEENVSDGSPNAGSVEQTPK.K	2	CDN1B_HUMAN
P46527	2	3.516	R.NLFGPVDHEELTR.D	2	CDN1B_HUMAN
P46527	3	5.3426	R.ANRTEENVSDGSPNAGSVEQTPK.K	7	CDN1B_HUMAN
P46527	2	3.0187	R.DMEEASQR.K	1	CDN1B_HUMAN
Cyclin-dependent kinase-like 5 - Homo sapiens (Human)					
O76039	2	3.7275	K.TYQASSQPGSTSK.D	3	CDKL5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Cyclin-T1 - Homo sapiens (Human)					
O60563	3	3.8108	K.HAEELAAQKR.Q	1	CCNT1_HUMAN
O60563	2	2.8281	K.HNSVEDSVTK.S	2	CCNT1_HUMAN
O60563	3	4.5536	K.HSSQTSNLAHK.T	2	CCNT1_HUMAN
Cyclin-T2 - Homo sapiens (Human)					
O60583	2	3.6774	R.SPVGLSSDGISSSSSSSR.K	1	CCNT2_HUMAN
Cystatin-B - Homo sapiens (Human)					
P04080	2	3.3679	K.AKHDELTYF.-	3	CYTB_HUMAN
P04080	3	3.9867	R.VFQSLPHENKPLTLSNYQTNK.A	2	CYTB_HUMAN
P04080	2	4.4867	K.SQVVAGTNYFIK.V	5	CYTB_HUMAN
Cysteine and glycine-rich protein 1 - Homo sapiens (Human)					
P21291	2	4.691	K.GFGFGQGAGALVHSE.-	14	CSRP1_HUMAN
P21291	2	4.1884	K.GLESTTLADKDGEIYCK.G	1	CSRP1_HUMAN
Cysteine and glycine-rich protein 2 - Homo sapiens (Human)					
Q16527	2	5.2824	K.GFGYGQGAGALVHAQ.-	8	CSRP2_HUMAN
Q16527	2	4.1435	K.GYGYGQGAGTLNMDR.G	1	CSRP2_HUMAN
Q16527	2	3.8468	K.SLESTTLTEKEGEIYCK.G	2	CSRP2_HUMAN
Cysteine-rich PDZ-binding protein. - Homo sapiens (Human)					
Q9P021	2	4.5695	K.KLGTVITPDTWK.D	2	Q9P021_HUMAN
Cysteine-rich protein 1 - Homo sapiens (Human)					
P50238	2	3.4738	R.GGAESHTFK.-	16	CRIP1_HUMAN
P50238	1	2.1049	K.EVYFAER.V	1	CRIP1_HUMAN
P50238	3	5.058	K.TLTSGGHAEHEGKPYCNHPCYAAMFGPK.G	1	CRIP1_HUMAN
Cysteine-rich protein 2 - Homo sapiens (Human)					
P52943	2	5.1918	K.PLAEGPQVTGPIEVPAAR.A	4	CRIP2_HUMAN
P52943	2	3.2102	R.KASGPPKGPSR.A	6	CRIP2_HUMAN
P52943	1	2.201	K.TVYFAEK.V	3	CRIP2_HUMAN
P52943	1	2.2656	K.VYFAEK.V	2	CRIP2_HUMAN
P52943	3	4.0408	K.TLTPGGHAEHDGQPYCHKPCYGILFGPK.G	1	CRIP2_HUMAN
P52943	3	3.7871	K.GVNTGAVGSYIYDRDPEGKVQP.-	4	CRIP2_HUMAN
P52943	3	6.9563	K.GVNIGGAGSYIYEKPLAEGPQVTGPIEVPAAR	3	CRIP2_HUMAN
P52943	2	3.7969	K.GVNIGGAGSYIYEK.P	1	CRIP2_HUMAN
P52943	2	2.9465	K.ASGPPKGPSR.A	1	CRIP2_HUMAN
P52943	2	2.7212	K.KVYFAEK.V	1	CRIP2_HUMAN
P52943	2	5.1114	R.ASSVTTFTGEPNTCPR.C	13	CRIP2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Cysteine-rich with EGF-like domain protein 2 precursor - Homo sapiens (Human)					
Q6UXH1	2	3.1604	K.TLSKYESSEIR.L	2	CREL2_HUMAN
Q6UXH1	3	3.9296	R.GLVDKFNQGMVDTAKK.N	1	CREL2_HUMAN
Cystin-1 - Homo sapiens (Human)					
Q717R9	2	3.6687	R.SPESLPAGPGAAALEGGTR.R	2	CYS1_HUMAN
Cytochrome b-245 heavy chain - Homo sapiens (Human)					
P04839	2	3.2618	K.TIASQHPNTR.I	2	CY24B_HUMAN
Cytochrome b-245 light chain - Homo sapiens (Human)					
P13498	3	4.2631	R.KKPSEEEAAAAAGGPPGGPQVNPIPVTDEVV	1	CY24A_HUMAN
Cytochrome b5 - Homo sapiens (Human)					
P00167	3	3.9066	R.EQAGGDATENFEDVGHSTDAR.E	1	CYB5_HUMAN
P00167	2	2.8748	K.YYTLEEIQK.H	1	CYB5_HUMAN
P00167	2	4.3466	K.TFIIGELHPDDRPK.L	2	CYB5_HUMAN
P00167	2	3.7594	K.FLEEHPGGEEVLR.E	2	CYB5_HUMAN
P00167	4	5.1622	K.FLEEHPGGEEVLREQAGGDATENFEDVGHS	2	CYB5_HUMAN
Cytochrome c - Homo sapiens (Human)					
P99999	2	3.699	K.TGQAPGYSYTAANK.N	2	CYC_HUMAN
P99999	3	5.5876	R.KTGQAPGYSYTAANKNK.G	1	CYC_HUMAN
Cytochrome c oxidase assembly protein COX19 - Homo sapiens (Human)					
Q49B96	2	3.6201	K.LGFGDLTSGK.S	2	COX19_HUMAN
Q49B96	2	2.9706	R.KLMLQEPLEK.L	1	COX19_HUMAN
Cytochrome c oxidase subunit 5A, mitochondrial precursor - Homo sapiens (Human)					
P20674	1	2.1392	K.IIDAALR.A	1	COX5A_HUMAN
P20674	3	4.1284	R.WVTYFNKPDIDAWELRK.G	1	COX5A_HUMAN
P20674	2	2.8853	R.ILEVVKDK.A	1	COX5A_HUMAN
P20674	2	4.023	K.GINTLVTYDM#VPEPK.I	3	COX5A_HUMAN
P20674	3	3.7739	R.KGINTLVTYDMVPEPK.I	2	COX5A_HUMAN
P20674	2	3.0035	K.GINTLVTYDMVPEPK.I	1	COX5A_HUMAN
Cytochrome c oxidase subunit 5B, mitochondrial precursor - Homo sapiens (Human)					
P10606	2	3.5209	K.GLDPYNVLAPK.G	4	COX5B_HUMAN
P10606	2	4.1731	K.KGLDPYNVLAPK.G	5	COX5B_HUMAN
P10606	1	2.1851	R.EIMLAAK.K	1	COX5B_HUMAN
Cytochrome c oxidase subunit VIb isoform 1 - Homo sapiens (Human)					
P14854	2	2.8786	K.GGDISVCEWYQR.V	1	CX6B1_HUMAN
P14854	2	2.9009	K.TAPFDSRFPNQNQTR.N	1	CX6B1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Cytochrome P450 19A1 - Homo sapiens (Human)					
P11511	2	3.0226	K.NMLEM#IFTPRNSDR.C	1	CP19A_HUMAN
Cytoplasmic dynein 1 intermediate chain 2 - Homo sapiens (Human)					
Q13409	2	2.7434	K.SELKAELER.K	1	DC1I2_HUMAN
Q13409	2	4.07	K.SVSTPSEAGSQDSGDGAVGSR.T	3	DC1I2_HUMAN
Q13409	2	3.397	K.EAVAPVQEESDLEK.K	2	DC1I2_HUMAN
Q13409	2	3.7499	R.TLAEINANRADAEEEAATRIPA.-	1	DC1I2_HUMAN
Cytoplasmic dynein 1 light intermediate chain 1 - Homo sapiens (Human)					
Q9Y6G9	2	5.0295	K.AGATSEGVLANFFNSLLSK.K	7	DC1L1_HUMAN
Q9Y6G9	3	4.7349	K.KTGSPGGPGVSGGSPAGGAGGGSSGLPPS	2	DC1L1_HUMAN
Q9Y6G9	3	6.7694	K.KTGSPGGPGVSGGSPAGGAGGGSSGLPPS	2	DC1L1_HUMAN
Q9Y6G9	2	4.6013	R.KPVTVSPTTPTSPTEGEAS.-	5	DC1L1_HUMAN
Q9Y6G9	2	4.0381	R.SVSSNVASVSPIPAGSK.K	1	DC1L1_HUMAN
Cytoplasmic dynein 1 light intermediate chain 2 - Homo sapiens (Human)					
O43237	3	5.0785	K.KPDPNIKNNAASEGVLASFFNSLLSK.K	1	DC1L2_HUMAN
O43237	2	4.2451	K.TVLSNVQEELDR.M	7	DC1L2_HUMAN
O43237	2	4.039	R.KPDSM#VTNSSTENEA.-	3	DC1L2_HUMAN
O43237	2	4.2765	R.KPDSMVTNSSTENEA.-	5	DC1L2_HUMAN
O43237	2	3.3693	R.GGPASVPSSSPGTSVK.K	1	DC1L2_HUMAN
O43237	2	3.9653	K.TGSPGSPGAGGVQSTAK.K	2	DC1L2_HUMAN
O43237	3	5.4091	K.NNAASEGVLASFFNSLLSK.K	1	DC1L2_HUMAN
O43237	2	5.3082	K.KTGSPGSPGAGGVQSTAK.K	12	DC1L2_HUMAN
O43237	3	4.9858	K.KTGSPGSPGAGGVQSTAKK.S	3	DC1L2_HUMAN
O43237	2	3.2283	K.TGSPGSPGAGGVQSTAKK.S	1	DC1L2_HUMAN
Cytoplasmic protein NCK2 - Homo sapiens (Human)					
O43639	2	3.2896	R.IYDLNIPAFVK.F	2	NCK2_HUMAN
O43639	2	2.9031	K.GSLVKNLKDTLGLGK.T	2	NCK2_HUMAN
O43639	2	2.7687	R.DASPTPSTDAEYPANGSGADR.I	1	NCK2_HUMAN
Cytoskeleton-associated protein 4 - Homo sapiens (Human)					
Q07065	2	5.0126	K.GAHPSGGADDVAK.K	5	CKAP4_HUMAN
Cytoskeleton-associated protein 5 - Homo sapiens (Human)					
Q14008	2	3.5768	R.KGPAEDMSSK.L	1	CKAP5_HUMAN
Cytosol aminopeptidase - Homo sapiens (Human)					
P28838	2	4.5181	K.GVLFASGQNLAR.Q	3	AMPL_HUMAN
P28838	3	4.6463	R.TFYGLHQDFPSVVLVGLGK.K	2	AMPL_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Cytosolic non-specific dipeptidase - Homo sapiens (Human)					
Q96KP4	2	4.7516	K.LPDGSEIPLPPILLGR.L	1	CNDP2_HUMAN
Cytospin-A - Homo sapiens (Human)					
Q69YQ0	2	3.4604	K.TPPAAAVSPM#QR.H	1	CYTSA_HUMAN
DDI1 homolog 2 - Homo sapiens (Human)					
Q5TDH0	2	2.7454	R.VLVEQQQDR.A	1	DDI2_HUMAN
Q5TDH0	2	4.1373	R.NPPLAEALLSGDLEK.F	1	DDI2_HUMAN
D-dopachrome decarboxylase - Homo sapiens (Human)					
P30046	2	3.7786	R.FFPLESWQIGK.I	3	DOPD_HUMAN
DEAD box protein 23 - Homo sapiens (Human)					
Q9BUQ8	3	3.9042	K.KAEEEAEAKPK.F	2	DDX23_HUMAN
Q9BUQ8	3	5.0899	R.MERETNGNEDEEGRQK.I	2	DDX23_HUMAN
Death-associated protein 1 - Homo sapiens (Human)					
P51397	2	3.3281	R.IVQKHPHTGDTK.E	4	DAP1_HUMAN
P51397	2	2.7349	R.TQHIQQPR.K	1	DAP1_HUMAN
Death-inducer obliterator 1 - Homo sapiens (Human)					
Q9BTC0	2	2.7712	R.GGAPFQFGGQR.R	1	DIDO1_HUMAN
Q9BTC0	2	2.7442	R.QAGPAPAAATAASK.K	1	DIDO1_HUMAN
Q9BTC0	2	2.9453	K.VGGSQPPFQGQR.E	1	DIDO1_HUMAN
Q9BTC0	2	2.7531	K.STAPLLDVFSSMLK.D	1	DIDO1_HUMAN
Q9BTC0	2	3.7774	K.IFQPVIEAPGASK.C	1	DIDO1_HUMAN
Q9BTC0	2	3.0855	R.SGPQSASTAVK.E	2	DIDO1_HUMAN
Dedicator of cytokinesis protein 1 - Homo sapiens (Human)					
Q14185	3	4.7402	K.GSVADYGNLMENQDLLGSPTPPPPPPHQR.H	2	DOCK1_HUMAN
Q14185	2	3.2751	K.SQVM#NVIGSER.R	2	DOCK1_HUMAN
Q14185	2	2.8623	K.SQVMNVIGSER.R	1	DOCK1_HUMAN
Q14185	2	2.8693	R.KQTSVDSGIVQ.-	3	DOCK1_HUMAN
Q14185	2	5.5337	R.RFSVSPSSPSSQQTPPPVTPR.A	3	DOCK1_HUMAN
Dedicator of cytokinesis protein 11 - Homo sapiens (Human)					
Q5JSL3	2	2.8317	R.EMLWGSSTQLASDGSPK.G	1	Q5JSL3_HUMAN
Dedicator of cytokinesis protein 2 - Homo sapiens (Human)					
Q92608	2	3.9052	K.SAEEGKQIPDSLSTDL.-	1	DOCK2_HUMAN
Dedicator of cytokinesis protein 5 - Homo sapiens (Human)					
Q9H7D0	2	4.0694	R.KSGIPTSEPGSQ.-	3	DOCK5_HUMAN
Q9H7D0	3	3.9971	R.LSPFHGSSPPQSTPLSPPPLTPK.A	1	DOCK5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Dedicator of cytokinesis protein 6 - Homo sapiens (Human)					
Q96HP0	2	2.7508	K.SISSSNPDLAVAPGSVDDEVSR.I	1	DOCK6_HUMAN
Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial precursor - Homo sapiens (Human)					
P30038	2	2.8696	K.STGSIVGQQPFGGAR.A	1	AL4A1_HUMAN
P30038	2	3.1007	R.NAAGNFYINDK.S	1	AL4A1_HUMAN
Dematin - Homo sapiens (Human)					
Q08495	2	4.0721	R.LQKQPLTSPGSVSPSR.D	1	DEMA_HUMAN
Q08495	4	4.867	K.AILDIERPDLM#IYEPHFTYSLLEHVELPR.S	1	DEMA_HUMAN
Q08495	2	2.8502	K.MDNQVLGYK.D	1	DEMA_HUMAN
Q08495	2	4.306	K.MDNQVLGYKDLAAIPK.D	6	DEMA_HUMAN
Q08495	2	3.3271	R.DSSVPGSPSSIVAK.M	3	DEMA_HUMAN
Density-regulated protein - Homo sapiens (Human)					
O43583	2	3.0837	K.LDADYPLR.V	2	DENR_HUMAN
O43583	3	4.0229	K.WPEVDDDSIEDLGEVKK.-	2	DENR_HUMAN
Deoxyhypusine hydroxylase - Homo sapiens (Human)					
Q9BU89	2	3.94	R.AFQYADGLEQLR.G	2	DOHH_HUMAN
Q9BU89	4	5.0801	R.RLEWLQQHGGEPAAGPYLSVDPAPPAEERD	1	DOHH_HUMAN
Deoxynucleotidyltransferase terminal-interacting protein 2 - Homo sapiens (Human)					
Q5TFJ4	1	2.1171	K.QQM#QQEKSSER.R	1	Q5TFJ4_HUMAN
Q5TFJ4	2	3.7803	K.ASIQAASAESSGQK.S	2	Q5TFJ4_HUMAN
Q5TFJ4	3	4.7666	K.ESYTEEIVSEAESHVSGISR.I	1	Q5TFJ4_HUMAN
Q5TFJ4	2	2.9914	K.LYTSAPNTSQGK.D	1	Q5TFJ4_HUMAN
Q5TFJ4	3	4.5161	K.NLSELQDTSLQQLVSQR.H	2	Q5TFJ4_HUMAN
Q5TFJ4	2	2.862	K.QIVGTPVNSEDSDTR.Q	1	Q5TFJ4_HUMAN
Deoxyribonuclease-2-alpha precursor - Homo sapiens (Human)					
O00115	2	3.7355	R.ALINSPEGAVGR.S	5	DNS2A_HUMAN
O00115	2	3.7552	K.YLDESSGGWR.D	4	DNS2A_HUMAN
DERPC - Homo sapiens (Human)					
Q71E72	2	3.3284	R.GAGSSAFSQSSGTLASNPATFQR.S	1	Q71E72_HUMAN
Q71E72	2	3.1397	R.GGGPM#GPGSGPNLR.A	2	Q71E72_HUMAN
Q71E72	2	3.7056	R.GGGPMGPGSGPNLR.A	3	Q71E72_HUMAN
Q71E72	2	2.7691	R.LGGLPGPGPMSNPR.A	1	Q71E72_HUMAN
Desmin - Homo sapiens (Human)					
P17661	3	4.5172	R.RIESLNEEIAFLK.K	2	DESM_HUMAN
P17661	2	4.5185	R.TFGGAPGFPLGSPLSSPVFPR.A	3	DESM_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P17661	2	3.1935	R.TSGGAGGLGSLR.A	2	DESM_HUMAN
Desmocollin-2 precursor - Homo sapiens (Human)					
Q02487	2	2.974	R.GPGVDQEPR.N	1	DSC2_HUMAN
Desmoglein-2 precursor - Homo sapiens (Human)					
Q14126	3	4.4557	K.SLQEANAEKVTQEIVTER.S	2	DSG2_HUMAN
Q14126	2	4.576	R.ILDVNDNIPVVENK.V	3	DSG2_HUMAN
Q14126	3	5.2906	R.ERESFLAPSSGVQPTLAMPNIAVGQNVTVTE	1	DSG2_HUMAN
Q14126	2	3.621	R.ATQFTGATGAIMTTETTK.T	2	DSG2_HUMAN
Q14126	2	3.2351	K.IDINKEIEQR.Q	1	DSG2_HUMAN
Q14126	2	3.8642	K.AASYTEEDENHTAK.D	1	DSG2_HUMAN
Q14126	2	3.8342	R.DGNGEVTDKPVK.Q	1	DSG2_HUMAN
Desmuslin - Homo sapiens (Human)					
O15061	2	3.3574	R.EVPVYIGEDSTIAR.E	1	DMN_HUMAN
O15061	3	4.9235	K.VALLYLDNEEEENDGHWF.-	2	DMN_HUMAN
O15061	2	3.8037	R.TQEAGALGVSDR.G	2	DMN_HUMAN
O15061	2	3.142	R.SPAPGSPDEEGGAEAPAAGIR.F	1	DMN_HUMAN
O15061	2	3.13	R.YSWQDEIVQGTR.R	3	DMN_HUMAN
O15061	2	4.0423	R.SGEFHAEPTVIEK.E	2	DMN_HUMAN
O15061	3	3.89	R.QFHAEKEIIFQGPISAAGK.V	1	DMN_HUMAN
O15061	2	3.9311	R.MREELSALTR.E	4	DMN_HUMAN
O15061	2	3.1287	R.HVTLGPGQSPLSR.E	1	DMN_HUMAN
O15061	2	3.1026	R.EVPISLEVSQDR.R	2	DMN_HUMAN
O15061	2	4.307	R.ELYIPSGESEVAGGASHSSGQR.T	2	DMN_HUMAN
O15061	2	3.2373	R.EGQGGPGSVSVDVK.K	2	DMN_HUMAN
O15061	2	3.9149	R.DKVAAGASESTR.S	2	DMN_HUMAN
O15061	2	3.7757	R.DADSRNDQAVGVSFK.A	3	DMN_HUMAN
O15061	2	3.4267	K.VGDYFATEESVGTQTSVR.Q	1	DMN_HUMAN
O15061	3	4.134	K.SAEQMIGDIINLGLK.G	1	DMN_HUMAN
O15061	3	5.6405	K.VVKPLDVPAPSLEGDLGSTHWK.E	3	DMN_HUMAN
Deubiquitinating protein VCIP135 - Homo sapiens (Human)					
Q96JH7	2	3.3394	K.TEPSVFTASSSNSELIR.I	2	VCIP1_HUMAN
Q96JH7	2	2.7332	R.DGPSSAPATPTK.A	1	VCIP1_HUMAN
Q96JH7	2	4.2289	R.DQSTEQSPSDLPQR.K	4	VCIP1_HUMAN
Development and differentiation-enhancing factor 2 - Homo sapiens (Human)					
O43150	2	3.2284	K.VQTASSANTLWK.T	2	DDEF2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43150	2	3.8784	R.VTSTNPLTPTPPPPVAK.T	5	DDEF2_HUMAN
O43150	2	2.703	K.GAFPVSFVHFIAD.-	1	DDEF2_HUMAN
Developmentally-regulated GTP-binding protein 1 - Homo sapiens (Human)					
Q9Y295	2	4.368	K.IAEIEAEMAR.T	2	DRG1_HUMAN
Differentially expressed in FDCP 6 - Homo sapiens (Human)					
Q9H4E7	2	2.8209	R.IKELEEMQQR.L	1	DEFI6_HUMAN
Q9H4E7	2	3.0711	R.TPSPNSNEQQK.S	3	DEFI6_HUMAN
Dihydropyrimidinase - Homo sapiens (Human)					
Q14117	2	3.3416	R.VTKEDATAGTR.K	1	DPYS_HUMAN
Q14117	3	4.0049	R.VTKEDATAGTRK.Q	2	DPYS_HUMAN
Dihydropyrimidinase-like 2 - Homo sapiens (Human)					
Q86U75	3	4.0355	K.AM#EGIFIKPSVEPSAGHDEL.-	2	Q86U75_HUMAN
Q86U75	3	4.1216	K.AMEGIFIKPSVEPSAGHDEL.-	1	Q86U75_HUMAN
Dihydropyrimidinase-related protein 2 - Homo sapiens (Human)					
Q16555	2	2.7608	R.GLYDGPVCEVSVTPK.T	1	DPYL2_HUMAN
Q16555	3	5.424	R.NLHQSGFSLSGAQIDDNIPR.R	3	DPYL2_HUMAN
DIP2B protein - Homo sapiens (Human)					
Q6B011	2	3.8064	R.NQTPAPSAAQTSAPSK.Y	2	Q6B011_HUMAN
Q6B011	3	5.7628	R.SKLLSPYSPQTQETDSAVQK.E	3	Q6B011_HUMAN
Q6B011	2	3.8546	K.GTSGSLADVFANTR.I	1	Q6B011_HUMAN
Disabled homolog 2 - Homo sapiens (Human)					
P98082	3	3.8781	K.KGPEKTDEYLLAR.F	2	DAB2_HUMAN
P98082	2	5.1914	R.KGEQTSSGTLSAFASYFNSK.V	5	DAB2_HUMAN
P98082	3	5.1022	R.AGPPKDISSDAFTALDPLGDKEIK.D	1	DAB2_HUMAN
P98082	2	3.7663	K.VGIPQENADHDDFDANQLLNK.I	1	DAB2_HUMAN
P98082	2	3.5876	K.TGQQAEPLVVDLK.D	1	DAB2_HUMAN
P98082	2	3.8211	K.STDNAFENPFFK.D	5	DAB2_HUMAN
P98082	2	3.7558	K.LIGIDDVPDARGDK.M	1	DAB2_HUMAN
P98082	2	2.7708	K.INEPPKPAPR.Q	1	DAB2_HUMAN
P98082	3	5.595	K.DSFGSSQASVASSQPVSSEMYR.D	8	DAB2_HUMAN
P98082	2	3.1416	K.DSFGSSQASVASSQPVSSEM#YR.D	1	DAB2_HUMAN
P98082	2	2.868	K.DLFQVIYNVK.K	1	DAB2_HUMAN
P98082	2	4.1169	K.DISSDAFTALDPLGDKEIK.D	2	DAB2_HUMAN
P98082	2	3.846	K.LIGIDDVPDAR.G	3	DAB2_HUMAN
Disabled homolog 2-interacting protein - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5VWQ8	3	4.8943	R.RIDQPPPPPPPPPAPR.G	2	DAB2P_HUMAN
Q5VWQ8	2	3.9067	K.LGSFSTAAEELAR.R	2	DAB2P_HUMAN
Discoidin, CUB and LCCL domain-containing protein 2 precursor - Homo sapiens (Human)					
Q96PD2	3	3.9961	K.AGKPGLPAPDELVYQVPQSTQEVSGAGR.D	1	DCBD2_HUMAN
Disco-interacting protein 2 homolog A - Homo sapiens (Human)					
Q14689	3	4.1938	R.FRSDVHTEAVQAALAK.Y	2	DIP2A_HUMAN
Disks large-associated protein 1 - Homo sapiens (Human)					
O14490	2	2.8528	K.ANNWKQMDPLDKK.E	1	DLGP1_HUMAN
DNA - Homo sapiens (Human)					
P26358	3	3.733	R.SKSDGEAKPEPSPSPR.I	1	DNMT1_HUMAN
P26358	2	2.9587	R.VPTLAVPAISLPDDVR.R	2	DNMT1_HUMAN
DNA damage-binding protein 2 - Homo sapiens (Human)					
Q92466	2	2.8766	K.TSEIVLRPR.N	1	DDB2_HUMAN
DNA fragmentation factor subunit alpha - Homo sapiens (Human)					
O00273	3	3.8201	K.QLLQLYLQALEKEGSLLSKQEESK.A	1	DFFA_HUMAN
O00273	2	3.6423	R.LQHTLQQVLDQR.E	1	DFFA_HUMAN
O00273	2	3.4203	R.LQQTQSLHSLR.S	1	DFFA_HUMAN
O00273	3	3.8848	R.EQHGVAASCLEDLR.S	2	DFFA_HUMAN
O00273	2	3.338	K.FVALASNEK.W	8	DFFA_HUMAN
O00273	2	3.3054	K.EGSLLSKQEESK.A	2	DFFA_HUMAN
O00273	2	4.3172	K.AAFGEEVDAVDTGISR.E	3	DFFA_HUMAN
O00273	2	4.9952	R.ETSSDVALASHILTALR.E	3	DFFA_HUMAN
O00273	2	3.5936	K.KTETVQEACER.E	1	DFFA_HUMAN
DNA polymerase epsilon subunit 3 - Homo sapiens (Human)					
Q9NRF9	2	2.8118	K.TLNASDVLSAMEEMEFQR.F	1	DPOE3_HUMAN
DNA polymerase subunit delta 3 - Homo sapiens (Human)					
Q15054	2	3.4164	K.IVEQPTVSVTEPK.L	2	DPOD3_HUMAN
Q15054	2	3.6104	K.EVTNASAAGNK.A	3	DPOD3_HUMAN
Q15054	2	4.0206	K.EVTNASAAGNKAPGK.G	3	DPOD3_HUMAN
DNA topoisomerase 1 - Homo sapiens (Human)					
P11387	3	4.9542	R.KLEEEEDGKLK.K	1	TOP1_HUMAN
P11387	2	3.7502	R.DEDDADYKPK.K	7	TOP1_HUMAN
P11387	2	2.8088	R.KLEEEEDGK.L	1	TOP1_HUMAN
DNA topoisomerase 2-binding protein 1 - Homo sapiens (Human)					
Q92547	2	2.8178	R.TGKRADESHFLIENSTK.E	1	TOPB1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
DNA-(apurinic or apyrimidinic site) lyase - Homo sapiens (Human)					
P27695	2	2.8506	K.GAVAEDGDELR.T	3	APEX1_HUMAN
P27695	2	5.3772	K.GAVAEDGDELRTEPEAK.K	14	APEX1_HUMAN
P27695	2	3.3779	K.KGAVAEDGDELR.T	1	APEX1_HUMAN
P27695	2	4.3911	K.KGAVAEDGDELRTEPEAK.K	3	APEX1_HUMAN
DNA-binding protein A - Homo sapiens (Human)					
P16989	3	5.1767	R.PAPAVGEAEDKENQQATSGPNQPSVR.R	2	DBPA_HUMAN
P16989	2	2.7049	R.RPYNYR.R	1	DBPA_HUMAN
P16989	4	6.3994	K.SPVGSGAPQAAAPAPAAHVAGNPGGDAAPA	1	DBPA_HUMAN
P16989	3	6.0152	R.IQAGEIGEMKDGVPEGAQLQGPVHR.N	2	DBPA_HUMAN
DNA-dependent protein kinase catalytic subunit - Homo sapiens (Human)					
P78527	3	4.6868	K.RLGLPGDEVDNKVK.G	2	PRKDC_HUMAN
DNA-directed RNA polymerase I subunit RPA34 - Homo sapiens (Human)					
O15446	3	7.3073	K.NQQLKEPEAAGPVGTEPTVETLEPLGVLFPS	2	RPA34_HUMAN
O15446	2	3.5841	K.QEQINTEPLEDTVLSPTK.K	1	RPA34_HUMAN
DNA-directed RNA polymerase II subunit RPB1 - Homo sapiens (Human)					
P24928	2	3.5002	K.YSPTSPTYSPTSPVYTPTSPK.Y	1	RPB1_HUMAN
DNA-directed RNA polymerase II subunit RPB9 - Homo sapiens (Human)					
P36954	2	3.6821	K.EAVFFQSHSAR.A	3	RPB9_HUMAN
DnaJ homolog subfamily A member 5 - Homo sapiens (Human)					
Q5F1R6	2	2.9545	R.DASEEELKK.A	1	DNJA5_HUMAN
DnaJ homolog subfamily B member 1 - Homo sapiens (Human)					
P25685	2	2.9878	R.GASDEEIKR.A	1	DNJB1_HUMAN
P25685	2	2.8998	R.KKQDPPVTHDLR.V	1	DNJB1_HUMAN
P25685	2	3.1324	R.DGSDVIYPAR.I	1	DNJB1_HUMAN
DnaJ homolog subfamily B member 12 - Homo sapiens (Human)					
Q9NXW2	2	3.3387	K.AGGTDAPSANGEAGGESTK.G	2	DJB12_HUMAN
DnaJ homolog subfamily B member 14 - Homo sapiens (Human)					
Q8TBM8	2	2.8246	K.DAGDEDLKK.A	1	DJB14_HUMAN
DnaJ homolog subfamily B member 2 - Homo sapiens (Human)					
P25686	2	3.7764	R.AEAGSGGPGFTFTFR.S	1	DNJB2_HUMAN
P25686	3	5.2272	R.EFFGSGDPFAELFDDLGPFSELQNR.G	2	DNJB2_HUMAN
P25686	2	2.9211	R.EGLTGTGTGPSR.A	2	DNJB2_HUMAN
P25686	2	3.4956	R.REQQPSVTSR.S	1	DNJB2_HUMAN
P25686	2	2.7493	R.SASADDIKK.A	1	DNJB2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
DnaJ homolog subfamily B member 6 - Homo sapiens (Human)					
O75190	2	4.0485	R.APGPWDPLASAAGLK.E	2	DNJB6_HUMAN
DnaJ homolog subfamily C member 1 - Homo sapiens (Human)					
Q96KC8	3	4.2029	R.EDAEGVAAEEEQEGDSGEQETGATDARPR.	2	DNJC1_HUMAN
Q96KC8	2	3.9876	R.LLEATAKPEPEEK.S	4	DNJC1_HUMAN
Q96KC8	2	3.9449	R.SAEEPWTQNQQK.L	4	DNJC1_HUMAN
Q96KC8	2	2.9644	K.STVQNSRPIK.T	1	DNJC1_HUMAN
DnaJ homolog subfamily C member 17 - Homo sapiens (Human)					
Q9NVM6	2	3.4276	R.QAQAQESEEEEESR.S	2	DJC17_HUMAN
Q9NVM6	2	3.373	K.ISWLEGQPQDAVGR.S	1	DJC17_HUMAN
DnaJ homolog subfamily C member 2 - Homo sapiens (Human)					
Q99543	2	3.5207	K.AAQEQVLNASR.A	1	DNJC2_HUMAN
DnaJ homolog subfamily C member 8 - Homo sapiens (Human)					
O75937	1	2.8393	R.NFQANTK.G	3	DNJC8_HUMAN
O75937	3	3.9414	R.QLSILVHPDKNQDDADRAQK.A	1	DNJC8_HUMAN
O75937	3	4.8804	R.LTRPGSSYFNLNPFEVLQIDPEVTDEEIKK.R	1	DNJC8_HUMAN
O75937	1	2.3005	R.DSVLTSK.N	2	DNJC8_HUMAN
O75937	3	5.5869	R.ALDVIQAGKEYVEHTVKER.K	2	DNJC8_HUMAN
O75937	3	4.3187	R.ALDVIQAGKEYVEHTVK.E	4	DNJC8_HUMAN
O75937	1	3.2324	K.LLLDQEQK.K	3	DNJC8_HUMAN
O75937	2	3.6393	K.LFAELEIK.R	3	DNJC8_HUMAN
O75937	3	4.3648	K.KEGKPTIVEEDDPELFK.Q	1	DNJC8_HUMAN
O75937	3	4.6151	K.EGKPTIVEEDDPELFK.Q	2	DNJC8_HUMAN
O75937	1	2.4807	K.AFEAVDK.A	1	DNJC8_HUMAN
O75937	2	3.519	K.AFEAVDKAYK.L	2	DNJC8_HUMAN
O75937	3	4.6497	R.QREEEIEAQEK.A	6	DNJC8_HUMAN
O75937	2	3.7144	R.ALDVIQAGK.E	4	DNJC8_HUMAN
DnaJ homolog subfamily C member 9 - Homo sapiens (Human)					
Q8WXX5	2	4.9082	K.KISLEDIQAFEK.T	1	DNJC9_HUMAN
Q8WXX5	2	4.6397	K.ISLEDIQAFEK.T	3	DNJC9_HUMAN
DNA-repair protein XRCC1 - Homo sapiens (Human)					
P18887	2	3.7415	R.AIGSTSKPQESPK.G	2	XRCC1_HUMAN
P18887	2	2.8757	R.KLDLNQEEK.K	1	XRCC1_HUMAN
Docking protein 1 - Homo sapiens (Human)					
Q99704	2	3.635	K.SHNSALYSQVQK.S	1	DOK1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q99704	3	3.9232	R.STKPLLAPKPQGPAFPEPGTATGSGIK.S	1	DOK1_HUMAN
Docking protein 3 - Homo sapiens (Human)					
Q7L591	3	3.9204	R.LLELDQVEGTGRPDPQAGFK.A	2	DOK3_HUMAN
Dolichol-phosphate mannosyltransferase subunit 3 - Homo sapiens (Human)					
Q9P2X0	2	2.7666	R.ELQSQIQEAR.A	2	DPM3_HUMAN
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 67 kDa subunit precursor - Homo					
P04843	2	3.2928	K.ALTSEIALLQSR.L	1	RIB1_HUMAN
P04843	2	3.8789	K.DTYIENEKLISGK.R	2	RIB1_HUMAN
Dolichyl-P-Man:Man(5)GlcNAc(2)-PP-dolichyl mannosyltransferase - Homo sapiens (Human)					
Q92685	2	3.0273	R.RLLLREPR.Y	3	ALG3_HUMAN
Doublecortin and CaM kinase-like 1 - Homo sapiens (Human)					
Q5VZY9	2	3.1628	R.DLYRPLSSDDLDSVGDSV.-	2	Q5VZY9_HUMA
Doublecortin domain-containing protein 2 - Homo sapiens (Human)					
Q9UHG0	2	4.12	K.STVGSSDNSSPQPLKR.K	3	DCDC2_HUMAN
Q9UHG0	3	3.9673	K.KEDVNSEKLTK.L	1	DCDC2_HUMAN
Q9UHG0	2	3.4436	K.STVGSSDNSSPQPLK.R	2	DCDC2_HUMAN
Double-strand break repair protein MRE11A - Homo sapiens (Human)					
P49959	2	3.2095	K.IM#SQSQVSK.G	2	MRE11_HUMAN
P49959	2	2.8992	K.IMSQSQVSK.G	3	MRE11_HUMAN
P49959	2	2.8738	K.NVQLSLLTER.G	1	MRE11_HUMAN
P49959	2	3.8255	R.ADTGLETSTR.S	4	MRE11_HUMAN
P49959	2	2.7737	R.NM#SIIDAFK.S	1	MRE11_HUMAN
Double-stranded RNA-specific adenosine deaminase - Homo sapiens (Human)					
P55265	2	3.0128	R.NTNSVPETAPAAIPETR.R	2	DSRAD_HUMAN
P55265	2	4.0008	R.DINAVLIDMER.Q	3	DSRAD_HUMAN
Double-stranded RNA-specific editase 1 - Homo sapiens (Human)					
P78563	3	5.3952	K.DGSTPGPGEGSQLSNGGGGGPGR.K	2	RED1_HUMAN
Dpy-30-like protein - Homo sapiens (Human)					
Q9C005	3	4.2066	K.ERPPNPIEFLASYLLK.N	6	DPY30_HUMAN
Q9C005	2	3.015	K.QKVDLQSLPTR.A	1	DPY30_HUMAN
Q9C005	3	6.5069	R.AYLDQTVVPILLQGLAVLAK.E	7	DPY30_HUMAN
Q9C005	2	4.038	R.IVENEKINAEK.S	6	DPY30_HUMAN
Q9C005	2	3.052	K.AQFEDRN.-	3	DPY30_HUMAN
DPYSL3 protein - Homo sapiens (Human)					
Q6DEN2	3	3.9136	R.RGWDSSHEDDLPVYLAR.P	1	Q6DEN2_HUMA

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q6DEN2	2	2.7889	K.RIVAPPGGR.S	1	Q6DEN2_HUMA
Q6DEN2	2	3.3131	R.PGTTDQVPR.Q	2	Q6DEN2_HUMA
Drebrin-like protein - Homo sapiens (Human)					
Q9UJU6	2	4.9137	K.FVLINWTGEGVNDVR.K	5	DBNL_HUMAN
Q9UJU6	2	3.4559	K.ASGANYSFHK.E	3	DBNL_HUMAN
Q9UJU6	2	2.741	K.DPNSGLPK.F	1	DBNL_HUMAN
Q9UJU6	2	4.8099	K.ESGRFQDVGPQAPVGSVYQK.T	4	DBNL_HUMAN
Q9UJU6	2	4.5808	R.FQDVGPQAPVGSVYQK.T	2	DBNL_HUMAN
Q9UJU6	2	4.328	R.AMSTTSISSPQPGK.L	9	DBNL_HUMAN
Q9UJU6	2	4.3288	K.SPTDWALFTYEGNSNDIR.V	8	DBNL_HUMAN
Q9UJU6	2	2.8119	K.TNAVSEIKR.V	1	DBNL_HUMAN
Q9UJU6	2	2.9473	R.AEEDVEPECIMEK.V	2	DBNL_HUMAN
Q9UJU6	2	3.5681	R.AM#STTSISSPQPGK.L	4	DBNL_HUMAN
Q9UJU6	1	2.2738	K.DSFWAK.A	1	DBNL_HUMAN
Q9UJU6	2	3.4382	R.NEQESAVHPR.E	3	DBNL_HUMAN
Q9UJU6	3	6.7982	R.VVTEKSPTDWALFTYEGNSNDIR.V	4	DBNL_HUMAN
Q9UJU6	2	4.0083	R.YQEQGGEASPQR.T	6	DBNL_HUMAN
Q9UJU6	3	4.1155	R.VAGTGEGGLEEMVEELNSGK.V	1	DBNL_HUMAN
Q9UJU6	2	4.6793	R.VAGTGEGGLEEM#VEELNSGK.V	6	DBNL_HUMAN
Q9UJU6	2	4.6521	R.TWEQQQEVVSR.N	7	DBNL_HUMAN
Q9UJU6	3	5.1162	R.NRNEQESAVHPR.E	9	DBNL_HUMAN
Q9UJU6	2	3.8678	R.NGPALQEAYVR.V	5	DBNL_HUMAN
Dynactin 2 variant - Homo sapiens (Human)					
Q53H88	2	4.091	K.LLGPDAAINLTDPDGALAK.R	1	Q53H88_HUMAN
Q53H88	2	4.3261	K.ESATEEKLTPVLLAK.Q	2	Q53H88_HUMAN
Q53H88	2	3.5732	K.ASVEDADTQSK.V	5	Q53H88_HUMAN
Q53H88	2	4.5913	K.LLGPDAAINLTDPDGALAKR.L	1	Q53H88_HUMAN
Q53H88	2	3.3911	K.RLLLQLEATK.N	2	Q53H88_HUMAN
Q53H88	2	4.7111	R.LLHEVQELTTEVEK.I	4	Q53H88_HUMAN
Q53H88	2	4.6162	R.WSPIASTLPELVQR.L	6	Q53H88_HUMAN
Q53H88	3	3.8992	R.TGYESGEYEMLGEGLGVKETPQQK.Y	1	Q53H88_HUMAN
Q53H88	2	3.612	R.TGYESGEYEM#LGEGLGVK.E	1	Q53H88_HUMAN
Q53H88	2	3.5869	R.LTELETAVR.C	2	Q53H88_HUMAN
Q53H88	2	2.9033	K.RLTELETAVR.C	1	Q53H88_HUMAN
Q53H88	2	2.7328	K.TTVKESATEEK.L	1	Q53H88_HUMAN
Q53H88	3	3.8374	R.LLHEVQELTTEVEKIK.T	1	Q53H88_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q53H88	3	3.9123	K.TTVKESATEEKLTPVLLAK.Q	1	Q53H88_HUMAN
Q53H88	2	4.132	K.VSALDLAVLDQVEAR.L	2	Q53H88_HUMAN
Q53H88	2	3.312	K.YADLPGIAR.N	2	Q53H88_HUMAN
Q53H88	2	3.0159	R.ENLATVEGNFASIDER.M	1	Q53H88_HUMAN
Dynamin-binding protein - Homo sapiens (Human)					
Q6XZF7	2	3.5101	R.SGDSADVAR.D	3	DNMBP_HUMAN
Dysbindin - Homo sapiens (Human)					
Q96EV8	3	3.7036	K.HM#QSQQLENYKK.N	1	DTBP1_HUMAN
Dystroglycan precursor - Homo sapiens (Human)					
Q14118	2	2.7901	R.GGEPNQRPELK.N	2	DAG1_HUMAN
Q14118	2	3.5074	K.ATSITVTGSGSCR.H	4	DAG1_HUMAN
Q14118	2	3.0613	K.LREQQLVGEK.S	3	DAG1_HUMAN
Q14118	2	3.7763	R.HLQFIPVVPPR.R	5	DAG1_HUMAN
Q14118	2	3.4972	R.IAEDDGKPRPAFSNALEPDFK.A	2	DAG1_HUMAN
Q14118	3	4.2893	K.IPSDTFYDHEDTTTDKLK.L	2	DAG1_HUMAN
E1A-binding protein p400 - Homo sapiens (Human)					
Q96L91	3	4.9367	R.HQPASASSTAASPAHPAK.L	1	EP400_HUMAN
E3 SUMO-protein ligase CBX4 - Homo sapiens (Human)					
O00257	2	3.6921	K.SGEVAEGEAR.S	2	CBX4_HUMAN
E3 SUMO-protein ligase RanBP2 - Homo sapiens (Human)					
P49792	2	2.7414	K.EIDTDSTSQGESK.I	1	RBP2_HUMAN
P49792	2	2.884	K.FGNTEQGFK.F	1	RBP2_HUMAN
E3 ubiquitin-protein ligase CBL - Homo sapiens (Human)					
P22681	2	3.2943	K.VPVSAPSSSDPWTGR.E	2	CBL_HUMAN
E3 ubiquitin-protein ligase HECW2 - Homo sapiens (Human)					
Q9P2P5	2	3.1689	R.ADDGSLTSQTK.L	2	HECW2_HUMAN
E3 ubiquitin-protein ligase NEDD4 - Homo sapiens (Human)					
P46934	2	2.9297	R.SYYVDHNSR.T	1	NEDD4_HUMAN
E3 ubiquitin-protein ligase NEDD4-like protein - Homo sapiens (Human)					
Q96PU5	2	4.554	R.SLSSPTVTLSAPLEGAK.D	2	NED4L_HUMAN
E3 ubiquitin-protein ligase Praja2 - Homo sapiens (Human)					
O43164	2	4.0031	R.YQESLGNTVFELENR.E	4	PJA2_HUMAN
O43164	2	2.8834	R.SSQDEM#VSTK.Q	2	PJA2_HUMAN
O43164	2	3.2125	R.SSQDEMVSTK.Q	2	PJA2_HUMAN
O43164	2	3.1572	R.ETENNQM#TSESGATAGR.Q	2	PJA2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
E3 ubiquitin-protein ligase RING2 - Homo sapiens (Human)					
Q99496	2	2.8937	R.PHPTLMEKDDSAQTR.Y	1	RING2_HUMAN
Early endosome antigen 1 - Homo sapiens (Human)					
Q15075	3	6.1463	R.ADLQNHLDTAQNALQDKQQELNK.I	3	EEA1_HUMAN
Q15075	2	3.9009	K.QQHQEQQALQQSTTAK.L	3	EEA1_HUMAN
Q15075	2	2.9556	K.SEFEKENQK.G	3	EEA1_HUMAN
Q15075	2	3.0756	K.SLGSADELFK.H	2	EEA1_HUMAN
Q15075	2	2.7096	K.SSVNELTQK.N	1	EEA1_HUMAN
Q15075	3	4.7556	K.TKQQHQEQQALQQSTTAK.L	1	EEA1_HUMAN
Q15075	2	4.862	K.LSLAQEDLISNR.N	4	EEA1_HUMAN
Q15075	3	6.2026	R.DDVTLLRQEVQDLQASLKEEK.W	3	EEA1_HUMAN
Q15075	2	2.972	R.IQTTVTELQK.V	2	EEA1_HUMAN
Q15075	2	2.897	R.NQQILKDQVK.K	1	EEA1_HUMAN
Q15075	3	4.5638	R.VLSLETSVNELNSQLNESK.E	1	EEA1_HUMAN
Q15075	3	6.0653	K.TELENKLQQQLTQAAQELAAEKEK.I	2	EEA1_HUMAN
Q15075	2	4.3323	K.NHTLQEQVTQLTEK.L	3	EEA1_HUMAN
Q15075	2	4.763	K.LREAQNDLEQVLR.Q	4	EEA1_HUMAN
Q15075	2	4.1347	K.LMDKEQQVADLQLK.L	2	EEA1_HUMAN
Q15075	2	3.0323	K.KSEALESIK.Q	3	EEA1_HUMAN
Q15075	2	3.0145	K.KLEADSLEVK.A	2	EEA1_HUMAN
Q15075	2	3.7736	K.IQNLEALLQK.S	4	EEA1_HUMAN
Q15075	2	3.2032	K.IQHEELNNR.I	1	EEA1_HUMAN
Q15075	2	4.8163	K.GPQEVAVYVQELQK.L	1	EEA1_HUMAN
Q15075	2	4.0656	K.ELVQVQTLMDNMTLER.E	2	EEA1_HUMAN
Q15075	2	5.4208	K.ASKEQALQDLQQQR.Q	1	EEA1_HUMAN
Q15075	2	4.8976	K.YLSLEQKTEELEGQIK.K	4	EEA1_HUMAN
Q15075	2	2.8176	K.NIQATLHQK.D	2	EEA1_HUMAN
Q15075	4	5.3051	R.ADLQNHLDTAQNALQDKQQELNKITTQLDQV	2	EEA1_HUMAN
Echinoderm microtubule-associated protein-like 1 - Homo sapiens (Human)					
O00423	2	4.6569	R.LNITEEQQAVLNR.K	2	EMAL1_HUMAN
O00423	2	4.1518	R.VQMQEDDIQLLK.S	4	EMAL1_HUMAN
O00423	2	3.8087	R.VQM#QEDDIQLLK.S	1	EMAL1_HUMAN
O00423	3	5.5025	R.RLNITEEQQAVLNR.K	5	EMAL1_HUMAN
O00423	1	2.466	K.ETAVPATK.S	3	EMAL1_HUMAN
O00423	2	3.0508	K.ARPLMQTLPLR.T	3	EMAL1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O00423	3	4.6489	R.RLNITEEQQAVLNRK.G	2	EMAL1_HUMAN
Echinoderm microtubule-associated protein-like 3 - Homo sapiens (Human)					
Q32P44	2	3.3663	R.GGKDPLSSPGGPGSR.R	1	EMAL3_HUMAN
Echinoderm microtubule-associated protein-like 4 - Homo sapiens (Human)					
Q9HC35	3	4.3502	R.RLAISEDHVASVK.K	2	EMAL4_HUMAN
Q9HC35	2	3.9521	R.KPSHTSAVSIAGK.E	2	EMAL4_HUMAN
Q9HC35	2	3.2887	R.EKKEESHSNDQSPQIR.A	2	EMAL4_HUMAN
Q9HC35	2	4.1436	R.ASPSPQPSSQPLQIHR.Q	5	EMAL4_HUMAN
Q9HC35	2	2.7821	K.SIKRPSPAEK.S	1	EMAL4_HUMAN
Q9HC35	2	4.2879	K.SHNSWENSDDSR.N	4	EMAL4_HUMAN
Q9HC35	3	4.8581	K.KEESHSNDQSPQIR.A	5	EMAL4_HUMAN
Q9HC35	2	3.1488	K.AALADVLR.R	2	EMAL4_HUMAN
EF-hand calcium-binding domain-containing protein 4B - Homo sapiens (Human)					
Q9BSW2	2	3.3456	R.GDEDLGDM#GEDEEAQFR.M	1	EFC4B_HUMAN
EF-hand domain-containing protein 1 - Homo sapiens (Human)					
Q5JVL4	2	3.1536	K.DPGVQELEALIDTIQK.Q	1	EFHC1_HUMAN
EF-hand domain-containing protein 2 - Homo sapiens (Human)					
Q96C19	2	3.4514	K.AAFKELQSTFK.-	2	EFHD2_HUMAN
Egl nine homolog 1 - Homo sapiens (Human)					
Q9GZT9	3	4.3381	R.AAAGGQGSAVAAEAEPGKEEPPAR.S	3	EGLN1_HUMAN
EH domain-binding protein 1 - Homo sapiens (Human)					
Q8NDI1	2	2.8983	K.AVTESSEQDM#K.S	1	EHBP1_HUMAN
EH domain-binding protein 1-like protein 1 - Homo sapiens (Human)					
Q8N3D4	2	3.9954	R.DSGVPGLEADTTGIQVK.E	2	EH1L1_HUMAN
Q8N3D4	3	4.3217	R.HVDTKGPEATGVMPEAR.C	1	EH1L1_HUMAN
Q8N3D4	2	3.5366	R.LGPSIEDKGSGDPFGR.Q	1	EH1L1_HUMAN
Q8N3D4	3	3.9081	R.LKAEEMDTEDRPEASGVDTEPR.S	1	EH1L1_HUMAN
EH domain-containing protein 1 - Homo sapiens (Human)					
Q9H4M9	3	3.7195	K.VKLEGHELPADLPPHLVPPSKR.R	1	EHD1_HUMAN
Q9H4M9	3	4.1538	K.DKPTYDEIFYTLSPVNGK.I	1	EHD1_HUMAN
Q9H4M9	3	6.7771	K.LADVDKDGLLDDEEFALANHLIK.V	2	EHD1_HUMAN
EH domain-containing protein 2 - Homo sapiens (Human)					
Q9NZN4	2	3.9811	K.LEGHGLPANLPR.R	4	EHD2_HUMAN
Q9NZN4	1	2.3995	K.TWMVGTK.L	1	EHD2_HUMAN
Q9NZN4	3	3.8406	K.TWM#VGTKLPNSVLGR.I	1	EHD2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9NZN4	2	5.417	K.SKYDEIFYNLAPADGK.L	5	EHD2_HUMAN
Q9NZN4	3	4.0314	K.DKSKYDEIFYNLAPADGK.L	2	EHD2_HUMAN
Q9NZN4	3	4.0166	K.LSDVDRDGMLDDEEFALASHLIEAK.L	2	EHD2_HUMAN
EH domain-containing protein 3 - Homo sapiens (Human)					
Q9NZN3	3	3.9274	R.DKPMYDEIFYTLSPVDGK.I	1	EHD3_HUMAN
EH domain-containing protein 4 - Homo sapiens (Human)					
Q9H223	3	4.0219	K.DKPVYDELFYTLSPINGK.I	1	EHD4_HUMAN
EHMT1 protein - Homo sapiens (Human)					
Q86X08	2	3.4825	K.TELLGEETPMAADEGSAEK.Q	2	Q86X08_HUMAN
EIF4G1 variant protein - Homo sapiens (Human)					
Q4LE58	2	2.7887	K.SDQWKPLNLEEK.K	1	Q4LE58_HUMAN
Q4LE58	1	2.9128	R.GPAGLGPR.R	4	Q4LE58_HUMAN
ELAV-like protein 1 - Homo sapiens (Human)					
Q15717	2	3.4751	R.VLVDQTTGLSR.G	2	ELAV1_HUMAN
Electron transfer flavoprotein subunit beta - Homo sapiens (Human)					
P38117	2	3.6229	K.IEVIKPGDLGVDLTSK.L	1	ETFB_HUMAN
P38117	2	3.1244	K.LSVISVEDPPQR.T	1	ETFB_HUMAN
ELKS/RAB6-interacting/CAST family member 1 - Homo sapiens (Human)					
Q8IUD2	2	4.5141	R.DNTIMDLQTQLK.E	7	RB6I2_HUMAN
Q8IUD2	3	4.0268	K.TLSMENIQSLNAAYATSGPM#YLSDHENVGS	2	RB6I2_HUMAN
Q8IUD2	2	3.5882	R.SVGKVEPSSQSPGR.S	2	RB6I2_HUMAN
Q8IUD2	3	3.9326	K.LSSTQQSLAEKETHLTNLR.A	1	RB6I2_HUMAN
Q8IUD2	1	2.5024	K.LSSSMNSIK.T	2	RB6I2_HUMAN
Q8IUD2	2	3.3866	K.LMADNYEDDHFK.S	2	RB6I2_HUMAN
Q8IUD2	3	3.9834	K.KTQEEVAALKR.E	1	RB6I2_HUMAN
Q8IUD2	2	2.7578	K.KAHEAALEAR.A	1	RB6I2_HUMAN
Q8IUD2	2	3.195	K.ALQTVIEMKDSK.I	1	RB6I2_HUMAN
Q8IUD2	2	4.4655	K.AAGLQAEIGQVK.Q	3	RB6I2_HUMAN
Q8IUD2	2	3.6066	R.GLTPPASYNLDDDQAAWENELQK.M	3	RB6I2_HUMAN
Q8IUD2	2	4.3215	R.DLNQLFQQDSSSR.T	2	RB6I2_HUMAN
Q8IUD2	2	2.8244	R.EMVLAQEESAR.T	1	RB6I2_HUMAN
Q8IUD2	2	3.0534	R.DNTIM#DLQTQLK.E	1	RB6I2_HUMAN
Q8IUD2	3	4.8411	R.TNAEKQVEELLMAMEK.V	2	RB6I2_HUMAN
Q8IUD2	2	3.2928	R.GLRDLEEEIQM#LK.S	1	RB6I2_HUMAN
Q8IUD2	2	3.2751	K.SAQMLEEAR.R	1	RB6I2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8IUD2	3	5.0717	K.SNGALSTEEREEEMKQMEVYR.S	2	RB6I2_HUMAN
Q8IUD2	3	5.1703	K.TLSM#ENIQSLNAAYATSGPM#YLSDHENVG	1	RB6I2_HUMAN
Q8IUD2	3	5.4708	K.TLSM#ENIQSLNAAYATSGPMYLSDHENVGS	2	RB6I2_HUMAN
Q8IUD2	2	4.8897	K.NKVEQLKEELSSK.E	1	RB6I2_HUMAN
Q8IUD2	4	4.749	K.VSLLQGDLSEKEASLLDLKEHASSLASSGLK.	1	RB6I2_HUMAN
Q8IUD2	2	5.0188	R.TNSTGGSSGSSVGGGSGK.T	13	RB6I2_HUMAN
Q8IUD2	3	3.9001	R.KDTELLALQTKLETLTNQFSDSK.Q	2	RB6I2_HUMAN
Q8IUD2	2	4.9889	R.GLRDLEEEIQMLK.S	3	RB6I2_HUMAN
Elongation factor 1-beta - Homo sapiens (Human)					
P24534	1	2.5172	R.LAQYESK.K	3	EF1B_HUMAN
P24534	2	3.0892	R.LAQYESKK.A	7	EF1B_HUMAN
Elongation factor 2 - Homo sapiens (Human)					
P13639	2	3.6136	K.EGIPALDNFLDKL.-	6	EF2_HUMAN
EMILIN-1 precursor - Homo sapiens (Human)					
Q9Y6C2	3	4.3337	R.LGQLEGLLQAHGDEGCGACGGVQEELGR.L	1	EMIL1_HUMAN
Q9Y6C2	2	3.5396	R.LDTVAGGLQGLR.E	2	EMIL1_HUMAN
Q9Y6C2	2	3.096	R.QATLEGLQEVVGR.L	1	EMIL1_HUMAN
Q9Y6C2	2	3.5431	R.VLLNDGGYYDPETGVFTAPLAGR.Y	3	EMIL1_HUMAN
Q9Y6C2	2	3.5652	R.APAPASAPPGPSEELLR.Q	2	EMIL1_HUMAN
EMILIN-2 precursor - Homo sapiens (Human)					
Q9BXX0	2	4.1905	R.KFQETEQTIQK.L	1	EMIL2_HUMAN
Q9BXX0	2	5.6764	R.TVLDLQSSLAGVSENLK.H	3	EMIL2_HUMAN
Ena/VASP-like protein - Homo sapiens (Human)					
Q9UI08	3	4.0261	R.RVQRPEDASGGSSPSGTSK.S	1	EVL_HUMAN
Q9UI08	2	3.6247	K.SPLQSQPHSR.M	2	EVL_HUMAN
Q9UI08	2	3.5258	K.EDESQM#EDPSTSPSPGTR.A	1	EVL_HUMAN
Q9UI08	2	5.6504	K.KEDESQMEDPSTSPSPGTR.A	2	EVL_HUMAN
Q9UI08	2	4.0587	R.MKQEILEEVVR.E	2	EVL_HUMAN
Q9UI08	2	4.4077	R.VQRPEDASGGSSPSGTSK.S	3	EVL_HUMAN
Q9UI08	2	3.9934	R.ASSGGGGGGLMEEMNK.L	2	EVL_HUMAN
Endoglin precursor - Homo sapiens (Human)					
P17813	2	3.2629	R.GEVTYTTSQVSK.G	2	EGLN_HUMAN
P17813	2	3.0332	K.TQILEWAAER.G	2	EGLN_HUMAN
Endomucin precursor - Homo sapiens (Human)					
Q9ULC0	2	3.7493	K.TISHESGEHSAQGK.T	2	MUCEN_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Endoplasmic reticulum protein ERp29 precursor - Homo sapiens (Human)					
P30040	2	3.7959	K.GALPLDTVTFYK.V	4	ERP29_HUMAN
P30040	2	3.1453	K.FDTQYPYGEK.Q	2	ERP29_HUMAN
P30040	2	4.2809	K.SLNILTAFQK.K	6	ERP29_HUMAN
P30040	3	4.5062	K.YKLDKESYPVFYLFR.D	8	ERP29_HUMAN
P30040	2	4.8139	K.ILDQGEDFPASEMTR.I	7	ERP29_HUMAN
P30040	2	3.007	K.KGAEKEEL.-	1	ERP29_HUMAN
P30040	2	3.3386	K.MSDGKKEELQK.S	3	ERP29_HUMAN
P30040	2	3.4889	K.SLNILTAFQKK.G	2	ERP29_HUMAN
Endoplasmin precursor - Homo sapiens (Human)					
P14625	3	6.2868	R.FQSSHHPTDITSLDQYVER.M	2	ENPL_HUMAN
P14625	3	5.0935	R.EAVEKEFEPLLNWM#KDK.A	1	ENPL_HUMAN
P14625	2	3.4139	R.EAVEKEFEPLLNWMK.D	1	ENPL_HUMAN
P14625	3	4.5919	R.EAVEKEFEPLLNWMKDK.A	4	ENPL_HUMAN
P14625	2	3.4901	R.KEAESSPFVER.L	1	ENPL_HUMAN
P14625	2	3.8154	R.LISLTDENALSGNEELTVK.I	1	ENPL_HUMAN
P14625	3	3.9515	R.RVFITDDFHDM#MPK.Y	1	ENPL_HUMAN
P14625	2	3.4832	K.SILFVPTSAPR.G	5	ENPL_HUMAN
P14625	2	3.6062	R.SGYLLPDTK.A	8	ENPL_HUMAN
P14625	2	2.7367	K.FDESEKTK.E	1	ENPL_HUMAN
P14625	3	4.0159	R.RVFITDDFHDMMPK.Y	1	ENPL_HUMAN
P14625	2	3.1699	K.EAESSPFVER.L	2	ENPL_HUMAN
P14625	3	4.1275	K.KYSQFINFPIYVWSSK.T	2	ENPL_HUMAN
P14625	2	3.8919	K.DISTNYYASQKK.T	4	ENPL_HUMAN
P14625	2	2.9104	K.SGTSEFLNK.M	1	ENPL_HUMAN
P14625	2	2.9219	K.EGVKFDESEKTK.E	1	ENPL_HUMAN
P14625	2	3.7018	K.EVEEDEYKAFYK.S	2	ENPL_HUMAN
P14625	2	3.1116	K.FAFQAEVNR.M	2	ENPL_HUMAN
P14625	3	5.0253	K.IADDKYNDTFWKEFGTNIK.L	2	ENPL_HUMAN
P14625	2	3.4606	K.LIINSLYK.N	2	ENPL_HUMAN
P14625	2	4.2357	K.NLLHVTDTGVGMTR.E	1	ENPL_HUMAN
P14625	2	3.8437	K.DISTNYYASQK.K	3	ENPL_HUMAN
Endothelial cell-selective adhesion molecule precursor - Homo sapiens (Human)					
Q96AP7	3	3.9931	R.ALRPPHGPPRPGALTPTPSLSSQALPSPR.L	1	ESAM_HUMAN
Q96AP7	2	6.1994	R.LPTTDGAHPQPISPIPGGVSSSGLSR.M	8	ESAM_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96AP7	2	3.5563	R.QLPSFQTFFAPALDVIR.G	1	ESAM_HUMAN
Endothelial cell-specific molecule 1 precursor - Homo sapiens (Human)					
Q9NQ30	2	2.86	K.FPFFQYSVTK.S	1	ESM1_HUMAN
Endothelial differentiation-related factor 1 - Homo sapiens (Human)					
O60869	3	5.9267	K.INEKPQVIADYESGR.A	13	EDF1_HUMAN
O60869	2	3.3416	R.ETEELHHDRVTLEVGK.V	2	EDF1_HUMAN
O60869	2	3.1471	R.ETEELHHDR.V	1	EDF1_HUMAN
Enhancer of mRNA-decapping protein 3 - Homo sapiens (Human)					
Q96F86	2	3.9524	K.KPASSSSAPQNIPK.R	1	EDC3_HUMAN
Enhancer of rudimentary homolog - Homo sapiens (Human)					
P84090	2	2.9632	R.ADTQTYQPYNKDWIK.E	2	ERH_HUMAN
Enigma homolog - Homo sapiens (Human)					
Q59FC9	3	5.7183	R.ISNSATYSGSVAPANSALGQTQPSDQDTLVQ	1	Q59FC9_HUMA
Ensconsin - Homo sapiens (Human)					
Q14244	2	3.9444	R.LSSSSATLLNSPDR.A	2	MAP7_HUMAN
Q14244	2	2.7216	R.VREEAERVR.Q	1	MAP7_HUMAN
Q14244	2	3.2344	R.SETAPDSYKVQDK.K	2	MAP7_HUMAN
Q14244	2	3.6313	R.RLEAEQAR.E	8	MAP7_HUMAN
Q14244	2	3.0399	R.RAVSPSNPK.A	1	MAP7_HUMAN
Q14244	2	2.8097	R.EIVWLEREER.A	2	MAP7_HUMAN
Q14244	3	5.1962	R.APLVKVEEATVEER.T	4	MAP7_HUMAN
Q14244	2	2.7463	R.ALREREEAER.A	2	MAP7_HUMAN
Q14244	2	3.6432	K.VEEATVEER.T	4	MAP7_HUMAN
Q14244	2	3.8386	K.TSAGTTDPEEATR.L	6	MAP7_HUMAN
Q14244	2	3.6844	K.STAALSGEAASCSPIIMPYK.A	1	MAP7_HUMAN
Q14244	3	4.4875	K.GRAPLVKVEEATVEER.T	2	MAP7_HUMAN
Q14244	2	3.0562	R.REQEELER.Q	3	MAP7_HUMAN
Envoplakin - Homo sapiens (Human)					
Q92817	2	3.1377	R.ILQQDWSDLM#ADPAGVRR.E	1	EVPL_HUMAN
Eosinophil cationic protein precursor - Homo sapiens (Human)					
P12724	2	4.3282	R.YPVVPVHLDTTI.-	3	ECP_HUMAN
Ephrin-B2 precursor - Homo sapiens (Human)					
P52799	2	4.3989	K.FQEFSPNLWGLEFQK.N	3	EFNB2_HUMAN
P52799	2	2.7946	K.TVGQYEYYK.V	1	EFNB2_HUMAN
P52799	2	4.6252	K.VGQDASSAGSTR.N	5	EFNB2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Epidermal growth factor receptor kinase substrate 8 - Homo sapiens (Human)					
Q12929	3	5.2236	R.KSQMEEVQDELIHR.L	2	EPS8_HUMAN
Q12929	3	4.2974	R.GSHLDQGEAAVAFKPTSNR.H	2	EPS8_HUMAN
Q12929	3	5.6053	R.MEYGPRPADTPPAPSPPPTPAPVPVPLPPST	1	EPS8_HUMAN
Q12929	2	5.3844	K.ISAAASDSGVESFDEGSSH.-	11	EPS8_HUMAN
Q12929	3	4.9888	K.AALEDSSGSSELQEIM#R.R	1	EPS8_HUMAN
Q12929	2	3.577	R.QNSSSSDSGGSIVR.D	3	EPS8_HUMAN
Q12929	2	4.5885	K.SQMEEVQDELIHR.L	2	EPS8_HUMAN
Epidermal growth factor receptor kinase substrate 8-like protein 2 - Homo sapiens (Human)					
Q9H6S3	2	2.9855	K.QQSGSELEELMNK.F	1	ES8L2_HUMAN
Q9H6S3	2	5.1303	R.SQPVSQPLTYESGPDEVR.A	4	ES8L2_HUMAN
Q9H6S3	2	3.1639	R.VYSQLTM#QK.A	2	ES8L2_HUMAN
Q9H6S3	2	3.3187	R.VYSQLTMQK.A	4	ES8L2_HUMAN
Epidermal growth factor receptor precursor - Homo sapiens (Human)					
P00533	3	4.7781	K.GSHQISLDNPDYQQDFFPK.E	5	EGFR_HUMAN
P00533	2	3.5694	K.GSTAENAEYLR.V	3	EGFR_HUMAN
Epithelial protein lost in neoplasm beta variant - Homo sapiens (Human)					
Q59FE8	2	3.4901	R.STPAEDDSPGDSQVK.S	5	Q59FE8_HUMAN
EPRS protein - Homo sapiens (Human)					
A0AVA9	2	2.7716	K.SLYDEVAAQGEVVR.K	1	A0AVA9_HUMA
A0AVA9	3	4.9566	K.TGQEYKPGNPPAEIGQNISSNSSASILESK.S	1	A0AVA9_HUMA
A0AVA9	2	3.3838	K.INEAVECLLSLK.A	2	A0AVA9_HUMA
A0AVA9	3	4.5347	K.APKDQVDIAVQELLQLK.A	2	A0AVA9_HUMA
A0AVA9	2	3.1727	R.DQDLEPGAPSM#GAK.S	2	A0AVA9_HUMA
A0AVA9	2	2.9538	K.NQGGGLSSSGAGEGQGPKK.Q	1	A0AVA9_HUMA
Epsin-1 - Homo sapiens (Human)					
Q9Y6I3	3	5.1256	K.TPESFLGPNAALVDLDSLVSRPGPTPPGAK.A	5	EPN1_HUMAN
Q9Y6I3	3	4.5772	R.KTPESFLGPNAALVDLDSLVSRPGPTPPGAK.	2	EPN1_HUMAN
Q9Y6I3	2	4.0805	R.TALPTSGSSAGELELLAGEVPAR.S	1	EPN1_HUMAN
ERO1-like protein alpha precursor - Homo sapiens (Human)					
Q96HE7	2	4.6539	R.LGAVDESLSEETQK.A	2	ERO1A_HUMAN
Erythrocyte membrane protein band 4.1 like 5 - Homo sapiens (Human)					
Q7Z5S1	3	4.1449	R.LPGLGEPEVEYETLKDTSEK.L	2	Q7Z5S1_HUMAN
ESF1 homolog - Homo sapiens (Human)					
Q9H501	2	3.5838	K.DGTSPEEEIEIER.Q	3	ESF1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Espin. - Homo sapiens (Human)					
Q5T8W7	2	3.2259	K.SFNMMSPTGDNSELLAEIK.A	1	Q5T8W7_HUMA
Q5T8W7	2	3.4236	K.SFNM#MSPTGDNSELLAEIK.A	1	Q5T8W7_HUMA
Ester hydrolase C11orf54 - Homo sapiens (Human)					
Q9H0W9	2	4.9755	R.IAEVGGVPYLLPLVNQK.K	7	CK054_HUMAN
Eukaryotic translation initiation factor 2 subunit 1 - Homo sapiens (Human)					
P05198	2	3.3056	K.VVTDTDETELAR.Q	3	IF2A_HUMAN
P05198	2	3.0302	R.GVFNVQMEPK.V	1	IF2A_HUMAN
P05198	2	4.9516	R.TEGLSVLSQAMAVIK.E	3	IF2A_HUMAN
Eukaryotic translation initiation factor 2 subunit 2 - Homo sapiens (Human)					
P20042	2	2.8306	K.IESDVQEPTEPEDDLDIMLGNK.K	1	IF2B_HUMAN
P20042	2	3.9972	K.DASDDLDDLNFFNQKK.K	1	IF2B_HUMAN
P20042	2	3.8543	K.IFDIDEAEEGVKDLK.I	1	IF2B_HUMAN
P20042	2	4.2417	K.KIFDIDEAEEGVK.D	2	IF2B_HUMAN
P20042	3	3.7039	K.KIFDIDEAEEGVKDLK.I	1	IF2B_HUMAN
P20042	3	4.7028	R.KKDASDDLDDLNFFNQK.K	3	IF2B_HUMAN
P20042	2	4.3076	K.IFDIDEAEEGVK.D	2	IF2B_HUMAN
Eukaryotic translation initiation factor 2A - Homo sapiens (Human)					
Q9BY44	2	4.8835	R.NTVSQSISGDPEIDKK.I	3	EIF2A_HUMAN
Q9BY44	2	2.8113	R.NVNNEVHFFENNNFNTIANK.L	1	EIF2A_HUMAN
Q9BY44	2	3.599	K.SPDLAPTPAPQSTPR.N	1	EIF2A_HUMAN
Q9BY44	2	4.4368	K.LKAIEQLKEQAATGK.Q	1	EIF2A_HUMAN
Q9BY44	2	4.2852	K.AIEQLKEQAATGK.Q	4	EIF2A_HUMAN
Q9BY44	2	4.7659	R.SDKSPDLAPTPAPQSTPR.N	2	EIF2A_HUMAN
Eukaryotic translation initiation factor 3 subunit 1 - Homo sapiens (Human)					
O75822	2	2.9901	K.KGVVPGGGLK.A	1	IF31_HUMAN
Eukaryotic translation initiation factor 3 subunit 10 - Homo sapiens (Human)					
Q14152	2	3.3197	K.AREESWGPPR.E	2	IF3A_HUMAN
Q14152	2	3.6077	R.RGPAEESSSWR.D	4	IF3A_HUMAN
Q14152	2	2.8902	R.RVPPPALSR.D	6	IF3A_HUMAN
Q14152	2	3.3002	R.SEREEVSSWR.R	2	IF3A_HUMAN
Eukaryotic translation initiation factor 3 subunit 4 - Homo sapiens (Human)					
O75821	2	3.1816	R.RADDNATIR.V	1	IF34_HUMAN
O75821	2	2.992	R.RGESMQPNR.R	2	IF34_HUMAN
O75821	2	4.2398	K.TVTEYKIDEDGKK.F	2	IF34_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Eukaryotic translation initiation factor 3 subunit 7 - Homo sapiens (Human)					
O15371	1	2.2734	R.DSSVEVR.S	1	IF37_HUMAN
O15371	2	3.3117	K.VADWTGATYQDK.R	2	IF37_HUMAN
Eukaryotic translation initiation factor 3 subunit 9 - Homo sapiens (Human)					
P55884	2	4.7845	R.AQAVSEDAGGNEGR.A	2	IF39_HUMAN
Eukaryotic translation initiation factor 4 gamma 1 - Homo sapiens (Human)					
Q04637	2	2.7627	R.GPQAGLGPR.R	1	IF4G1_HUMAN
Eukaryotic translation initiation factor 4 gamma 2 - Homo sapiens (Human)					
P78344	3	5.1464	K.KGQLNADEISLRPAQSFLMNK.N	2	IF4G2_HUMAN
P78344	3	4.0694	K.SQGLSQLYHNQSQGLLSQLQGQSK.D	1	IF4G2_HUMAN
Eukaryotic translation initiation factor 4B - Homo sapiens (Human)					
Q4G0E3	2	3.7874	R.STPKEDDSSASTSQSTR.A	7	Q4G0E3_HUMA
Q4G0E3	2	3.8334	K.EDDSSASTSQSTR.A	3	Q4G0E3_HUMA
Eukaryotic translation initiation factor 4E-binding protein 1 - Homo sapiens (Human)					
Q13541	3	5.7719	R.RVVLGDGVQLPPGDYSTTPGGTLFSTTPGGT	1	4EBP1_HUMAN
Q13541	3	6.9691	K.TPPRDLPTIPGVTSPSSDEPPMEASQSHLR.N	2	4EBP1_HUMAN
Q13541	2	3.5891	K.RAGGEESQFEM#DI.-	2	4EBP1_HUMAN
Q13541	2	3.2121	K.RAGGEESQFEMDI.-	1	4EBP1_HUMAN
Eukaryotic translation initiation factor 4E-binding protein 2 - Homo sapiens (Human)					
Q13542	2	2.8527	K.HAVGDDAQFEMDI.-	1	4EBP2_HUMAN
Q13542	2	3.2079	R.KHAVGDDAQFEMDI.-	1	4EBP2_HUMAN
Eukaryotic translation initiation factor 4H - Homo sapiens (Human)					
Q15056	2	3.84	R.PREEVVQKEQE.-	6	IF4H_HUMAN
Q15056	1	2.182	R.AYSSFGGGR.G	1	IF4H_HUMAN
Q15056	2	3.2475	R.EEVVQKEQE.-	3	IF4H_HUMAN
Q15056	2	3.2417	R.GSNMDFREPTEEER.A	3	IF4H_HUMAN
Eukaryotic translation initiation factor 5A-1 - Homo sapiens (Human)					
P63241	2	4.5806	R.NDFQLIGIQDGYLSLLQDSGEVR.E	2	IF5A1_HUMAN
P63241	3	5.5021	K.RNDFQLIGIQDGYLSLLQDSGEVR.E	1	IF5A1_HUMAN
Eukaryotic translation initiation factor 5B - Homo sapiens (Human)					
O60841	3	5.0405	K.KQDFDEDDILKELEELSLEAQGIK.A	1	IF2P_HUMAN
O60841	2	4.2553	K.LLQAQGVEVPSKDSLPK.K	3	IF2P_HUMAN
O60841	2	3.8603	K.WDGSEEDEDNSK.K	2	IF2P_HUMAN
O60841	2	3.1741	R.ETVAVKPTENNEEEFTSK.D	2	IF2P_HUMAN
O60841	2	2.8871	R.KFEEETVK.S	4	IF2P_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
EVH1 domain binding protein - Homo sapiens (Human)					
Q9NZI9	2	2.8185	K.SNTWSWGILK.M	1	Q9NZI9_HUMAN
Q9NZI9	2	3.7929	K.VSDSDNNEGSSFPAPPK.Q	4	Q9NZI9_HUMAN
Extracellular superoxide dismutase [Cu-Zn] precursor - Homo sapiens (Human)					
P08294	2	3.9631	R.GGNQASVENGNAGR.R	2	SODE_HUMAN
Ezrin-radixin-moesin-binding phosphoprotein 50 - Homo sapiens (Human)					
O14745	2	3.3518	R.IVEVNGVCM#EGK.Q	2	NHERF_HUMAN
O14745	2	4.2621	R.IVEVNGVCMEGK.Q	7	NHERF_HUMAN
O14745	3	4.0588	R.LCCLEKGPNGYGFHLHGEK.G	1	NHERF_HUMAN
O14745	2	3.7728	R.LLVVDPETDEQLQK.L	8	NHERF_HUMAN
O14745	1	2.7944	R.LVEPGSPAEK.A	2	NHERF_HUMAN
O14745	2	2.8051	R.LVEVNGENVEK.E	2	NHERF_HUMAN
O14745	3	5.3847	R.EELLRAQEAPGQAEPPAAAEVQGAGNENEP	2	NHERF_HUMAN
O14745	2	4.8236	R.SASSDTSEELNSQDSPPK.Q	3	NHERF_HUMAN
O14745	2	3.9834	K.LLVVDRETDEFFKK.C	5	NHERF_HUMAN
O14745	2	4.0251	R.SVDPDSPAEASGLR.A	4	NHERF_HUMAN
O14745	2	5.1244	R.LVEVNGENVEKETHQQVVSR.I	1	NHERF_HUMAN
O14745	3	5.5623	K.AGLLAGDRLVEVNGENVEK.E	5	NHERF_HUMAN
O14745	3	6.4321	R.AQEAPGQAEPPAAAEVQGAGNENEPR.E	3	NHERF_HUMAN
O14745	2	3.2102	K.AGLLAGDR.L	2	NHERF_HUMAN
O14745	2	3.4051	R.EALAEAALESPRPALVR.S	5	NHERF_HUMAN
O14745	3	8.3202	K.AGLLAGDRLVEVNGENVEKETHQQVVSR.I	10	NHERF_HUMAN
O14745	2	4.2386	K.GPSGYGFNLHSDK.S	2	NHERF_HUMAN
O14745	2	5.3923	K.KGPSGYGFNLHSDK.S	13	NHERF_HUMAN
O14745	2	2.7358	K.LGVQVREELLR.A	2	NHERF_HUMAN
O14745	2	3.0941	K.LLVVDRETDEFFK.K	2	NHERF_HUMAN
O14745	2	2.8123	R.APQMDWSK.K	2	NHERF_HUMAN
FABP1 protein - Homo sapiens (Human)					
Q05CP7	2	2.9352	K.YQLQSQENFEAFM#K.A	1	Q05CP7_HUMA
Q05CP7	2	3.0729	K.YQLQSQENFEAFMK.A	2	Q05CP7_HUMA
FACT complex subunit SSRP1 - Homo sapiens (Human)					
Q08945	3	3.93	K.IKSDHPGISITDLSK.K	1	SSRP1_HUMAN
Q08945	2	2.7515	K.NM#SGSLYEM#VSRVMK.A	1	SSRP1_HUMAN
Q08945	2	2.8886	K.SDHPGISITDLSK.K	1	SSRP1_HUMAN
Q08945	1	2.2619	R.QLSESFK.S	1	SSRP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
F-actin-capping protein subunit alpha-1 - Homo sapiens (Human)					
P52907	3	5.5663	K.FITHAPPGEFNEVFNDVR.L	4	CAZA1_HUMAN
P52907	2	3.8265	K.EASDPQPEEADGGLK.S	4	CAZA1_HUMAN
F-actin-capping protein subunit alpha-2 - Homo sapiens (Human)					
P47755	3	4.6056	K.FIIHAPPGEFNEVFNDVR.L	1	CAZA2_HUMAN
FAM39B protein - Homo sapiens (Human)					
Q2TAC3	2	4.7906	R.QDDGSSSASPSVQGAPR.E	3	Q2TAC3_HUMA
FAM44A protein - Homo sapiens (Human)					
Q96AL1	2	4.2121	K.VMQTDESNKETANLQER.S	1	Q96AL1_HUMAN
Q96AL1	2	2.7061	R.SISNDDGEEK.I	1	Q96AL1_HUMAN
Q96AL1	2	3.6855	R.SLTVSDDAESSEPER.K	2	Q96AL1_HUMAN
Family with sequence similarity 39, member B - Homo sapiens (Human)					
Q86TK9	2	3.3701	R.ATSQGGDLM#SDLFNK.L	1	Q86TK9_HUMAN
Far upstream element-binding protein 1 - Homo sapiens (Human)					
Q96AE4	3	3.995	K.RPLEDGDQPDAKK.V	1	FUBP1_HUMAN
Q96AE4	3	4.5761	R.SVMTEEYKVPDGM#VGFIIGR.G	2	FUBP1_HUMAN
Q96AE4	3	4.3803	R.SVM#TEEYKVPDGM#VGFIIGR.G	1	FUBP1_HUMAN
Q96AE4	2	3.8397	K.RPLEDGDQPDAK.K	3	FUBP1_HUMAN
Q96AE4	2	2.7016	K.RLLDQIVEK.G	1	FUBP1_HUMAN
Q96AE4	2	4.3606	K.IQIAPDSGGLPER.S	4	FUBP1_HUMAN
Q96AE4	2	4.0232	K.IGGDAGTSLNSNDYGYGGQK.R	4	FUBP1_HUMAN
Q96AE4	3	4.7837	R.SVM#TEEYKVPDGMVGFIIGR.G	1	FUBP1_HUMAN
Far upstream element-binding protein 2 - Homo sapiens (Human)					
Q92945	2	3.0155	K.DAFADAVQR.A	3	FUBP2_HUMAN
Q92945	3	3.7861	R.IIGDPYKVQQACEMVM#DILR.E	1	FUBP2_HUMAN
Q92945	3	5.7806	R.TSMTEEYRVPDGMVGLIIGR.G	3	FUBP2_HUMAN
Q92945	3	4.9774	R.TSMTEEYRVPDGM#VGLIIGR.G	3	FUBP2_HUMAN
Q92945	3	5.6904	R.TSM#TEEYRVPDGMVGLIIGR.G	3	FUBP2_HUMAN
Q92945	3	3.9962	R.TSM#TEEYRVPDGM#VGLIIGR.G	1	FUBP2_HUMAN
Q92945	2	4.0859	R.SVSLTGAPESVQK.A	5	FUBP2_HUMAN
Q92945	2	3.0047	R.QLEDGDQPESK.K	4	FUBP2_HUMAN
Q92945	2	3.9224	R.KDAFADAVQR.A	2	FUBP2_HUMAN
Q92945	2	4.3127	R.IQFKQDDGTGPEK.I	1	FUBP2_HUMAN
Q92945	3	3.9642	R.IIGDPYKVQQACEM#VMDILR.E	1	FUBP2_HUMAN
Q92945	2	4.2446	R.IGGGIDVPVPR.H	6	FUBP2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q92945	2	4.0486	K.VQISPDSGGLPER.S	2	FUBP2_HUMAN
Q92945	2	3.5588	K.MMLDDIVSR.G	3	FUBP2_HUMAN
Q92945	2	4.3587	K.MILIQDGSQNTNVDKPLR.I	2	FUBP2_HUMAN
Q92945	2	3.8647	K.LASQGDSISSQLGPIHPPPR.T	2	FUBP2_HUMAN
Q92945	3	4.6158	K.IGGDAATTVNNSTPDFGFGGQKR.Q	4	FUBP2_HUMAN
Q92945	2	4.054	K.AINQQTGAFVEISR.Q	2	FUBP2_HUMAN
Q92945	3	5.1517	K.KLASQGDSISSQLGPIHPPPR.T	5	FUBP2_HUMAN
Fatty acid-binding protein, epidermal - Homo sapiens (Human)					
Q01469	1	2.1573	K.GFDEYMK.E	1	FABPE_HUMAN
Q01469	3	4.5367	R.LVDSKGFDEYMKELGVGIALR.K	2	FABPE_HUMAN
Q01469	2	2.7627	K.FEETTADGRK.T	1	FABPE_HUMAN
Q01469	3	3.9703	K.GFDEYMKELGVGIALRK.M	1	FABPE_HUMAN
Q01469	2	4.6262	K.GFDEYMKELGVGIALR.K	4	FABPE_HUMAN
Fatty acid-binding protein, heart - Homo sapiens (Human)					
P05413	2	2.8983	K.WDGQETTLVR.E	1	FABPH_HUMAN
P05413	2	3.1167	R.QVASMTKPTTIIEK.N	1	FABPH_HUMAN
FERM, RhoGEF and pleckstrin domain-containing protein 1 - Homo sapiens (Human)					
Q9Y4F1	3	3.8658	K.QADGAASAPTEEEEEVVKDR.T	1	FARP1_HUMAN
Q9Y4F1	2	3.3765	K.VSAGEPGSHPSPAPR.R	1	FARP1_HUMAN
FERM, RhoGEF and pleckstrin domain-containing protein 2 - Homo sapiens (Human)					
O94887	2	3.7626	K.DSSSSLTDPQVSYVK.S	2	FARP2_HUMAN
Ferritin heavy chain - Homo sapiens (Human)					
P02794	3	4.1383	K.AIKELGDHVTNLR.K	3	FRIH_HUMAN
P02794	2	4.2803	K.NVNQSLLELHK.L	4	FRIH_HUMAN
P02794	3	4.0269	K.MGAPESGLAEYLFDKHTLGDSDNES.-	1	FRIH_HUMAN
P02794	2	4.6006	K.AIKELGDHVTNLRK.M	4	FRIH_HUMAN
P02794	2	4.2668	K.M#GAPESGLAEYLFDK.H	3	FRIH_HUMAN
Ferritin light chain - Homo sapiens (Human)					
P02792	2	4.8461	R.LGGPEAGLGEYLFER.L	11	FRIL_HUMAN
FGFR1 oncogene partner - Homo sapiens (Human)					
O95684	3	5.0422	K.IGSLGLGTGEDDDYVDDFNSTSHR.S	6	FR1OP_HUMAN
FGFR1 oncogene partner 2 - Homo sapiens (Human)					
Q9NVK5	2	3.1657	R.TSLEEHQSALELIMSK.Y	2	FGOP2_HUMAN
Q9NVK5	2	3.4462	K.QYQEEIQELNEVAR.H	1	FGOP2_HUMAN
Q9NVK5	2	3.1196	R.STLVMGIQQENR.Q	1	FGOP2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
FH1/FH2 domain-containing protein 1 - Homo sapiens (Human)					
Q9Y613	2	5.1642	K.SGLGDDLVQALGLSK.G	1	FHOD1_HUMAN
Q9Y613	2	3.7661	K.SGLGDDLVQALGLSKGPGLEV.-	2	FHOD1_HUMAN
Fibrillin-1 precursor - Homo sapiens (Human)					
P35555	2	4.0502	R.YEDEECTLPIAGR.H	3	FBN1_HUMAN
P35555	1	3.5431	K.GYILQEDGR.S	6	FBN1_HUMAN
P35555	1	3.3943	R.AGYQSTLTR.T	3	FBN1_HUMAN
P35555	3	4.232	R.IGQGHCVSGM#GM#GR.G	3	FBN1_HUMAN
P35555	3	4.0238	R.IGQGHCVSGM#GMGR.G	1	FBN1_HUMAN
P35555	3	4.0593	R.IGQGHCVSGMGM#GR.G	1	FBN1_HUMAN
P35555	2	2.8336	R.NTPEYEELCPR.G	1	FBN1_HUMAN
Fibrinogen alpha chain precursor [Contains: Fibrinopeptide A] - Homo sapiens (Human)					
P02671	2	2.7296	K.VTSGSTTTTR.R	1	FIBA_HUMAN
P02671	2	2.9056	R.GDSTFESK.S	1	FIBA_HUMAN
P02671	2	3.5577	R.GGSTSYGTGSETESPR.N	4	FIBA_HUMAN
P02671	2	2.9722	R.GSESGIFTNTK.E	3	FIBA_HUMAN
P02671	3	5.3934	R.HRHPDEAAFFDTASTGK.T	6	FIBA_HUMAN
P02671	2	3.2286	R.MELERPGGNEITR.G	2	FIBA_HUMAN
P02671	2	5.6046	R.NPSSAGSWNSGSSGPGSTGNR.N	6	FIBA_HUMAN
P02671	3	7.1025	R.GGSTSYGTGSETESPRNPSSAGSWNSGSS	4	FIBA_HUMAN
P02671	2	3.9943	K.TVIGPDGHKEVTK.E	5	FIBA_HUMAN
P02671	2	4.7947	R.PGSTGTWNPGSSER.G	4	FIBA_HUMAN
P02671	3	4.8114	K.TFPGFFSPM#LGEFVSETESR.G	6	FIBA_HUMAN
P02671	2	4.7247	K.MADEAGSEADHEGTHSTKR.G	4	FIBA_HUMAN
P02671	2	4.1411	K.MADEAGSEADHEGTHSTK.R	5	FIBA_HUMAN
P02671	2	3.8702	K.M#ADEAGSEADHEGTHSTKR.G	3	FIBA_HUMAN
P02671	2	2.7717	K.ALTDM#PQMR.M	2	FIBA_HUMAN
P02671	2	3.574	K.LVTSKGDKELR.T	4	FIBA_HUMAN
P02671	2	4.5934	R.HPDEAAFFDTASTGK.T	5	FIBA_HUMAN
P02671	3	4.4398	K.EVTKEVVTSEDGSDCPEAMDLGTLSGIGTLD	1	FIBA_HUMAN
P02671	3	4.1314	K.ESSSHHPGIAEFPSR.G	10	FIBA_HUMAN
P02671	2	3.404	K.EKVTSGSTTTTR.R	3	FIBA_HUMAN
P02671	1	2.2535	K.ALTDMPQMR.M	2	FIBA_HUMAN
P02671	3	4.7208	K.TFPGFFSPMLGEFVSETESR.G	17	FIBA_HUMAN
Fibrinogen beta chain precursor [Contains: Fibrinopeptide B] - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P02675	3	4.2279	K.GGETSEMYLIQPDSSVKPYR.V	1	FIBB_HUMAN
P02675	3	6.864	R.KGGETSEMYLIQPDSSVKPYR.V	3	FIBB_HUMAN
Fibrinogen gamma chain precursor - Homo sapiens (Human)					
P02679	2	4.6127	K.AIQLTYNPDESSKPNMIDAATLK.S	1	FIBG_HUMAN
P02679	2	3.446	R.FGSYCPTTCGIADFLSTYQTK.V	1	FIBG_HUMAN
P02679	3	3.7637	K.AIQLTYNPDESSKPNM#IDAATLK.S	1	FIBG_HUMAN
Fibroleukin precursor - Homo sapiens (Human)					
Q14314	2	3.1049	K.LQADDNGDPGR.N	2	FGL2_HUMAN
Fibronectin precursor - Homo sapiens (Human)					
P02751	2	4.1452	K.GLAFTDVDVDSIK.I	2	FINC_HUMAN
P02751	2	3.2775	K.IAWESPQGQVSR.Y	2	FINC_HUMAN
P02751	2	6.4671	R.RPGGEPSPEGTTGQSYNQYSQR.Y	4	FINC_HUMAN
Fibronectin type III and SPRY domain containing 2 - Homo sapiens (Human)					
A1L4K1	1	2.2344	R.AGPSPSSER.A	1	A1L4K1_HUMAN
Fibulin-1 precursor - Homo sapiens (Human)					
P23142	2	3.8623	K.TGYYFDGISR.M	5	FBLN1_HUMAN
P23142	2	3.4066	K.DCSLPYATESK.E	3	FBLN1_HUMAN
P23142	2	2.7967	R.SAATLQQEK.T	1	FBLN1_HUMAN
P23142	2	2.9754	R.GYHLNEEGTR.C	1	FBLN1_HUMAN
P23142	2	3.8657	K.SQETGDLDVGGLQETDK.I	1	FBLN1_HUMAN
P23142	2	3.5463	K.IIEVEEEQEDPYLNDR.C	2	FBLN1_HUMAN
P23142	3	6.1287	K.SQETGDLDVGGLQETDKIIEVEEEQEDPYLND	3	FBLN1_HUMAN
Filamin-C - Homo sapiens (Human)					
Q14315	2	3.025	R.AWGPGLETGQVGK.S	2	FLNC_HUMAN
FK506-binding protein 10 precursor - Homo sapiens (Human)					
Q96AY3	2	3.5568	K.GGTYDTYVGSGWLIK.G	1	FKB10_HUMAN
FK506-binding protein 15 - Homo sapiens (Human)					
Q5T1M5	3	3.7687	K.KDDVTSSTGPHK.E	2	Q5T1M5_HUMA
Q5T1M5	2	2.7119	R.EVAPDGPLQESSTR.L	1	Q5T1M5_HUMA
Q5T1M5	3	3.8852	R.LSLTSDPEEGDPLALGPESPGEPQPPQLKK.D	1	Q5T1M5_HUMA
FK506-binding protein 2 precursor - Homo sapiens (Human)					
P26885	2	2.819	K.LVIPSELGYGER.G	2	FKBP2_HUMAN
P26885	2	3.1328	R.KLVIPSELGYGER.G	1	FKBP2_HUMAN
P26885	2	3.1388	K.GWDQGLLGMCEGEK.R	1	FKBP2_HUMAN
FK506-binding protein 3 - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q00688	2	4.6433	K.SEETLDEGPPKYTK.S	3	FKBP3_HUMAN
Q00688	2	4.5247	R.GWDEALLTMSK.G	9	FKBP3_HUMAN
Q00688	2	3.6294	R.GWDEALLTM#SK.G	2	FKBP3_HUMAN
Q00688	2	4.6113	R.AWTVEQLRSEQLPK.K	5	FKBP3_HUMAN
Q00688	2	3.54	K.ETKSEETLDEGPPK.Y	3	FKBP3_HUMAN
Q00688	1	2.282	R.AWTVEQLR.S	1	FKBP3_HUMAN
Q00688	3	5.2783	K.TANKDHLVTAYNHLFETKR.F	4	FKBP3_HUMAN
Q00688	3	4.8959	K.TANKDHLVTAYNHLFETK.R	2	FKBP3_HUMAN
Q00688	2	4.1292	R.GWDEALLTMSKGEK.A	2	FKBP3_HUMAN
Q00688	3	4.2204	K.ARLEIEPEWAYGK.K	5	FKBP3_HUMAN
Q00688	2	2.793	K.LNEDKPKETK.S	2	FKBP3_HUMAN
Q00688	1	2.1575	K.LLGNIK.N	3	FKBP3_HUMAN
Q00688	2	3.2389	K.KGDKTNFPK.K	1	FKBP3_HUMAN
Q00688	3	4.8264	K.FLQEHGSDSFLAEHK.L	5	FKBP3_HUMAN
Q00688	3	5.4461	K.FLQEHGSDSFLAEHKLLGNIK.N	2	FKBP3_HUMAN
Q00688	3	4.7818	K.ARLEIEPEWAYGKK.G	6	FKBP3_HUMAN
Q00688	2	4.2726	K.SEETLDEGPPK.Y	4	FKBP3_HUMAN
FK506-binding protein 4 - Homo sapiens (Human)					
Q02790	2	3.6501	K.AEASSGDHPTDTEMKEEQK.S	2	FKBP4_HUMAN
Q02790	2	3.3948	R.LAEEENKAK.A	7	FKBP4_HUMAN
FK506-binding protein 7 precursor - Homo sapiens (Human)					
Q9Y680	3	4.1235	K.KGDLLNAHYDGYLAK.D	2	FKBP7_HUMAN
Q9Y680	2	3.3165	K.WFVLGVGQVIK.G	2	FKBP7_HUMAN
FKSG42 - Homo sapiens (Human)					
Q9BZ74	2	3.0248	-.MDTTMLNMRNLFEQLVR.G	1	Q9BZ74_HUMAN
FLJ00412 protein - Homo sapiens (Human)					
Q86YV0	2	5.1139	R.TSQTQPTATSPLTSYR.W	5	Q86YV0_HUMA
FLJ10357 protein - Homo sapiens (Human)					
A5PL07	2	3.4177	K.QISLAPETLDSSGDVSPGPR.N	2	A5PL07_HUMAN
FLJ43692 protein - Homo sapiens (Human)					
A5YM69	2	3.1928	K.QKQEQVQDVMLGR.Q	1	A5YM69_HUMA
A5YM69	2	3.8129	K.SEEVTGKQEDHGIKEK.G	2	A5YM69_HUMA
A5YM69	3	4.6528	R.KAQGQGGPEQGEER.K	7	A5YM69_HUMA
A5YM69	2	5.2169	K.GVPVSGQEAKEPESWDGGR.L	3	A5YM69_HUMA
Flotillin-2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q14254	2	3.4736	K.TAEAQLAYELQGAR.E	1	FLOT2_HUMAN
Follistatin-related protein 1 precursor - Homo sapiens (Human)					
Q12841	2	3.7402	K.GAQTQTEEEM#TR.Y	3	FSTL1_HUMAN
Q12841	2	3.2519	K.GAQTQTEEEMTR.Y	1	FSTL1_HUMAN
Q12841	2	3.0302	K.LSFQEFLK.C	2	FSTL1_HUMAN
Q12841	1	2.3213	R.DACLTGSK.I	1	FSTL1_HUMAN
Forkhead box protein K1 - Homo sapiens (Human)					
P85037	2	5.876	R.SRVEEPSGAVTTPAGVIAAAGPQGPGTGE.-	2	FOXK1_HUMAN
Forkhead box protein N3 - Homo sapiens (Human)					
O00409	2	3.1988	K.DSLGDSGYASQHK.K	1	FOXN3_HUMAN
Forkhead box protein O1A - Homo sapiens (Human)					
Q12778	3	4.522	K.ELLTSDSPPHNDIMTPVDPGVAQPNSR.V	2	FOXO1_HUMAN
Forkhead box protein O3A - Homo sapiens (Human)					
O43524	2	3.4103	R.NAWGNLSYADLITR.A	2	FOXO3_HUMAN
Formin-binding protein 1-like - Homo sapiens (Human)					
Q5T0N5	3	3.917	R.MEIHKNEAWLSEVEGK.T	1	FBP1L_HUMAN
Formin-binding protein 4 - Homo sapiens (Human)					
Q8N3X1	2	4.1142	R.NANFEALPEDWR.A	3	FNBP4_HUMAN
Forty-two-three domain-containing protein 1 - Homo sapiens (Human)					
Q96QD9	2	3.1182	K.GVPLQFDINSVGK.Q	1	FYTD1_HUMAN
Q96QD9	2	3.0444	R.LVGATATSSPPPK.A	1	FYTD1_HUMAN
Fos-related antigen 2 - Homo sapiens (Human)					
P15408	3	3.8763	R.GSSGSPAHAESYSSGGGGQQK.F	1	FOSL2_HUMAN
Four and a half LIM domains protein 3 - Homo sapiens (Human)					
Q13643	2	3.34	K.RPIVGLGGGK.Y	2	FHL3_HUMAN
Fragile X mental retardation syndrome-related protein 1 - Homo sapiens (Human)					
P51114	2	3.6509	K.AINGPTSASGDDISK.L	4	FXR1_HUMAN
P51114	2	3.2494	K.DVIEEHGPSEK.A	2	FXR1_HUMAN
Fragile X mental retardation syndrome-related protein 2 - Homo sapiens (Human)					
P51116	3	4.2527	R.TDGSISGDRQPVTVADYISR.A	1	FXR2_HUMAN
Frataxin, mitochondrial precursor - Homo sapiens (Human)					
Q16595	2	4.1487	K.LDLSSLAYSGKDA.-	1	FRDA_HUMAN
Q16595	2	3.5674	K.LGGDLGTYVINK.Q	2	FRDA_HUMAN
Q16595	3	4.5593	K.SGTLGHPGSLDETTYER.L	4	FRDA_HUMAN
Q16595	2	5.0139	K.TKLDLSSLAYSGKDA.-	2	FRDA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Frizzled-6 precursor - Homo sapiens (Human)					
O60353	2	2.7514	K.GQAGSVSESAR.S	1	FZD6_HUMAN
O60353	3	4.4486	R.LSGEQVDGKGQAGSVSESAR.S	2	FZD6_HUMAN
Fructose-1,6-bisphosphatase 1 - Homo sapiens (Human)					
P09467	2	3.8775	K.EAVLDVIPTDIHQR.A	1	F16P1_HUMAN
P09467	2	4.5092	R.APVILGSPDDVLEFLK.V	4	F16P1_HUMAN
Fructose-bisphosphate aldolase A - Homo sapiens (Human)					
P04075	2	4.8207	K.GILAADESTGSIAK.R	2	ALDOA_HUMAN
P04075	2	4.6548	K.GILAADESTGSIAKR.L	3	ALDOA_HUMAN
P04075	3	4.8828	K.GVVPLAGTNGETTTQGLDGLSER.C	2	ALDOA_HUMAN
P04075	3	4.161	K.RLQSIGTENTEENRR.F	4	ALDOA_HUMAN
P04075	2	3.7339	K.YTPSGQAGAAASESLFVSNHAY.-	2	ALDOA_HUMAN
P04075	2	3.911	R.ALANSLACQGK.Y	1	ALDOA_HUMAN
P04075	2	2.8258	R.LQSIGTENTEENRR.F	1	ALDOA_HUMAN
P04075	2	3.4031	K.AAQEEYVKR.A	4	ALDOA_HUMAN
Fructose-bisphosphate aldolase C - Homo sapiens (Human)					
P09972	2	3.7219	K.YEGSGEDGGAAAQSLYIANHAY.-	2	ALDOC_HUMAN
P09972	3	5.893	K.VDKGVVPLAGTDGETTTQGLDGLSER.C	1	ALDOC_HUMAN
P09972	3	6.2886	K.GVVPLAGTDGETTTQGLDGLSER.C	6	ALDOC_HUMAN
P09972	2	3.6531	R.DNAGAATEEFIK.R	1	ALDOC_HUMAN
FYVE and coiled-coil domain-containing protein 1 - Homo sapiens (Human)					
Q9BQS8	2	3.1315	K.M#LADLDDLNRTKK.Y	1	FYCO1_HUMAN
Q9BQS8	2	2.9562	R.EKKQQQEEK.E	1	FYCO1_HUMAN
Q9BQS8	2	3.6792	R.ESAIQGSLASLEAEQASIR.H	1	FYCO1_HUMAN
FYVE, RhoGEF and PH domain-containing protein 4 - Homo sapiens (Human)					
Q96M96	2	2.9929	R.HGLTTTPQQK.L	2	FGD4_HUMAN
FYVE, RhoGEF and PH domain-containing protein 5 - Homo sapiens (Human)					
Q6ZNL6	2	4.4408	R.VPLREDEPKDEGSVGNK.A	4	FGD5_HUMAN
Q6ZNL6	2	2.9686	K.GLESEQAPK.L	2	FGD5_HUMAN
Q6ZNL6	2	4.0978	K.TLLSLEGKPLEASR.A	2	FGD5_HUMAN
Q6ZNL6	3	4.2596	R.SRPPFLPLPLTKPR.S	2	FGD5_HUMAN
G patch domain and KOW motifs-containing protein - Homo sapiens (Human)					
Q92917	2	5.9599	K.EGVLPLTAASTAPISFGFTR.T	2	GPKOW_HUMA
Q92917	2	3.0057	R.ELQSVKPQEAPK.E	1	GPKOW_HUMA
G patch domain-containing protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BRR8	2	3.2006	R.QLAAATAPIPGATLLDDLITPAK.L	1	GPTC1_HUMAN
Q9BRR8	2	3.1569	R.SLAQNAQSSR.A	1	GPTC1_HUMAN
G patch domain-containing protein 8 - Homo sapiens (Human)					
Q9UKJ3	2	3.3272	K.ATGPPSQNSNIGTGR.G	1	GPTC8_HUMAN
G protein-regulated inducer of neurite outgrowth 1 - Homo sapiens (Human)					
Q7Z2K8	2	4.7025	K.ADAGPSGQGDSVSIGK.V	2	GRIN1_HUMAN
G protein-regulated inducer of neurite outgrowth 3 - Homo sapiens (Human)					
Q6ZVF9	2	2.8702	K.KLAGTNSSSLK.A	1	GRIN3_HUMAN
Q6ZVF9	2	2.9279	R.SVSTSPSILTAFLK.E	1	GRIN3_HUMAN
Q6ZVF9	2	3.9137	R.APEHFEQEQLR.V	3	GRIN3_HUMAN
Q6ZVF9	3	4.5642	R.VVSHSSSPVGGPEGER.Q	2	GRIN3_HUMAN
Q6ZVF9	3	6.1984	K.TSLIAASGKEDDLGEPQAASPR.H	3	GRIN3_HUMAN
Q6ZVF9	2	2.762	K.NQESTLEENR.Q	1	GRIN3_HUMAN
Q6ZVF9	2	2.7792	K.FETRPSEFAEK.T	1	GRIN3_HUMAN
Q6ZVF9	2	4.4277	K.ADHSGSLDPTNKGDAR.E	1	GRIN3_HUMAN
Q6ZVF9	2	3.5916	K.SQESGGTESAANPTPSPIR.K	1	GRIN3_HUMAN
Q6ZVF9	2	4.7064	K.SQESGGTESAANPTPSPIRK.N	2	GRIN3_HUMAN
Galectin-1 - Homo sapiens (Human)					
P09382	1	2.7228	K.LPDGYEFK.F	3	LEG1_HUMAN
P09382	2	2.8151	R.VRGEVAPDAK.S	3	LEG1_HUMAN
P09382	2	3.9692	R.LNLEAINYMAADGDFKIK.C	4	LEG1_HUMAN
P09382	2	5.4494	R.LNLEAINYMAADGDFK.I	30	LEG1_HUMAN
P09382	2	5.3484	R.LNLEAINYM#AADGDFK.I	4	LEG1_HUMAN
P09382	1	2.2702	R.GEVAPDAK.S	2	LEG1_HUMAN
P09382	2	5.5314	R.FNAHGDANTIVCNSK.D	8	LEG1_HUMAN
P09382	2	4.11	K.LPDGYEFKFPNR.L	4	LEG1_HUMAN
P09382	4	5.7458	K.FPNRLNLEAINYMAADGDFKIK.C	1	LEG1_HUMAN
P09382	3	4.4802	K.FPNRLNLEAINYMAADGDFK.I	1	LEG1_HUMAN
P09382	2	3.643	K.DGGAWGTEQR.E	8	LEG1_HUMAN
P09382	1	2.7655	K.SFVLNLGK.D	9	LEG1_HUMAN
Galectin-2 - Homo sapiens (Human)					
P05162	2	3.0887	R.GGFNM#SSFK.L	2	LEG2_HUMAN
P05162	2	3.442	R.GGFNMSSFK.L	4	LEG2_HUMAN
P05162	3	3.9553	K.LPDGHELTFPNR.L	2	LEG2_HUMAN
Galectin-3-binding protein precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q08380	2	4.5011	R.ELSEALGQIFDSQR.G	3	LG3BP_HUMAN
Q08380	2	4.3492	R.LADGGATNQGR.V	17	LG3BP_HUMAN
Q08380	2	3.1367	R.SDLAVPSELALLK.A	1	LG3BP_HUMAN
Q08380	1	2.3237	R.STHTLDLSR.E	1	LG3BP_HUMAN
Gamma-adducin - Homo sapiens (Human)					
Q9UEY8	3	4.0426	K.SPEKIEEVLSPEGSPSK.S	3	ADDG_HUMAN
Q9UEY8	3	4.3534	K.VSGGTPIKIEDPNQFVPLNTNPNEVLEKR.N	2	ADDG_HUMAN
Q9UEY8	2	3.152	R.WLNSPNTYMK.V	1	ADDG_HUMAN
Gamma-enolase - Homo sapiens (Human)					
P09104	2	4.107	K.DATNVGDEGGFAPNILENSEALELVK.E	1	ENOG_HUMAN
Gamma-glutamyl hydrolase precursor - Homo sapiens (Human)					
Q92820	2	3.7405	K.KPIIGILM#QK.C	2	GGH_HUMAN
Q92820	3	3.7165	K.KPIIGILMQK.C	2	GGH_HUMAN
Gamma-glutamyltranspeptidase 1 precursor - Homo sapiens (Human)					
P19440	2	3.0639	R.LFQPSIQLAR.Q	1	GGT1_HUMAN
P19440	2	2.8993	R.TVIEQQPVLCEVFCR.D	1	GGT1_HUMAN
P19440	2	2.8009	K.GLAAALENKR.T	1	GGT1_HUMAN
Gamma-interferon-inducible protein Ifi-16 - Homo sapiens (Human)					
Q16666	2	3.055	K.TSLSAPPNSSSTENPK.T	1	IF16_HUMAN
Q16666	2	3.1677	R.TPQMPPTTPSSSFFTK.K	1	IF16_HUMAN
Gamma-synuclein - Homo sapiens (Human)					
O76070	3	4.0181	K.TKEGVMYVGAK.T	1	SYUG_HUMAN
O76070	3	4.3633	K.TKEQANAVSEAVVSSVNTVATK.T	2	SYUG_HUMAN
O76070	2	3.7127	K.TVEEAENIAVTSGVVR.K	2	SYUG_HUMAN
O76070	2	4.734	K.TKENVVQSVTSVAEK.T	3	SYUG_HUMAN
O76070	2	4.4065	K.EKEEVAEEAQSGGD.-	5	SYUG_HUMAN
O76070	2	4.9792	K.TVEEAENIAVTSGVVRK.E	1	SYUG_HUMAN
O76070	1	2.4014	K.EGVVGAVEK.T	2	SYUG_HUMAN
O76070	1	2.7905	K.EGVMYVGAK.T	2	SYUG_HUMAN
O76070	2	3.1389	K.EQANAVSEAVVSSVNTVATK.T	1	SYUG_HUMAN
GAS2-like protein 3 - Homo sapiens (Human)					
Q86XJ1	3	4.2825	K.TIATGLGTQSQPSDGAPQAKPVPAQK.L	4	GA2L3_HUMAN
Q86XJ1	2	2.9962	K.VIPAQNSADLPESTLLPNK.C	1	GA2L3_HUMAN
GatC-like protein - Homo sapiens (Human)					
O43716	2	2.749	R.LALVDFGSR.E	1	GATCL_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43716	2	3.1379	K.LDEQEPFPHS.-	2	GATCL_HUMAN
O43716	2	2.7414	R.AVDTDGVEPMESVLEDR.C	1	GATCL_HUMAN
GC-rich promoter-binding protein 1-like 1 - Homo sapiens (Human)					
Q9H751	2	3.2899	K.LQFEEEDFPSLNPEAGK.Q	1	Q9H751_HUMAN
Q9H751	2	2.9118	R.HDSVDSGVSK.G	3	Q9H751_HUMAN
GC-rich sequence DNA-binding factor - Homo sapiens (Human)					
P16383	2	4.1041	K.IPDAAFIQAAR.R	4	GCF_HUMAN
GDP-L-fucose synthetase - Homo sapiens (Human)					
Q13630	2	3.4301	K.DADLTDTAQTR.A	1	FCL_HUMAN
Gelsolin precursor - Homo sapiens (Human)					
P06396	2	3.121	R.GASQAGAPQGR.V	2	GELS_HUMAN
General transcription factor II-I - Homo sapiens (Human)					
P78347	3	5.1192	K.VPEIEVTVEGPNNNNPQTSAVR.T	3	GTF2I_HUMAN
Gephyrin - Homo sapiens (Human)					
Q9NQX3	2	3.7415	R.DTASLSTTPSESPR.A	2	GEPH_HUMAN
GH3 domain-containing protein precursor - Homo sapiens (Human)					
Q8N2G8	2	2.7724	R.VAWGALVWAATWQRRR.L	1	GHDC_HUMAN
Girdin - Homo sapiens (Human)					
Q3V6T2	3	5.8924	R.SDSSEGFLQLPHQDSQDSSSVGSNSLEDGQ	2	GRDN_HUMAN
Q3V6T2	3	4.9799	R.LPISVDSPPAAADSNTTAASNVDKVQESR.N	1	GRDN_HUMAN
Q3V6T2	3	4.0818	R.KTEDTYFISSAGKPTPGTQGK.I	1	GRDN_HUMAN
Glia maturation factor beta - Homo sapiens (Human)					
P60983	2	3.131	K.NKLVQTAELTK.V	2	GMFB_HUMAN
Glia maturation factor gamma - Homo sapiens (Human)					
O60234	2	3.7758	R.TTDDLTEAWLQEK.L	2	GMFG_HUMAN
Glial fibrillary acidic protein - Homo sapiens (Human)					
P14136	2	3.9177	K.ALAAELNQLR.A	1	GFAP_HUMAN
P14136	2	2.9412	R.DGEVIKESK.Q	1	GFAP_HUMAN
Glucocorticoid receptor - Homo sapiens (Human)					
P04150	3	4.259	K.THSDVSSEQQHLK.G	2	GCR_HUMAN
P04150	3	3.9904	K.VMGNDLGFPQQGQISLSSGETDLKLLEESIAN	1	GCR_HUMAN
P04150	2	3.613	K.SSASTAVSAAPTEKEFPK.T	2	GCR_HUMAN
P04150	2	3.0923	K.SSASTAVSAAPTEK.E	1	GCR_HUMAN
P04150	2	3.7603	K.LLEESIANLNR.S	2	GCR_HUMAN
P04150	2	4.006	K.GSVSNAQQPDLSK.A	2	GCR_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P04150	2	3.962	K.VSASSPSLAVASQSDSK.Q	3	GCR_HUMAN
Glucocorticoid receptor DNA-binding factor 1 - Homo sapiens (Human)					
Q9NRY4	2	3.4605	R.NEEENIYSVPHDSTQGK.I	1	GRLF1_HUMAN
Q9NRY4	2	3.1289	R.TSFSVGSDDELGPIR.K	1	GRLF1_HUMAN
Glucocorticoid-induced transcript 1 protein - Homo sapiens (Human)					
Q86VQ1	2	3.7737	K.LLGPLLPASDLMLK.N	3	GLCI1_HUMAN
Q86VQ1	2	4.8182	R.GPSPSSPTPPAAAAPAEQAPR.A	2	GLCI1_HUMAN
Glucose-6-phosphate isomerase - Homo sapiens (Human)					
P06744	3	4.0088	K.KIEPELDGSAQVTSHDASTNGLINFIK.Q	1	G6PI_HUMAN
Glucosidase 2 subunit beta precursor - Homo sapiens (Human)					
P14314	2	3.6539	K.ETMVTSTTEPSR.C	2	GLU2B_HUMAN
P14314	2	3.6265	R.SLKDMEESIR.N	9	GLU2B_HUMAN
P14314	2	2.7216	R.NKFEEAER.S	1	GLU2B_HUMAN
P14314	4	4.7795	K.LGGSPTSLGTWGSWIGPDHDKFSAMK.Y	2	GLU2B_HUMAN
P14314	2	3.1524	K.ETM#VTSTTEPSR.C	2	GLU2B_HUMAN
P14314	2	3.6786	K.YEQGTGCWQGPNR.S	3	GLU2B_HUMAN
Glutamate dehydrogenase 1, mitochondrial precursor - Homo sapiens (Human)					
P00367	2	4.3921	R.GASIVEDKLVEDLR.T	2	DHE3_HUMAN
Glutaminase kidney isoform, mitochondrial precursor - Homo sapiens (Human)					
O94925	2	3.4594	K.DGPGETDAFGNSEGK.E	2	GLSK_HUMAN
Glutamyl aminopeptidase - Homo sapiens (Human)					
Q07075	2	3.2366	R.ASLIDDAFALAR.A	2	AMPE_HUMAN
Q07075	3	3.9304	R.VNYEVATWDSIATALSLNHK.T	1	AMPE_HUMAN
Glutaredoxin-1 - Homo sapiens (Human)					
P35754	2	3.6449	R.LKQIGALQ.-	6	GLRX1_HUMAN
P35754	2	3.5876	R.RAQEILSQLPIK.Q	2	GLRX1_HUMAN
P35754	2	4.2529	R.AQEILSQLPIK.Q	4	GLRX1_HUMAN
P35754	2	3.6318	K.DCIGGCSDLVSLQQSGELLTR.L	2	GLRX1_HUMAN
Glutaredoxin-related protein 5 - Homo sapiens (Human)					
Q86SX6	3	4.8678	R.DYAAYNVLDDPELR.Q	3	GLRX5_HUMAN
Q86SX6	2	3.9284	K.LGIHSALLDEKKDQDSK.-	1	GLRX5_HUMAN
Glutathione reductase, mitochondrial precursor - Homo sapiens (Human)					
P00390	3	4.1159	K.ADFDNTVAIHPTSSEELVTLR.-	1	GSHR_HUMAN
Glutathione S-transferase P - Homo sapiens (Human)					
P09211	3	4.9566	K.ALPGQLKPFETLLSQNQGGK.T	2	GSTP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Glyceraldehyde-3-phosphate dehydrogenase - Homo sapiens (Human)					
P04406	2	4.5706	R.VVDLMAHMASKE.-	2	G3P_HUMAN
P04406	2	4.5323	K.LISWYDNEFGYSNR.V	12	G3P_HUMAN
P04406	2	3.8913	R.VVDLM#AHM#ASKE.-	4	G3P_HUMAN
P04406	2	4.2421	R.VVDLMAHM#ASKE.-	3	G3P_HUMAN
Glycine cleavage system H protein, mitochondrial precursor - Homo sapiens (Human)					
P23434	2	2.9195	K.SCYEDGWLIK.M	2	GCSH_HUMAN
P23434	3	3.9008	K.MTLSNPSELDELMSEEAYEKYIKSIEE.-	2	GCSH_HUMAN
P23434	2	3.3324	K.M#TLSNPSELDELM#SEEAYEK.Y	2	GCSH_HUMAN
P23434	3	3.878	K.MTLSNPSELDELM#SEEAYEK.Y	2	GCSH_HUMAN
P23434	3	3.784	K.MTLSNPSELDELMSEEAYEK.Y	1	GCSH_HUMAN
Glycogen phosphorylase, liver form - Homo sapiens (Human)					
P06737	3	4.3399	R.TIKEYAQNIWNVEPSDLK.I	1	PYGL_HUMAN
Glycogenin-1 - Homo sapiens (Human)					
P46976	2	2.8303	R.KLDTYLQ.-	1	GLYG_HUMAN
P46976	2	3.4819	R.LVVLATPQVSDSMR.K	1	GLYG_HUMAN
P46976	2	2.8742	K.RPELGVTLTK.L	2	GLYG_HUMAN
P46976	2	2.9427	K.GALVLGSSLK.Q	2	GLYG_HUMAN
P46976	3	6.7271	K.ERWEQGQADYMGADSFDNIKR.K	2	GLYG_HUMAN
P46976	3	3.9754	R.RLVVLATPQVSDSMRK.V	1	GLYG_HUMAN
Glycosyltransferase 25 domain-containing protein 1 - Homo sapiens (Human)					
Q8NBJ5	2	4.3204	K.NSDVLQSPLDSAARDEL.-	2	Q8NBJ5_HUMA
Glycylpeptide N-tetradecanoyltransferase 1 - Homo sapiens (Human)					
P30419	2	4.2968	K.AIELFSVGQGPAK.T	2	NMT1_HUMAN
P30419	2	4.186	K.EKGSETDSAQDQPVK.M	3	NMT1_HUMAN
P30419	2	3.9569	K.GSETDSAQDQPVK.M	4	NMT1_HUMAN
P30419	1	3.1728	R.GGLSPANDTGAK.K	7	NMT1_HUMAN
GMP reductase 1 - Homo sapiens (Human)					
P36959	3	4.3713	K.TVEVPYKGDVENTILDILGGLR.S	1	GMPR1_HUMAN
Golgi phosphoprotein 3 - Homo sapiens (Human)					
Q9H4A6	2	4.2239	R.AAGGGAGSSEDDAQSR.R	2	GOLP3_HUMAN
Golgi phosphoprotein 4 - Homo sapiens (Human)					
O00461	2	3.0702	R.QQEQQQQQVAR.E	3	GOLP4_HUMAN
O00461	2	3.9936	K.FQSPYEEQLEQQR.L	2	GOLP4_HUMAN
Golgi reassembly-stacking protein 2 - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9H8Y8	2	3.2801	K.ADAASSLTVDVTPPTAK.A	2	GORS2_HUMAN
Q9H8Y8	3	5.7696	K.APTTVEDRVGDSTPVSEKPVSAAVDANASES	2	GORS2_HUMAN
Q9H8Y8	2	3.2956	R.VGDSTPVSEKPVSAAVDANASESP.-	1	GORS2_HUMAN
Golgin subfamily A member 1 - Homo sapiens (Human)					
Q92805	2	2.7235	K.NILTAQLQEMK.N	1	GOGA1_HUMAN
Q92805	2	2.9677	R.AADQTTAEQGM#R.Q	2	GOGA1_HUMAN
Q92805	2	3.883	R.TQALEAQIVALER.M	2	GOGA1_HUMAN
Golgin subfamily A member 2 - Homo sapiens (Human)					
Q08379	2	3.0099	K.TFSSTESLR.Q	3	GOGA2_HUMAN
Q08379	2	4.2461	R.VQELETSLAELR.N	3	GOGA2_HUMAN
Q08379	2	3.6824	R.ELKEQLAELQSGFVK.L	2	GOGA2_HUMAN
Q08379	2	2.9303	R.AAELWGEQAEAR.R	2	GOGA2_HUMAN
Q08379	2	3.5069	K.AGMQLNLEELQK.K	2	GOGA2_HUMAN
Q08379	3	3.7686	K.NTQSNEDLKQEKSELEEK.L	1	GOGA2_HUMAN
Q08379	2	3.9054	K.NTQSNEDLKQEK.S	1	GOGA2_HUMAN
Q08379	2	4.6317	K.LLELQELVLR.L	4	GOGA2_HUMAN
Q08379	2	3.6685	K.SQEAQSLQQQR.D	1	GOGA2_HUMAN
Q08379	2	2.7024	R.DNPTAQQIM#QLLR.E	1	GOGA2_HUMAN
Golgin subfamily A member 3 - Homo sapiens (Human)					
Q08378	3	4.4186	R.IQALEAELQAVSHSK.T	1	GOGA3_HUMAN
Q08378	2	3.4375	R.LFSTLDPELM#LNPENLPR.A	2	GOGA3_HUMAN
Q08378	3	4.7369	R.LFSTLDPELMLNPENLPR.A	2	GOGA3_HUMAN
Q08378	2	2.84	R.LGSDLTSAQK.E	1	GOGA3_HUMAN
Q08378	2	3.5626	R.MADSAASLEQQLEQVK.L	1	GOGA3_HUMAN
Q08378	3	3.8665	R.MADSAASLEQQLEQVKLTLLQR.D	1	GOGA3_HUMAN
Q08378	2	4.0429	R.QQMTALQSQLQQVQLER.T	1	GOGA3_HUMAN
Q08378	2	3.1104	R.RLQEALAAK.E	3	GOGA3_HUMAN
Q08378	2	2.8486	R.EQSLDALQTHYDELQAR.L	1	GOGA3_HUMAN
Q08378	2	4.4984	R.QWYQQQLALAQEAR.V	3	GOGA3_HUMAN
Q08378	2	2.9711	R.QDSLSSEVDTLK.Q	1	GOGA3_HUMAN
Q08378	3	3.7417	K.DNTVHDLRQQMTALQSQLQQVQLER.T	1	GOGA3_HUMAN
Q08378	2	4.075	R.DATSKDQLISELK.A	2	GOGA3_HUMAN
Q08378	2	3.1651	K.ASQAEISSLQSVR.Q	2	GOGA3_HUMAN
Q08378	2	2.9139	K.EAADAELGQLR.A	1	GOGA3_HUMAN
Q08378	2	4.1981	K.LTGLGQSNAALR.E	5	GOGA3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q08378	2	3.783	K.NASLASSNNDLQVAEEQYQR.L	1	GOGA3_HUMAN
Q08378	2	3.7064	K.QLTLTQEALQSR.E	2	GOGA3_HUMAN
Q08378	3	3.9829	K.SGQVEHLQQETAALKK.Q	2	GOGA3_HUMAN
Q08378	2	3.3907	K.TLLQQNQQLK.L	5	GOGA3_HUMAN
Q08378	3	4.7955	K.TRLFSTLDPELMLNPENLPR.A	3	GOGA3_HUMAN
Q08378	2	3.0949	R.AMTDLQNMLEAK.N	2	GOGA3_HUMAN
Golgin subfamily A member 5 - Homo sapiens (Human)					
Q8TBA6	2	3.7776	R.SKETQEELNK.A	2	GOGA5_HUMAN
Q8TBA6	2	2.8569	R.KDNASNIYSK.N	1	GOGA5_HUMAN
Golgin subfamily B member 1 - Homo sapiens (Human)					
Q14789	2	2.7252	R.SSSSQTQPLK.V	1	GOGB1_HUMAN
Q14789	3	4.1737	R.KLEEHEESLVGR.A	1	GOGB1_HUMAN
Q14789	3	6.5244	R.APLDPELHQESDMEFNNTTQEDVQER.L	2	GOGB1_HUMAN
Q14789	3	5.8488	R.AQVVDLLQQELTAAEQR.N	1	GOGB1_HUMAN
Q14789	2	3.4883	R.IVGDYQQLEER.H	2	GOGB1_HUMAN
Q14789	3	4.4124	R.KFSDAIQSKEEEIR.L	1	GOGB1_HUMAN
Q14789	3	5.4185	R.APLDPELHQESDM#EFNNTTQEDVQER.L	1	GOGB1_HUMAN
Q14789	2	2.805	R.KLQAALISR.K	1	GOGB1_HUMAN
Q14789	3	4.1362	R.LALLQEERDKLITEMDR.S	2	GOGB1_HUMAN
Q14789	3	4.3318	R.LSGLANVVLHELSGDDDTDQNMR.A	1	GOGB1_HUMAN
Q14789	2	2.9527	R.NETETAEER.V	2	GOGB1_HUMAN
Q14789	2	3.3776	K.TLKEQLNLLSR.A	2	GOGB1_HUMAN
Q14789	4	6.0456	R.RLDYESQTAHDNLLTEQIHSLSIEAK.S	1	GOGB1_HUMAN
Q14789	2	3.6593	K.LSSQITLLEAQNR.T	2	GOGB1_HUMAN
Q14789	3	5.3149	R.TGEADREVSEISIVDIANKR.S	2	GOGB1_HUMAN
Q14789	3	3.7312	R.VKMEYETLSK.K	1	GOGB1_HUMAN
Q14789	2	3.4885	R.NTVETEREESK.I	2	GOGB1_HUMAN
Q14789	3	4.67	K.KEEDVSYLSGQLSEKEAALTK.I	2	GOGB1_HUMAN
Q14789	3	5.0379	K.AKEISNLNQLIEEFKK.N	2	GOGB1_HUMAN
Q14789	2	3.2544	K.AQTEVQLQQK.V	2	GOGB1_HUMAN
Q14789	3	4.0755	K.DVQLQQKDEALQEER.K	2	GOGB1_HUMAN
Q14789	2	4.1394	K.MNLLNQQIQEELSR.V	2	GOGB1_HUMAN
Q14789	2	3.7289	K.IQTEIIEQEDLIK.A	2	GOGB1_HUMAN
Q14789	3	3.8924	K.SLADVESQVSAQNKEKDTVLGR.L	1	GOGB1_HUMAN
Q14789	2	2.9983	K.KLQEALTSR.K	1	GOGB1_HUMAN
Q14789	2	4.139	K.LAAEEQFQALVK.Q	2	GOGB1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q14789	2	4.1313	K.LITSNTDASDGDSVALVK.E	1	GOGB1_HUMAN
Q14789	3	4.4447	K.LKETAEEEKDDLEER.L	2	GOGB1_HUMAN
Q14789	3	6.9884	K.LLQDKNEQAVQSAQTIQQLEDQLQQK.S	2	GOGB1_HUMAN
Q14789	2	3.6001	K.M#NLLNQQIQEELSR.V	2	GOGB1_HUMAN
Q14789	3	4.7898	K.NKEFSQTLENEKNTLLSQISTK.D	1	GOGB1_HUMAN
Q14789	3	3.8555	K.QM#NQTLQDKTNQIDLLQAEISENQAIIQK.L	1	GOGB1_HUMAN
Q14789	2	3.1226	K.GLTAQIQSFGR.S	1	GOGB1_HUMAN
G-protein coupled receptor 98 precursor - Homo sapiens (Human)					
Q8WXG9	2	3.1444	R.FTIQLISIDEVEISPVK.G	1	GPR98_HUMAN
Grainyhead-like protein 1 homolog - Homo sapiens (Human)					
Q9NZI5	2	2.9399	K.ESEEVFDALM#LKTPSLK.G	1	GRHL1_HUMAN
GRAM domain-containing protein 3 - Homo sapiens (Human)					
Q96HH9	3	4.3985	R.SPTAQSPTPSVEADSPDQKK.I	2	GRAM3_HUMAN
GRB2-associated-binding protein 1 - Homo sapiens (Human)					
Q13480	3	5.3549	K.SSGSGSSVADERVDYVVVDQQK.T	1	GAB1_HUMAN
Q13480	2	3.2596	R.SNTISTVDLNK.L	1	GAB1_HUMAN
Q13480	2	4.5571	R.APSASVDSSLYNLPR.S	1	GAB1_HUMAN
GRB2-associated-binding protein 2 - Homo sapiens (Human)					
Q9UQC2	2	4.1776	K.TQALQNTMQEWTDVR.Q	2	GAB2_HUMAN
GRB2-related adapter protein 2 - Homo sapiens (Human)					
O75791	2	3.7595	K.LGLFPANYVAPMTR.-	3	GRAP2_HUMAN
GRINL1A downstream protein Gdown3 - Homo sapiens (Human)					
Q6EEV7	3	4.0116	R.GLGAAEFGGAAGNVEAPGETFAQRK.I	1	Q6EEV7_HUMA
Growth factor receptor-bound protein 2 - Homo sapiens (Human)					
P62993	2	2.824	R.ESESAPGDFSLSVK.F	2	GRB2_HUMAN
P62993	2	2.797	R.NQQIFLR.D	1	GRB2_HUMAN
P62993	2	2.7572	R.AKAEEMLSK.Q	1	GRB2_HUMAN
P62993	3	3.8858	K.FNSLNELVDYHR.S	1	GRB2_HUMAN
GrpE protein homolog 1, mitochondrial precursor - Homo sapiens (Human)					
Q9HAV7	2	3.2433	K.DLLEVADVLEK.A	1	GRPE1_HUMAN
GTPase IMAP family member 1 - Homo sapiens (Human)					
Q8WWP7	2	2.7452	K.MATDEENVYGLEENAQSR.Q	1	GIMA1_HUMAN
GTPase-activating protein ZNF289 - Homo sapiens (Human)					
Q8N6H7	2	4.6503	K.YKDNPFSLGESFGSR.W	6	ZN289_HUMAN
Q8N6H7	2	4.2548	R.LAYQELQIDR.K	8	ZN289_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8N6H7	3	5.0429	R.SSVSHSVLSEMQVIEQETPVSAK.S	3	ZN289_HUMAN
Q8N6H7	2	3.2396	R.SSGLESSEAR.Q	4	ZN289_HUMAN
Q8N6H7	3	4.4603	R.SQLDLFDDVGTFASGPPK.Y	1	ZN289_HUMAN
Q8N6H7	2	2.9874	R.LGMGLVSR.S	2	ZN289_HUMAN
Q8N6H7	2	2.7674	R.LGM#GLVSR.S	1	ZN289_HUMAN
Q8N6H7	2	3.5864	K.QAEESMVASMR.L	3	ZN289_HUMAN
Q8N6H7	2	3.0928	K.QAEESM#VASMR.L	2	ZN289_HUMAN
Q8N6H7	2	4.105	K.LREQQAADAKK.Q	2	ZN289_HUMAN
Q8N6H7	2	3.3714	K.LREQQAADAK.K	1	ZN289_HUMAN
Q8N6H7	2	2.947	K.KLQNLEGK.K	2	ZN289_HUMAN
Q8N6H7	2	3.2686	K.AISSDMFFGR.E	3	ZN289_HUMAN
Q8N6H7	2	3.3776	K.VSSQSFSEIER.Q	3	ZN289_HUMAN
GTP-binding nuclear protein Ran - Homo sapiens (Human)					
P62826	3	4.3475	K.SNYNFEKPFLWLAR.K	1	RAN_HUMAN
GTP-binding protein 1 - Homo sapiens (Human)					
O00178	3	7.9849	K.RDEGGPSGGPAVGAPPPGDEASSVGAGQP	1	GTPB1_HUMAN
Guanine nucleotide-binding protein G(I)/G(S)/G(O) gamma-12 subunit precursor - Homo sapiens (					
Q9UBI6	2	4.0199	K.TASTNNIAQAR.R	3	GBG12_HUMAN
Q9UBI6	2	2.8898	R.RTVQQLR.L	1	GBG12_HUMAN
Guanylate cyclase soluble subunit alpha-3 - Homo sapiens (Human)					
Q02108	2	3.5918	K.DVEDGNANFLGK.A	1	GCYA3_HUMAN
H/ACA ribonucleoprotein complex subunit 2 - Homo sapiens (Human)					
Q9NX24	3	3.8122	R.TYQELLVNQNPIAQPLASR.R	1	NOLA2_HUMAN
H/ACA ribonucleoprotein complex subunit 4 - Homo sapiens (Human)					
O60832	2	4.5263	K.AGLESGAEPGDGDSDTTK.K	9	DKC1_HUMAN
O60832	3	3.8989	K.AGLESGAEPGDGDSDTTKK.K	1	DKC1_HUMAN
O60832	1	2.7191	K.APQVVAEAAK.T	1	DKC1_HUMAN
Haptoglobin precursor [Contains: Haptoglobin alpha chain; Haptoglobin beta chain] - Homo sapie					
P00738	2	4.9684	K.LRTEGDGVYTLNNEK.Q	1	HPT_HUMAN
P00738	3	4.263	K.LRTEGDGVYTLNNEKQWINK.A	1	HPT_HUMAN
P00738	2	3.5436	R.TEGDGVYTLNNEK.Q	4	HPT_HUMAN
P00738	2	3.0903	R.TEGDGVYTLNNEKQWINK.A	1	HPT_HUMAN
HBS1-like protein - Homo sapiens (Human)					
Q9Y450	2	4.7207	K.GPPIEDAIASSDVLETASK.S	2	HBS1L_HUMAN
Q9Y450	3	4.1667	K.SANPPHTIQASEEQSSTPAPVKK.S	1	HBS1L_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
HCLS1-binding protein 3. - Homo sapiens (Human)					
Q53T59	2	3.8051	R.LTIFDEEVDPDEGLFGPGR.K	1	Q53T59_HUMAN
HCV F-transactivated protein 1 - Homo sapiens (Human)					
Q6J203	2	2.7316	R.RGFSEGGR.Q	1	Q6J203_HUMAN
HD domain-containing protein 3 - Homo sapiens (Human)					
Q8N4P3	3	4.5256	R.RLVEEVTDDKTLPK.L	1	HDDC3_HUMAN
Headcase protein homolog - Homo sapiens (Human)					
Q9UBI9	3	6.0385	R.SPGGSPGQSPPTGYSILSPAHFSGPR.S	1	HDC_HUMAN
Heat shock 70 kDa protein 4 - Homo sapiens (Human)					
P34932	3	3.8031	K.IRFQESEERPK.L	1	HSP74_HUMAN
P34932	3	4.3938	K.NKEDQYDHLDAADMTK.V	2	HSP74_HUMAN
P34932	1	2.7129	K.NLGQPIK.I	3	HSP74_HUMAN
P34932	2	3.1297	K.STNEAMEWMNNK.L	2	HSP74_HUMAN
P34932	2	3.8051	K.STNEAMEWMNNKLNLQNK.Q	1	HSP74_HUMAN
P34932	2	2.9117	R.DKLSGEYEK.F	1	HSP74_HUMAN
Heat shock cognate 71 kDa protein - Homo sapiens (Human)					
P11142	2	4.3039	K.NSLESYAFNM#K.A	6	HSP7C_HUMAN
P11142	2	4.173	K.NSLESYAFNMK.A	8	HSP7C_HUMAN
Heat shock factor-binding protein 1 - Homo sapiens (Human)					
O75506	3	6.5321	R.IDDLEKNIADLMTQAGVEELESENKIPATQK.S	2	HSBP1_HUMAN
O75506	3	4.2432	K.TVQDLTSVVQTLLQQMQDK.F	1	HSBP1_HUMAN
Heat shock protein 90Ae - Homo sapiens (Human)					
Q58FG0	3	3.984	R.M#IKLGLGVDEYDPTANDINAAITKEMPPLR.G	1	Q58FG0_HUMA
Heat shock protein beta-1 - Homo sapiens (Human)					
P04792	2	6.0532	R.AQLGGPEAAKSDETAAK.-	14	HSPB1_HUMAN
P04792	2	2.9477	R.QLSSGVSEIR.H	2	HSPB1_HUMAN
P04792	2	5.4446	R.PLPPAAIESPAVAAPAYSR.A	12	HSPB1_HUMAN
P04792	2	4.2523	R.LFDQAFGLPR.L	7	HSPB1_HUMAN
P04792	2	4.395	K.HEERQDEHGYISR.C	3	HSPB1_HUMAN
P04792	3	7.6576	R.KYTLPPGVDPTQVSSSLSPEGTLTVEAPMPK.	6	HSPB1_HUMAN
P04792	2	5.2546	R.VSLDVNHFAPDELTVK.T	13	HSPB1_HUMAN
P04792	3	5.9862	R.KYTLPPGVDPTQVSSSLSPEGTLTVEAPM#P	4	HSPB1_HUMAN
P04792	3	3.7004	R.GPSWDPFRDWYPHSR.L	1	HSPB1_HUMAN
P04792	1	2.5629	R.AQLGGPEAAK.S	2	HSPB1_HUMAN
P04792	3	6.4212	K.YTLPPGVDPTQVSSSLSPEGTLTVEAPM#PK.	1	HSPB1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P04792	3	4.9323	K.TKDGVVEITGKHEER.Q	8	HSPB1_HUMAN
P04792	2	4.2001	K.LATQSNEITIPVTFESR.A	54	HSPB1_HUMAN
P04792	3	3.9608	K.DGVVEITGKHEERQDEHGYISR.C	1	HSPB1_HUMAN
P04792	2	4.2216	K.DGVVEITGKHEER.Q	4	HSPB1_HUMAN
P04792	2	3.9372	K.DGVVEITGK.H	10	HSPB1_HUMAN
P04792	2	4.0239	K.TKDGVVEITGK.H	12	HSPB1_HUMAN
Heat shock protein beta-6 - Homo sapiens (Human)					
O14558	2	2.9841	K.HFSPEEIAVK.V	2	HSPB6_HUMAN
O14558	3	5.7427	R.YRLPPGVDPAAVTSALSPEGVLSIQAAPASA	1	HSPB6_HUMAN
Heat shock protein beta-8 - Homo sapiens (Human)					
Q9UJY1	3	4.4725	K.HEEKQQEGGIVSK.N	2	HSPB8_HUMAN
Heat shock protein HSP 90-beta - Homo sapiens (Human)					
P08238	3	4.0822	K.HLEINPDHPIVETLR.Q	1	HS90B_HUMAN
P08238	2	3.2323	R.NPDDITQEEYGEFYK.S	1	HS90B_HUMAN
Heat shock-related 70 kDa protein 2 - Homo sapiens (Human)					
P54652	2	2.8432	K.NALESYTYNIK.Q	2	HSP72_HUMAN
Hematological and neurological expressed 1 protein - Homo sapiens (Human)					
Q9UK76	2	4.1996	K.M#ASNIFGTPEENQASWAK.S	1	HN1_HUMAN
Q9UK76	2	4.4695	K.SSGGREDLESSGLQR.R	7	HN1_HUMAN
Q9UK76	3	6.5639	R.VLRPPGGGSNFSLGFDEPTEQPVRK.N	6	HN1_HUMAN
Hematological and neurological expressed 1-like protein - Homo sapiens (Human)					
Q9H910	3	4.4255	R.M#ASNIFGPTEEPQNIPKR.T	1	HN1L_HUMAN
Q9H910	2	2.7066	R.SIPAGAEPGEKGSAR.K	1	HN1L_HUMAN
Q9H910	2	3.4604	R.MASNIFGPTEEPQNIPK.R	4	HN1L_HUMAN
Q9H910	2	4.1181	R.M#ASNIFGPTEEPQNIPK.R	6	HN1L_HUMAN
Q9H910	1	2.4033	R.LAHPNKPK.D	1	HN1L_HUMAN
Q9H910	3	5.5024	R.KAGPAKEQEPMPTVDSHEPR.L	4	HN1L_HUMAN
Q9H910	2	4.8778	K.TSDIFGSPVTATSR.L	7	HN1L_HUMAN
Q9H910	2	2.964	K.EQEPMPTVDSHEPR.L	1	HN1L_HUMAN
Q9H910	2	2.8366	K.EQEPM#PTVDSHEPR.L	1	HN1L_HUMAN
Q9H910	3	3.8101	K.DHVFLCEGEEPKSDLK.A	1	HN1L_HUMAN
Q9H910	3	4.343	K.AGPAKEQEPMPTVDSHEPR.L	2	HN1L_HUMAN
Q9H910	1	2.6283	R.SIPAGAEPGEK.G	1	HN1L_HUMAN
Hematopoietic lineage cell-specific protein - Homo sapiens (Human)					
P14317	3	4.8686	R.VDKSAVGFNEMEAPTTAYK.K	3	HCLS1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P14317	3	4.8078	K.SAVGHEYVAEVEKHSSQTDAAK.G	3	HCLS1_HUMAN
P14317	3	4.8356	K.SVVGHDVSVSVETQGDDWDTDPDFVNDISE	2	HCLS1_HUMAN
P14317	3	6.0584	K.SVVGHDVSVSVETQGDDWDTDPDFVNDISE	2	HCLS1_HUMAN
P14317	2	3.8209	K.TTPIEAASSGAR.G	5	HCLS1_HUMAN
P14317	3	3.8406	R.ADKSAVGFDYKGEVEK.H	1	HCLS1_HUMAN
P14317	2	5.0697	K.SAVGHEYVAEVEK.H	5	HCLS1_HUMAN
P14317	2	2.9191	R.EHPVPLLPIR.Q	1	HCLS1_HUMAN
P14317	3	5.7341	R.MDKSAVGHEYVAEVEKHSSQTDAAK.G	1	HCLS1_HUMAN
P14317	2	3.8353	R.NKVSEEHDVLR.K	3	HCLS1_HUMAN
P14317	2	4.1485	R.QTLPEDNEEPPALPPR.T	14	HCLS1_HUMAN
P14317	2	3.3808	R.TEHINIHQLR.N	1	HCLS1_HUMAN
P14317	2	4.6876	K.SAVGFNEMEAPTTAYKK.T	8	HCLS1_HUMAN
P14317	3	4.4352	R.VDKSAVGFNEM#EAPTTAYKK.T	2	HCLS1_HUMAN
P14317	2	3.4886	R.EEEEKAQQVAR.R	5	HCLS1_HUMAN
P14317	3	4.9575	R.VDKSAVGFNEMEAPTTAYKK.T	4	HCLS1_HUMAN
P14317	3	4.7127	R.VDKSAVGFNEM#EAPTTAYK.K	1	HCLS1_HUMAN
P14317	1	3.8566	K.GFGGQYGIQK.D	12	HCLS1_HUMAN
P14317	3	5.6171	R.MDKSAVGHEYVAEVEK.H	6	HCLS1_HUMAN
P14317	2	4.9548	K.AALGYDYKGETEKHESQR.D	3	HCLS1_HUMAN
P14317	2	3.1187	K.AKFESM#AEEK.R	2	HCLS1_HUMAN
P14317	2	4.064	K.AKFESMAEEKR.K	2	HCLS1_HUMAN
P14317	2	3.6711	K.SAVGFNEMEAPTTAYK.K	1	HCLS1_HUMAN
P14317	2	2.8144	K.FESMAEEKR.K	1	HCLS1_HUMAN
P14317	2	3.1564	K.GFGGQYGIQKDR.V	1	HCLS1_HUMAN
P14317	2	2.7741	K.ISSEAWPPVGTPPSSESEPVR.T	1	HCLS1_HUMAN
P14317	3	4.6225	K.KISSEAWPPVGTPPSSESEPVR.T	5	HCLS1_HUMAN
P14317	2	3.3473	K.KTTPIEAASSGAR.G	1	HCLS1_HUMAN
P14317	3	4.3999	K.REEEEKAQQVAR.R	2	HCLS1_HUMAN
P14317	3	5.1579	K.RSPEAPQPVIAM#EEPAVPAPLPK.K	4	HCLS1_HUMAN
P14317	3	5.37	K.RSPEAPQPVIAMEEPAVPAPLPK.K	2	HCLS1_HUMAN
P14317	3	5.0768	K.RSPEAPQPVIAMEEPAVPAPLPKK.I	2	HCLS1_HUMAN
P14317	2	4.6694	K.SAVGFDYKGEVEK.H	5	HCLS1_HUMAN
P14317	2	3.8095	K.SAVGFNEM#EAPTTAYKK.T	3	HCLS1_HUMAN
P14317	2	3.0307	K.FESM#AEEK.R	1	HCLS1_HUMAN
Hematopoietic stem/progenitor cell induced protein 2 - Homo sapiens (Human)					
Q6PN88	2	3.2799	R.RTAGAVGKAMGLEVR.A	1	Q6PN88_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Heme oxygenase 1 - Homo sapiens (Human)					
P09601	2	2.8774	K.VQDSAPVETPR.G	2	HMOX1_HUMAN
Heme-binding protein 1 - Homo sapiens (Human)					
Q9NRV9	2	3.1806	K.EADYVAQATR.L	1	HEBP1_HUMAN
Q9NRV9	2	2.7907	K.FATVEVTDKPVDEALR.E	1	HEBP1_HUMAN
Heme-binding protein 2 - Homo sapiens (Human)					
Q9Y5Z4	2	4.5201	R.NNEVWLIQKNEPTKENE.-	4	HEBP2_HUMAN
Hemoglobin subunit beta - Homo sapiens (Human)					
P68871	2	3.0175	K.EFTPPVQAAYQK.V	3	HBB_HUMAN
Hemoglobin subunit delta - Homo sapiens (Human)					
P02042	2	4.7245	R.NFGKEFTPQMQAAYQK.V	2	HBD_HUMAN
P02042	2	3.9102	R.LLGNVLVCVLAR.N	2	HBD_HUMAN
Hemopexin precursor - Homo sapiens (Human)					
P02790	3	3.9193	K.GDKVWVYPPEKK.E	1	HEMO_HUMAN
P02790	2	3.812	K.GGYTLVSGYPK.R	3	HEMO_HUMAN
P02790	3	4.4558	K.SGAQATWTELPWPHEK.V	1	HEMO_HUMAN
P02790	3	4.4576	R.WKNFPSPVDAAFR.Q	6	HEMO_HUMAN
P02790	2	3.5237	R.YYCFQGNQFLR.F	5	HEMO_HUMAN
Hepatitis B virus X-interacting protein - Homo sapiens (Human)					
O43504	2	4.3327	K.HDGITVAVHK.M	5	XIP_HUMAN
O43504	3	3.7687	K.LTSDPTDIPVVCLESDNGNIM#IQK.H	1	XIP_HUMAN
O43504	2	7.3582	R.GTLSDEHAGVISVLAQQAAK.L	5	XIP_HUMAN
Hepatocyte nuclear factor 1-alpha - Homo sapiens (Human)					
P20823	3	5.203	K.LSQLQTELLAALLESGLSK.E	1	HNF1A_HUMAN
P20823	2	3.3341	K.AVVETLLQEDPWR.V	1	HNF1A_HUMAN
Hepatocyte nuclear factor 1-beta - Homo sapiens (Human)					
P35680	2	3.882	K.ELQALNTEEAAEQR.A	1	HNF1B_HUMAN
P35680	3	4.624	K.LTSLQQELLSALLSSGVTK.E	2	HNF1B_HUMAN
Hepatoma-derived growth factor - Homo sapiens (Human)					
P51858	3	3.735	K.MKGYPHWPARIDEMPEAAVK.S	1	HDGF_HUMAN
P51858	2	3.3835	R.IDEM#PEAAVK.S	3	HDGF_HUMAN
P51858	1	3.2505	R.IDEMPEAAVK.S	10	HDGF_HUMAN
P51858	2	3.0677	R.IDEMPEAAVKSTANK.Y	1	HDGF_HUMAN
Hepatoma-derived growth factor-related protein 3 - Homo sapiens (Human)					
Q9Y3E1	1	2.7575	R.NTTSDLQK.T	4	HDGR3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y3E1	2	3.4213	K.AGDLVFAK.M	3	HDGR3_HUMAN
Q9Y3E1	3	4.1221	K.EEENKSSSEGGDAGNDTR.N	1	HDGR3_HUMAN
Q9Y3E1	2	5.1909	K.GFNEGLWEIENNPGVK.F	12	HDGR3_HUMAN
Q9Y3E1	3	4.8788	K.RKGFNEGLWEIENNPGVK.F	2	HDGR3_HUMAN
Q9Y3E1	3	5.2002	R.KGFNEGLWEIENNPGVK.F	6	HDGR3_HUMAN
Hepatopoietin protein - Homo sapiens (Human)					
Q9H290	2	6.469	R.DAAASASTPAQAPTSDSPVAEDASR.R	6	Q9H290_HUMAN
Heterogeneous nuclear ribonucleoprotein A/B - Homo sapiens (Human)					
Q99729	1	2.3145	K.DAASVEK.V	4	ROAA_HUMAN
Q99729	1	2.1936	K.DLKDYFTK.F	1	ROAA_HUMAN
Q99729	2	4.5342	K.EVYQQQQYGSGGR.G	4	ROAA_HUMAN
Q99729	2	3.2569	K.M#FVGGLSWDTSKK.D	2	ROAA_HUMAN
Q99729	2	4.2236	K.MFVGGLSWDTSK.K	5	ROAA_HUMAN
Q99729	2	3.7963	K.MFVGGLSWDTSKK.D	3	ROAA_HUMAN
Q99729	2	3.2043	K.VLDQKEHR.L	18	ROAA_HUMAN
Q99729	1	3.0177	R.GFGFILFK.D	5	ROAA_HUMAN
Q99729	2	3.8761	R.GFVFITFKEEEPVKK.V	1	ROAA_HUMAN
Heterogeneous nuclear ribonucleoprotein A0 - Homo sapiens (Human)					
Q13151	2	3.5467	K.AEIIADKQSGK.K	4	ROA0_HUMAN
Q13151	2	5.1732	K.SGGGGGGGGSSWGGR.S	4	ROA0_HUMAN
Q13151	3	4.3221	K.RGFGFVYFQNHDAADK.A	1	ROA0_HUMAN
Q13151	3	4.3291	K.KAVPKEDIYSGGGGGGSR.S	1	ROA0_HUMAN
Q13151	3	4.5521	K.AVPKEDIYSGGGGGGSR.S	4	ROA0_HUMAN
Q13151	2	4.0462	K.GDVAEGDLIEHFSQFGTVEK.A	2	ROA0_HUMAN
Heterogeneous nuclear ribonucleoprotein A3 - Homo sapiens (Human)					
P51991	2	3.9644	R.GFAFVTFDDHDTVDK.I	1	ROA3_HUMAN
Heterogeneous nuclear ribonucleoprotein D0 - Homo sapiens (Human)					
Q14103	2	5.6964	K.IDASKNEEDEGHSNSSPR.H	5	HNRPD_HUMAN
Q14103	2	4.2535	R.HSEAATAQREEWK.M	2	HNRPD_HUMAN
Heterogeneous nuclear ribonucleoprotein D-like - Homo sapiens (Human)					
O14979	3	4.2695	K.NQQDDGKMFIGGLSWDTSKK.D	1	HNRDL_HUMAN
O14979	3	4.0581	K.NQQDDGKM#FIGGLSWDTSKK.D	2	HNRDL_HUMAN
O14979	2	2.9539	R.QLAPLLPSLAPSSAR.Q	1	HNRDL_HUMAN
O14979	2	3.8383	K.DAASVDKVLELK.E	4	HNRDL_HUMAN
O14979	2	2.9285	K.KDLTEYLSR.F	2	HNRDL_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O14979	2	3.1804	K.DLTEYLSR.F	4	HNRDL_HUMAN
O14979	3	4.5255	R.FGEVVDCTIKTDPVTGR.S	1	HNRDL_HUMAN
O14979	2	4.0234	K.MFIGGLSWDTSKK.D	3	HNRDL_HUMAN
Heterogeneous nuclear ribonucleoprotein F - Homo sapiens (Human)					
P52597	2	3.7123	K.HSGPNSADSANDGFVR.L	2	HNRPF_HUMAN
P52597	3	4.4831	R.M#RPGAYSTGYGGYEEYSGLSDGYGFTTDL	1	HNRPF_HUMAN
P52597	3	4.9876	R.MRPGAYSTGYGGYEEYSGLSDGYGFTTDLF	1	HNRPF_HUMAN
P52597	2	3.7339	R.QSGEAFVELGSEDDVK.M	2	HNRPF_HUMAN
Heterogeneous nuclear ribonucleoprotein H' - Homo sapiens (Human)					
P55795	2	4.8813	R.ATENDIYNFFSPLNPMR.V	4	HNRH2_HUMAN
Heterogeneous nuclear ribonucleoprotein H3 - Homo sapiens (Human)					
P31942	2	4.2492	R.ATGEADVEFVTHEDAVAAMSK.D	1	HNRH3_HUMAN
P31942	2	3.0463	R.VHIDIGADGR.A	1	HNRH3_HUMAN
P31942	2	3.7537	R.DGM#DNQGGYGSVGR.M	7	HNRH3_HUMAN
P31942	3	4.2664	R.ATGEADVEFVTHEDAVAAM#SK.D	2	HNRH3_HUMAN
P31942	2	3.2225	K.HNGPNDASDGTVR.L	2	HNRH3_HUMAN
P31942	2	3.992	R.DGMDNQGGYGSVGR.M	4	HNRH3_HUMAN
Heterogeneous nuclear ribonucleoprotein L - Homo sapiens (Human)					
P14866	1	3.1161	R.YYGGGSEGGR.A	3	HNRPL_HUMAN
Heterogeneous nuclear ribonucleoprotein M - Homo sapiens (Human)					
P52272	2	3.6867	R.MGAGM#GFGLER.M	3	HNRPM_HUMAN
P52272	2	3.8174	R.AFITNIPFDVK.W	3	HNRPM_HUMAN
P52272	1	2.2704	R.VGQTIER.M	3	HNRPM_HUMAN
P52272	2	5.3533	R.MGPLGLDHMASSIER.M	3	HNRPM_HUMAN
P52272	2	3.1601	R.MGPAMGPALGAGIER.M	1	HNRPM_HUMAN
P52272	2	4.2621	R.MGPAM#GPALGAGIER.M	4	HNRPM_HUMAN
P52272	2	4.282	R.MGLAMGGGGGASFDR.A	3	HNRPM_HUMAN
P52272	2	4.2951	R.MGLAM#GGGGGASFDR.A	2	HNRPM_HUMAN
P52272	2	3.2214	R.MGANNLER.M	3	HNRPM_HUMAN
P52272	2	4.2258	R.MGAGMGFGLER.M	3	HNRPM_HUMAN
P52272	2	3.0209	K.MGGMEGPFGGGMENM#GR.F	1	HNRPM_HUMAN
P52272	2	3.6609	R.M#GLAMGGGGGASFDR.A	2	HNRPM_HUMAN
P52272	2	3.6741	R.M#GLAM#GGGGGASFDR.A	2	HNRPM_HUMAN
P52272	2	2.9551	R.M#GANNLER.M	1	HNRPM_HUMAN
P52272	2	3.4371	R.M#GAGMGFGLER.M	2	HNRPM_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P52272	2	3.527	K.GIGMGNIGPAGM#GMEGIGFGINK.M	1	HNRPM_HUMAN
P52272	3	4.0002	R.GNFGGSFAGSFGGAGGHAPGVAR.K	2	HNRPM_HUMAN
P52272	2	2.7018	R.INEILSNALK.R	1	HNRPM_HUMAN
P52272	2	3.2258	K.MGGMEGPFGGGMENMGR.F	2	HNRPM_HUMAN
P52272	2	3.9944	R.M#GPAM#GPALGAGIER.M	2	HNRPM_HUMAN
Heterogeneous nuclear ribonucleoprotein Q - Homo sapiens (Human)					
O60506	2	4.6959	R.NLANTVTEEILEK.A	4	HNRPQ_HUMAN
O60506	2	3.1515	R.DLFEDELVPLFEK.A	1	HNRPQ_HUMAN
O60506	3	5.32	R.AIEALKEFNEDGALAVLQQFK.D	2	HNRPQ_HUMAN
O60506	2	3.5881	K.DSDLSHVQNK.S	4	HNRPQ_HUMAN
O60506	2	3.6758	K.VADSSKGPDEAK.I	6	HNRPQ_HUMAN
Heterogeneous nuclear ribonucleoprotein U - Homo sapiens (Human)					
Q00839	2	3.7939	K.VSELKEELKK.R	6	HNRPU_HUMAN
Q00839	2	2.7192	R.LSDKGLK.A	1	HNRPU_HUMAN
Q00839	3	6.6643	R.LQAALDDEEAGGRPAMEPGNGSLDLGGDSA	3	HNRPU_HUMAN
Q00839	2	2.8998	K.GLKAELMER.L	3	HNRPU_HUMAN
Q00839	3	5.6538	R.LQAALDDEEAGGRPAM#EPGNGSLDLGGDS	2	HNRPU_HUMAN
Q00839	2	4.4609	R.LSDKGLKAELMER.L	5	HNRPU_HUMAN
Q00839	2	4.2165	K.LKVSELKEELK.K	4	HNRPU_HUMAN
Q00839	2	3.0536	K.GLKAELM#ER.L	5	HNRPU_HUMAN
Q00839	3	4.3031	K.AEGGGGGGRPGAPAGDGKTEQK.G	4	HNRPU_HUMAN
Q00839	3	3.7652	K.SSGPTSLFAVTVAPPGAR.Q	4	HNRPU_HUMAN
Q00839	2	3.5898	K.VSELKEELK.K	4	HNRPU_HUMAN
Heterogeneous nuclear ribonucleoprotein U-like protein 1 - Homo sapiens (Human)					
Q9BUJ2	3	4.5062	R.RPLEMEQQQAYRPEMK.T	1	HNRL1_HUMAN
Q9BUJ2	1	2.3362	R.NPPGASTYNK.N	1	HNRL1_HUMAN
Q9BUJ2	3	4.4229	R.QNQFYDTQVIKQENESGYER.R	5	HNRL1_HUMAN
Heterogeneous nuclear ribonucleoprotein U-like protein 2 - Homo sapiens (Human)					
Q1KMD3	2	3.5487	K.AAEEQGDDQDSEK.S	8	HNRL2_HUMAN
Q1KMD3	3	4.2486	R.SGDETPGSEVPGDKAAEEQGDDQDSEK.S	1	HNRL2_HUMAN
Q1KMD3	2	3.1409	K.REEDEPEER.S	2	HNRL2_HUMAN
Heterogeneous nuclear ribonucleoproteins A2/B1 - Homo sapiens (Human)					
P22626	2	3.3774	R.GGNFGFGDSR.G	6	ROA2_HUMAN
P22626	2	3.0654	R.QEMQEVQSSR.S	2	ROA2_HUMAN
P22626	2	5.103	R.NMGGPYGGGNYGPGGSGGSGGYGGR.S	6	ROA2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P22626	2	5.1	R.NM#GGPYGGGNYGPGGSGGSGGYGGR.S	3	ROA2_HUMAN
P22626	3	4.6957	R.GGNFGFGDSRGGGGNFGPGPGSNFR.G	1	ROA2_HUMAN
P22626	2	4.1698	R.GFGDGYNGYGGGPGGGNFGGSPGYGGGR.	2	ROA2_HUMAN
P22626	1	3.1151	K.SGNFGGSR.N	12	ROA2_HUMAN
P22626	2	4.942	K.ALSRQEMQEVQSSR.S	2	ROA2_HUMAN
P22626	3	3.9735	K.ALSRQEM#QEVQSSR.S	1	ROA2_HUMAN
P22626	2	3.4781	R.GGGGNFGPGPGSNFR.G	12	ROA2_HUMAN
Heterogeneous nuclear ribonucleoproteins C1/C2 - Homo sapiens (Human)					
P07910	2	2.888	K.QAVEMKNDKSEEEQSSSSVK.K	1	HNRPC_HUMAN
P07910	2	2.8729	K.SEEEQSSSSVK.K	3	HNRPC_HUMAN
P07910	3	4.0097	K.QAVEM#KNDKSEEEQSSSSVK.K	2	HNRPC_HUMAN
P07910	3	4.8388	K.NDKSEEEQSSSSVKK.D	3	HNRPC_HUMAN
P07910	2	4.3373	K.NDKSEEEQSSSSVK.K	6	HNRPC_HUMAN
P07910	2	4.3302	K.GFAFVQYVNER.N	4	HNRPC_HUMAN
P07910	3	4.5601	R.MIAGQVLDINLAAEPK.V	2	HNRPC_HUMAN
Hexamethylene bis-acetamide-inducible protein 1. - Homo sapiens (Human)					
O94992	2	5.141	K.LGAPAAGGEEEWGQQQR.Q	5	O94992_HUMAN
O94992	2	4.0934	K.QELIKEYLELEK.C	2	O94992_HUMAN
O94992	2	4.9218	R.AENLQLLTENELHR.Q	2	O94992_HUMAN
O94992	3	5.4037	R.LRAENLQLLTENELHR.Q	1	O94992_HUMAN
O94992	2	3.3893	R.YHTESLQNMSK.Q	2	O94992_HUMAN
Hexamethylene bis-acetamide-inducible protein 2. - Homo sapiens (Human)					
Q96MH2	2	3.6375	R.QVEELAAEVQR.L	2	Q96MH2_HUMA
High mobility group nucleosome-binding domain-containing protein 3 - Homo sapiens (Human)					
Q15651	2	3.3545	K.EGTAPSENGETKAEEAQK.T	2	HMGN3_HUMAN
Q15651	1	2.4883	K.TESVDNEGE.-	4	HMGN3_HUMAN
Q15651	2	4.2498	K.AEEAQKTESVDNEGE.-	3	HMGN3_HUMAN
Q15651	3	4.7866	K.QEAGKEGTAPSENGETKAEEAQK.T	1	HMGN3_HUMAN
Q15651	3	4.7563	K.EGTAPSENGETKAEEAQKTESVDNEGE.-	5	HMGN3_HUMAN
High mobility group nucleosome-binding domain-containing protein 4 - Homo sapiens (Human)					
O00479	1	2.1664	K.KGEKLPK.G	1	HMGN4_HUMAN
O00479	2	4.0167	K.NRDASTLQSQK.A	11	HMGN4_HUMAN
O00479	3	5.4865	K.NRDASTLQSQKAEGTGDAK.-	11	HMGN4_HUMAN
O00479	2	3.2199	R.DASTLQSQK.A	7	HMGN4_HUMAN
O00479	3	5.1593	R.DASTLQSQKAEGTGDAK.-	20	HMGN4_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O00479	3	4.5441	R.LSAKPAPPKPEPR.P	6	HMGN4_HUMAN
High mobility group protein 20A - Homo sapiens (Human)					
Q9NP66	3	4.3406	R.LHSIILANPQDNENFIATVR.E	1	HM20A_HUMAN
High mobility group protein B1 - Homo sapiens (Human)					
P09429	3	6.1545	K.KKEEEEDEEDEEDEEEEEDEEDEDEEEDDD	1	HMGB1_HUMAN
High mobility group protein B2 - Homo sapiens (Human)					
P26583	2	3.3999	K.KHPDSSVNFAEFSK.K	2	HMGB2_HUMAN
P26583	2	3.7976	K.SEHPGLSIGDTAK.K	5	HMGB2_HUMAN
P26583	3	4.9085	K.LGEMWSEQSAKDKQPYEQK.A	5	HMGB2_HUMAN
P26583	2	3.7766	K.LGEMWSEQSAKDK.Q	2	HMGB2_HUMAN
P26583	1	3.3277	K.LGEMWSEQSAK.D	8	HMGB2_HUMAN
P26583	2	3.9594	K.LGEM#WSEQSAK.D	6	HMGB2_HUMAN
P26583	2	4.8687	K.KLGEMWSEQSAK.D	8	HMGB2_HUMAN
P26583	2	3.8205	K.KLGEMWSEQSAKDK.Q	1	HMGB2_HUMAN
P26583	1	2.7301	K.SKFEDMAK.S	6	HMGB2_HUMAN
P26583	3	4.2351	K.SEHPGLSIGDTAKK.L	8	HMGB2_HUMAN
P26583	2	3.993	K.KHPDSSVNFAEFSKK.C	2	HMGB2_HUMAN
P26583	3	4.0179	K.IKSEHPGLSIGDTAKK.L	3	HMGB2_HUMAN
P26583	2	4.2758	K.IKSEHPGLSIGDTAK.K	4	HMGB2_HUMAN
P26583	3	5.1437	K.HPDSSVNFAEFSKK.C	1	HMGB2_HUMAN
P26583	3	3.8929	K.HPDSSVNFAEFSK.K	2	HMGB2_HUMAN
P26583	2	3.2526	K.EKSKFEDMAK.S	1	HMGB2_HUMAN
P26583	2	3.006	K.DKQPYEQK.A	4	HMGB2_HUMAN
P26583	3	4.6458	K.KLGEM#WSEQSAK.D	9	HMGB2_HUMAN
P26583	3	3.8366	K.KLGEM#WSEQSAKDK.Q	2	HMGB2_HUMAN
High mobility group protein B3 - Homo sapiens (Human)					
O15347	3	4.5297	K.KLGEM#WNNLNDSEKQPYITK.A	1	HMGB3_HUMAN
O15347	3	6.7895	K.KLGEMWNNLNDSEKQPYITK.A	2	HMGB3_HUMAN
O15347	3	4.2536	K.LGEMWNNLNDSEKQPYITK.A	2	HMGB3_HUMAN
High mobility group protein HMG-I/HMG-Y - Homo sapiens (Human)					
P17096	2	4.2517	R.KQPPVSPGTALVGSQK.E	3	HMGA1_HUMAN
P17096	2	4.0696	R.KQPPVSPGTALVGSQKEPSEVPTPK.R	1	HMGA1_HUMAN
HIRA-interacting protein 3 - Homo sapiens (Human)					
Q9BW71	2	3.6243	R.EESEESEAEPVQR.T	2	HIRP3_HUMAN
Q9BW71	3	4.6718	R.GRPDLSTLTHSIVR.R	2	HIRP3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BW71	2	2.9226	R.SHLEPEEK.Q	2	HIRP3_HUMAN
Histidine triad nucleotide-binding protein 1 - Homo sapiens (Human)					
P49773	4	4.9923	K.KHISQISVAEDDDESLLGHLMIVGK.K	1	HINT1_HUMAN
P49773	3	4.367	R.MVVNEGSDGGQSVYHVHLHVLGGR.Q	2	HINT1_HUMAN
P49773	2	3.2063	R.PGGDTIFGK.I	4	HINT1_HUMAN
P49773	3	3.854	K.KHISQISVAEDDDESLLGHLMIVGKK.C	1	HINT1_HUMAN
P49773	4	5.9841	K.IIFEDDRCLAFHDISPQAPTHFLVIPK.K	2	HINT1_HUMAN
P49773	2	2.7947	K.IIFEDDR.C	1	HINT1_HUMAN
P49773	2	3.9409	K.AQVARPGGDTIFGK.I	4	HINT1_HUMAN
P49773	3	6.3475	K.HISQISVAEDDDESLLGHLMIVGK.K	1	HINT1_HUMAN
Histidine triad nucleotide-binding protein 2 - Homo sapiens (Human)					
Q9BX68	2	4.998	K.AQQATPGGAAPTIFSR.I	8	HINT2_HUMAN
Q9BX68	2	2.8081	R.DVAPQAPVHFLVIPK.K	1	HINT2_HUMAN
Q9BX68	2	5.3737	R.ISQAEEEDQQLLGHLLLVAK.Q	2	HINT2_HUMAN
Histidine-rich glycoprotein precursor - Homo sapiens (Human)					
P04196	2	2.9319	R.KYWNDCEPPDSR.R	1	HRG_HUMAN
P04196	3	4.9195	R.VRGGEGTGYFVDFSVR.N	8	HRG_HUMAN
P04196	2	2.9596	R.SSTTKPPFKPHGSR.D	5	HRG_HUMAN
P04196	2	2.9799	K.DSPVLIDFFEDTERYR.K	2	HRG_HUMAN
P04196	3	4.2623	R.RRDGYLFQLLR.I	2	HRG_HUMAN
P04196	2	3.7802	R.RPSEIVIGQCK.V	3	HRG_HUMAN
P04196	2	4.0623	R.RDGYLFQLLR.I	5	HRG_HUMAN
P04196	2	4.404	R.GGEGTGYFVDFSVR.N	7	HRG_HUMAN
P04196	2	3.8237	R.DGYLFQLLR.I	16	HRG_HUMAN
P04196	2	2.8955	K.YWNDCEPPDSR.R	1	HRG_HUMAN
P04196	2	4.5637	K.YKEENDDFASFR.V	9	HRG_HUMAN
P04196	3	4.3924	K.HPLKPDNQPFPQSVSESCPGK.F	1	HRG_HUMAN
P04196	2	2.9686	K.SGFPQVSMFFTHTFPK.-	1	HRG_HUMAN
P04196	3	5.9357	R.KGEVLPLPEANFPSFPLPHHK.H	4	HRG_HUMAN
Histocompatibility - Homo sapiens (Human)					
Q6P189	2	4.993	R.KDGADAVFPGPSLEPPAGSSGVK.A	2	Q6P189_HUMAN
Q6P189	2	3.3091	R.AGSPSPQPSGELPR.K	3	Q6P189_HUMAN
Histone deacetylase 4 - Homo sapiens (Human)					
P56524	2	3.0964	K.GKESAVASTEVK.M	1	HDAC4_HUMAN
Histone H1.0 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07305	2	2.7886	K.RLVTTGVLK.Q	1	H10_HUMAN
P07305	3	4.4333	K.STDHPKYSDMIVAAIQAEK.N	2	H10_HUMAN
P07305	2	3.6993	K.VGENADSQIK.L	15	H10_HUMAN
P07305	2	4.0046	K.YSDM#IVAAIQAEK.N	5	H10_HUMAN
P07305	2	4.6736	K.YSDMIVAAIQAEK.N	6	H10_HUMAN
P07305	1	2.624	R.LVTTGVLK.Q	4	H10_HUMAN
P07305	1	3.0983	K.GVGASGSFR.L	3	H10_HUMAN
Histone H1.1 - Homo sapiens (Human)					
Q02539	3	4.358	K.KKPAGPSVSELIVQAASSSK.E	1	H11_HUMAN
Q02539	2	2.8564	K.ASSVETKPGASK.V	1	H11_HUMAN
Q02539	2	2.7199	K.KASSVETKPGASK.V	1	H11_HUMAN
Histone H1.2 - Homo sapiens (Human)					
P16403	3	4.3798	K.AKKPAAATVTK.K	6	H12_HUMAN
P16403	2	2.8358	K.KPAAATVTK.K	3	H12_HUMAN
Histone H1.3 - Homo sapiens (Human)					
P16402	2	2.9862	K.KVAGAATPK.K	3	H13_HUMAN
P16402	2	2.7137	K.VAGAATPK.K	2	H13_HUMAN
Histone H1.5 - Homo sapiens (Human)					
P16401	2	3.3075	K.ATGPPVSELITK.A	2	H15_HUMAN
P16401	2	3.3592	R.NGLSLAALK.K	2	H15_HUMAN
P16401	3	4.595	R.KATGPPVSELITK.A	9	H15_HUMAN
P16401	3	4.2722	K.RKATGPPVSELITK.A	6	H15_HUMAN
P16401	2	4.62	K.KALAAGGYDVEK.N	6	H15_HUMAN
P16401	1	3.6408	K.ALAAGGYDVEK.N	11	H15_HUMAN
P16401	3	4.7594	K.KALAAGGYDVEKNNSR.I	3	H15_HUMAN
Histone H1x - Homo sapiens (Human)					
Q92522	2	4.8653	K.YSQLVVETIR.R	7	H1X_HUMAN
Q92522	2	3.6653	R.RGAPAAATAPAPTAHK.A	2	H1X_HUMAN
Q92522	2	3.9342	R.GAPAAATAPAPTAHK.A	1	H1X_HUMAN
Q92522	2	2.7465	K.KVPWFDQQNGR.T	1	H1X_HUMAN
Q92522	2	3.0492	K.YSQLVVETIRR.L	2	H1X_HUMAN
Q92522	1	2.1758	K.GTGANGSFK.L	1	H1X_HUMAN
Q92522	2	4.7029	K.ALVQNDTLLQVK.G	9	H1X_HUMAN
Q92522	2	3.9303	K.AGGSAALSPSKK.R	6	H1X_HUMAN
Q92522	1	4.2305	K.AGGSAALSPSK.K	7	H1X_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q92522	2	2.8832	K.KAAKPSVPK.V	1	H1X_HUMAN
Histone H2A - Homo sapiens (Human)					
Q92646	2	3.1733	K.ILELAGNAAR.D	1	Q92646_HUMAN
Histone H2A.x - Homo sapiens (Human)					
P16104	1	2.9796	K.ATQASQEY.-	4	H2AX_HUMAN
P16104	1	3.1673	K.KATQASQEY.-	4	H2AX_HUMAN
Histone H4 - Homo sapiens (Human)					
P62805	2	3.1484	K.RISGLIYEETR.G	2	H4_HUMAN
P62805	3	5.371	R.KTVTAMDVVYALKR.Q	2	H4_HUMAN
P62805	2	4.1873	R.ISGLIYEETR.G	8	H4_HUMAN
P62805	2	3.7515	R.DNIQGITKPAIR.R	4	H4_HUMAN
P62805	2	3.6568	R.DAVTYTEHAK.R	8	H4_HUMAN
P62805	3	5.0156	K.VFLENVIRDAVTYTEHAK.R	6	H4_HUMAN
P62805	2	3.3882	K.VFLENVIR.D	2	H4_HUMAN
P62805	2	4.1951	K.TVTAMDVVYALK.R	3	H4_HUMAN
P62805	2	3.799	K.TVTAMDVVYALKR.Q	3	H4_HUMAN
Histone RNA hairpin-binding protein - Homo sapiens (Human)					
Q14493	3	4.8813	R.RPESFTTPEGPKPR.S	2	SLBP_HUMAN
Histone-binding protein RBBP4 - Homo sapiens (Human)					
Q09028	3	4.7123	K.IGEEQSPEDAEDGPPELLFIHGGHTAK.I	1	RBBP4_HUMAN
Histone-lysine N-methyltransferase NSD3 - Homo sapiens (Human)					
Q9BZ95	3	4.9397	R.NEKPTQSVSSPEATSGSTGSVEK.K	1	NSD3_HUMAN
Histone-lysine N-methyltransferase, H3 lysine-4 specific SET7 - Homo sapiens (Human)					
Q8WTS6	2	2.9451	K.IAYVYPDER.T	3	SETD7_HUMAN
Q8WTS6	4	5.1364	K.LATLMSTEEGRPHFELMPGNSVYHFDK.S	2	SETD7_HUMAN
Q8WTS6	2	4.3163	K.VAVGPNTVMSFYNGVR.I	3	SETD7_HUMAN
HIV Tat-specific factor 1 - Homo sapiens (Human)					
O43719	2	4.7033	R.LRGWEAFLNAPEANR.G	2	HTSF1_HUMAN
O43719	2	3.4785	R.RSDSVSASER.A	3	HTSF1_HUMAN
O43719	2	3.6178	R.HFSEHPSTSK.M	3	HTSF1_HUMAN
O43719	2	3.138	R.GWEAFLNAPEANR.G	1	HTSF1_HUMAN
O43719	2	4.4182	K.TEDGGEFEEGASENNAK.E	2	HTSF1_HUMAN
O43719	2	3.8849	K.MNAQETATGMAFEEPIDEK.K	2	HTSF1_HUMAN
O43719	2	3.6553	K.M#NAQETATGMAFEEPIDEK.K	1	HTSF1_HUMAN
HLA class II DR-beta - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q29890	2	3.1039	R.RVYPEVTVYPAK.T	2	Q29890_HUMAN
HMGA1 protein - Homo sapiens (Human)					
Q6IPL9	2	2.909	R.KQPPKEPSEVPTPK.R	1	Q6IPL9_HUMAN
Homeobox protein cut-like 1 - Homo sapiens (Human)					
P39880	2	2.9909	K.TAEPAQPSSASGSGNSDDAIR.S	2	CUTL1_HUMAN
Homeobox protein Hox-A5 - Homo sapiens (Human)					
P20719	2	3.0343	R.SYAASASAAPAEPR.Y	1	HXA5_HUMAN
Homeodomain-only protein - Homo sapiens (Human)					
Q9BPY8	2	2.9589	R.RSEGLPSECR.S	1	HOP_HUMAN
Hook homolog 2 - Homo sapiens (Human)					
Q96ED9	2	3.3341	R.LNQQQLSELR.A	1	HOOK2_HUMAN
Hook homolog 3 - Homo sapiens (Human)					
Q86VS8	3	3.764	K.LNQEGSDNEKIALLQSLLDDANLR.K	1	HOOK3_HUMAN
Q86VS8	2	3.5516	R.LASTGSGQSFLAR.Q	2	HOOK3_HUMAN
Hornerin - Homo sapiens (Human)					
Q86YZ3	3	5.8334	R.GEQHGSSSGSSSSYGQHGSGSR.Q	1	HORN_HUMAN
Q86YZ3	2	4.0504	R.YGQQGSGSGQSPSR.G	2	HORN_HUMAN
Q86YZ3	3	3.9524	R.QGSSAGSSSSYGQHGSGSR.Q	2	HORN_HUMAN
Host cell factor - Homo sapiens (Human)					
P51610	2	2.7832	K.GAPGQPGTILR.T	1	HCFC1_HUMAN
P51610	2	4.6486	K.IATGHGQQGVTQVVLK.G	3	HCFC1_HUMAN
P51610	2	3.5418	K.SPISVPGGSALISNLGK.V	3	HCFC1_HUMAN
P51610	2	3.4078	K.VASSPVM#VSNPATR.M	1	HCFC1_HUMAN
P51610	2	3.7238	K.VASSPVMVSNPATR.M	2	HCFC1_HUMAN
P51610	2	2.7484	K.VM#TSGTGAPAK.I	1	HCFC1_HUMAN
P51610	2	3.7644	R.SPAFVQLAPLSSK.V	2	HCFC1_HUMAN
Hsp90 co-chaperone Cdc37 - Homo sapiens (Human)					
Q16543	3	4.1482	K.ASEAKEGEEAGPGDPLLEAVPK.T	1	CDC37_HUMAN
Q16543	2	3.4314	R.SWEQKLEEMR.K	1	CDC37_HUMAN
Q16543	2	3.1892	R.LQAEAQQLR.K	3	CDC37_HUMAN
Q16543	3	3.8914	K.LKELEVAEGGKAELER.L	1	CDC37_HUMAN
Q16543	2	2.9384	K.TGDEKDVSV.-	1	CDC37_HUMAN
HSPA5 protein - Homo sapiens (Human)					
Q2KHP4	2	4.7509	K.VTHAVVTVPAYFNDAQR.Q	4	Q2KHP4_HUMA
Q2KHP4	3	3.7427	K.SQIFSTASDNQPTVTIK.V	2	Q2KHP4_HUMA

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q2KHP4	2	4.051	K.TFAPEEISAM#VLTK.M	4	Q2KHP4_HUMA
Q2KHP4	2	2.7528	R.NELESYAYSLK.N	1	Q2KHP4_HUMA
Q2KHP4	2	4.6874	K.TKPYIQVDIGGGQTK.T	7	Q2KHP4_HUMA
Q2KHP4	2	3.6236	K.SDIDEIVLVGGSTR.I	3	Q2KHP4_HUMA
Q2KHP4	3	5.8041	K.VLEDSDLKKSDIDEIVLVGGSTR.I	4	Q2KHP4_HUMA
Q2KHP4	2	3.2637	K.VYEGERPLTK.D	3	Q2KHP4_HUMA
Q2KHP4	2	3.6287	R.AKFEELNM#DLFR.S	1	Q2KHP4_HUMA
Q2KHP4	3	5.2128	R.AKFEELNMDLFR.S	7	Q2KHP4_HUMA
Q2KHP4	3	4.1768	R.IDTRNELESYAYSLK.N	7	Q2KHP4_HUMA
Q2KHP4	2	3.8384	R.IINEPTAAAIAYGLDKR.E	4	Q2KHP4_HUMA
Q2KHP4	2	3.2242	K.RALSSQHQAR.I	5	Q2KHP4_HUMA
Q2KHP4	2	4.6956	R.MVNDAEKFAEEDKK.L	3	Q2KHP4_HUMA
Q2KHP4	3	5.0467	K.TFAPEEISAMVLTK.M	10	Q2KHP4_HUMA
Q2KHP4	2	3.7612	R.TWNDPSVQQDIK.F	5	Q2KHP4_HUMA
Q2KHP4	2	4.4978	R.ITPSYVAFTPEGER.L	4	Q2KHP4_HUMA
Q2KHP4	2	4.8481	K.KSDIDEIVLVGGSTR.I	2	Q2KHP4_HUMA
Q2KHP4	2	2.8796	K.VLEDSDLKK.S	1	Q2KHP4_HUMA
Q2KHP4	3	5.5892	K.AVEEKIEWLESHQDADIEDFKAK.K	2	Q2KHP4_HUMA
Q2KHP4	3	4.5944	K.DAGTIAGLNVM#R.I	7	Q2KHP4_HUMA
Q2KHP4	2	4.6581	K.DAGTIAGLNVMR.I	3	Q2KHP4_HUMA
Q2KHP4	2	2.752	K.EFFNGKEPSR.G	1	Q2KHP4_HUMA
Q2KHP4	2	3.847	K.FEELNMDLFR.S	3	Q2KHP4_HUMA
Q2KHP4	3	4.8924	K.IEWLESHQDADIEDFKAK.K	1	Q2KHP4_HUMA
Q2KHP4	2	3.9989	K.NQLTSNPENTVFDAKR.L	3	Q2KHP4_HUMA
Q2KHP4	2	5.4213	K.KKELEEIVQPIISK.L	5	Q2KHP4_HUMA
Q2KHP4	2	4.5095	K.KSQIFSTASDNQPTVTIK.V	2	Q2KHP4_HUMA
Q2KHP4	3	4.3519	K.KTKPYIQVDIGGGQTK.T	1	Q2KHP4_HUMA
Q2KHP4	2	4.1022	K.KVTHAVVTVPAYFNDAQR.Q	3	Q2KHP4_HUMA
Q2KHP4	2	3.9091	K.MKETAEAYLGK.K	2	Q2KHP4_HUMA
Q2KHP4	3	4.5696	K.MKETAEAYLGKK.V	3	Q2KHP4_HUMA
Q2KHP4	3	5.0312	K.NGRVEIIANDQGNR.I	5	Q2KHP4_HUMA
Q2KHP4	2	3.0497	K.NKITITNDQNR.L	1	Q2KHP4_HUMA
Q2KHP4	2	4.5825	K.NQLTSNPENTVFDAK.R	1	Q2KHP4_HUMA
Q2KHP4	2	3.0065	K.ITITNDQNR.L	4	Q2KHP4_HUMA
HSRBC protein - Homo sapiens (Human)					
Q969G5	2	3.0662	R.RQGGLAGSVR.R	2	Q969G5_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q969G5	2	4.9742	R.SHDTTSNTLAQLLAK.A	2	Q969G5_HUMA
Q969G5	3	4.942	R.GKLHVLLFKEEGEVPASAFQK.A	1	Q969G5_HUMA
Q969G5	2	4.3877	R.KGPAAPPPTPVKPPR.L	5	Q969G5_HUMA
Huntingtin-interacting protein HYPK - Homo sapiens (Human)					
Q9NX55	3	4.1985	K.VTIKKEDLELIMTEMEISR.A	2	HYPK_HUMAN
Q9NX55	2	4.0533	R.EHMGNVVEALIALTN.-	4	HYPK_HUMAN
Q9NX55	2	4.5114	R.KHDSGAADLER.V	6	HYPK_HUMAN
Q9NX55	3	5.3623	R.VTDYAEEKEIQSSNLETAMSVIGDRR.S	1	HYPK_HUMAN
Hypothetical gene supported by AK027125 - Homo sapiens (Human)					
A4D104	3	4.7658	K.TSDIEANQPLETNKENSSSVTVSDPEMENK.A	2	A4D104_HUMAN
Hypothetical LOC284297 - Homo sapiens (Human)					
A1L4H1	2	4.2092	R.QALLLGLTQLVEAAR.G	2	A1L4H1_HUMAN
Hypothetical LOC51149 - Homo sapiens (Human)					
Q6NTE8	2	5.1583	K.LNLLQGQVSELPLR.S	4	Q6NTE8_HUMA
Hypoxia up-regulated protein 1 precursor - Homo sapiens (Human)					
Q9Y4L1	2	2.7874	K.AVGKEELGK.N	1	HYOU1_HUMAN
Q9Y4L1	2	3.4714	K.NINADEAAAMGAVYQAAALSK.A	1	HYOU1_HUMAN
Q9Y4L1	2	4.2723	K.AEAGPEGVAPAPEGEKK.Q	2	HYOU1_HUMAN
Q9Y4L1	2	3.147	R.LIPEMDQIFTEVEM#TTLEK.V	1	HYOU1_HUMAN
ICOS ligand precursor - Homo sapiens (Human)					
O75144	2	3.1463	R.GLYDVVSVLR.I	2	ICOSL_HUMAN
Ig kappa chain V-I region Mev - Homo sapiens (Human)					
P01612	2	4.1958	-.DVQM#TQSPSSLSASVGDR.V	2	KV120_HUMAN
Ig kappa chain V-I region Wes - Homo sapiens (Human)					
P01611	2	4.0595	-.DIQM#TQSPSSVSASVGDR.V	2	KV119_HUMAN
P01611	2	3.381	-.DIQMTQSPSSVSASVGDR.V	1	KV119_HUMAN
Ig lambda chain V-III region SH - Homo sapiens (Human)					
P01714	2	4.6204	-.SELTQDPAVSVALGQTVR.I	2	LV301_HUMAN
IGKC protein - Homo sapiens (Human)					
Q569I9	2	3.1982	R.GTVAAPSVFIFPPSDEQLK.S	2	Q569I9_HUMAN
Immunoglobulin J chain - Homo sapiens (Human)					
P01591	3	4.5046	K.KCDPTEVELDNQIVTATQSNICDEDSATETCY	2	IGJ_HUMAN
P01591	2	3.5857	R.FVYHLSDLCK.K	5	IGJ_HUMAN
P01591	2	3.011	R.SSEDPNEDIVER.N	2	IGJ_HUMAN
Immunoglobulin-binding protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P78318	3	5.9641	K.AAQQQEEQEEKEEEDDEQTLHR.A	2	IGBP1_HUMAN
P78318	2	3.2193	K.YGALPDQGIAK.A	1	IGBP1_HUMAN
Import inner membrane translocase subunit TIM44, mitochondrial precursor - Homo sapiens (Hu					
O43615	3	4.1951	R.KGFLSGLLDNVKQELAK.N	1	TIM44_HUMAN
Inactive phospholipase C-like protein 2 - Homo sapiens (Human)					
Q9UPR0	1	2.204	K.PGGLPR.R	1	PLCL2_HUMAN
Q9UPR0	2	4.1347	R.GGAAGGALPTSPGPALGAK.G	1	PLCL2_HUMAN
InaD-like protein - Homo sapiens (Human)					
Q8NI35	2	2.9452	R.KTSSSTSPLEPPSDR.G	1	INADL_HUMAN
Inner nuclear membrane protein Man1 - Homo sapiens (Human)					
Q9Y2U8	2	3.592	R.ISASGPESLLGGPGGASAAPAAGSK.V	1	MAN1_HUMAN
Q9Y2U8	2	3.7098	K.KLREEEQQQHR.S	2	MAN1_HUMAN
Q9Y2U8	3	3.9581	K.LREEEQQQHR.S	4	MAN1_HUMAN
Q9Y2U8	2	2.9188	K.VLLGFSSDESDVEASPR.D	1	MAN1_HUMAN
Q9Y2U8	2	4.6065	R.ENYSDSEEEDDDDVASSR.Q	4	MAN1_HUMAN
Inositol polyphosphate phosphatase-like protein 1 - Homo sapiens (Human)					
O15357	2	4.7101	K.TLSEVDYAPAGPAR.S	4	O15357_HUMAN
O15357	2	3.2923	R.SALLPGPLELQPPR.G	1	O15357_HUMAN
Inositol-trisphosphate 3-kinase B - Homo sapiens (Human)					
P27987	2	3.9741	K.SGGGPGPSGSETPPPPR.R	2	IP3KB_HUMAN
P27987	2	3.1585	R.GASFLFPPAESLSPEEPR.S	1	IP3KB_HUMAN
Insulin-like growth factor-binding protein 4 precursor - Homo sapiens (Human)					
P22692	3	4.0047	K.TGVKLPGGLEPK.G	1	IBP4_HUMAN
Insulin-like growth factor-binding protein 5 precursor - Homo sapiens (Human)					
P24593	2	2.9024	R.HM#EASLQELK.A	1	IBP5_HUMAN
P24593	2	2.8258	R.HMEASLQELK.A	2	IBP5_HUMAN
Insulin-like growth factor-binding protein 7 precursor - Homo sapiens (Human)					
Q16270	2	4.1361	K.ITVVDALHEIPVK.K	2	IBP7_HUMAN
Q16270	2	3.8821	R.TELLPGDRDNLAIQTR.G	4	IBP7_HUMAN
Q16270	3	4.7447	R.GGPEKHEVTGWVLVSPLSK.E	2	IBP7_HUMAN
Q16270	2	3.8595	R.GEGEPCGGGGAGR.G	13	IBP7_HUMAN
Q16270	3	4.2232	K.EDAGEYECHASNSQGQASASAK.I	3	IBP7_HUMAN
Q16270	2	4.848	K.AGAAAGGPGVSGVCVCK.S	2	IBP7_HUMAN
Q16270	3	3.985	K.ITVVDALHEIPVKKGEGAEL.-	1	IBP7_HUMAN
Integrator complex subunit 12 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96CB8	3	3.8189	K.TTSESSSSPSASLKGPTSQESQLNAMK.R	1	INT12_HUMAN
Integrin alpha-1 precursor - Homo sapiens (Human)					
P56199	2	3.3858	K.VIQDCEDENIQR.F	2	ITA1_HUMAN
Integrin alpha-5 precursor - Homo sapiens (Human)					
P08648	2	4.4779	R.LLESSLSSSEGEEPVEYK.S	2	ITA5_HUMAN
Integrin alpha-7 precursor [Contains: Integrin alpha-7 heavy chain; Integrin alpha-7 light chain] -					
Q13683	4	4.784	R.REGPDAHPILAADGHPELGPDGHPGPGTA.-	1	ITA7_HUMAN
Integrin alpha-V precursor - Homo sapiens (Human)					
P06756	2	3.8383	R.STGLNAVPSQILEGQWAAR.S	1	ITAV_HUMAN
Integrin beta-4 precursor - Homo sapiens (Human)					
P16144	2	3.7564	R.LVFSALGPTSLR.V	3	ITB4_HUMAN
Integrin, beta 1 - Homo sapiens (Human)					
Q8WUM6	3	4.27	K.LKPEDITQIQPQQLVLR.L	4	Q8WUM6_HUMA
Inter-alpha-trypsin inhibitor heavy chain H1 precursor - Homo sapiens (Human)					
P19827	2	3.6707	K.ILGDM#QPGDYFDLVLFGTR.V	1	ITIH1_HUMAN
P19827	2	3.0038	R.EVAFDLEIPK.T	2	ITIH1_HUMAN
P19827	3	4.2941	R.FAHYVVTSQVVNTANEAR.E	2	ITIH1_HUMAN
P19827	2	4.1997	K.GSLVQASEANLQAAQDFVR.G	2	ITIH1_HUMAN
Intercellular adhesion molecule 1 precursor - Homo sapiens (Human)					
P05362	2	3.1367	K.LLGIETPLPK.K	2	ICAM1_HUMAN
P05362	2	2.7674	K.LLGIETPLPKK.E	1	ICAM1_HUMAN
P05362	2	4.1933	K.REPAVGEPAEVTTTVLVR.R	1	ICAM1_HUMAN
P05362	2	3.5743	K.VTLNGVPAQPLGPR.A	2	ICAM1_HUMAN
Intercellular adhesion molecule 2 precursor - Homo sapiens (Human)					
P13598	2	4.0347	R.QVILTLQPTLVAVGK.S	3	ICAM2_HUMAN
Interferon regulatory factor 2-binding protein 1 - Homo sapiens (Human)					
Q53EL7	2	2.8724	K.DPGGGGGPVR.A	2	Q53EL7_HUMAN
Interferon-induced protein with tetratricopeptide repeats 3 - Homo sapiens (Human)					
O14879	4	4.7253	K.STDKEEIKDQPQNVSENLLPQNAPNYWYLQG	1	IFIT3_HUMAN
Interleukin enhancer-binding factor 3 - Homo sapiens (Human)					
Q12906	3	5.2568	K.AYAALAALEKLFPDTPLALDANKK.K	1	ILF3_HUMAN
Q12906	1	2.5145	K.FNYSGSGGR.S	2	ILF3_HUMAN
Q12906	2	3.8569	K.VLQDM#GLPTGAEGR.D	1	ILF3_HUMAN
Q12906	2	4.3427	K.VLQDMGLPTGAEGR.D	1	ILF3_HUMAN
Q12906	2	3.3074	R.LNQLKPGLQYK.L	1	ILF3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Intracellular hyaluronan-binding protein 4 - Homo sapiens (Human)					
Q5JVS0	2	3.1647	R.RAENYGPR.A	2	HABP4_HUMAN
Q5JVS0	4	4.8139	K.YRDDMVKDDYEDDSHVFR.K	1	HABP4_HUMAN
Q5JVS0	3	4.231	R.FHQLLDDESDPFDILR.E	3	HABP4_HUMAN
Q5JVS0	2	2.9116	R.KPESTVPSK.A	4	HABP4_HUMAN
Iron-responsive element-binding protein 2 - Homo sapiens (Human)					
P48200	2	2.7941	R.RGNDAVMTRGTFANIK.L	1	IREB2_HUMAN
Iron-sulfur cluster assembly 2 homolog, mitochondrial precursor - Homo sapiens (Human)					
Q86U28	2	3.7685	R.LQVEGGGCSGFQYK.F	2	ISCA2_HUMAN
Q86U28	2	3.8929	R.VVVDSDSLAFVK.G	2	ISCA2_HUMAN
Q86U28	2	4.0593	R.LLEITEGSEFLR.L	2	ISCA2_HUMAN
Isocitrate dehydrogenase [NADP] cytoplasmic - Homo sapiens (Human)					
O75874	2	2.8422	R.DATNDQVTK.D	1	IDHC_HUMAN
IWS1 homolog - Homo sapiens (Human)					
Q96ST2	2	3.1928	R.DLEQMPQR.R	1	IWS1_HUMAN
Q96ST2	2	3.4505	R.RMNSTGGQTPR.R	2	IWS1_HUMAN
Q96ST2	2	3.649	R.PIFGLTSNYK.G	1	IWS1_HUMAN
Q96ST2	3	4.99	R.NRDGGTFISDADDVVSAMIVK.M	1	IWS1_HUMAN
Q96ST2	2	3.1925	K.MNEAAEEDR.Q	2	IWS1_HUMAN
Q96ST2	2	2.8825	R.MNSTGGQTPR.R	2	IWS1_HUMAN
Q96ST2	2	2.7228	R.VSDSESEGPQK.G	1	IWS1_HUMAN
Q96ST2	2	3.3151	R.DGGTFISDADDVVSAMIVK.M	1	IWS1_HUMAN
Q96ST2	2	4.4915	K.MNEAAEEDRQLNNQK.K	1	IWS1_HUMAN
Q96ST2	3	4.5297	K.LINEWSRPIFGLTSNYK.G	2	IWS1_HUMAN
Q96ST2	3	4.8431	K.KLTLLPAVVMHLK.K	1	IWS1_HUMAN
Q96ST2	2	3.0614	K.GPASDSETEDASR.H	1	IWS1_HUMAN
Q96ST2	2	3.4514	K.WNVEMESSR.F	2	IWS1_HUMAN
Q96ST2	2	3.0712	R.DLEQM#PQR.R	2	IWS1_HUMAN
JmjC domain-containing histone demethylation protein 2B - Homo sapiens (Human)					
Q7LBC6	2	3.3638	K.GGNASGEPGLDQR.A	2	JHD2B_HUMAN
JmjC domain-containing protein 3 - Homo sapiens (Human)					
O15054	2	3.1735	R.SATDPADPVDTAEPADSGTER.L	1	JMJD3_HUMAN
JUB protein - Homo sapiens (Human)					
Q96IF1	2	3.3118	R.YEGSFPAGPPPTR.A	1	Q96IF1_HUMAN
Q96IF1	2	2.8986	R.SGSDGTPGPGK.G	4	Q96IF1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Junctophilin-2 - Homo sapiens (Human)					
Q9BR39	2	3.8617	K.DGLLSPGAWNGEPSGEGSR.S	1	JPH2_HUMAN
Kelch repeat and BTB domain-containing protein 11 - Homo sapiens (Human)					
O94819	2	3.7933	R.QWEAGSAGAASPEELASPEER.A	1	KBTBB_HUMAN
Keratin, type I cytoskeletal 10 - Homo sapiens (Human)					
P13645	2	5.0648	R.GSSGGGCFGGSSGGYGGLGGFGGGSFR.G	4	K1C10_HUMAN
P13645	2	3.5373	R.SLLEGEGSSGGGGR.G	1	K1C10_HUMAN
P13645	2	3.7306	R.SQYEQLAEQNRK.D	2	K1C10_HUMAN
P13645	3	3.7505	R.NVSTGDVNVEM#NAAPGVDLTQLLNNMR.S	1	K1C10_HUMAN
P13645	3	4.8778	K.TIDDLKNQILNLTTDNANILLQIDNAR.L	1	K1C10_HUMAN
P13645	2	5.7683	K.GSLGGGFSSGGFSGGSFSR.G	7	K1C10_HUMAN
P13645	3	4.6518	K.SKELTTEIDNNIEQISSYK.S	1	K1C10_HUMAN
P13645	2	3.8871	R.ALEESNYELEGK.I	3	K1C10_HUMAN
Keratin, type I cytoskeletal 14 - Homo sapiens (Human)					
P02533	2	4.6076	R.APSTYGGGLSVSSSR.F	1	K1C14_HUMAN
Keratin, type I cytoskeletal 16 - Homo sapiens (Human)					
P08779	2	3.9504	R.APSTYGGGLSVSSR.F	2	K1C16_HUMAN
Keratin, type I cytoskeletal 18 - Homo sapiens (Human)					
P05783	2	4.0322	R.IVDGKVVSETNDTK.V	1	K1C18_HUMAN
P05783	3	5.6324	R.SLGSVQAPSYGARPVSSAASVYAGAGGSGS	19	K1C18_HUMAN
P05783	2	4.0018	R.SLGSVQAPSYGAR.P	11	K1C18_HUMAN
P05783	3	4.6501	R.RLLEDGEDFNLGDALDSSNSMQTIQK.T	1	K1C18_HUMAN
P05783	1	2.1603	R.LASYLDR.V	1	K1C18_HUMAN
P05783	3	4.5797	R.YALQMEQLNGILLHLESELAQTR.A	1	K1C18_HUMAN
P05783	3	4.2668	R.GGMGSGGLATGIAGGLAGMGGIQNEKETMQ	1	K1C18_HUMAN
P05783	2	4.5772	R.AQIFANTVDNAR.I	1	K1C18_HUMAN
P05783	1	2.909	K.VVSETNDTK.V	10	K1C18_HUMAN
P05783	2	6.6003	R.PVSSAASVYAGAGGSGSR.I	28	K1C18_HUMAN
Keratin, type I cytoskeletal 19 - Homo sapiens (Human)					
P08727	3	4.0874	R.FGAQLAHIQALISGIEAQLGDVR.A	1	K1C19_HUMAN
P08727	2	4.0456	R.SLLEGQEDHYNNLSASK.V	1	K1C19_HUMAN
P08727	2	4.2492	R.QSSATSSFGGLGGGSVR.F	2	K1C19_HUMAN
P08727	2	3.3462	R.GQVGGQVSVEVDSAPGTDLAK.I	1	K1C19_HUMAN
P08727	2	4.2628	K.SRLEQEIATYR.S	2	K1C19_HUMAN
P08727	2	2.7963	K.NHEEEISTLR.G	1	K1C19_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P08727	2	4.1544	K.AALEDTLAETEAR.F	1	K1C19_HUMAN
P08727	2	2.9087	K.DAEAWFTSR.T	2	K1C19_HUMAN
Keratin, type I cytoskeletal 9 - Homo sapiens (Human)					
P35527	2	5.3968	R.GGSGGSYGGGGSGGGYGGGSGSR.G	2	K1C9_HUMAN
P35527	2	2.7546	K.KGPAAIQK.N	1	K1C9_HUMAN
P35527	2	5.328	R.SGGGGGGGLGSGGSIR.S	6	K1C9_HUMAN
P35527	2	3.2146	R.QGVDADINGLR.Q	2	K1C9_HUMAN
P35527	3	5.2681	R.LASYLDKVQALEEANNDLENK.I	2	K1C9_HUMAN
P35527	3	3.9508	R.IKFEMEQNLR.Q	2	K1C9_HUMAN
P35527	3	6.6454	R.HGVQELEIELQSQLSKK.A	2	K1C9_HUMAN
P35527	3	4.093	R.HGVQELEIELQSQLSK.K	1	K1C9_HUMAN
P35527	3	6.315	K.EIETYHNLLEGGQEDFESSGAGK.I	1	K1C9_HUMAN
P35527	2	4.8374	R.GGGGSFGYSYGGGSGGGFSASSLGGGFGG	2	K1C9_HUMAN
P35527	2	4.3039	R.FSSSSGYGGGSSR.V	4	K1C9_HUMAN
P35527	2	3.0632	K.TLLDIDNTR.M	3	K1C9_HUMAN
P35527	1	3.3394	K.STMQELNSR.L	6	K1C9_HUMAN
P35527	3	4.6152	K.NYSPYYNTIDDLKDQIVDLTVGNNK.T	1	K1C9_HUMAN
P35527	3	7.5795	R.GGSGGSHGGGSGFGGESGGSYGGGEEAS	1	K1C9_HUMAN
P35527	2	3.2784	K.STM#QELNSR.L	5	K1C9_HUMAN
Keratin, type II cytoskeletal 1 - Homo sapiens (Human)					
P04264	3	5.1743	R.GSYGSGGSSYGSGGGSYGSGGGGGGHGS	1	K2C1_HUMAN
P04264	2	3.8736	R.LLRDYQELMNTK.L	3	K2C1_HUMAN
P04264	3	3.7338	R.LRSEIDNVKK.Q	1	K2C1_HUMAN
P04264	2	4.4706	R.SGGGFSSGSAGIINYQR.R	2	K2C1_HUMAN
P04264	2	5.0739	R.SLDLDSIIAEVK.A	4	K2C1_HUMAN
P04264	1	2.8828	R.SLVNLGGSK.S	5	K2C1_HUMAN
P04264	2	3.983	R.TNAENEFVTIKK.D	3	K2C1_HUMAN
P04264	2	2.8933	R.TLLEGEESR.M	1	K2C1_HUMAN
P04264	2	4.086	R.TNAENEFVTIK.K	3	K2C1_HUMAN
P04264	2	2.9417	K.DVDGAYMTK.V	3	K2C1_HUMAN
P04264	3	5.2217	R.GGSGGGGGGSSGGRGSGGGSSGGSIGGR.	2	K2C1_HUMAN
P04264	3	4.1591	R.THNLEPYFESFINNLR.R	3	K2C1_HUMAN
P04264	2	4.6423	R.FSSCGGGGGSFGAGGGFGSR.S	3	K2C1_HUMAN
P04264	3	5.4235	R.FLEQQNQVLQTKWELLQQVDTSTR.T	3	K2C1_HUMAN
P04264	2	3.8803	K.YEELQITAGR.H	3	K2C1_HUMAN
P04264	2	4.1552	K.SLNNQFASFIDKVR.F	3	K2C1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P04264	2	4.547	K.SLNNQFASFIDK.V	5	K2C1_HUMAN
P04264	2	4.8502	K.SKAEAESLYQSK.Y	2	K2C1_HUMAN
P04264	3	3.893	K.LNDLEDALQQAKEDLAR.L	1	K2C1_HUMAN
P04264	2	2.8476	K.DVDGAYM#TK.V	4	K2C1_HUMAN
P04264	2	2.9705	K.AEAESLYQSK.Y	2	K2C1_HUMAN
P04264	3	4.2476	K.NKLNDLEDALQQAKEDLAR.L	1	K2C1_HUMAN
P04264	2	5.0812	R.GGGGGGYGSGGSSYGSGGGSYGSGGGGG	5	K2C1_HUMAN
Keratin, type II cytoskeletal 2 epidermal - Homo sapiens (Human)					
P35908	2	3.9665	R.GSSSGGGYSSGSSSYGSGGR.Q	1	K22E_HUMAN
P35908	2	4.8142	R.HGGGGGGFGGGGFGSR.S	1	K22E_HUMAN
P35908	2	5.3581	R.GGSSSGGGYGSGGGGSSSVK.G	2	K22E_HUMAN
P35908	2	4.0381	R.GFSSGSAVVSGGSR.R	2	K22E_HUMAN
P35908	2	3.7802	K.VDLLNQEIEFLK.V	3	K22E_HUMAN
P35908	2	5.1291	K.GGSISGGGYGSGGGK.H	4	K22E_HUMAN
P35908	2	5.6525	R.GGSGGGGSISGGGYGSGGGSGGR.Y	2	K22E_HUMAN
P35908	3	4.2524	R.NKLNDLEEALQQAKEDLAR.L	1	K22E_HUMAN
Keratin, type II cytoskeletal 5 - Homo sapiens (Human)					
P13647	2	3.1823	K.AQYEEIANR.S	1	K2C5_HUMAN
P13647	2	2.8695	R.ISISTSGGSFR.N	1	K2C5_HUMAN
Keratin, type II cytoskeletal 8 - Homo sapiens (Human)					
P05787	3	4.9281	K.LLEGEESRLESGM#QNMSIHTK.T	2	K2C8_HUMAN
P05787	2	3.1742	R.LQAEIEGLK.G	3	K2C8_HUMAN
P05787	2	3.9008	R.LQAEIEGLKGQR.A	3	K2C8_HUMAN
P05787	3	4.4949	R.QLYEEEIRELQSQISDTSVVLSM#DNSR.S	2	K2C8_HUMAN
P05787	2	3.0092	R.SLDM#DSIIAEVK.A	2	K2C8_HUMAN
P05787	1	4.4679	R.SLDMDSIIAEVK.A	4	K2C8_HUMAN
P05787	3	5.0734	R.SLDMDSIIAEVKAQYEDIANR.S	2	K2C8_HUMAN
P05787	2	3.0519	R.SNM#DNMFESYINNLR.R	1	K2C8_HUMAN
P05787	3	4.9135	R.SRAEAESM#YQIKYEELQSLAGK.H	1	K2C8_HUMAN
P05787	3	5.0873	R.SRAEAESMYQIKYEELQSLAGK.H	2	K2C8_HUMAN
P05787	1	2.3737	R.SYTSGPGSR.I	2	K2C8_HUMAN
P05787	2	4.622	R.LEGLTDEINFLR.Q	4	K2C8_HUMAN
P05787	2	3.4523	R.TEMENEFVLIKK.D	3	K2C8_HUMAN
P05787	1	2.7503	K.YEELQSLAGK.H	1	K2C8_HUMAN
P05787	3	3.9241	R.KLLEGEESRLESGMQNMSIHTK.T	1	K2C8_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P05787	2	4.4687	K.DVDEAYMNKVELESR.L	6	K2C8_HUMAN
P05787	2	3.9292	K.LVSESSDVLPK.-	9	K2C8_HUMAN
P05787	3	5.0706	K.LLEGEESRLESGMQNM#SIHTK.T	3	K2C8_HUMAN
P05787	3	4.6203	R.AEAESM#YQIKYEELQSLAGK.H	2	K2C8_HUMAN
P05787	3	4.0574	R.AEAESMYQIKYEELQSLAGK.H	1	K2C8_HUMAN
P05787	2	3.6379	R.ASLEAAIADAEQR.G	2	K2C8_HUMAN
P05787	3	3.9196	R.ASLEAAIADAEQRGELAIK.D	3	K2C8_HUMAN
P05787	3	5.901	R.ASLEAAIADAEQRGELAIKDANAK.L	5	K2C8_HUMAN
P05787	3	4.7786	R.DGKLVSESSDVLPK.-	20	K2C8_HUMAN
P05787	1	2.4252	R.ISSSSFSR.V	2	K2C8_HUMAN
P05787	3	4.0016	K.IETRDGKLVSESSDVLPK.-	1	K2C8_HUMAN
Keratin-71 - Homo sapiens (Human)					
Q3SY84	2	2.776	K.DTLGKGSSLSAPSK.K	1	Q3SY84_HUMA
Keratin-75 - Homo sapiens (Human)					
O95678	2	2.8377	R.NTKQEISEMNRMIQR.L	1	O95678_HUMAN
KH domain-containing, RNA-binding, signal transduction-associated protein 1 - Homo sapiens (H					
Q07666	2	4.3763	R.SGSMDPSGAHPSVR.Q	2	SAM68_HUMAN
Q07666	2	2.8042	K.GAYREHPYGR.Y	1	SAM68_HUMAN
Q07666	2	3.3628	R.SGSM#DPSGAHPSVR.Q	2	SAM68_HUMAN
KIAA0515 - Homo sapiens (Human)					
Q5JSZ9	3	4.2478	R.DLFEERGEEYLSAFDKK.A	1	Q5JSZ9_HUMAN
KIAA1155 protein - Homo sapiens (Human)					
Q9ULR5	3	4.007	R.DLPQAM#GQLQQQLNGLSVSEGHDSEDILSK	1	Q9ULR5_HUMA
KIAA1276 protein - Homo sapiens (Human)					
Q9ULE4	2	2.9925	K.SEKEIKQLEEEK.A	1	Q9ULE4_HUMA
KIAA1430 - Homo sapiens (Human)					
Q8N6E7	2	4.1188	K.STETQPSSTAPK.C	4	Q8N6E7_HUMA
KIAA1843 protein - Homo sapiens (Human)					
Q96JI4	1	2.1168	K.STAQNVEGIIVSAMFK.S	1	Q96JI4_HUMAN
KIDINS220 protein - Homo sapiens (Human)					
A1L4N4	2	4.172	R.TPSTVTLNNNSAPANR.A	2	A1L4N4_HUMAN
Kidney and brain protein - Homo sapiens (Human)					
Q8IX03	2	3.2256	R.RGDSQPYQALK.Y	1	WWC1_HUMAN
Q8IX03	2	3.1708	R.SDSDSSTLSK.K	3	WWC1_HUMAN
Kinectin - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q86UP2	2	5.3459	K.ALKEEIGNVQLEK.A	7	KTN1_HUMAN
Q86UP2	1	3.107	K.AQQLSITSK.V	3	KTN1_HUMAN
Q86UP2	2	2.7146	R.ELKDLLTELQK.K	1	KTN1_HUMAN
Kinesin heavy chain - Homo sapiens (Human)					
P33176	2	3.1268	K.LKTQMLDQEELLASTR.R	1	KINH_HUMAN
P33176	2	3.4703	K.SLTEYLQNVEQK.K	2	KINH_HUMAN
P33176	2	5.1371	R.GGGAFVQNSQPVAVR.G	5	KINH_HUMAN
Kinesin light chain 2 - Homo sapiens (Human)					
Q9H0B6	2	4.3027	K.SVEEPTQPGGTGLSDSR.T	3	KLC2_HUMAN
Q9H0B6	2	2.7035	R.TLSSSSMDLSR.R	1	KLC2_HUMAN
Kinesin-like protein KIF13B - Homo sapiens (Human)					
Q9NQT8	2	4.1642	R.MEEAQPEMGPDVLVQTMGAPALK.I	1	KI13B_HUMAN
Q9NQT8	2	5.314	K.GRWESQQDVSQTTVSR.G	3	KI13B_HUMAN
Q9NQT8	2	3.2372	R.WESQQDVSQTTVSR.G	1	KI13B_HUMAN
Q9NQT8	2	3.4493	R.IM#VQSASPDIR.V	2	KI13B_HUMAN
Q9NQT8	3	4.6353	R.M#EEAQPEMGPDVLVQTMGAPALK.I	1	KI13B_HUMAN
Kinesin-like protein KIF21A - Homo sapiens (Human)					
Q7Z4S6	2	4.599	R.YLLDHFLSMGINK.G	2	KI21A_HUMAN
Q7Z4S6	3	4.7824	K.LSSSDAPAQDTGSSAAAVETDASR.T	3	KI21A_HUMAN
Q7Z4S6	2	3.6224	K.MTISNMEADMNR.L	2	KI21A_HUMAN
Q7Z4S6	3	4.6187	R.KLSSSDAPAQDTGSSAAAVETDASR.T	2	KI21A_HUMAN
Q7Z4S6	3	3.7416	R.LKQTEITSATQNQLLFHMLK.E	1	KI21A_HUMAN
Q7Z4S6	2	2.7401	R.RVTDIIMQK.M	1	KI21A_HUMAN
Kinesin-like protein KIF3C - Homo sapiens (Human)					
O14782	2	2.7365	R.GVIPNAFEHIFTHISR.S	1	KIF3C_HUMAN
Krueppel-like factor 12 - Homo sapiens (Human)					
Q9Y4X4	2	4.3593	R.IPVVVQSVPVVYTAVR.S	3	KLF12_HUMAN
Krueppel-like factor 16 - Homo sapiens (Human)					
Q9BXK1	2	4.6749	R.GGPGAAPGGASPASSSSAASSPSSGR.A	2	KLF16_HUMAN
L antigen family member 3 - Homo sapiens (Human)					
Q14657	2	4.2718	R.DADADAGGGADGGDGR.G	1	LAGE3_HUMAN
Lactoylglutathione lyase - Homo sapiens (Human)					
Q04760	2	5.2338	K.GLAFIQDPDGYWIEILNPNK.M	4	LGUL_HUMAN
Q04760	2	2.7058	K.RFEELGVK.F	1	LGUL_HUMAN
Ladinin 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96GD8	3	3.7121	R.GPWALEEESLVGREPEER.K	1	Q96GD8_HUMA
Lamin-A/C - Homo sapiens (Human)					
P02545	3	6.616	K.ASASGSGAQVGGPISSGSSASSVTVTR.S	8	LMNA_HUMAN
P02545	2	4.9454	R.SVGGSGGGSFGDNLVTR.S	9	LMNA_HUMAN
Lamina-associated polypeptide 2, isoforms beta/gamma - Homo sapiens (Human)					
P42167	2	3.1528	R.AEVGEKTEER.R	5	LAP2B_HUMAN
P42167	3	4.5412	R.RVEHNQSYSQAGITETEWTSGSSK.G	1	LAP2B_HUMAN
P42167	3	4.1357	K.HASPILPITEFSDIPR.R	1	LAP2B_HUMAN
Lamin-B receptor - Homo sapiens (Human)					
Q14739	2	2.7138	K.FADGEVVR.G	1	LBR_HUMAN
Q14739	2	3.3251	R.SASASHQADIK.E	2	LBR_HUMAN
Lamin-B1 - Homo sapiens (Human)					
P20700	2	3.9819	K.NSQGEEVAQR.S	3	LMNB1_HUMAN
P20700	3	7.1877	R.EMAEIRDQMQQQLNDYEQLLDVK.L	2	LMNB1_HUMAN
P20700	3	6.1389	K.SLEGDLEDLKDQIAQLEASLAAAKK.Q	1	LMNB1_HUMAN
P20700	2	3.6768	K.NQNSWGTGEDVK.V	2	LMNB1_HUMAN
P20700	3	5.4397	R.EM#AEIRDQMQQQLNDYEQLLDVK.L	1	LMNB1_HUMAN
Lamin-B2 - Homo sapiens (Human)					
Q03252	2	2.7912	K.SVFEEEVRETR.R	1	LMNB2_HUMAN
Q03252	2	2.8039	R.RVLDETAR.E	1	LMNB2_HUMAN
Q03252	2	3.5227	R.RLVEVDSSR.Q	2	LMNB2_HUMAN
Q03252	2	3.4332	R.AGGPATPLSPTR.L	1	LMNB2_HUMAN
Q03252	2	4.3694	R.ATSSSSGSLSATGR.L	4	LMNB2_HUMAN
Q03252	3	3.83	R.VKDLESLFHR.S	2	LMNB2_HUMAN
Q03252	2	3.2123	K.MAQALEELR.S	3	LMNB2_HUMAN
Q03252	2	4.3346	K.LSSDQNDKAASAAR.E	2	LMNB2_HUMAN
Q03252	2	2.7053	K.LRAELDEVNK.S	1	LMNB2_HUMAN
Q03252	2	3.1607	K.ISEKEEVTTR.E	2	LMNB2_HUMAN
Q03252	2	2.8434	K.GQSSWGTGESFR.T	1	LMNB2_HUMAN
Q03252	2	2.9529	K.ALYESELADAR.R	1	LMNB2_HUMAN
Q03252	3	4.2358	K.MAQALEELRSQHDEQVR.L	1	LMNB2_HUMAN
Q03252	2	5.1062	R.MRLESLSYQLSGLQK.Q	5	LMNB2_HUMAN
Laminin subunit alpha-4 precursor - Homo sapiens (Human)					
Q16363	3	4.481	K.ESM#DTINHASQLVEQAHDMR.D	2	LAMA4_HUMAN
Q16363	2	2.737	R.ELVDEEADEAYELLSQAESWQR.L	1	LAMA4_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q16363	2	3.2856	R.DKIQEINNK.M	4	LAMA4_HUMAN
Q16363	2	4.2445	K.MLYYGEEHELSPK.E	1	LAMA4_HUMAN
Q16363	2	3.6416	K.GKIEQTQASEK.K	3	LAMA4_HUMAN
Q16363	3	4.7118	K.ESMDTINHASQLVEQAHDMR.D	2	LAMA4_HUMAN
Q16363	3	5.0095	R.TLFPVVLEQLDDYNAK.L	2	LAMA4_HUMAN
Q16363	2	3.6426	K.AESSSDEAVADTSR.R	7	LAMA4_HUMAN
Q16363	4	5.746	K.ESMDTINHASQLVEQAHDMRDKIQEINNK.M	1	LAMA4_HUMAN
Laminin subunit alpha-5 precursor - Homo sapiens (Human)					
O15230	4	5.0777	R.HETAQQLEVLEQQSTSLGQDARR.L	1	LAMA5_HUMAN
O15230	2	3.1162	R.RLGGQAVGTR.D	3	LAMA5_HUMAN
O15230	2	3.4789	R.LGLVWAALQGAR.T	2	LAMA5_HUMAN
O15230	4	6.2554	R.LGGQAVGTRDQASQLLAGTEATLGHAK.T	1	LAMA5_HUMAN
O15230	2	3.182	R.GQLQLVEGNFR.H	1	LAMA5_HUMAN
O15230	2	5.8021	R.DLGAPQAAAEAELAAAQR.L	4	LAMA5_HUMAN
O15230	2	4.8271	K.AVAAEAQDTATR.V	4	LAMA5_HUMAN
O15230	2	3.4512	R.AIEASNAYSR.I	1	LAMA5_HUMAN
Laminin subunit beta-1 precursor - Homo sapiens (Human)					
P07942	2	3.1973	R.DRVEDVMMER.E	2	LAMB1_HUMAN
P07942	2	3.0677	R.SLLKDISQK.V	2	LAMB1_HUMAN
P07942	2	3.5851	R.RKAEMLQNEAK.T	1	LAMB1_HUMAN
P07942	2	3.2104	R.RKAEM#LQNEAK.T	2	LAMB1_HUMAN
P07942	3	5.183	R.NFLTQDSADLDSIEAVANEVLK.M	2	LAMB1_HUMAN
P07942	1	2.1085	R.LLDELAGK.L	1	LAMB1_HUMAN
P07942	2	4.0072	R.KAAQNSGEAEYIEK.V	2	LAMB1_HUMAN
P07942	2	3.8134	R.KAEM#LQNEAK.T	4	LAMB1_HUMAN
P07942	2	3.6481	R.YLEDKAQELAR.L	2	LAMB1_HUMAN
P07942	2	2.8789	R.IPSWTGAGFVR.V	2	LAMB1_HUMAN
P07942	3	4.9519	K.AM#DLDQDVLSALAEVEQLSK.M	9	LAMB1_HUMAN
P07942	3	5.9733	K.VTADMVKEALEEAEKAQVAAEK.A	2	LAMB1_HUMAN
P07942	2	3.4537	R.KAEMLQNEAK.T	2	LAMB1_HUMAN
P07942	2	3.5679	K.AAQNSGEAEYIEK.V	1	LAMB1_HUMAN
P07942	3	5.3803	K.AMDLDQDVLSALAEVEQLSK.M	5	LAMB1_HUMAN
P07942	2	5.2829	K.EALEEAEKAQVAAEK.A	3	LAMB1_HUMAN
P07942	2	2.8018	K.ISGVIGPYR.E	1	LAMB1_HUMAN
P07942	2	3.0314	K.KVENLIAK.K	1	LAMB1_HUMAN
P07942	2	2.7333	K.MDKSNEELR.N	1	LAMB1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07942	2	3.3524	K.TLLAQANSK.L	6	LAMB1_HUMAN
P07942	3	4.3273	K.VTADM#VKEALEEAEKAQVAAEK.A	1	LAMB1_HUMAN
Laminin subunit beta-2 precursor - Homo sapiens (Human)					
P55268	2	2.7048	R.LVTPGETPSWTGSGFVR.L	1	LAMB2_HUMAN
P55268	1	2.1911	R.AGYTGLR.C	1	LAMB2_HUMAN
Laminin subunit gamma-1 precursor - Homo sapiens (Human)					
P11047	3	4.6678	R.LQNIRNTIEETGNLAEQAR.A	2	LAMC1_HUMAN
P11047	2	5.0365	R.EAQQALGSAAADATEAK.N	6	LAMC1_HUMAN
P11047	2	3.1825	R.KVSDLENEAK.K	2	LAMC1_HUMAN
P11047	2	3.7877	R.LNTFGDEVFNDPK.V	3	LAMC1_HUMAN
P11047	2	4.3951	R.DTLQEANDILNNLK.D	1	LAMC1_HUMAN
P11047	2	4.3668	R.LSAEDLVLEGAGLR.V	6	LAMC1_HUMAN
P11047	2	4.9328	R.NTIEETGNLAEQAR.A	7	LAMC1_HUMAN
P11047	2	4.3705	R.SAGYLDDVTLASAR.P	3	LAMC1_HUMAN
P11047	2	3.0705	R.DIEEIMKDIR.N	2	LAMC1_HUMAN
P11047	3	5.1179	R.VKLQELESLIANLGTGDEM#VTDQAFEDR.L	2	LAMC1_HUMAN
P11047	3	3.8966	K.KQEAAIMDYNRDIEEIMK.D	1	LAMC1_HUMAN
P11047	3	5.253	R.VKLQELESLIANLGTGDEM#VTDQAFEDRLK.	1	LAMC1_HUMAN
P11047	2	5.0478	R.TFAEVTDLDNEVNNMLK.Q	4	LAMC1_HUMAN
P11047	1	2.9249	K.VSDLENEAK.K	3	LAMC1_HUMAN
P11047	2	4.4597	K.TREAQQALGSAAADATEAK.N	1	LAMC1_HUMAN
P11047	2	3.6877	K.TLPSGCFNTPSIEKP.-	4	LAMC1_HUMAN
P11047	2	3.4037	K.TEQQTADQLLAR.A	2	LAMC1_HUMAN
P11047	2	3.7974	K.SYYYAISDFAVGGR.C	3	LAMC1_HUMAN
P11047	2	2.8114	K.LKDYEDLREDMR.G	1	LAMC1_HUMAN
P11047	3	5.9521	K.GRDTLQEANDILNNLKDFDR.R	2	LAMC1_HUMAN
P11047	2	4.9785	K.GKTEQQTADQLLAR.A	6	LAMC1_HUMAN
P11047	2	4.4728	K.DGFFGNPLAPNPADK.C	2	LAMC1_HUMAN
P11047	2	3.1564	K.ALAEEAAKK.G	3	LAMC1_HUMAN
P11047	1	2.2465	K.ALAEEAAK.K	3	LAMC1_HUMAN
P11047	2	3.3022	K.AFDITYVR.L	2	LAMC1_HUMAN
P11047	2	3.2244	K.MEAENLEQLIDQK.L	2	LAMC1_HUMAN
La-related protein 1 - Homo sapiens (Human)					
Q6PKG0	2	3.6319	K.SVQPQSHKPQPTR.K	2	LARP1_HUMAN
Q6PKG0	3	4.6414	K.VGDFGDAINWPTPGEIAHK.S	5	LARP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6PKG0	2	3.1015	R.SLPTTVPESPNYR.N	1	LARP1_HUMAN
Q6PKG0	2	4.1029	K.WTSQHSNTQTLGK.-	2	LARP1_HUMAN
Q6PKG0	2	3.1523	R.DFVEAPPPK.V	2	LARP1_HUMAN
Q6PKG0	3	4.5863	R.EQFDTLTPEPPVDPNQEVPPGPPR.F	3	LARP1_HUMAN
Q6PKG0	4	5.4088	R.ESPRPLQLPGAEGPAISDGEEGGGEPGAGG	1	LARP1_HUMAN
Q6PKG0	2	2.9208	R.FQQVPTDALANK.L	1	LARP1_HUMAN
Q6PKG0	3	4.7525	R.FQQVPTDALANKLFGAPEPSTIAR.S	1	LARP1_HUMAN
Q6PKG0	3	3.9928	R.HSSNPPLESHVGWVMDSR.E	2	LARP1_HUMAN
Q6PKG0	2	2.7702	R.KFDGVEGPR.T	1	LARP1_HUMAN
Q6PKG0	3	4.1612	K.SDESGEEKNGDEDCQR.G	2	LARP1_HUMAN
Large proline-rich protein BAT3 - Homo sapiens (Human)					
P46379	2	3.1172	R.DLEAPEVQESYR.Q	1	BAT3_HUMAN
Lariat debranching enzyme - Homo sapiens (Human)					
Q9UK59	2	2.8669	R.LSDEHEPEQR.K	2	DBR1_HUMAN
Latent-transforming growth factor beta-binding protein 2 precursor - Homo sapiens (Human)					
Q14767	2	4.3089	K.QSTFTLPLSNQLASVNPSLVK.V	2	LTBP2_HUMAN
Q14767	2	4.3221	R.EQDAPVAGLQPVER.A	4	LTBP2_HUMAN
Q14767	3	4.2657	R.GAGGQSM#SEAPTGDHAPAPTR.M	2	LTBP2_HUMAN
Q14767	2	4.2432	R.GAGGQSMSEAPTGDHAPAPTR.M	2	LTBP2_HUMAN
Q14767	2	3.4465	R.RSSAAGEGTLAR.A	1	LTBP2_HUMAN
Q14767	2	3.714	R.SLGPGTCTLPLAQR.I	1	LTBP2_HUMAN
Q14767	2	4.8938	R.SSEVYAQLCNVAR.I	3	LTBP2_HUMAN
Q14767	2	3.6107	R.STPLGQQQPAPR.T	3	LTBP2_HUMAN
Latent-transforming growth factor beta-binding protein 3 precursor - Homo sapiens (Human)					
Q9NS15	2	4.2405	R.FCQVPAGGAGGGTGGSGPGLSR.T	1	LTBP3_HUMAN
Q9NS15	2	3.3751	R.GAGGGGALAR.E	3	LTBP3_HUMAN
Latent-transforming growth factor beta-binding protein, isoform 1L precursor - Homo sapiens (Hu					
Q14766	2	3.9475	R.SKVPQETQSGGGSR.L	3	LTB1L_HUMAN
Latent-transforming growth factor beta-binding protein, isoform 1S precursor - Homo sapiens (Hu					
P22064	2	3.3509	K.LYQHSQQPGK.A	2	LTB1S_HUMAN
Layilin precursor - Homo sapiens (Human)					
Q6UX15	3	3.7587	R.DGGQLVSIESEDEQKLIEK.F	1	LAYN_HUMAN
Lck-interacting transmembrane adapter 1 - Homo sapiens (Human)					
Q9H400	2	2.7403	R.SPQEPQQGK.T	1	LIME1_HUMAN
Leiomodin 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5VUU2	3	4.0584	K.KGGAPAAPPPPPPPLAPPLIM#ENLK.N	1	Q5VUU2_HUMA
Q5VUU2	3	3.8672	K.KGGAPAAPPPPPPPLAPPLIMENLK.N	1	Q5VUU2_HUMA
LEM domain-containing protein 2 - Homo sapiens (Human)					
Q8NC56	2	3.3121	R.LREDAPLR.A	2	LEMD2_HUMAN
Leucine zipper protein 1 - Homo sapiens (Human)					
Q86V48	2	3.7202	R.VGPSSGDAPEPSSR.R	3	LUZP1_HUMAN
Q86V48	2	3.3624	K.SREQPSVLSR.Y	1	LUZP1_HUMAN
Q86V48	2	3.8145	K.VTSSITIYPSDSSSPR.A	1	LUZP1_HUMAN
Q86V48	2	3.6646	R.EFALNNENYSLSNR.Q	3	LUZP1_HUMAN
Q86V48	3	5.0149	R.EKPDSDDDLDIASLVTAK.L	5	LUZP1_HUMAN
Q86V48	2	3.5892	R.IEDGISSTLPSK.E	3	LUZP1_HUMAN
Q86V48	2	4.65	R.MAELEKLEEAFSR.S	3	LUZP1_HUMAN
Q86V48	2	3.1023	R.SSIIIKPSDPVER.N	1	LUZP1_HUMAN
Q86V48	2	3.2384	R.SSTDFSELEQPR.S	2	LUZP1_HUMAN
Q86V48	2	5.2462	R.TSLFENDKDAGMENESVK.S	3	LUZP1_HUMAN
Q86V48	2	3.0998	K.IM#GGSGTETTLEK.Q	2	LUZP1_HUMAN
Q86V48	2	2.9198	K.SLTLSFVSER.K	1	LUZP1_HUMAN
Q86V48	2	3.3844	R.TFSDTTHGSVPSDPLGR.A	1	LUZP1_HUMAN
Q86V48	2	4.0435	K.SHSAPSEVGFSDAR.H	2	LUZP1_HUMAN
Q86V48	2	3.4751	K.NVESTNSNAYTQR.S	2	LUZP1_HUMAN
Q86V48	2	4.3967	K.LVNTTITPEPEPKPQPNSR.E	2	LUZP1_HUMAN
Q86V48	2	3.0936	K.KISSELEMLR.V	2	LUZP1_HUMAN
Q86V48	2	5.1118	K.ASHM#GVSTDSGTQETK.K	4	LUZP1_HUMAN
Q86V48	2	3.9564	K.IMGGSGTETTLEK.Q	2	LUZP1_HUMAN
Q86V48	3	5.1972	K.HFESLEEELKK.M	3	LUZP1_HUMAN
Q86V48	2	3.4924	K.EKLEEEENLTR.E	1	LUZP1_HUMAN
Q86V48	2	2.7414	K.DLNQEIEKLK.T	1	LUZP1_HUMAN
Q86V48	2	4.2673	K.ASHMGVSTDSGTQETK.K	4	LUZP1_HUMAN
Q86V48	3	5.7102	K.SKNNDLQDNYLSEQNKNK.L	2	LUZP1_HUMAN
Q86V48	2	2.9712	K.ISSELEMLR.V	1	LUZP1_HUMAN
Leucine zipper putative tumor suppressor 1 - Homo sapiens (Human)					
Q9Y250	2	3.9668	K.VNLLEQELQELR.A	3	LZTS1_HUMAN
Q9Y250	2	3.9681	K.QLQQSYVAMYQR.N	4	LZTS1_HUMAN
Leucine-rich alpha-2-glycoprotein precursor - Homo sapiens (Human)					
P02750	2	4.0333	R.VAAGAFQGLR.Q	2	A2GL_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P02750	3	3.757	R.NALTGLPPGLFQASATLDTLVLK.E	1	A2GL_HUMAN
P02750	2	2.7177	R.TLDLGENQLETLPPDLLR.G	1	A2GL_HUMAN
Leucine-rich repeat and calponin homology domain-containing protein 3 precursor - Homo sapien					
Q96II8	2	3.2301	R.EAQLAALQYEEEKIR.T	1	LRCH3_HUMAN
Leucine-rich repeat and calponin homology domain-containing protein 4 - Homo sapiens (Human)					
O75427	2	2.8139	R.REEPAGEER.R	2	LRCH4_HUMAN
Leucine-rich repeat flightless-interacting protein 1 - Homo sapiens (Human)					
Q32MZ4	2	5.2539	R.KSAVEAQNEVTENPK.Q	2	LRRF1_HUMAN
Q32MZ4	2	4.9336	K.SAVEAQNEVTENPK.Q	5	LRRF1_HUMAN
Q32MZ4	2	5.4081	R.NMPGLSAATLASLGGTSSR.R	2	LRRF1_HUMAN
Q32MZ4	3	4.7039	R.KSAVEAQNEVTENPKQK.I	1	LRRF1_HUMAN
Q32MZ4	3	4.8381	R.KALDSNSLENDDLSAPGREPGHFNPESR.E	3	LRRF1_HUMAN
Q32MZ4	2	3.3354	R.EIDCLSPEAQK.L	3	LRRF1_HUMAN
Q32MZ4	3	4.9965	R.ASEVEVKNEIVANVGKR.E	2	LRRF1_HUMAN
Q32MZ4	2	5.0529	R.AGGEELDEGVAKDNAK.I	6	LRRF1_HUMAN
Q32MZ4	2	4.1568	R.AGGEELDEGVAK.D	4	LRRF1_HUMAN
Q32MZ4	2	5.9506	K.SAVEAQNEVTENPKQK.I	3	LRRF1_HUMAN
Q32MZ4	2	3.6546	R.NM#PGLSAATLASLGGTSSR.R	1	LRRF1_HUMAN
Q32MZ4	2	2.7001	K.NEIVANVGKR.E	1	LRRF1_HUMAN
Q32MZ4	2	4.192	K.ELTYQNTDLSEIKEEEQVK.S	2	LRRF1_HUMAN
Q32MZ4	3	5.3224	K.HAHSILQFQFAEVK.E	3	LRRF1_HUMAN
Q32MZ4	2	3.4188	K.EALKQREEMLEK.H	2	LRRF1_HUMAN
Q32MZ4	2	3.9665	K.IAAESSENVDCPENPK.I	4	LRRF1_HUMAN
Q32MZ4	2	3.6739	K.IDGATQSSPAEPK.S	7	LRRF1_HUMAN
Q32MZ4	2	3.6636	K.IDGATQSSPAEPKSEDADR.C	2	LRRF1_HUMAN
Q32MZ4	3	7.0015	K.IKLDGKLDQEGDDVQTAAEEVLADGDTLDFE	1	LRRF1_HUMAN
Q32MZ4	3	7.1045	K.LDGKLDQEGDDVQTAAEEVLADGDTLDFEDD	1	LRRF1_HUMAN
Q32MZ4	2	3.7548	K.M#TKEELNALK.S	2	LRRF1_HUMAN
Leucine-rich repeat flightless-interacting protein 2 - Homo sapiens (Human)					
Q9Y608	3	5.0093	R.TALDKIEEMEMTNSHLAK.R	2	LRRF2_HUMAN
Q9Y608	2	3.5476	K.ELKESLSEVEEK.Y	2	LRRF2_HUMAN
Q9Y608	3	4.5136	K.LSKAEQDITTLEQSISR.L	2	LRRF2_HUMAN
Q9Y608	4	5.8758	K.LSKAEQDITTLEQSISRLEGQVLR.Y	2	LRRF2_HUMAN
Q9Y608	2	3.2044	K.MEELKEGLR.Q	2	LRRF2_HUMAN
Q9Y608	2	4.4053	R.FSAEDEALSNIAR.E	4	LRRF2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y608	3	4.8458	R.KLAGEKEELLSQIR.K	5	LRRF2_HUMAN
Q9Y608	2	4.1006	R.NSASATTPLSGNSSR.R	4	LRRF2_HUMAN
Q9Y608	3	4.4211	R.TALDKIEEMEM#TNSHLAK.R	1	LRRF2_HUMAN
Q9Y608	2	4.8823	K.AEQDITTLEQSISR.L	3	LRRF2_HUMAN
Leucine-rich repeat neuronal protein 3 precursor - Homo sapiens (Human)					
Q9H3W5	2	2.884	K.PTFALGELYPPLINLWEAGKEK.S	1	LRRN3_HUMAN
Leucine-rich repeat protein SHOC-2 - Homo sapiens (Human)					
Q9UQ13	2	2.8493	K.RPNPAPGTR.K	1	SHOC2_HUMAN
Leucine-rich repeat-containing protein 1 - Homo sapiens (Human)					
Q9BTT6	2	4.1427	R.FVEDEKDEEDNETR.T	3	LRRC1_HUMAN
Leucine-rich repeat-containing protein 16. - Homo sapiens (Human)					
Q5VZK9	2	3.0607	R.SDGGGAVPK.L	1	Q5VZK9_HUMA
Leucine-rich repeat-containing protein 47 - Homo sapiens (Human)					
Q8N1G4	2	4.1642	R.EGGDGEEQDVGDAGR.L	8	LRC47_HUMAN
Q8N1G4	2	4.7906	R.NALGPGLSPELGPLPALR.V	4	LRC47_HUMAN
Leucine-rich repeat-containing protein 59 - Homo sapiens (Human)					
Q96AG4	2	2.7841	K.AVQADQERER.Q	1	LRC59_HUMAN
Q96AG4	3	5.4959	K.LDGNELDLSLSDLNEVPVKELAALPK.A	2	LRC59_HUMAN
Q96AG4	3	6.6307	R.DKLDGNELDLSLSDLNEVPVKELAALPK.A	2	LRC59_HUMAN
Leucyl-cystinyl aminopeptidase - Homo sapiens (Human)					
Q9UIQ6	3	5.2854	R.GLGEHEM#EEDEEDYESSAK.L	3	LCAP_HUMAN
Q9UIQ6	2	4.7949	R.GLGEHEMEEDEEDYESSAK.L	4	LCAP_HUMAN
Leukocyte receptor cluster member 1 - Homo sapiens (Human)					
Q96BZ8	2	3.3057	R.ALQEGQPEEDETDDRR.R	2	LENG1_HUMAN
Q96BZ8	2	3.0156	R.SRAEALLAR.V	1	LENG1_HUMAN
Q96BZ8	2	4.048	R.YNSQFNPQLAR.R	4	LENG1_HUMAN
Q96BZ8	2	3.048	R.ALQEGQPEEDETDDR.R	1	LENG1_HUMAN
Leukocyte-associated immunoglobulin-like receptor 1 precursor - Homo sapiens (Human)					
Q6GTX8	3	4.8003	R.AVSPQSTKPMAESITYAAVAR.H	1	LAIR1_HUMAN
Q6GTX8	3	4.9569	R.FRIDSVSEGNAGPYR.C	4	LAIR1_HUMAN
Leupaxin - Homo sapiens (Human)					
O60711	3	3.8362	R.STLQDSDEYSNPAPLPLDQHSR.K	1	LPXN_HUMAN
Leydig cell tumor 10 kDa protein homolog - Homo sapiens (Human)					
Q9UNZ5	2	2.9326	K.NLEVGIRK.K	1	L10K_HUMAN
Q9UNZ5	2	3.1234	R.KKIEHDVVMK.A	2	L10K_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9UNZ5	2	3.8548	R.KKIEHDVVM#K.A	2	L10K_HUMAN
Q9UNZ5	2	3.4132	K.KIEHDVVMK.A	3	L10K_HUMAN
Q9UNZ5	2	3.1518	K.KIEHDVVM#K.A	2	L10K_HUMAN
Q9UNZ5	1	2.4897	K.NLEVGIR.K	1	L10K_HUMAN
Ligand-dependent corepressor - Homo sapiens (Human)					
Q96JN0	2	3.5316	K.NSSTQDPSQPNSTK.N	1	LCOR_HUMAN
Ligand-dependent nuclear receptor corepressor-like protein - Homo sapiens (Human)					
Q8N3X6	2	3.2637	R.ERMAAAAAAAAAAAAAAQCRSPR.C	1	Q8N3X6_HUMA
LIM and calponin homology domains-containing protein 1 - Homo sapiens (Human)					
Q9UPQ0	2	3.322	R.ASVLDTSMSAGSGSPSK.T	2	LIMC1_HUMAN
LIM and cysteine-rich domains protein 1 - Homo sapiens (Human)					
Q9NZU5	2	3.0394	K.MSLGQLQSAR.G	1	LMCD1_HUMAN
LIM and SH3 domain protein 1 - Homo sapiens (Human)					
Q14847	1	2.3172	K.MTLNMK.N	2	LASP1_HUMAN
Q14847	2	3.8803	R.YKEEFEKNK.G	6	LASP1_HUMAN
Q14847	1	3.0587	R.YKEEFEK.N	4	LASP1_HUMAN
Q14847	4	6.6878	R.RPLEQQQPHHIPTSAPVYQQPQQQPVAQSY	2	LASP1_HUMAN
Q14847	2	4.7941	R.LKQQSELQSQVR.Y	8	LASP1_HUMAN
Q14847	3	4.2319	K.TQDQISNIKYHEEFEKSR.M	2	LASP1_HUMAN
Q14847	2	5.5397	K.TQDQISNIKYHEEFEK.S	5	LASP1_HUMAN
Q14847	2	3.3927	K.TQDQISNIK.Y	5	LASP1_HUMAN
Q14847	2	4.5237	K.QSFTMVADTPENLR.L	10	LASP1_HUMAN
Q14847	2	2.8704	K.QQSELQSQVR.Y	2	LASP1_HUMAN
Q14847	2	5.4791	K.KTQDQISNIKYHEEFEK.S	1	LASP1_HUMAN
Q14847	2	4.1612	K.KTQDQISNIK.Y	4	LASP1_HUMAN
Q14847	3	5.2813	K.GKGFSVVADTPELQR.I	8	LASP1_HUMAN
Q14847	2	4.6219	K.GFSVVADTPELQR.I	15	LASP1_HUMAN
Q14847	2	3.931	K.QSFTM#VADTPENLR.L	8	LASP1_HUMAN
LIM domain and actin-binding protein 1 - Homo sapiens (Human)					
Q9UHB6	2	3.6948	R.STPAEDDSRDSQVK.S	4	LIMA1_HUMAN
LIM domain only protein 7 - Homo sapiens (Human)					
Q8WWI1	2	3.5131	R.NPSSSVPPPSAGSVK.T	4	LMO7_HUMAN
Q8WWI1	2	2.8861	K.EQVPSGAELER.Q	1	LMO7_HUMAN
Q8WWI1	2	5.1136	K.TSTTGVATTQSPTPR.S	3	LMO7_HUMAN
LIM domain-containing protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9UGP4	2	4.079	R.SSEGSLGGQNSGIGGR.S	2	LIMD1_HUMAN
Q9UGP4	1	2.2459	R.WEVVGSK.L	2	LIMD1_HUMAN
LIM-nebulette - Homo sapiens (Human)					
Q70I54	2	3.8093	K.QSFTTVADTPENLR.L	1	Q70I54_HUMAN
Q70I54	1	2.356	R.DFEESK.G	3	Q70I54_HUMAN
Q70I54	2	4.3051	R.LKQQSELQSQVK.Y	7	Q70I54_HUMAN
Q70I54	2	3.2182	R.TQEQISNVK.Y	2	Q70I54_HUMAN
Lin-7 homolog C - Homo sapiens (Human)					
Q9NUP9	2	3.3781	R.VLQSEFCNAVR.E	2	LIN7C_HUMAN
Lipin-2 - Homo sapiens (Human)					
Q92539	4	4.9924	R.SQHQGPDDIYLDDLKGLEPEVAALYFPK.S	1	LPIN2_HUMAN
Q92539	3	4.7667	K.SGGDETPSQSSDISHVLETETIFTPSSVK.K	2	LPIN2_HUMAN
Lipolysis-stimulated lipoprotein receptor - Homo sapiens (Human)					
Q86X29	2	4.9605	R.SSSAGGQGSYVPLLR.D	4	LSR_HUMAN
Q86X29	2	3.2526	R.IQASQQDDSM#R.V	2	LSR_HUMAN
Q86X29	2	3.8202	R.IQASQQDDSMR.V	2	LSR_HUMAN
Lipoma-preferred partner - Homo sapiens (Human)					
Q93052	2	3.5829	R.M#VIPNQPPLTATK.K	3	LPP_HUMAN
Q93052	2	4.7517	R.YYEGYYAAGPGYGGR.N	6	LPP_HUMAN
Q93052	2	2.9678	R.NDSDPTYGQQGHPNTWK.R	2	LPP_HUMAN
Q93052	2	3.6518	R.MVIPNQPPLTATK.K	2	LPP_HUMAN
Q93052	3	6.1953	R.METTHSFGNPSISVSTQQPPKK.F	7	LPP_HUMAN
Q93052	2	2.721	K.KFAPVVAPK.P	1	LPP_HUMAN
Q93052	3	5.7819	R.METTHSFGNPSISVSTQQPPK.K	5	LPP_HUMAN
Q93052	2	2.9135	R.M#VIPNQPPLTATKK.S	2	LPP_HUMAN
Q93052	3	3.824	R.M#ETTHSFGNPSISVSTQQPPK.K	1	LPP_HUMAN
Q93052	2	3.5948	K.STGEPLGHVPAR.M	2	LPP_HUMAN
Q93052	3	5.0937	K.REPGYTPPGAGNQNPPGMYPVTGPK.K	2	LPP_HUMAN
Q93052	3	4.2269	K.KFAPVVAPKPK.Y	4	LPP_HUMAN
Q93052	2	2.9126	K.MLYDM#ENPPADEYFGR.C	2	LPP_HUMAN
Q93052	3	4.866	K.GGHSGQLGPSSVAPSFRPEDELEHLTKK.M	2	LPP_HUMAN
Q93052	2	2.8064	K.FAPVVAPKPK.Y	1	LPP_HUMAN
Q93052	3	5.2134	R.M#ETTHSFGNPSISVSTQQPPKK.F	4	LPP_HUMAN
Lipopolysaccharide-responsive and beige-like anchor protein - Homo sapiens (Human)					
P50851	2	5.6011	R.DINVSVGSQQPDTK.D	4	LRBA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Liprin-alpha-1 - Homo sapiens (Human)					
Q13136	2	3.2878	K.VQTLNEQDWER.A	2	LIPA1_HUMAN
Q13136	2	2.8955	R.VTSSM#SSPSM#QPK.K	2	LIPA1_HUMAN
Q13136	2	3.6974	R.LLDTLRETQETLALTQGK.L	1	LIPA1_HUMAN
Q13136	3	4.1149	R.LIQEEKENTEQRAEEIESR.V	1	LIPA1_HUMAN
Q13136	2	5.1968	R.GLAAGSAETLPANFR.V	6	LIPA1_HUMAN
Q13136	3	4.4582	R.LAALRDEPSKVQTLNEQDWER.A	1	LIPA1_HUMAN
Liprin-beta-1 - Homo sapiens (Human)					
Q86W92	2	5.35	R.GLLEMMETDEKEGLR.C	6	LIPB1_HUMAN
Q86W92	3	5.047	R.TSLETQKLDLMAEISNLK.L	2	LIPB1_HUMAN
Q86W92	3	5.0768	R.TSLETQKLDLM#AEISNLK.L	3	LIPB1_HUMAN
Q86W92	3	5.5299	R.LYEEDDLDRLEQMEDSEGTVR.Q	2	LIPB1_HUMAN
Q86W92	2	3.1483	K.ETSDGEKETIQK.T	1	LIPB1_HUMAN
Q86W92	3	5.0027	R.LYEEDDLDRLEQM#EDSEGTVR.Q	1	LIPB1_HUMAN
Q86W92	2	3.6799	R.DTEGLIQEINDLR.L	2	LIPB1_HUMAN
Q86W92	2	2.7599	K.SSSLGNLKK.E	1	LIPB1_HUMAN
Q86W92	3	5.7309	K.KAVESLMAANEEKDR.K	2	LIPB1_HUMAN
Q86W92	3	4.3641	K.AVESLMAANEEKDR.K	4	LIPB1_HUMAN
Q86W92	2	3.4182	K.MQDTVVLAQGK.D	2	LIPB1_HUMAN
Liprin-beta-2 - Homo sapiens (Human)					
Q8ND30	2	3.4389	K.SPPTICQPDATGSSLLR.L	1	LIPB2_HUMAN
Q8ND30	2	2.8763	R.RIEELTGLLNQYR.K	1	LIPB2_HUMAN
Q8ND30	3	4.0914	R.RIEELTGLLNQYRK.V	1	LIPB2_HUMAN
L-lactate dehydrogenase A chain - Homo sapiens (Human)					
P00338	2	2.8814	K.DYNVTANSK.L	1	LDHA_HUMAN
P00338	2	4.1632	K.SADTLWGIQK.E	6	LDHA_HUMAN
P00338	2	3.6613	K.VTLTSEEEAR.L	4	LDHA_HUMAN
L-lactate dehydrogenase B chain - Homo sapiens (Human)					
P07195	2	3.7406	K.LIAPVAEEEATVPNNK.I	2	LDHB_HUMAN
P07195	2	3.3674	K.SADTLWDIQK.D	4	LDHB_HUMAN
LOC387763 protein - Homo sapiens (Human)					
Q7Z7L8	2	3.5463	K.LRNSLDSSDSDSAL.-	2	Q7Z7L8_HUMAN
Q7Z7L8	1	3.1928	R.STQSLSLQR.E	5	Q7Z7L8_HUMAN
Q7Z7L8	2	3.868	R.RSTQSLSLQR.E	3	Q7Z7L8_HUMAN
LOC649305 protein - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q4G118	2	2.7326	R.QVATAEERSGSHM.-	1	Q4G118_HUMA
Low-density lipoprotein receptor-related protein 2 precursor - Homo sapiens (Human)					
P98164	2	2.89	K.QPIIFENPM#YSAR.D	1	LRP2_HUMAN
P98164	2	3.1737	R.DPTPTYSATEDTFK.D	3	LRP2_HUMAN
P98164	2	3.3027	R.TVVQYLNNPR.S	2	LRP2_HUMAN
LRRFIP1 protein - Homo sapiens (Human)					
Q9BSL6	3	5.1344	R.SALDKTEELEVSNGHLVK.R	3	Q9BSL6_HUMA
Q9BSL6	3	4.2314	K.SAAENAEKIEDELKAEK.R	2	Q9BSL6_HUMA
L-selectin precursor - Homo sapiens (Human)					
Q9UJ43	2	2.8175	K.SLTEEAENWGDGEPNNKK.N	1	Q9UJ43_HUMAN
LSM14 protein homolog A - Homo sapiens (Human)					
Q8ND56	2	3.183	K.SFFDNISCDDNR.E	2	LS14A_HUMAN
Q8ND56	2	3.1705	K.VSRPENEQLR.N	2	LS14A_HUMAN
Q8ND56	3	5.3811	K.DFDFESANAQFNKEEIDREFHNK.L	1	LS14A_HUMAN
Lumican precursor - Homo sapiens (Human)					
P51884	1	3.3595	R.FNALQYLR.L	6	LUM_HUMAN
P51884	4	5.4073	R.LPSGLPVSLLTLYLDNNKISNIPDEYFK.R	2	LUM_HUMAN
P51884	2	5.4336	R.LPSGLPVSLLTLYLDNNK.I	3	LUM_HUMAN
P51884	2	3.5169	R.NNQIDHIDEK.A	2	LUM_HUMAN
P51884	2	3.3789	R.ISETSLPPDMYECLR.V	2	LUM_HUMAN
P51884	2	4.2406	K.SLEDLQLTHNK.I	5	LUM_HUMAN
P51884	2	3.3774	K.RFNALQYLR.L	2	LUM_HUMAN
P51884	2	2.8855	K.ISNIPDEYFKR.F	1	LUM_HUMAN
P51884	1	2.4816	K.ILGPLSYSK.I	1	LUM_HUMAN
P51884	2	3.7736	K.SLEDLQLTHNKITK.L	1	LUM_HUMAN
P51884	2	3.8163	R.LKEDAVSAAFK.G	5	LUM_HUMAN
Lupus La protein - Homo sapiens (Human)					
P05455	2	3.0962	K.GKVQFQGKK.T	2	LA_HUMAN
P05455	2	4.6916	R.LTTDFNVIVEALSK.S	4	LA_HUMAN
P05455	2	2.797	R.EETDKEEPASK.Q	1	LA_HUMAN
P05455	3	4.7174	R.AREETDKEEPASK.Q	18	LA_HUMAN
P05455	1	2.52	K.VQFQGK.K	3	LA_HUMAN
P05455	2	4.4406	K.IIEDQQESLNK.W	4	LA_HUMAN
P05455	2	2.9117	K.DANNGNLQLR.N	1	LA_HUMAN
P05455	2	4.2746	K.AKDANNGNLQLR.N	2	LA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P05455	2	3.4774	K.IIEDQQESLNKWK.S	2	LA_HUMAN
Lutheran blood group glycoprotein precursor - Homo sapiens (Human)					
P50895	2	3.0859	R.LEVPVEM#NPEGYM#TSR.T	1	LU_HUMAN
P50895	2	2.8832	R.VAYLDPLELSEGK.V	1	LU_HUMAN
P50895	2	3.9223	R.SPPYQLDSQGR.L	4	LU_HUMAN
P50895	2	3.3463	R.LEVPVEM#NPEGYMTSR.T	1	LU_HUMAN
P50895	2	3.8572	R.GRSPPYQLDSQGR.L	1	LU_HUMAN
P50895	2	2.7725	R.GDGSPSPEYTLFR.L	2	LU_HUMAN
P50895	2	3.8716	R.AGAAGTAEATAR.L	5	LU_HUMAN
P50895	1	2.2347	K.VTSALSR.D	1	LU_HUMAN
P50895	2	3.1525	R.LASAEMQGSELQVTMHDTR.G	1	LU_HUMAN
Lymphocyte cytosolic protein 2 - Homo sapiens (Human)					
Q13094	2	3.4354	R.SLAPFDREPFTLGK.K	2	LCP2_HUMAN
Q13094	2	2.9028	K.LSQEINKNEER.R	1	LCP2_HUMAN
Q13094	2	2.7832	R.SLGEHLPK.I	1	LCP2_HUMAN
Q13094	2	2.7293	R.SEVLGWDPDSLADYFKK.L	1	LCP2_HUMAN
Q13094	2	4.8108	R.FLNLTENDIQKFPK.L	2	LCP2_HUMAN
Q13094	2	2.9947	K.LSQEINKNEERR.S	2	LCP2_HUMAN
Lymphocyte-specific protein 1 - Homo sapiens (Human)					
P33241	4	7.3761	R.SPEGEQEDRPGLHAYEKEDSDEVHLEELSLS	5	LSP1_HUMAN
P33241	2	3.0714	K.FVATGHGKYEK.V	2	LSP1_HUMAN
P33241	2	4.6566	K.IDQWLEQYTQAIETAGR.T	9	LSP1_HUMAN
P33241	2	3.2381	K.KSQPDLPISK.I	4	LSP1_HUMAN
P33241	2	3.0158	K.LIDRTESLNR.S	5	LSP1_HUMAN
P33241	3	3.8826	K.QEMLLSLKPSEAPELDEDEGFGDWSQRPEQ	1	LSP1_HUMAN
P33241	1	2.2219	K.SQPDLPISK.I	1	LSP1_HUMAN
P33241	3	4.1344	R.QQHEGAQGALDSGEPPQCR.S	2	LSP1_HUMAN
P33241	3	5.3057	K.EGPGPEDTVQDNLGAAGAEEEQEEHQK.C	6	LSP1_HUMAN
LysM and putative peptidoglycan-binding domain-containing protein 2 - Homo sapiens (Human)					
Q8IV50	1	2.15	R.SYGSTASVR.A	1	LYSM2_HUMAN
Q8IV50	2	3.5741	R.APLGAGVIER.H	4	LYSM2_HUMAN
Lysosomal alpha-glucosidase precursor - Homo sapiens (Human)					
P10253	2	3.2006	R.AGYIIPLQGPGLTTTESR.Q	1	LYAG_HUMAN
Lysosomal alpha-mannosidase precursor - Homo sapiens (Human)					
O00754	2	2.8364	R.GVSEPLMENGSGAWVR.G	1	MA2B1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O00754	2	2.957	R.IYITDGNMQLTVLTDR.S	1	MA2B1_HUMAN
Lysosomal-associated membrane protein 1 - Homo sapiens (Human)					
Q8WU33	2	3.2043	R.ALQATVGNSYK.C	2	Q8WU33_HUMA
Lysozyme C precursor - Homo sapiens (Human)					
P61626	1	2.1056	R.QYVQGCGV.-	1	LYSC_HUMAN
P61626	2	4.5298	R.STDYGIFQINSR.Y	5	LYSC_HUMAN
Macrophage migration inhibitory factor - Homo sapiens (Human)					
P14174	2	3.6327	-.MPMFIVNTNVPR.A	2	MIF_HUMAN
MAGUK p55 subfamily member 5 - Homo sapiens (Human)					
Q8N3R9	2	3.6893	K.ILEIEDLFSSLK.H	2	MPP5_HUMAN
MAK16-like protein RBM13 - Homo sapiens (Human)					
Q9BXY0	3	4.7853	K.ALIAAQLDNAIEKELLER.L	2	RBM13_HUMAN
Malate dehydrogenase, cytoplasmic - Homo sapiens (Human)					
P40925	2	2.9457	K.FVEGLPINDFSR.E	1	MDHC_HUMAN
Malate dehydrogenase, mitochondrial precursor - Homo sapiens (Human)					
P40926	2	2.8393	K.KGEDFVK.T	1	MDHM_HUMAN
Male-enhanced antigen 1 - Homo sapiens (Human)					
Q16626	2	3.5869	R.EISDAQWEDVVQK.A	2	MEA1_HUMAN
Q16626	2	4.1318	R.TM#AGVSLPAPGVPAWAR.E	1	MEA1_HUMAN
Q16626	2	3.7213	R.TMAGVSLPAPGVPAWAR.E	3	MEA1_HUMAN
Mannose-6-phosphate receptor-binding protein 1 - Homo sapiens (Human)					
O60664	2	3.1997	K.SVVTGGVQSVM#GSR.L	1	M6PBP_HUMAN
O60664	3	3.845	R.GLDKLEENLPILQQPTEK.V	1	M6PBP_HUMAN
O60664	2	3.2651	R.GAVQSGVDKTK.S	4	M6PBP_HUMAN
O60664	3	5.7882	R.TLTAAAVSGAQPILSK.L	7	M6PBP_HUMAN
O60664	2	2.7873	R.GAVQSGVDK.T	2	M6PBP_HUMAN
O60664	2	4.5451	K.SVVTGGVQSVMGSR.L	2	M6PBP_HUMAN
O60664	2	3.5502	K.LEPQIASASEYAHR.G	2	M6PBP_HUMAN
O60664	2	3.2621	K.DTVATQLSEAVDATR.G	1	M6PBP_HUMAN
O60664	2	4.5508	K.VSGAQEMVSSAK.D	3	M6PBP_HUMAN
O60664	2	3.8416	K.VLADTKELVSSK.V	1	M6PBP_HUMAN
Mannosyl-oligosaccharide glucosidase - Homo sapiens (Human)					
Q13724	2	2.8714	R.RAVPAEGVR.T	1	GCS1_HUMAN
MAP kinase-activating death domain protein - Homo sapiens (Human)					
Q8WXG6	2	3.427	R.LASDSDAESDSR.A	4	MADD_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
MARCKS-related protein - Homo sapiens (Human)					
P49006	2	3.655	K.AAATPESQEPQAK.G	5	MRP_HUMAN
P49006	3	7.0992	R.KEGGGDSSASSPTEEEQEQGEIGACSDEGT	1	MRP_HUMAN
Matrin-3 - Homo sapiens (Human)					
P43243	3	4.4436	K.TDGSQKTESSTEGKEQEEK.S	1	MATR3_HUMAN
P43243	2	2.9068	R.TEEGPTLSYGR.D	2	MATR3_HUMAN
P43243	3	6.4348	R.DLSAAGIGLLAAATQSLSMPASLGR.M	1	MATR3_HUMAN
P43243	3	4.3036	R.DLDELSRYPEDKITPENLPQILLQLK.R	1	MATR3_HUMAN
P43243	3	6.3238	R.DLSAAGIGLLAAATQSLSM#PASLGR.M	2	MATR3_HUMAN
Matrix Gla protein precursor - Homo sapiens (Human)					
P08493	2	4.2368	R.NANTFISPQQR.W	4	MGP_HUMAN
P08493	2	3.629	R.YAMVYGYNAAYNR.Y	2	MGP_HUMAN
Matrix-remodeling-associated protein 5 precursor - Homo sapiens (Human)					
Q9NR99	2	3.0556	R.VHASHQLTR.V	1	MXRA5_HUMAN
Q9NR99	3	4.4517	K.HSEKEPETNVAEGR.R	1	MXRA5_HUMAN
Q9NR99	2	3.0135	K.SSLSTQDTLLIK.K	2	MXRA5_HUMAN
Matrix-remodeling-associated protein 7 - Homo sapiens (Human)					
P84157	2	2.8437	R.RGAAASPEPAR.A	1	MXRA7_HUMAN
P84157	3	5.2398	R.QEEEQDLDGEKGPSSEGPEEEDGEGFSFK.Y	4	MXRA7_HUMAN
P84157	3	5.2263	K.DNKETFGEMSDGDVQEQLR.L	1	MXRA7_HUMAN
P84157	2	3.1216	R.GAAASPEPAR.A	1	MXRA7_HUMAN
Matrix-remodelling associated protein 7 - Homo sapiens (Human)					
Q6ZR64	2	4.3176	K.EELEEEQRTEE.-	4	Q6ZR64_HUMA
Q6ZR64	2	4.078	K.MMTKEELEEEQRTEE.-	1	Q6ZR64_HUMA
Max-binding protein MNT - Homo sapiens (Human)					
Q99583	2	3.9168	R.FLEWQAQQQQR.A	2	MNT_HUMAN
Mediator of DNA damage checkpoint protein 1 - Homo sapiens (Human)					
Q14676	3	4.7293	R.DTQRGEPEGGSQDQK.G	1	MDC1_HUMAN
Mediator of RNA polymerase II transcription subunit 28 - Homo sapiens (Human)					
Q9H204	2	3.0671	K.EDVSELRNELQR.K	1	MED28_HUMAN
Melanoma antigen family B, 16 - Homo sapiens (Human)					
A2A368	1	2.1191	K.EAPAAK.A	2	A2A368_HUMAN
Melanoma inhibitory activity protein 3 precursor - Homo sapiens (Human)					
Q5JRA6	3	6.2215	R.WSAEASGKPSPSDPGSGTATM#M#NSSSR.	2	MIA3_HUMAN
Q5JRA6	2	4.6306	R.TQTAISVVEEDLK.L	3	MIA3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5JRA6	2	3.9904	R.SEFGSVDGPLPHPR.W	3	MIA3_HUMAN
Q5JRA6	2	3.7663	R.GWPEVWAGSVGR.T	3	MIA3_HUMAN
Q5JRA6	1	2.2175	R.DVAATASK.Q	1	MIA3_HUMAN
Q5JRA6	2	3.4829	R.ASESQILSIAEK.M	2	MIA3_HUMAN
Q5JRA6	2	3.1693	K.KGDPVYVYYK.L	3	MIA3_HUMAN
Q5JRA6	3	6.4837	R.WSAEASGKPSPSDPGSGTATMMNSSSR.G	2	MIA3_HUMAN
Q5JRA6	3	4.4693	K.YFNVHELEALLQEMSSK.L	1	MIA3_HUMAN
Melanoma-associated antigen D2 - Homo sapiens (Human)					
Q9UNF1	1	2.9861	K.ALEVSEDVK.V	4	MAGD2_HUMAN
Q9UNF1	2	4.0449	K.ATEVSKTPEAR.E	5	MAGD2_HUMAN
Melanoma-associated antigen G1 - Homo sapiens (Human)					
Q96MG7	2	2.7408	R.ARPQPSGPAPSS.-	1	MAGG1_HUMAN
Q96MG7	2	3.8254	R.DGFAEEAPSTSR.G	4	MAGG1_HUMAN
Q96MG7	2	5.0407	R.GPGGSQGSQGPSPQGAR.R	2	MAGG1_HUMAN
Membrane-associated progesterone receptor component 1 - Homo sapiens (Human)					
O00264	3	5.4354	K.LLKEGEEPTVYSDEEEPKDESAR.K	4	PGRC1_HUMAN
O00264	2	3.0557	R.GDQPAASGDSDDDEPPPLPR.L	1	PGRC1_HUMAN
O00264	3	4.3205	K.IVRGDQPAASGDSDDDEPPPLPR.L	1	PGRC1_HUMAN
O00264	2	4.0647	K.FYGPEGPYGVFAGR.D	4	PGRC1_HUMAN
O00264	2	3.5332	K.EGEEPTVYSDEEEPKDESAR.K	1	PGRC1_HUMAN
O00264	2	4.8762	R.KFYGPEGPYGVFAGR.D	9	PGRC1_HUMAN
Membrane-associated progesterone receptor component 2 - Homo sapiens (Human)					
O15173	2	3.881	K.FYGPAGPYGIFAGR.D	1	PGRC2_HUMAN
Mesoderm development candidate 2 - Homo sapiens (Human)					
Q14696	3	4.3588	K.DIRDYNDADM#AR.L	3	MESD2_HUMAN
Q14696	3	3.9824	K.DIRDYNDADMAR.L	4	MESD2_HUMAN
Q14696	2	3.316	R.DYNDADMAR.L	4	MESD2_HUMAN
Metal transporter CNNM3 - Homo sapiens (Human)					
Q8NE01	3	3.8547	K.TTTAAGSSHSRPGVPVEGSPGRNPGV.-	2	CNNM3_HUMAN
Metallothionein-3 - Homo sapiens (Human)					
P25713	2	3.3058	K.GGEAAEAEAEK.C	3	MT3_HUMAN
Metastasis suppressor protein 1 - Homo sapiens (Human)					
O43312	3	5.2185	R.LSSVSSHDSGFISQDAFQSK.S	3	MTSS1_HUMAN
O43312	2	2.9434	R.SSNLAQQAPVR.L	1	MTSS1_HUMAN
O43312	2	2.8657	R.DTPQGEDM#LNAIR.R	1	MTSS1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Methionine aminopeptidase 2 - Homo sapiens (Human)					
P50579	3	5.4377	K.GPSAAGEQEPDKESGASVDEVAR.Q	1	AMPM2_HUMAN
P50579	3	5.7172	K.SKGPSAAGEQEPDKESGASVDEVAR.Q	3	AMPM2_HUMAN
Methionine-R-sulfoxide reductase B2 - Homo sapiens (Human)					
Q9Y3D2	2	3.7663	R.RLDTSLGSAR.T	2	MSRB2_HUMAN
Methionyl-tRNA synthetase, cytoplasmic - Homo sapiens (Human)					
P56192	2	3.7397	K.ADKNEVAAEVAK.L	2	SYMC_HUMAN
P56192	3	4.2474	K.AQKADKNEVAAEVAK.L	2	SYMC_HUMAN
Methylated-DNA--protein-cysteine methyltransferase - Homo sapiens (Human)					
P16455	2	3.3895	K.GAGATSGSPPAGRN.-	2	MGMT_HUMAN
Methyl-CpG-binding domain protein 2 - Homo sapiens (Human)					
Q9UBB5	2	2.7867	R.YLGNTVDLSSFDFR.T	1	MBD2_HUMAN
Q9UBB5	3	5.1584	R.KKLEEALMADILSR.A	2	MBD2_HUMAN
Methyl-CpG-binding domain protein 3 - Homo sapiens (Human)					
O95983	3	5.6309	K.RLEEALMADMLAHVEELAR.D	1	MBD3_HUMAN
Methyl-CpG-binding protein 2 - Homo sapiens (Human)					
P51608	2	4.7611	K.VGDTSLDPNDFDFTVTGR.G	8	MECP2_HUMAN
P51608	2	3.774	R.GPMYDDPTLPEGWTR.K	1	MECP2_HUMAN
P51608	2	3.8939	R.SVQETVLPIKK.R	2	MECP2_HUMAN
P51608	1	2.2354	R.SVQETVLPIK.K	1	MECP2_HUMAN
P51608	2	5.0626	R.SAGKYDVYLINPQGK.A	4	MECP2_HUMAN
P51608	3	4.0954	R.KPGSVVAAAAAEAKK.K	1	MECP2_HUMAN
P51608	2	4.9989	R.KPGSVVAAAAAEAK.K	5	MECP2_HUMAN
P51608	2	3.9627	R.KAEADPQAIPK.K	4	MECP2_HUMAN
P51608	2	4.5704	R.GRKPGSVVAAAAAEAK.K	3	MECP2_HUMAN
P51608	2	3.4158	K.AATSEGVQVKR.V	2	MECP2_HUMAN
P51608	3	3.9191	R.DRGPM#YDDPTLPEGWTR.K	1	MECP2_HUMAN
P51608	2	3.1627	K.AATSEGVQVK.R	3	MECP2_HUMAN
P51608	2	2.9458	K.MPFQTSPGGK.A	2	MECP2_HUMAN
P51608	2	3.8979	K.EVVKPLLVSTLGEK.S	4	MECP2_HUMAN
P51608	2	4.2822	K.AEGGGATTSTQVMVIK.R	5	MECP2_HUMAN
P51608	2	3.6856	K.AEGGGATTSTQVM#VIK.R	2	MECP2_HUMAN
P51608	2	2.8172	K.AEADPQAIPK.K	1	MECP2_HUMAN
P51608	2	4.0459	R.DRGPMYDDPTLPEGWTR.K	3	MECP2_HUMAN
Methylosome subunit pICln - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P54105	2	4.7773	K.GLGTGTLYIAESR.L	7	ICLN_HUMAN
MGC11257 protein - Homo sapiens (Human)					
Q9BRJ6	3	4.7674	K.GEAVLRPGLDAEPELSPEEQR.V	1	Q9BRJ6_HUMA
Mi2-beta (Human)					
Q14839	2	2.9277	R.AYHLVC*LDPELEK.A	2	CHD4_HUMAN
Q14839	2	4.5646	R.LANRAPEPTPQQVAQQQ.-	2	CHD4_HUMAN
MICAL-like protein 1 - Homo sapiens (Human)					
Q8N3F8	3	4.8519	K.QQHQQQLAEDAKDVPGGGPSSSAPAGAEAD	1	MILK1_HUMAN
Q8N3F8	3	5.9321	K.SSSEPAVHAPGTPGNPVSLSTNSSLASSGEL	2	MILK1_HUMAN
Q8N3F8	2	2.9778	R.SSLQQENLVEQAGSSSLVNGR.L	1	MILK1_HUMAN
Q8N3F8	2	4.2557	R.VEQMPQASPGLAPR.T	4	MILK1_HUMAN
MICAL-like protein 2 - Homo sapiens (Human)					
Q8IY33	2	3.7702	K.TEAPQASPLAKPLQSSSPR.V	1	MILK2_HUMAN
Q8IY33	2	2.8401	R.ASEDSEEEPSGK.K	1	MILK2_HUMAN
Microfibrillar-associated protein 1 - Homo sapiens (Human)					
P55081	2	2.9848	R.NLTEEERR.A	1	MFAP1_HUMAN
P55081	3	3.9487	K.AKEQEAEPEEQEEDSSSDPR.L	3	MFAP1_HUMAN
P55081	2	3.7842	K.EQEAEPEEQEEDSSSDPR.L	3	MFAP1_HUMAN
P55081	2	3.9613	R.DFSAPTLEDHFNK.T	2	MFAP1_HUMAN
P55081	2	3.7438	R.GAFFMDEDEEVYKR.D	3	MFAP1_HUMAN
P55081	2	3.1069	R.ISEDVEER.L	4	MFAP1_HUMAN
P55081	2	2.8443	R.MRNLTEEER.R	1	MFAP1_HUMAN
P55081	1	2.2588	R.NLTEEER.R	1	MFAP1_HUMAN
Microphthalmia-associated transcription factor - Homo sapiens (Human)					
O75030	2	3.532	K.LQAAQFMQQR.V	2	MITF_HUMAN
Microtubule associated serine/threonine kinase family member 4 - Homo sapiens (Human)					
Q8N4X4	2	3.227	R.APAAWAPASVLLER.G	1	Q8N4X4_HUMA
Microtubule-actin cross-linking factor 1, isoform 4 - Homo sapiens (Human)					
Q96PK2	2	3.2249	K.IEESSSQVVPQGISVK.H	1	MACF4_HUMAN
Microtubule-associated protein 1A - Homo sapiens (Human)					
P78559	3	3.8282	K.LAKREEVVEEGAK.E	1	MAP1A_HUMAN
P78559	2	3.3356	K.REEVVEEGAK.E	1	MAP1A_HUMAN
P78559	3	4.4197	R.FHTSTYDLPGPEGAGPFEASQPADSAVPATS	2	MAP1A_HUMAN
Microtubule-associated protein 2 - Homo sapiens (Human)					
Q8IUX2	2	2.8619	K.VSNSTLSK.I	1	Q8IUX2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P11137	3	5.3037	K.DMQGTEEEKAPLALFGHTLVASLEDMK.Q	2	MAP2_HUMAN
P11137	2	3.1423	R.LASVSADAEVAR.R	1	MAP2_HUMAN
P11137	2	2.939	R.FAALEQPEVER.R	2	MAP2_HUMAN
P11137	3	3.9248	K.APQEADAFMGVESGHMK.E	1	MAP2_HUMAN
P11137	2	3.5892	K.HAALVSQPETTK.T	2	MAP2_HUMAN
P11137	2	3.2274	K.DLSIPTDASSEKAEK.G	1	MAP2_HUMAN
P11137	3	4.7222	K.DAHIPVVEEHVMGK.V	3	MAP2_HUMAN
P11137	3	5.239	K.VLEEEKEAINQETVQQR.D	2	MAP2_HUMAN
Microtubule-associated protein 4 isoform 1 variant - Homo sapiens (Human)					
Q59FT2	3	4.1108	K.GISEDSHLESLQDVGQSAAPTFMISPETVTGT	1	Q59FT2_HUMAN
Q59FT2	3	7.0591	K.GISEDSHLESLQDVGQSAAPTFMISPETVTGT	3	Q59FT2_HUMAN
Microtubule-associated protein 8 - Homo sapiens (Human)					
Q27QB1	3	4.629	R.HDPLPDPLKVPPPLPDPSSICMVDPEMLPPK.	1	Q27QB1_HUMA
Microtubule-associated protein RP/EB family member 1 - Homo sapiens (Human)					
Q15691	3	3.8321	R.QGQETAVAPSLVAPALNKPK.K	1	MARE1_HUMAN
Q15691	2	2.9979	R.NIELICQENEGENDPVLQR.I	1	MARE1_HUMAN
Q15691	3	4.9675	K.NPGVGNGDDEAAELM#QQVNVLK.L	3	MARE1_HUMAN
Q15691	2	4.6152	K.LTVEDLEKERDFYFGK.L	2	MARE1_HUMAN
Q15691	2	3.1465	K.LTVEDLEKER.D	2	MARE1_HUMAN
Q15691	2	3.6376	K.KPLTSSSAAPQRPISTQR.T	1	MARE1_HUMAN
Q15691	3	3.7183	K.LRNIELICQENEGENDPVLQR.I	1	MARE1_HUMAN
Microtubule-associated protein RP/EB family member 2 - Homo sapiens (Human)					
Q15555	2	3.1333	K.LALEGVEKER.D	2	MARE2_HUMAN
Q15555	3	5.9646	K.SDKDLETQVIQLNEQVHSLK.L	3	MARE2_HUMAN
Q15555	2	2.7573	K.SSPAAKPGSTPSRPSSAK.R	1	MARE2_HUMAN
Microtubule-associated protein tau - Homo sapiens (Human)					
P10636	2	5.1361	R.GPSLGEDTKEADLPEPSEK.Q	6	TAU_HUMAN
P10636	2	3.0882	K.SRLQTAPVPM#PDLK.N	1	TAU_HUMAN
P10636	2	4.0946	K.STPTAEDVTAPLVDEGAPGK.Q	5	TAU_HUMAN
P10636	1	2.421	K.TDHGAEIVYK.S	2	TAU_HUMAN
P10636	2	4.0449	R.SGYSSPGSPGTPGSR.S	5	TAU_HUMAN
P10636	2	4.0354	K.VSTEIPASEPDGPSVGR.A	7	TAU_HUMAN
P10636	2	3.9797	K.TPPSSGEPPKSGDR.S	6	TAU_HUMAN
P10636	2	2.8358	R.GPSLGEDTK.E	1	TAU_HUMAN
P10636	4	5.1091	R.GPSLGEDTKEADLPEPSEKQPAAAPR.G	1	TAU_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P10636	3	3.9768	R.HLSNVSSTGSIDMVDSPQLATLADEVSASLAK	1	TAU_HUMAN
P10636	2	3.446	R.LQTAPVPMPDLK.N	4	TAU_HUMAN
P10636	2	3.3209	R.QEFEVMEDHAGTYGLGDR.K	2	TAU_HUMAN
P10636	3	4.0218	R.QEFEVMEDHAGTYGLGDRK.D	1	TAU_HUMAN
P10636	2	3.9625	K.SPVVSGDTSPR.H	4	TAU_HUMAN
P10636	2	3.976	R.EATRQPSGTGPEDTEGGR.H	6	TAU_HUMAN
P10636	3	3.8234	K.HQLLGDLHQEGPPLK.G	1	TAU_HUMAN
P10636	2	2.708	R.AAFPGAPGEGPEAR.G	1	TAU_HUMAN
P10636	2	4	K.AKTDHGAEIVYK.S	4	TAU_HUMAN
P10636	2	3.3583	K.ASPAQDGRPPQTAAR.E	1	TAU_HUMAN
P10636	3	3.9739	K.DQGGYTM#HQDQEGDTDAGLK.E	2	TAU_HUMAN
P10636	2	4.0699	K.DQGGYTMHQDQEGDTDAGLK.E	2	TAU_HUMAN
P10636	2	3.0254	K.SEKLDFKDR.V	1	TAU_HUMAN
P10636	2	3.2096	K.GQANATRIPAK.T	2	TAU_HUMAN
P10636	2	5.2742	K.HQLLGDLHQEGPPLKGAGGK.E	4	TAU_HUMAN
P10636	2	4.4253	K.IGSLDNITHVPGGGNK.K	3	TAU_HUMAN
P10636	2	4.6499	K.IGSLDNITHVPGGGNKK.I	2	TAU_HUMAN
P10636	1	2.3408	K.IGSTENLK.H	2	TAU_HUMAN
P10636	2	3.2376	K.KLDLSNVQSK.C	3	TAU_HUMAN
P10636	2	2.7795	K.LDLSNVQSK.C	2	TAU_HUMAN
P10636	2	3.0508	K.LTFRENAK.A	4	TAU_HUMAN
P10636	3	3.7653	K.ESPLQTPTEDGSEEPGSETSDAK.S	3	TAU_HUMAN
Microtubule-associated serine/threonine-protein kinase 2 - Homo sapiens (Human)					
Q6P0Q8	2	2.9211	R.SLGPMVPSLLTGITLGPPR.M	2	MAST2_HUMAN
Microtubule-associated serine/threonine-protein kinase 4 - Homo sapiens (Human)					
O15021	2	3.6624	R.EASAASSDTSSAK.A	6	MAST4_HUMAN
O15021	2	3.8757	R.SPGTVM#ESNPQQR.E	2	MAST4_HUMAN
O15021	2	3.3563	K.SGPDVFPATPGSQNK.A	2	MAST4_HUMAN
O15021	2	3.6459	K.GKEPATQSLGGSSR.E	1	MAST4_HUMAN
O15021	3	3.9221	R.ASEGAM#SDGPVPAEHR.Q	1	MAST4_HUMAN
Mimitin, mitochondrial precursor - Homo sapiens (Human)					
Q8N183	3	3.9566	K.GHASAPYFGKEEPSVAPSSTGK.T	1	MIMIT_HUMAN
MIR-interacting saposin-like protein precursor - Homo sapiens (Human)					
Q9Y2B0	1	3.1956	R.IDSDISGTLK.F	4	MSAP_HUMAN
Q9Y2B0	2	4.8635	R.SEAHLTELLEEICDR.M	6	MSAP_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y2B0	3	3.7912	R.NGESSELDLQGIRIDSDISGTLK.F	1	MSAP_HUMAN
Q9Y2B0	2	3.686	R.NGESSELDLQGIR.I	2	MSAP_HUMAN
Q9Y2B0	2	3.1738	R.MKEYGEQIDPSTHRK.N	1	MSAP_HUMAN
Q9Y2B0	3	4.2642	R.MKEYGEQIDPSTHR.K	2	MSAP_HUMAN
Q9Y2B0	3	4.0161	R.INPDGSQSVVEVPYAR.S	3	MSAP_HUMAN
Q9Y2B0	2	2.8371	R.EADNVKDKLCSK.R	1	MSAP_HUMAN
Q9Y2B0	2	4.5504	R.ALVDELEWEIAQVDPKK.T	1	MSAP_HUMAN
Q9Y2B0	3	4.4933	R.ALVDELEWEIAQVDPK.K	2	MSAP_HUMAN
Q9Y2B0	1	2.2096	K.TIQMGSFR.I	2	MSAP_HUMAN
Q9Y2B0	2	2.8417	K.KTIQMGSFR.I	1	MSAP_HUMAN
Q9Y2B0	2	2.7135	K.EYGEQIDPSTHRK.N	1	MSAP_HUMAN
Q9Y2B0	2	2.9839	K.EYGEQIDPSTHR.K	1	MSAP_HUMAN
Q9Y2B0	3	3.8732	R.M#KEYGEQIDPSTHR.K	1	MSAP_HUMAN
Misshapen-like kinase 1 - Homo sapiens (Human)					
Q8N4C8	2	3.352	R.SQSLQDQPTR.N	2	MINK1_HUMAN
Mitochondrial 28S ribosomal protein S36 - Homo sapiens (Human)					
P82909	2	3.9912	R.KLVSQEEM#EFIQR.G	1	RT36_HUMAN
P82909	2	2.7011	R.DNPKPNVSEALR.S	1	RT36_HUMAN
P82909	2	5.2031	R.KLVSQEEMEFIQR.G	8	RT36_HUMAN
P82909	2	3.7873	R.SAGLPSHSSVISQHSK.G	1	RT36_HUMAN
P82909	2	4.9582	K.SPDLLM#YQGPPDTAEIIK.T	6	RT36_HUMAN
P82909	2	3.9977	K.LVSQEEMEFIQR.G	5	RT36_HUMAN
P82909	2	2.8733	K.LVSQEEM#EFIQR.G	1	RT36_HUMAN
P82909	3	5.2026	K.GSKSPDLLMYQGPPDTAEIIK.T	1	RT36_HUMAN
P82909	3	3.9767	R.RDNPKPNVSEALR.S	1	RT36_HUMAN
P82909	3	4.0108	K.SPDLLMYQGPPDTAEIIK.T	2	RT36_HUMAN
Mitochondrial 39S ribosomal protein L49 - Homo sapiens (Human)					
Q13405	2	3.6666	R.FVESVDEYQFVER.L	1	RM49_HUMAN
Mitochondrial antiviral-signaling protein - Homo sapiens (Human)					
Q7Z434	2	3.8285	K.VPANPASVSTVPSK.L	1	MAVS_HUMAN
Q7Z434	2	4.1906	R.GPVSPSVSFQPLAR.S	2	MAVS_HUMAN
Q7Z434	3	6.5806	R.EKEPSYPMPVQETQAPESPGENSEQALQTL	2	MAVS_HUMAN
Q7Z434	2	2.711	K.VSASTVPTDGSSR.N	1	MAVS_HUMAN
Q7Z434	3	6.3116	R.ASRLPGPTGSVVSTGTSFSSSSPGLASAGAA	2	MAVS_HUMAN
Mitochondrial import inner membrane translocase subunit Tim10 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P62072	2	3.8886	K.LTELSMQDEELM#KR.V	3	TIM10_HUMAN
P62072	2	2.8861	K.YLDIHER.M	1	TIM10_HUMAN
P62072	2	4.1525	K.LTELSMQDEELMKR.V	3	TIM10_HUMAN
P62072	2	3.4864	K.EAELSKGESVCLDR.C	1	TIM10_HUMAN
P62072	2	3.5452	K.LTELSMQDEELMK.R	3	TIM10_HUMAN
P62072	2	3.7349	K.LTELSMQDEELM#K.R	2	TIM10_HUMAN
P62072	3	4.7189	K.KLTELSMQDEELMKR.V	4	TIM10_HUMAN
P62072	3	4.9945	K.KLTELSMQDEELM#KR.V	2	TIM10_HUMAN
P62072	2	5.4114	K.KLTELSMQDEELMK.R	3	TIM10_HUMAN
Mitochondrial import inner membrane translocase subunit Tim13 - Homo sapiens (Human)					
Q9Y5L4	2	5.5036	K.VQIAVANAQELLQR.M	4	TIM13_HUMAN
Q9Y5L4	2	3.0386	R.YMDAWNTVSR.A	1	TIM13_HUMAN
Mitochondrial import inner membrane translocase subunit Tim8 A - Homo sapiens (Human)					
O60220	2	3.849	K.SKPVFSESLSD.-	1	TIM8A_HUMAN
O60220	2	4.0873	R.FIDTSQFILNR.L	4	TIM8A_HUMAN
O60220	2	4.0553	R.FIDTSQFILNRLEQTQK.S	1	TIM8A_HUMAN
Mitochondrial import inner membrane translocase subunit Tim8 B - Homo sapiens (Human)					
Q9Y5J9	2	3.0296	R.FAQIVQK.G	1	TIM8B_HUMAN
Mitochondrial import inner membrane translocase subunit Tim9 - Homo sapiens (Human)					
Q9Y5J7	3	4.1293	R.FQEYHIQQNEALAAK.A	2	TIM9_HUMAN
Mitochondrial import receptor subunit TOM34 - Homo sapiens (Human)					
Q15785	2	3.5355	K.NRVPSAGDVEK.A	2	OM34_HUMAN
Mitochondrial intermembrane space import and assembly protein 40 - Homo sapiens (Human)					
Q8N4Q1	3	3.9358	K.KPAEQAEETAPIEATATKEEEGSS.-	1	MIA40_HUMAN
Q8N4Q1	1	2.3735	R.IIFVTK.E	2	MIA40_HUMAN
Mitogen-activated protein kinase 1 - Homo sapiens (Human)					
P28482	3	3.9128	K.LKELIFEETAR.F	1	MK01_HUMAN
Mitogen-activated protein kinase kinase kinase 7-interacting protein 2 - Homo sapiens (Human)					
Q9NYJ8	2	3.972	R.TSSTSSSVNSQTLNR.N	1	TAB2_HUMAN
Mitogen-activated protein kinase kinase kinase MLT - Homo sapiens (Human)					
Q9NYL2	2	3.9812	R.DSGFSSGNTDTSSER.G	2	MLTK_HUMAN
MKI67 FHA domain-interacting nucleolar phosphoprotein - Homo sapiens (Human)					
Q9BYG3	2	2.8396	K.VSGTLDTPEK.T	1	MK67I_HUMAN
Q9BYG3	2	4.1201	K.GIDYDFPSLILQK.T	3	MK67I_HUMAN
MKL/myocardin-like protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q969V6	2	4.3832	K.EAIIVGQVNYPK.V	3	MKL1_HUMAN
MKL/myocardin-like protein 2 - Homo sapiens (Human)					
Q9ULH7	2	2.8676	K.NSNSGNSALNNATPNTPR.Q	1	MKL2_HUMAN
Moesin - Homo sapiens (Human)					
P26038	2	5.9498	R.DQKKTQEQLALEMAELTAR.I	6	MOES_HUMAN
P26038	3	4.4753	K.TQEQLALEMAELTAR.I	3	MOES_HUMAN
P26038	1	3.0475	R.ALELEQER.K	5	MOES_HUMAN
P26038	2	3.6268	K.ALTSELANAR.D	3	MOES_HUMAN
P26038	2	3.0624	R.AMLENEKK.K	3	MOES_HUMAN
P26038	2	4.4984	K.TANDMIHAENMR.L	3	MOES_HUMAN
P26038	2	3.3718	R.AQSEAEKLAK.E	4	MOES_HUMAN
P26038	3	4.1803	R.DQKKTQEQLALEM#AELTAR.I	1	MOES_HUMAN
P26038	2	3.2935	R.ISQLEM#AR.Q	1	MOES_HUMAN
P26038	2	3.0472	R.ISQLEMAR.Q	1	MOES_HUMAN
P26038	2	2.7985	R.LKQIEEQTK.K	1	MOES_HUMAN
P26038	2	3.2795	R.LKQIEEQTKK.A	3	MOES_HUMAN
P26038	2	3.303	R.RALELEQER.K	2	MOES_HUMAN
P26038	3	3.9566	R.RALELEQERK.R	1	MOES_HUMAN
P26038	3	4.2866	R.SEEERTTEAEKNER.V	4	MOES_HUMAN
P26038	2	3.3123	K.TANDM#IHAENMR.L	1	MOES_HUMAN
P26038	2	2.7158	R.QEAEEAKEALLQASR.D	1	MOES_HUMAN
P26038	2	4.2533	K.AQQELEEQTR.R	4	MOES_HUMAN
P26038	2	4.1533	K.ALTSELANARDESK.K	2	MOES_HUMAN
P26038	2	3.671	K.AQM#VQEDLEK.T	2	MOES_HUMAN
P26038	2	4.5176	K.ALTSELANARDESKK.T	2	MOES_HUMAN
P26038	1	3.7519	K.AQMVQEDLEK.T	3	MOES_HUMAN
P26038	3	8.0435	K.TAMSTPHVAEPAENEQDEQDENGAEASADL	3	MOES_HUMAN
P26038	3	5.4986	K.ERQEAEEAKEALLQASR.D	8	MOES_HUMAN
P26038	1	3.3547	K.ESEAVEWQQK.A	5	MOES_HUMAN
P26038	3	6.2851	K.TAM#STPHVAEPAENEQDEQDENGAEASAD	4	MOES_HUMAN
P26038	2	3.1025	K.KAQQELEEQTRR.A	2	MOES_HUMAN
P26038	2	3.642	K.KTANDM#IHAENMR.L	2	MOES_HUMAN
P26038	2	3.6417	K.KTANDMIHAENM#R.L	2	MOES_HUMAN
P26038	2	4.6101	K.KTANDMIHAENMR.L	3	MOES_HUMAN
P26038	2	3.3522	K.KTQEQLALEM#AELTAR.I	2	MOES_HUMAN
P26038	2	2.947	K.QRIDEFESM#.-	2	MOES_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P26038	2	4.2251	K.KAQQELEEQTR.R	2	MOES_HUMAN
P26038	2	3.4901	K.AQM#VQEDLEKTR.A	1	MOES_HUMAN
Mono-ADP-ribosyltransferase sirtuin-6 - Homo sapiens (Human)					
Q8N6T7	2	2.8961	K.EESPTRINGSIPAGPK.Q	1	SIRT6_HUMAN
Monocarboxylate transporter 1 - Homo sapiens (Human)					
P53985	2	3.1589	K.AAESPDQKDTEGGPKEEESPV.-	1	MOT1_HUMAN
P53985	2	2.8047	K.DTEGGPKEEESPV.-	1	MOT1_HUMAN
Monocarboxylate transporter 4 - Homo sapiens (Human)					
O15427	2	2.7664	K.NGEVVHTPETSV.-	1	MOT4_HUMAN
MORC family CW-type zinc finger protein 3 - Homo sapiens (Human)					
Q14149	3	4.8817	R.INAELLFRPTALSTPSFSSPK.E	2	MORC3_HUMAN
MORF4 family-associated protein 1 - Homo sapiens (Human)					
Q9Y605	2	3.2871	K.MAEMLVELVR.R	1	Q9Y605_HUMAN
Q9Y605	3	3.912	K.TQVEASEESALNHLQNPGDAAEGR.A	1	Q9Y605_HUMAN
Mortality factor 4-like protein 1 - Homo sapiens (Human)					
Q9UBU8	2	2.7877	K.ANQEQYAEGK.M	1	MO4L1_HUMAN
Mothers against decapentaplegic homolog 2 - Homo sapiens (Human)					
Q15796	2	3.7634	K.SAGGSGGAGGGEQNGQEEK.W	1	SMAD2_HUMAN
M-phase phosphoprotein 6 - Homo sapiens (Human)					
Q99547	3	4.0885	R.RYETLVGTIGKK.F	1	MPH6_HUMAN
M-phase phosphoprotein 8 - Homo sapiens (Human)					
Q99549	3	4.6477	K.FVESQVESESSVLNDSPFPEDDSEGLHSDSR	1	MPP8_HUMAN
Q99549	2	4.5402	K.SKPDLESSLESLVFDLR.T	2	MPP8_HUMAN
MRG-binding protein - Homo sapiens (Human)					
Q9NV56	2	4.0473	K.VLTANSNPSSPSAAK.R	2	MRGBP_HUMAN
MSTP116 - Homo sapiens (Human)					
Q7Z2S7	3	3.7696	R.SLLKEEDPAVLISEVLR.R	1	Q7Z2S7_HUMAN
Msx2-interacting protein - Homo sapiens (Human)					
Q96T58	3	4.7115	K.VDATRPEATTEVGPQIGVK.E	2	MINT_HUMAN
Multidrug resistance-associated protein 9 - Homo sapiens (Human)					
Q96J65	2	3.0039	R.EEDAGIIVLAPGNEKDEGK.E	1	MRP9_HUMAN
Multimerin-2 precursor - Homo sapiens (Human)					
Q9H8L6	4	5.2282	R.QLDGSSLQALQNAVDAVSLAVDAHKAEGER.	2	MMRN2_HUMAN
Q9H8L6	2	4.333	R.REEELQYTLEDMR.A	3	MMRN2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9H8L6	1	2.1497	R.VILMEK.S	1	MMRN2_HUMAN
Q9H8L6	2	5.3763	R.VALQDAASGLQEQALGWDELAAR.V	2	MMRN2_HUMAN
Q9H8L6	2	3.3798	R.VADSLPGLWK.A	2	MMRN2_HUMAN
Q9H8L6	3	4.3947	R.SQVQALDDEVGALKAAAAEAR.H	1	MMRN2_HUMAN
Q9H8L6	2	5.0389	R.SQVQALDDEVGALK.A	2	MMRN2_HUMAN
Q9H8L6	2	4.1198	R.SISELQADVDTKLK.R	2	MMRN2_HUMAN
Q9H8L6	1	3.7273	R.SISELQADVDTK.L	4	MMRN2_HUMAN
Q9H8L6	2	2.7717	R.VQDSAVAR.A	1	MMRN2_HUMAN
Q9H8L6	2	3.0417	R.ALEETQVSLDERR.Q	2	MMRN2_HUMAN
Q9H8L6	2	4.2343	K.AEAEDTSKDPVGR.N	2	MMRN2_HUMAN
Q9H8L6	2	2.8115	R.REEELQYTLEDM#R.A	1	MMRN2_HUMAN
Q9H8L6	2	3.4253	R.ADFQELGAK.F	6	MMRN2_HUMAN
Q9H8L6	2	2.9957	R.ELQSLSNDVK.N	1	MMRN2_HUMAN
Q9H8L6	3	5.4287	R.HVDEIKELYSESDETFDQISK.V	2	MMRN2_HUMAN
Q9H8L6	2	5.3524	R.LRSQVQALDDEVGALK.A	3	MMRN2_HUMAN
Q9H8L6	3	5.832	R.QLDGSSLQALQNAVDAVSLAVDAHK.A	1	MMRN2_HUMAN
Q9H8L6	3	4.1743	R.QLHSAFAALLEDALR.H	1	MMRN2_HUMAN
Q9H8L6	2	2.9092	K.LYLDLDVIR.E	1	MMRN2_HUMAN
Multiple coagulation factor deficiency protein 2 precursor - Homo sapiens (Human)					
Q8NI22	4	5.1874	K.NTVHDQEHIMEHLEGVINKPEAEM#SPQELQL	1	MCFD2_HUMAN
Q8NI22	4	6.765	K.NTVHDQEHIMEHLEGVINKPEAEMSPQELQL	3	MCFD2_HUMAN
Multiple myeloma tumor-associated protein 2 - Homo sapiens (Human)					
Q9BU76	2	4.1655	R.VESGGPGTSAASAR.R	2	MMTA2_HUMAN
Muted protein homolog - Homo sapiens (Human)					
Q8TDH9	3	4.1386	K.IHSDHLVASEK.Q	1	MUTED_HUMAN
Myc box-dependent-interacting protein 1 - Homo sapiens (Human)					
O00499	2	3.1102	R.KGPPVPPPPK.H	2	BIN1_HUMAN
O00499	2	2.7351	R.LDLPPGFMFK.V	2	BIN1_HUMAN
O00499	2	4.9635	R.VNHEPEPAGGATPGATLPK.S	2	BIN1_HUMAN
O00499	2	5.8901	K.VQAQHDYTATDTDELQLK.A	12	BIN1_HUMAN
O00499	2	4.3442	K.SPSPPDGSPAATPEIR.V	1	BIN1_HUMAN
O00499	3	3.8675	K.LVDQALLTMDTYLGQFPDIK.S	1	BIN1_HUMAN
O00499	2	3.0026	R.GVFPENFTERVP.-	2	BIN1_HUMAN
Myc target protein 1. - Homo sapiens (Human)					
Q8N699	2	3.4033	R.SNLSLASLTFQR.Q	1	Q8N699_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Myelin basic protein - Homo sapiens (Human)					
P02686	1	2.252	R.NLGELSR.T	1	MBP_HUMAN
Myelin expression factor 2 - Homo sapiens (Human)					
Q9P2K5	2	2.9318	R.REPHPAEAEK.Q	1	MYEF2_HUMAN
Myeloid cell nuclear differentiation antigen - Homo sapiens (Human)					
P41218	2	4.5615	K.INQEEVGLAAPAPTAR.N	4	MNDA_HUMAN
P41218	2	5.3405	K.KINQEEVGLAAPAPTAR.N	3	MNDA_HUMAN
MYL6 protein - Homo sapiens (Human)					
Q6IBG5	3	4.5228	K.MTEEEVEM#LVAGHEDSNGCINYEELVR.M	1	Q6IBG5_HUMAN
Myocyte-specific enhancer factor 2C - Homo sapiens (Human)					
Q06413	2	3.0356	R.NSPGLLVSPGNLNK.N	2	MEF2C_HUMAN
MyoD family inhibitor domain-containing protein - Homo sapiens (Human)					
Q9P1T7	2	6.2413	R.VAEAGGGQLGSTAQGK.C	3	MDFIC_HUMAN
Myosin light chain kinase, smooth muscle - Homo sapiens (Human)					
Q15746	3	4.8745	K.KKLPAENGSSSAETLNAK.A	2	MYLK_HUMAN
Q15746	2	4.1949	K.SSLPPVLGTESDATVK.K	9	MYLK_HUMAN
Q15746	3	3.7737	K.TLSEDDLKEIPAEQMDFR.A	1	MYLK_HUMAN
Q15746	2	3.0171	K.VHSPQQVDFR.S	1	MYLK_HUMAN
Q15746	2	4.1667	R.APGLGVLSPSGEER.K	2	MYLK_HUMAN
Q15746	2	3.5602	R.LSSMAM#ISGLSGR.K	1	MYLK_HUMAN
Q15746	2	4.4699	R.LSSMAMISGLSGR.K	3	MYLK_HUMAN
Myosin light polypeptide 4 - Homo sapiens (Human)					
P12829	2	3.2058	R.ALGQNPTNAEVLR.V	1	MYL4_HUMAN
P12829	2	4.1258	R.NKEQGTYEDFVEGLR.V	2	MYL4_HUMAN
P12829	3	4.0776	K.PAPAPAPAPAPAPAPAPEAPKEPAFDPK.S	2	MYL4_HUMAN
Myosin light polypeptide 6 - Homo sapiens (Human)					
P60660	2	3.6472	K.EAFQLFDRTGDGK.I	2	MYL6_HUMAN
P60660	3	4.1755	K.MTEEEVEM#LVAGHEDSNGCINYEAFVR.H	1	MYL6_HUMAN
P60660	2	3.0677	R.TGDGKILYSQCGDVMR.A	1	MYL6_HUMAN
P60660	2	2.7169	K.EAFQLFDR.T	1	MYL6_HUMAN
Myosin light polypeptide 6B - Homo sapiens (Human)					
P14649	2	4.9338	R.RVDFETFLPMLQAVAK.N	1	MYL6B_HUMAN
P14649	3	4.3095	K.NRGQGTYEDYLEGFR.V	5	MYL6B_HUMAN
P14649	2	2.7281	K.PAAAGAPPAK.T	1	MYL6B_HUMAN
P14649	2	3.9298	K.TKAEPAVPQAPQK.T	2	MYL6B_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P14649	2	3.1123	R.GQGTYEDYLEGFR.V	1	MYL6B_HUMAN
P14649	2	2.7446	R.HVLTTLGEK.M	1	MYL6B_HUMAN
Myosin phosphatase Rho-interacting protein - Homo sapiens (Human)					
Q6WCQ1	2	3.3844	R.SKSVIEQVSWDT.-	1	MRIP_HUMAN
Q6WCQ1	2	2.7116	R.FGM#LDATDGPGTEDAALR.M	1	MRIP_HUMAN
Q6WCQ1	3	3.8037	K.HVHPTTAPDVTSSLPEEK.N	2	MRIP_HUMAN
Q6WCQ1	3	4.4218	K.TFDWAEFRPIQQALAQER.V	1	MRIP_HUMAN
Myosin-10 - Homo sapiens (Human)					
P35580	3	4.0156	R.ALEQQVEEMRTQLEELEDELQATEDAK.L	1	MYH10_HUMAN
P35580	3	5.3033	R.HATALEELSEQLEQAKR.F	2	MYH10_HUMAN
P35580	2	3.0888	R.QLEEAEEEATR.A	2	MYH10_HUMAN
P35580	2	3.232	R.GGPISFSSSR.S	2	MYH10_HUMAN
P35580	2	4.3991	R.ELDDATEANEGLSR.E	10	MYH10_HUMAN
P35580	2	3.8045	R.DLQTRDEQNEEK.K	2	MYH10_HUMAN
P35580	2	2.8023	R.ASRDEIFAQSK.E	1	MYH10_HUMAN
P35580	2	3.601	K.LQNELDNVSTLLEEAEK.K	1	MYH10_HUMAN
P35580	3	4.3803	K.KLDAQVQELHAK.V	2	MYH10_HUMAN
P35580	3	3.9019	K.IGQLEEQLEQEAKER.A	3	MYH10_HUMAN
P35580	2	5.0266	K.IGQLEEQLEQEAK.E	2	MYH10_HUMAN
P35580	3	5.4733	K.NRLQQELDDLTVDLDHQR.Q	1	MYH10_HUMAN
P35580	2	3.8664	K.DAASLESQLQDTQELLQEETR.Q	2	MYH10_HUMAN
P35580	2	3.3987	K.AKLQELEGAVK.S	1	MYH10_HUMAN
P35580	2	3.5114	R.RGGPISFSSSR.S	2	MYH10_HUMAN
Myosin-14 - Homo sapiens (Human)					
Q7Z406	2	3.038	R.QEEEAGALEAGEEARR.R	1	MYH14_HUMAN
Myosin-6 - Homo sapiens (Human)					
P13533	2	3.0614	K.KALQEAHQQALDDLQVEEDK.V	3	MYH6_HUMAN
Myosin-9 - Homo sapiens (Human)					
P35579	3	3.892	K.REQEVNILKK.T	2	MYH9_HUMAN
P35579	2	4.2916	K.VLLQGKGDSEHKR.K	4	MYH9_HUMAN
P35579	1	2.2154	K.VLLQGK.G	4	MYH9_HUMAN
P35579	2	4.0922	K.VEAQLQELQVK.F	1	MYH9_HUMAN
P35579	1	2.8999	K.VAAYDKLEK.T	2	MYH9_HUMAN
P35579	3	6.9569	K.TRLQQELDDLLVDLDHQR.Q	3	MYH9_HUMAN
P35579	3	5.6134	K.TLEEEAKTHEAQIQEMR.Q	6	MYH9_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P35579	2	3.9459	K.TLEEEAKTHEAQIQEM#R.Q	2	MYH9_HUMAN
P35579	4	5.5793	R.DLGEELEALKTELEDTLDSTAAQQELR.S	2	MYH9_HUMAN
P35579	2	3.4915	K.THEAQIQEM#R.Q	1	MYH9_HUMAN
P35579	2	5.3429	R.KKVEAQLQELQVK.F	4	MYH9_HUMAN
P35579	2	3.3293	K.THEAQIQEMR.Q	2	MYH9_HUMAN
P35579	2	4.1002	R.ELESQISELQEDLESER.A	2	MYH9_HUMAN
P35579	1	2.2691	R.EQEVNILK.K	2	MYH9_HUMAN
P35579	2	3.4273	R.EQLEEEEEAKHNLEK.Q	1	MYH9_HUMAN
P35579	2	5.1785	R.IAEFTTNLTEEEEKSK.S	3	MYH9_HUMAN
P35579	3	5.0749	K.NKHEAMITDLEER.L	4	MYH9_HUMAN
P35579	3	4.0685	R.LQQELDDLLVDLDHQR.Q	2	MYH9_HUMAN
P35579	3	4.2559	R.QKHSQAVEELAEQLEQTK.R	2	MYH9_HUMAN
P35579	2	3.4528	R.QKHSQAVEELAEQLEQTKR.V	2	MYH9_HUMAN
P35579	3	4.319	R.RKLEGDSTDLSDQIAELQAQIAELK.M	1	MYH9_HUMAN
P35579	1	3.2099	R.TELADKVTK.L	5	MYH9_HUMAN
P35579	2	3.0908	R.VEEEAAQK.N	2	MYH9_HUMAN
P35579	2	3.2379	R.IAEFTTNLTEEEEK.S	2	MYH9_HUMAN
P35579	2	5.7831	K.IRELESQISELQEDLESER.A	5	MYH9_HUMAN
P35579	2	5.0552	K.AKQTLENERGELANEVK.V	3	MYH9_HUMAN
P35579	3	5.683	K.DFSALESQLQDTQELLQEENR.Q	2	MYH9_HUMAN
P35579	2	4.5139	K.DFSALESQLQDTQELLQEENRQK.L	1	MYH9_HUMAN
P35579	2	3.0345	K.DLEGLSQR.H	2	MYH9_HUMAN
P35579	2	3.2333	K.DLEGLSQRHEEK.V	3	MYH9_HUMAN
P35579	1	2.2493	K.FNEGER.V	3	MYH9_HUMAN
P35579	2	3.9297	K.HEAMITDLEER.L	3	MYH9_HUMAN
P35579	2	3.2161	K.QIATLHAQVADMKK.K	2	MYH9_HUMAN
P35579	3	5.641	K.HSQAVEELAEQLEQTKR.V	5	MYH9_HUMAN
P35579	2	3.3382	K.QTLENERGELANEVK.V	2	MYH9_HUMAN
P35579	3	4.9523	K.KEEELQAALARVEEEAAQK.N	2	MYH9_HUMAN
P35579	3	4.9395	K.LTKDFSALESQLQDTQELLQEENRQK.L	1	MYH9_HUMAN
P35579	2	3.6121	K.QIATLHAQVADMK.K	3	MYH9_HUMAN
P35579	2	5.8314	K.HSQAVEELAEQLEQTK.R	3	MYH9_HUMAN
P35579	3	4.0624	K.NKHEAM#ITDLEER.L	3	MYH9_HUMAN
P35579	2	4.6106	K.KLEEEQIILEDQNCK.L	4	MYH9_HUMAN
P35579	3	7.5245	K.LTKDFSALESQLQDTQELLQEENR.Q	3	MYH9_HUMAN
P35579	2	5.9252	K.LQVELDNVTGLLSQSDSK.S	7	MYH9_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P35579	3	6.4594	K.LLEDRIAEFTTNLTEEEEKSK.S	2	MYH9_HUMAN
P35579	3	4.6443	K.LKNKHEAMITDLEER.L	4	MYH9_HUMAN
P35579	2	5.3882	K.KVEAQLQELQVK.F	5	MYH9_HUMAN
P35579	2	5.9619	K.KTLEEEAKTHEAQIQEMR.Q	5	MYH9_HUMAN
P35579	3	5.2244	K.KMEDSVGCLETAEEVKR.K	4	MYH9_HUMAN
Myosin-Ie - Homo sapiens (Human)					
Q12965	2	3.8894	K.VLQVSIGPGLPK.N	2	MYO1E_HUMAN
Q12965	2	4.0885	R.NTTQNTGYSSGTQNANYPVR.A	3	MYO1E_HUMAN
Myosin-If - Homo sapiens (Human)					
O00160	3	4.1693	R.ARPPSEHNTEFLNVPDQGMAGMQR.K	1	MYO1F_HUMAN
O00160	2	3.9013	R.GPPSTSLGASR.R	2	MYO1F_HUMAN
Myosin-IXA - Homo sapiens (Human)					
Q9UNJ2	2	3.4014	R.ELEQAIFSLELLK.V	1	Q9UNJ2_HUMA
Myosin-Va - Homo sapiens (Human)					
Q9Y4I1	2	2.9645	K.QETEQLVSNLKEENTLLK.Q	1	MYO5A_HUMAN
Myosin-XVB - Homo sapiens (Human)					
Q96JP2	2	3.3297	K.EAEAEPAKETAAK.G	4	MO15B_HUMAN
Q96JP2	3	6.7626	K.VHIPQGEAQEEEEEEEEEEQEEQEVETR.A	2	MO15B_HUMAN
Myosin-XVIIIa - Homo sapiens (Human)					
Q92614	2	2.9546	K.EMESRDEEVEEAR.Q	1	MY18A_HUMAN
Q92614	3	4.267	K.VKDQEEELDEQAGTIQMLEQAK.L	1	MY18A_HUMAN
Q92614	3	5.4012	R.RFDSELSQAHEEAQR.E	2	MY18A_HUMAN
Myotrophin - Homo sapiens (Human)					
P58546	3	4.0753	K.GADINAPDKHHITPLLSAVYEGHVSCVK.L	1	MTPN_HUMAN
P58546	3	5.6008	K.GADKTVKGPDGLTAFEATDNQAIK.A	1	MTPN_HUMAN
P58546	2	5.508	K.GPDGLTAFEATDNQAIK.A	14	MTPN_HUMAN
P58546	2	4.2464	K.NGDLDEVKDYVAK.G	4	MTPN_HUMAN
P58546	2	5.1479	K.NGDLDEVKDYVAKGEDVNR.T	6	MTPN_HUMAN
P58546	2	5.7346	K.TVKGPDGLTAFEATDNQAIK.A	7	MTPN_HUMAN
Myristoylated alanine-rich C-kinase substrate - Homo sapiens (Human)					
P29966	1	2.1761	K.LSGFSFK.K	2	MARCS_HUMAN
P29966	3	4.5859	K.TAAKGEAAAERPGEAAVASSPSK.A	1	MARCS_HUMAN
P29966	2	5.4504	K.GEPAAAAAPEAGASPVEK.E	3	MARCS_HUMAN
P29966	3	4.7172	K.EELQANGSAPAADKEEPAAAGSGAASPSAA	1	MARCS_HUMAN
P29966	2	5.3038	K.EAGEGGEAEAPAAEGGK.D	6	MARCS_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P29966	2	3.5516	K.AEDGATPSPSNETPKK.K	1	MARCS_HUMAN
P29966	2	3.2868	K.AAEEPSKVEEK.K	3	MARCS_HUMAN
P29966	2	3.9403	K.AEDGATPSPSNETPK.K	12	MARCS_HUMAN
P29966	2	4.7779	K.EAPAEGEAAEPGSPTAAEGEAASAASSTSSP	5	MARCS_HUMAN
N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 - Homo sapiens (Human)					
O94760	2	3.5625	R.ALPESLGQHALR.S	3	DDAH1_HUMAN
O94760	3	4.6836	R.QHQLYVGVLGSK.L	1	DDAH1_HUMAN
O94760	2	3.7738	R.SAKGEEVDVAR.A	6	DDAH1_HUMAN
O94760	2	3.2909	R.TPEEYPESAK.V	4	DDAH1_HUMAN
O94760	3	3.7907	R.TPEEYPESAKVYEK.L	1	DDAH1_HUMAN
N(G),N(G)-dimethylarginine dimethylaminohydrolase 2 - Homo sapiens (Human)					
O95865	3	4.0413	K.AGAGLSSLCLVLSTRPHS.-	1	DDAH2_HUMAN
N-acetylglucosamine-6-sulfatase precursor - Homo sapiens (Human)					
P15586	2	2.8574	K.IQEPNTFPAILR.S	1	GNS_HUMAN
P15586	2	2.9353	R.WQTLLSVDDLVEK.L	1	GNS_HUMAN
N-acetylmuramoyl-L-alanine amidase precursor - Homo sapiens (Human)					
Q96PD5	2	4.3789	R.DGSPDVTTADIGANTPDATK.G	2	PGRP2_HUMAN
N-acetyltransferase 12 - Homo sapiens (Human)					
Q147X3	2	4.2389	R.SPAGGESATVAAK.G	3	Q147X3_HUMAN
N-acylneuraminate cytidylyltransferase - Homo sapiens (Human)					
Q8NFW8	1	3.2501	K.GAATSVSNPR.G	3	NEUA_HUMAN
NADH dehydrogenase - Homo sapiens (Human)					
Q8WU60	2	5.3549	R.GKLLATQTAAELSK.N	3	Q8WU60_HUMA
Q8WU60	2	4.4929	R.KVASPSPSGSVLFTDEGVPK.F	2	Q8WU60_HUMA
Q8WU60	2	3.4174	K.LLATQTAAELSK.N	2	Q8WU60_HUMA
Q8WU60	2	3.1597	R.KTLVEFPQK.V	3	Q8WU60_HUMA
Q8WU60	3	4.2115	R.LNEIDKESQKPFEVK.G	1	Q8WU60_HUMA
Q8WU60	3	4.9883	R.KETSGTQGIEGHLK.G	6	Q8WU60_HUMA
Q8WU60	2	3.5624	R.GGLRKPEASHSFENR.A	2	Q8WU60_HUMA
Q8WU60	3	4.0485	R.AEGQLQASPPGAAEGHLEKPVPEPQR.K	2	Q8WU60_HUMA
Q8WU60	2	4.6173	K.NLSSPSSYPPAVNK.G	4	Q8WU60_HUMA
Q8WU60	2	3.6813	K.GSPAPAVLAEEAR.A	5	Q8WU60_HUMA
Q8WU60	2	4.9225	R.GGTQEPAPVPAEPFDNTTYK.N	1	Q8WU60_HUMA
Q8WU60	3	4.0901	K.TLLQKPHVDITDPEKPHQPK.K	2	Q8WU60_HUMA
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43678	2	3.4706	R.ALENVLSGKA.-	3	NDUA2_HUMAN
O43678	2	3.5124	R.YAFGQETNVPLNNFSADQVTR.A	1	NDUA2_HUMAN
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 3 - Homo sapiens (Human)					
O95167	1	2.2202	K.YSVM#INK.A	1	NDUA3_HUMAN
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 - Homo sapiens (Human)					
O00483	2	3.3481	K.FYSVNVDYSK.L	2	NDUA4_HUMAN
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 - Homo sapiens (Human)					
O95182	2	2.9481	K.PAESSAVAATEK.K	1	NDUA7_HUMAN
O95182	2	5.7817	K.ALVSGKPAESSAVAATEK.K	3	NDUA7_HUMAN
O95182	3	5.8704	K.ALVSGKPAESSAVAATEKK.A	4	NDUA7_HUMAN
O95182	2	2.9471	K.KAVTPAPPIKR.W	2	NDUA7_HUMAN
NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial precursor - Homo sapiens (Hum					
P19404	2	3.3809	R.QNGWLPISAMNK.V	2	NDUV2_HUMAN
P19404	2	3.8116	K.AAAVLPVLDLAQR.Q	4	NDUV2_HUMAN
P19404	2	2.8066	R.DTPENNPDTPFDFTPENYKR.I	1	NDUV2_HUMAN
NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial precursor - Homo sapiens					
O43181	2	3.0815	K.NGWSYDIEER.K	2	NDUS4_HUMAN
O43181	2	2.8991	K.SYGANFSWNK.R	1	NDUS4_HUMAN
O43181	2	2.7786	R.NNMQSGVNNTK.K	1	NDUS4_HUMAN
O43181	2	3.7218	R.NNMQSGVNNTKK.W	2	NDUS4_HUMAN
NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial precursor - Homo sapiens					
O75380	2	3.2982	K.VTHTGQVYDDKDYRR.I	2	NDUS6_HUMAN
O75380	2	3.7883	K.VYINLDKETK.T	3	NDUS6_HUMAN
O75380	3	4.0412	R.QKEVNENFAIDLIAEQPVSEVETR.V	2	NDUS6_HUMAN
NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial precursor - Homo sapiens					
O00217	2	2.8939	R.RYPSGEER.C	1	NDUS8_HUMAN
NADPH oxidase 5 - Homo sapiens (Human)					
Q96PH1	2	2.7164	K.NDMKAIGLQMALDLLANK.E	1	NOX5_HUMAN
Nance-Horan syndrome protein - Homo sapiens (Human)					
Q6T4R5	2	2.7263	R.ATTPSLPSVDNEFK.L	1	NHS_HUMAN
Q6T4R5	2	4.4264	K.IIQYGPGPDETLEQVQK.A	1	NHS_HUMAN
Q6T4R5	2	4.6146	K.SPESSESQTSQSESR.A	3	NHS_HUMAN
Nanos homolog 3 - Homo sapiens (Human)					
P60323	2	2.7396	R.LSPQPEPEPMLEPDQK.R	1	NANO3_HUMAN
Nascent polypeptide-associated complex subunit alpha - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q13765	3	4.7186	K.SPASDTYIVFGEAKIEDLSQQAQLAAAEK.F	3	NACA_HUMAN
Q13765	2	5.3162	R.ALKNNSNDIVNAIMELTM#.-	10	NACA_HUMAN
Q13765	2	5.2891	R.ALKNNSNDIVNAIM#ELTM#.-	8	NACA_HUMAN
Q13765	2	4.8768	K.SPASDTYIVFGEAK.I	4	NACA_HUMAN
Q13765	2	4.063	K.NNSNDIVNAIMELTM#.-	3	NACA_HUMAN
Q13765	2	3.346	K.LGLRQVTGVTR.V	2	NACA_HUMAN
Q13765	2	3.0429	K.IEDLSQQAQLAAAEKFK.V	1	NACA_HUMAN
Q13765	3	6.1223	K.IEDLSQQAQLAAAEK.F	15	NACA_HUMAN
Q13765	2	3.8084	K.DIELVM#SQANVSR.A	1	NACA_HUMAN
Q13765	2	3.7455	R.ALKNNSNDIVNAIM#ELTM.-	3	NACA_HUMAN
NEDD4-like E3 ubiquitin-protein ligase WWP2 - Homo sapiens (Human)					
O00308	2	3.3663	R.DSSGTAVAPENR.H	3	WWP2_HUMAN
NEDD8 precursor - Homo sapiens (Human)					
Q15843	2	2.7709	R.IKERVEEK.E	1	NEDD8_HUMAN
Q15843	2	3.7198	K.EIEIDIEPTDKVER.I	3	NEDD8_HUMAN
Q15843	2	4.4879	K.ILGGSVLHLVLALR.G	1	NEDD8_HUMAN
Q15843	2	2.7074	K.QMNDEKTAADYK.I	1	NEDD8_HUMAN
Q15843	3	5.1279	K.TLTGKEIEIDIEPTDKVER.I	7	NEDD8_HUMAN
NEDD8 ultimate buster 1 - Homo sapiens (Human)					
Q9Y5A7	2	3.1494	R.KNFQLEEEEQNEAKLK.E	1	NUB1_HUMAN
Negative elongation factor E - Homo sapiens (Human)					
P18615	2	4.0915	K.SVWGSLAVQNSPK.G	5	NELFE_HUMAN
Neogenin precursor - Homo sapiens (Human)					
Q92859	2	3.2911	R.TFTPFYFLVEPVDTLSVR.G	2	NEO1_HUMAN
Nephronectin precursor - Homo sapiens (Human)					
Q6UXI9	2	2.7768	R.QPSNDLFEIFEIER.G	1	NPNT_HUMAN
Nesprin-2 - Homo sapiens (Human)					
Q8WXH0	2	2.9891	K.QLVASVSC*PELEGQIAK.L	1	SYNE2_HUMAN
Nestin - Homo sapiens (Human)					
P48681	2	3.5129	R.SPEEVDKESQR.N	4	NEST_HUMAN
P48681	2	4.1628	R.LLEEENQESLR.S	3	NEST_HUMAN
P48681	2	6.0867	R.NLEEEENLGKGEYQESLR.S	6	NEST_HUMAN
P48681	2	3.3948	R.SLEDENKEAFR.S	2	NEST_HUMAN
P48681	2	4.7467	R.SLEEEGQELPQSADVQR.W	7	NEST_HUMAN
P48681	2	3.9451	R.SLEEQDQETLR.T	4	NEST_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P48681	2	2.9892	R.SLEKENQEPLK.T	2	NEST_HUMAN
P48681	3	4.4479	R.SLGAWNLENLRSPEEVDKESQR.N	2	NEST_HUMAN
P48681	2	2.8699	R.TLEKETQQR.R	2	NEST_HUMAN
P48681	3	4.8261	K.SLDQEIARPLENENQEFLK.S	4	NEST_HUMAN
P48681	2	4.0358	R.GSPFQEEEGSALK.T	1	NEST_HUMAN
P48681	2	4.0927	R.SLGAWNLENLR.S	4	NEST_HUMAN
P48681	3	4.8026	K.ENQEPLRSPEVGDEEALRPLTK.E	1	NEST_HUMAN
P48681	4	5.1552	K.SQHVNGGVMNGLEQSEEVGQGMPLVSEGD	1	NEST_HUMAN
P48681	2	2.8113	R.EGDRESWSSGED.-	1	NEST_HUMAN
P48681	2	3.122	K.EAVGQLKPTGK.E	1	NEST_HUMAN
P48681	2	3.2286	K.EDTQTLQSLQK.E	2	NEST_HUMAN
P48681	2	3.8084	K.SAGQENLETLK.S	2	NEST_HUMAN
P48681	4	5.4827	K.SAGQENLETLKSPETQAPLWTPEEINQGAMN	1	NEST_HUMAN
P48681	2	3.2247	K.SLETEILESLK.S	1	NEST_HUMAN
P48681	2	2.9428	K.DVEVVRPLEK.E	2	NEST_HUMAN
P48681	2	4.425	K.TSWAGAPVHLGQGQFLK.F	2	NEST_HUMAN
P48681	3	6.271	R.EDKVPGPLEIPSGM#EDAGPGADIIGVNGQG	1	NEST_HUMAN
P48681	3	5.2124	K.EAVGQLKPTGKEDTQTLQSLQK.E	3	NEST_HUMAN
Netrin-4 precursor - Homo sapiens (Human)					
Q9HB63	2	2.9996	R.M#GNLALGR.K	1	NET4_HUMAN
Neudesin precursor - Homo sapiens (Human)					
Q9UMX5	2	4.2205	R.YGGEEEDQPIYLAVK.G	5	NENF_HUMAN
Q9UMX5	3	6.9323	K.MSLDPADLTHDTTGLTAKELEALDEVFTK.V	1	NENF_HUMAN
Q9UMX5	3	4.8069	R.GPPVRLFTEEELAR.Y	3	NENF_HUMAN
Q9UMX5	2	3.0939	R.GAPYNALTGKDSTR.G	1	NENF_HUMAN
Q9UMX5	2	2.825	K.YPIVGYTAR.R	4	NENF_HUMAN
Q9UMX5	3	4.9754	K.M#SLDPADLTHDTTGLTAKELEALDEVFTK.V	2	NENF_HUMAN
Q9UMX5	2	2.9259	K.M#SLDPADLTHDTTGLTAK.E	1	NENF_HUMAN
Q9UMX5	2	5.1658	K.GVVFDVTSGKEFYGR.G	4	NENF_HUMAN
Q9UMX5	2	4.0181	K.AKYPIVGYTAR.R	4	NENF_HUMAN
Q9UMX5	2	3.2278	K.ELEALDEVFTK.V	1	NENF_HUMAN
Neurabin-2 - Homo sapiens (Human)					
Q96SB3	2	3.0972	R.KLQSLEQEK.G	2	NEB2_HUMAN
Q96SB3	2	3.8248	R.SAYEAGIQALKPPDAPGPDEAPK.G	3	NEB2_HUMAN
Q96SB3	3	3.9506	R.FDSKPAPSAQPAPPPHPPSR.L	2	NEB2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96SB3	3	3.9681	R.FNGSTEALDKLDADAVSPTVSQLSAVFEK.A	1	NEB2_HUMAN
Q96SB3	3	5.4354	K.HAVTEAEIQQLKR.K	3	NEB2_HUMAN
Q96SB3	2	3.199	K.EAAAVAPPER.G	2	NEB2_HUMAN
Q96SB3	2	4.0388	K.AQLEQSVEENKER.M	2	NEB2_HUMAN
Q96SB3	2	4.6344	K.AQAAPEKEAAAVAPPER.G	1	NEB2_HUMAN
Q96SB3	2	2.8249	K.TVTEGGAAHR.D	4	NEB2_HUMAN
Q96SB3	2	3.356	R.VLEESELAR.K	2	NEB2_HUMAN
Neural Wiskott-Aldrich syndrome protein - Homo sapiens (Human)					
O00401	1	2.7533	K.VIYDFIEK.T	2	WASL_HUMAN
O00401	2	3.2653	R.DALLDQIR.Q	2	WASL_HUMAN
O00401	2	2.7489	K.AALLDQIR.E	2	WASL_HUMAN
O00401	2	3.9611	R.FYGPQVNNISHTK.E	1	WASL_HUMAN
Neuroblast differentiation-associated protein AHNAK - Homo sapiens (Human)					
Q09666	3	7.288	K.SKGHYEVTGSDDETGKLQGSGVSLASK.K	3	AHNK_HUMAN
Q09666	2	2.9205	K.MPSMNIQTHK.I	2	AHNK_HUMAN
Q09666	3	4.7485	K.MPTFSTPGAKGEGPDVHMTLPK.G	2	AHNK_HUMAN
Q09666	2	3.8069	K.MSLPDVDLDLKGPK.M	2	AHNK_HUMAN
Q09666	2	3.1439	K.MPSM#NIQTHK.I	1	AHNK_HUMAN
Q09666	3	5.5971	K.SAKM#DIDVPDVEVQGPDWHLK.M	2	AHNK_HUMAN
Q09666	2	2.751	K.TPEM#IIQKPK.I	1	AHNK_HUMAN
Q09666	2	3.5266	K.VDIDAPDVEVHDPDWHLK.M	3	AHNK_HUMAN
Q09666	2	3.2095	K.MPEMSIKPQK.I	1	AHNK_HUMAN
Q09666	3	4.5887	K.VDIKAPDVEGQGLDWSLK.I	4	AHNK_HUMAN
Q09666	2	4.1179	K.MDAEVPDVNIEGPDAK.L	7	AHNK_HUMAN
Q09666	3	4.9647	K.VDIKAPDVEGQGLDWSLKIPK.M	4	AHNK_HUMAN
Q09666	2	4.3145	K.VDIDVPDVNLEAPEGK.L	5	AHNK_HUMAN
Q09666	2	3.8724	K.M#KGNVDISAPK.I	3	AHNK_HUMAN
Q09666	3	4.2485	K.LKTDVDVSLPK.V	2	AHNK_HUMAN
Q09666	2	2.9086	K.LNIKAPK.V	1	AHNK_HUMAN
Q09666	2	4.7298	K.M#DAEVPDVNIEGPDAK.L	4	AHNK_HUMAN
Q09666	3	4.1106	K.M#DIDAPDVEVQGPDWHLK.M	1	AHNK_HUMAN
Q09666	3	3.7478	K.M#DIDVPDVEVQGPDWHLK.M	3	AHNK_HUMAN
Q09666	3	4.0409	K.MDIDVPDVEVQGPDWHLK.M	3	AHNK_HUMAN
Q09666	3	3.8342	K.M#KGDYDVTVPKVEGEIK.A	1	AHNK_HUMAN
Q09666	2	4.1734	K.MKGNVDISAPK.I	4	AHNK_HUMAN
Q09666	2	4.4713	K.M#SLPDVDLDLKGPK.M	4	AHNK_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q09666	3	4.0262	K.VEGEM#KVPDVDIKGPK.V	1	AHNK_HUMAN
Q09666	2	4.721	K.MDIDAPDVEVQGPDWHLK.M	3	AHNK_HUMAN
Q09666	2	4.1906	K.VDINAPDVEVHGPDWHLK.M	2	AHNK_HUMAN
Q09666	2	3.0014	K.MDVNVGDIDIEGPEGK.L	1	AHNK_HUMAN
Q09666	2	3.447	K.MKGDYDVTVPK.V	2	AHNK_HUMAN
Q09666	2	3.604	K.M#KGDYDVTVPK.V	3	AHNK_HUMAN
Q09666	2	4.377	K.VKGEYDVTVPK.L	6	AHNK_HUMAN
Q09666	2	5.0281	K.LKGPQITGPSLEGDLGLK.G	6	AHNK_HUMAN
Q09666	2	4.5543	R.VDIETPNLEGTLTGPR.L	1	AHNK_HUMAN
Q09666	3	6.9511	R.SNSFSDEREFSGPSTPTGTLEFEGGEVSLEG	1	AHNK_HUMAN
Q09666	2	5.1549	R.GGVQVPAVDISSSLGGR.P	5	AHNK_HUMAN
Q09666	3	5.1523	R.GPKVDVSAPDVEAHGPEWNLK.M	1	AHNK_HUMAN
Q09666	2	3.2272	R.VSAPEVSVGHK.G	4	AHNK_HUMAN
Q09666	1	2.5013	R.EVDVNLPK.A	2	AHNK_HUMAN
Q09666	2	3.2049	R.EFSGPSTPTGTLEFEGGEVSLEGGK.V	1	AHNK_HUMAN
Q09666	2	3.4684	K.VSMPDVDLNLKGPK.L	4	AHNK_HUMAN
Q09666	2	4.9297	K.VSAPGVQGDVKGPQVALK.G	4	AHNK_HUMAN
Q09666	2	3.2963	K.VSAPGVQGDVK.G	2	AHNK_HUMAN
Q09666	2	4.4744	K.VNVEAPDVNLEGLGGK.L	6	AHNK_HUMAN
Q09666	2	3.9495	K.VEGEIKAPDVDIK.G	2	AHNK_HUMAN
Q09666	2	4.0394	K.VKGGVDVTLPR.V	4	AHNK_HUMAN
Q09666	2	4.3386	K.VDISAPDVDVHGPDWHLK.M	3	AHNK_HUMAN
Q09666	2	4.0803	K.VKGEYDMTVPK.L	6	AHNK_HUMAN
Q09666	2	2.8518	K.VKGDVDVSVPK.V	1	AHNK_HUMAN
Q09666	3	3.7282	K.VKGDMDVSVPKVEGEMK.V	1	AHNK_HUMAN
Q09666	3	3.9803	K.VKGDMDVSVPK.V	5	AHNK_HUMAN
Q09666	2	4.6192	K.VGVEVPDVNIEGPEGK.L	9	AHNK_HUMAN
Q09666	1	4.8164	K.VGGSGVNVNAK.G	8	AHNK_HUMAN
Q09666	2	4.8274	K.VESEIKVPDVELK.S	1	AHNK_HUMAN
Q09666	3	3.9532	K.VEGEMKVPDVDIKGPK.V	2	AHNK_HUMAN
Q09666	2	4.0723	K.VEGEMKVPDVDIK.G	1	AHNK_HUMAN
Q09666	2	4.5998	K.VEGEIKAPDVDIKGPK.V	1	AHNK_HUMAN
Q09666	2	4.1712	K.VDVSAPDVEAHGPEWNLK.M	3	AHNK_HUMAN
Q09666	2	2.9306	K.VDVEVPDVSLEGPEGK.L	1	AHNK_HUMAN
Q09666	2	3.8827	K.VDTNAPDLSLEGPEGK.L	3	AHNK_HUMAN
Q09666	2	2.7311	K.VKVPEVDVR.G	1	AHNK_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q09666	3	3.7483	K.FSM#PGFKGEGPEVDM#NLPK.A	1	AHNK_HUMAN
Q09666	2	3.0688	K.GEGPEVDVKLPK.A	1	AHNK_HUMAN
Q09666	2	3.355	K.GEGPDVHMTLPK.G	3	AHNK_HUMAN
Q09666	2	2.9015	K.GDVDVSVPK.V	1	AHNK_HUMAN
Q09666	2	2.7537	K.GDISISGPK.V	2	AHNK_HUMAN
Q09666	3	4.7967	K.FSVSGAKGEETGIDVTLPTGEVTVPGVSGDV	2	AHNK_HUMAN
Q09666	2	3.6106	K.FSMPGFKGEGPEVDVK.L	3	AHNK_HUMAN
Q09666	2	3.196	K.FSMPGFKGEGPEVDMNLPK.A	1	AHNK_HUMAN
Q09666	2	3.6809	K.FSMPGFKGEGPEVDM#NLPK.A	3	AHNK_HUMAN
Q09666	2	5.4794	K.FSMPGFKAEGPEVDVNLPK.A	8	AHNK_HUMAN
Q09666	2	3.0542	K.FSM#PSLKGEGPEVDVNLPK.A	1	AHNK_HUMAN
Q09666	2	3.7803	K.FSM#PSLKGEGPEFDVNLSK.A	3	AHNK_HUMAN
Q09666	2	3.0757	K.FSM#PGFKGEGR.E	1	AHNK_HUMAN
Q09666	2	2.728	K.GEYDVTMPK.V	1	AHNK_HUMAN
Q09666	2	3.8978	K.FSM#PGFKGEGPEVDMNLPK.A	2	AHNK_HUMAN
Q09666	2	2.8663	K.FKMPSMNIQTHK.I	2	AHNK_HUMAN
Q09666	2	3.9041	K.FSM#PGFKAEGPEVDVNLPK.A	5	AHNK_HUMAN
Q09666	3	3.8536	K.FKMPEMNIKVPK.I	1	AHNK_HUMAN
Q09666	3	3.716	K.FKM#PDMHFK.A	1	AHNK_HUMAN
Q09666	1	2.3114	K.FGVSTGR.E	2	AHNK_HUMAN
Q09666	1	2.3206	K.FGTFGGLGSK.S	2	AHNK_HUMAN
Q09666	3	4.4118	K.ASLGSLEGEAEAEASSPK.G	3	AHNK_HUMAN
Q09666	1	3.594	K.ANVDISAPK.V	4	AHNK_HUMAN
Q09666	3	4.2456	K.AEASIQAGAGDGEWEESEVKLK.K	1	AHNK_HUMAN
Q09666	3	4.3783	K.ADVDVSGPKM#DAEVPDVNIEGPDAK.L	1	AHNK_HUMAN
Q09666	2	3.055	K.ADLGVSGPK.V	8	AHNK_HUMAN
Q09666	2	4.6346	K.ADIDVSGPSVDTDAPDLDIEGPEGK.L	2	AHNK_HUMAN
Q09666	2	4.142	R.ISMSEVDLNVAAPK.V	2	AHNK_HUMAN
Q09666	3	4.0962	K.LKGEIDASVPELEGDLRGPQVDVK.G	1	AHNK_HUMAN
Q09666	2	3.2475	K.FSM#PGFKGEGPEVDVK.L	2	AHNK_HUMAN
Q09666	2	4.0107	K.IEGEMQVPDVDIRGPK.V	2	AHNK_HUMAN
Q09666	3	5.4454	K.LKGDVDVSLPEVEGEMKVPDVDIK.G	3	AHNK_HUMAN
Q09666	2	4.8471	K.ISMQDVDLSLGSPK.L	3	AHNK_HUMAN
Q09666	2	3.321	K.ISMPDVGLNLKAPK.L	1	AHNK_HUMAN
Q09666	2	3.2335	K.ISMPDVGLNLK.A	2	AHNK_HUMAN
Q09666	2	3.0554	K.ISMPDVDLNLKGPK.L	2	AHNK_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q09666	2	3.1966	K.ISMPDVDLNLK.G	2	AHNK_HUMAN
Q09666	2	2.7923	K.ISMPDVDLHMK.G	1	AHNK_HUMAN
Q09666	2	2.7395	K.ISMPDVDLHLKGPNVK.G	1	AHNK_HUMAN
Q09666	2	5.2217	K.ISM#QDVDLSLGSPK.L	5	AHNK_HUMAN
Q09666	2	3.7047	K.ISM#PDVDLNLKGPK.L	3	AHNK_HUMAN
Q09666	3	3.9836	K.FKM#PEMNIKVPK.I	1	AHNK_HUMAN
Q09666	3	3.9311	K.ISM#PDVDLHVK.G	1	AHNK_HUMAN
Q09666	2	5.144	K.GGKPGLTIQAPQLEVSVPSANIEGLEGK.L	2	AHNK_HUMAN
Q09666	2	3.1046	K.ISIPDVGLHLK.G	1	AHNK_HUMAN
Q09666	1	2.1227	K.GNVDISAPK.I	2	AHNK_HUMAN
Q09666	2	3.4972	K.GGVDVTLPR.V	4	AHNK_HUMAN
Q09666	2	4.203	K.GGVTGSPEASISGSK.G	3	AHNK_HUMAN
Q09666	2	4.5483	K.GGVTGSPEASISGSKGDLK.S	5	AHNK_HUMAN
Q09666	3	5.4713	K.GHYEVTGSDDETGKLQGSGVSLASK.K	2	AHNK_HUMAN
Q09666	2	3.4866	K.LKGEIDASVPELEGDLR.G	1	AHNK_HUMAN
Q09666	2	5.1951	K.GKGGVTGSPEASISGSK.G	1	AHNK_HUMAN
Q09666	2	5.6308	K.GPQITGPSLEGDLGLK.G	9	AHNK_HUMAN
Q09666	3	6.7492	K.GSLGATGEIKGPTVGGGLPGIGVQGLEGNLQ	3	AHNK_HUMAN
Q09666	2	5.9544	K.GSRVDIETPNLEGTLTGPR.L	3	AHNK_HUMAN
Q09666	2	3.2444	K.IDVTAPDVSIEEPEGK.L	1	AHNK_HUMAN
Q09666	2	3.9389	K.IEGEM#QVPDVDIR.G	5	AHNK_HUMAN
Q09666	2	3.412	K.IEGEM#QVPDVDIRGPK.V	2	AHNK_HUMAN
Q09666	2	4.1346	K.IEGEMQVPDVDIR.G	6	AHNK_HUMAN
Q09666	3	5.2579	K.GHYEVTGSDDETGKLQGSGVSLASKK.S	4	AHNK_HUMAN
Neuroblastoma breakpoint family member 7 - Homo sapiens (Human)					
P0C2Y1	2	2.9025	K.QKFLVTQMAGFLANQQNKYK.Y	1	NBPF7_HUMAN
P0C2Y1	2	2.9415	R.EQLAEGC*RLAR.H	2	NBPF7_HUMAN
Neurogranin - Homo sapiens (Human)					
Q92686	2	4.9901	R.KGPGPGGPGGAGVAR.G	6	NEUG_HUMAN
Neuromodulin - Homo sapiens (Human)					
P17677	3	5.3386	K.ESARQDEGKEEEPEADQEHA.-	1	NEUM_HUMAN
Neutral alpha-glucosidase AB precursor - Homo sapiens (Human)					
Q14697	2	4.6439	K.DPAEGDGAQPEETPR.D	2	GANAB_HUMAN
Q14697	2	3.3194	K.MMDYLQGSGETPQTDVR.W	3	GANAB_HUMAN
Q14697	3	3.8442	R.QYASLTGTQALPPLFSLGYHQSR.W	1	GANAB_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q14697	3	5.5698	R.VSQGSKDPAEGDGAQPEETPR.D	3	GANAB_HUMAN
Neutral amino acid transporter A - Homo sapiens (Human)					
P43007	3	4.2563	K.SEEETSPLVTHQNPAGPVASAPELESK.E	1	SATT_HUMAN
P43007	3	4.8905	K.SEEETSPLVTHQNPAGPVASAPELESKESVL.	1	SATT_HUMAN
Neutral and basic amino acid transport protein rBAT - Homo sapiens (Human)					
Q07837	3	4.2311	R.GILGSQEPDFKGVQPYAGMPK.E	1	SLC31_HUMAN
NEXN protein - Homo sapiens (Human)					
Q7Z2X0	2	2.8267	K.RAEQEGDDSLLITVVPVK.S	1	Q7Z2X0_HUMAN
NFATC2-interacting protein - Homo sapiens (Human)					
Q8NCF5	2	4.2809	R.LVLDPGEAPLVPVYSGK.V	4	NF2IP_HUMAN
Q8NCF5	2	2.746	R.RLVLDPGEAPLVPVYSGK.V	1	NF2IP_HUMAN
Q8NCF5	2	3.0395	R.GAADEVEVEPPEPPGPVASR.D	2	NF2IP_HUMAN
NHS-like protein 1. - Homo sapiens (Human)					
Q5SYE7	2	4.1304	K.DFAVEPAENVSEALR.A	1	Q5SYE7_HUMA
Q5SYE7	3	3.8091	R.SPGAPSAGEAEARPSPSTTPLPDSSPSR.K	1	Q5SYE7_HUMA
Nibrin - Homo sapiens (Human)					
O60934	2	3.2835	K.VQKQEEDVNVR.K	1	NBN_HUMAN
O60934	1	2.3331	R.NYFQPSTK.K	1	NBN_HUMAN
O60934	2	2.8575	R.ASQQQQTNSIR.N	1	NBN_HUMAN
O60934	2	2.7532	K.TICALICGRPIVK.P	1	NBN_HUMAN
O60934	2	3.9893	R.MDIETNDTFSDEAVPESSK.I	4	NBN_HUMAN
O60934	2	3.9237	K.TSSNNNSM#VSNTLAK.M	2	NBN_HUMAN
Nicastrin precursor - Homo sapiens (Human)					
Q92542	1	2.1668	R.LPRC*VRSTAR.L	1	NICA_HUMAN
Nidogen-1 precursor - Homo sapiens (Human)					
P14543	2	3.7326	R.RVLFETDLVNPR.G	1	NID1_HUMAN
P14543	2	3.2945	R.VLFETDLVNPR.G	2	NID1_HUMAN
Nipped-B-like protein - Homo sapiens (Human)					
Q6KC79	2	3.0767	R.LSSDDGDSSTM#R.N	2	NIPBL_HUMAN
Q6KC79	2	3.2227	R.DKDGNVTQETK.K	3	NIPBL_HUMAN
Nitric oxide synthase-interacting protein - Homo sapiens (Human)					
Q9Y314	2	2.8526	K.ESAIVSRPLNPFTAK.A	1	NOSIP_HUMAN
Q9Y314	2	3.4165	K.VLPSFWIPSLTPEAK.A	3	NOSIP_HUMAN
NK-tumor recognition protein - Homo sapiens (Human)					
P30414	2	3.207	R.DM#PVVTAEPEPK.I	2	NKTR_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
NMDA receptor-regulated protein 2 - Homo sapiens (Human)					
Q659A1	2	2.7205	K.M#QDELLKPISRK.V	1	NARG2_HUMAN
Nonhistone chromosomal protein HMG-14 - Homo sapiens (Human)					
P05114	2	4.4512	K.TEESPASDEAGEKEAKSD.-	6	HMGN1_HUMAN
P05114	2	4.0671	K.GKQAEVANQETK.E	2	HMGN1_HUMAN
P05114	2	4.5696	R.KVSSAEGAAKEEPK.R	3	HMGN1_HUMAN
P05114	2	4.0621	K.VSSAEGAAKEEPKR.R	1	HMGN1_HUMAN
P05114	2	3.9322	K.VSSAEGAAKEEPK.R	9	HMGN1_HUMAN
P05114	2	3.5734	K.TEESPASDEAGEK.E	3	HMGN1_HUMAN
P05114	3	4.4948	K.EDLPAENGETKTEESPASDEAGEK.E	1	HMGN1_HUMAN
P05114	3	6.0189	K.GKQAEVANQETKEDLPAENGETK.T	1	HMGN1_HUMAN
P05114	2	3.2386	K.QAEVANQETKEDLPAENGETK.T	1	HMGN1_HUMAN
P05114	2	4.2309	K.TEESPASDEAGEKEAK.S	2	HMGN1_HUMAN
Nonhistone chromosomal protein HMG-17 - Homo sapiens (Human)					
P05204	2	5.3446	K.TDQAQKAEGAGDAK.-	39	HMGN2_HUMAN
P05204	2	4.551	K.ADAGKEGNNPAENGDAK.T	7	HMGN2_HUMAN
P05204	3	6.2044	K.ADAGKEGNNPAENGDAKTDQAQK.A	2	HMGN2_HUMAN
P05204	2	3.1626	K.EGNNPAENGDAK.T	8	HMGN2_HUMAN
P05204	2	4.2786	K.EGNNPAENGDAKTDQAQK.A	6	HMGN2_HUMAN
P05204	3	4.075	K.EGNNPAENGDAKTDQAQKAEGAGDAK.-	2	HMGN2_HUMAN
P05204	3	4.8284	K.GKADAGKEGNNPAENGDAK.T	9	HMGN2_HUMAN
Non-POU domain-containing octamer-binding protein - Homo sapiens (Human)					
Q15233	2	3.1605	R.RQQEEMMR.R	3	NONO_HUMAN
Q15233	2	3.6674	R.MGQMAMGGAMGINNR.G	2	NONO_HUMAN
Q15233	2	3.5019	R.MGQM#AMGGAMGINNR.G	1	NONO_HUMAN
Q15233	2	3.5928	R.FGQAATMEGIGAIGGTPPAFNR.A	1	NONO_HUMAN
Q15233	2	5.0353	K.ALIEMEKQQQDQVDR.N	4	NONO_HUMAN
Q15233	2	3.3429	R.FAQPGSFEYEYAMR.W	3	NONO_HUMAN
Nonsecretory ribonuclease precursor - Homo sapiens (Human)					
P10153	3	4.4237	R.RDPPQYPVVPVHLDR.I	1	RNAS2_HUMAN
Nonspecific lipid-transfer protein - Homo sapiens (Human)					
P22307	2	2.8075	K.IGGIFAFK.V	1	NLTP_HUMAN
P22307	3	4.0944	K.KLEEEGEQFVKK.I	1	NLTP_HUMAN
P22307	2	4.3514	K.LQNLQLQPGNAKL.-	4	NLTP_HUMAN
P22307	2	3.3793	K.M#NPQSAFFQGK.L	2	NLTP_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Novel protein - Homo sapiens (Human)					
Q5T992	2	3.7859	K.WQNVSLESWNQPR.K	1	Q5T992_HUMAN
Q5T377	2	3.4734	K.LLGSENDTGSPDFYTPR.T	3	Q5T377_HUMAN
Q5T992	3	4.0232	K.KPVWEEELR.M	4	Q5T992_HUMAN
Q5T992	2	4.0672	R.GGADSSDSQDSQQM#DAYVPR.H	3	Q5T992_HUMAN
Novel protein similar to multidomain presynaptic cytomatrix protein Piccolo; piccolo - Homo sapie					
Q5HYW2	2	3.292	R.ISLQSQEEAEK.K	2	Q5HYW2_HUMA
NSFL1 cofactor p47 - Homo sapiens (Human)					
Q9UNZ2	2	3.7988	K.SPNELVDDLFK.G	5	NSF1C_HUMAN
Q9UNZ2	2	4.098	R.DLIHDQDEDEEEEEGQR.F	2	NSF1C_HUMAN
Q9UNZ2	1	2.2232	R.FYAGGSER.S	1	NSF1C_HUMAN
Q9UNZ2	2	3.0233	R.SGQQIVGPPR.K	1	NSF1C_HUMAN
Nuclear autoantigen Sp-100 - Homo sapiens (Human)					
P23497	3	4.5492	R.VIGQDHDFSESSEEEAPAEASSGALR.S	1	SP100_HUMAN
P23497	2	3.2943	R.VVYNVLSELEK.T	1	SP100_HUMAN
P23497	2	5.3889	R.SGLQLSLEQGTGENSFR.S	4	SP100_HUMAN
P23497	2	2.7176	R.LLYDIVFK.H	1	SP100_HUMAN
P23497	2	2.8716	K.TFPFLEGLR.D	2	SP100_HUMAN
P23497	3	5.0445	K.GFENVIHDKLPLQESEEEEREER.S	2	SP100_HUMAN
P23497	3	4.7169	R.NKVEISNAIKK.T	4	SP100_HUMAN
Nuclear factor 1 - Homo sapiens (Human)					
Q8TA97	3	4.4433	K.VAASHHPPDRPPDPFSTL.-	1	Q8TA97_HUMAN
Nuclear factor NF-kappa-B p100 subunit - Homo sapiens (Human)					
Q00653	2	3.2093	R.DSGEEAAEPSAPSR.T	2	NFKB2_HUMAN
Nuclear factor of activated T-cells, cytoplasmic 2 - Homo sapiens (Human)					
Q13469	2	3.0216	R.LSPGSYPTVIQQQNATSQR.A	1	NFAC2_HUMAN
Q13469	3	3.942	R.IEITPSHELIQAVGPLR.M	1	NFAC2_HUMAN
Nuclear factor of activated T-cells, cytoplasmic 3 - Homo sapiens (Human)					
Q12968	2	4.7342	R.DM#SQISVSQGAGVSR.Q	3	NFAC3_HUMAN
Q12968	2	2.8067	R.DMSQISVSQGAGVSR.Q	1	NFAC3_HUMAN
Nuclear fragile X mental retardation-interacting protein 2 - Homo sapiens (Human)					
Q7Z417	2	2.7381	R.VDGSKPIWK.Y	1	NUFP2_HUMAN
Q7Z417	2	4.1395	R.VLNGNQQVVDTSLK.Q	2	NUFP2_HUMAN
Q7Z417	2	2.8726	R.TSPQVLGSILK.S	1	NUFP2_HUMAN
Q7Z417	2	2.8691	K.YETGPGGTSR.G	2	NUFP2_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q7Z417	2	3.4805	K.TGYGELNGNAGER.E	2	NUFP2_HUMAN
Q7Z417	2	2.8453	K.QGLETFKPDYSEQK.G	1	NUFP2_HUMAN
Q7Z417	2	4.0703	K.IMQQETSVPTLK.Q	1	NUFP2_HUMAN
Q7Z417	2	3.6001	K.IM#QQETSVPTLK.Q	1	NUFP2_HUMAN
Q7Z417	4	6.1227	K.HASAVASKEDSWTLFKPPPVFPVDNSSAK.I	2	NUFP2_HUMAN
Q7Z417	3	4.2341	K.EDSWTLFKPPPVFPVDNSSAK.I	2	NUFP2_HUMAN
Q7Z417	2	3.4023	K.GADNDGSGSESGYTTPK.K	3	NUFP2_HUMAN
Q7Z417	2	4.6162	K.NLSSDEATNPISR.V	5	NUFP2_HUMAN
Nuclear mitotic apparatus protein 1 - Homo sapiens (Human)					
Q14980	3	4.5683	R.KNSLISSLEEEVSILNR.Q	3	NUMA1_HUMAN
Q14980	3	4.9131	R.VAQEKDQLQEQLQALKESLK.V	2	NUMA1_HUMAN
Q14980	2	3.1973	R.QFLEVELDQAR.E	1	NUMA1_HUMAN
Q14980	2	3.8753	R.LLQAETASNSAR.A	2	NUMA1_HUMAN
Q14980	2	2.8162	R.HREELEQSK.Q	2	NUMA1_HUMAN
Q14980	2	2.7723	R.ERKELEEER.A	1	NUMA1_HUMAN
Q14980	2	3.0174	R.ELGELIPLR.Q	2	NUMA1_HUMAN
Q14980	2	4.0892	R.DSAQTSVTQAQR.E	2	NUMA1_HUMAN
Q14980	2	4.3734	R.AALMESQGQQQEER.G	3	NUMA1_HUMAN
Q14980	2	3.806	R.AALM#ESQGQQQEER.G	2	NUMA1_HUMAN
Q14980	2	3.1947	K.YVQELAAVR.A	2	NUMA1_HUMAN
Q14980	2	3.2137	K.QAQLAQTLQQQEQASQGLR.H	1	NUMA1_HUMAN
Q14980	2	3.8518	K.LADDLSTLQEK.M	3	NUMA1_HUMAN
Q14980	3	7.8144	K.EKQAQLAQTLQQQEQASQGLR.H	2	NUMA1_HUMAN
Q14980	3	5.7769	K.EKEHASGSGAQSEAAGR.T	3	NUMA1_HUMAN
Q14980	3	3.8404	K.EHASGSGAQSEAAGR.T	1	NUMA1_HUMAN
Q14980	3	3.8942	R.QELTSQAERAEELGQELK.A	1	NUMA1_HUMAN
Q14980	4	7.2172	R.AERDSALETLQGQLEEKAQELGHSQSALASA	1	NUMA1_HUMAN
Nuclear pore complex protein Nup153 - Homo sapiens (Human)					
P49790	2	3.627	K.QLSAQSYGVTSSTAR.R	2	NU153_HUMAN
P49790	2	2.9899	R.SGIDITDFQAK.R	2	NU153_HUMAN
P49790	2	3.4742	R.IPSIVSSPLNSPLDR.S	1	NU153_HUMAN
P49790	3	4.4293	K.RIPSIVSSPLNSPLDR.S	2	NU153_HUMAN
P49790	2	3.4796	K.NVFSSSGTSFSGR.K	2	NU153_HUMAN
P49790	2	3.0853	K.NTSLPPLWSPEAER.S	1	NU153_HUMAN
P49790	2	3.0188	K.TSQLGDSPFYPGK.T	1	NU153_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Nuclear pore complex protein Nup205 - Homo sapiens (Human)					
Q92621	2	2.8645	K.RCIANSLKALIQSR.R	1	NU205_HUMAN
Nuclear pore complex protein Nup93 - Homo sapiens (Human)					
Q8N1F7	2	3.1025	R.NLQEIQQAGER.L	1	NUP93_HUMAN
Nuclear pore complex protein Nup98-Nup96 precursor [Contains: Nuclear pore complex protein N					
P52948	2	4.2403	K.NLNNSNLFSPVNR.D	4	NUP98_HUMAN
P52948	2	2.918	K.RISEACSLAQQSGDHR.L	1	NUP98_HUMAN
P52948	2	2.8518	K.VGYYTIPSMDDLAK.I	1	NUP98_HUMAN
Nuclear protein Hcc-1 - Homo sapiens (Human)					
P82979	2	3.6658	K.GLSSDNKPMVNLDK.L	1	HCC1_HUMAN
P82979	2	3.0114	R.FNVPVSLESK.K	1	HCC1_HUMAN
P82979	2	3.8764	R.FNVPVSLESKK.A	2	HCC1_HUMAN
P82979	4	7.6614	R.LQAYLEEHAEEEANEEDVLGDETEEEETKPIE	1	HCC1_HUMAN
P82979	2	2.9793	K.SEDDEKLK.K	4	HCC1_HUMAN
P82979	2	3.7117	K.ITSEIPQTER.M	7	HCC1_HUMAN
P82979	2	3.8047	K.LAELKQECLAR.G	4	HCC1_HUMAN
P82979	2	2.9574	K.SEDDEKLKK.R	2	HCC1_HUMAN
Nuclear protein localization protein 4 homolog - Homo sapiens (Human)					
Q8TAT6	2	3.2849	K.TGEITASSNK.S	1	NPL4_HUMAN
Nuclear receptor coactivator 5 - Homo sapiens (Human)					
Q9HCD5	2	2.7234	R.DPYGFGDSR.D	1	NCOA5_HUMAN
Nuclear receptor coactivator 6 - Homo sapiens (Human)					
Q14686	2	3.123	R.GFDQQGLNPTTLK.A	2	NCOA6_HUMAN
Q14686	2	2.7369	R.SIVTTLVPSELISAVPTTK.S	1	NCOA6_HUMAN
Q14686	2	3.4736	R.VLSSTSEEDEPGVVK.F	1	NCOA6_HUMAN
Nuclear receptor interacting protein 2 - Homo sapiens (Human)					
A2RRE3	2	3.4674	K.DSADLLPLDSLKR.L	3	A2RRE3_HUMA
A2RRE3	2	3.2701	R.TRGQEAQLR.D	2	A2RRE3_HUMA
A2RRE3	2	4.0189	K.DSADLLPLDSLK.R	4	A2RRE3_HUMA
Nuclear transcription factor Y subunit alpha - Homo sapiens (Human)					
P23511	3	6.8179	K.EKDSPHMQDPNQADEEAM#TQIIR.V	2	NFYA_HUMAN
P23511	3	5.2707	K.EKDSPHMQDPNQADEEAMTQIIR.V	1	NFYA_HUMAN
P23511	2	4.4131	R.IPLPGAEM#LEEEPLYVNAK.Q	4	NFYA_HUMAN
P23511	3	5.0637	K.EKDSPHM#QDPNQADEEAMTQIIR.V	1	NFYA_HUMAN
Nuclear ubiquitous casein and cyclin-dependent kinases substrate - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9H1E3	2	2.8698	K.TSTSPPPEKSGDEGSEDEAPSGED.-	1	NUCKS_HUMAN
Q9H1E3	3	6.5825	K.VVDYSQFQESDDADEDYGR.D	4	NUCKS_HUMAN
Q9H1E3	3	5.0403	R.SGKNSQEDSEDSEDKDVK.T	4	NUCKS_HUMAN
Q9H1E3	2	3.2652	R.LKATVTPSPVK.G	5	NUCKS_HUMAN
Q9H1E3	4	5.6338	R.KVVDYSQFQESDDADEDYGRDSGPPTK.K	5	NUCKS_HUMAN
Q9H1E3	2	7.6377	R.KVVDYSQFQESDDADEDYGR.D	15	NUCKS_HUMAN
Q9H1E3	3	4.2143	K.VVDYSQFQESDDADEDYGRDSGPPTKK.I	1	NUCKS_HUMAN
Q9H1E3	3	4.9249	K.TPSPKEEDEEPESPPEKK.T	6	NUCKS_HUMAN
Q9H1E3	3	4.3859	K.TPSPKEEDEEPESPPEK.K	3	NUCKS_HUMAN
Q9H1E3	2	4.4953	K.NSQEDSEDSEDKDVK.T	14	NUCKS_HUMAN
Q9H1E3	2	2.937	K.EEDEEPESPPEKK.T	1	NUCKS_HUMAN
Q9H1E3	2	2.8091	K.ATVTPSPVKGK.G	1	NUCKS_HUMAN
Q9H1E3	1	2.1079	K.ATVTPSPVK.G	1	NUCKS_HUMAN
Q9H1E3	3	5.1118	K.VVDYSQFQESDDADEDYGRDSGPPTK.K	1	NUCKS_HUMAN
Nucleobindin-1 precursor - Homo sapiens (Human)					
Q02818	2	2.9275	R.YLESLGEEQR.K	1	NUCB1_HUMAN
Q02818	2	4.0209	R.DLELLIQTATR.D	6	NUCB1_HUMAN
Q02818	2	3.4614	R.EKLQAANAEDIK.S	2	NUCB1_HUMAN
Q02818	3	3.9048	R.EKLQAANAEDIKSGK.L	1	NUCB1_HUMAN
Q02818	2	3.3384	R.ELDFVSHHVR.T	3	NUCB1_HUMAN
Q02818	2	3.565	R.ELQQAVLHM#EQR.K	2	NUCB1_HUMAN
Q02818	2	4.5565	R.ELQQAVLHMEQR.K	4	NUCB1_HUMAN
Q02818	2	3.8842	R.LPEVEVPQHL.-	4	NUCB1_HUMAN
Q02818	2	3.8534	R.LSQETEALGR.S	4	NUCB1_HUMAN
Q02818	2	4.437	R.LVTLEEFLASTQRK.E	3	NUCB1_HUMAN
Q02818	2	2.9108	R.YEMLKEHER.R	2	NUCB1_HUMAN
Q02818	2	5.5998	K.MDAEQDPNVQVDHLNLLK.Q	5	NUCB1_HUMAN
Q02818	2	3.2658	R.YLESLGEEQRK.E	3	NUCB1_HUMAN
Q02818	2	3.1539	R.YLESLGEEQRKEAER.K	1	NUCB1_HUMAN
Q02818	3	5.6392	R.YLQEVIDVLETDGHFR.E	4	NUCB1_HUMAN
Q02818	4	4.798	R.YLQEVIDVLETDGHFREK.L	2	NUCB1_HUMAN
Q02818	2	3.0862	R.SQGRLEAQKR.E	5	NUCB1_HUMAN
Q02818	2	3.0619	K.EVWEELDGLDPNR.F	2	NUCB1_HUMAN
Q02818	2	2.8644	K.VNVPGSQAQLK.E	1	NUCB1_HUMAN
Q02818	2	3.4082	R.DLAQYDAAHHEEFK.R	1	NUCB1_HUMAN
Q02818	3	4.9275	K.AKM#DAEQDPNVQVDHLNLLK.Q	2	NUCB1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q02818	3	5.7097	K.AKMDAEQDPNVQVDHLNLLK.Q	5	NUCB1_HUMAN
Q02818	2	3.0801	K.APAAHPEGQLK.F	2	NUCB1_HUMAN
Q02818	2	3.8278	K.EVWEELDGLDPNRFNPK.T	3	NUCB1_HUMAN
Q02818	2	4.8325	K.FHPDTDDVPVPAPAGDQK.E	6	NUCB1_HUMAN
Q02818	3	4.6073	K.FHPDTDDVPVPAPAGDQKEVDTSEK.K	1	NUCB1_HUMAN
Q02818	3	3.8679	K.KLLERLPEVEVPQHL.-	2	NUCB1_HUMAN
Q02818	2	4.1784	K.LLERLPEVEVPQHL.-	8	NUCB1_HUMAN
Q02818	3	5.4003	K.NVDTNQDRLVTLEEFLASTQRK.E	2	NUCB1_HUMAN
Q02818	1	3.3644	K.LQAANAEDIK.S	4	NUCB1_HUMAN
Q02818	4	5.018	K.AKMDAEQDPNVQVDHLNLLKQFEHLDPQNQ	1	NUCB1_HUMAN
Q02818	2	4.6044	K.LQAANAEDIKSGK.L	3	NUCB1_HUMAN
Q02818	3	4.7551	K.M#DAEQDPNVQVDHLNLLK.Q	1	NUCB1_HUMAN
Q02818	3	4.3035	K.FHPDTDDVPVPAPAGDQKEVDTSEKK.L	2	NUCB1_HUMAN
Nucleobindin-2 precursor - Homo sapiens (Human)					
P80303	2	2.78	R.LVTLEEFLK.A	3	NUCB2_HUMAN
P80303	3	4.0327	R.MREHVM#NEVDTNKDR.L	1	NUCB2_HUMAN
P80303	2	5.6741	R.MREHVMNEVDTNKDR.L	4	NUCB2_HUMAN
Nucleolar MIF4G domain-containing protein 1 - Homo sapiens (Human)					
Q5C9Z4	3	3.9323	R.SGAEEASGHRQDTEER.A	2	NOM1_HUMAN
Nucleolar phosphoprotein p130 - Homo sapiens (Human)					
Q14978	2	2.9439	R.GAAGDWGER.A	1	NOLC1_HUMAN
Q14978	3	4.3466	R.VVPSDLYPLVLGFLR.D	3	NOLC1_HUMAN
Q14978	2	3.4108	R.VREEEIEVDSR.V	1	NOLC1_HUMAN
Q14978	2	2.7292	K.AALSLPAK.Q	1	NOLC1_HUMAN
Q14978	2	3.0967	R.VADNSFDAKR.G	1	NOLC1_HUMAN
Q14978	2	3.0957	R.VADNSFDAK.R	5	NOLC1_HUMAN
Q14978	2	3.2419	K.VAGGAAPSKPASAK.K	2	NOLC1_HUMAN
Q14978	2	2.9212	K.SPAVKPAAAPK.Q	2	NOLC1_HUMAN
Q14978	2	3.3665	K.RGAAGDWGER.A	2	NOLC1_HUMAN
Q14978	1	2.1375	K.AAVVVSK.S	2	NOLC1_HUMAN
Q14978	2	3.6871	K.NSSNKPAVTTK.S	4	NOLC1_HUMAN
Q14978	2	2.7806	R.DNQLSEVANK.F	1	NOLC1_HUMAN
Nucleolar protein 3 - Homo sapiens (Human)					
O60936	2	5.2421	R.GVLTGPEYEALDALPDAER.R	6	NOL3_HUMAN
O60936	2	2.824	R.LLLLVQGK.G	5	NOL3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O60936	2	2.9245	K.GEAACQELLR.C	1	NOL3_HUMAN
O60936	3	4.9557	R.LVETLQADSGLLLDALLAR.G	7	NOL3_HUMAN
Nucleolar protein 5 - Homo sapiens (Human)					
Q9Y2X3	2	3.3895	K.TYDPSGDSTLPTCSK.K	1	NOL5_HUMAN
Q9Y2X3	3	6.08	K.VEEEEEEKVAEEEETSVK.K	1	NOL5_HUMAN
Nucleolar protein 7 - Homo sapiens (Human)					
Q9UMY1	2	3.1498	R.KLLPDTILEK.L	3	NOL7_HUMAN
Nucleolar RNA helicase 2 - Homo sapiens (Human)					
Q9NR30	3	4.0735	K.KKAEPSEVDMNSPK.S	2	DDX21_HUMAN
Q9NR30	3	4.233	K.KKEEPSQNDISPK.T	1	DDX21_HUMAN
Nucleoporin-like protein RIP - Homo sapiens (Human)					
P52594	2	4.4588	K.SLLGDSAPTLHLNK.G	2	NUPL_HUMAN
P52594	2	4.8239	K.SSSADFGTFNTSQSHQTASAVSK.V	2	NUPL_HUMAN
P52594	3	6.7363	K.AGLQTADKYAALANLDNIFSAGQGGDQGSGF	2	NUPL_HUMAN
Nucleosome assembly protein 1-like 1 - Homo sapiens (Human)					
P55209	2	4.7504	K.NVDLLSDMVQEHDEPILK.H	1	NP1L1_HUMAN
P55209	2	3.1703	K.YAVLYQPLFDK.R	2	NP1L1_HUMAN
Nucleosome assembly protein 1-like 4 - Homo sapiens (Human)					
Q99733	2	3.6289	K.AAATAEEPDPK.G	3	NP1L4_HUMAN
Q99733	2	3.669	K.LTDQVM#QNPR.V	2	NP1L4_HUMAN
Q99733	2	4.6234	K.LTDQVMQNPR.V	3	NP1L4_HUMAN
Q99733	3	4.5151	K.NASNTEKLTDQVMQNPR.V	3	NP1L4_HUMAN
Q99733	2	2.746	K.QVPNESFFNFFNPLK.A	1	NP1L4_HUMAN
Q99733	2	3.3538	R.KYAALYQPLFDK.R	2	NP1L4_HUMAN
Q99733	3	5.0399	R.LDNVPHTPSSYIETLPK.A	5	NP1L4_HUMAN
Q99733	2	3.2605	R.VLAALQER.L	1	NP1L4_HUMAN
Nucleosome assembly protein 1-like 5 - Homo sapiens (Human)					
Q96NT1	2	4.1063	K.NDFIESLPNSVK.C	5	Q96NT1_HUMA
Q96NT1	2	3.0825	K.FDKEFQALEK.K	1	Q96NT1_HUMA
Q96NT1	3	4.8591	K.FDKEFQALEKK.Y	2	Q96NT1_HUMA
Q96NT1	3	4.1796	K.HDDAHAEM#PDDAKK.-	1	Q96NT1_HUMA
NudC domain-containing protein 2 - Homo sapiens (Human)					
Q8WVJ2	2	2.72	K.GGPDFSNLEK.-	1	NUDC2_HUMAN
OCIA domain-containing protein 2 - Homo sapiens (Human)					
Q56VL3	2	3.2722	K.HGLSEKGDSQPSAS.-	2	Q56VL3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Opioid growth factor receptor - Homo sapiens (Human)					
Q9NZT2	2	3.918	R.DADAGDEDEESEEPR.A	6	OGFR_HUMAN
Optineurin - Homo sapiens (Human)					
Q96CV9	2	4.3032	R.AVLKELSEKLELAEK.A	1	OPTN_HUMAN
Q96CV9	2	3.7106	R.TSDSDQQAYLVQR.G	5	OPTN_HUMAN
Q96CV9	2	3.3033	R.SSEDPTDDSR.L	2	OPTN_HUMAN
Q96CV9	2	2.7295	R.QSLMEMQSR.H	1	OPTN_HUMAN
Q96CV9	2	4.3396	R.MAEGEAEGSVKEIK.H	2	OPTN_HUMAN
Q96CV9	1	3.5566	R.MAEGEAEGSVK.E	4	OPTN_HUMAN
Q96CV9	2	2.7792	K.LSEAELM#K.K	1	OPTN_HUMAN
Q96CV9	2	2.9156	R.M#AEGEAEGSVK.E	1	OPTN_HUMAN
Q96CV9	2	4.1926	R.LQAEKADLLGIVSELQLK.L	2	OPTN_HUMAN
Q96CV9	2	3.5469	R.LMALSHENEKLKEELGK.L	1	OPTN_HUMAN
Q96CV9	2	3.043	R.LMALSHENEK.L	1	OPTN_HUMAN
Q96CV9	3	4.8541	R.KNSAIPSELNEKQELVYTNK.K	2	OPTN_HUMAN
Q96CV9	3	3.7669	R.AQMEVYCSDFHAER.A	1	OPTN_HUMAN
Q96CV9	2	3.7021	R.AEAEQEKDQLR.T	2	OPTN_HUMAN
Q96CV9	2	2.8432	K.TIEELTRK.E	1	OPTN_HUMAN
Q96CV9	3	4.0025	K.QTIAKQEEDLETMTILR.A	1	OPTN_HUMAN
Q96CV9	3	4.1813	K.QLQMDEMKQTIAKQEEDLETMTILR.A	2	OPTN_HUMAN
Q96CV9	2	2.7682	K.NSAIPSELNEKQELVYTNK.K	1	OPTN_HUMAN
Q96CV9	2	2.9576	K.LSEAELMK.K	1	OPTN_HUMAN
Q96CV9	2	2.7119	K.LLQEHNNALK.T	1	OPTN_HUMAN
Q96CV9	2	4.7427	K.GRFEELSAWTEK.Q	3	OPTN_HUMAN
Q96CV9	2	3.3656	K.ELSEKLELAEK.A	1	OPTN_HUMAN
Q96CV9	2	3.4721	K.LTVLQMTHNK.L	2	OPTN_HUMAN
Organic-cation transporter-like 3 - Homo sapiens (Human)					
Q9Y226	2	2.7593	R.WLLTRGRMDEAIQLIQK.A	1	ORCT3_HUMAN
Origin recognition complex subunit 2 - Homo sapiens (Human)					
Q13416	2	2.909	R.AQLLVNPK.K	1	ORC2_HUMAN
Osteoclast-stimulating factor 1 - Homo sapiens (Human)					
Q92882	2	2.866	R.ALYTFEPR.T	1	OSTF1_HUMAN
Q92882	2	3.8481	K.GYADIVQLLLAK.G	2	OSTF1_HUMAN
Osteopontin precursor - Homo sapiens (Human)					
P10451	2	2.8706	R.ISHELDSASSEVN.-	1	OSTP_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P10451	3	4.7789	K.ANDESNEHSDVIDSQELSK.V	1	OSTP_HUMAN
OTTHUMP00000017057 - Homo sapiens (Human)					
Q5TFE7	2	2.7951	K.AGQEDAPPSTK.G	1	Q5TFE7_HUMA
OTTHUMP00000028696 - Homo sapiens (Human)					
Q5W189	2	5.2831	K.LASSDTGESDQSSTETDSTVK.S	10	Q5W189_HUMA
OTU domain-containing protein 4 - Homo sapiens (Human)					
Q01804	2	4.6438	K.TAADVVSPGANSVDSR.V	2	OTUD4_HUMAN
Q01804	2	3.9571	R.QHAFSSHSSGSQSQK.F	1	OTUD4_HUMAN
Oxidation resistance protein 1 - Homo sapiens (Human)					
Q8N573	2	3.6773	K.FDTTPNELVQLNK.L	2	OXR1_HUMAN
Q8N573	2	3.4546	K.MTGSNTEEIDSR.I	2	OXR1_HUMAN
Q8N573	2	3.7442	K.SSGASSESVQTVNQAEVESLTVK.S	1	OXR1_HUMAN
Q8N573	2	4.1777	R.DAGNDSASTAPR.S	3	OXR1_HUMAN
Oxysterol-binding protein-related protein 10 - Homo sapiens (Human)					
Q9BXB5	2	4.0842	R.AVQGTDGGGGSNSSSR.S	5	OSB10_HUMAN
P37 AUF1 - Homo sapiens (Human)					
Q12771	2	4.1864	K.IDASKNEEDEGK.M	4	Q12771_HUMAN
Q12771	3	5.8837	K.IDASKNEEDEGKMFIGGLSWDTTK.K	2	Q12771_HUMAN
Palmdelphin - Homo sapiens (Human)					
Q9NP74	2	3.8446	K.EINEEKEDDEQNR.K	1	PALMD_HUMAN
Q9NP74	2	3.3899	K.EINEEKEDDEQNRK.A	2	PALMD_HUMAN
Q9NP74	2	3.3335	K.IQEEISQKR.L	1	PALMD_HUMAN
Q9NP74	3	4.0219	K.SPTEYHEPVYANPFYRPTTPQR.E	2	PALMD_HUMAN
Q9NP74	2	3.0863	K.TGESTVLSSIPLPSDDFKGTGIK.V	1	PALMD_HUMAN
Q9NP74	2	3.6683	R.AEESIEDIYANIPDLPK.S	2	PALMD_HUMAN
Q9NP74	2	2.7265	R.SVKVEREER.A	1	PALMD_HUMAN
Pantothenate kinase 2, mitochondrial precursor - Homo sapiens (Human)					
Q9BZ23	2	4.4517	R.ASATSVSSAGEQAAGDPEGR.R	4	PANK2_HUMAN
Paralemmin - Homo sapiens (Human)					
O75781	3	4.7627	R.KTEVVMNSQQTPVGTPK.D	2	PALM_HUMAN
O75781	2	3.1189	R.WLLEGTPSSASEGDEDLRR.Q	1	PALM_HUMAN
O75781	2	2.8778	R.APAPSPAKEER.K	1	PALM_HUMAN
O75781	2	3.5786	R.VLSSTTLLPR.Q	1	PALM_HUMAN
O75781	3	3.9682	R.LLEDSVSRLEKEIEVLER.G	2	PALM_HUMAN
Paraspeckle component 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8WXF1	3	4.9781	K.NLSPVVSNELLEQAFSQFGPVEK.A	1	PSPC1_HUMAN
Q8WXF1	2	4.4852	R.FAQPGTFEFEYASR.W	2	PSPC1_HUMAN
Q8WXF1	3	5.0586	R.TGSETPQAPM#SGVGPVSGGPGGFGR.G	2	PSPC1_HUMAN
Parathymosin - Homo sapiens (Human)					
P20962	2	4.571	K.SVEAAAELSAKDLK.E	2	PTMS_HUMAN
P20962	1	4.3814	K.SVEAAAELSAK.D	116	PTMS_HUMAN
Parvalbumin alpha - Homo sapiens (Human)					
P20472	2	2.7407	K.FFQMVGLK.K	1	PRVA_HUMAN
PC4 and SFRS1-interacting protein - Homo sapiens (Human)					
O75475	2	3.6766	K.NMFLVGEGDSVITQVLNK.S	2	PSIP1_HUMAN
PCDH12 protein - Homo sapiens (Human)					
Q7Z738	2	3.0919	R.NQGNQGAPAESR.E	1	Q7Z738_HUMAN
PDZ and LIM domain protein 1 - Homo sapiens (Human)					
O00151	2	4.0535	R.SAMPFTASPASSTTAR.V	7	PDLI1_HUMAN
O00151	3	3.9002	R.SEHKVWSPLVTEEGK.R	1	PDLI1_HUMAN
O00151	2	4.5448	R.SAM#PFTASPASSTTAR.V	10	PDLI1_HUMAN
O00151	4	6.8904	K.TAASGVEANSRPLDHAQPPSSLVIDKESEVY	4	PDLI1_HUMAN
O00151	2	3.1197	K.VWSPLVTEEGKR.H	3	PDLI1_HUMAN
O00151	3	4.6183	R.ERVTPPEGYEVVTVFPK.-	4	PDLI1_HUMAN
O00151	2	2.7781	K.VWSPLVTEEGK.R	1	PDLI1_HUMAN
O00151	2	3.7412	K.VAASIGNAQK.L	5	PDLI1_HUMAN
O00151	2	5.7295	K.MNLASEPQEVLHIGSAHNR.S	2	PDLI1_HUMAN
O00151	3	4.2241	K.MLQEKQELNEPPK.Q	1	PDLI1_HUMAN
O00151	3	4.1388	K.M#NLASEPQEVLHIGSAHNR.S	1	PDLI1_HUMAN
O00151	2	3.8339	K.GHFFVEDQIYCEK.H	6	PDLI1_HUMAN
O00151	2	4.2024	K.DFEQPLAISR.V	4	PDLI1_HUMAN
O00151	2	4.8616	R.LVGGKDFEQPLAISR.V	7	PDLI1_HUMAN
O00151	2	3.6396	K.GCTDNLTLTVAR.S	2	PDLI1_HUMAN
PDZ and LIM domain protein 2 - Homo sapiens (Human)					
Q96JY6	4	5.0896	K.AKDADLRPGDIIVAINGESAEGMLHAEAQSK.I	2	PDLI2_HUMAN
Q96JY6	3	4.5464	R.AGSPFSPPPSSSSLTGEAAISR.S	6	PDLI2_HUMAN
Q96JY6	2	4.3791	R.SFQSLACSPGLPAADR.L	2	PDLI2_HUMAN
Q96JY6	2	4.1606	R.SQATSPGQTNGDSSLEVLATR.F	5	PDLI2_HUMAN
Q96JY6	2	3.0813	R.SSYSSPTSLSPR.A	1	PDLI2_HUMAN
Q96JY6	2	3.4653	R.TYTESQSSLR.S	3	PDLI2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
PDZ and LIM domain protein 3 - Homo sapiens (Human)					
Q53GG5	2	3.7291	R.VLQGMVDDGSDDRPAGTR.S	3	PDLI3_HUMAN
PDZ and LIM domain protein 5 - Homo sapiens (Human)					
Q96HC4	1	2.1572	K.TAVNVPR.Q	1	PDLI5_HUMAN
Q96HC4	2	3.9634	R.ASAAPKPEPVPVQKGEPK.E	2	PDLI5_HUMAN
Q96HC4	2	3.3618	K.VTSTNNM#AYNK.A	2	PDLI5_HUMAN
Q96HC4	2	4.0771	K.EVVKPVPITSPAVSK.V	7	PDLI5_HUMAN
Q96HC4	3	3.9587	K.APRPFGSVSSPK.V	1	PDLI5_HUMAN
Q96HC4	2	3.6986	K.VTSTNNMAYNK.A	4	PDLI5_HUMAN
PDZ and LIM domain protein 7 - Homo sapiens (Human)					
Q9NR12	2	3.1376	R.YTFAPSVSLNK.T	3	PDLI7_HUMAN
Q9NR12	2	4.065	K.VVLEGPAPWGFR.L	4	PDLI7_HUMAN
Q9NR12	2	2.8357	R.AQPVQSKPQK.A	1	PDLI7_HUMAN
Q9NR12	2	3.8717	R.HSQPATPTPLQSR.T	1	PDLI7_HUMAN
PDZ domain-containing protein 1 - Homo sapiens (Human)					
Q5T2W1	2	4.8713	K.KIPIVSSLADPLDTPPDSK.E	4	PDZD1_HUMAN
Q5T2W1	3	5.5399	R.LAHFSPFLYYQSQELPNGSVK.E	1	PDZD1_HUMAN
Q5T2W1	3	4.8921	R.KGGDQTSLLVVDKETDNMYR.L	2	PDZD1_HUMAN
Q5T2W1	2	3.4356	R.IEKDTEGHLVR.V	2	PDZD1_HUMAN
Q5T2W1	3	9.0236	R.AGVLADDHLIEVNGENVEDASHEEVVEK.V	1	PDZD1_HUMAN
Q5T2W1	2	3.4423	K.VVDRIQSSGK.N	1	PDZD1_HUMAN
Q5T2W1	3	4.4276	K.LSKQEGQNYGFFLR.I	2	PDZD1_HUMAN
Q5T2W1	2	5.2963	K.KGVYMTDITPQGVAMR.A	1	PDZD1_HUMAN
Q5T2W1	2	2.9916	K.KAYDYFQAK.K	1	PDZD1_HUMAN
Q5T2W1	2	5.4692	K.GQIIKDIDSGSPAEEAGLK.N	3	PDZD1_HUMAN
Q5T2W1	3	4.236	K.GGDQTSLLVVDKETDNMYR.L	1	PDZD1_HUMAN
Q5T2W1	2	4.3337	K.DIDSGSPAEEAGLK.N	7	PDZD1_HUMAN
Q5T2W1	2	2.9871	K.AGLQDGDR.V	2	PDZD1_HUMAN
Q5T2W1	2	2.9445	K.AYDYFQAK.K	1	PDZD1_HUMAN
PDZ domain-containing protein 11 - Homo sapiens (Human)					
Q5EBL8	2	3.8731	K.KPPGAQLGFNIR.G	3	PDZ11_HUMAN
Q5EBL8	3	5.4635	R.AGLQEGDQVLAVNDVDFQDIEHSK.A	1	PDZ11_HUMAN
Q5EBL8	3	3.9778	R.VHHPDYNNELTQFLPR.T	1	PDZ11_HUMAN
PDZ domain-containing protein 2 - Homo sapiens (Human)					
O15018	3	3.9206	K.TSVDTGQVSRPENPSQPASPR.V	1	PDZD2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O15018	2	3.4541	R.TDTQSPTNTGSPSSPQQK.S	2	PDZD2_HUMAN
PDZ domain-containing protein 8 - Homo sapiens (Human)					
Q8NEN9	3	3.9047	R.AGIEDIETLESLSLDQHSK.K	1	PDZD8_HUMAN
PDZ domain-containing protein GIPC1 - Homo sapiens (Human)					
O14908	2	2.7173	R.DTELAATMVELGKDKR.N	1	GIPC1_HUMAN
O14908	2	4.0998	R.NPDELAEALDER.L	1	GIPC1_HUMAN
O14908	2	4.7185	R.GPATVEDLPSAFEEK.A	2	GIPC1_HUMAN
PDZ domain-containing protein GIPC2 - Homo sapiens (Human)					
Q8TF65	2	3.5763	K.GPATVEEM#PSETK.A	1	GIPC2_HUMAN
PDZK1-interacting protein 1 - Homo sapiens (Human)					
Q13113	3	4.3041	R.SSEHENAYENVPEEEGKVR.S	4	PDZ1I_HUMAN
Q13113	2	2.8377	R.YSSM#AASFR.S	1	PDZ1I_HUMAN
Q13113	2	3.2106	R.YSSMAASFR.S	3	PDZ1I_HUMAN
Pentatricopeptide repeat protein 1 - Homo sapiens (Human)					
O75127	1	2.4557	K.NFELNLK.T	1	PTCD1_HUMAN
Peptidase inhibitor 16 precursor - Homo sapiens (Human)					
Q6UXB8	2	4.1213	R.WDEELAAFAK.A	3	PI16_HUMAN
Peptidyl-prolyl cis-trans isomerase - Homo sapiens (Human)					
Q6IBH5	2	3.2216	K.DTNGSQFFITTVK.T	2	Q6IBH5_HUMAN
Q6IBH5	3	4.1141	R.VIKDFMIQGGDFTR.G	2	Q6IBH5_HUMAN
Q6IBH5	2	2.7131	R.IGDEDVGR.V	1	Q6IBH5_HUMAN
Q6IBH5	2	3.2554	K.VLEGMEVVR.K	4	Q6IBH5_HUMAN
Q6IBH5	2	2.7184	K.VLEGM#EVVR.K	1	Q6IBH5_HUMAN
Q6IBH5	2	3.1415	K.HYGPGWVSMANAGK.D	1	Q6IBH5_HUMAN
Q6IBH5	2	3.9135	K.TVDNFVALATGEK.G	2	Q6IBH5_HUMAN
Peptidyl-prolyl cis-trans isomerase A - Homo sapiens (Human)					
P62937	2	4.1428	R.VSFELFADKVPK.T	7	PPIA_HUMAN
Peptidyl-prolyl cis-trans isomerase G - Homo sapiens (Human)					
Q13427	2	2.9474	K.NKEDEKIR.S	2	PPIG_HUMAN
Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1 - Homo sapiens (Human)					
Q13526	3	4.2458	R.GQMQKPFEDASFALR.T	5	PIN1_HUMAN
Q13526	2	5.5875	R.TKEEALELINGYIQK.I	4	PIN1_HUMAN
Q13526	3	4.8015	K.IKSGEEDFESLASQFSDCSSAK.A	1	PIN1_HUMAN
Q13526	3	5.1352	R.VYYFNHITNASQWERPSGNSSSGGK.N	2	PIN1_HUMAN
Q13526	3	4.107	R.GQM#QKPFEDASFALR.T	2	PIN1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Peptidyl-prolyl cis-trans isomerase-like 2 - Homo sapiens (Human)					
Q13356	2	3.8545	K.SSQPQAGSQGPQTFR.Q	2	PPIL2_HUMAN
Q13356	2	3.314	R.AAEEEPSTSATVPM#SK.K	2	PPIL2_HUMAN
Q13356	2	3.4605	R.AAEEEPSTSATVPMSK.K	2	PPIL2_HUMAN
Pericentriolar material 1 protein - Homo sapiens (Human)					
Q15154	3	4.7073	K.LLGELHTLRDQHLNNSSSSPQR.S	1	PCM1_HUMAN
Q15154	2	2.7027	R.INFSDLDQR.S	1	PCM1_HUMAN
Q15154	2	3.6466	R.KASAQASLASK.D	3	PCM1_HUMAN
Q15154	3	4.2261	R.TPWLYEQEGEVEKPFIK.T	1	PCM1_HUMAN
Periphilin-1 - Homo sapiens (Human)					
Q8NEY8	2	3.639	K.MLIEKDPSLEK.S	2	PPHLN_HUMAN
Q8NEY8	2	3.8089	R.DTSPSSGSAVSSSK.V	5	PPHLN_HUMAN
Q8NEY8	2	4.23	R.LTEKELAEAASK.W	2	PPHLN_HUMAN
Periplakin - Homo sapiens (Human)					
O60437	2	3.4003	R.ASQEEQIAR.K	2	PEPL_HUMAN
O60437	3	3.9084	K.LHSEGDQLLAAEHPGR.N	1	PEPL_HUMAN
O60437	3	4.207	R.YVNKDMSIQELAVLVSGQK.-	1	PEPL_HUMAN
O60437	3	4.4602	R.VVQQEVVRYEEEPGLR.A	1	PEPL_HUMAN
O60437	2	2.8709	R.QLLEGELETLR.R	2	PEPL_HUMAN
O60437	2	2.8594	R.LQSEINMAATETR.D	2	PEPL_HUMAN
O60437	2	5.0562	R.LQNLEFALNLLR.Q	4	PEPL_HUMAN
O60437	2	3.1175	R.KVDSDLNQK.Y	3	PEPL_HUMAN
O60437	2	3.1326	R.DGGQEYVVK.E	4	PEPL_HUMAN
O60437	2	3.0848	K.VPDPVLEESFQQLQR.T	1	PEPL_HUMAN
O60437	2	3.1923	K.TENPGDASDLQGR.Q	1	PEPL_HUMAN
O60437	3	3.7679	K.DKDLEIDELQKR.L	1	PEPL_HUMAN
O60437	2	3.0748	K.NLLDEIASR.E	2	PEPL_HUMAN
O60437	2	4.7945	K.SLLGEVEQNLQAAK.Q	2	PEPL_HUMAN
O60437	3	4.4623	K.KVPDPVLEESFQQLQR.T	2	PEPL_HUMAN
O60437	2	2.8166	K.FTEVYAINR.Q	1	PEPL_HUMAN
O60437	2	4.158	K.VVLQQDPQQAR.E	4	PEPL_HUMAN
Peroxiredoxin-1 - Homo sapiens (Human)					
Q06830	3	4.6745	K.KQGGLGPMNIPLVSDPKR.T	1	PRDX1_HUMAN
Q06830	2	4.5328	R.TIAQDYGVLKADEGISFR.G	1	PRDX1_HUMAN
Q06830	2	3.789	R.LVQAFQFTDK.H	4	PRDX1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q06830	2	3.0147	R.TIAQDYGVLK.A	1	PRDX1_HUMAN
Peroxiredoxin-2 - Homo sapiens (Human)					
P32119	3	4.7354	R.RLSEDYGVLKTDEGIAYR.G	2	PRDX2_HUMAN
P32119	2	3.3545	K.TDEGIAYR.G	3	PRDX2_HUMAN
P32119	1	2.5993	R.GLFIIDGK.G	2	PRDX2_HUMAN
P32119	3	3.9056	R.KEGGLGPLNIPLLADVTR.R	1	PRDX2_HUMAN
P32119	2	4.8592	R.LSEDYGVLKTDEGIAYR.G	5	PRDX2_HUMAN
P32119	2	3.6476	R.RLSEDYGVLK.T	2	PRDX2_HUMAN
Peroxiredoxin-4 - Homo sapiens (Human)					
Q13162	2	3.1954	R.QITLNDLPVGR.S	4	PRDX4_HUMAN
Q13162	3	4.2183	R.QGGLGPIRIPLLSDLTHQISK.D	1	PRDX4_HUMAN
Peroxiredoxin-5, mitochondrial precursor - Homo sapiens (Human)					
P30044	2	3.8597	K.GVLFGVPGAFTPGCSK.T	3	PRDX5_HUMAN
P30044	2	4.2744	K.THLPGFVEQAEALK.A	4	PRDX5_HUMAN
P30044	3	4.4934	K.THLPGFVEQAEALKAK.G	2	PRDX5_HUMAN
P30044	2	3.7852	K.VGDAIPAVEVFEGEPGNK.V	1	PRDX5_HUMAN
P30044	3	3.9815	R.LLADPTGAFGKETDLLLDDSLVSIFGNR.R	1	PRDX5_HUMAN
Peroxisomal membrane protein PEX14 - Homo sapiens (Human)					
O75381	2	3.773	R.RGGDGQINEQVEK.L	1	PEX14_HUMAN
O75381	2	3.4713	R.GGDGQINEQVEK.L	3	PEX14_HUMAN
Peroxisomal targeting signal 1 receptor - Homo sapiens (Human)					
P50542	2	3.5617	K.LAGHFTQDK.A	2	PEX5_HUMAN
P50542	3	4.0866	K.MDDLLAEMQQIEQSNFR.Q	1	PEX5_HUMAN
PERQ amino acid-rich with GYF domain-containing protein 1 - Homo sapiens (Human)					
O75420	2	3.1709	R.SIEEGDGAFGR.S	2	PERQ1_HUMAN
PERQ amino acid-rich with GYF domain-containing protein 2 - Homo sapiens (Human)					
Q6Y7W6	2	4.8269	K.ALQQQQQQQQQK.L	4	PERQ2_HUMAN
Q6Y7W6	2	3.6202	K.KVEEEEKLLK.L	2	PERQ2_HUMAN
Q6Y7W6	2	3.4454	K.NSNM#GFWDDAVK.E	1	PERQ2_HUMAN
Q6Y7W6	2	3.3233	K.NSNMGFWDDAVK.E	2	PERQ2_HUMAN
Q6Y7W6	2	4.2194	K.SLLEIQQEEAR.Q	4	PERQ2_HUMAN
Pescadillo homolog 1 - Homo sapiens (Human)					
O00541	2	2.7407	R.LAQEEESEAK.R	1	PESC_HUMAN
O00541	2	3.6471	R.LAQEEESEAKR.L	1	PESC_HUMAN
PEST proteolytic signal-containing nuclear protein - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8WW12	2	4.8273	R.AGAAGGPEEEAEKPVK.T	5	PCNP_HUMAN
Q8WW12	2	3.5803	R.DTPTSAGPNSFNK.G	4	PCNP_HUMAN
Q8WW12	3	3.7893	K.TLSVAAAFNEDEDSEPEEMPPEAK.M	1	PCNP_HUMAN
Q8WW12	2	3.193	K.SHLGNVHDQDN.-	2	PCNP_HUMAN
Q8WW12	3	5.018	K.RSAEEEAADLPTKPTK.I	3	PCNP_HUMAN
Q8WW12	2	4.879	K.NIGRDTPTSAGPNSFNK.G	2	PCNP_HUMAN
Q8WW12	2	3.1085	K.KASAISIK.L	3	PCNP_HUMAN
Q8WW12	2	3.3891	K.FGFAIGSQTTKK.A	2	PCNP_HUMAN
Q8WW12	2	4.229	K.FGFAIGSQTTK.K	5	PCNP_HUMAN
Q8WW12	2	5.1345	R.SAEEEAADLPTKPTK.I	6	PCNP_HUMAN
PH domain leucine-rich repeat-containing protein phosphatase - Homo sapiens (Human)					
O60346	3	3.7739	R.ASAPAAAAAAAAAAAAAAAALAAAAGGGR.S	1	PHLPP_HUMAN
PHD finger protein 3 - Homo sapiens (Human)					
Q92576	4	5.5189	K.KTLQDQTLVQIFKPLTHSLSDK.S	1	PHF3_HUMAN
Q92576	2	3.2881	R.KVEQDSKETVK.L	2	PHF3_HUMAN
PHD finger-like domain-containing protein 5A - Homo sapiens (Human)					
Q7RTV0	1	2.3886	K.IVNLGSSK.T	1	PHF5A_HUMAN
Phenylalanyl-tRNA synthetase alpha chain - Homo sapiens (Human)					
Q9Y285	2	5.2631	K.SLQALGEVIEAELR.S	4	SYFA_HUMAN
Phosphatase and actin regulator 2 - Homo sapiens (Human)					
O75167	2	3.2277	R.TTLYSGTGLSVNR.E	1	PHAR2_HUMAN
O75167	3	4.2387	R.LSQRPTTEELEQR.N	2	PHAR2_HUMAN
Phosphatase and actin regulator 4 - Homo sapiens (Human)					
Q8IZ21	3	4.0001	R.IQQALTSPLPMTPILEGSHR.A	1	PHAR4_HUMAN
Q8IZ21	3	6.1862	R.AHSLLFENSDSFSEDSSTLGR.T	2	PHAR4_HUMAN
Phosphatidylethanolamine-binding protein 1 - Homo sapiens (Human)					
P30086	3	4.6218	K.NRPTSISWDGLDSGKLYTLVLTDPDAPSR.K	1	PEBP1_HUMAN
P30086	2	4.3648	R.YVWLVYEQDRPLK.C	4	PEBP1_HUMAN
P30086	3	4.8705	R.APVAGTCYQAEWDDYVPKLYEQLSGK.-	1	PEBP1_HUMAN
P30086	4	7.1379	K.WSGPLSLQEVDEQPQHPLHVTYAGAAVDEL	7	PEBP1_HUMAN
P30086	3	4.9374	K.NRPTSISWDGLDSGK.L	6	PEBP1_HUMAN
P30086	2	3.9985	K.LYTLVLTDPDAPSRK.D	1	PEBP1_HUMAN
P30086	2	4.024	K.LYTLVLTDPDAPSR.K	2	PEBP1_HUMAN
P30086	1	2.8961	K.LYEQLSGK.-	4	PEBP1_HUMAN
P30086	3	3.7334	K.GNDISSGTVLSDYVGSGPPKGTGLHR.Y	1	PEBP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P30086	2	4.7679	K.GNDISSGTVLSDYVGSGPPK.G	2	PEBP1_HUMAN
P30086	2	4.4026	R.APVAGTCYQAEWDDYVPK.L	1	PEBP1_HUMAN
Phosphatidylinositol-4-phosphate 3-kinase C2 domain-containing beta polypeptide - Homo sapiens					
O00750	3	4.6756	K.ISQPSDINTFSLVEQLPGK.L	1	P3C2B_HUMAN
Phosphatidylinositol-4-phosphate 5-kinase type-2 alpha - Homo sapiens (Human)					
P48426	2	2.8648	R.FLDFIGHILT.-	2	PI52A_HUMAN
Phosphofurin acidic cluster sorting protein 1 - Homo sapiens (Human)					
Q6VY07	3	4.3518	K.TDLQGSASPSKVEGVHTPR.Q	1	PACS1_HUMAN
Q6VY07	3	4.53	R.GSGVAQSPQQPPPQQQQQQPPQQPTPPK.L	1	PACS1_HUMAN
Q6VY07	2	4.3339	K.VSDEVGFGLEHVSR.E	2	PACS1_HUMAN
Q6VY07	2	3.3464	K.TDLQGSASPSK.V	2	PACS1_HUMAN
Q6VY07	2	4.2921	K.GSLGKDTTSPMELAALEK.I	2	PACS1_HUMAN
Q6VY07	2	3.5531	K.GSLGKDTTSPM#ELAALEK.I	1	PACS1_HUMAN
Q6VY07	2	3.2824	K.DTTSPM#ELAALEK.I	3	PACS1_HUMAN
Q6VY07	3	4.7964	R.FKVSDEVGFGLEHVSR.E	2	PACS1_HUMAN
Phosphofurin acidic cluster sorting protein 2 - Homo sapiens (Human)					
Q86VP3	3	5.0168	R.EHPGQPEDSPEAEASTLDVFTER.L	1	PACS2_HUMAN
Q86VP3	3	3.7912	R.SHKEPPSPADVPEK.T	1	PACS2_HUMAN
Phosphoglycerate kinase 1 - Homo sapiens (Human)					
P00558	2	3.0315	K.VLNNMEIGTSLFDEEGAK.I	1	PGK1_HUMAN
P00558	1	2.1443	K.YAEAVTR.A	1	PGK1_HUMAN
P00558	2	3.4696	K.NNQITNNQR.I	3	PGK1_HUMAN
P00558	3	4.2833	R.AHSSMVGVNLPQK.A	1	PGK1_HUMAN
Phospholipase C X domain-containing protein 2 - Homo sapiens (Human)					
Q0VAA5	2	2.7608	R.IQEAFGNKLCPACSVESLTLR.T	1	Q0VAA5_HUMA
Phospholipase C, beta 4 - Homo sapiens (Human)					
Q5JYT3	2	3.0854	R.DGPQTSNSSMK.L	2	Q5JYT3_HUMAN
Phosphoprotein associated with glycosphingolipid-enriched microdomains 1 - Homo sapiens (Hum					
Q9NWQ8	2	4.1684	K.SGQSLTVPESTYTSIQGDPQR.S	2	PAG1_HUMAN
Q9NWQ8	3	5.7476	K.SREEDPTLTEEEISAMYSSVNKPGQLVNK.S	1	PAG1_HUMAN
Pigment epithelium-derived factor precursor - Homo sapiens (Human)					
P36955	2	4.4175	K.LAAAVSNFGYDLYR.V	2	PEDF_HUMAN
Pinin - Homo sapiens (Human)					
Q9H307	2	3.1877	R.DLIQDQNMDEK.G	2	PININ_HUMAN
Q9H307	2	3.9011	R.DLIQDQNMDEKGK.Q	2	PININ_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9H307	1	2.9817	R.TLQEQLEK.A	3	PININ_HUMAN
Q9H307	2	2.9585	R.DLIQDQNM#DEK.G	1	PININ_HUMAN
Pitrilysin metallopeptidase 1 - Homo sapiens (Human)					
Q5JRW8	2	2.8346	K.RPVC*GSGDLGLISPR.A	2	Q5JRW8_HUMA
Plakophilin-4 - Homo sapiens (Human)					
Q99569	3	3.8001	R.QEAASTGPGMEPETTATTILASVKEQELQFQ	1	PKP4_HUMAN
Plasma membrane calcium-transporting ATPase 4 - Homo sapiens (Human)					
P23634	3	4.5443	R.TPLLDEEEEENPDKASK.F	2	AT2B4_HUMAN
Plasma serine protease inhibitor precursor - Homo sapiens (Human)					
P05154	2	4.1833	K.AVVEVDESGTR.A	5	IPSP_HUMAN
P05154	2	4.1374	R.AAAATGTIFTFR.S	3	IPSP_HUMAN
Plasmalemma vesicle-associated protein - Homo sapiens (Human)					
Q9BX97	3	4.3288	R.PMGPVPNPQPIDPASLEEFKR.K	2	PLVAP_HUMAN
Plasminogen activator inhibitor 1 RNA-binding protein - Homo sapiens (Human)					
Q8NC51	2	3.2375	R.TDKSSASAPDVDDPEAFPALA.-	1	PAIRB_HUMAN
Q8NC51	2	5.5292	R.GGSGSHNWGTVKDELTESPK.Y	5	PAIRB_HUMAN
Q8NC51	3	4.4495	R.KNPLPPSVGVVDKK.E	2	PAIRB_HUMAN
Q8NC51	2	3.5469	R.KPNEGADGQWK.K	7	PAIRB_HUMAN
Q8NC51	2	3.5523	R.PDQQLQGEGK.I	2	PAIRB_HUMAN
Q8NC51	2	2.9152	R.RFEKPLEEK.G	1	PAIRB_HUMAN
Q8NC51	2	2.8973	R.SSFSHYSGLK.H	2	PAIRB_HUMAN
Q8NC51	2	5.2238	R.KPANDITSQLEINFGDLGR.P	4	PAIRB_HUMAN
Q8NC51	2	3.1958	R.FDQLFDDESDPFEVLK.A	4	PAIRB_HUMAN
Q8NC51	2	4.1674	R.RPDQQLQGEGK.I	5	PAIRB_HUMAN
Q8NC51	3	4.6038	K.SKSEEAHAEDSVM#DHHFR.K	2	PAIRB_HUMAN
Q8NC51	3	5.1206	K.SEEAHAEDSVMDHHFR.K	4	PAIRB_HUMAN
Q8NC51	2	5.0516	K.SKSEEAHAEDSVMDHHFR.K	4	PAIRB_HUMAN
Q8NC51	2	5.2163	K.SAAQAAAQTNSNAAGK.Q	9	PAIRB_HUMAN
Q8NC51	2	4.8495	K.KKEAGGGGVGGPGAK.S	11	PAIRB_HUMAN
Q8NC51	1	2.402	K.KGFVLHK.S	2	PAIRB_HUMAN
Q8NC51	3	4.0152	K.KEETQPPVALKK.E	3	PAIRB_HUMAN
Q8NC51	1	2.3464	K.EMTLDEWK.A	1	PAIRB_HUMAN
Q8NC51	2	2.78	K.EETQPPVALKK.E	1	PAIRB_HUMAN
Q8NC51	2	3.2895	K.EAGGGGVGGPGAK.S	9	PAIRB_HUMAN
Q8NC51	2	4.2565	R.KPNEGADGQWKK.G	5	PAIRB_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Plasminogen precursor - Homo sapiens (Human)					
P00747	2	3.5964	K.TPENYPNAGLTM#NYCR.N	1	PLMN_HUMAN
P00747	2	3.5525	R.NPDGDVGGPWCYTTNPR.K	1	PLMN_HUMAN
P00747	2	4.4977	K.TPENYPNAGLTMNYCR.N	2	PLMN_HUMAN
P00747	2	3.3955	R.HSIFTPETNPR.A	2	PLMN_HUMAN
Platelet basic protein precursor - Homo sapiens (Human)					
P02775	2	6.0096	K.GKEESLDSDLYAELR.C	2	SCYB7_HUMAN
P02775	2	2.8032	K.ICLDPDAPR.I	2	SCYB7_HUMAN
P02775	1	3.5889	K.KLAGDESAD.-	12	SCYB7_HUMAN
P02775	1	2.2146	K.LAGDESAD.-	1	SCYB7_HUMAN
P02775	2	2.9205	K.NIQSLEVIGK.G	6	SCYB7_HUMAN
Platelet endothelial cell adhesion molecule precursor - Homo sapiens (Human)					
P16284	2	4.5588	K.DTETVYSEVR.K	7	PECA1_HUMAN
P16284	1	2.1448	R.TEGSLDGT.-	1	PECA1_HUMAN
P16284	2	3.1029	R.YSRTEGSLDGT.-	2	PECA1_HUMAN
P16284	2	3.3817	R.KAVPDAVESR.Y	4	PECA1_HUMAN
P16284	2	3.693	K.DTETVYSEVRK.A	3	PECA1_HUMAN
P16284	1	2.365	K.AVPDAVESR.Y	2	PECA1_HUMAN
P16284	2	3.2459	K.STESYFIPEVR.I	1	PECA1_HUMAN
P16284	3	4.4683	K.KDTETVYSEVRK.A	2	PECA1_HUMAN
Platelet-activating factor acetylhydrolase IB subunit beta - Homo sapiens (Human)					
P68402	2	3.5562	K.IIVLGLLPR.G	4	PA1B2_HUMAN
P68402	2	3.4179	R.ELFSPLHALNFGIGGDTTR.H	1	PA1B2_HUMAN
Pleckstrin homology domain-containing family A member 1 - Homo sapiens (Human)					
Q9HB21	2	3.6511	K.VTEQALLRPQSK.N	2	PKHA1_HUMAN
Q9HB21	2	3.3335	R.SNSLVSTFTMEK.R	2	PKHA1_HUMAN
Pleckstrin homology domain-containing family A member 2 - Homo sapiens (Human)					
Q9HB19	3	4.0423	K.EKPFM#FNLDDENIRTSDV.-	1	PKHA2_HUMAN
Pleckstrin homology domain-containing family A member 5 - Homo sapiens (Human)					
Q9HAU0	2	4.023	R.TNSM#QQLEQWIK.I	3	PKHA5_HUMAN
Q9HAU0	2	4.3609	R.TNSMQQLEQWIK.I	4	PKHA5_HUMAN
Q9HAU0	2	3.7299	R.GTTEIGM#IGSKPFSTVK.Y	2	PKHA5_HUMAN
Q9HAU0	2	4.4315	R.GTTEIGMIGSKPFSTVK.Y	3	PKHA5_HUMAN
Q9HAU0	2	2.8122	R.GVISYQTLPR.N	1	PKHA5_HUMAN
Q9HAU0	2	3.1308	R.IQDVMEGLSK.H	1	PKHA5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9HAU0	2	2.803	R.MTVEEQM#ER.I	1	PKHA5_HUMAN
Q9HAU0	2	4.9841	R.NQMQEQLDHLGEVQTESAGIQR.A	1	PKHA5_HUMAN
Q9HAU0	2	2.789	R.SVPAGLTLQSVSPQSLQGK.T	1	PKHA5_HUMAN
Q9HAU0	3	4.3304	R.GGNRPNTGPLYTEADR.V	3	PKHA5_HUMAN
Q9HAU0	3	4.7779	R.RSVPAGLTLQSVSPQSLQGK.T	2	PKHA5_HUMAN
Q9HAU0	2	2.894	K.DGQDRPLTK.I	1	PKHA5_HUMAN
Q9HAU0	2	2.7699	R.GNQTM#AVK.S	1	PKHA5_HUMAN
Q9HAU0	2	2.9571	R.GNQTMAVK.S	2	PKHA5_HUMAN
Q9HAU0	2	3.0235	K.GLNVIGASDQSPLQSPSNLR.D	1	PKHA5_HUMAN
Q9HAU0	2	4.5746	K.IVNVSLADLR.G	2	PKHA5_HUMAN
Q9HAU0	3	3.7218	K.KGLNVIGASDQSPLQSPSNLR.D	1	PKHA5_HUMAN
Q9HAU0	3	4.8278	K.KKGLNVIGASDQSPLQSPSNLR.D	1	PKHA5_HUMAN
Q9HAU0	2	3.5576	K.TLSQDEGRGTLYK.Y	2	PKHA5_HUMAN
Q9HAU0	3	5.9007	K.TMVNISDQTMHSIPTSPSHGSIAAYQGYSPQ	2	PKHA5_HUMAN
Q9HAU0	3	3.842	K.TRPESICSVTPSTHDK.T	2	PKHA5_HUMAN
Q9HAU0	2	3.2871	K.YRPEEVDIDAK.L	2	PKHA5_HUMAN
Pleckstrin homology domain-containing family A member 7 - Homo sapiens (Human)					
Q6IQ23	2	3.7361	R.VFFINDQLR.C	2	PKHA7_HUMAN
Q6IQ23	2	3.6128	R.SPYSPAEEDALFMDLPTGPR.G	2	PKHA7_HUMAN
Pleckstrin homology domain-containing family F member 2 - Homo sapiens (Human)					
Q9H8W4	2	2.7126	R.LANSEANTR.R	1	PKHF2_HUMAN
Pleckstrin homology domain-containing family G member 1 - Homo sapiens (Human)					
Q9ULL1	2	3.1407	K.TSETAQDIQK.V	1	PKHG1_HUMAN
Pleckstrin homology-like domain family B member 1 - Homo sapiens (Human)					
Q86UU1	2	5.2747	R.GLLTDSPAATVLAEAR.R	2	PHLB1_HUMAN
Q86UU1	3	5.1284	R.ADGGPEAGELPSIGEATAALALAGR.R	3	PHLB1_HUMAN
Q86UU1	2	3.9208	R.DKEAEALETETK.L	2	PHLB1_HUMAN
Q86UU1	4	5.0634	R.DKEAEALETETKLFEDLEFQQLER.E	1	PHLB1_HUMAN
Pleckstrin homology-like domain family B member 2 - Homo sapiens (Human)					
Q86SQ0	2	3.0886	R.LQEETSQR.Q	1	PHLB2_HUMAN
Q86SQ0	3	3.9492	R.GYNHQQMSEGHR.Q	1	PHLB2_HUMAN
Q86SQ0	2	2.9245	K.EGLYLSDTLPR.K	1	PHLB2_HUMAN
Q86SQ0	2	3.7442	K.RLQEETSQR.Q	4	PHLB2_HUMAN
Q86SQ0	2	3.3988	K.NNLIMMLQR.E	1	PHLB2_HUMAN
Pleiotropic regulator 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43660	2	3.2206	K.GPQNATDSYVHK.Q	1	PLRG1_HUMAN
O43660	3	4.7893	K.LRNEYGPVLHMPTSK.E	1	PLRG1_HUMAN
O43660	2	5.0553	R.MPSESAAQSLAVALPLQTK.A	3	PLRG1_HUMAN
PLEKHG2 protein - Homo sapiens (Human)					
Q96D18	2	4.5905	R.PGFPEPLLILEDSDLGGDSGSGK.A	1	Q96D18_HUMAN
Plexin domain-containing protein 2 precursor - Homo sapiens (Human)					
Q6UX71	2	2.7294	R.ASVGQDSPEPR.S	1	PXDC2_HUMAN
Plexin-A2 precursor - Homo sapiens (Human)					
O75051	2	2.8405	R.RDKC*QQAWEPNR.F	1	PLXA2_HUMAN
PML-RARA-regulated adapter molecule 1 - Homo sapiens (Human)					
Q96QH2	4	5.1196	K.KSVPQPEFSEAAQTPLWKPQSSEPK.R	1	PRAM_HUMAN
Q96QH2	3	4.1977	K.KPPQPELGGLPR.T	1	PRAM_HUMAN
Q96QH2	2	3.1823	K.FQASQPEPSDLPK.K	2	PRAM_HUMAN
Q96QH2	3	3.9625	K.KPAQPEFNVYPK.K	2	PRAM_HUMAN
Poliovirus receptor precursor - Homo sapiens (Human)					
P15151	2	4.4826	R.ENSSSQDPQTEGTR.-	2	PVR_HUMAN
Poliovirus receptor-related protein 1 precursor - Homo sapiens (Human)					
Q15223	2	2.7595	K.ITQVTWQK.S	1	PVRL1_HUMAN
Q15223	2	2.9871	K.QNVAIYNPSMGVSVLAPYR.E	1	PVRL1_HUMAN
Q15223	3	4.3269	R.TLGYQYDPEQLDLAENMVSQNDGSFISKK.E	1	PVRL1_HUMAN
Poly [ADP-ribose] polymerase 1 - Homo sapiens (Human)					
P09874	2	2.8839	K.GQVKEEGINK.S	1	PARP1_HUMAN
P09874	2	3.2138	K.TLGDFAAEYAK.S	2	PARP1_HUMAN
P09874	3	4.1562	K.GQVKEEGINKSEK.R	2	PARP1_HUMAN
Poly [ADP-ribose] polymerase 14 - Homo sapiens (Human)					
Q460N5	2	2.8605	K.ATLPDTAAPPGLPPAAAGPGKTSWEK.G	1	PAR14_HUMAN
Q460N5	2	3.3359	R.FLVFFYPEDVR.Q	2	PAR14_HUMAN
Poly(A) polymerase alpha - Homo sapiens (Human)					
P51003	3	3.7942	K.TSSTDLSDIPALPANPIPVIK.N	2	PAPOA_HUMAN
Poly(rC)-binding protein 1 - Homo sapiens (Human)					
Q15365	2	3.4647	K.IANPVEGSSGR.Q	4	PCBP1_HUMAN
Polyadenylate-binding protein 1 - Homo sapiens (Human)					
P11940	3	4.4359	R.VANTSTQTM#GPRPAAAAAAATPAVR.T	4	PABP1_HUMAN
P11940	3	4.9289	R.VANTSTQTMGPRPAAAAAAATPAVR.T	2	PABP1_HUMAN
Polyadenylate-binding protein 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q86U42	2	3.5612	K.GFAYIEFSDKESVR.T	2	PABP2_HUMAN
Q86U42	4	6.7798	R.HLVPGAGGEAGEGAPGGAGDYGNGLESEEL	1	PABP2_HUMAN
Q86U42	2	3.9254	R.TSLALDESLFR.G	1	PABP2_HUMAN
Q86U42	2	3.9245	R.VREMEEEAEK.L	4	PABP2_HUMAN
Polycystin-2 - Homo sapiens (Human)					
Q13563	2	3.5263	R.SAASSAVSSVGAR.S	2	PKD2_HUMAN
Q13563	2	4.9797	R.DPPAGAAASPSPPLSSCSR.Q	4	PKD2_HUMAN
Polyglutamine-binding protein 1 - Homo sapiens (Human)					
O60828	2	2.7093	R.REELAPYPK.S	1	PQBP1_HUMAN
O60828	3	4.4934	R.KDEELDPM#DPSSYSDAPR.G	2	PQBP1_HUMAN
O60828	3	5.561	R.KDEELDPMDPSSYSDAPR.G	5	PQBP1_HUMAN
Polymerase I and transcript release factor - Homo sapiens (Human)					
Q6NZI2	3	4.5346	K.TAVYKVPPFTFHVK.K	7	PTRF_HUMAN
Q6NZI2	2	3.5285	R.KSFTPDHVVYAR.S	2	PTRF_HUMAN
Q6NZI2	1	2.8196	R.EGQVEVLK.A	8	PTRF_HUMAN
Q6NZI2	2	5.4266	K.ATEMVEVGADDDEGGAER.G	5	PTRF_HUMAN
Q6NZI2	1	2.135	K.VSVNVK.T	1	PTRF_HUMAN
Q6NZI2	2	3.0589	R.LKTKENLEK.T	3	PTRF_HUMAN
Q6NZI2	2	2.9962	K.VPPFTFHVK.K	3	PTRF_HUMAN
Q6NZI2	2	4.6556	K.VMIYQDEVKLPAK.L	5	PTRF_HUMAN
Q6NZI2	2	2.8766	K.VMIYQDEVK.L	3	PTRF_HUMAN
Q6NZI2	2	4.2932	K.VM#IYQDEVKLPAK.L	6	PTRF_HUMAN
Q6NZI2	2	3.8069	K.TAVYKVPPFTFHVKK.I	2	PTRF_HUMAN
Q6NZI2	2	2.9744	K.SFTPDHVVYAR.S	1	PTRF_HUMAN
Q6NZI2	2	4.2205	K.LGKAHATTSNTVSK.L	1	PTRF_HUMAN
Q6NZI2	2	3.9811	K.LEVNEAELLR.R	4	PTRF_HUMAN
Q6NZI2	2	3.2166	K.KLEVNEAELLRR.R	1	PTRF_HUMAN
Q6NZI2	2	4.5524	K.KLEVNEAELLR.R	7	PTRF_HUMAN
Q6NZI2	3	4.1257	K.IREGQVEVLK.A	4	PTRF_HUMAN
Q6NZI2	3	6.7875	K.IIGAVDQIQLTQAQLEERQAEM#EGAVQSIQG	1	PTRF_HUMAN
Q6NZI2	3	5.4037	K.ATEMVEVGADDDEGGAERGEAGDLR.R	39	PTRF_HUMAN
Q6NZI2	3	4.2967	K.ATEM#VEVGADDDEGGAERGEAGDLR.R	6	PTRF_HUMAN
Q6NZI2	2	4.7552	K.ATEM#VEVGADDDEGGAER.G	9	PTRF_HUMAN
Q6NZI2	2	5.0631	R.QAEMEGAVQSIQGELSK.L	2	PTRF_HUMAN
Q6NZI2	3	5.4839	K.IIGAVDQIQLTQAQLEER.Q	2	PTRF_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Polymeric-immunoglobulin receptor precursor - Homo sapiens (Human)					
P01833	2	5.0782	R.ASVDSGSSEEQGGSSR.A	1	PIGR_HUMAN
Polypeptide N-acetylgalactosaminyltransferase 6 - Homo sapiens (Human)					
Q8NCL4	3	4.0119	R.VWQCGGQLEIIPC*SVVGHVFRTK.S	1	GALT6_HUMAN
Porphobilinogen deaminase - Homo sapiens (Human)					
P08397	3	5.1212	R.GPQLAAQNLGISLANLLLSK.G	1	HEM3_HUMAN
Potassium channel modulatory factor 1 - Homo sapiens (Human)					
Q9BWK2	2	2.8758	R.DLDESSGVR.H	1	Q9BWK2_HUMA
PRA1 family protein 3 - Homo sapiens (Human)					
O75915	2	4.2875	R.TPMGIVLDALEQQEEGINR.L	2	PRAF3_HUMAN
Prefoldin subunit 1 - Homo sapiens (Human)					
O60925	1	2.4843	K.AFTELQAK.V	3	PFD1_HUMAN
O60925	1	3.1085	K.EAIHSQLLEK.Q	6	PFD1_HUMAN
O60925	2	3.0407	K.IAEEKIKELEQK.K	1	PFD1_HUMAN
O60925	2	4.0963	K.LADIQIEQLNR.T	5	PFD1_HUMAN
O60925	3	4.993	K.VKLADIQIEQLNR.T	2	PFD1_HUMAN
O60925	2	2.9078	R.MFILQSK.E	1	PFD1_HUMAN
Prefoldin subunit 2 - Homo sapiens (Human)					
Q9UHV9	3	4.2081	R.LMGEDEKPAAKENSEGAGAK.A	1	PFD2_HUMAN
Q9UHV9	2	3.0704	R.M#VGGVLVER.T	3	PFD2_HUMAN
Q9UHV9	2	3.7487	R.LMGEDEKPAAK.E	2	PFD2_HUMAN
Q9UHV9	2	3.0183	R.LM#GEDEKPAAK.E	2	PFD2_HUMAN
Q9UHV9	2	3.0574	K.GKELNEFREK.H	2	PFD2_HUMAN
Q9UHV9	2	4.0385	K.EVLPALENNKEQIQK.I	3	PFD2_HUMAN
Q9UHV9	2	4.3255	K.GAVSAEQVIAGFNR.L	6	PFD2_HUMAN
Q9UHV9	2	3.9362	R.MVGGVLVER.T	6	PFD2_HUMAN
Q9UHV9	2	5.1111	K.IIETLTQQLQAK.G	4	PFD2_HUMAN
Prefoldin subunit 3 - Homo sapiens (Human)					
P61758	2	3.2343	K.KKESTNSMETR.F	1	PFD3_HUMAN
P61758	2	3.4271	K.KLDEQYQK.Y	4	PFD3_HUMAN
Prefoldin subunit 4 - Homo sapiens (Human)					
Q9NQP4	2	4.2864	K.NLQEEIDALESR.V	6	PFD4_HUMAN
Q9NQP4	2	3.8811	R.ITELKEEIEVK.K	3	PFD4_HUMAN
Q9NQP4	2	3.7169	K.NLQEEIDALESRVESIQR.V	2	PFD4_HUMAN
Q9NQP4	2	5.1032	K.KAAAEDVNVTFEDQQK.I	2	PFD4_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9NQP4	2	4.351	K.AAAEDVNVTFEDQQKINK.F	1	PFD4_HUMAN
Q9NQP4	2	4.5229	K.AAAEDVNVTFEDQQK.I	5	PFD4_HUMAN
Q9NQP4	1	2.1625	K.VQLYAK.F	1	PFD4_HUMAN
Prefoldin subunit 6 - Homo sapiens (Human)					
O15212	2	2.969	K.RLDYITAEIK.R	3	PFD6_HUMAN
O15212	3	4.5308	K.RLDYITAEIKR.Y	1	PFD6_HUMAN
O15212	2	2.7495	K.RYESQLR.D	1	PFD6_HUMAN
O15212	3	4.6625	R.ETLAQLQQEFQR.A	5	PFD6_HUMAN
O15212	3	4.2521	R.QSEQQRETLAQLQQEFQR.A	3	PFD6_HUMAN
Pre-mRNA 3'-end-processing factor FIP1 - Homo sapiens (Human)					
Q6UN15	3	4.3583	R.KPGADLSDYFNYGFNEDTWK.A	1	FIP1_HUMAN
Q6UN15	2	3.5849	R.M#GLEVIPVTSTTNK.I	2	FIP1_HUMAN
Q6UN15	2	5.002	R.TGNSEKETALPSTK.A	4	FIP1_HUMAN
Q6UN15	2	5.0189	K.TGAPQYGSYGTAPVNLNIK.T	5	FIP1_HUMAN
Q6UN15	2	3.3572	K.AEFTSPPSLFK.T	2	FIP1_HUMAN
Q6UN15	2	3.7736	R.ANENSNIQVLSER.S	3	FIP1_HUMAN
Pre-mRNA cleavage complex 2 protein Pcf11 - Homo sapiens (Human)					
O94913	1	2.3766	K.ELDQLDSK.S	2	PCF11_HUMAN
O94913	3	4.4069	K.TEEERPQETTNQHSTK.S	1	PCF11_HUMAN
Pre-mRNA-processing factor 40 homolog A - Homo sapiens (Human)					
O75400	3	4.3363	R.ALEKEEEEEKQK.S	2	PRP40_HUMAN
Pre-mRNA-processing factor 6 - Homo sapiens (Human)					
O94906	2	2.7882	R.DANDPVDDR.H	3	PRP6_HUMAN
Pre-mRNA-splicing factor ATP-dependent RNA helicase PRP16 - Homo sapiens (Human)					
Q92620	2	3.2492	R.RNEPESPR.H	5	PRP16_HUMAN
Q92620	3	4.1005	R.VETPSHPGGVSEEFWER.S	1	PRP16_HUMAN
Q92620	2	4.2506	R.SQWESPSPTPSYR.D	2	PRP16_HUMAN
Q92620	2	2.8489	K.SAASEQHVFK.A	2	PRP16_HUMAN
Q92620	3	5.2109	K.APAPRPSLLGLDLLASLK.R	5	PRP16_HUMAN
Q92620	2	3.433	R.STWEEEDSGYGSSR.R	1	PRP16_HUMAN
Pre-mRNA-splicing factor ISY1 homolog - Homo sapiens (Human)					
Q9ULR0	2	4.5216	K.VAQIQNAGLGEFR.I	3	ISY1_HUMAN
Pre-mRNA-splicing factor RBM22 - Homo sapiens (Human)					
Q9NW64	2	3.2855	R.EISNSDGTRPVGMLGK.A	1	RBM22_HUMAN
Pre-mRNA-splicing factor SLU7 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O95391	2	3.6087	R.QRPDDPMASFLGQ.-	1	SLU7_HUMAN
Pre-mRNA-splicing factor SYF2 - Homo sapiens (Human)					
O95926	3	4.0565	R.KLNHQEVVEEDKR.L	1	SYF2_HUMAN
O95926	3	3.7961	R.LKLPANWEAK.K	1	SYF2_HUMAN
O95926	3	3.8473	R.RRPYNDDADIDYINER.N	1	SYF2_HUMAN
Pre-rRNA-processing protein TSR2 homolog - Homo sapiens (Human)					
Q969E8	1	2.1247	K.VTATALK.T	1	TSR2_HUMAN
Q969E8	2	4.245	K.WLGGAVEDYFMR.N	5	TSR2_HUMAN
Q969E8	2	4.0959	K.WLGGAVEDYFM#R.N	3	TSR2_HUMAN
PRKC apoptosis WT1 regulator protein - Homo sapiens (Human)					
Q96IZ0	2	4.3774	R.LMQDKEEMIGK.L	2	PAWR_HUMAN
Q96IZ0	3	4.361	R.YNRDANVSGTLVSSSTLEK.K	1	PAWR_HUMAN
Q96IZ0	2	4.9138	R.YKSTTSVSEEDVSSR.Y	1	PAWR_HUMAN
Q96IZ0	3	4.8855	R.RSEDEPPAASASAAPPPQRDEEEPDGVPEK.	1	PAWR_HUMAN
Q96IZ0	2	3.5967	R.LMQDKEEM#IGK.L	1	PAWR_HUMAN
Q96IZ0	2	3.7833	R.LM#QDKEEMIGK.L	1	PAWR_HUMAN
Q96IZ0	3	5.2555	R.KREDAITQQNTIQNEAVNLLDPGSSYLLQEPP	2	PAWR_HUMAN
Q96IZ0	2	3.6562	R.DLDDIEDENEQLKQENK.T	2	PAWR_HUMAN
Q96IZ0	2	5.8277	R.DANVSGTLVSSSTLEK.K	10	PAWR_HUMAN
Q96IZ0	2	3.5072	K.STTSVSEEDVSSR.Y	3	PAWR_HUMAN
Q96IZ0	2	3.8135	K.KIEDLEKEVVR.E	2	PAWR_HUMAN
Q96IZ0	3	4.1994	R.SEDEPPAASASAAPPPQRDEEEPDGVPEK.G	2	PAWR_HUMAN
PRKR interacting protein 1 - Homo sapiens (Human)					
Q96CF8	2	2.9049	K.IAAEEQTAK.R	1	Q96CF8_HUMA
Q96CF8	2	3.5825	K.IAAEEQTAKR.R	2	Q96CF8_HUMA
Probable asparaginyl-tRNA synthetase, mitochondrial precursor - Homo sapiens (Human)					
Q96I59	2	3.5497	R.DALGAQNASGER.I	2	SYNM_HUMAN
Probable ATP-dependent RNA helicase DDX10 - Homo sapiens (Human)					
Q13206	2	3.8248	R.SNSEVEDVGPTSHNR.K	1	DDX10_HUMAN
Probable ATP-dependent RNA helicase DDX17 - Homo sapiens (Human)					
Q92841	2	3.3893	K.GTAYTFFTPGNLK.Q	2	DDX17_HUMAN
Q92841	2	3.4446	K.VLEEANQAINPK.L	1	DDX17_HUMAN
Probable ATP-dependent RNA helicase DDX27 - Homo sapiens (Human)					
Q96GQ7	2	2.9744	R.AATTLDEKIEK.V	1	DDX27_HUMAN
Probable ATP-dependent RNA helicase DDX49 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y6V7	2	3.0602	R.TPSGSHSGPVPSQGLV.-	2	DDX49_HUMAN
Probable ATP-dependent RNA helicase DDX6 - Homo sapiens (Human)					
P26196	3	6.2287	R.GPVKPTGGPGGGGTQTQQQM#NQLK.N	2	DDX6_HUMAN
P26196	3	5.2788	R.GPVKPTGGPGGGGTQTQQQMNQLK.N	1	DDX6_HUMAN
P26196	2	4.2345	R.TENPVIM#GLSSQNGQLR.G	1	DDX6_HUMAN
P26196	2	4.8853	R.TENPVIMGLSSQNGQLR.G	3	DDX6_HUMAN
P26196	4	5.2081	K.NTNTINNGTQQQAQSM#TTTIKPGDDWKK.T	1	DDX6_HUMAN
Probable D-tyrosyl-tRNA(Tyr) deacylase 1 - Homo sapiens (Human)					
Q8TEA8	2	4.4577	R.SASSGAEGDVSSEREP.-	7	DTD1_HUMAN
Probable G-protein coupled receptor 124 precursor - Homo sapiens (Human)					
Q96PE1	1	2.4217	R.DSLKGGGALEK.E	2	GP124_HUMAN
Probable helicase senataxin - Homo sapiens (Human)					
Q7Z333	2	2.7859	R.SLDYVAQLR.D	1	SETX_HUMAN
Q7Z333	2	3.3382	K.SLETSSALSPSLK.N	1	SETX_HUMAN
Q7Z333	2	3.24	R.LLTDSSTDALEK.V	1	SETX_HUMAN
Probable helicase with zinc finger domain - Homo sapiens (Human)					
P42694	3	3.7146	R.AIAQPGPILPSHLNSFIDENPSGLPIGEALDR.I	1	HELZ_HUMAN
Probable histone acetyltransferase MYST1 - Homo sapiens (Human)					
Q9H7Z6	3	4.6512	R.RPDSTWHSAEVIQSR.V	1	MYST1_HUMAN
Probable palmitoyltransferase ZDHHC5 - Homo sapiens (Human)					
Q9C0B5	2	5.1634	K.SAQGTGFELGQLQSIR.S	4	ZDHC5_HUMAN
Q9C0B5	2	3.4622	K.SLGSASPGPGQPPLSSPTR.G	2	ZDHC5_HUMAN
Probable RNA-binding protein 25 - Homo sapiens (Human)					
P49756	2	4.5323	R.DREEDEEDAYER.R	2	RBM25_HUMAN
Probable rRNA-processing protein EBP2 - Homo sapiens (Human)					
Q99848	2	2.9127	K.KAVNDVNGLK.Q	1	EBP2_HUMAN
Q99848	2	3.5584	R.QAQAAVLAVLPR.L	2	EBP2_HUMAN
Q99848	2	3.215	K.KVQTEVLQK.R	2	EBP2_HUMAN
Profilin-1 - Homo sapiens (Human)					
P07737	2	2.9204	K.TLVLLMGK.E	3	PROF1_HUMAN
P07737	2	4.4214	R.DSLLQDGEFSM#DLR.T	3	PROF1_HUMAN
P07737	2	5.3726	K.TFVNITPAEVGVLVGKDR.S	5	PROF1_HUMAN
P07737	2	4.464	K.TFVNITPAEVGVLVGK.D	1	PROF1_HUMAN
P07737	2	4.0807	K.TDKTLVLLMGK.E	3	PROF1_HUMAN
P07737	2	3.3027	K.STGGAPTFNVTVTK.T	4	PROF1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07737	2	2.8794	K.EGVHGGLINKK.C	1	PROF1_HUMAN
P07737	2	3.434	K.DSPSVWAAVPGK.T	2	PROF1_HUMAN
P07737	2	4.1609	R.SSFYVNGLTLGGQK.C	2	PROF1_HUMAN
Programmed cell death protein 10 - Homo sapiens (Human)					
Q9BUL8	2	3.3437	R.VNLSAAQTLR.A	2	PDC10_HUMAN
Programmed cell death protein 5 - Homo sapiens (Human)					
O14737	2	2.9381	R.KVMDSDEDDDY.-	1	PDCD5_HUMAN
O14737	2	2.7227	R.RKVMDSDEDDDY.-	1	PDCD5_HUMAN
O14737	1	2.9461	R.YGQLSEK.V	11	PDCD5_HUMAN
O14737	3	4.6627	R.YGQLSEKVSEQGLIEILKK.V	2	PDCD5_HUMAN
O14737	2	5.1859	R.NSILAQVLDQSAR.A	7	PDCD5_HUMAN
O14737	2	3.863	R.LSNLALVKPEK.T	4	PDCD5_HUMAN
O14737	2	4.2403	K.VSEQGLIEILKK.V	6	PDCD5_HUMAN
O14737	2	4.2151	K.VSEQGLIEILK.K	4	PDCD5_HUMAN
O14737	3	4.6124	K.TKAVENYLIQMAR.Y	5	PDCD5_HUMAN
O14737	3	4.0556	K.TKAVENYLIQM#AR.Y	1	PDCD5_HUMAN
O14737	2	3.7031	K.AVENYLIQM#AR.Y	4	PDCD5_HUMAN
O14737	3	5.8287	R.YGQLSEKVSEQGLIEILK.K	3	PDCD5_HUMAN
O14737	3	4.9532	R.ARLSNLALVKPEK.T	1	PDCD5_HUMAN
Proline-rich acidic protein 1 - Homo sapiens (Human)					
Q5VWY4	3	5.0508	R.VVEPPEKDDQLVVLFPVQKPK.L	1	Q5VWY4_HUMA
Proline-rich AKT1 substrate 1 - Homo sapiens (Human)					
Q96B36	2	2.817	K.SLPVSVPVWGFK.E	1	AKTS1_HUMAN
Proline-rich protein 8 - Homo sapiens (Human)					
Q9NSV0	2	4.4606	K.NQDVSISNVQPK.T	4	PRR8_HUMAN
Q9NSV0	2	4.2082	K.VLPIKPADVEEPAVPQTPR.V	2	PRR8_HUMAN
Q9NSV0	2	3.9212	R.TNSGGGDGPHISSK.V	1	PRR8_HUMAN
Proline-rich protein PRCC - Homo sapiens (Human)					
Q92733	1	2.5582	K.SAALQVTK.Q	2	PRCC_HUMAN
Q92733	2	2.8579	K.TILQGSSEGTGLSALLPQPK.N	2	PRCC_HUMAN
Q92733	2	4.7075	K.TSSLAPVVGTTTTTPSPSAIK.A	3	PRCC_HUMAN
Proline-serine-threonine phosphatase-interacting protein 1 - Homo sapiens (Human)					
O43586	2	3.3717	R.GFVPGSYLEKL.-	3	PPIP1_HUMAN
Prolyl 4-hydroxylase subunit alpha-1 precursor - Homo sapiens (Human)					
P13674	2	2.9239	K.SASDDQSDQKTTPK.K	2	P4HA1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P13674	2	4.0563	R.LQDTYNLDTDTISK.G	1	P4HA1_HUMAN
ProSAAS precursor - Homo sapiens (Human)					
Q9UHG2	2	3.6095	R.NSDPALGLDDDPDAPAAQLAR.A	1	PCSK1_HUMAN
Prostaglandin E synthase 3 - Homo sapiens (Human)					
Q15185	2	3.5907	K.LTFSCLGGSDNFK.H	2	TEBP_HUMAN
Q15185	2	3.9966	R.KGESGQSWPR.L	4	TEBP_HUMAN
Q15185	3	4.2652	K.HLNEIDLFHCIDPNDSK.H	3	TEBP_HUMAN
Proteasome - Homo sapiens (Human)					
Q5VWC4	3	5.6868	R.AAAASAAEAGIATTGTEGERDSDDALLK.M	3	Q5VWC4_HUMA
Proteasome activator complex subunit 1 - Homo sapiens (Human)					
Q06323	4	6.9024	K.KISELDAFLKEPALNEANLSNLK.A	3	PSME1_HUMAN
Q06323	2	4.3494	K.TENLLGSYFPK.K	4	PSME1_HUMAN
Proteasome subunit alpha type 1 - Homo sapiens (Human)					
P25786	2	3.0704	K.AQPAQPADEPAEKADEPM#EH.-	1	PSA1_HUMAN
P25786	2	2.8462	K.AQPAQPADEPAEKADEPMEH.-	1	PSA1_HUMAN
P25786	3	5.1459	R.KAQPAQPADEPAEKADEPMEH.-	2	PSA1_HUMAN
Proteasome subunit alpha type 5 - Homo sapiens (Human)					
P28066	2	3.7823	R.LFQVEYAIEAIK.L	1	PSA5_HUMAN
Proteasome subunit beta type 4 precursor - Homo sapiens (Human)					
P28070	2	4.0043	R.TQNPM#VTGTSVLGVK.F	2	PSB4_HUMAN
P28070	2	4.0605	R.TQNPMVTGTSVLGVK.F	1	PSB4_HUMAN
Proteasome subunit beta type 6 precursor - Homo sapiens (Human)					
P28072	2	2.8957	R.DGSSGGVIR.L	2	PSB6_HUMAN
P28072	2	3.8654	R.LAAIAESGVER.Q	1	PSB6_HUMAN
P28072	2	2.7965	R.TTTGSYIANR.V	1	PSB6_HUMAN
Proteasome subunit beta type 9 precursor - Homo sapiens (Human)					
P28065	2	3.5296	R.VILGNELPKFYDE.-	2	PSB9_HUMAN
Protein 4.1 - Homo sapiens (Human)					
P11171	3	4.5413	K.KEDEPPEQAEPEPTEAWKVEK.T	1	41_HUMAN
P11171	2	3.0441	R.ERLDGENIYIR.H	2	41_HUMAN
P11171	3	4.8014	R.HSNLMLEDLDKSQEEIKK.H	3	41_HUMAN
Protein ADRM1 - Homo sapiens (Human)					
Q16186	2	3.1261	K.AMQNNAKPEQK.E	4	ADRM1_HUMAN
Q16186	2	3.6672	K.DKKDEEEDMSLD.-	1	ADRM1_HUMAN
Protein AF-9 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P42568	2	3.5401	K.RPPISDSEELSAK.K	1	AF9_HUMAN
Protein ALO17 - Homo sapiens (Human)					
Q9HCF4	2	3.0433	K.TKDEMAAAEEK.V	1	ALO17_HUMAN
Q9HCF4	2	4.623	R.GLSQEGTGPPTSAGEGHSR.T	3	ALO17_HUMAN
Q9HCF4	2	2.9632	K.ASWTVQESK.K	3	ALO17_HUMAN
Q9HCF4	2	3.9843	K.TKDEM#AAAEEK.V	3	ALO17_HUMAN
Protein ARMET precursor - Homo sapiens (Human)					
P55145	2	4.6232	K.IINEVSKPLAHHIPVEK.I	2	ARMET_HUMAN
P55145	3	6.1812	R.FYQDLKDRDVTFSPATIENELIK.F	2	ARMET_HUMAN
P55145	2	2.7269	R.KINELMPK.Y	1	ARMET_HUMAN
P55145	3	4.3356	K.DRDVTFSPATIENELIK.F	1	ARMET_HUMAN
PRotein Associated with Tlr4 - Homo sapiens (Human)					
Q8N129	2	3.1508	R.ILDYSVHAER.K	1	Q8N129_HUMAN
Q8N129	3	5.7632	R.SREVLELGQVLDTGKR.K	3	Q8N129_HUMAN
Q8N129	2	3.3718	R.EVLELGQVLDTGKR.K	2	Q8N129_HUMAN
Q8N129	2	3.7234	R.EVLELGQVLDTGK.R	1	Q8N129_HUMAN
Q8N129	2	4.9535	K.LLSTELQAELSR.T	7	Q8N129_HUMAN
Protein bicaudal C homolog 1 - Homo sapiens (Human)					
Q9H694	2	3.628	R.TPTNTWSGLGFSK.S	2	BICC1_HUMAN
Q9H694	3	4.8216	K.VLSANHGDPSIQTSGSEQTSPK.S	2	BICC1_HUMAN
Protein bicaudal D homolog 2 - Homo sapiens (Human)					
Q8TD16	2	4.296	R.LVMEAQPEWLR.A	1	BICD2_HUMAN
Q8TD16	2	3.6357	R.NVLTNTQSENER.L	3	BICD2_HUMAN
Protein BUD31 homolog - Homo sapiens (Human)					
P41223	3	3.7428	R.KAPPDGWELIEPTLDELDQK.M	1	BUD31_HUMAN
Protein C10 - Homo sapiens (Human)					
Q99622	3	5.7773	K.M#LQFVLPVATQIQQEVIK.A	1	C10_HUMAN
Q99622	3	6.5946	K.MLQFVLPVATQIQQEVIK.A	6	C10_HUMAN
Q99622	2	4.4381	K.SYEAQDPEIASLSGK.L	3	C10_HUMAN
Protein C20orf11 - Homo sapiens (Human)					
Q9NWU2	2	3.8443	K.LLLWAQNELDQK.K	2	CT011_HUMAN
Q9NWU2	3	3.8833	R.TLALLAFDSPEESPFGDLLHTM#QR.Q	1	CT011_HUMAN
Q9NWU2	2	3.2189	R.LIMNYLVTEGFK.E	1	CT011_HUMAN
Protein capicua homolog - Homo sapiens (Human)					
Q96RK0	3	4.8347	K.AQESGQGSTAGPLRPPPPGAGGPATPSK.A	2	CIC_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Protein CASC3 - Homo sapiens (Human)					
O15234	3	5.0607	R.AGFRPVEAGGQHGGR.S	2	CASC3_HUMAN
O15234	2	3.2706	R.GQTQEEEVRPK.G	1	CASC3_HUMAN
O15234	2	4.0447	R.QSGDGQESTEPVENK.V	3	CASC3_HUMAN
O15234	3	3.7362	R.RTPQPVTIKPPPPEVVSR.G	1	CASC3_HUMAN
O15234	2	2.7301	R.TGALHLR.R	1	CASC3_HUMAN
Protein CDV3 homolog - Homo sapiens (Human)					
Q9UKY7	2	3.777	R.GRDEVSKNQALK.L	2	Q9UKY7_HUMA
Protein cordon-bleu - Homo sapiens (Human)					
O75128	2	3.5757	R.GPPSTPVPTQTQNPESR.L	2	COBL_HUMAN
Protein CutA precursor - Homo sapiens (Human)					
O60888	2	5.5726	K.GKIEEDSEVLMMIK.T	4	CUTA_HUMAN
O60888	2	3.0535	K.TQSSLVPALTDFVR.S	1	CUTA_HUMAN
Protein CWC15 homolog - Homo sapiens (Human)					
Q9P013	2	3.3598	K.GEGDLSQLSK.Q	2	CWC15_HUMAN
Q9P013	3	4.0822	R.GKGEGDLSQLSK.Q	2	CWC15_HUMAN
Q9P013	2	3.2008	R.RWDDDVVFK.N	3	CWC15_HUMAN
Protein DEK - Homo sapiens (Human)					
P35659	1	2.116	K.SLIVEGK.R	1	DEK_HUMAN
P35659	2	2.983	R.LTMQVSSLQREPFTIAQGK.G	1	DEK_HUMAN
P35659	2	3.9111	R.LTMQVSSLQR.E	3	DEK_HUMAN
P35659	2	3.4026	R.LTM#QVSSLQR.E	4	DEK_HUMAN
P35659	2	4.357	K.NVGQFSGFPFEK.G	4	DEK_HUMAN
P35659	2	3.4839	K.LLYNRPGTVSSLK.K	1	DEK_HUMAN
P35659	2	4.3044	K.LLASANLEEVTM#K.Q	2	DEK_HUMAN
P35659	2	4.7454	K.KNVGQFSGFPFEK.G	3	DEK_HUMAN
P35659	2	5.2156	K.KLLASANLEEVTMK.Q	1	DEK_HUMAN
P35659	1	2.5155	R.EPFTIAQGK.G	2	DEK_HUMAN
Protein DGCR14 - Homo sapiens (Human)					
Q96DF8	2	3.0666	R.EAGEAGAATSK.Q	2	DGC14_HUMAN
Q96DF8	3	5.1437	R.HAWLYQAEEEFEKR.Q	1	DGC14_HUMAN
Q96DF8	2	3.0782	K.REAGEAGAATSK.Q	1	DGC14_HUMAN
Q96DF8	2	3.1748	K.GLSPAMSPALQR.L	1	DGC14_HUMAN
Q96DF8	3	3.821	R.VLDEEEYIEGLQTVIQR.D	1	DGC14_HUMAN
Protein disulfide-isomerase A3 precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P30101	2	4.1279	R.YLKSEPIPESNDGPVK.V	3	PDIA3_HUMAN
P30101	2	2.8197	R.FLQDYFDGNLK.R	1	PDIA3_HUMAN
P30101	2	4.1158	R.DGEEAGAYDGPR.T	12	PDIA3_HUMAN
P30101	2	3.6687	R.EATNPPVIQEEKPK.K	4	PDIA3_HUMAN
P30101	2	2.8583	R.EATNPPVIQEEKPKK.K	2	PDIA3_HUMAN
P30101	2	3.8483	R.ELSDFISYLQR.E	3	PDIA3_HUMAN
P30101	2	3.8321	R.FLQDYFDGNLKR.Y	4	PDIA3_HUMAN
P30101	2	3.305	R.GFPTIYFSPANKK.L	3	PDIA3_HUMAN
P30101	3	4.1765	R.KTFSHELSDFGLESTAGEIPVVAIR.T	2	PDIA3_HUMAN
P30101	2	3.6416	K.YGVSGYPTLK.I	6	PDIA3_HUMAN
P30101	2	3.949	R.TADGIVSHLKK.Q	2	PDIA3_HUMAN
P30101	3	4.5321	K.RLAPEYEAAATRLK.G	2	PDIA3_HUMAN
P30101	2	3.2085	R.LAPEYEAAATR.L	7	PDIA3_HUMAN
P30101	2	3.1874	K.LNFAVASR.K	3	PDIA3_HUMAN
P30101	3	5.9548	K.FISDKDASIVGFFDDSFSEAHSEFLK.A	2	PDIA3_HUMAN
P30101	2	5.6333	K.TFSHELSDFGLESTAGEIPVVAIR.T	1	PDIA3_HUMAN
P30101	2	5.3003	K.IFRDGEEAGAYDGPR.T	9	PDIA3_HUMAN
P30101	2	3.4329	K.VDCTANTNTCNK.Y	2	PDIA3_HUMAN
P30101	2	4.469	K.LSKDPNIVIAK.M	4	PDIA3_HUMAN
P30101	2	4.2748	K.M#DATANDVPSPYEVR.G	2	PDIA3_HUMAN
P30101	3	4.3602	K.M#DATANDVPSPYEVRGFPTIYFSPANK.K	1	PDIA3_HUMAN
P30101	2	4.3975	K.MDATANDVPSPYEVR.G	5	PDIA3_HUMAN
P30101	2	4.6644	K.RLAPEYEAAATR.L	4	PDIA3_HUMAN
P30101	2	3.2064	K.SEPIPESNDGPVK.V	2	PDIA3_HUMAN
P30101	2	3.2204	K.FVMQEEFSR.D	4	PDIA3_HUMAN
Protein disulfide-isomerase A4 precursor - Homo sapiens (Human)					
P13667	2	3.6493	K.M#DATANDVPSDR.Y	3	PDIA4_HUMAN
P13667	3	4.8459	R.YKVEGFPTIYFAPSGDK.K	1	PDIA4_HUMAN
P13667	2	3.7069	K.YGIVDYMIEQSGPPSK.E	2	PDIA4_HUMAN
P13667	2	3.1608	K.VDATAETDLAKR.F	2	PDIA4_HUMAN
P13667	3	5.0038	R.YKVEGFPTIYFAPSGDKK.N	2	PDIA4_HUMAN
P13667	2	4.0203	K.VDATAETDLAK.R	4	PDIA4_HUMAN
P13667	3	4.2569	K.KGQAVDYEGSRTQEEIVAK.V	1	PDIA4_HUMAN
P13667	2	3.7675	K.KGQAVDYEGSR.T	2	PDIA4_HUMAN
P13667	2	4.8611	K.IDATSASVLASR.F	4	PDIA4_HUMAN
P13667	2	3.3023	K.RFDVSGYPTLK.I	2	PDIA4_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P13667	2	3.5448	K.MDATANDVPSDRYK.V	1	PDIA4_HUMAN
Protein disulfide-isomerase A6 precursor - Homo sapiens (Human)					
Q15084	2	3.6757	K.NRPEDYQGGR.T	2	PDIA6_HUMAN
Q15084	2	2.7821	R.TRSDIVSR.A	1	PDIA6_HUMAN
Q15084	3	6.1545	K.VGAVDADKHHSLGGQYGVQGFPTIK.I	2	PDIA6_HUMAN
Q15084	2	4.1305	K.NLEPEWAAAASEVKEQTK.G	1	PDIA6_HUMAN
Q15084	2	3.2492	K.NLEPEWAAAASEVK.E	2	PDIA6_HUMAN
Q15084	2	5.366	K.LAAVDATVNQVLASR.Y	3	PDIA6_HUMAN
Q15084	2	3.9857	K.GSFSEQGINEFLR.E	2	PDIA6_HUMAN
Q15084	2	3.33	K.AATALKDVVK.V	2	PDIA6_HUMAN
Q15084	2	4.6644	R.TGEAIVDAALSALR.Q	6	PDIA6_HUMAN
Protein disulfide-isomerase precursor - Homo sapiens (Human)					
P07237	2	3.9599	R.NNFEGEVTKENLLDFIK.H	1	PDIA1_HUMAN
P07237	2	3.1338	R.ILEFFGLK.K	2	PDIA1_HUMAN
P07237	2	5.2602	K.VDATEESDLAQQYGVR.G	24	PDIA1_HUMAN
P07237	3	4.0537	K.VLVGKNFEDVAFDEKK.N	1	PDIA1_HUMAN
P07237	2	3.1516	K.THILLFLPK.S	5	PDIA1_HUMAN
P07237	3	4.5617	K.YQLDKDGVVLFK.K	9	PDIA1_HUMAN
P07237	3	4.9157	K.YQLDKDGVVLFKK.F	6	PDIA1_HUMAN
P07237	2	2.8274	R.EADDIVNWLK.K	4	PDIA1_HUMAN
P07237	2	3.2074	R.EADDIVNWLKK.R	2	PDIA1_HUMAN
P07237	2	2.9503	R.ILEFFGLKK.E	1	PDIA1_HUMAN
P07237	2	5.29	R.KSNFAEALAAHK.Y	7	PDIA1_HUMAN
P07237	2	5.8809	R.LAKVDATEESDLAQQYGVR.G	5	PDIA1_HUMAN
P07237	2	3.6736	R.LITLEEEM#TK.Y	4	PDIA1_HUMAN
P07237	2	2.9211	K.TAAESFKGK.I	2	PDIA1_HUMAN
P07237	2	4.3143	R.NGDTASPKEYTAGR.E	1	PDIA1_HUMAN
P07237	3	3.8313	K.VDATEESDLAQQYGVRGYPTIK.F	1	PDIA1_HUMAN
P07237	3	3.9492	R.TGPAATTLPDGAAAESLVESSEVAVIGFFKDV	1	PDIA1_HUMAN
P07237	1	2.4718	R.TLDGFKK.F	3	PDIA1_HUMAN
P07237	2	2.7984	R.TVIDYNGER.T	4	PDIA1_HUMAN
P07237	1	3.32	R.LITLEEEMTK.Y	6	PDIA1_HUMAN
P07237	3	4.0476	K.IKPHLMSQELPEDWDKQPVK.V	1	PDIA1_HUMAN
P07237	3	4.128	K.YKPESEELTAER.I	3	PDIA1_HUMAN
P07237	1	2.2461	K.TAAESFK.G	1	PDIA1_HUMAN
P07237	2	3.5457	K.AAGKLKAEGSEIR.L	2	PDIA1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07237	2	2.7319	K.EYTAGREADDIVNWLK.K	1	PDIA1_HUMAN
P07237	3	5.4606	K.FDEGRNNFEGEVTKENLLDFIK.H	4	PDIA1_HUMAN
P07237	2	3.255	K.FFPASADRTVIDYNGER.T	1	PDIA1_HUMAN
P07237	3	4.7608	K.FLESGGQDGAGDDDDLEDLEEAEEPDMEED	1	PDIA1_HUMAN
P07237	3	3.9634	K.IKPHLM#SQELPEDWDKQPVK.V	1	PDIA1_HUMAN
P07237	2	3.3228	K.ILFIFIDSDHTDNQR.I	2	PDIA1_HUMAN
P07237	2	3.8417	K.NFEDVAFDEKK.N	4	PDIA1_HUMAN
P07237	2	4.2668	K.SVSDYDGKLSNFK.T	7	PDIA1_HUMAN
P07237	1	2.6015	K.SVSDYDGK.L	6	PDIA1_HUMAN
P07237	3	4.8099	K.HNQLPLVIEFTEQTAPK.I	1	PDIA1_HUMAN
P07237	3	4.1383	K.QLAPIWDKLGETYKDHENIVIAK.M	1	PDIA1_HUMAN
P07237	2	4.3787	K.LGETYKDHENIVIAK.M	3	PDIA1_HUMAN
P07237	2	2.7023	K.NFEDVAFDEK.K	1	PDIA1_HUMAN
P07237	2	4.0777	K.MDSTANEVEAVK.V	8	PDIA1_HUMAN
P07237	2	3.9896	K.M#DSTANEVEAVK.V	6	PDIA1_HUMAN
P07237	2	2.7893	K.LKAEGSEIR.L	2	PDIA1_HUMAN
P07237	2	4.1626	K.SNFAEALAAHK.Y	7	PDIA1_HUMAN
Protein DJ-1 - Homo sapiens (Human)					
Q99497	2	5.2153	K.VTVAGLAGKDPVQCSR.D	9	PARK7_HUMAN
Q99497	1	2.231	K.APLVLKD.-	3	PARK7_HUMAN
Q99497	3	7.6725	R.KGLIAAICAGPTALLAHEIGFGSK.V	3	PARK7_HUMAN
Q99497	3	5.6919	R.GPGTSFEFALAIVEALNGKEVAAQVK.A	5	PARK7_HUMAN
Q99497	3	4.3587	R.GPGTSFEFALAIVEALNGK.E	3	PARK7_HUMAN
Q99497	2	3.9174	R.DVVICPDASLEDAKK.E	2	PARK7_HUMAN
Q99497	2	3.5171	K.GAEEM#ETVIPVDVM#R.R	2	PARK7_HUMAN
Q99497	1	2.1465	K.EVAAQVK.A	1	PARK7_HUMAN
Q99497	2	3.0498	K.EILKEQENRK.G	3	PARK7_HUMAN
Q99497	2	3.8921	K.EGPYDVVVLPGGNLGAQNLSESAAVK.E	3	PARK7_HUMAN
Q99497	2	3.3857	K.DGLILTSR.G	4	PARK7_HUMAN
Q99497	2	3.4824	K.GAEEMETVIPVDVMRR.A	2	PARK7_HUMAN
Protein ENL - Homo sapiens (Human)					
Q03111	3	3.7176	K.RPATADSPKPSAK.K	1	ENL_HUMAN
Protein FAM101B - Homo sapiens (Human)					
Q8N5W9	2	3.3793	R.LSLQDVPELVDAK.K	2	F101B_HUMAN
Protein FAM103A1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BTL3	3	5.1193	K.RPPESPPIVEEWNSR.A	6	F103A_HUMAN
Q9BTL3	2	4.7287	R.FTENDKEYQEYLK.R	4	F103A_HUMAN
Protein FAM110C - Homo sapiens (Human)					
Q1W6H9	2	3.4669	R.SAAPSSVPAAPPGPEPR.V	3	F110C_HUMAN
Q1W6H9	3	4.8464	R.TGDEGKAGNPETVPTTPGPAADPATPETPAP	2	F110C_HUMAN
Protein FAM114A2 - Homo sapiens (Human)					
Q9NRY5	2	4.3614	K.NSESVDQGAKPESK.S	3	F1142_HUMAN
Protein FAM122A - Homo sapiens (Human)					
Q96E09	1	2.1738	R.NSTTFPSR.H	1	F122A_HUMAN
Q96E09	2	3.3927	R.HGLLLPASPVR.M	5	F122A_HUMAN
Q96E09	3	3.9698	R.LHQIKQEEGMDLINR.E	1	F122A_HUMAN
Protein FAM122B - Homo sapiens (Human)					
Q7Z309	2	3.1164	R.KGEMETESQPK.R	1	F122B_HUMAN
Protein FAM125A - Homo sapiens (Human)					
Q96EY5	2	5.2248	R.DM#QGLSLDAASQPSK.G	6	F125A_HUMAN
Protein FAM126B - Homo sapiens (Human)					
Q8IXS8	3	4.5911	K.QYVQQPTDLSVDSVELTPMKK.H	2	F126B_HUMAN
Q8IXS8	3	3.9504	R.KQYVQQPTDLSVDSVELTPMKK.H	1	F126B_HUMAN
Protein FAM136A - Homo sapiens (Human)					
Q96C01	2	2.9809	R.VQEAVESM#VK.S	2	F136A_HUMAN
Protein FAM21C - Homo sapiens (Human)					
Q9Y4E1	2	4.3335	K.TSLFEEDKEDDLFAIAK.D	5	FA21C_HUMAN
Q9Y4E1	3	6.7097	R.KVQSTADIFGDEEGDLFKEK.A	2	FA21C_HUMAN
Q9Y4E1	3	4.275	K.KTSLFEEDKEDDLFAIAK.D	1	FA21C_HUMAN
Q9Y4E1	2	4.5846	K.KQTLSLQAQREEK.A	1	FA21C_HUMAN
Q9Y4E1	4	4.8403	K.IPAGAVSVFLGDTDVFGAASVPSLKEPQKPE	1	FA21C_HUMAN
Q9Y4E1	2	3.3206	K.GLFSDEEDSEDLFSSQSASNLK.G	3	FA21C_HUMAN
Protein FAM32A - Homo sapiens (Human)					
Q9Y421	2	2.7728	K.LLEAM#GTSK.K	1	FA32A_HUMAN
Q9Y421	2	2.7077	R.TPAQAAFEK.M	1	FA32A_HUMAN
Q9Y421	2	3.09	K.GVAELGVTKR.K	3	FA32A_HUMAN
Protein FAM43A - Homo sapiens (Human)					
Q8N2R8	2	3.6527	R.QADGASADEPHSG.-	3	FA43A_HUMAN
Protein FAM44A - Homo sapiens (Human)					
Q8NFC6	2	2.8764	K.DVQQESSEQKNK.S	1	FA44A_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q8NFC6	3	3.7186	K.KLPSQPTTDTSTDKER.T	1	FA44A_HUMAN
Q8NFC6	2	3.5422	R.TSEDMADKEK.S	3	FA44A_HUMAN
Protein FAM50A - Homo sapiens (Human)					
Q14320	2	2.7846	R.IAEENIM#K.S	1	FA50A_HUMAN
Q14320	2	2.9945	R.IAEENIMK.S	3	FA50A_HUMAN
Q14320	3	4.4877	K.KFSAHYDAVEAELK.S	2	FA50A_HUMAN
Protein FAM65B - Homo sapiens (Human)					
Q9Y4F9	2	3.5853	R.SQSFAGFSGLQER.R	2	FA65B_HUMAN
Protein FAM76B - Homo sapiens (Human)					
Q5HYJ3	2	3.3555	K.ADFQYQESNLR.T	1	FA76B_HUMAN
Q5HYJ3	2	2.9679	K.ISNLSPEEEQGLWK.Q	1	FA76B_HUMAN
Q5HYJ3	2	2.804	K.SSATIQNETPK.K	1	FA76B_HUMAN
Protein FAM83H - Homo sapiens (Human)					
Q6ZRV2	2	3.2183	K.EEASSPGAGEGPAEEGTR.D	2	Q6ZRV2_HUMA
Q6ZRV2	2	5.4256	R.DPGGGAGAITVASHSK.A	4	Q6ZRV2_HUMA
Protein fosB - Homo sapiens (Human)					
P53539	2	4.068	K.IPYEEGPGPGPLAEVR.D	1	FOSB_HUMAN
Protein fucU homolog - Homo sapiens (Human)					
A2VDF0	2	3.9991	R.ADGLGIPQLLEAVLK.L	1	CJ125_HUMAN
Protein HEG homolog 1 precursor - Homo sapiens (Human)					
Q9ULI3	2	3.2095	K.SGTASEMGTER.A	1	HEG1_HUMAN
Protein KIAA0284 - Homo sapiens (Human)					
Q9Y4F5	2	3.1515	R.SHTSTATQTPR.A	1	K0284_HUMAN
Q9Y4F5	2	3.6985	R.SGPGPPELDSEQPSR.L	1	K0284_HUMAN
Q9Y4F5	2	3.3008	R.RKPAAPPPSPAAR.E	1	K0284_HUMAN
Q9Y4F5	2	4.345	R.LGDASDTEAADGER.G	2	K0284_HUMAN
Q9Y4F5	3	4.3276	R.GPVLAHLPSSDVMASNHETPEATGAGR.L	1	K0284_HUMAN
Q9Y4F5	3	4.0658	R.GDRDESDDGGVAQR.M	2	K0284_HUMAN
Q9Y4F5	3	3.7418	K.QLEVINAIVDPSGSLDLLTGNR.S	1	K0284_HUMAN
Q9Y4F5	2	3.7092	R.APGEPTPASFFIGDQNGDAVLSR.K	1	K0284_HUMAN
Protein KIAA1107 - Homo sapiens (Human)					
Q9UPP5	2	5.0781	R.SEDYDAGGSQDDDGSNDR.G	1	Q9UPP5_HUMA
Q9UPP5	2	3.5942	K.TQGSQGESPNSVK.S	2	Q9UPP5_HUMA
Protein KIAA1191 - Homo sapiens (Human)					
Q6IA24	3	3.747	R.GWFTSGSSTALPGPNPSTMDSGSGDKDR.N	1	Q6IA24_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Protein KIAA1430 - Homo sapiens (Human)					
Q9P2B7	2	3.8147	K.STETQPSSTTPK.C	2	Q9P2B7_HUMA
Protein KIAA1522 - Homo sapiens (Human)					
Q9P206	3	4.5637	K.GGGPPREDVGAPLVTPSLLQM#VR.L	1	Q9P206_HUMAN
Q9P206	3	4.4375	K.GGGPPREDVGAPLVTPSLLQMVR.L	1	Q9P206_HUMAN
Q9P206	2	3.1618	R.FSSVSSPQPR.S	1	Q9P206_HUMAN
Q9P206	2	3.2017	R.SGPQILTPLGDR.F	1	Q9P206_HUMAN
Protein KIAA1967 - Homo sapiens (Human)					
Q8N163	2	3.414	K.AAYNPGQAVPWNAVK.V	2	K1967_HUMAN
Protein kinase C and casein kinase substrate in neurons protein 2 - Homo sapiens (Human)					
Q9UNF0	2	2.8764	R.AADAVEDLR.W	1	PACN2_HUMAN
Q9UNF0	2	5.2414	R.ALYDYEGQEHDELSFK.A	3	PACN2_HUMAN
Q9UNF0	1	2.1934	K.NVSSYEK.T	1	PACN2_HUMAN
Q9UNF0	2	3.6374	K.HLDLSNVAGYK.A	2	PACN2_HUMAN
Q9UNF0	3	4.7452	K.AGDELTKMEDEDEQGWCK.G	1	PACN2_HUMAN
Protein KRI1 homolog - Homo sapiens (Human)					
Q8N9T8	2	3.5321	K.AQEEADYIEWLK.G	1	KRI1_HUMAN
Protein LAP2 - Homo sapiens (Human)					
Q96RT1	2	3.3743	K.LITNDTFQPEIM#ER.S	1	LAP2_HUMAN
Q96RT1	2	2.8081	K.STEDLSPQK.S	1	LAP2_HUMAN
Q96RT1	3	3.9796	R.LKDEETNEDSGR.D	1	LAP2_HUMAN
Q96RT1	2	2.925	R.SESTENQSYAK.H	1	LAP2_HUMAN
Protein LAP4 - Homo sapiens (Human)					
Q14160	2	3.0838	R.QSPASPPPLGGGAPVR.T	1	LAP4_HUMAN
Q14160	2	2.9796	R.VSLVGADDLR.K	2	LAP4_HUMAN
Protein LTV1 homolog. - Homo sapiens (Human)					
Q96GA3	3	4.7326	K.LPSSVFASEFEEDVGLLNK.A	4	Q96GA3_HUMA
Q96GA3	3	4.6523	R.SQRDPLAADESAPQR.V	2	Q96GA3_HUMA
Q96GA3	3	4.0677	R.REEKEETLVIPSTGIK.L	2	Q96GA3_HUMA
Q96GA3	2	4.5385	R.DPLAADESAPQR.V	4	Q96GA3_HUMA
Q96GA3	2	2.7674	K.TGIPLNVLPK.K	1	Q96GA3_HUMA
Protein LYRIC - Homo sapiens (Human)					
Q86UE4	2	3.8526	K.TISTSDPAEVLVK.N	5	LYRIC_HUMAN
Q86UE4	2	3.6773	R.TVEVAEGEAVR.T	3	LYRIC_HUMAN
Q86UE4	3	4.7878	R.HDGKEVDEGAWETK.I	2	LYRIC_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q86UE4	2	3.2302	K.TLPPATSTEPSVILSK.S	5	LYRIC_HUMAN
Q86UE4	2	3.067	K.SETSWESPK.Q	6	LYRIC_HUMAN
Q86UE4	3	4.3491	K.SDSDKSSSQVPPILQETDKSK.S	1	LYRIC_HUMAN
Q86UE4	3	4.4036	K.NKGDSHLNVQVSNFK.S	2	LYRIC_HUMAN
Q86UE4	3	4.9364	K.LSSQISAGEEKWNSVSPASAGKR.K	2	LYRIC_HUMAN
Q86UE4	3	4.0113	K.LSSQISAGEEKWNSVSPASAGK.R	3	LYRIC_HUMAN
Q86UE4	2	3.6284	K.LSSQISAGEEK.W	1	LYRIC_HUMAN
Q86UE4	2	2.8847	K.WNSVSPASAGK.R	1	LYRIC_HUMAN
Protein LZIC - Homo sapiens (Human)					
Q8WZA0	2	4.1835	R.DLYTQQKVEILTALR.K	2	LZIC_HUMAN
Q8WZA0	2	2.9822	K.ILALASFEVEK.T	1	LZIC_HUMAN
Q8WZA0	3	4.5319	R.LAEMDRDLMVGKLER.D	4	LZIC_HUMAN
Q8WZA0	3	4.2924	R.LAEMDRDLM#VGKLER.D	2	LZIC_HUMAN
Q8WZA0	3	4.0552	R.DLYTQQKVEILTALRK.L	1	LZIC_HUMAN
Q8WZA0	3	4.407	K.VSTDLGSGDKILALASFEVEK.T	1	LZIC_HUMAN
Q8WZA0	1	2.8947	K.VSTDLGSGDK.I	2	LZIC_HUMAN
Q8WZA0	3	5.355	K.LTADDEAFLSANAGAILSQFEK.V	13	LZIC_HUMAN
Q8WZA0	2	4.4213	K.LKQNLEEQLDR.L	6	LZIC_HUMAN
Q8WZA0	2	2.8405	R.DLYTQQK.V	3	LZIC_HUMAN
Protein max - Homo sapiens (Human)					
P61244	2	3.6191	R.QNALLEQQVR.A	5	MAX_HUMAN
P61244	2	4.7194	R.SSAQLQTNYPSSDNSLYTNAK.G	2	MAX_HUMAN
Protein MTO1 homolog, mitochondrial precursor - Homo sapiens (Human)					
Q9Y2Z2	2	3.017	R.IC*DQSGVHYKVLNRR.K	1	MTO1_HUMAN
Protein NDRG1 - Homo sapiens (Human)					
Q92597	2	3.4498	K.SIIGMGTGAGAYILTR.F	1	NDRG1_HUMAN
Protein NEDD1 - Homo sapiens (Human)					
Q8NHV4	2	4.3369	K.LVTSGAESGNLNTSPSSNQTR.N	2	NEDD1_HUMAN
Q8NHV4	2	5.0026	R.SVNVNAASGGVQNSGIVR.E	2	NEDD1_HUMAN
Protein NOXP20 - Homo sapiens (Human)					
Q8IWE2	2	5.0333	K.SLLSSASATVGHGLTAVK.E	2	NXP20_HUMAN
Protein numb homolog - Homo sapiens (Human)					
P49757	2	2.8416	R.RHAPIEQLAR.Q	1	NUMB_HUMAN
P49757	2	3.5192	R.TNPSPTNPFSSDLQK.T	5	NUMB_HUMAN
P49757	3	3.8221	R.TPSEADRWLEEVSK.S	1	NUMB_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Protein OSCP1 - Homo sapiens (Human)					
Q8WVF1	2	4.1639	R.LSTSKGDDLLAMMDEL.-	1	OSCP1_HUMAN
Protein phosphatase 1 regulatory subunit 11 - Homo sapiens (Human)					
O60927	3	5.3918	K.KVEWTSDTVDNEHM#GR.R	1	PP1RB_HUMAN
O60927	3	5.808	K.KVEWTSDTVDNEHMGR.R	3	PP1RB_HUMAN
Protein phosphatase 1 regulatory subunit 12A - Homo sapiens (Human)					
O14974	3	4.0924	R.RISEMEEELKMLPDLK.A	1	MYPT1_HUMAN
O14974	2	2.9669	R.KDESPATWR.L	1	MYPT1_HUMAN
O14974	2	3.1584	R.YETSSTSAGDR.Y	2	MYPT1_HUMAN
O14974	2	4.7038	R.KTGSYGALAEITASK.E	3	MYPT1_HUMAN
O14974	2	4.3101	R.LASTSDIEEKENR.D	4	MYPT1_HUMAN
O14974	2	3.4761	R.LAYVAPTIPR.R	1	MYPT1_HUMAN
O14974	2	3.3892	R.LEKDDSTDFKK.L	2	MYPT1_HUMAN
O14974	2	3.0158	R.LSSSLDNKEK.E	2	MYPT1_HUMAN
O14974	2	3.8889	R.LSSSLDNKEKEK.D	1	MYPT1_HUMAN
O14974	3	5.5175	R.RLASTSDIEEKENR.D	5	MYPT1_HUMAN
O14974	3	5.1588	R.RSTQGVTLTDLQEAEK.T	6	MYPT1_HUMAN
O14974	3	5.6388	R.RSTQGVTLTDLQEAEKTIGR.S	1	MYPT1_HUMAN
O14974	2	3.7726	R.SGSYSYLEER.K	4	MYPT1_HUMAN
O14974	2	4.6068	R.STQGVTLTDLQEAEK.T	4	MYPT1_HUMAN
O14974	2	3.9448	R.STQGVTLTDLQEAEKTIGR.S	1	MYPT1_HUMAN
O14974	2	2.831	R.SYLTPVRDEESESQRK.A	1	MYPT1_HUMAN
O14974	3	4.7148	R.ISEMEEELKMLPDLK.A	3	MYPT1_HUMAN
O14974	2	3.875	R.WIGSETDLEPPVVKR.Q	2	MYPT1_HUMAN
O14974	2	5.1569	K.TGSYGALAEITASK.E	3	MYPT1_HUMAN
O14974	2	3.1817	R.YETSSTSAGDRYDSLLGR.S	1	MYPT1_HUMAN
O14974	2	3.3468	R.WIGSETDLEPPVVK.R	1	MYPT1_HUMAN
O14974	2	2.755	K.DTAGVTR.S	1	MYPT1_HUMAN
O14974	3	4.3634	K.AQLHDTNM#ELTDLKLQLEK.A	1	MYPT1_HUMAN
O14974	2	4.2521	K.AQLHDTNMELTDLK.L	4	MYPT1_HUMAN
O14974	3	4.9831	K.VKFDDGAVFLAACSSGDTDEVLK.L	2	MYPT1_HUMAN
O14974	2	2.9448	K.DDSTDFKK.L	2	MYPT1_HUMAN
O14974	2	3.1017	R.ISEMEEELK.M	1	MYPT1_HUMAN
O14974	2	2.8286	K.ESETSREDEYKQK.Y	1	MYPT1_HUMAN
O14974	2	2.908	K.FDDGAVFLAACSSGDTDEVLK.L	1	MYPT1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O14974	2	5.9184	K.ITTGSSSAGTQSSTSNR.L	4	MYPT1_HUMAN
O14974	2	3.0154	K.KQNLLHSEKR.D	2	MYPT1_HUMAN
O14974	2	3.5917	K.LYEQILAENEK.L	2	MYPT1_HUMAN
O14974	2	4.6588	K.LYEQILAENEKLK.A	3	MYPT1_HUMAN
O14974	2	2.7351	K.MLPDLKADNQR.L	2	MYPT1_HUMAN
O14974	2	4.0002	K.NKETLIIEPEK.N	3	MYPT1_HUMAN
O14974	2	4.1005	K.SGGTALHVAAAK.G	5	MYPT1_HUMAN
O14974	2	3.3778	K.SLLSSTSTTTK.I	2	MYPT1_HUMAN
O14974	2	5.206	K.SPLIESTANMDNNQSQK.T	4	MYPT1_HUMAN
O14974	3	4.2785	K.VGQTAFDVADEDILGYLEELQKK.Q	1	MYPT1_HUMAN
O14974	2	4.2152	R.ILVDNLCDMEMVNK.V	10	MYPT1_HUMAN
O14974	3	4.8862	K.KSPLIESTANMDNNQSQK.T	1	MYPT1_HUMAN
O14974	3	5.1975	K.AQLHDTNMELTDLKLQLEK.A	2	MYPT1_HUMAN
Protein phosphatase 1 regulatory subunit 12B - Homo sapiens (Human)					
O60237	2	3.6542	K.LYESALTENQK.L	2	MYPT2_HUMAN
O60237	2	3.0845	R.KMSEMEEEMK.V	1	MYPT2_HUMAN
O60237	2	3.5024	K.VLTELKSDNQR.L	2	MYPT2_HUMAN
O60237	2	5.1735	K.TKLQEAQLELADIK.S	2	MYPT2_HUMAN
Protein phosphatase 1 regulatory subunit 12C - Homo sapiens (Human)					
Q9BZL4	2	4.0266	R.VPGVENSDSPAQR.A	3	Q9BZL4_HUMAN
Q9BZL4	3	5.6057	R.VPGVENSDSPAQRAEAPDGQGPGPQAAR.E	2	Q9BZL4_HUMAN
Q9BZL4	2	3.8159	R.TGASALHVAAAK.G	2	Q9BZL4_HUMAN
Q9BZL4	2	3.6692	R.TAEGAPGAGLQR.S	2	Q9BZL4_HUMAN
Q9BZL4	3	3.8562	R.SYQMPVRDEESESQRK.A	1	Q9BZL4_HUMAN
Q9BZL4	2	3.0091	R.SYQMPVRDEESESQR.K	1	Q9BZL4_HUMAN
Q9BZL4	2	4.7477	R.SASSSWLEGTSTQAK.E	1	Q9BZL4_HUMAN
Q9BZL4	3	5.0466	R.RPGGAGGPPIQDEDEGEEGPTEPPPAEPR.T	2	Q9BZL4_HUMAN
Q9BZL4	2	3.4507	R.GQEPQAPSSSK.H	2	Q9BZL4_HUMAN
Q9BZL4	3	6.0046	R.GPAEGEEAEPADRSQESSTLEGGPSAR.R	2	Q9BZL4_HUMAN
Q9BZL4	2	3.3608	R.GPAEGEEAEPADR.S	1	Q9BZL4_HUMAN
Q9BZL4	3	4.087	R.AVLDSTNADGISALHQACIDENLEVVR.F	1	Q9BZL4_HUMAN
Q9BZL4	2	4.7473	R.AEAPDGQGPGPQAAR.E	7	Q9BZL4_HUMAN
Q9BZL4	2	2.9882	K.APESEKPAQSLDPSR.R	1	Q9BZL4_HUMAN
Q9BZL4	2	3.9536	K.SPVQLEEAPFSR.R	2	Q9BZL4_HUMAN
Q9BZL4	3	4.4725	R.IPEPESPAKPNVPTASTAPPADSR.D	2	Q9BZL4_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Protein phosphatase 1 regulatory subunit 14A - Homo sapiens (Human)					
Q96A00	2	3.0639	K.WIDGRLEELYR.G	2	PP14A_HUMAN
Q96A00	2	2.7229	R.GPGGSPGGLQK.R	1	PP14A_HUMAN
Protein phosphatase 1 regulatory subunit 14B - Homo sapiens (Human)					
Q96C90	2	3.9238	R.VYFQSPPGAAGEGPGGADDEGPVRR.Q	3	PP14B_HUMAN
Protein phosphatase 1 regulatory subunit 14C - Homo sapiens (Human)					
Q8TAE6	3	5.0058	R.EDSAPVATAAAAGQVQQQQQR.R	2	PP14C_HUMAN
Q8TAE6	3	7.012	R.GGAGGSPGSSSGSGSSREDSAPVATAAAAG	2	PP14C_HUMAN
Protein phosphatase 1 regulatory subunit 14D - Homo sapiens (Human)					
Q9NXH3	3	4.968	R.PTEAFISELLSQLKK.L	2	PP14D_HUMAN
Protein phosphatase 1 regulatory subunit 7 - Homo sapiens (Human)					
Q15435	2	4.231	R.GAGQQQSQEMMEVDR.R	2	PP1R7_HUMAN
Protein phosphatase 1A - Homo sapiens (Human)					
P35813	2	3.4127	R.NVIEAVYNR.L	2	PPM1A_HUMAN
Protein phosphatase 1G - Homo sapiens (Human)					
O15355	2	3.7977	R.LPLPYGFSAMQGWR.V	4	PPM1G_HUMAN
O15355	2	2.9009	R.ETPSQENGPTAK.A	1	PPM1G_HUMAN
O15355	2	3.4047	K.AYTGFSSNSER.G	4	PPM1G_HUMAN
O15355	2	2.9675	K.ALEDAFLAIDAK.L	2	PPM1G_HUMAN
Protein phosphatase methylesterase 1 - Homo sapiens (Human)					
Q9Y570	3	3.762	R.LPSRPPLPGSGGSQSGAK.M	1	PPME1_HUMAN
Protein phosphatase Slingshot homolog 2 - Homo sapiens (Human)					
Q76I76	3	4.0947	R.SSSLSNTPHASEESSMDEEQSK.A	1	SSH2_HUMAN
Protein phosphatase Slingshot homolog 3 - Homo sapiens (Human)					
Q8TE77	2	2.7422	R.QASVHDSGEEGEA.-	1	SSH3_HUMAN
Protein Red - Homo sapiens (Human)					
Q13123	1	3.1079	K.AAFQYGIK.M	4	RED_HUMAN
Q13123	2	2.8508	K.KMEADGVEVK.R	2	RED_HUMAN
Q13123	2	4.2053	R.AVGPTAEADKSAAEK.R	2	RED_HUMAN
Q13123	3	3.8125	R.DGVNKDYEETELISTTANYR.A	1	RED_HUMAN
Protein RMI1 homolog - Homo sapiens (Human)					
Q9H9A7	2	2.8662	K.AMVLALQDVNM#EHLENLKKR.L	1	RMI1_HUMAN
Protein S100-A10 - Homo sapiens (Human)					
P60903	2	3.0954	K.FAGDKGYLTK.E	1	S10AA_HUMAN
P60903	3	4.876	R.VLMEKEFPGFLENQKDPLAVDK.I	1	S10AA_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Protein S100-A11 - Homo sapiens (Human)					
P31949	1	3.5561	K.DGYNYTLSK.T	7	S10AB_HUMAN
P31949	3	3.9372	K.ISSPTETERCIESLIAVFQK.Y	1	S10AB_HUMAN
P31949	2	3.866	K.YAGKDGYNYTLSK.T	2	S10AB_HUMAN
Protein S100-A12 - Homo sapiens (Human)					
P80511	2	3.6935	K.GHFDTLSKGELK.Q	1	S10AC_HUMAN
Protein S100-A13 - Homo sapiens (Human)					
Q99584	2	3.3281	K.DVGSLDEKMK.S	3	S10AD_HUMAN
Q99584	3	4.7003	K.SLDVNQDSELKFNEYWR.L	2	S10AD_HUMAN
Q99584	1	2.2945	R.LIGELAK.E	2	S10AD_HUMAN
Q99584	3	4.4271	R.KDSLSVNEFKELVTQQLPHLLK.D	2	S10AD_HUMAN
Q99584	2	3.1232	R.KDSLSVNEFK.E	2	S10AD_HUMAN
Q99584	1	3.0688	K.DSLSVNEFK.E	2	S10AD_HUMAN
Q99584	2	3.0479	K.SLDVNQDSELK.F	2	S10AD_HUMAN
Q99584	2	3.5111	K.DVGSLDEKM#K.S	6	S10AD_HUMAN
Q99584	1	2.9049	K.DVGSLDEK.M	4	S10AD_HUMAN
Q99584	4	5.446	K.DSLSVNEFKELVTQQLPHLLKDVGSLDEK.M	3	S10AD_HUMAN
Q99584	2	4.2359	K.ELVTQQLPHLLK.D	1	S10AD_HUMAN
Protein S100-A4 - Homo sapiens (Human)					
P26447	2	3.3078	K.RTDEAAFQK.L	2	S10A4_HUMAN
Protein S100-A6 - Homo sapiens (Human)					
P06703	2	3.0948	K.LQDAEIAR.L	4	S10A6_HUMAN
Protein S100-A8 - Homo sapiens (Human)					
P05109	2	3.492	K.LLETECPQYIR.K	2	S10A8_HUMAN
Protein S100-A9 - Homo sapiens (Human)					
P06702	2	4.779	R.NIETIINTFHQYSVK.L	6	S10A9_HUMAN
P06702	1	2.2378	R.LTWASHEK.M	1	S10A9_HUMAN
P06702	2	3.0675	R.KDLQNFLK.K	3	S10A9_HUMAN
P06702	2	5.1988	K.VIEHIMEDLDTNADK.Q	3	S10A9_HUMAN
P06702	2	3.6178	K.MHEGDEGPGHHHKPGLGEGTP.-	1	S10A9_HUMAN
P06702	2	4.5312	K.LGHPDTLNQGEFKELVRK.D	2	S10A9_HUMAN
P06702	2	4.5322	K.LGHPDTLNQGEFKELVR.K	1	S10A9_HUMAN
P06702	2	2.8088	K.DLQNFLK.K	3	S10A9_HUMAN
P06702	2	4.2145	K.LGHPDTLNQGEFK.E	13	S10A9_HUMAN
Protein SET - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q01105	3	5.7374	K.KPRPPPALGPEETSASAGLPK.K	5	SET_HUMAN
Q01105	2	2.8056	K.RQSPLPPQK.K	3	SET_HUMAN
Protein Shroom4 - Homo sapiens (Human)					
Q9ULL8	2	4.994	R.ASSVDSNPLNEASAELAK.A	2	SHRM4_HUMAN
Q9ULL8	3	3.9637	K.APSGRPNVAETSGGSR.R	1	SHRM4_HUMAN
Protein SLC7A6OS - Homo sapiens (Human)					
Q96CW6	2	3.624	R.SDAVESAAQK.T	3	S7A6O_HUMAN
Protein TMED8 - Homo sapiens (Human)					
Q6PL24	3	4.3218	K.QDLLPADQAQVLNEM#AK.Y	1	TMED8_HUMAN
Q6PL24	2	4.0834	K.QDLLPADQAQVLNEMAK.Y	3	TMED8_HUMAN
Protein transport protein Sec31A - Homo sapiens (Human)					
O94979	3	3.7257	K.KPIPDEHLILK.T	1	SC31A_HUMAN
O94979	3	3.8192	K.RLEFLYDKLR.E	2	SC31A_HUMAN
Protein TSSC4 - Homo sapiens (Human)					
Q9Y5U2	2	2.9149	R.GM#SSTFSQR.S	1	TSSC4_HUMAN
Q9Y5U2	2	3.3072	R.SRDIFDCLEGAAR.R	1	TSSC4_HUMAN
Q9Y5U2	2	3.1516	R.GMSSTFSQR.S	3	TSSC4_HUMAN
Q9Y5U2	2	2.9378	K.RPLAPSGR.S	1	TSSC4_HUMAN
Q9Y5U2	2	3.4185	R.NKSSSPEDPGAEV.-	2	TSSC4_HUMAN
Protein tyrosine phosphatase receptor type C-associated protein - Homo sapiens (Human)					
Q14761	2	3.4845	R.AAGGQGLHVTAL.-	2	PTCA_HUMAN
Protein UNQ6126/PRO20091 precursor - Homo sapiens (Human)					
Q6UXV3	3	3.754	K.INC*FFYLEKQLC*QLPSPLCLSSLLTLK.L	1	U6126_HUMAN
Protein wibg homolog - Homo sapiens (Human)					
Q9BRP8	3	4.1151	R.RQQQEKGEAEALSR.T	2	WIBG_HUMAN
Q9BRP8	3	4.9031	R.TLDKVSLEETAQLPSAPQGSR.A	4	WIBG_HUMAN
Q9BRP8	2	3.9753	R.IQAGEVSQPSK.E	5	WIBG_HUMAN
Q9BRP8	2	3.9188	R.QQQEKGEAEALSR.T	2	WIBG_HUMAN
Q9BRP8	2	5.9795	R.IQAGEVSQPSKEQLEK.L	4	WIBG_HUMAN
Q9BRP8	2	2.7168	K.YIASTQRPDGTWR.K	1	WIBG_HUMAN
Q9BRP8	2	4.4481	K.VSLEETAQLPSAPQGSR.A	2	WIBG_HUMAN
Q9BRP8	3	3.7385	K.SKPELPPGLSPEATAPVTPSRPEGGEPGLSK.	1	WIBG_HUMAN
Q9BRP8	3	4.1247	K.LRQVEELQQR.I	4	WIBG_HUMAN
Q9BRP8	2	3.76	K.EGYVPQEEVPVYENK.Y	2	WIBG_HUMAN
Q9BRP8	2	4.7976	R.VKEGYVPQEEVPVYENK.Y	3	WIBG_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BRP8	2	4.3237	R.AAPTAASDQPDSAATTEK.A	7	WIBG_HUMAN
Protein Z-dependent protease inhibitor precursor - Homo sapiens (Human)					
Q9UK55	3	3.911	R.GLHLQALKPTKPGLLPSLFK.G	2	ZPI_HUMAN
Q9UK55	2	5.3551	R.IFSPFADLSELSATGR.N	3	ZPI_HUMAN
Protein-L-isoaspartate(D-aspartate) O-methyltransferase - Homo sapiens (Human)					
P22061	4	4.7806	R.M#GYAEEAPYDAIHVGAAAPVVPQALIDQLKP	1	PIMT_HUMAN
P22061	4	4.9551	R.MGYAEEAPYDAIHVGAAAPVVPQALIDQLKP	1	PIMT_HUMAN
P22061	2	2.815	K.DDPTLLSSGR.V	1	PIMT_HUMAN
Prothrombin precursor - Homo sapiens (Human)					
P00734	3	4.909	K.SLEDKTERELLESYIDGR.I	2	THRB_HUMAN
P00734	1	2.1312	K.YTACETAR.T	1	THRB_HUMAN
P00734	2	2.8513	R.ELLESYIDGR.I	1	THRB_HUMAN
P00734	3	3.7261	R.LAVTTHGLPCLAWASAQAK.A	1	THRB_HUMAN
P00734	2	3.4814	R.TATSEYQTFFNPR.T	4	THRB_HUMAN
P00734	3	4.4589	K.HQDFNSAVQLVENFCR.N	2	THRB_HUMAN
Prothymosin a14 - Homo sapiens (Human)					
Q9UMZ1	2	3.7318	R.AAEDDEDNDVDTK.K	9	Q9UMZ1_HUMA
Q9UMZ1	2	3.9524	R.AAEDDEDNDVDTKK.Q	2	Q9UMZ1_HUMA
Protocadherin Fat 2 precursor - Homo sapiens (Human)					
Q9NYQ8	2	3.0927	K.VM#ARDGGGRVAFC*TVK.I	1	FAT2_HUMAN
Protocadherin-17 precursor - Homo sapiens (Human)					
O14917	2	3.6301	R.DPPFM#ASDQM#AR.V	2	PCD17_HUMAN
Proto-oncogene C-crk - Homo sapiens (Human)					
P46108	3	4.6803	R.VPNAYDKTALALEVGELVK.V	1	CRK_HUMAN
P46108	2	3.2842	K.TALALEVGELVK.V	2	CRK_HUMAN
P46108	3	5.687	K.RVPNAYDKTALALEVGELVK.V	1	CRK_HUMAN
Proto-oncogene protein c-fos - Homo sapiens (Human)					
P01100	2	3.0547	R.KGSSSNEPSSDSLSSPTLLAL.-	1	FOS_HUMAN
Proto-oncogene tyrosine-protein kinase ABL1 - Homo sapiens (Human)					
P00519	3	5.2789	K.AGGKPSQSPSQEAAGEAVLGAK.T	2	ABL1_HUMAN
P00519	3	5.1031	K.TKATSLVDAVNSDAAKPSQPGEGLK.K	1	ABL1_HUMAN
P00519	2	3.617	R.LATGEEEGGGSSSK.R	2	ABL1_HUMAN
Pseudouridylate synthase 7 homolog - Homo sapiens (Human)					
Q96PZ0	2	4.0111	K.HGLTEADVGITK.F	1	PUS7_HUMAN
Pterin-4-alpha-carbinolamine dehydratase - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P61457	2	4.3413	R.LSAEERDQLLPNLR.A	5	PHS_HUMAN
P61457	3	4.6366	R.VALQAEKLDHHPEWFNVYNK.V	2	PHS_HUMAN
P61457	2	3.7272	R.AVGWNELEGR.D	4	PHS_HUMAN
P61457	3	3.7412	K.LDHHPEWFNVYNK.V	1	PHS_HUMAN
PTPN18 protein - Homo sapiens (Human)					
Q7Z637	2	5.2366	R.GAPAGAGSGTQTGTGTGAR.S	5	Q7Z637_HUMAN
Pumilio homolog 2 - Homo sapiens (Human)					
Q8TB72	2	3.7748	R.DAETDGPEKGDQK.G	3	PUM2_HUMAN
Putative adenosylhomocysteinase 2 - Homo sapiens (Human)					
O43865	3	4.43	K.EIEDAEKYSFMATVTK.A	2	SAHH2_HUMAN
O43865	2	3.3543	K.KQIQFADDMQEFTK.F	1	SAHH2_HUMAN
O43865	3	5.2435	K.KQIQFADDMQEFTKFPTK.T	2	SAHH2_HUMAN
Putative adenosylhomocysteinase 3 - Homo sapiens (Human)					
Q96HN2	3	4.6001	R.HRDGGEALVSPDGTVTEAPR.T	2	SAHH3_HUMAN
Putative ATP-dependent RNA helicase DHX29 - Homo sapiens (Human)					
Q7Z478	2	3.8476	K.SAEAGIAGEAQSK.K	2	DHX29_HUMAN
Q7Z478	2	3.2597	R.PATAAAAAAGSR.E	2	DHX29_HUMAN
Putative ATP-dependent RNA helicase DHX57 - Homo sapiens (Human)					
Q6P158	2	4.6018	R.DLQEQDADAGSER.G	5	DHX57_HUMAN
Putative GTP-binding protein 9 - Homo sapiens (Human)					
Q9NTK5	2	3.1571	R.NYIVEDGDIIFFK.F	1	GTPB9_HUMAN
Putative GTP-binding protein Parf - Homo sapiens (Human)					
Q3YEC7	3	4.2268	R.TAADELEAFLGGGAPGGR.H	2	PARF_HUMAN
Putative MAPK activating protein - Homo sapiens (Human)					
Q7Z422	1	2.1806	R.EAEYAEAR.K	1	Q7Z422_HUMAN
Q7Z422	2	4.151	R.ILGSASPEEEQEKPILDRPTR.I	1	Q7Z422_HUMAN
Q7Z422	2	3.8593	R.ISQPEDSRQPNNVIR.Q	2	Q7Z422_HUMAN
Q7Z422	3	4.8705	K.SPPKVPIVIQDDSLPAGPPPQIR.I	3	Q7Z422_HUMAN
Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 - Homo sapiens (Human					
O43143	2	5.1539	R.HRLDLGEDYPSGK.K	3	DHX15_HUMAN
Putative RNA methyltransferase NOL1 - Homo sapiens (Human)					
P46087	2	4.7953	K.TQASSSFQDSSQPAGK.A	4	NOL1_HUMAN
Putative RNA-binding protein 15 - Homo sapiens (Human)					
Q96T37	2	3.2997	R.RLPEESGGR.H	2	RBM15_HUMAN
Q96T37	2	3.2969	R.SDGSAPSTSTASSK.L	2	RBM15_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96T37	2	3.5953	R.EYDTGGGSSSSR.L	2	RBM15_HUMAN
Q96T37	2	3.3084	K.ISHLSGSGSGDER.V	2	RBM15_HUMAN
Q96T37	4	5.0129	K.ISELGSQLSDEAVEDGLFHEFKR.F	1	RBM15_HUMAN
Putative RNA-binding protein 3 - Homo sapiens (Human)					
P98179	2	3.3399	R.AM#NGESLDGR.Q	4	RBM3_HUMAN
P98179	1	3.1913	R.AMNGESLDGR.Q	8	RBM3_HUMAN
P98179	2	5.6466	R.GFGFITFTNPEHASVAM#R.A	2	RBM3_HUMAN
P98179	2	5.6206	R.GFGFITFTNPEHASVAMR.A	13	RBM3_HUMAN
P98179	2	3.8314	R.PGGYGYGYGR.S	2	RBM3_HUMAN
P98179	3	5.9917	R.SRGFGFITFTNPEHASVAMR.A	4	RBM3_HUMAN
P98179	2	4.3163	R.YSGGNYRDNYDN.-	3	RBM3_HUMAN
P98179	2	3.3525	R.YYDSRPGGYGYGYGR.S	2	RBM3_HUMAN
P98179	4	4.7543	K.LFVGGLNFNTDEQALEDHFSSFGPISEVVVVK	3	RBM3_HUMAN
Putative RNA-binding protein Luc7-like 2 - Homo sapiens (Human)					
Q9Y383	2	4.0108	R.AMLDQLMGTSR.D	3	LC7L2_HUMAN
Putative rRNA methyltransferase 3 - Homo sapiens (Human)					
Q8IY81	3	4.1656	R.LTEVQDDKEEEEEENPLLVPLEEK.A	1	RRMJ3_HUMAN
Q8IY81	2	3.0158	K.SDDDGFEIVPIEDPAK.H	1	RRMJ3_HUMAN
Q8IY81	2	3.1725	R.ILDPEGLALGAVIASSK.K	1	RRMJ3_HUMAN
Putative thiosulfate sulfurtransferase KAT - Homo sapiens (Human)					
Q8NFU3	2	3.2216	K.RGLQATQLAR.S	3	KAT_HUMAN
Q8NFU3	1	3.5255	R.GLQATQLAR.S	6	KAT_HUMAN
Q8NFU3	1	2.5393	R.NYAGAYR.E	3	KAT_HUMAN
Q8NFU3	2	4.2408	R.NYAGAYREWLEK.E	2	KAT_HUMAN
Q8NFU3	2	4.1359	R.NYAGAYREWLEKES.-	4	KAT_HUMAN
Q8NFU3	1	2.4358	R.SLGYTGAR.N	2	KAT_HUMAN
Q8NFU3	1	2.5043	R.SLLASGR.A	4	KAT_HUMAN
Putative uncharacterized protein - Homo sapiens (Human)					
Q9Y520	2	4.1236	R.SVSHGSNHTQKPDEQR.S	4	Q9Y520_HUMAN
Q05CP8	3	6.0938	R.KLMQLQHEKAELEQHLEQEQEFQVNK.L	1	Q05CP8_HUMA
Q9Y520	2	3.0367	R.MLWGSDPYPHAEPQQATTPK.A	1	Q9Y520_HUMAN
A0JLQ0	2	4.6923	R.YSLTYIYTGLSK.H	7	A0JLQ0_HUMAN
Q9Y520	2	3.0234	R.PAVLSGYFK.Q	2	Q9Y520_HUMAN
Q9Y520	2	3.2207	R.DHAISLSEPR.M	2	Q9Y520_HUMAN
Q05DR4	3	4.7844	R.RSLAALSQIAYQR.N	7	Q05DR4_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96F87	2	4.7183	R.SFQPLGPGYGISQSSR.L	5	Q96F87_HUMAN
Q05DR4	2	4.1898	R.SLAALSQIAYQR.N	7	Q05DR4_HUMA
Q05DR4	3	3.7851	R.SLAALSQIAYQRNDDDEEEAARER.R	1	Q05DR4_HUMA
Q05DE8	1	2.9767	R.TYAGGTASATK.V	3	Q05DE8_HUMA
Q05CP8	4	5.0215	R.KLM#QLQHEKAELEQHLEQEQEFQVNK.L	2	Q05CP8_HUMA
Q05DR4	2	4.0159	R.SLAALSQIAYQRNDDDEEEAAR.E	2	Q05DR4_HUMA
Q9Y520	3	3.7313	K.EATPVVHETEPESGSQPRPAVLSGYFK.Q	1	Q9Y520_HUMAN
Q9Y520	2	4.0302	R.ISAVESQPSR.K	3	Q9Y520_HUMAN
Q05DE8	2	3.801	K.AASLGSSQSSR.T	11	Q05DE8_HUMA
Q05DE8	2	3.4302	K.KAASLGSSQSSR.T	4	Q05DE8_HUMA
Q05DE8	2	2.8511	K.SSSM#NPTETK.A	3	Q05DE8_HUMA
Q05DE8	2	3.2385	K.SSSMNPTETK.A	3	Q05DE8_HUMA
Q05DE8	2	3.2215	K.TSESPSKPGEK.K	13	Q05DE8_HUMA
A2RQF4	2	3.5343	R.AQQLLDAVEQR.Q	1	A2RQF4_HUMA
Q9HBR0	3	6.2378	R.DLGLAADLPGGAEGAAAQPQAVLR.Q	1	Q9HBR0_HUMA
Q7Z2U7	2	3.3017	R.FSGSNSGNTATLSISR.V	1	Q7Z2U7_HUMA
A5PL36	2	4.6705	R.GEVGAGAGPGAQAGPSAK.R	2	A5PL36_HUMAN
Q9Y520	2	2.7552	R.ESSEAQVQK.F	1	Q9Y520_HUMAN
Putative uncharacterized protein CAPG - Homo sapiens (Human)					
Q53SA7	3	3.8424	R.M#QYAPNTQVEILPQGHESPIFK.Q	1	Q53SA7_HUMA
Putative uncharacterized protein DKFZp434M1616 - Homo sapiens (Human)					
Q9NSM8	2	3.0401	K.M#NSIVYQK.Q	1	Q9NSM8_HUMA
Q9NSM8	2	3.6787	K.QRAEVLQSTQR.F	1	Q9NSM8_HUMA
Q9NSM8	2	2.7671	R.AEVLQSTQR.F	1	Q9NSM8_HUMA
Q9NSM8	2	2.9862	R.FFSEQQQSK.Q	1	Q9NSM8_HUMA
Putative uncharacterized protein DKFZp451J0218 - Homo sapiens (Human)					
Q86T83	2	3.2912	K.LEEELDDLLVDLDHQR.Q	1	Q86T83_HUMAN
Putative uncharacterized protein DKFZp564G0422 - Homo sapiens (Human)					
Q5JXL8	2	2.7828	K.REQAEEER.Y	1	Q5JXL8_HUMAN
Q5JXL8	1	2.2757	R.EQLAALKK.H	4	Q5JXL8_HUMAN
Q5JXL8	1	2.5194	R.EQLAALK.K	4	Q5JXL8_HUMAN
Q5JXL8	2	2.9993	K.REQAEEERYFR.R	1	Q5JXL8_HUMAN
Q5JXL8	1	2.2086	R.EAGGAFGK.R	2	Q5JXL8_HUMAN
Q5JXL8	1	2.1556	R.EAGGAFGKR.E	1	Q5JXL8_HUMAN
Putative uncharacterized protein DKFZp667I133 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q8ND71	2	2.9244	K.NVQEM#SQAEK.L	1	Q8ND71_HUMA
Putative uncharacterized protein DKFZp686I04144 - Homo sapiens (Human)					
Q5HYK7	2	2.9501	R.ASGEWDSGTENR.L	1	Q5HYK7_HUMA
Q5HYK7	3	4.1388	R.IKVFEGQTNIETSGLPK.K	2	Q5HYK7_HUMA
Putative uncharacterized protein DKFZp686L07201 - Homo sapiens (Human)					
Q63Z42	3	3.8521	R.WDYPEGTPNGGSTTLPSAPPPASAGLK.S	2	Q63Z42_HUMAN
Putative uncharacterized protein DKFZp761F0123 - Homo sapiens (Human)					
Q7L8J4	3	4.1655	R.IQEELEHLNQASEEINQVELQLDEAR.T	1	Q7L8J4_HUMAN
Q7L8J4	2	3.658	K.VTELEQQVAQAK.T	2	Q7L8J4_HUMAN
Putative uncharacterized protein DKFZp779L1068 - Homo sapiens (Human)					
Q629K1	3	3.888	K.DAATIKLPVDQYRK.Q	1	Q629K1_HUMAN
Putative uncharacterized protein FLJ20035 - Homo sapiens (Human)					
Q8IY21	2	3.1084	R.FYGNSLETVSSK.I	1	Q8IY21_HUMAN
Putative uncharacterized protein FLJ20628 - Homo sapiens (Human)					
Q9BVS5	2	2.7108	R.ELEDSSGDQGR.C	1	Q9BVS5_HUMA
Putative uncharacterized protein LBH - Homo sapiens (Human)					
Q53QV2	2	3.5833	R.KDGLSYQIFPDPSDFDR.C	2	Q53QV2_HUMA
Putative uncharacterized protein LOC114984 - Homo sapiens (Human)					
Q96CP2	2	3.4961	K.ALLQTHPEAQR.A	2	Q96CP2_HUMA
Q96CP2	2	4.3216	K.ASQEPSPKPGTEVIPAAPR.K	3	Q96CP2_HUMA
Q96CP2	2	3.5483	R.AIEAAPQEPEQKR.S	1	Q96CP2_HUMA
Putative uncharacterized protein LOC134121 - Homo sapiens (Human)					
A4QMS7	1	2.3315	R.LDYDQK.L	3	A4QMS7_HUMA
Putative uncharacterized protein LOC144097 - Homo sapiens (Human)					
Q9BUA3	3	4.1775	R.MPAEIVVLLDSEDNPSLPKR.S	2	Q9BUA3_HUMA
Putative uncharacterized protein LOC255374 - Homo sapiens (Human)					
A4D2B0	2	3.8589	R.GSGGAEAALEEAAR.G	1	A4D2B0_HUMAN
Putative uncharacterized protein LOC388900 - Homo sapiens (Human)					
Q5TG08	2	2.8359	K.ANHMFIPPSAVNEESPDKKTK.G	1	YV009_HUMAN
Putative uncharacterized protein MGC29506 - Homo sapiens (Human)					
Q8WU39	2	3.9947	R.AVAYQM#WQNLAK.A	8	Q8WU39_HUMA
Q8WU39	2	4.1911	R.AVAYQMWQNLAK.A	6	Q8WU39_HUMA
Q8WU39	2	4.1903	R.ELSELVYTDVLDR.S	4	Q8WU39_HUMA
Q8WU39	1	2.4729	R.NWQDYGVR.E	2	Q8WU39_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Putative uncharacterized protein PTPN18 - Homo sapiens (Human)					
Q53P42	2	3.8034	R.GAPAGAGSGTQTGTGTGTGAR.S	1	Q53P42_HUMAN
PX domain-containing protein kinase-like protein - Homo sapiens (Human)					
Q7Z7A4	3	4.7509	K.RSALENSEEHSAK.Y	1	PXK_HUMAN
Q7Z7A4	2	3.9003	R.GALLSSIQNFQK.G	4	PXK_HUMAN
Q7Z7A4	2	3.2491	R.SALENSEEHSAK.Y	1	PXK_HUMAN
Pyridoxal-dependent decarboxylase domain-containing protein 1 - Homo sapiens (Human)					
Q6P996	2	3.3763	K.AQGAGVTLPPTPSGSR.T	1	PDXD1_HUMAN
Q6P996	2	3.9937	K.GVPHPEDDHSQVEGPESLR.-	2	PDXD1_HUMAN
Pyridoxine-5'-phosphate oxidase - Homo sapiens (Human)					
Q9NVS9	2	3.9218	R.EAFEETHLTSLDPVK.Q	3	PNPO_HUMAN
Pyruvate kinase isozymes 2/3 - Homo sapiens (Human)					
P14618	2	2.8432	R.NTGIICTIGPASR.S	1	KPYM_HUMAN
P14618	2	3.6427	K.IYVDDGLISLQVK.Q	1	KPYM_HUMAN
P14618	2	3.95	R.LDIDSPPITAR.N	5	KPYM_HUMAN
Quinone oxidoreductase - Homo sapiens (Human)					
Q08257	2	3.0091	R.VFEFGGPEVLK.L	1	QOR_HUMAN
R3H domain-containing protein 1 - Homo sapiens (Human)					
Q15032	2	4.2109	R.LQDEDASSTQQR.R	2	R3HD1_HUMAN
Rab effector MyRIP - Homo sapiens (Human)					
Q8NFW9	2	2.8299	K.EEERLSELKQK.L	1	MYRIP_HUMAN
Rab GTPase-activating protein 1 - Homo sapiens (Human)					
Q9Y3P9	2	3.3622	K.IVGNGSEQQLQK.E	2	RBGP1_HUMAN
Rab GTPase-binding effector protein 1 - Homo sapiens (Human)					
Q15276	3	5.2642	R.AQASEILLEELQQGLSQAK.R	2	RABE1_HUMAN
Q15276	3	4.597	R.QNAVLQAAQDDLGHLR.T	4	RABE1_HUMAN
Q15276	2	4.5005	R.TQLWEAQAEMENIK.A	5	RABE1_HUMAN
Q15276	2	3.2545	R.MEIVLTSEQLR.Q	2	RABE1_HUMAN
Q15276	2	4.5711	R.LVSETEWNLLQK.E	3	RABE1_HUMAN
Q15276	2	4.1069	R.AQQQLEQEFNQK.R	1	RABE1_HUMAN
Q15276	2	3.5248	K.QLQGIQIQEAETR.D	2	RABE1_HUMAN
Q15276	2	2.7563	K.LSQTLQVQLER.I	1	RABE1_HUMAN
Q15276	2	3.3209	K.DQEDDEQQRLNK.R	1	RABE1_HUMAN
Q15276	2	3.5274	K.ATVEQLMFEEK.N	2	RABE1_HUMAN
Q15276	2	3.2182	K.ANDQFLESQR.L	2	RABE1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q15276	3	3.7896	K.AIATVSENTKQEAIDEVKR.Q	1	RABE1_HUMAN
Q15276	1	2.2645	K.ALGYNYK.A	1	RABE1_HUMAN
Rab GTPase-binding effector protein 2 - Homo sapiens (Human)					
Q9H5N1	2	3.0431	R.LQAQEHGAER.L	1	RABE2_HUMAN
Rab11 family-interacting protein 1 - Homo sapiens (Human)					
Q6WKZ4	2	4.0167	K.TPLSQSMSVLPTSKPEK.V	2	RFIP1_HUMAN
Q6WKZ4	3	5.8882	K.ASDHEGLLSDPLSDLQLVSDFK.S	2	RFIP1_HUMAN
Q6WKZ4	2	3.2061	K.DTAAVVSR.Q	2	RFIP1_HUMAN
Q6WKZ4	3	3.776	K.EGMLMGVKPGEDASGPAEDLVR.R	1	RFIP1_HUMAN
Rab11 family-interacting protein 3 - Homo sapiens (Human)					
O75154	2	3.6494	R.LQQLDEENSELR.S	1	RFIP3_HUMAN
O75154	2	3.6891	R.ANALEEQLKEQELR.A	1	RFIP3_HUMAN
O75154	2	3.3916	K.SIEIENLQTR.L	2	RFIP3_HUMAN
O75154	2	3.8237	K.LLDEIESLTLR.L	2	RFIP3_HUMAN
Rab11 family-interacting protein 5 - Homo sapiens (Human)					
Q9BXF6	4	4.8199	K.KKYDLESASAILPSSAIEDPDLGSLGK.M	2	RFIP5_HUMAN
Q9BXF6	4	5.3533	K.VLAPGASHPGEEEGARLPEGKPVQVATPIVA	2	RFIP5_HUMAN
Q9BXF6	2	4.6691	R.ALLDLQGHLDAASR.S	2	RFIP5_HUMAN
Q9BXF6	3	4.3612	R.KSSLTQSNTSLGSDSTLSSASGSLAYQGPGA	1	RFIP5_HUMAN
Q9BXF6	2	3.9384	R.SSISGSLPSSGSLQAVSSR.F	4	RFIP5_HUMAN
Q9BXF6	2	3.534	R.SSWLSTEGGR.D	3	RFIP5_HUMAN
RAB3A-interacting protein-like 1 - Homo sapiens (Human)					
Q8TBN0	2	4.2761	R.GKIDMLQAEVTALK.T	1	Q8TBN0_HUMA
Rabenosyn-5 - Homo sapiens (Human)					
Q9H1K0	2	3.1641	K.TPSLSSTQPTR.V	2	RBNS5_HUMAN
Q9H1K0	2	2.892	R.VWSGPPAVGQER.L	2	RBNS5_HUMAN
Rab-interacting lysosomal protein - Homo sapiens (Human)					
Q96NA2	2	3.9153	K.AESSEDETSSPAPSK.L	2	RILP_HUMAN
Radixin - Homo sapiens (Human)					
P35241	2	3.2063	R.AKEEAERLEK.E	1	RADI_HUMAN
P35241	2	3.2644	R.SEEERVTETQK.N	2	RADI_HUMAN
P35241	2	4.452	K.TQNDVLHAENVK.A	1	RADI_HUMAN
P35241	2	4.6018	K.QLQALSSELAQAR.D	4	RADI_HUMAN
P35241	2	4.6731	K.KTQNDVLHAENVK.A	1	RADI_HUMAN
P35241	2	3.0082	K.AQKELEEQTR.K	1	RADI_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P35241	2	2.8776	K.QRIDEFEAM.-	1	RADI_HUMAN
P35241	2	3.9846	R.SEEERVTETQKNER.V	4	RADI_HUMAN
Raftlin - Homo sapiens (Human)					
Q14699	2	2.9032	K.QPSSPSGEGDGGELSPQGVSK.T	2	RFTN1_HUMAN
RalA-binding protein 1 - Homo sapiens (Human)					
Q15311	2	2.9766	R.RVEHGSGLTR.T	1	RBP1_HUMAN
Ran-binding protein 3 - Homo sapiens (Human)					
Q9H6Z4	2	3.2081	R.SRVEQEQEAK.M	1	RANB3_HUMAN
Q9H6Z4	3	4.0237	R.SVLRPAVLQAPQPK.A	2	RANB3_HUMAN
Ran-specific GTPase-activating protein - Homo sapiens (Human)					
P43487	2	3.9566	K.TLEEDEEELFK.M	3	RANG_HUMAN
P43487	1	3.5914	R.FLNAENAQK.F	13	RANG_HUMAN
P43487	2	4.6548	R.FASENDLPEWKER.G	3	RANG_HUMAN
P43487	2	3.5362	K.TLEEDEEELFKM#R.A	4	RANG_HUMAN
P43487	2	3.297	K.EETKEDAEEKQ.-	4	RANG_HUMAN
P43487	2	3.0269	R.FASENDLPEWK.E	2	RANG_HUMAN
Rap guanine nucleotide exchange factor 2 - Homo sapiens (Human)					
Q9Y4G8	2	4.4711	K.ILSLSEEGSLER.H	1	RPGF2_HUMAN
Q9Y4G8	2	3.2768	R.ESLEQAQSR.A	1	RPGF2_HUMAN
Q9Y4G8	2	3.1638	R.KPPDYNVALQR.S	2	RPGF2_HUMAN
Q9Y4G8	3	4.7371	R.YSIPDLAVDVEQVIGLEK.V	1	RPGF2_HUMAN
Ras and Rab interactor 2 - Homo sapiens (Human)					
Q8WYP3	3	4.1782	K.TLSGGRPGAGPELELGTAGSPGGAPPEAAP	1	RIN2_HUMAN
Ras association domain-containing protein 8 - Homo sapiens (Human)					
Q8NHQ8	2	4.1178	R.QVNLQQFIQQTGTK.V	1	RASF8_HUMAN
Ras GTPase-activating protein-binding protein 1 - Homo sapiens (Human)					
Q13283	2	4.2773	K.SSSPAPADIAQTVQEDLR.T	1	G3BP1_HUMAN
Q13283	2	3.3808	R.TFSWASVTSK.N	2	G3BP1_HUMAN
Ras GTPase-activating protein-binding protein 2 - Homo sapiens (Human)					
Q9UN86	3	4.4549	K.GVGGKLPNFGFVVFDDSEPVQR.I	1	G3BP2_HUMAN
Q9UN86	2	3.6836	R.VEAKPEVQSQPPR.V	2	G3BP2_HUMAN
Ras GTPase-activating-like protein IQGAP1 - Homo sapiens (Human)					
P46940	2	2.8142	K.ATFYGEQVDYYK.S	1	IQGA1_HUMAN
P46940	3	4.4763	K.SVKEDSNLTLQEKK.E	1	IQGA1_HUMAN
Ras-associated and pleckstrin homology domains-containing protein 1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q70E73	2	4.0877	K.SSLSVQPGFLADLNR.T	1	RAPH1_HUMAN
Q70E73	2	3.4218	K.SSSGAEHPEPK.R	10	RAPH1_HUMAN
Q70E73	2	3.0806	K.RPSVDSLVSK.F	1	RAPH1_HUMAN
Q70E73	3	4.7276	K.KQPAFPASYIPPSPPTPPVPVPPPTLPK.Q	2	RAPH1_HUMAN
Q70E73	3	5.113	K.AGYGGSHISGYATLR.R	5	RAPH1_HUMAN
Q70E73	3	4.5156	R.LTQAEISEQPTM#ATVVPQVPTSPK.S	1	RAPH1_HUMAN
Ras-interacting protein 1 - Homo sapiens (Human)					
Q5U651	2	4.1828	K.IFGAGLASGANYK.S	2	RAIN_HUMAN
Q5U651	2	3.9784	K.KLPELAAGVAPEPPLATR.A	2	RAIN_HUMAN
Q5U651	3	4.6366	R.AASGGAALASPGPGTGSGAPAGSGGK.E	5	RAIN_HUMAN
Q5U651	3	5.1279	R.AASGGAALASPGPGTGSGAPAGSGGKER.S	2	RAIN_HUMAN
Q5U651	2	5.0149	R.GSGTGTTGSSGAGGPGTPGGAQR.W	6	RAIN_HUMAN
Q5U651	3	6.5914	R.RLEQEAFGAADSEGTGAPSWRPQK.N	1	RAIN_HUMAN
Ras-related protein Rab-10 - Homo sapiens (Human)					
P61026	2	3.0886	K.AFLTLAEDILR.K	1	RAB10_HUMAN
Ras-related protein Rab-1A - Homo sapiens (Human)					
P62820	2	2.9371	R.MGPGATAGGAEK.S	1	RAB1A_HUMAN
Ras-related protein Rab-7a - Homo sapiens (Human)					
P51149	2	4.0456	K.EAINVEQAFQTIAR.N	1	RAB7A_HUMAN
RAS-responsive element-binding protein 1 - Homo sapiens (Human)					
Q92766	2	4.2248	K.LAEGDGEAGAGGAASQEQK.L	2	RREB1_HUMAN
RBMY2FP protein - Homo sapiens (Human)					
Q8TC46	1	2.6396	K.DGTIGLKENK.F	2	Q8TC46_HUMA
Receptor expression-enhancing protein 5 - Homo sapiens (Human)					
Q00765	2	2.9125	K.KATVNLLGEEK.K	1	REEP5_HUMAN
Receptor-interacting serine/threonine-protein kinase 2 - Homo sapiens (Human)					
O43353	2	2.9239	R.LQPGIAQQWIQSK.R	2	RIPK2_HUMAN
O43353	2	2.9573	R.SPSLNLLQNK.S	1	RIPK2_HUMAN
Regulator of G-protein signaling 10 - Homo sapiens (Human)					
O43665	2	3.2187	K.MQDKTQMQEK.A	2	RGS10_HUMAN
O43665	2	3.1938	R.TEEEEEDLPDAQTAAKR.A	1	RGS10_HUMAN
O43665	2	3.7748	K.WAASLENLLEDPEGVKR.F	2	RGS10_HUMAN
O43665	2	3.5416	K.ILEEPHPLMFQK.L	1	RGS10_HUMAN
O43665	2	2.8035	K.ASSQVNVEGQSR.L	1	RGS10_HUMAN
O43665	2	3.9507	R.TEEEEEDLPDAQTAAK.R	3	RGS10_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Regulator of G-protein signaling 12 - Homo sapiens (Human)					
O14924	2	5.008	R.LPPGSTELTLPTPAAVAK.G	2	RGS12_HUMAN
Regulator of nonsense transcripts 1 - Homo sapiens (Human)					
Q92900	2	3.8255	R.AYQHGGVTGLSQY.-	6	RENT1_HUMAN
Regulator of nonsense transcripts 3A - Homo sapiens (Human)					
Q9H1J1	2	3.2978	K.AEGSGTGPEKREEAE.-	1	REN3A_HUMAN
Q9H1J1	2	3.5905	K.GSQDSGAPGEAM#ER.L	3	REN3A_HUMAN
Q9H1J1	2	2.9526	R.SQEQESEAQR.Y	1	REN3A_HUMAN
Regulator of nonsense transcripts 3B - Homo sapiens (Human)					
Q9BZI7	2	4.4868	K.KAESTESIGSSEK.T	2	REN3B_HUMAN
Q9BZI7	3	4.9346	K.KAESTESIGSSEKTEK.K	2	REN3B_HUMAN
Q9BZI7	2	2.9516	K.FLESYATDNEK.M	1	REN3B_HUMAN
Q9BZI7	2	3.8086	K.NQEDIILFR.D	1	REN3B_HUMAN
Remodeling and spacing factor 1 - Homo sapiens (Human)					
Q96T23	2	2.9418	K.DADSSISVLEIHSQK.A	1	RSF1_HUMAN
Q96T23	3	4.4045	R.FDEFDEAIDEAIEDDIKEADGGGVGR.G	1	RSF1_HUMAN
Replication factor C subunit 1 - Homo sapiens (Human)					
P35251	2	4.5058	K.DTEAGETFSSVQANLSK.A	2	RFC1_HUMAN
P35251	2	2.9577	K.ELSQNTDESGLNDEAIAK.Q	1	RFC1_HUMAN
P35251	2	2.8645	K.TKSDEETLK.A	2	RFC1_HUMAN
Replication protein A 14 kDa subunit - Homo sapiens (Human)					
P35244	3	4.0003	K.IIHDFPQFYPLGIVQHD.-	1	RFA3_HUMAN
Response gene to complement 32 protein - Homo sapiens (Human)					
Q9H4X1	2	2.8122	K.ELEAFIADLDK.T	1	RGC32_HUMAN
Q9H4X1	2	3.1609	R.SSASVSDSSGFSDSESADSLYR.N	1	RGC32_HUMAN
Q9H4X1	3	5.1191	K.LGDTKELEAFIADLDK.T	2	RGC32_HUMAN
Q9H4X1	2	3.9118	K.LNSPTDSTPALLSATVTPQK.A	1	RGC32_HUMAN
Reticulocalbin-1 precursor - Homo sapiens (Human)					
Q15293	2	3.85	R.EQFNEFRDLNK.D	2	RCN1_HUMAN
Q15293	2	3.8577	K.TFDQLTPDESKER.L	3	RCN1_HUMAN
Q15293	1	2.6193	R.YIFDNVAK.V	5	RCN1_HUMAN
Q15293	2	2.9807	R.IDNDGDGFVTTEELK.T	2	RCN1_HUMAN
Q15293	3	4.2383	R.HWILPQDYDHAQAEAR.H	5	RCN1_HUMAN
Q15293	2	4.7945	R.HLVYESDKNKDEK.L	2	RCN1_HUMAN
Q15293	2	4.8636	K.AADLNGDLTATR.E	2	RCN1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q15293	2	2.7805	K.RYIFDNVAK.V	1	RCN1_HUMAN
Q15293	4	6.1303	K.LTKEEILENWNMFVGSQATNYGEDLTKNHDE	2	RCN1_HUMAN
Q15293	2	5.0078	K.IVDRIDNDGDGFVTTEELK.T	22	RCN1_HUMAN
Q15293	2	2.8042	K.DGKLDKDEIR.H	2	RCN1_HUMAN
Q15293	2	4.5175	R.FKAADLNGDLTATR.E	1	RCN1_HUMAN
Reticulocalbin-2 precursor - Homo sapiens (Human)					
Q14257	3	4.2843	R.SDYDREALLGVQEDVDEYVK.L	1	RCN2_HUMAN
Q14257	2	3.1295	K.HYAMQEAK.Q	2	RCN2_HUMAN
Q14257	2	4.6199	K.LSEEEILENPDLFLTSEATDYGR.Q	2	RCN2_HUMAN
Reticulocalbin-3 precursor - Homo sapiens (Human)					
Q96D15	3	4.983	R.VHQAAPLSDAPHDDAHGNFQYDHEAFLGR.E	2	RCN3_HUMAN
Q96D15	2	3.225	R.VADQDGDSMATR.E	1	RCN3_HUMAN
Q96D15	2	3.415	R.VADQDGDSM#ATR.E	3	RCN3_HUMAN
Q96D15	2	5.115	R.EVAKEFDQLTPEESQAR.L	3	RCN3_HUMAN
Q96D15	2	5.185	R.DIVIAETLEDLDRNK.D	1	RCN3_HUMAN
Q96D15	2	3.7092	K.EFDQLTPEESQAR.L	1	RCN3_HUMAN
Q96D15	2	4.4691	R.AGDGDGWVSLAELR.A	5	RCN3_HUMAN
Reticulon-1 - Homo sapiens (Human)					
Q16799	3	4.4635	R.HRGEGENEAVTPK.G	4	RTN1_HUMAN
Q16799	2	3.0756	R.GLFSSDSGIEM#TPAESTEVNK.I	1	RTN1_HUMAN
Q16799	2	2.9982	R.RAPQITTPVK.I	1	RTN1_HUMAN
Q16799	3	3.8561	K.GLSYETAENPRPVGQLADRPEVK.A	1	RTN1_HUMAN
Q16799	2	3.5409	K.GATPAPQAGEPSPGLGAR.A	2	RTN1_HUMAN
Q16799	2	4.2664	K.ILADPLDQMKAEAYK.Y	2	RTN1_HUMAN
Retinal dehydrogenase 1 - Homo sapiens (Human)					
P00352	2	4.5062	K.GYFVQPTVFSNVTDEMR.I	2	AL1A1_HUMAN
Retinoblastoma-associated protein - Homo sapiens (Human)					
P06400	2	3.134	K.ISEGLPTPTK.M	2	RB_HUMAN
Retinoblastoma-binding protein 6 - Homo sapiens (Human)					
Q7Z6E9	3	4.2431	K.DVSHEIIQHEVK.S	2	RBBP6_HUMAN
Q7Z6E9	2	3.458	K.EHQETKPVKEEK.V	2	RBBP6_HUMAN
Rho GDP-dissociation inhibitor 1 - Homo sapiens (Human)					
P52565	2	4.8268	K.SIQEIQELDKDDESLR.K	1	GDIR_HUMAN
P52565	3	3.9063	K.SIQEIQELDKDDESLRK.Y	3	GDIR_HUMAN
Rho GDP-dissociation inhibitor 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P52566	3	4.2295	K.ATFMVGSYGPRPEEYEFLTPVEEAPK.G	1	GDIS_HUMAN
P52566	2	4.0339	K.TLLGDGPVVTDPK.A	4	GDIS_HUMAN
P52566	2	5.3977	K.SLKELQEMDKDDESLIK.Y	7	GDIS_HUMAN
P52566	2	4.7325	K.ELQEMDKDDESLIK.Y	4	GDIS_HUMAN
P52566	3	3.9669	K.SLKELQEM#DKDDESLIK.Y	2	GDIS_HUMAN
Rho GTPase-activating protein 17 - Homo sapiens (Human)					
Q68EM7	2	4.3144	K.DPVSAAVPAPGR.N	3	RHG17_HUMAN
Q68EM7	2	2.8883	K.TEVLSEDLLQIER.R	1	RHG17_HUMAN
Q68EM7	3	4.4925	R.KHISPAFQPPLPPTDGSTVVPAGPEPPPQSS	2	RHG17_HUMAN
Q68EM7	3	5.3472	R.NRPSVPPPPQPPGVHSAGDSSLTNTAPTASK	1	RHG17_HUMAN
Q68EM7	3	5.3913	R.SPSPPTQHTGQPPGQPSAPSQLSAPR.R	3	RHG17_HUMAN
Rho GTPase-activating protein 18 - Homo sapiens (Human)					
Q8N392	2	3.3415	K.STNDADVPQGVIR.V	2	RHG18_HUMAN
Q8N392	2	3.1589	R.YGQYTMNQESTTIK.V	2	RHG18_HUMAN
Q8N392	2	3.715	K.RVETVSQTLR.K	3	RHG18_HUMAN
Q8N392	2	3.9275	R.FLSQESGVAQTLK.K	2	RHG18_HUMAN
Q8N392	2	4.2011	R.DDEASNLVGEEK.L	3	RHG18_HUMAN
Rho GTPase-activating protein 21 - Homo sapiens (Human)					
Q5T5U3	2	2.7674	R.TDSAPDQQVETGK.S	1	Q5T5U3_HUMA
Rho GTPase-activating protein 23 - Homo sapiens (Human)					
Q9P227	2	2.7824	R.WQDLNVISSLLKSFFRK.L	1	RHG23_HUMAN
Rho GTPase-activating protein 24 - Homo sapiens (Human)					
Q8N264	2	3.1412	R.M#GILNSDTLGNPTNVR.N	1	RHG24_HUMAN
Rho GTPase-activating protein 25 - Homo sapiens (Human)					
P42331	2	2.8492	K.SAALEISLR.N	1	RHG25_HUMAN
P42331	3	4.9976	R.TDSFSSMTSDSDTTSPTGQQPSDAFPEDSSK	2	RHG25_HUMAN
P42331	2	4.5051	K.NSGEEEIDSLQR.M	3	RHG25_HUMAN
P42331	3	4.6959	K.MEIFKNEFWSPSSEAK.A	2	RHG25_HUMAN
P42331	2	3.936	K.ALEEEVKEFVK.S	3	RHG25_HUMAN
Rho GTPase-activating protein 29 - Homo sapiens (Human)					
O15463	3	5.618	K.GVTTSLQISGDHSINATQPSKPYAEPVR.S	1	O15463_HUMAN
O15463	3	3.9967	K.IQDKQYEQNSLTAK.T	1	O15463_HUMAN
O15463	2	3.0111	K.FNGFDQQTLQK.I	2	O15463_HUMAN
Rho GTPase-activating protein 30 - Homo sapiens (Human)					
Q7Z6I6	3	4.1749	K.ASEDRGEAGGSQETK.V	9	RHG30_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q7Z6I6	2	3.4455	R.ESGDGEAEGDQR.A	6	RHG30_HUMAN
Q7Z6I6	2	3.5554	R.LASTLVQVQQVR.S	1	RHG30_HUMAN
Q7Z6I6	2	3.0596	R.SYAFETQANPGKGEGL.-	1	RHG30_HUMAN
Rho GTPase-activating protein 6 - Homo sapiens (Human)					
O43182	2	3.423	R.LGDAGWLDWQR.E	1	RHG06_HUMAN
Rho guanine nucleotide exchange factor 12 - Homo sapiens (Human)					
Q9NZN5	3	3.7319	R.TDCSSGDASRPSSDNADSPK.S	1	ARHGC_HUMAN
Q9NZN5	3	4.5634	R.QSGLANEGTDAGYLPANSMSSVASGASFSQ	1	ARHGC_HUMAN
Q9NZN5	2	3.1405	K.QLLVQQLGLTEK.S	2	ARHGC_HUMAN
Q9NZN5	2	3.0556	K.QVGETSAPGDTLDGTPR.T	1	ARHGC_HUMAN
Q9NZN5	2	2.9048	R.GFPSILGPPR.R	1	ARHGC_HUMAN
Rho guanine nucleotide exchange factor 15 - Homo sapiens (Human)					
O94989	2	4.389	R.AQDADAPEPGLQAR.A	2	ARHGF_HUMAN
Rho guanine nucleotide exchange factor 16 - Homo sapiens (Human)					
Q5VV41	2	5.202	R.LDAGGNPASGLPMVR.G	4	ARHGG_HUMAN
Q5VV41	3	4.2382	K.GLGKPGGQGDAIQLSPK.L	1	ARHGG_HUMAN
Q5VV41	1	2.7479	K.SLAVASK.A	2	ARHGG_HUMAN
Q5VV41	2	3.632	R.LDAGGNPASGLPM#VR.G	3	ARHGG_HUMAN
Rho guanine nucleotide exchange factor 17 - Homo sapiens (Human)					
Q96PE2	2	5.2818	R.MGAQQDDGSDAPPGSPDWAGDVTR.G	7	ARHGH_HUMAN
Q96PE2	2	5.117	R.RDEGSQDQTGSLSR.A	2	ARHGH_HUMAN
Q96PE2	3	4.5559	R.RPSADSESPGTPSPDGAAWEPPAR.E	2	ARHGH_HUMAN
Q96PE2	2	3.4809	R.SARPLTGPETEGR.L	3	ARHGH_HUMAN
Q96PE2	2	2.9213	R.SLSDPIPQR.H	1	ARHGH_HUMAN
Q96PE2	2	3.0758	R.LDDGSAGTR.D	2	ARHGH_HUMAN
Q96PE2	2	3.9011	R.YSSTETLKDDDLWSSR.G	2	ARHGH_HUMAN
Q96PE2	3	4.2859	R.ALRDGGFEPEK.S	1	ARHGH_HUMAN
Q96PE2	2	4.5922	R.SPSFGAGEGLLR.S	5	ARHGH_HUMAN
Q96PE2	2	3.0786	R.GSGGWGVYR.S	2	ARHGH_HUMAN
Q96PE2	2	5.412	R.DGGVLPAAAEEAAEGPAR.G	3	ARHGH_HUMAN
Q96PE2	3	3.778	R.AQRPADGLHSWHIFSQPQAGAR.A	1	ARHGH_HUMAN
Q96PE2	2	3.1816	R.ALPEALPPPATAHR.N	1	ARHGH_HUMAN
Q96PE2	3	4.4907	R.AATSEEPTGFSVDSNLLGSLSPK.T	3	ARHGH_HUMAN
Q96PE2	2	4.1082	K.VSFPSYLASPAGSR.G	3	ARHGH_HUMAN
Q96PE2	2	3.8137	K.SLSNPDIASETLTLLSFLR.S	3	ARHGH_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96PE2	2	3.594	K.LADILSPR.L	2	ARHGH_HUMAN
Q96PE2	2	5.3118	R.DEGSQDQTGSLSR.A	10	ARHGH_HUMAN
Rho-associated protein kinase 2 - Homo sapiens (Human)					
O75116	2	2.9636	K.IQQNQSIR.R	2	ROCK2_HUMAN
Ribonucleases P/MRP protein subunit POP1 - Homo sapiens (Human)					
Q99575	2	3.7055	K.AGPEGTSQEIPK.Y	1	POP1_HUMAN
Q99575	3	4.2387	R.VNPHSLPDPEVNEQSSSK.G	2	POP1_HUMAN
Q99575	2	2.7887	K.YITASTFAQAR.A	1	POP1_HUMAN
Ribonucleoprotein PTB-binding 2 - Homo sapiens (Human)					
Q9HCJ3	3	4.5265	K.TGIASSILDAISQGSESQHALEK.C	2	RAVR2_HUMAN
Ribosomal L1 domain-containing protein 1 - Homo sapiens (Human)					
O76021	3	3.744	K.IKEEAVKEK.S	4	RL1D1_HUMAN
Ribosomal protein S6 kinase alpha-3 - Homo sapiens (Human)					
P51812	2	2.8136	R.NQSPVLEPVGR.S	1	KS6A3_HUMAN
Ribosome biogenesis protein BMS1 homolog - Homo sapiens (Human)					
Q14692	3	3.9646	R.FLVEAPHDWDLEEVMNSIR.D	1	BMS1_HUMAN
Ribosome biogenesis regulatory protein homolog - Homo sapiens (Human)					
Q15050	2	3.6029	K.AEQDEAEKLQR.I	1	RRS1_HUMAN
Ribosome production factor 1 - Homo sapiens (Human)					
Q9H9Y2	2	2.9316	K.VGIQELGPR.F	2	RPF1_HUMAN
Ribosome recycling factor, mitochondrial precursor - Homo sapiens (Human)					
Q96E11	2	2.8824	K.SKDTVSEDTIR.L	1	RRFM_HUMAN
Ribosome-binding protein 1 - Homo sapiens (Human)					
Q9P2E9	2	4.7139	K.LLATEQEDAAVAK.S	5	RRBP1_HUMAN
Q9P2E9	2	3.7587	K.LKGELESSDQVR.E	3	RRBP1_HUMAN
Q9P2E9	2	3.3004	K.KTESASVQGR.N	6	RRBP1_HUMAN
Q9P2E9	2	5.1691	K.KLQEQLEKAEDGSSSK.E	4	RRBP1_HUMAN
Q9P2E9	2	2.8706	K.LQEQLEK.A	1	RRBP1_HUMAN
Q9P2E9	3	5.8756	K.KKGEPGPPDADGPLYLPYK.T	3	RRBP1_HUMAN
Q9P2E9	2	3.4093	K.QVLQLQASHR.E	4	RRBP1_HUMAN
Q9P2E9	2	4.4949	K.KGEPGPPDADGPLYLPYK.T	5	RRBP1_HUMAN
Q9P2E9	2	2.8688	K.KLQEQLEK.A	1	RRBP1_HUMAN
Q9P2E9	2	5.2276	K.LQEQLEKAEDGSSSK.E	4	RRBP1_HUMAN
Q9P2E9	3	4.8723	K.LQEQLEKAEDGSSSKEGTSV.-	4	RRBP1_HUMAN
Q9P2E9	2	3.115	K.LQSSEAEVR.S	4	RRBP1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9P2E9	3	4.2068	K.LRELNKEMAAEK.A	1	RRBP1_HUMAN
Q9P2E9	2	3.4767	K.RLDEVSR.E	3	RRBP1_HUMAN
Q9P2E9	2	3.7553	K.SVEEEEQVWR.A	5	RRBP1_HUMAN
Q9P2E9	3	4.8214	K.SEAQKQSDELALVR.Q	3	RRBP1_HUMAN
Q9P2E9	2	5.8657	K.ADSVANQGTKVEGITNQGK.K	5	RRBP1_HUMAN
Q9P2E9	3	4.9646	K.KGEGAPIQGKK.A	4	RRBP1_HUMAN
Q9P2E9	2	5.8066	K.LREAEETQSTLQAECDQYR.S	3	RRBP1_HUMAN
Q9P2E9	2	3.621	K.EAIELREAVEQQK.V	1	RRBP1_HUMAN
Q9P2E9	3	4.8767	K.SHVEDGDIAGAPASSPEAPPAEQDPVQLK.T	2	RRBP1_HUMAN
Q9P2E9	2	4.6947	K.AEDGSSSKEGTSV.-	10	RRBP1_HUMAN
Q9P2E9	3	4.1348	K.AEGAQNQGQKGEGAQNQGKK.T	2	RRBP1_HUMAN
Q9P2E9	2	3.009	K.AGIIQDTWHK.A	2	RRBP1_HUMAN
Q9P2E9	2	4.0884	K.AM#EALATAEQACK.E	2	RRBP1_HUMAN
Q9P2E9	2	4.8176	K.AMEALATAEQACK.E	5	RRBP1_HUMAN
Q9P2E9	2	4.9097	K.AQEQQQQMAELHSK.L	5	RRBP1_HUMAN
Q9P2E9	3	4.9002	K.AEGAQNQGQKGEGAQNQGK.K	2	RRBP1_HUMAN
Q9P2E9	3	4.1534	K.ATQKGDPVAILKR.Q	1	RRBP1_HUMAN
Q9P2E9	2	3.7446	K.KGEGAPIQGK.K	9	RRBP1_HUMAN
Q9P2E9	2	2.7969	K.GDPVAILKR.Q	1	RRBP1_HUMAN
Q9P2E9	1	2.2766	K.GEGAPIQGK.K	1	RRBP1_HUMAN
Q9P2E9	2	2.773	K.GEPGPPDADGPLYLPYK.T	1	RRBP1_HUMAN
Q9P2E9	4	5.4076	K.HLEEIVEKLKGELESSDQVR.E	1	RRBP1_HUMAN
Q9P2E9	2	2.8972	K.HPPAPAEPSSDLASK.L	1	RRBP1_HUMAN
Q9P2E9	3	4.2515	K.IRTLQEQLENGPNTQLAR.L	1	RRBP1_HUMAN
Q9P2E9	3	5.5276	K.KADSVANQGTKVEGITNQGK.K	3	RRBP1_HUMAN
Q9P2E9	3	7.0508	K.KAEGAQNQGQKGEGAQNQGK.K	11	RRBP1_HUMAN
Q9P2E9	2	3.1332	K.ATQKGDPVAILK.R	2	RRBP1_HUMAN
Q9P2E9	2	4.15	R.SIEALLEAGQAR.D	5	RRBP1_HUMAN
Q9P2E9	2	3.1278	R.LIEILSEK.A	1	RRBP1_HUMAN
Q9P2E9	2	3.1649	K.SEAVRQDEQQR.K	3	RRBP1_HUMAN
Q9P2E9	3	5.9689	R.LQQENSILRDALNQATSQVESK.Q	3	RRBP1_HUMAN
Q9P2E9	2	3.5477	R.NTDVAQSPEAPK.Q	2	RRBP1_HUMAN
Q9P2E9	2	5.61	R.NTDVAQSPEAPKQEAPAK.K	8	RRBP1_HUMAN
Q9P2E9	3	4.1279	R.QKLTAEFEEAQTSACR.L	1	RRBP1_HUMAN
Q9P2E9	2	3.0203	R.EQEITAVQAR.M	2	RRBP1_HUMAN
Q9P2E9	3	4.4082	R.QLLLESQSQLDAAKSEAQK.Q	2	RRBP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9P2E9	1	2.6921	R.ESEEALQK.R	4	RRBP1_HUMAN
Q9P2E9	2	3.1711	R.SILAETEGM#LR.D	1	RRBP1_HUMAN
Q9P2E9	2	4.2053	R.SILAETEGMLR.D	8	RRBP1_HUMAN
Q9P2E9	2	4.8326	R.SILAETEGMLRDLQK.S	2	RRBP1_HUMAN
Q9P2E9	3	4.3331	R.SKCEELSGLHGQLQEAR.A	2	RRBP1_HUMAN
Q9P2E9	2	4.9255	R.TAGPLESSETEEASQLKER.L	6	RRBP1_HUMAN
Q9P2E9	2	5.3428	R.TLQEQLENGPNTQLAR.L	3	RRBP1_HUMAN
Q9P2E9	2	4.253	R.QLLLESQSQLDAAK.S	1	RRBP1_HUMAN
Q9P2E9	2	4.0812	K.VGAAEEELQK.S	7	RRBP1_HUMAN
Q9P2E9	2	2.976	K.TESASVQGR.N	3	RRBP1_HUMAN
Q9P2E9	3	5.7604	K.TLVSTVGSM#VFNEGEAQR.L	1	RRBP1_HUMAN
Q9P2E9	3	6.6944	K.TLVSTVGSMVFNEGEAQR.L	12	RRBP1_HUMAN
Q9P2E9	3	5.2303	K.TQLEWTEAILEDEQTQR.Q	6	RRBP1_HUMAN
Q9P2E9	2	4.8454	R.LKELESQVSGLEK.E	3	RRBP1_HUMAN
Q9P2E9	2	2.8938	K.VEGITNQGK.K	1	RRBP1_HUMAN
Q9P2E9	2	3.0709	R.ELNKEMAAEK.A	3	RRBP1_HUMAN
Q9P2E9	3	6.1335	R.ADAEKAQEQQQQM#AELHSK.L	2	RRBP1_HUMAN
Q9P2E9	2	6.2753	R.ADAEKAQEQQQQMAELHSK.L	4	RRBP1_HUMAN
Q9P2E9	2	3.1755	R.AENSQLTER.I	3	RRBP1_HUMAN
Q9P2E9	2	4.9917	R.AKVGAAEEELQK.S	8	RRBP1_HUMAN
Q9P2E9	2	4.5325	R.DALNQATSQVESK.Q	4	RRBP1_HUMAN
Q9P2E9	2	5.6586	R.DAQDVQASQAEADQQQTR.L	12	RRBP1_HUMAN
Q9P2E9	3	4.1742	R.EAEETQSTLQAECDQYR.S	2	RRBP1_HUMAN
Q9P2E9	2	4.0864	R.EHVKEVQQLQGK.I	5	RRBP1_HUMAN
Q9P2E9	1	2.8763	K.TTQEQLAR.E	3	RRBP1_HUMAN
Ribulose-phosphate 3-epimerase - Homo sapiens (Human)					
Q96AT9	2	3.6845	K.IGPSILNSDLANLGAECLR.M	2	RPE_HUMAN
RING finger protein 10 - Homo sapiens (Human)					
Q8N5U6	2	3.0275	R.SASAGPAGESKPK.S	2	RNF10_HUMAN
RING finger protein 113A - Homo sapiens (Human)					
O15541	2	3.1139	K.AVDQVCTFLFK.K	1	R113A_HUMAN
RING finger protein 214 - Homo sapiens (Human)					
Q8ND24	2	3.4142	R.VDQDDDQDSSSLK.L	2	RN214_HUMAN
RING finger protein C13orf7 - Homo sapiens (Human)					
Q5W0B1	2	2.8665	K.EYEDEIDC*LQKEVEELK.S	1	CM007_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
RING finger protein C14orf4 - Homo sapiens (Human)					
Q9H1B7	2	3.3521	K.DGSSVHSTTASAR.R	1	CN004_HUMAN
RING1 and YY1-binding protein - Homo sapiens (Human)					
Q8N488	3	3.707	K.DKEISPSVTKK.N	1	RYBP_HUMAN
Q8N488	3	5.0456	K.SDILKDPPSEANSIQSANATTK.T	1	RYBP_HUMAN
Q8N488	3	3.9134	R.INSQLVAQQVAQQYATPPPPKK.E	1	RYBP_HUMAN
Q8N488	3	4.5116	R.SSSTSSSTVTSSAGSEQQNQSSSGSESTDK	1	RYBP_HUMAN
RNA 3'-terminal phosphate cyclase-like protein - Homo sapiens (Human)					
Q9Y2P8	2	2.8738	K.QFGIDGESFELK.I	1	RCL1_HUMAN
RNA polymerase II-associated protein 1 - Homo sapiens (Human)					
Q9BWH6	2	3.2179	R.LLAQLDPSLVAFLR.S	1	RPAP1_HUMAN
RNA polymerase-associated protein LEO1 - Homo sapiens (Human)					
Q8WVC0	2	3.6821	K.LTSDEEGEPSGK.R	4	LEO1_HUMAN
Q8WVC0	2	3.4493	R.TEMIKKEEER.L	1	LEO1_HUMAN
Q8WVC0	3	4.1148	R.RDEEGNEIKESNAR.I	2	LEO1_HUMAN
Q8WVC0	3	4.2632	R.IEVEIPKVNTDLGNDLYFVK.L	1	LEO1_HUMAN
Q8WVC0	2	4.1491	K.VNTDLGNDLYFVK.L	3	LEO1_HUMAN
Q8WVC0	3	4.3518	K.KLTSDEEGEPSGKR.K	2	LEO1_HUMAN
Q8WVC0	2	4.5798	R.DEEGNEIKESNAR.I	2	LEO1_HUMAN
RNA polymerase-associated protein RTF1 homolog - Homo sapiens (Human)					
Q92541	2	2.7205	K.YGSGVLPDAPK.E	1	RTF1_HUMAN
RNA U small nuclear RNA export adapter protein - Homo sapiens (Human)					
Q9H814	2	2.7175	R.QSETYNYLLAK.K	1	RNUXA_HUMAN
Q9H814	2	4.4398	R.TPGGVFLNLLK.N	4	RNUXA_HUMAN
RNA-binding motif protein, X-linked 2 - Homo sapiens (Human)					
Q9Y388	2	2.7159	R.TAYSGGAEDLER.E	1	RBMX2_HUMAN
RNA-binding protein 14 - Homo sapiens (Human)					
Q96PK6	2	3.7099	R.LSESQLSFR.R	5	RBM14_HUMAN
Q96PK6	2	3.1429	R.RLPDAHSDYAR.Y	1	RBM14_HUMAN
Q96PK6	2	2.7793	R.YSGSYNDYLR.A	1	RBM14_HUMAN
RNA-binding protein 26 - Homo sapiens (Human)					
Q5T8P6	3	5.0806	K.TRAEAEAAAVHGAR.F	3	RBM26_HUMAN
Q5T8P6	2	3.6002	R.AEAEAAAVHGAR.F	2	RBM26_HUMAN
RNA-binding protein 27 - Homo sapiens (Human)					
Q9P2N5	2	3.2744	K.LSQLQVEAAR.L	1	RBM27_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9P2N5	2	3.1865	K.TSSAVSTPSK.V	1	RBM27_HUMAN
Q9P2N5	2	3.3019	R.DQPGTSAVPNLASVGTR.L	2	RBM27_HUMAN
Q9P2N5	2	2.7802	R.LQLGTPPPLLAAR.L	1	RBM27_HUMAN
RNA-binding protein 28 - Homo sapiens (Human)					
Q9NW13	2	3.1812	R.FNQLVEQYK.Q	2	RBM28_HUMAN
RNA-binding protein 35B - Homo sapiens (Human)					
Q9H6T0	2	2.8653	R.PTGDAFALFAC*EELAQAALRR.H	1	RB35B_HUMAN
RNA-binding protein 8A - Homo sapiens (Human)					
Q9Y5S9	3	5.5181	R.MREDYDSVEQDGDEPGPQR.S	12	RBM8A_HUMAN
Q9Y5S9	2	5.1125	R.M#REDYDSVEQDGDEPGPQR.S	6	RBM8A_HUMAN
Q9Y5S9	2	3.7555	R.EDYDSVEQDGDEPGPQR.S	3	RBM8A_HUMAN
Q9Y5S9	2	2.7379	K.NIHLNLDRR.T	1	RBM8A_HUMAN
Q9Y5S9	2	3.2181	K.GRGFGSEEGSR.A	2	RBM8A_HUMAN
Q9Y5S9	2	3.4034	K.GYTLVEYETYK.E	1	RBM8A_HUMAN
RNA-binding protein MEX3C - Homo sapiens (Human)					
Q5U5Q3	2	3.7741	R.APAAAAQGQAR.R	1	MEX3C_HUMAN
RNA-binding protein NOB1 - Homo sapiens (Human)					
Q9ULX3	2	2.8065	R.RLNPNASR.K	1	NOB1_HUMAN
RNA-binding protein with multiple splicing - Homo sapiens (Human)					
Q93062	2	3.3412	R.TLFVSGLPLDIKPR.E	1	RBPMS_HUMAN
Q93062	3	4.4772	K.NKLVGTPNPSTPLPNTVPQFIAR.E	2	RBPMS_HUMAN
Q93062	3	4.0019	K.NALNGIRFDPEIPQTLR.L	1	RBPMS_HUMAN
RNA-binding protein with serine-rich domain 1 - Homo sapiens (Human)					
Q15287	3	4.2261	R.NVTKDHIMEIFSTYGK.I	2	RNPS1_HUMAN
Q15287	2	4.3374	K.GYAYVEFENPDEAEK.A	2	RNPS1_HUMAN
Round spermatid basic protein 1. - Homo sapiens (Human)					
Q5VWQ0	2	3.9406	R.AVAAQEEPDKEGK.E	2	Q5VWQ0_HUMA
RPL14 protein - Homo sapiens (Human)					
Q6IPH7	2	6.313	K.GTAAAAAAAAAAAAAAAK.V	7	Q6IPH7_HUMAN
RRP15-like protein - Homo sapiens (Human)					
Q9Y3B9	2	2.7208	K.KDFISVLR.G	1	RRP15_HUMAN
Q9Y3B9	2	3.1051	K.SEEGPGWTILR.D	2	RRP15_HUMAN
Q9Y3B9	2	3.7725	R.GM#DGSTNETASSR.K	6	RRP15_HUMAN
Q9Y3B9	2	4.0858	R.GMDGSTNETASSR.K	4	RRP15_HUMAN
RRP1-like protein - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P56182	2	3.0773	K.AANVQEPEKK.K	2	RRP1_HUMAN
P56182	2	4.5761	R.AEAGEQPGTAER.A	4	RRP1_HUMAN
RRP1-like protein B - Homo sapiens (Human)					
Q14684	2	3.3454	K.TPTSSPASSPLVAK.K	1	RRP1B_HUMAN
RUN and FYVE domain-containing protein 1 - Homo sapiens (Human)					
Q96T51	3	5.3577	R.GRELEPELEPGPGPGSALEPGEEFEIVDR.S	2	RUFY1_HUMAN
RUN and FYVE domain-containing protein 4 - Homo sapiens (Human)					
Q6ZNE9	2	2.709	R.APWIEIFLGNSTPSTQGQGK.G	1	RUFY4_HUMAN
RUN domain-containing protein 2A - Homo sapiens (Human)					
Q9HA26	2	4.0181	K.TPGAGESSEDNSDR.S	2	RUN2A_HUMAN
RWD domain-containing protein 1 - Homo sapiens (Human)					
Q9H446	2	4.1995	K.AKFDAELLEIK.K	4	RWDD1_HUMAN
Q9H446	3	4.0123	K.AKFDAELLEIKK.K	4	RWDD1_HUMAN
Q9H446	3	4.3179	K.QLFHGTPVTIENFLNWK.A	3	RWDD1_HUMAN
Q9H446	2	3.0725	K.RMKEEEQAGK.N	1	RWDD1_HUMAN
RWD domain-containing protein 4A - Homo sapiens (Human)					
Q6NW29	2	3.0674	R.GWNWVDVVK.H	1	RWDD4_HUMAN
SAM and SH3 domain-containing protein 1 - Homo sapiens (Human)					
O94885	2	3.5768	R.NQLGNYPTLPLMK.S	2	SASH1_HUMAN
SAM domain and HD domain-containing protein 1 - Homo sapiens (Human)					
Q9Y3Z3	3	4.3931	R.NFTKPQDGDVIAPLITPQKK.E	2	SAMH1_HUMAN
Q9Y3Z3	2	2.8142	R.GGFEEPVLLK.N	1	SAMH1_HUMAN
SAM domain-containing protein SAMSN-1 - Homo sapiens (Human)					
Q9NSI8	2	4.3078	K.TPMGMWTGMLNNK.V	2	SAMN1_HUMAN
SAP30-binding protein - Homo sapiens (Human)					
Q9UHR5	2	4.9212	K.GGLVSDAYGEDDFSR.L	5	S30BP_HUMAN
Q9UHR5	2	3.1674	K.TTVISAVGTIVKK.A	1	S30BP_HUMAN
SAPK substrate protein 1 - Homo sapiens (Human)					
Q04323	2	4.0277	R.RQGQELSAAR.Q	4	SAKS1_HUMAN
Q04323	3	3.8098	K.YGGSVGSQPPPVAPEPGPVPSSPSQEPPTK	1	SAKS1_HUMAN
Q04323	2	3.293	R.LPDGTSLTQTFR.A	2	SAKS1_HUMAN
Q04323	1	3.7992	R.MLELVAQK.Q	5	SAKS1_HUMAN
SAPS domain family member 3 - Homo sapiens (Human)					
Q5H9R7	4	5.6916	R.IQQFDDGGSDEEDIWEEKHIAFTPESQR.R	1	SAPS3_HUMAN
Q5H9R7	2	3.4258	K.DAEECPETAEAK.C	2	SAPS3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5H9R7	2	3.9097	K.ETGWASFSEFTSSLSTK.D	1	SAPS3_HUMAN
SAPS1 protein - Homo sapiens (Human)					
Q2M2H3	2	3.0882	R.SQDPTPPSAPQEATEGSK.V	1	Q2M2H3_HUMA
Sarcomeric tropomyosin kappa - Homo sapiens (Human)					
Q6DV90	2	4.0273	K.C*AELEEELKTVTNDLK.S	1	Q6DV90_HUMA
Sarcoplasmic reticulum histidine-rich calcium-binding protein precursor - Homo sapiens (Human)					
P23327	3	4.2824	K.VGDEGVSGEEVFAEHGGQAR.G	1	SRCH_HUMAN
P23327	4	5.9609	R.SQDHKVGDEGVSGEEVFAEHGGQAR.G	1	SRCH_HUMAN
Scaffold attachment factor B2 - Homo sapiens (Human)					
Q14151	3	5.63	K.AVKEEGQDPDEIGIELEATSKK.S	2	SAFB2_HUMAN
Q14151	2	4.0149	R.NLWVSGLSSTTR.A	4	SAFB2_HUMAN
Q14151	2	3.5725	R.AWQGAMDAGAASR.E	2	SAFB2_HUMAN
Q14151	2	3.214	K.NTLETSSLNFK.V	1	SAFB2_HUMAN
Q14151	3	5.7331	K.AVKEEGQDPDEIGIELEATSK.K	2	SAFB2_HUMAN
Q14151	1	2.1553	K.EYVAAQLR.Q	1	SAFB2_HUMAN
Scavenger receptor class F member 2 precursor - Homo sapiens (Human)					
Q96GP6	2	3.2161	K.SAHTVEHGSPR.T	2	SREC2_HUMAN
Q96GP6	3	3.9247	R.GRGPGLLEPTDAGGPPR.S	2	SREC2_HUMAN
Q96GP6	2	3.5161	R.SAPEAASMLAAELR.G	2	SREC2_HUMAN
Scavenger receptor cysteine-rich type 1 protein M130 precursor - Homo sapiens (Human)					
Q86VB7	2	3.9655	K.EAEFGQGTGPIWLNEVK.C	2	C163A_HUMAN
SEC16 homolog A - Homo sapiens (Human)					
O15027	3	4.5608	K.NEHRPASALVNPLAR.G	2	SC16A_HUMAN
O15027	2	3.1516	R.YGPLPGPAVPR.H	3	SC16A_HUMAN
O15027	2	4.0158	R.NPSSAAPVQSR.G	3	SC16A_HUMAN
O15027	3	6.4898	R.LSGSARPQELVGTFIQQEVGKPEDEASGSFF	2	SC16A_HUMAN
O15027	2	3.4744	R.IVNHWASPELR.Q	1	SC16A_HUMAN
O15027	3	5.8407	R.GSVSQPSTPSPPKPTGIFQTSANSSFEPVK.S	2	SC16A_HUMAN
O15027	2	4.2222	K.QIDSSPVGGETDETTVSQNYR.G	3	SC16A_HUMAN
O15027	2	3.562	K.DAQGQPGLER.A	5	SC16A_HUMAN
O15027	2	3.4614	R.GGIGASENLENPPK.M	2	SC16A_HUMAN
SEC6-like protein C14orf73 - Homo sapiens (Human)					
Q17RC7	2	3.7228	K.EAEEPQTPAQGSR.R	2	CN073_HUMAN
Secretory carrier membrane protein 3 - Homo sapiens (Human)					
Q4VX17	2	3.381	R.QYATLDVYNPFETR.E	1	Q4VX17_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Segment polarity protein dishevelled homolog DVL-2 - Homo sapiens (Human)					
O14641	2	4.6548	R.GGEASGTSDGGPPPSR.G	2	DVL2_HUMAN
O14641	3	4.9464	R.VVSWLVSSDNPQPEMAPPVHEPR.A	2	DVL2_HUMAN
Selenium-binding protein 1 - Homo sapiens (Human)					
Q13228	4	5.9362	K.GGLKLNPNFLVDFGKEPLGPALAHELR.Y	1	SBP1_HUMAN
Q13228	3	3.7114	K.SPQYCQVIHR.L	1	SBP1_HUMAN
Q13228	2	4.1082	R.NTGTEAPDYLATVDVDPK.S	3	SBP1_HUMAN
Selenoprotein H - Homo sapiens (Human)					
Q8IZQ5	2	2.8012	K.AEAAVVAVAEK.R	1	SELH_HUMAN
Q8IZQ5	3	4.6444	K.FPEPQEVVEELKK.Y	1	SELH_HUMAN
Q8IZQ5	3	4.3506	R.KAEAAVVAVAEKR.E	2	SELH_HUMAN
Q8IZQ5	2	4.259	R.NAAALSQALR.L	4	SELH_HUMAN
Q8IZQ5	2	3.243	R.NAAALSQALRLEAPELPVK.V	2	SELH_HUMAN
Selenoprotein P precursor - Homo sapiens (Human)					
P49908	2	3.6825	R.DM#PASEDLQDLQK.K	7	SEPP1_HUMAN
Semaphorin-4D precursor - Homo sapiens (Human)					
Q92854	1	2.1286	K.FADSDADGD.-	1	SEM4D_HUMAN
Semaphorin-6B precursor - Homo sapiens (Human)					
Q9H3T3	2	3.0676	R.APEQPPAPGEPTPDGR.L	2	SEM6B_HUMAN
Sentrin-specific protease 6 - Homo sapiens (Human)					
Q9GZR1	2	4.8118	K.SGGSAGEITFLEALAR.S	2	SENP6_HUMAN
Septin-11 - Homo sapiens (Human)					
Q9NVA2	2	4.6059	K.AAAQLLQSQAQQSGAQQTK.K	1	SEP11_HUMAN
Septin-2 - Homo sapiens (Human)					
Q15019	2	3.3268	R.RMQEMIAR.M	2	SEPT2_HUMAN
Septin-5 - Homo sapiens (Human)					
Q99719	2	2.7817	K.LATPEDKQDIDK.Q	1	SEPT5_HUMAN
Septin-7 - Homo sapiens (Human)					
Q16181	2	4.2087	K.LKDSEAELQR.R	4	SEPT7_HUMAN
Q16181	2	2.7519	K.DSEAELQR.R	1	SEPT7_HUMAN
Septin-9 - Homo sapiens (Human)					
Q9UHD8	2	2.7013	R.SFEVEEVETPNSTPPRR.V	1	SEPT9_HUMAN
Q9UHD8	2	2.9851	R.SFEVEEVETPNSTPPR.R	1	SEPT9_HUMAN
Q9UHD8	2	3.1054	R.LGDSSGPALKR.S	2	SEPT9_HUMAN
Q9UHD8	3	3.7133	R.RLGDSSGPALKR.S	1	SEPT9_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9UHD8	2	2.8537	R.RLGDSSGPALK.R	1	SEPT9_HUMAN
Sequestosome-1 - Homo sapiens (Human)					
Q13501	2	3.2365	R.LTPVSPESSSTEEK.S	1	SQSTM_HUMAN
Serine arginine-rich pre-mRNA splicing factor SR-A1 - Homo sapiens (Human)					
Q7Z5V7	2	2.7559	K.DAASAGLGSIGVK.F	1	Q7Z5V7_HUMAN
Q7Z5V7	3	4.8939	R.AARPPPAASATPTAQPLPQPPAPR.A	4	Q7Z5V7_HUMAN
Q7Z5V7	3	4.3197	R.FDIYDPFHPTDEAYSPPPAPEQK.Y	1	Q7Z5V7_HUMAN
Serine hydroxymethyltransferase, cytosolic - Homo sapiens (Human)					
P34896	3	4.9006	K.MLAQPLKDSDVEVYNIIKK.E	1	GLYC_HUMAN
Serine/arginine repetitive matrix protein 2 - Homo sapiens (Human)					
Q9UQ35	2	2.9012	R.TSVPENHAQSR.I	1	SRRM2_HUMAN
Q9UQ35	2	3.5131	R.SSTGPEPPAPTPLLAER.H	2	SRRM2_HUMAN
Q9UQ35	2	3.1422	R.TAAALAPASLTSAR.M	2	SRRM2_HUMAN
Q9UQ35	2	3.5521	R.TPAAAAAM#NLASPR.T	2	SRRM2_HUMAN
Q9UQ35	2	3.9728	R.TPAAAAAMNLASPR.T	3	SRRM2_HUMAN
Q9UQ35	3	4.465	R.TPAALAALSLTGSGTPPTAANYPSSSR.T	1	SRRM2_HUMAN
Q9UQ35	2	3.8946	R.TPAIPTAVNLADSR.T	4	SRRM2_HUMAN
Q9UQ35	2	2.8156	R.SAHATAPVNIAGSR.T	1	SRRM2_HUMAN
Q9UQ35	2	3.2952	R.TPTAPAVNLAGAR.T	3	SRRM2_HUMAN
Q9UQ35	2	3.2515	R.MAPALSGANLTSPR.V	4	SRRM2_HUMAN
Q9UQ35	2	3.6629	R.TPQAPASANLVGPR.S	6	SRRM2_HUMAN
Q9UQ35	2	3.995	K.SSTPPGESYFGVSSLQLK.G	3	SRRM2_HUMAN
Q9UQ35	2	2.8929	R.SPVPSAFSDQSR.C	2	SRRM2_HUMAN
Q9UQ35	2	2.727	K.AGMSSNQSISSPVLDAVPR.T	1	SRRM2_HUMAN
Q9UQ35	2	2.711	R.AQSGSDSSPEPK.A	1	SRRM2_HUMAN
Q9UQ35	2	4.9153	R.FQSDSSSYPTVDSNSLLGQSR.L	3	SRRM2_HUMAN
Q9UQ35	2	3.1115	R.GEFSASPMLK.S	2	SRRM2_HUMAN
Q9UQ35	2	2.8041	R.HSLSGSSPGMK.D	1	SRRM2_HUMAN
Q9UQ35	2	5.2694	R.IPAASAAAM#NLASAR.T	4	SRRM2_HUMAN
Q9UQ35	2	5.07	R.IPAASAAAMNLASAR.T	2	SRRM2_HUMAN
Q9UQ35	2	2.7798	R.LETAESKEK.M	2	SRRM2_HUMAN
Q9UQ35	2	2.864	K.AGM#SSNQSISSPVLDAVPR.T	1	SRRM2_HUMAN
Serine/threonine protein kinase kkialre-like 1 - Homo sapiens (Human)					
Q8WXQ5	2	2.8038	K.IPDPEDMSLCLSVTLTEGGLLASGAVK.R	1	Q8WXQ5_HUMA
Serine/threonine-protein kinase 3 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q13188	2	3.0904	K.NLSLEELQM#R.L	1	STK3_HUMAN
Q13188	2	3.2185	K.NLSLEELQMR.L	2	STK3_HUMAN
Q13188	3	4.0922	K.VPQDGDFDFLKNLSLEELQMR.L	1	STK3_HUMAN
Serine/threonine-protein kinase 32C - Homo sapiens (Human)					
Q86UX6	2	2.8399	R.DAAEPVEDEAER.S	1	ST32C_HUMAN
Serine/threonine-protein kinase 36 - Homo sapiens (Human)					
Q9NRP7	1	2.1125	K.EAALIALR.S	1	STK36_HUMAN
Serine/threonine-protein kinase 4 - Homo sapiens (Human)					
Q13043	2	3.0629	K.RQPILDAIEAK.K	2	STK4_HUMAN
Serine/threonine-protein kinase DCLK2 - Homo sapiens (Human)					
Q8N568	2	3.2586	R.GAPSSSGGSSSSGPK.G	3	DCLK2_HUMAN
Serine/threonine-protein kinase Duet - Homo sapiens (Human)					
Q9Y2A5	2	2.914	K.ATAAESSDGSIK.K	1	DUET_HUMAN
Serine/threonine-protein kinase LMTK2 precursor - Homo sapiens (Human)					
Q8IWU2	3	5.0521	K.IFDLM#ELNGVQADFKPATLSSSLDNPK.E	1	LMTK2_HUMAN
Q8IWU2	2	4.6359	K.NLEAVETLNQLNSK.D	2	LMTK2_HUMAN
Q8IWU2	3	3.7245	R.SLLKPTAANAPDPLPEDWKK.E	1	LMTK2_HUMAN
Q8IWU2	2	3.4455	R.SQDSPGESEETLR.L	2	LMTK2_HUMAN
Q8IWU2	3	4.8749	R.SQDSPGESEETLRLTESDSVLADDILASR.V	1	LMTK2_HUMAN
Q8IWU2	2	3.5339	R.TGPELSQLTALR.S	3	LMTK2_HUMAN
Serine/threonine-protein kinase MARK2 - Homo sapiens (Human)					
Q7KZI7	3	3.9041	R.KKTTPTPSTNSVLSTSTNR.S	1	MARK2_HUMAN
Q7KZI7	2	6.1953	R.VPVASPSAHNISSSGGAPDR.T	6	MARK2_HUMAN
Q7KZI7	2	3.0302	K.VPASPLPGLER.K	2	MARK2_HUMAN
Serine/threonine-protein kinase MRCK beta - Homo sapiens (Human)					
Q9Y5S2	2	3.713	R.SM#SDPDQDFDKEPDSDSTK.H	2	MRCKB_HUMAN
Q9Y5S2	2	2.9334	R.SMSDPDQDFDKEPDSDSTK.H	1	MRCKB_HUMAN
Q9Y5S2	2	5.8241	R.NKPYISWPSSGGSEPSVTVPLR.S	7	MRCKB_HUMAN
Serine/threonine-protein kinase Nek9 - Homo sapiens (Human)					
Q8TD19	3	4.1966	K.ELENAEFIPMPDSPSPLSAAFSESEKDTLPYE	1	NEK9_HUMAN
Q8TD19	3	3.7786	K.KLEGGQQVGMHSK.G	1	NEK9_HUMAN
Serine/threonine-protein kinase OSR1 - Homo sapiens (Human)					
O95747	2	2.9004	K.IPISLVLR.L	2	OXSR1_HUMAN
O95747	2	5.0645	K.TAQALSSGSGSQETK.I	6	OXSR1_HUMAN
O95747	2	5.6241	K.TAQALSSGSGSQETKIPISLVLR.L	6	OXSR1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Serine/threonine-protein kinase PAK 1 - Homo sapiens (Human)					
Q13153	3	4.3114	K.DAGTLNHGSKPLPPNPEEK.K	4	PAK1_HUMAN
Q13153	2	2.9494	R.NTSTMIGAGSK.D	4	PAK1_HUMAN
Q13153	2	3.2792	R.NTSTM#IGAGSK.D	2	PAK1_HUMAN
Serine/threonine-protein kinase PAK 4 - Homo sapiens (Human)					
O96013	3	3.9989	K.RVEISAPSNFEHR.V	1	PAK4_HUMAN
O96013	2	4.1039	R.ARQENGMPEEPATTAR.G	1	PAK4_HUMAN
O96013	3	4.6808	R.DKRPLSGPDVGTPQPAGLASGAK.L	3	PAK4_HUMAN
O96013	2	4.98	R.FAGHSEAGGGSGDR.R	3	PAK4_HUMAN
O96013	2	2.9488	R.QWQSLIEESAR.R	2	PAK4_HUMAN
Serine/threonine-protein kinase PRP4 homolog - Homo sapiens (Human)					
Q13523	3	4.2441	R.TKLDDLALLEDLEKQR.A	2	PRP4B_HUMAN
Serine/threonine-protein kinase RIO2 - Homo sapiens (Human)					
Q9BVS4	3	5.2706	R.SSGDPEQIKEDSLSEESADAR.S	1	RIOK2_HUMAN
Serine/threonine-protein kinase TAO1 - Homo sapiens (Human)					
Q7L7X3	2	2.8046	R.ASDPQSPPQVSR.H	1	TAOK1_HUMAN
Serine/threonine-protein kinase TAO3 - Homo sapiens (Human)					
Q9H2K8	3	3.8828	R.MGFGNLVTLDFPKEDYR.-	1	TAOK3_HUMAN
Serine/threonine-protein kinase tousled-like 2 - Homo sapiens (Human)					
Q86UE8	2	3.1738	R.KAEPYETSQGK.G	1	TLK2_HUMAN
Serine/threonine-protein kinase WNK1 - Homo sapiens (Human)					
Q9H4A3	3	4.2844	K.EGPVASPPFM#DLEQAVLPAVIPK.K	1	WNK1_HUMAN
Q9H4A3	2	3.7245	K.EGPVASPPFMDLEQAVLPAVIPK.K	2	WNK1_HUMAN
Q9H4A3	2	4.8461	K.LGAAAADAVTGR.T	2	WNK1_HUMAN
Serine/threonine-protein phosphatase 1 regulatory subunit 10 - Homo sapiens (Human)					
Q96QC0	2	3.4907	K.GILQELFLNK.E	1	PP1RA_HUMAN
Q96QC0	2	4.2854	R.STGLELETPSLVPVKK.N	2	PP1RA_HUMAN
Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit delta isoform - Homo sapiens (					
Q14738	2	3.3434	K.RAEEFLTASQEAL.-	2	2A5D_HUMAN
Serine/threonine-protein phosphatase 2B catalytic subunit alpha isoform - Homo sapiens (Human)					
Q08209	2	3.7233	R.DAM#PSDANLNSINK.A	5	PP2BA_HUMAN
Serine/threonine-protein phosphatase 5 - Homo sapiens (Human)					
P53041	2	3.7022	R.AALRDYETVVK.V	3	PPP5_HUMAN
P53041	1	2.7218	K.TQANDYFK.A	3	PPP5_HUMAN
P53041	3	5.3824	K.FYSQAIELNPSNAIYYGNR.S	2	PPP5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Serologically defined colon cancer antigen 1 - Homo sapiens (Human)					
O60524	2	3.3203	K.LLGSAGSNKEEK.G	2	SDCG1_HUMAN
O60524	2	2.7641	K.NVAAVQPM#KR.G	1	SDCG1_HUMAN
O60524	2	3.1851	K.TALNSFMHSK.E	3	SDCG1_HUMAN
O60524	2	3.3659	K.VSAPNLLNVK.R	3	SDCG1_HUMAN
Serologically defined colon cancer antigen 8 - Homo sapiens (Human)					
Q86SQ7	2	4.6012	K.SPENSTLEEILGQYQR.S	2	SDCG8_HUMAN
Serpin H1 precursor - Homo sapiens (Human)					
P50454	3	5.0429	K.HLAGLGLTEAIDKNK.A	2	SERPH_HUMAN
P50454	2	3.4835	K.GVVEVTHDLQK.H	2	SERPH_HUMAN
SERPINC1 protein - Homo sapiens (Human)					
Q8TCE1	3	4.3561	K.KATEDEGSEQKIPEATNR.R	1	Q8TCE1_HUMA
Q8TCE1	2	4.2096	K.ATEDEGSEQKIPEATNR.R	11	Q8TCE1_HUMA
Q8TCE1	1	2.3169	K.IPEATNR.R	1	Q8TCE1_HUMA
Serum amyloid A-4 protein precursor - Homo sapiens (Human)					
P35542	2	3.0795	R.GPGGVWAAK.L	2	SAA4_HUMAN
Serum deprivation-response protein - Homo sapiens (Human)					
O95810	3	4.7211	K.VRYEGSYALTSEEAERSDGDPVQPAVLQVH	1	SDPR_HUMAN
O95810	1	2.2567	K.VSPLTFGR.K	1	SDPR_HUMAN
O95810	3	6.8573	R.YEGSYALTSEEAERSDGDPVQPAVLQVHQT	7	SDPR_HUMAN
O95810	2	4.5129	R.YEGSYALTSEEAER.S	2	SDPR_HUMAN
O95810	1	2.883	K.YQASTSNTVSK.L	2	SDPR_HUMAN
O95810	3	3.7597	K.VREGESHAENETK.S	2	SDPR_HUMAN
O95810	1	2.2389	K.GIQNDLTK.L	1	SDPR_HUMAN
O95810	2	4.017	K.VLIFQEENEIPASVFVK.Q	6	SDPR_HUMAN
O95810	1	2.2762	R.QISLEGSVK.G	2	SDPR_HUMAN
O95810	2	3.913	K.RLENNHAQLLR.R	4	SDPR_HUMAN
O95810	2	4.9938	K.LVNMLDAVQENQHK.M	2	SDPR_HUMAN
O95810	2	4.8529	K.VRYEGSYALTSEEAER.S	2	SDPR_HUMAN
Serum response factor binding protein 1 - Homo sapiens (Human)					
Q8NEF9	3	4.3436	K.AVTIANSPSKPSEKDSVVSLESQK.T	1	Q8NEF9_HUMA
Q8NEF9	2	2.9192	R.SLDFPQNEPQIK.N	1	Q8NEF9_HUMA
SET domain-containing protein 3 - Homo sapiens (Human)					
Q86TU7	2	3.4846	R.SENESLNQESKR.A	2	SETD3_HUMAN
Q86TU7	2	3.4611	K.GSSSDSTAGVKE.-	3	SETD3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q86TU7	2	4.6724	R.AVEDAKGSSSDSTAGVKE.-	1	SETD3_HUMAN
Seven transmembrane helix receptor - Homo sapiens (Human)					
Q8IXE5	2	3.0854	R.GANQERAEKVK.L	2	Q8IXE5_HUMAN
SF3A2 protein - Homo sapiens (Human)					
Q05DF2	2	3.7679	K.TGSGGVASSSESNR.D	2	Q05DF2_HUMA
SFRS2-interacting protein - Homo sapiens (Human)					
Q99590	2	3.4021	R.KPLSGNSNSSGSESFK.F	1	SFRIP_HUMAN
SH2 domain-containing adapter protein B - Homo sapiens (Human)					
Q15464	3	4.9288	K.DKVTIADDYSDPFDAKNDLK.S	1	SHB_HUMAN
SH2 domain-containing adapter protein E - Homo sapiens (Human)					
Q5VZ18	2	3.6497	R.DSLQGLIQAAAGK.G	2	SHE_HUMAN
SH2B adapter protein 3 - Homo sapiens (Human)					
Q9UQQ2	2	2.8295	R.DSDYEM#DSSSR.S	1	SH2B3_HUMAN
SH3 and multiple ankyrin repeat domains protein 2 - Homo sapiens (Human)					
Q9UPX8	2	3.9285	K.ANVISELNSILQQMNR.E	3	SHAN2_HUMAN
Q9UPX8	3	4.3759	K.AQGPESSPAVPSASSGTAGPGNYVHPLTGR.	1	SHAN2_HUMAN
SH3 and multiple ankyrin repeat domains protein 3 - Homo sapiens (Human)					
Q9BYB0	2	4.0169	R.LFSSLGELSSISAQR.S	3	SHAN3_HUMAN
Q9BYB0	2	4.7342	R.SASDINLKGEAQPAASPGPSLR.S	1	SHAN3_HUMAN
Q9BYB0	3	4.177	R.SRSPSPSPLPSPASGPGPGAPGPR.R	1	SHAN3_HUMAN
Q9BYB0	3	3.9137	R.LGAGAAGLYEPGAALGPLPYPER.Q	1	SHAN3_HUMAN
Q9BYB0	2	3.7302	R.GSLASPAFSPR.S	4	SHAN3_HUMAN
Q9BYB0	3	6.8728	R.GPRPGGLDYGAGDGPGLAFGGPGPAK.D	2	SHAN3_HUMAN
Q9BYB0	2	2.8219	K.QLQVEDAQER.A	1	SHAN3_HUMAN
Q9BYB0	2	4.4621	R.AALAVGSPGPGGGSFAR.E	3	SHAN3_HUMAN
Q9BYB0	2	4.1057	R.ITPAEISSLFER.Q	2	SHAN3_HUMAN
SH3 and PX domain containing 3 - Homo sapiens (Human)					
Q8WV41	2	3.9558	R.GETGLFPASYVEIVR.S	4	Q8WV41_HUMA
Q8WV41	3	6.743	R.SGISTNHADYSSSPAGSPGAQVSLYNSPSVA	4	Q8WV41_HUMA
SH3 and PX domain-containing protein 2A - Homo sapiens (Human)					
Q5TCZ1	2	2.8726	K.EAEEGPTGASESQDSPR.K	1	SPD2A_HUMAN
Q5TCZ1	2	3.082	R.AASQGSDSPLLPAQR.N	1	SPD2A_HUMAN
SH3 domain GRB2-like protein B1 - Homo sapiens (Human)					
Q9Y371	2	4.2013	K.GKVPITYLELLN.-	8	SHLB1_HUMAN
SH3 domain-binding glutamic acid-rich-like protein - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O75368	3	5.3474	K.KQQDVLGFLEANK.I	4	SH3L1_HUMAN
O75368	1	4.764	R.VYIASSSGSTAIK.K	11	SH3L1_HUMAN
O75368	2	3.0391	R.GDYDAFFEAR.E	4	SH3L1_HUMAN
O75368	3	5.6174	R.ENVPENSRPATGYPLPPQIFNESQYR.G	5	SH3L1_HUMAN
O75368	2	3.7519	R.ENNAVYAFLGLTAPPGSK.E	5	SH3L1_HUMAN
O75368	2	3.8222	R.VYIASSSGSTAIKK.K	4	SH3L1_HUMAN
O75368	3	5.0853	K.KQQDVLGFLEANKIGFEEK.D	1	SH3L1_HUMAN
O75368	3	4.7804	K.IGFEEKDIAANEENRK.W	8	SH3L1_HUMAN
O75368	3	5.0746	K.IGFEEKDIAANEENR.K	3	SH3L1_HUMAN
O75368	2	2.8739	K.EAEVQAKQQA.-	1	SH3L1_HUMAN
O75368	2	3.0498	K.DIAANEENRK.W	5	SH3L1_HUMAN
O75368	2	3.4418	K.DIAANEENR.K	3	SH3L1_HUMAN
O75368	2	3.4356	K.QQDVLGFLEANK.I	6	SH3L1_HUMAN
O75368	3	5.8226	K.KKQQDVLGFLEANK.I	6	SH3L1_HUMAN
SH3 domain-binding protein 1 - Homo sapiens (Human)					
Q9Y3L3	2	2.7971	R.RLVGSSLR.A	1	3BP1_HUMAN
SH3 domain-containing RING finger protein 1 - Homo sapiens (Human)					
Q7Z6J0	2	3.2814	K.DLQSSQGGQQPR.V	4	Q7Z6J0_HUMAN
SH3-containing GRB2-like protein 1 - Homo sapiens (Human)					
Q99961	3	4.335	K.VGGAEGTKLDDDFKEMEK.K	2	SH3G1_HUMAN
Shootin-1 - Homo sapiens (Human)					
A0MZ66	3	4.6804	R.RKVTAEADSSSPTGILATSESK.S	1	SHOT1_HUMAN
A0MZ66	2	4.0344	K.SLDPENSETELER.I	1	SHOT1_HUMAN
A0MZ66	2	3.8183	K.SM#PVLGSVSSVTK.T	3	SHOT1_HUMAN
A0MZ66	2	4.5619	K.SMPVLGSVSSVTK.T	2	SHOT1_HUMAN
A0MZ66	2	4.2465	K.TLEAEFNSPSPPTPEPGEGPR.K	3	SHOT1_HUMAN
A0MZ66	2	5.2529	K.VTAEADSSSPTGILATSESK.S	3	SHOT1_HUMAN
A0MZ66	2	6.3417	R.KVTAEADSSSPTGILATSESK.S	3	SHOT1_HUMAN
Sialate O-acetylesterase precursor - Homo sapiens (Human)					
Q9HAT2	2	5.1969	R.FFPFGLVQLSSDLSK.K	2	SIAE_HUMAN
Sialic acid-binding Ig-like lectin 9 precursor - Homo sapiens (Human)					
Q9Y336	2	3.6519	R.EGANTDQDAPVATNNPAR.A	1	SIGL9_HUMAN
Sickle tail protein homolog - Homo sapiens (Human)					
Q5T5P2	2	3.4237	K.QLENTISEMSPK.A	3	SKT_HUMAN
Q5T5P2	2	3.7721	R.TLDSLEQTIK.Q	2	SKT_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5T5P2	2	5.4868	R.RQEQPSIESTSPISR.T	3	SKT_HUMAN
Q5T5P2	3	3.8917	R.PLLVPDEGPTALEPPTSIPSASR.K	1	SKT_HUMAN
Q5T5P2	2	3.9726	R.NPGGQEMNRTELNK.F	2	SKT_HUMAN
Q5T5P2	2	3.1341	R.NPGGQEMNR.T	2	SKT_HUMAN
Q5T5P2	2	2.7511	K.RFEIAR.S	1	SKT_HUMAN
Q5T5P2	3	4.5695	R.TLDSLEQTIKQLENTISEM#SPK.A	1	SKT_HUMAN
Q5T5P2	2	2.8522	K.QLAALTQAIR.T	1	SKT_HUMAN
Q5T5P2	2	3.2759	K.KQLAALTQAIR.T	1	SKT_HUMAN
Q5T5P2	3	3.8282	K.KPVIIIFDEPMDIR.S	1	SKT_HUMAN
Q5T5P2	2	2.7583	K.ENSISNMSLLR.D	1	SKT_HUMAN
Q5T5P2	2	3.6667	K.ASFGFSGISPLEDEINK.G	2	SKT_HUMAN
Q5T5P2	3	4.2799	R.KSDVEYENGPQMEFQK.V	1	SKT_HUMAN
Sideroflexin-1 - Homo sapiens (Human)					
Q9H9B4	2	3.2483	K.SSMSVTSLEAELQAK.I	1	SFXN1_HUMAN
Q9H9B4	2	3.1347	K.SSM#SVTSLEAELQAK.I	1	SFXN1_HUMAN
Signal recognition particle 14 kDa protein - Homo sapiens (Human)					
P37108	2	3.1251	K.EVNKFQMAYSNLLR.A	2	SRP14_HUMAN
P37108	2	2.8976	K.FQM#AYSNLLR.A	2	SRP14_HUMAN
P37108	2	3.4166	K.FQMAYSNLLR.A	2	SRP14_HUMAN
P37108	2	4.1402	K.KGTVEGFEPADNK.C	3	SRP14_HUMAN
Signal recognition particle 19 kDa protein - Homo sapiens (Human)					
P09132	2	4.3765	K.TGGADQSLQQGEGSKK.G	2	SRP19_HUMAN
P09132	2	3.6318	K.TGGADQSLQQGEGSK.K	1	SRP19_HUMAN
Signal recognition particle 54 kDa protein - Homo sapiens (Human)					
P61011	2	2.8853	R.QFQQGAAGNM#K.G	2	SRP54_HUMAN
Signal recognition particle 68 kDa protein - Homo sapiens (Human)					
Q9UHB9	3	4.2403	K.QVPGGGGGGGSGGGGGSGGGGSGGGR.G	2	SRP68_HUMAN
Signal recognition particle 72 kDa protein - Homo sapiens (Human)					
O76094	2	3.8583	R.HQKPAGAPATK.K	4	SRP72_HUMAN
O76094	2	3.0723	R.HQKPAGAPATKK.K	1	SRP72_HUMAN
Signal recognition particle receptor subunit alpha - Homo sapiens (Human)					
P08240	3	4.2011	K.KEGSDGPLATSKPVPAEK.S	1	SRPR_HUMAN
P08240	1	2.1108	K.SDAPKEKGK.K	1	SRPR_HUMAN
P08240	2	4.1583	K.SGLPVGPENGVELSKEELIR.R	1	SRPR_HUMAN
Signal recognition particle receptor subunit beta - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y5M8	2	3.5663	R.GDVGSADIQDLEK.W	1	SRPRB_HUMAN
Signal transducer and activator of transcription 1-alpha/beta - Homo sapiens (Human)					
P42224	2	3.6303	K.ELSAVTFPDIIR.N	2	STAT1_HUMAN
P42224	2	3.5623	K.VMAAENIPENPLK.Y	1	STAT1_HUMAN
Signal transducer and activator of transcription 2 - Homo sapiens (Human)					
P52630	2	3.036	R.HLNTEPMEIFR.N	1	STAT2_HUMAN
P52630	2	3.4178	R.PSHFYTDGPLMPSDF.-	1	STAT2_HUMAN
Signal transducing adapter molecule 2 - Homo sapiens (Human)					
O75886	2	2.8064	K.VLEALELYNK.L	1	STAM2_HUMAN
Signal-induced proliferation-associated 1-like protein 1 - Homo sapiens (Human)					
O43166	2	3.7801	R.LQEESQNASDK.L	3	SI1L1_HUMAN
O43166	2	4.507	K.LIDLESPTPESQK.S	4	SI1L1_HUMAN
O43166	2	3.367	K.QGTSGESFFDLLK.G	2	SI1L1_HUMAN
O43166	2	2.9868	R.FLMPEAYPSSPR.K	1	SI1L1_HUMAN
Signal-induced proliferation-associated 1-like protein 3 - Homo sapiens (Human)					
O60292	2	3.2514	K.VVLQSEVASLR.Q	2	SI1L3_HUMAN
Signal-induced proliferation-associated protein 1 - Homo sapiens (Human)					
Q96FS4	2	3.0256	R.LQAESESAATR.L	1	SIPA1_HUMAN
Q96FS4	2	4.5057	R.RLQAESESAATR.L	3	SIPA1_HUMAN
Q96FS4	2	3.7799	K.SDAEPEPGNLSEK.V	2	SIPA1_HUMAN
Signaling threshold-regulating transmembrane adapter 1 precursor - Homo sapiens (Human)					
Q9Y3P8	3	3.9404	R.LSQDPEPDQQDPTLGGPAR.A	3	SIT1_HUMAN
Single Ig IL-1-related receptor - Homo sapiens (Human)					
Q6IA17	2	2.9111	R.ALDSEVDPDPEGDLGVR.G	1	SIGIR_HUMAN
Single-stranded DNA-binding protein, mitochondrial precursor - Homo sapiens (Human)					
Q04837	2	4.365	K.NPVTIFSLATNEMWR.S	1	SSB_HUMAN
Sister chromatid cohesion protein PDS5 homolog A - Homo sapiens (Human)					
Q29RF7	3	4.3719	K.TDEKVDESGPPAPSKPR.R	1	PDS5A_HUMAN
Q29RF7	2	3.9811	R.STGTETGSNINVNSELNPSTGNR.S	1	PDS5A_HUMAN
Q29RF7	2	3.4359	K.NDDLNKPINK.G	2	PDS5A_HUMAN
Q29RF7	3	4.0493	R.SREQSSEAAETGVSENEENPVR.I	2	PDS5A_HUMAN
Q29RF7	2	3.4046	R.EQSSEAAETGVSENEENPVR.I	1	PDS5A_HUMAN
Sjoegren syndrome/scleroderma autoantigen 1 - Homo sapiens (Human)					
O60232	2	3.2808	R.LMGDYLLR.G	2	SSA27_HUMAN
Ski oncogene - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P12755	2	3.8512	R.ARPEAAGSEGAAELEP.-	1	SKI_HUMAN
SLAIN motif family member 2 - Homo sapiens (Human)					
Q9P270	2	5.6645	R.SGAVQGAGSLGPGSPVR.A	4	Q9P270_HUMAN
Q9P270	2	3.0773	R.AGASIPSSGAASPR.G	1	Q9P270_HUMAN
Q9P270	2	3.1521	R.KLQELVK.K	1	Q9P270_HUMAN
SLAM family member 5 precursor - Homo sapiens (Human)					
Q9UIB8	2	2.9	R.IYDEILQSK.V	1	SLAF5_HUMAN
SLIT and NTRK-like protein 3 precursor - Homo sapiens (Human)					
O94933	2	2.7856	K.QWIETISSVSVVGDVLC*R.S	1	SLIK3_HUMAN
SLIT-ROBO Rho GTPase-activating protein 1 - Homo sapiens (Human)					
Q7Z6B7	2	4.4974	K.HAPDVVLDTLEQVK.N	1	SRGP1_HUMAN
Q7Z6B7	2	2.7092	K.TNPTIGPAPPPQGPTDK.S	1	SRGP1_HUMAN
SLIT-ROBO Rho GTPase-activating protein 2 - Homo sapiens (Human)					
O75044	2	3.9011	K.TNATSPGVNSSTSPQSTDK.S	2	FNBP2_HUMAN
Smad nuclear-interacting protein 1 - Homo sapiens (Human)					
Q8TAD8	2	3.502	R.DGDVVLPAGVVVK.Q	4	SNIP1_HUMAN
Small acidic protein - Homo sapiens (Human)					
O00193	2	2.8148	R.LVIGDHK.S	1	SMAP_HUMAN
O00193	2	5.178	R.SASPDDDLGSSNWEAADLGNEER.K	10	SMAP_HUMAN
O00193	3	6.0527	R.TGEEDKKINEELESQYQQSM#DSK.L	9	SMAP_HUMAN
O00193	2	3.2172	K.INEELESQYQQSM#DSK.L	1	SMAP_HUMAN
O00193	2	4.6263	K.INEELESQYQQSMDSK.L	3	SMAP_HUMAN
O00193	1	2.1873	R.LMGAGK.K	1	SMAP_HUMAN
O00193	3	6.0967	R.TGEEDKKINEELESQYQQSMDSK.L	11	SMAP_HUMAN
Small EDRK-rich factor 2 - Homo sapiens (Human)					
P84101	2	3.6354	R.DDGLSAAAR.K	3	SERF2_HUMAN
P84101	2	3.599	R.DSEIM#QQK.Q	9	SERF2_HUMAN
P84101	2	3.3778	R.DSEIMQQK.Q	12	SERF2_HUMAN
Small glutamine-rich tetratricopeptide repeat-containing protein A - Homo sapiens (Human)					
O43765	2	3.6405	R.TPPSEEDSAEAER.L	1	SGTA_HUMAN
O43765	2	3.5426	R.SRTPSASNDDQQE.-	2	SGTA_HUMAN
Small nuclear ribonucleoprotein F - Homo sapiens (Human)					
P62306	2	3.0286	R.GVEEEEEDGEM#RE.-	2	RUXF_HUMAN
P62306	2	3.2982	R.GVEEEEEDGEMRE.-	5	RUXF_HUMAN
Small nuclear ribonucleoprotein polypeptide C - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5TAL2	3	4.1974	-.M#EEQAQSLIDKTTAAFQQGK.I	1	Q5TAL2_HUMAN
Small nuclear ribonucleoprotein Sm D2 - Homo sapiens (Human)					
P62316	1	2.3005	R.NPLIAGK.-	2	SMD2_HUMAN
Small nuclear ribonucleoprotein Sm D3 - Homo sapiens (Human)					
P62318	2	3.199	R.VAQLEQVYIR.G	2	SMD3_HUMAN
Small ubiquitin-related modifier 1 precursor - Homo sapiens (Human)					
P63165	2	3.039	R.IADNHTPK.E	3	SUMO1_HUMAN
Small ubiquitin-related modifier 2 precursor - Homo sapiens (Human)					
P61956	2	3.3412	K.EGVKTENNDHINLK.V	4	SUMO2_HUMAN
P61956	2	3.4024	K.TENNDHINLK.V	5	SUMO2_HUMAN
Small ubiquitin-related modifier 3 precursor - Homo sapiens (Human)					
P55854	2	3.5713	K.EGVKTENDHINLK.V	4	SUMO3_HUMAN
P55854	2	2.7369	K.TENDHINLK.V	1	SUMO3_HUMAN
SNARE-associated protein Snapin - Homo sapiens (Human)					
O95295	2	4.4392	R.AMLDSGIYPPGSPGK.-	1	S25BP_HUMAN
SNW domain-containing protein 1 - Homo sapiens (Human)					
Q13573	2	3.5815	K.THVEKEDGEAR.E	3	SNW1_HUMAN
Q13573	2	2.938	R.YTPSQQGVAFNSGAK.Q	1	SNW1_HUMAN
Solute carrier family 12 member 2 - Homo sapiens (Human)					
P55011	2	3.0467	K.QTPADGEASGESEPAK.G	2	S12A2_HUMAN
P55011	2	6.2418	R.AAAAAAAAAAAAAAAGAGAGAK.Q	4	S12A2_HUMAN
P55011	2	2.7307	R.VELPGTAVPSVPEDAAPASR.D	1	S12A2_HUMAN
Solute carrier family 2, facilitated glucose transporter member 1 - Homo sapiens (Human)					
P11166	3	3.7186	R.QGGASQSDKTPEELFHPLGADSQV.-	1	GTR1_HUMAN
Something about silencing protein 10 - Homo sapiens (Human)					
Q9NQZ2	2	2.943	R.YSGELSGIR.A	1	SAS10_HUMAN
Son of sevenless homolog 1 - Homo sapiens (Human)					
Q07889	2	4.2651	R.DGPPLLENAHSS.-	4	SOS1_HUMAN
Son of sevenless homolog 2 - Homo sapiens (Human)					
Q07890	2	2.9596	R.DISTCPNSPSTPPSTPSPR.V	1	SOS2_HUMAN
SON protein - Homo sapiens (Human)					
P18583	3	4.3328	K.RLTDLDKAQLLEIAK.A	1	SON_HUMAN
P18583	2	3.5389	K.SVESTSPEPSK.I	3	SON_HUMAN
Sorbin and SH3 domain-containing protein 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O94875	3	4.3406	K.SSILQHERPASLYQSSIDR.S	3	O94875_HUMAN
Sorting nexin-11 - Homo sapiens (Human)					
Q9Y5W9	3	4.698	R.AVGGDHAVPLDPGQLETVLEK.-	1	SNX11_HUMAN
Sorting nexin-2 - Homo sapiens (Human)					
O60749	2	4.3423	R.AVNTQALSGAGILR.M	2	SNX2_HUMAN
O60749	2	3.0632	K.SMSAPVIFDR.S	2	SNX2_HUMAN
Sorting nexin-3 - Homo sapiens (Human)					
O60493	2	2.9652	R.YSDFEWLR.S	1	SNX3_HUMAN
O60493	3	4.2365	R.LITKPQNLNDAYGPPSNFLEIDVSNPQTVGVG	1	SNX3_HUMAN
O60493	3	3.8615	R.KQGLEQFINK.V	7	SNX3_HUMAN
O60493	3	3.7191	K.VAGHPLAQNER.C	1	SNX3_HUMAN
Sorting nexin-5 - Homo sapiens (Human)					
Q9Y5X3	2	3.9893	R.NNVSLLQSCIDLFK.N	2	SNX5_HUMAN
Sorting nexin-9 - Homo sapiens (Human)					
Q9Y5X1	2	4.2299	K.SGNWESSEGWGAQPEGAGAQR.N	5	SNX9_HUMAN
SOX-12 protein - Homo sapiens (Human)					
O15370	2	3.6483	R.DGGPPPPGPGPAEEGAR.E	2	SOX12_HUMAN
SPARC precursor - Homo sapiens (Human)					
P09486	2	4.9179	K.NVLVTLYERDEDNNLLTEK.Q	2	SPRC_HUMAN
P09486	2	4.0778	R.LEAGDHPVELLAR.D	1	SPRC_HUMAN
SPARC-like protein 1 precursor - Homo sapiens (Human)					
Q14515	3	3.7948	K.TVSEALLM#EPTDDGNTTPR.N	2	SPRL1_HUMAN
Q14515	3	5.2104	K.VHENENIGTTEPGEHQEAK.K	3	SPRL1_HUMAN
Q14515	4	5.3811	R.DQGNQEQDPNISNGEEEEEKEPGEVGTHND	1	SPRL1_HUMAN
Q14515	2	3.8758	R.LLAGDHPIDLLLR.D	1	SPRL1_HUMAN
Spartin - Homo sapiens (Human)					
Q8N0X7	2	3.6961	R.IQPEEKPVEVSPAVTK.G	4	SPG20_HUMAN
Spastin - Homo sapiens (Human)					
Q9UBP0	2	2.8314	K.TGSAGLSGHHR.A	3	SPAST_HUMAN
Q9UBP0	2	4.0507	R.APSYSGLSMVSGVK.Q	2	SPAST_HUMAN
Spectrin alpha chain, brain - Homo sapiens (Human)					
Q13813	2	3.2528	R.ELPTAFDYVEFTR.S	2	SPTA2_HUMAN
Q13813	2	3.7956	R.DVEDEETWIR.E	3	SPTA2_HUMAN
Q13813	3	4.1329	R.DVEDEETWIREKEPIAASTNR.G	1	SPTA2_HUMAN
Q13813	2	5.2294	R.LIQSHPESAEDLQEK.C	3	SPTA2_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q13813	3	4.7177	R.EAFLNTEDKGDSLDSVEALIK.K	2	SPTA2_HUMAN
Q13813	3	5.7781	R.DLAALGDKVNSLGETAER.L	3	SPTA2_HUMAN
Q13813	2	3.9969	R.EANELQQWINEK.E	4	SPTA2_HUMAN
Q13813	2	2.9207	R.EANQQQQFNR.N	1	SPTA2_HUMAN
Q13813	2	3.9796	R.EELITNWEQIR.T	1	SPTA2_HUMAN
Q13813	2	2.7631	R.EKEPIAASTNR.G	1	SPTA2_HUMAN
Q13813	3	4.3841	R.FLADFRDLTSWVTEMK.A	2	SPTA2_HUMAN
Q13813	2	3.7387	R.FNRDVDETISWIK.E	2	SPTA2_HUMAN
Q13813	2	4.3182	R.GKDLIGVQNLLK.K	4	SPTA2_HUMAN
Q13813	3	4.1333	R.GLVSSDELAKDVTGAEALLER.H	1	SPTA2_HUMAN
Q13813	2	3.9786	R.GVIDMGNSLIER.G	3	SPTA2_HUMAN
Q13813	3	4.3234	R.MKQVEELYHSLLELGEK.R	2	SPTA2_HUMAN
Q13813	2	3.3208	R.SLQQLAEER.S	1	SPTA2_HUMAN
Q13813	2	4.1985	R.SQLLGSAHEVQR.F	2	SPTA2_HUMAN
Q13813	2	3.4433	R.DLAALEDKVK.A	2	SPTA2_HUMAN
Q13813	2	3.7063	R.YEALKEPMVAR.K	3	SPTA2_HUMAN
Q13813	3	3.92	K.KHQALQAEIAGHEPR.I	1	SPTA2_HUMAN
Q13813	2	4.3985	R.DAEELEKWIQEK.L	1	SPTA2_HUMAN
Q13813	3	3.7223	R.NTTGVTEEALKEFSMMFK.H	1	SPTA2_HUMAN
Q13813	2	5.4529	K.EAALTSEEVGADLEQVEVLQK.K	7	SPTA2_HUMAN
Q13813	3	3.7077	K.GNAMVEEGHFAAEDVK.A	1	SPTA2_HUMAN
Q13813	3	3.8793	K.IAALQAFADQLIAAGHYAK.G	1	SPTA2_HUMAN
Q13813	4	4.7375	K.KFEEFQTDMAAHEERVNEVNQFAAK.L	1	SPTA2_HUMAN
Q13813	3	4.7983	K.KHEALMSDLSAYGSSIQALR.E	3	SPTA2_HUMAN
Q13813	2	4.096	K.LFGAAEVQR.F	3	SPTA2_HUMAN
Q13813	2	4.2205	K.LGESQTLQQFSR.D	3	SPTA2_HUMAN
Q13813	2	3.2925	K.LHELNQKWEALK.A	1	SPTA2_HUMAN
Q13813	2	3.7511	K.SSEEIESAFR.A	3	SPTA2_HUMAN
Q13813	4	5.3289	R.DADETKEWIEEKNQALNTDNYGHDLASVQAL	1	SPTA2_HUMAN
Q13813	3	4.6564	K.KGDILTLLNSTNKDWWK.V	5	SPTA2_HUMAN
Q13813	2	3.9555	K.VLETAEDIQER.R	4	SPTA2_HUMAN
Q13813	2	3.9686	K.LIQEQHPEEELIK.T	2	SPTA2_HUMAN
Q13813	3	4.2074	K.SADESGQALLAAGHYASDEVREK.L	1	SPTA2_HUMAN
Q13813	4	5.0247	K.RLEAELAAHEPAIQGVLDTGKK.L	1	SPTA2_HUMAN
Q13813	2	4.3793	K.REELITNWEQIR.T	1	SPTA2_HUMAN
Q13813	3	4.3838	K.NQALNTDNYGHDLASVQALQR.K	2	SPTA2_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q13813	3	4.1638	K.LSDDNTIGKEEIQQR.L	4	SPTA2_HUMAN
Q13813	2	3.4343	K.VLETAEDIQERR.Q	2	SPTA2_HUMAN
Spectrin beta chain, brain 1 - Homo sapiens (Human)					
Q01082	2	3.3785	R.DQNTVETLQR.M	2	SPTB2_HUMAN
Q01082	2	3.8555	K.FM#ELLEPLNER.K	3	SPTB2_HUMAN
Q01082	2	3.817	R.TQETPSAQMEGFLNR.K	5	SPTB2_HUMAN
Q01082	2	4.1245	R.LTTLELLEVR.R	2	SPTB2_HUMAN
Q01082	3	4.8631	R.KKEIEELQSQAQALSQEGK.S	2	SPTB2_HUMAN
Q01082	3	4.7994	R.EVVAGSHELGQDYEHVTMLQER.F	1	SPTB2_HUMAN
Q01082	2	3.1626	R.EVDDLEQWIAER.E	1	SPTB2_HUMAN
Q01082	2	2.8328	R.ETASELLMRLKDNR.D	1	SPTB2_HUMAN
Q01082	2	4.7956	R.AQTLPTSVVTITSESSPGKR.E	4	SPTB2_HUMAN
Q01082	2	5.1109	R.AQTLPTSVVTITSESSPGK.R	3	SPTB2_HUMAN
Q01082	2	4.3484	K.TALPAQSAATLPAR.T	4	SPTB2_HUMAN
Q01082	2	3.4846	K.MWEVLESTTQTK.A	2	SPTB2_HUMAN
Q01082	2	3.7139	K.FMELLEPLNER.K	2	SPTB2_HUMAN
Q01082	3	6.1888	R.DASVAEAWLLGQEPYLSSR.E	1	SPTB2_HUMAN
Q01082	3	6.9194	K.HQILEQAVEDYAETVHQLSK.T	2	SPTB2_HUMAN
Spectrin beta chain, erythrocyte - Homo sapiens (Human)					
P11277	2	4.1607	R.LLSGEDVGQDEGATR.A	2	SPTB1_HUMAN
Sperm antigen with calponin homology and coiled-coil domains 1 - Homo sapiens (Human)					
Q5M775	2	3.8934	K.SVSSPTSSNTPTPTK.H	3	SPEC1_HUMAN
Q5M775	2	4.3437	R.ASSEDTLNKPGSTAASGVVR.L	4	SPEC1_HUMAN
Q5M775	3	4.1762	R.GVTQRLDLPDLPLSDILK.G	2	SPEC1_HUMAN
Q5M775	2	5.0494	R.KSVSSPTSSNTPTPTK.H	2	SPEC1_HUMAN
Q5M775	2	2.7923	R.LDLPDLPLSDILK.G	1	SPEC1_HUMAN
Q5M775	2	4.0111	R.VKNEEPTTQEGK.I	3	SPEC1_HUMAN
Q5M775	3	4.0526	R.AEQLSQENEKLMNLLQER.V	1	SPEC1_HUMAN
Q5M775	3	4.6925	R.LKEENEKLNEFLELER.H	1	SPEC1_HUMAN
Q5M775	2	3.7071	K.TATAGAISELTESR.L	2	SPEC1_HUMAN
Q5M775	2	3.7781	R.AEQLSQENEK.L	3	SPEC1_HUMAN
Q5M775	2	2.9038	K.LMNLLQER.V	1	SPEC1_HUMAN
Q5M775	2	4.5078	K.LLNIQQQLTCSLR.K	1	SPEC1_HUMAN
Q5M775	4	5.0931	K.LHNNQLISELESSVIKLEEQK.S	1	SPEC1_HUMAN
Q5M775	4	4.7421	K.ASESDAEIKDMKETIFELEDQVEQHR.A	1	SPEC1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5M775	3	3.8083	R.AEQLSQENEKLM#NLLQER.V	1	SPEC1_HUMAN
Sperm-associated antigen 1 - Homo sapiens (Human)					
Q07617	2	2.7536	K.C*SDVKHLEKILC*VLR.S	1	SPAG1_HUMAN
Q07617	2	3.0007	R.SISALPTVVAYNNR.A	1	SPAG1_HUMAN
Sperm-associated antigen 7 - Homo sapiens (Human)					
O75391	2	3.0574	K.DAAHM#LQANK.T	2	SPAG7_HUMAN
O75391	2	3.5325	R.SIEEAMNEIR.A	2	SPAG7_HUMAN
O75391	2	2.8602	R.SIEEAM#NEIR.A	1	SPAG7_HUMAN
O75391	2	3.8915	K.DAAHMLQANK.T	3	SPAG7_HUMAN
O75391	2	3.644	K.EVSDFIQDSGQIK.K	1	SPAG7_HUMAN
Spermatogenesis-associated serine-rich protein 2. - Homo sapiens (Human)					
Q86XZ4	2	2.9136	K.SVSIQEEQSAPSSEK.G	1	Q86XZ4_HUMAN
Q86XZ4	2	2.8758	R.ELEDPESAM#LDTLDR.T	1	Q86XZ4_HUMAN
Q86XZ4	1	2.1424	R.TSQTEAVNS.-	1	Q86XZ4_HUMAN
Q86XZ4	2	3.65	R.YQSAPSQAPGNTIER.G	3	Q86XZ4_HUMAN
Sperm-specific antigen 2 - Homo sapiens (Human)					
P28290	2	2.911	R.ASVALTPTAPSR.T	1	SSFA2_HUMAN
P28290	2	4.8764	R.EIVSGLLAAVSSSK.A	3	SSFA2_HUMAN
P28290	3	4.2914	R.IGSMSSVTSNKETDPPPPLTR.S	1	SSFA2_HUMAN
P28290	2	3.2673	K.NVEQDELQQVIR.E	2	SSFA2_HUMAN
S-phase kinase-associated protein 1A - Homo sapiens (Human)					
P63208	2	2.769	K.GKTPEEIRK.T	1	SKP1_HUMAN
P63208	2	3.3999	K.NDFTEEEEAQVR.K	3	SKP1_HUMAN
P63208	2	4.8563	K.TFNIKNDFTEEEEAQVR.K	4	SKP1_HUMAN
P63208	3	5.4507	K.TFNIKNDFTEEEEAQVRK.E	2	SKP1_HUMAN
P63208	3	5.1981	K.TM#LEDLGM#DDEGDDDPVPLPNVNAAILKK.	2	SKP1_HUMAN
P63208	3	4.3129	K.TMLEDLGMDDEGDDDPVPLPNVNAAILKK.V	1	SKP1_HUMAN
Splicing factor 1 - Homo sapiens (Human)					
Q15637	3	4.3037	K.VMIPQDEYPEINFVGLLIGPR.G	3	SF01_HUMAN
Q15637	3	4.0312	R.VSDKVM#IPQDEYPEINFVGLLIGPR.G	1	SF01_HUMAN
Q15637	4	6.1131	R.HNLITEMVALNPDFKPPADYKPPATR.V	2	SF01_HUMAN
Q15637	3	4.0294	K.DGQM#LPGEDEPLHALVTANTMENVK.K	1	SF01_HUMAN
Q15637	2	3.8284	R.ILRPWQSSETR.S	3	SF01_HUMAN
Splicing factor 3B subunit 1 - Homo sapiens (Human)					
O75533	2	3.5116	K.THEDIEAQIR.E	4	SF3B1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O75533	3	3.9052	R.NRPLSDEELDAMFPEGYK.V	1	SF3B1_HUMAN
O75533	2	2.7949	R.LDPFADGGKTPDPK.M	1	SF3B1_HUMAN
O75533	2	3.4381	R.GGDSIGETPTPGASK.R	2	SF3B1_HUMAN
O75533	3	4.0008	R.GDTPGHATPGHGGATSSAR.K	1	SF3B1_HUMAN
O75533	2	3.7213	R.AKGSETPGATPGSK.I	2	SF3B1_HUMAN
O75533	2	4.9528	R.WDQTADQTPGATPK.K	3	SF3B1_HUMAN
O75533	3	5.2516	K.KPGYHAPVALLNDIPQSTEQYDPFAEHRPPK.	2	SF3B1_HUMAN
O75533	3	4.994	K.KLSSWDQAETPGHTPSLR.W	2	SF3B1_HUMAN
O75533	3	5.1268	K.KAALDEAQGVGLDSTGYYDQEIYGGSDSR.F	1	SF3B1_HUMAN
O75533	3	3.9224	K.IADREDEYKK.H	1	SF3B1_HUMAN
O75533	2	3.0252	K.GSETPGATPGSK.I	3	SF3B1_HUMAN
O75533	2	2.8894	K.AALDEAQGVGLDSTGYYDQEIYGGSDSR.F	1	SF3B1_HUMAN
O75533	2	3.4875	K.VVNGAAASQPPSK.R	2	SF3B1_HUMAN
O75533	2	3.5849	K.VVNGAAASQPPSKR.K	1	SF3B1_HUMAN
Splicing factor 3B subunit 2 - Homo sapiens (Human)					
Q13435	3	7.4529	K.ESRQEEMNSQQEEEEMETDAR.S	3	SF3B2_HUMAN
Q13435	3	4.7143	R.QEEMNSQQEEEEMETDAR.S	3	SF3B2_HUMAN
Q13435	3	5.3215	R.QEEMNSQQEEEEM#ETDAR.S	1	SF3B2_HUMAN
Q13435	2	3.1736	R.QEEM#NSQQEEEEMETDAR.S	1	SF3B2_HUMAN
Q13435	2	3.5694	R.GVSSESSGDREKDSTR.S	2	SF3B2_HUMAN
Q13435	2	2.8722	R.GPPPPPGDENREMDDPSVGPK.I	1	SF3B2_HUMAN
Q13435	2	4.4223	R.AAVLLEQERQQEIAK.M	4	SF3B2_HUMAN
Q13435	1	2.3795	K.IPQALEK.I	1	SF3B2_HUMAN
Q13435	3	5.8383	K.ESRQEEMNSQQEEEEM#ETDAR.S	1	SF3B2_HUMAN
Q13435	3	6.8698	K.ESRQEEM#NSQQEEEEMETDAR.S	2	SF3B2_HUMAN
Q13435	2	3.0171	K.MGTPVPRPPQDMGQIGVR.T	2	SF3B2_HUMAN
Splicing factor 4 - Homo sapiens (Human)					
Q8IWZ8	2	4.0987	R.DVDASPSPLSVQDLK.G	4	SF04_HUMAN
Q8IWZ8	3	3.8481	K.AQTSTDAPTSAPSAPPSTPTPSAGKR.S	1	SF04_HUMAN
Q8IWZ8	3	3.8165	K.MNMNILHQEELIAQK.K	1	SF04_HUMAN
Splicing factor U2AF 35 kDa subunit - Homo sapiens (Human)					
Q01081	2	3.9628	R.NPQNSSQSADGLR.C	2	U2AF1_HUMAN
Splicing factor, arginine/serine-rich 1 - Homo sapiens (Human)					
Q07955	2	4.17	R.IYVGNLPPDIR.T	26	SFRS1_HUMAN
Q07955	3	4.3374	R.TKDIEDVFYK.Y	5	SFRS1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q07955	2	3.3102	R.SHEGETAYIR.V	8	SFRS1_HUMAN
Q07955	4	5.1012	R.RGGPPFAFVEFEDPRDAEDAVYGR.D	2	SFRS1_HUMAN
Q07955	3	3.9028	R.KEDMTYAVRK.L	3	SFRS1_HUMAN
Q07955	1	3.3108	K.DIEDVFYK.Y	2	SFRS1_HUMAN
Q07955	2	2.9623	R.KEDM#TYAVR.K	3	SFRS1_HUMAN
Q07955	2	2.8106	R.GGPPFAFVEFEDPR.D	1	SFRS1_HUMAN
Q07955	2	3.6863	R.EAGDVCYADVYR.D	3	SFRS1_HUMAN
Q07955	2	3.7299	R.DGYDYDGYR.L	10	SFRS1_HUMAN
Q07955	2	2.9126	R.DGTGVVEFVRK.E	4	SFRS1_HUMAN
Q07955	2	3.2492	R.DGTGVVEFVR.K	6	SFRS1_HUMAN
Q07955	3	4.1168	R.DAEDAVYGRDGYDYDGYR.L	3	SFRS1_HUMAN
Q07955	2	3.0342	R.DAEDAVYGR.D	2	SFRS1_HUMAN
Q07955	3	4.3416	K.FRSHEGETAYIR.V	4	SFRS1_HUMAN
Q07955	2	3.2764	R.KEDMTYAVR.K	7	SFRS1_HUMAN
Splicing factor, arginine/serine-rich 15 - Homo sapiens (Human)					
O95104	3	5.9037	K.GVSEAAVLKPSEELPAEATSSVEPEKDSGSA	2	SFR15_HUMAN
O95104	2	6.1165	K.TVEPPISQVGNVDTASELEK.G	5	SFR15_HUMAN
Splicing factor, arginine/serine-rich 16 - Homo sapiens (Human)					
Q8N2M8	2	3.1285	K.QATTYGMADGDFVR.M	1	SFR16_HUMAN
Q8N2M8	3	3.7281	K.LTRPAASPAVGEK.L	1	SFR16_HUMAN
Splicing factor, arginine/serine-rich 2 - Homo sapiens (Human)					
Q01130	2	3.2472	R.YGRPPDSHHSR.R	4	SFRS2_HUMAN
Q01130	1	2.5722	R.YGGGGYGR.R	4	SFRS2_HUMAN
Q01130	2	3.3558	R.RYGGGGYGR.R	6	SFRS2_HUMAN
Splicing factor, arginine/serine-rich 2B - Homo sapiens (Human)					
Q9BRL6	2	3.3748	R.DAQDAEAAM#DGAELDGR.E	2	Q9BRL6_HUMA
Q9BRL6	2	4.3278	R.DAQDAEAAMDGAELDGR.E	1	Q9BRL6_HUMA
Splicing factor, arginine/serine-rich 3 - Homo sapiens (Human)					
P84103	3	4.8896	R.VRVELSNGEKR.S	1	SFRS3_HUMAN
P84103	2	3.9218	K.VYVGNLGNNGNK.T	4	SFRS3_HUMAN
P84103	2	4.2894	K.VYVGNLGNNGNKTELER.A	2	SFRS3_HUMAN
P84103	2	2.9265	R.AFGYYGPLR.S	2	SFRS3_HUMAN
P84103	1	2.5805	R.DAADAVR.E	2	SFRS3_HUMAN
P84103	2	2.9116	R.VELSNGEKR.S	1	SFRS3_HUMAN
Splicing factor, arginine/serine-rich 4 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q08170	2	3.0446	R.LVEDKPGSR.R	2	SFRS4_HUMAN
Q08170	3	5.1471	R.KNEGVIEFVSYSDMKR.A	2	SFRS4_HUMAN
Q08170	2	3.7071	R.GESENAGTNQETR.S	3	SFRS4_HUMAN
Q08170	2	4.2698	R.EGRGESENAGTNQETR.S	7	SFRS4_HUMAN
Q08170	3	4.1394	R.ALEKLDGTEVNGRK.I	1	SFRS4_HUMAN
Q08170	2	3.2458	R.ALEKLDGTEVNGR.K	1	SFRS4_HUMAN
Q08170	2	2.7106	K.SKPNLPSESR.S	1	SFRS4_HUMAN
Q08170	2	4.5623	K.NEGVIEFVSYSDMKR.A	3	SFRS4_HUMAN
Q08170	2	3.7471	K.IQNNDNVGKPK.S	2	SFRS4_HUMAN
Q08170	2	3.0211	K.LDGTEVNGR.K	2	SFRS4_HUMAN
Splicing factor, arginine/serine-rich 5 - Homo sapiens (Human)					
Q13243	2	2.9234	R.VSWQDLK.D	2	SFRS5_HUMAN
Q13243	2	3.4644	R.VSWQDLKDFMR.Q	3	SFRS5_HUMAN
Splicing factor, arginine/serine-rich 7 - Homo sapiens (Human)					
Q16629	2	3.1237	R.AFSYYGPLR.T	2	SFRS7_HUMAN
Q16629	2	2.7671	R.RPFDPNDR.C	2	SFRS7_HUMAN
Q16629	2	2.8333	R.VELSTGMPR.R	1	SFRS7_HUMAN
Q16629	2	3.321	R.VRVELSTGMPR.R	2	SFRS7_HUMAN
Q16629	3	4.2283	R.VRVELSTGMPRR.S	1	SFRS7_HUMAN
Q16629	2	4.6135	K.VYVGNLGTGAGKGELER.A	2	SFRS7_HUMAN
Splicing factor, arginine/serine-rich 9 - Homo sapiens (Human)					
Q13242	2	3.3133	R.HGLVPFAFVR.F	3	SFRS9_HUMAN
Q13242	2	3.1654	R.SHEGETSYIR.V	2	SFRS9_HUMAN
Q13242	2	3.9973	R.IYVGNLPTDVR.E	5	SFRS9_HUMAN
Q13242	3	3.7001	R.FEDPRDAEDAIYGR.N	1	SFRS9_HUMAN
Q13242	2	3.3371	R.EKDLEDLFYK.Y	1	SFRS9_HUMAN
Q13242	2	3.1992	R.KEDMEYALR.K	2	SFRS9_HUMAN
Splicing factor, proline- and glutamine-rich - Homo sapiens (Human)					
P23246	2	2.8816	R.GM#GPGTPAGYGR.G	1	SFPQ_HUMAN
P23246	3	3.7387	R.RMEELHNQEM#QK.R	1	SFPQ_HUMAN
P23246	2	3.6881	R.RMEELHNQEMQK.R	1	SFPQ_HUMAN
P23246	3	4.352	R.PVIVEPLEQLDDEDGLPEKLAQK.N	1	SFPQ_HUMAN
P23246	3	4.6089	R.NLSPYVSNELLEEAFSQFGPIER.A	2	SFPQ_HUMAN
P23246	2	3.2479	R.MGGGGAMNMGDPYGSGGQK.F	2	SFPQ_HUMAN
P23246	2	2.9167	R.MGGGGAMNM#GDPYGSGGQK.F	1	SFPQ_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P23246	2	3.7051	R.MEELHNQEMQK.R	1	SFPQ_HUMAN
P23246	4	5.4671	K.DAKDKLESEMEDAYHEHQANLLR.Q	2	SFPQ_HUMAN
P23246	2	5.1842	R.FAQHGTFEYEYSQR.W	6	SFPQ_HUMAN
P23246	2	2.8853	R.EEYEGPNKKPR.F	1	SFPQ_HUMAN
P23246	2	3.163	K.SLDEMEKQQR.E	2	SFPQ_HUMAN
P23246	3	4.8149	K.MPGGPKPGGGPGLSTPGGHPKPPHR.G	2	SFPQ_HUMAN
P23246	3	3.8415	K.M#PGGPKPGGGPGLSTPGGHPKPPHR.G	2	SFPQ_HUMAN
P23246	2	2.7168	R.SPPPGMGLNQNR.G	1	SFPQ_HUMAN
P23246	2	3.5324	K.LAQKNPMYQK.E	2	SFPQ_HUMAN
P23246	2	4.451	R.FGQGGAGPVGGQGPR.G	3	SFPQ_HUMAN
Squamous cell carcinoma antigen recognized by T-cells 3 - Homo sapiens (Human)					
Q15020	1	2.4191	K.MSNADFAK.L	1	SART3_HUMAN
Q15020	3	6.7294	R.ALQRPSAAAPQAENGPAAAPAVAAPAATEAP	1	SART3_HUMAN
SRA stem-loop-interacting RNA-binding protein, mitochondrial precursor - Homo sapiens (Human					
Q9GZT3	2	4.1138	K.LPQTSDDEKKDF.-	4	SLIRP_HUMAN
Q9GZT3	2	4.3128	R.GLGWVQFSSEEGLR.N	2	SLIRP_HUMAN
Q9GZT3	2	4.5824	R.NALQQENHIIDGVK.V	3	SLIRP_HUMAN
Q9GZT3	2	3.1593	R.SINQPVAFVR.R	2	SLIRP_HUMAN
Src substrate cortactin - Homo sapiens (Human)					
Q14247	3	3.7775	R.QDSAAVGFDYKEK.L	7	SRC8_HUMAN
Q14247	3	5.0284	K.FGVQSERQDSAAVGFDYKEK.L	1	SRC8_HUMAN
Q14247	3	5.1925	K.LAKHESQQDYSK.G	9	SRC8_HUMAN
Q14247	2	3.979	K.LQLHESQKDYK.T	3	SRC8_HUMAN
StAR-related lipid transfer protein 13 - Homo sapiens (Human)					
Q9Y3M8	2	5.7019	R.TGGLVISGPMLQQEPESFK.A	1	STA13_HUMAN
Stathmin - Homo sapiens (Human)					
P16949	2	3.7078	K.LTHKMEANKENR.E	2	STMN1_HUMAN
P16949	2	2.9235	K.M#AEEKLTHK.M	1	STMN1_HUMAN
P16949	2	3.3655	K.MAEEKLTHK.M	4	STMN1_HUMAN
P16949	2	3.5602	K.NKESKDPADETEAD.-	3	STMN1_HUMAN
P16949	2	3.3766	K.ESKDPADETEAD.-	3	STMN1_HUMAN
Stathmin-3 - Homo sapiens (Human)					
Q9NZ72	2	2.9259	K.ALEENNNFSR.Q	1	STMN3_HUMAN
Q9NZ72	2	2.8112	K.SPSDLSPESPMLSSPPK.K	1	STMN3_HUMAN
STE20-like serine/threonine-protein kinase - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9H2G2	1	3.7344	K.ALGSEVQDASK.V	6	SLK_HUMAN
Q9H2G2	1	2.3509	R.SVVADTDQK.A	1	SLK_HUMAN
Q9H2G2	2	4.1509	R.KLQEQEVFFK.M	4	SLK_HUMAN
Q9H2G2	2	3.1646	R.INSTATPDQDRDKIK.Q	1	SLK_HUMAN
Q9H2G2	2	3.7791	R.INSTATPDQDRDK.I	2	SLK_HUMAN
Q9H2G2	2	3.0929	R.EAAIWELEER.H	1	SLK_HUMAN
Q9H2G2	2	2.7141	K.VTTQIDKEK.K	1	SLK_HUMAN
Q9H2G2	2	4.0979	K.MTGESECLNPSTQSR.I	4	SLK_HUMAN
Q9H2G2	2	3.6532	K.DSGSISLQETR.R	2	SLK_HUMAN
Q9H2G2	3	4.2359	K.LKELDEEHSQELKEWR.E	1	SLK_HUMAN
Sterile alpha motif domain-containing protein 14 - Homo sapiens (Human)					
Q8IZD0	2	3.0548	K.SASQESTLSDDSTPPSSSPK.I	1	SAM14_HUMAN
Q8IZD0	2	2.7	R.DSASSAEDGEGSDGPGGK.V	1	SAM14_HUMAN
Sterile alpha motif domain-containing protein 9 - Homo sapiens (Human)					
Q5K651	3	4.1748	K.EHLVDMGITHGPAIQIEELFK.E	2	SAMD9_HUMAN
Q5K651	3	4.6187	K.HREILTEQDVNGAVLK.W	2	SAMD9_HUMAN
Q5K651	2	3.3854	K.TAIEDSIQTSK.M	3	SAMD9_HUMAN
Steroid receptor RNA activator 1 - Homo sapiens (Human)					
Q9HD15	2	2.9548	K.RVAAPQDGSPR.V	2	SRA1_HUMAN
Q9HD15	2	3.2296	R.SLFSEEAANEEK.S	2	SRA1_HUMAN
Q9HD15	3	4.7907	R.VPASETSPGPPPM#GPPPPSSK.A	2	SRA1_HUMAN
Q9HD15	2	4.739	R.VPASETSPGPPPMGPPPPSSK.A	3	SRA1_HUMAN
STIP1 homology and U box-containing protein 1 - Homo sapiens (Human)					
Q9UNE7	2	2.7757	R.LGAGGGSPEK.S	2	STUB1_HUMAN
Q9UNE7	3	3.8601	R.NHEGDEDDSHVR.A	3	STUB1_HUMAN
Stomatin-like protein 2 - Homo sapiens (Human)					
Q9UJZ1	2	2.9265	R.DVQGTDASLDEELDRVK.M	1	STML2_HUMAN
Stress-70 protein, mitochondrial precursor - Homo sapiens (Human)					
P38646	2	3.2762	K.DSETGENIR.Q	1	GRP75_HUMAN
P38646	2	3.1976	K.VLENAEGAR.T	2	GRP75_HUMAN
P38646	2	3.2353	K.VQQTVQDLFGR.A	2	GRP75_HUMAN
P38646	2	3.4594	R.AQFEGIVTDLIR.R	1	GRP75_HUMAN
P38646	2	3.165	R.KDSETGENIR.Q	2	GRP75_HUMAN
P38646	2	2.8627	R.RYDDPEVQK.D	1	GRP75_HUMAN
P38646	2	3.0178	R.TTPSVVAFTADGER.L	1	GRP75_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Stress-induced-phosphoprotein 1 - Homo sapiens (Human)					
P31948	2	3.6668	K.AAALEFLNR.F	2	STIP1_HUMAN
P31948	2	3.4672	K.AAALEFLNRFEEAK.R	1	STIP1_HUMAN
Stromal interaction molecule 1 precursor - Homo sapiens (Human)					
Q13586	3	3.7472	R.DLTHSDSESSLHMSDR.Q	1	STIM1_HUMAN
Q13586	3	5.116	R.LIEGVHPGSLVEKLPDSPALAK.K	1	STIM1_HUMAN
Q13586	2	4.3127	R.SHSPSSPDPDTPSPVGDSR.A	4	STIM1_HUMAN
Q13586	3	4.2892	K.KAVAEEDNGSIGEETDSSPGR.K	2	STIM1_HUMAN
Structural maintenance of chromosomes protein 1A - Homo sapiens (Human)					
Q14683	2	3.3034	R.MEEESQSQGR.D	2	SMC1A_HUMAN
Structural maintenance of chromosomes protein 5 - Homo sapiens (Human)					
Q8IY18	2	2.7658	R.DPSSEVPSK.R	1	SMC5_HUMAN
Succinyl-CoA ligase [ADP-forming] beta-chain, mitochondrial precursor - Homo sapiens (Human)					
Q9P2R7	2	3.1193	K.SPDEAYAIAK.K	1	SUCB1_HUMAN
Succinyl-CoA ligase [GDP-forming] beta-chain, mitochondrial precursor - Homo sapiens (Human)					
Q96I99	2	4.5516	R.FFVADTANEALEAAK.R	3	SUCB2_HUMAN
Succinyl-CoA ligase [GDP-forming] subunit alpha, mitochondrial precursor - Homo sapiens (Hum					
P53597	3	3.8978	K.ISALQSAGVVVSM#SPAQLGTTIYKEFEK.R	1	SUCA_HUMAN
Sulfatase-modifying factor 2 precursor - Homo sapiens (Human)					
Q8NBJ7	2	3.2593	R.MGNTPDSASDNLGFR.C	1	SUMF2_HUMAN
Q8NBJ7	2	3.2354	K.SVLWWLPVEK.A	1	SUMF2_HUMAN
Q8NBJ7	2	3.8624	R.LPTEEEWEFAAR.G	2	SUMF2_HUMAN
Superoxide dismutase [Cu-Zn] - Homo sapiens (Human)					
P00441	3	9.3334	K.GLTEGLHGFHVHEFGDNTAGCTSAGPHFNPL	60	SODC_HUMAN
P00441	3	6.8109	R.TLVVHEKADDLGKGGNEESTK.T	7	SODC_HUMAN
P00441	2	4.8468	R.TLVVHEKADDLGK.G	5	SODC_HUMAN
P00441	1	2.7661	R.LACGVIGIAQ.-	9	SODC_HUMAN
P00441	1	2.2613	K.VWGSIK.G	5	SODC_HUMAN
P00441	2	3.7954	R.HVGDLGNVTADK.D	2	SODC_HUMAN
P00441	3	5.164	K.GLTEGLHGFHVHEFGDNTAGCTSAGPHFNPL	1	SODC_HUMAN
P00441	3	5.2402	K.ADDLGKGGNEESTKTGNAGSR.L	4	SODC_HUMAN
P00441	4	7.9798	R.HVGDLGNVTADKDGVADVSIEDSVISLSGDH	4	SODC_HUMAN
P00441	2	4.3014	K.ADDLGKGGNEESTK.T	28	SODC_HUMAN
P00441	3	6.2233	K.AVCVLKGDGPVQGIINFEQK.E	54	SODC_HUMAN
P00441	4	5.4549	K.AVCVLKGDGPVQGIINFEQKESNGPVK.V	4	SODC_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P00441	2	3.4697	K.ESNGPVKVWGSIK.G	5	SODC_HUMAN
P00441	3	7.7668	K.GLTEGLHGFHVHEFGDNTAGC*TSAGPHFNP	4	SODC_HUMAN
P00441	2	4.6622	K.GDGPVQGIINFEQK.E	11	SODC_HUMAN
P00441	2	4.7785	K.GDGPVQGIINFEQKESNGPVK.V	3	SODC_HUMAN
Supervillin - Homo sapiens (Human)					
O95425	2	3.5785	R.RGQELSATR.Q	5	SVIL_HUMAN
O95425	2	3.7804	K.TPTGEGLLDSPSK.T	2	SVIL_HUMAN
O95425	2	4.7138	R.SNEEEETSDSSLEK.Q	9	SVIL_HUMAN
O95425	2	3.5417	R.NSPELASESVTQR.R	2	SVIL_HUMAN
O95425	2	3.5949	R.DASSLYPGTETMGLR.T	3	SVIL_HUMAN
O95425	2	3.5401	R.DASSLYPGTETM#GLR.T	1	SVIL_HUMAN
O95425	2	4.4822	R.YQTQPVTLGEVEQVQSGK.L	3	SVIL_HUMAN
O95425	2	4.1783	K.SQAWQPLVEGSENK.G	2	SVIL_HUMAN
O95425	2	2.7524	K.SFDEQNVPK.R	1	SVIL_HUMAN
O95425	2	2.9382	K.MNAQGNLDLR.D	2	SVIL_HUMAN
O95425	2	2.791	R.SPVEMDEDFDVIFDPYAPK.L	1	SVIL_HUMAN
O95425	3	4.0562	K.HAHSSSLQQAASR.S	3	SVIL_HUMAN
O95425	2	3.651	K.FSSSIENSDSPVR.S	2	SVIL_HUMAN
O95425	3	5.088	K.DQTNEGKELAEQGEPDSSTLSLAEK.L	1	SVIL_HUMAN
O95425	2	3.2456	K.TLLALEGDGLVR.S	1	SVIL_HUMAN
Suppressor of hairy wing homolog 4 - Homo sapiens (Human)					
Q6N043	2	2.9254	K.SNTSKPNTIKSNASK.P	4	SUHW4_HUMAN
Surfeit locus protein 6 - Homo sapiens (Human)					
O75683	2	3.5977	K.TQGSETAGPPK.K	3	SURF6_HUMAN
Survival motor neuron protein - Homo sapiens (Human)					
Q16637	2	3.2179	R.GTGQSDDSDIWDDTALIK.A	1	SMN_HUMAN
Survival of motor neuron-related-splicing factor 30 - Homo sapiens (Human)					
O75940	2	4.0709	K.RSIFASPESVTGK.V	1	SPF30_HUMAN
O75940	2	3.0176	R.SIFASPESVTGK.V	3	SPF30_HUMAN
O75940	2	2.9024	K.KEMIAQQR.E	2	SPF30_HUMAN
SUV3-like protein 1 - Homo sapiens (Human)					
O43630	2	2.8646	R.LVQQGLLTPDMLK.Q	1	O43630_HUMAN
SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily A containin					
Q9H4L7	3	6.4181	R.RNDDISELEDLSELEDLKDAK.L	2	SMRCD_HUMAN
SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily A-like protei					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9NZC9	2	2.9217	K.LLAEQHQR.T	2	SMAL1_HUMAN
Q9NZC9	2	4.0084	R.TSSGTSIAGNPFQAK.Q	2	SMAL1_HUMAN
Switch-associated protein 70 - Homo sapiens (Human)					
Q9UH65	2	3.1339	K.SSELEQYLQR.V	1	SWP70_HUMAN
Synaptophysin-like protein 1 - Homo sapiens (Human)					
Q16563	2	3.2369	K.TVTATFGYPFR.L	1	SYPL1_HUMAN
Synaptopodin - Homo sapiens (Human)					
Q8N3V7	2	4.1811	K.TGILEESMAR.R	4	SYNPO_HUMAN
Q8N3V7	2	4.8741	R.SPPSYSVLYPSSDPK.S	2	SYNPO_HUMAN
Q8N3V7	3	7.4243	R.QRDQGEVGVEEEPFALGAEASNFQQEPAPR.	2	SYNPO_HUMAN
Q8N3V7	3	3.9934	R.KSMFTFVEKPK.V	3	SYNPO_HUMAN
Q8N3V7	2	4.4658	R.AASPAKPSSLDLVPNLPK.G	5	SYNPO_HUMAN
Q8N3V7	2	4.5654	K.YVIESSSHTPELAR.C	2	SYNPO_HUMAN
Q8N3V7	2	5.2617	R.TARPFGIQAPGGTSQMER.S	4	SYNPO_HUMAN
Q8N3V7	2	2.7583	K.TGILEESM#AR.R	1	SYNPO_HUMAN
Q8N3V7	2	2.7258	K.SMFTFVEKPK.V	1	SYNPO_HUMAN
Q8N3V7	2	3.8796	K.PKPNQNLSEASGK.G	4	SYNPO_HUMAN
Q8N3V7	2	3.9738	K.ENAALLTANGLHLSQNR.E	2	SYNPO_HUMAN
Q8N3V7	2	3.2206	K.APAPQPPSLPDR.S	2	SYNPO_HUMAN
Q8N3V7	2	2.8541	K.YPTNAPGAFR.V	2	SYNPO_HUMAN
Q8N3V7	2	3.0536	K.VASEEEEVPLVVYLK.E	1	SYNPO_HUMAN
Synaptopodin-2 - Homo sapiens (Human)					
Q9UMS6	2	3.2743	K.VNSALAM#K.Q	2	SYNP2_HUMAN
Q9UMS6	2	3.7618	R.VIQESEAGDAGLPR.V	1	SYNP2_HUMAN
Q9UMS6	2	3.5867	R.TGILQEAK.R	1	SYNP2_HUMAN
Q9UMS6	1	2.1811	R.GAQLFAK.R	2	SYNP2_HUMAN
Q9UMS6	2	3.3979	R.AQSPTPSLPASWK.Y	1	SYNP2_HUMAN
Q9UMS6	2	4.6411	R.APPPVAYNPIHSPSYPLAALK.S	2	SYNP2_HUMAN
Q9UMS6	2	4.0724	K.VSPNPELLSLLQNSEGKR.G	3	SYNP2_HUMAN
Q9UMS6	1	2.1014	K.VNSALAMK.Q	1	SYNP2_HUMAN
Q9UMS6	2	3.3942	K.SGVTIQVWKPSVVEE.-	1	SYNP2_HUMAN
Q9UMS6	4	6.0951	K.GPQAAVASQNYTPKPTVSTPTVNAVQPGAV	1	SYNP2_HUMAN
Q9UMS6	2	3.146	K.VSPNPELLSLLQNSEGK.R	3	SYNP2_HUMAN
Synaptosomal-associated protein 23 - Homo sapiens (Human)					
O00161	2	4.6368	R.ILGLAIESQDAGIK.T	1	SNP23_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O00161	3	5.8733	K.DMALNIGNEIDAQNPQIK.R	2	SNP23_HUMAN
O00161	3	4.2794	K.DMALNIGNEIDAQNPQIKR.I	1	SNP23_HUMAN
Synaptosomal-associated protein 25 - Homo sapiens (Human)					
P60880	3	4.3084	R.RADQLADESLESTRR.M	2	SNP25_HUMAN
Synaptosomal-associated protein 29 - Homo sapiens (Human)					
O95721	2	3.5302	K.EAISTSKEQEAK.Y	2	SNP29_HUMAN
O95721	2	3.3089	K.VGVASSEELAR.Q	2	SNP29_HUMAN
Synaptotagmin-like protein 2 - Homo sapiens (Human)					
Q9HCH5	2	3.192	K.LPEGHSSQQTK.N	2	SYTL2_HUMAN
Q9HCH5	2	3.8067	K.STVADTSIQKLEK.S	1	SYTL2_HUMAN
Q9HCH5	2	2.7136	R.DRQQGSEEEPSPVLK.T	1	SYTL2_HUMAN
Q9HCH5	3	3.7277	R.KPSLFHQSTSSPYVSK.S	1	SYTL2_HUMAN
Synaptotagmin-like protein 4 - Homo sapiens (Human)					
Q96C24	2	3.3047	K.SALEAESESLDSFTADSDSTSR.R	1	SYTL4_HUMAN
Syndecan-1 precursor - Homo sapiens (Human)					
P18827	2	4.2274	K.EGEAVVLPEVEPGLTAR.E	2	SDC1_HUMAN
Syndecan-4 precursor - Homo sapiens (Human)					
P31431	2	3.5786	K.KLEENEVIPK.R	5	SDC4_HUMAN
P31431	2	3.4	R.AGSGSQVPTEPK.K	2	SDC4_HUMAN
P31431	2	2.8248	R.ISPVEESEDVSNK.V	1	SDC4_HUMAN
Syntaxin-3 - Homo sapiens (Human)					
Q13277	2	2.7569	K.HIEEDEVR.S	1	STX3_HUMAN
Q13277	2	3.732	K.LYSIILSAPIPEPK.T	3	STX3_HUMAN
Syntaxin-6 - Homo sapiens (Human)					
O43752	2	3.216	K.AVNTAQGLFQR.W	2	STX6_HUMAN
Syntaxin-7 - Homo sapiens (Human)					
O15400	2	3.2724	R.VSGSFPEDSSKER.N	2	STX7_HUMAN
O15400	3	3.8269	K.ETDKYIKEFGSLPTTPSEQR.Q	1	STX7_HUMAN
O15400	2	3.232	K.ITQCSVEIQR.T	2	STX7_HUMAN
O15400	1	2.3042	R.ISSNIQK.I	3	STX7_HUMAN
O15400	2	4.9416	R.LVAEFTTSLTNFQK.V	5	STX7_HUMAN
O15400	2	2.7772	R.TLNQLGTPQDSPELR.Q	1	STX7_HUMAN
Syntaxin-binding protein 3 - Homo sapiens (Human)					
O00186	2	3.4457	K.TSLSALTQLMK.K	1	STXB3_HUMAN
Talin-1 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y490	3	4.6592	R.SGASGPENFQVGSM#PPAQQQITSGQM#HR.	1	TLN1_HUMAN
Q9Y490	2	3.8294	K.ADAEGESDLENSR.K	2	TLN1_HUMAN
Q9Y490	3	5.8393	R.SGASGPENFQVGSM#PPAQQQITSGQMHR.	2	TLN1_HUMAN
Q9Y490	2	3.5981	K.TLAESALQLLYTAK.E	1	TLN1_HUMAN
Q9Y490	3	3.9906	K.QAAASATQTIAAAQHAASTPK.A	1	TLN1_HUMAN
Q9Y490	2	3.6933	K.MVGGIAQIIAAQEEMLR.K	1	TLN1_HUMAN
Q9Y490	2	4.3273	K.KSTVLQQQYNR.V	1	TLN1_HUMAN
Q9Y490	2	3.9715	K.GAAAHPDSEEQQQR.L	3	TLN1_HUMAN
Q9Y490	2	2.7099	K.EADESLNFEEQILEAAK.S	1	TLN1_HUMAN
Q9Y490	3	4.3426	R.VGKVEHGSVALPAIMR.S	1	TLN1_HUMAN
Target of myb1 - Homo sapiens (Human)					
Q86X74	3	7.2495	K.AADRLPNLSSPSAEGPPGPPSGPAPR.K	4	Q86X74_HUMAN
Q86X74	3	4.4278	K.EVKYEAPQATDGLAGALDAR.Q	6	Q86X74_HUMAN
Q86X74	2	3.1424	K.TQEKDDDMLFAL.-	2	Q86X74_HUMAN
Q86X74	2	4.0561	K.YEAPQATDGLAGALDAR.Q	3	Q86X74_HUMAN
Q86X74	2	3.188	R.LEDEFDMFALTR.G	2	Q86X74_HUMAN
TATA element modulatory factor - Homo sapiens (Human)					
P82094	2	2.8381	K.SPVVSKPPAK.S	1	TMF1_HUMAN
P82094	2	2.8727	R.TPETTESQVK.D	2	TMF1_HUMAN
P82094	3	4.5651	R.QIENLQATLGSQTSSWEKLEK.N	2	TMF1_HUMAN
P82094	3	3.7053	R.AKDKENENMVAK.L	1	TMF1_HUMAN
P82094	2	3.2231	K.IQMSSMESQNSLLR.Q	2	TMF1_HUMAN
P82094	2	3.6341	K.DSEAQEAALSR.E	1	TMF1_HUMAN
P82094	2	3.2515	K.TLLNSQLEMER.M	2	TMF1_HUMAN
TATA-binding protein-associated factor 2N - Homo sapiens (Human)					
Q92804	2	3.7585	K.AAIDWFDGKEFHGNIIK.V	2	RBP56_HUMAN
Q92804	3	4.0797	K.QSSYSQQPYNNQGQQQNM#ESSGSQGGR.	1	RBP56_HUMAN
Q92804	3	4.9819	R.YGEDNRGYGGSQGGGR.G	1	RBP56_HUMAN
TATA-binding protein-associated phosphoprotein - Homo sapiens (Human)					
Q01658	2	2.9413	R.LENLGIPEEELLRQQQELFAK.A	1	TBAP_HUMAN
Tax1-binding protein 3 - Homo sapiens (Human)					
O14907	2	5.0872	R.VSEGGPAEIAGLQIGDK.I	2	TX1B3_HUMAN
TBC1 domain family member 1 - Homo sapiens (Human)					
Q86TI0	2	3.7559	R.SLTESLESILSR.G	1	TBCD1_HUMAN
TBC1 domain family member 10A - Homo sapiens (Human)					

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9BXI6	2	2.8166	R.APAAGESLSGTR.E	2	TB10A_HUMAN
TBC1 domain family member 12 - Homo sapiens (Human)					
O60347	3	4.6057	R.LRLEGPGDEDADGAGSPSDWASPLEDPLR.S	1	TBC12_HUMAN
TBC1 domain family member 22B - Homo sapiens (Human)					
Q9NU19	2	2.7842	R.SQSTTSDVPANYK.V	1	TB22B_HUMAN
TBC1 domain family member 5 - Homo sapiens (Human)					
Q92609	2	3.1244	K.VSNSLINFGR.K	2	TBCD5_HUMAN
Q92609	2	3.4493	R.GQGQSVQM#SGAIK.Q	1	TBCD5_HUMAN
Q92609	2	4.7107	R.GQGQSVQMSGAIK.Q	2	TBCD5_HUMAN
Q92609	2	3.5205	R.GSFSGQAQPLR.T	5	TBCD5_HUMAN
Q92609	3	3.9195	R.KLISPAM#APGSAGGPVPGGNSSSSSSVVIP	1	TBCD5_HUMAN
Q92609	3	4.1952	R.TSAEAPSHHLQQQQQQQR.L	2	TBCD5_HUMAN
T-box transcription factor TBX2 - Homo sapiens (Human)					
Q13207	2	2.9936	R.LVSGLESQR.A	2	TBX2_HUMAN
T-cell surface antigen CD2 precursor - Homo sapiens (Human)					
P06729	2	2.9653	R.VQPKPPHGAAENSLSPSSN.-	1	CD2_HUMAN
P06729	3	5.1542	K.RPPAPSGTQVHQQK.G	3	CD2_HUMAN
T-complex protein 1 subunit alpha - Homo sapiens (Human)					
P17987	2	4.4343	R.SQNVMAAASIANIVK.S	4	TCPA_HUMAN
T-complex protein 1 subunit delta - Homo sapiens (Human)					
P50991	1	2.5976	R.FSNISAAK.A	4	TCPD_HUMAN
TDP43 - Homo sapiens (Human)					
A4GUK4	2	4.8251	R.FGGNPGGFGNQGGFGNSR.G	2	A4GUK4_HUMA
Telomerase-binding protein EST1A - Homo sapiens (Human)					
Q86US8	2	3.7779	R.GILATLAPQAGSR.E	1	EST1A_HUMAN
Telomeric repeat-binding factor 2 - Homo sapiens (Human)					
Q15554	2	3.3549	K.YGEGNWAAISK.N	2	TERF2_HUMAN
Telomeric repeat-binding factor 2-interacting protein 1 - Homo sapiens (Human)					
Q9NYB0	2	3.1089	K.FGAQNVAR.R	4	TE2IP_HUMAN
Q9NYB0	2	4.2083	R.SPSSVTGNALWK.A	6	TE2IP_HUMAN
Q9NYB0	3	4.4474	R.LGPASAADTGSEAKPGALAEGAAEPEPQR.H	2	TE2IP_HUMAN
Q9NYB0	3	4.2286	R.KAEEDPEAADSGEPQNKR.T	3	TE2IP_HUMAN
Q9NYB0	2	2.9105	R.ADGYPIWSR.Q	1	TE2IP_HUMAN
Q9NYB0	2	4.9813	K.FNLDLSTVTQAFLK.N	3	TE2IP_HUMAN
Q9NYB0	2	3.6528	K.AEEDPEAADSGEPQNK.R	1	TE2IP_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9NYB0	2	3.1404	K.NSGELEATSAFLASGQR.A	1	TE2IP_HUMAN
Tenascin precursor - Homo sapiens (Human)					
P24821	2	3.7809	R.GLEPGQEYNVLLTAEK.G	4	TENA_HUMAN
P24821	2	3.1783	R.RPETSYR.Q	3	TENA_HUMAN
P24821	2	3.2424	R.RGDMSSNPAK.E	10	TENA_HUMAN
P24821	2	4.5998	R.VPGDQTSTIIQELEPGVEYFIR.V	3	TENA_HUMAN
P24821	2	4.1311	K.YAPISGGDHAEVDVPK.S	3	TENA_HUMAN
P24821	2	3.0821	K.ETFTTGLDAPR.N	1	TENA_HUMAN
P24821	2	3.5454	K.AATPYTVSIYGVIQGYR.T	2	TENA_HUMAN
P24821	2	4.155	R.LEELENLVSSLR.E	2	TENA_HUMAN
P24821	2	4.136	R.NMNKEDEGEITK.S	3	TENA_HUMAN
Teneurin-3 - Homo sapiens (Human)					
Q9P273	2	3.0563	R.QDGMFDLVANGGASLTLVFER.S	1	TEN3_HUMAN
Tensin-1 - Homo sapiens (Human)					
Q9HBL0	2	2.7369	R.SYVESVAR.T	1	TENS1_HUMAN
Q9HBL0	3	4.0545	R.HPGAHQGNLASGLHSNAIASPGSPSLGR.H	1	TENS1_HUMAN
Q9HBL0	2	2.8897	R.VAGVQAR.E	4	TENS1_HUMAN
Q9HBL0	2	3.3156	R.LLSGFGLER.E	2	TENS1_HUMAN
Q9HBL0	2	3.7477	R.HVAYGGYSTPEDR.R	2	TENS1_HUMAN
Q9HBL0	2	5.4605	R.RAASDGQYENQSPEATSPR.S	3	TENS1_HUMAN
Q9HBL0	2	3.3775	R.SGSLGQPSPSAQR.N	3	TENS1_HUMAN
Q9HBL0	2	4.6059	R.SGYIPSGHSLGTPEPAPR.A	5	TENS1_HUMAN
Q9HBL0	2	3.3804	R.SRPAGGSAVPSSGR.H	6	TENS1_HUMAN
Q9HBL0	3	5.032	R.TPEEEPLNLEGLVAHR.V	5	TENS1_HUMAN
Q9HBL0	2	3.6088	R.TPTQPLLESGFR.S	7	TENS1_HUMAN
Q9HBL0	2	3.7151	R.TVGTNTPPSPGFGR.R	4	TENS1_HUMAN
Q9HBL0	2	5.766	R.HPAGVYQVSGLHNK.V	5	TENS1_HUMAN
Q9HBL0	2	4.0433	R.SQSFSEAEPQLPPAPVR.G	4	TENS1_HUMAN
Q9HBL0	2	2.7023	K.YSM#PDNSPETR.A	1	TENS1_HUMAN
Q9HBL0	3	4.4448	R.GLNSWQQQQQQQQQPRPPPR.Q	3	TENS1_HUMAN
Q9HBL0	2	2.7748	R.QGSPTPALPEK.R	1	TENS1_HUMAN
Q9HBL0	3	5.9155	K.EATSDPSRTPEEEPLNLEGLVAHR.V	2	TENS1_HUMAN
Q9HBL0	2	4.6954	K.SFSAPATQAYGHEIPLR.N	3	TENS1_HUMAN
Q9HBL0	2	4.2662	K.SQSVPGAWPGASPLSSQPLSGSSR.Q	2	TENS1_HUMAN
Q9HBL0	2	2.9296	K.VATTPGSPSLGR.H	2	TENS1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9HBL0	2	4.5804	R.AASDGQYENQSPEATSPR.S	4	TENS1_HUMAN
Q9HBL0	3	5.0611	R.AASQQEIEQSIETLNMLMLDLEPASAAAPLHK.	1	TENS1_HUMAN
Q9HBL0	3	3.8835	R.AGPAHAGHTAPM#RPSYSAQEGLAGYQR.E	1	TENS1_HUMAN
Q9HBL0	3	4.5316	R.AGPAHAGHTAPMRPSYSAQEGLAGYQR.E	3	TENS1_HUMAN
Q9HBL0	2	4.3292	R.AGSLPNYATINGK.V	2	TENS1_HUMAN
Q9HBL0	2	4.0502	R.AQFSVAGVHTVPGSPQAR.H	3	TENS1_HUMAN
Q9HBL0	2	2.8253	R.EKQPAEPPAPLR.R	2	TENS1_HUMAN
Q9HBL0	3	5.0423	K.TDKTDEPVPGASSATAALSPQEKR.E	2	TENS1_HUMAN
Tensin-3 - Homo sapiens (Human)					
Q68CZ2	3	4.1304	R.QQQM#VVAHQYSFAPDGEAR.L	1	TENS3_HUMAN
Q68CZ2	2	3.0155	R.LPDTGEGPSR.A	2	TENS3_HUMAN
Q68CZ2	3	3.8097	R.QQQMVVAHQYSFAPDGEAR.L	1	TENS3_HUMAN
Q68CZ2	2	3.2434	K.ETM#TPGYPQDLDIIDGR.I	1	TENS3_HUMAN
Q68CZ2	2	3.1811	K.AELDQLLSGFGLEDPGSSLK.E	1	TENS3_HUMAN
Q68CZ2	2	3.9925	R.GVGSGPHPPDTQQPSPSK.A	2	TENS3_HUMAN
Tensin-like C1 domain-containing phosphatase - Homo sapiens (Human)					
Q63HR2	2	3.0141	K.LALPTAALYGLR.L	2	TENC1_HUMAN
Q63HR2	3	5.2964	R.HLPGPGQQPGPWGPEQASSPAR.G	1	TENC1_HUMAN
Q63HR2	2	2.869	R.SPVPTTLPGLR.H	1	TENC1_HUMAN
Testin - Homo sapiens (Human)					
Q9UGI8	2	3.4047	R.NVMILTNPVAAK.K	2	TES_HUMAN
Testis-expressed sequence 15 protein - Homo sapiens (Human)					
Q9BXT5	1	2.1273	K.IYDTLSK.D	1	TEX15_HUMAN
Testis-specific Y-encoded-like protein 1 - Homo sapiens (Human)					
Q9H0U9	3	3.7983	R.SASELTAGAEAEAEEVKTGK.C	1	TSYL1_HUMAN
Q9H0U9	3	3.8041	K.AGQEEGQPPAEGLAAASVVMAADR.S	1	TSYL1_HUMAN
Q9H0U9	3	5.2694	R.MDPLEAIQLELDTVNAQADR.A	1	TSYL1_HUMAN
Tetratricopeptide repeat protein 1 - Homo sapiens (Human)					
Q99614	2	3.7369	K.AIQLNPSYIR.A	2	TTC1_HUMAN
Q99614	4	4.9454	K.LLRDDEAHLQEDQGEEECFHDCSASFEEEPG	1	TTC1_HUMAN
Tetratricopeptide repeat, ankyrin repeat and coiled-coil-containing protein 1 - Homo sapiens (Hum					
Q9C0D5	2	3.6263	K.SLREPVAQPGLLLQPSK.Q	2	Q9C0D5_HUMA
Q9C0D5	2	3.3585	K.TVSHLYQESISK.Q	1	Q9C0D5_HUMA
Q9C0D5	2	3.7865	R.FQQQSNPPSR.S	2	Q9C0D5_HUMA
Thiamine-triphosphatase - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BU02	2	3.3446	R.LSSMLGVPAQETAPAK.L	1	THTPA_HUMAN
Thioredoxin - Homo sapiens (Human)					
P10599	2	3.8921	K.M#IKPFFHSLSEK.Y	4	THIO_HUMAN
P10599	2	3.8878	K.MIKPFFHSLSEK.Y	4	THIO_HUMAN
Thioredoxin domain-containing protein 1 precursor - Homo sapiens (Human)					
Q9H3N1	2	3.4118	K.VEEEQEADEEDVSEEEAESK.E	1	TXND1_HUMAN
Thioredoxin domain-containing protein 12 precursor - Homo sapiens (Human)					
O95881	3	4.0432	K.YFYVSAEQVVQGMKEAQER.L	1	TXD12_HUMAN
O95881	3	4.2726	R.ILFLDPSGKVHPEIINENGNPSYK.Y	1	TXD12_HUMAN
O95881	2	2.7315	K.YFYVSAEQVVQGM#K.E	1	TXD12_HUMAN
O95881	2	4.266	K.EAAASGLPLMVIIHK.S	4	TXD12_HUMAN
O95881	2	3.2382	K.YFYVSAEQVVQGMK.E	2	TXD12_HUMAN
Thioredoxin domain-containing protein 4 precursor - Homo sapiens (Human)					
Q9BS26	3	4.0482	K.SDPIQEIRDLAEITTLDR.S	1	TXND4_HUMAN
Q9BS26	3	4.9402	R.HMYVFGDFKDVLIPGK.L	2	TXND4_HUMAN
Thioredoxin domain-containing protein 6 - Homo sapiens (Human)					
Q86XW9	1	2.2724	K.AGEEAFEK.L	2	TXND6_HUMAN
Thioredoxin, mitochondrial precursor - Homo sapiens (Human)					
Q99757	3	4.8131	K.NGDVVDKFVGIKDEDQLEAFLK.K	2	THIOM_HUMAN
Thioredoxin-like protein 5 - Homo sapiens (Human)					
Q9BRA2	2	3.5401	K.NLKVTAVPTLLK.Y	2	TXNL5_HUMAN
Q9BRA2	2	3.3374	K.TIFAYFTGSK.D	3	TXNL5_HUMAN
Thioredoxin-like selenoprotein M precursor - Homo sapiens (Human)					
Q8WWX9	2	2.7666	K.AFVTQDIPFYHNLVMK.H	1	SELM_HUMAN
THO complex subunit 4 - Homo sapiens (Human)					
Q86V81	2	4.2235	R.SLGTADVHFER.K	6	THOC4_HUMAN
Q86V81	2	3.2251	R.PMNIQLVTSQIDAQR.R	1	THOC4_HUMAN
Q86V81	2	3.3778	R.GAGGFGGGGGTR.R	4	THOC4_HUMAN
Q86V81	3	5.2282	K.WQHDLFDSGFGGGAGVETGGK.L	3	THOC4_HUMAN
Q86V81	2	5.4065	K.QQLSAEELDAQLDAYNAR.M	4	THOC4_HUMAN
Q86V81	2	2.7593	K.KAAVHYDR.S	1	THOC4_HUMAN
Q86V81	3	5.0388	K.QLPDKWQHDLFDSGFGGGAGVETGGK.L	4	THOC4_HUMAN
Thrombomodulin precursor - Homo sapiens (Human)					
P07204	2	3.0232	R.SSVAADVISLLLNGDGGVGR.R	1	TRBM_HUMAN
Thrombospondin-1 precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07996	2	4.3285	R.FTGSQPFGQGVEHATANK.Q	3	TSP1_HUMAN
P07996	2	2.7392	R.TNYIGHK.T	1	TSP1_HUMAN
P07996	2	4.6897	R.KVTEENKELANELR.R	3	TSP1_HUMAN
P07996	2	2.887	K.GTSQNDPNWVVR.H	2	TSP1_HUMAN
P07996	2	2.9915	K.GGVNDNFQGVLQNVR.F	2	TSP1_HUMAN
P07996	2	3.7019	R.TIVTTLQDSIR.K	4	TSP1_HUMAN
P07996	3	3.8242	R.DTDMDGVGDQCDNCPLEHNPDQLDSDSDR.I	1	TSP1_HUMAN
Thymocyte nuclear protein 1 - Homo sapiens (Human)					
Q9P016	2	2.8972	K.TENSGEALAK.V	1	THYN1_HUMAN
Q9P016	3	3.8485	K.TENSGEALAKVEDSNPQK.T	1	THYN1_HUMAN
Q9P016	3	4.0496	R.TKTENSGEALAK.V	4	THYN1_HUMAN
Thymosin beta-10 - Homo sapiens (Human)					
P63313	2	4.0279	K.NTLPTKETIEQEKR.S	13	TYB10_HUMAN
P63313	2	4.0896	K.TETQEKNTLPTK.E	4	TYB10_HUMAN
P63313	2	4.0701	K.NTLPTKETIEQEK.R	10	TYB10_HUMAN
P63313	1	2.4055	K.NTLPTK.E	7	TYB10_HUMAN
P63313	2	4.7269	K.KTETQEKNTLPTK.E	4	TYB10_HUMAN
P63313	2	5.1986	K.TETQEKNTLPTKETIEQEK.R	3	TYB10_HUMAN
Thyroid hormone receptor-associated protein 3 - Homo sapiens (Human)					
Q9Y2W1	2	3.0496	K.SDSFAPK.T	2	TR150_HUMAN
Q9Y2W1	3	5.1046	R.KTEELEEESFPER.S	5	TR150_HUMAN
Q9Y2W1	2	3.2107	R.ASAVSELSPR.E	2	TR150_HUMAN
Q9Y2W1	2	3.1115	K.EESAASGGAAYTKR.Y	2	TR150_HUMAN
Q9Y2W1	2	3.5392	K.WEGLVYAPPGK.E	1	TR150_HUMAN
Q9Y2W1	2	3.571	R.NREEEWDPEYTPK.S	4	TR150_HUMAN
Q9Y2W1	2	2.8352	K.TDSEKPFR.G	2	TR150_HUMAN
Q9Y2W1	3	5.2379	K.SPPSTGSTYGSSQKEESAASGGAAYTKR.Y	1	TR150_HUMAN
Q9Y2W1	3	4.4033	K.SPPSTGSTYGSSQKEESAASGGAAYTK.R	1	TR150_HUMAN
Q9Y2W1	2	3.4196	K.SPLQSVVVR.R	4	TR150_HUMAN
Q9Y2W1	2	3.0088	K.MKSDSFAPK.T	4	TR150_HUMAN
Q9Y2W1	2	3.0538	K.M#KSDSFAPK.T	2	TR150_HUMAN
Q9Y2W1	2	2.9928	K.GSFSDTGLGDGK.M	5	TR150_HUMAN
Q9Y2W1	2	3.125	K.GRKESEFDDEPK.F	1	TR150_HUMAN
Q9Y2W1	3	4.3883	K.FSGEEGEIEDDESGTENREEKDNIQPTTE.-	2	TR150_HUMAN
Q9Y2W1	2	3.3783	K.EQTFSGGTSQDTK.A	2	TR150_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y2W1	2	3.022	K.EESAASGGAAYTK.R	1	TR150_HUMAN
Q9Y2W1	3	4.9453	K.ASESSKPWPDATYGTGSASR.A	6	TR150_HUMAN
Q9Y2W1	2	3.2451	K.ESEFDDEPKFMSK.V	1	TR150_HUMAN
Thyroid receptor-interacting protein 11 - Homo sapiens (Human)					
Q15643	2	2.957	K.IIEDQNQSK.M	3	TRIPB_HUMAN
Q15643	2	3.8482	R.ILAQSASVEEVFR.L	2	TRIPB_HUMAN
Thyroid transcription factor 1-associated protein 26 - Homo sapiens (Human)					
Q9P031	2	3.536	K.AQEEYEQIQAK.R	2	TAP26_HUMAN
Q9P031	2	3.0225	K.HLYLAEEER.H	1	TAP26_HUMAN
Tight junction protein ZO-1 - Homo sapiens (Human)					
Q07157	2	3.8488	R.LEEPTPAPSTSYSPQADSLR.T	2	ZO1_HUMAN
Q07157	4	5.716	R.SFENKPPAHIAASHLSEPAKPAHSQNQSNFS	1	ZO1_HUMAN
Tight junction protein ZO-2 - Homo sapiens (Human)					
Q9UDY2	2	4.3832	R.MQELQEAQNAR.I	4	ZO2_HUMAN
Q9UDY2	2	3.0787	R.RAASSDQLR.D	1	ZO2_HUMAN
Q9UDY2	3	3.9578	R.SILKPSTPIPPQEGEEVGESSEEQDNAPK.S	1	ZO2_HUMAN
Q9UDY2	2	4.315	K.SNPSAVAGNETPGASTK.G	4	ZO2_HUMAN
Q9UDY2	3	4.3943	R.IEIAQKHPDIYAVPIK.T	2	ZO2_HUMAN
Q9UDY2	2	4.3584	R.M#QELQEAQNAR.I	2	ZO2_HUMAN
Tight junction-associated protein 1 - Homo sapiens (Human)					
Q5JTD0	2	3.8353	R.GTEEGPGTSHTEGR.A	2	TJAP1_HUMAN
Q5JTD0	3	4.6961	R.KHLHSGQEAASPGPAPSLAPGAVVPTSVIAR.	2	TJAP1_HUMAN
Q5JTD0	2	3.9198	R.LEIPGSRLEQEEPLTDAER.M	2	TJAP1_HUMAN
Q5JTD0	2	3.422	R.QAISLSLVEEGSER.A	1	TJAP1_HUMAN
Q5JTD0	3	4.5383	R.RVIEFSEDKVR.I	2	TJAP1_HUMAN
TOM1-like protein 2 - Homo sapiens (Human)					
Q6ZVM7	2	3.0944	R.DGFDMFAQTR.G	1	TM1L2_HUMAN
Q6ZVM7	3	4.9893	R.KTVTYEDPQAVGGLASALDNRK.Q	2	TM1L2_HUMAN
Q6ZVM7	2	3.6166	R.DGFDM#FAQTR.G	3	TM1L2_HUMAN
Q6ZVM7	3	4.9761	K.TVTYEDPQAVGGLASALDNR.K	4	TM1L2_HUMAN
Torsin family protein C9orf167 - Homo sapiens (Human)					
Q9NXH8	2	3.1411	R.GQPSLEPAAAAPR.A	2	CI167_HUMAN
Torsin-1A-interacting protein 1 - Homo sapiens (Human)					
Q5JTV8	3	5.1439	K.VNFSEEGETEEDDQDSSHSSVTTVK.A	1	TOIP1_HUMAN
Q5JTV8	3	3.9671	R.LQQQHSEQPPLQPSPVM#TR.R	1	TOIP1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q5JTV8	2	3.5843	R.SIQEAPVSEDLVIR.L	1	TOIP1_HUMAN
Torsin-1A-interacting protein 2 - Homo sapiens (Human)					
Q8NFQ8	3	5.1253	K.EASDGTGASQEPPTTDSQEAQSPGHSSAGQ	2	TOIP2_HUMAN
TRAF family member-associated NF-kappa-B activator - Homo sapiens (Human)					
Q92844	2	2.851	R.RQEVSSPR.K	2	TANK_HUMAN
TRAF2 and NCK-interacting protein kinase - Homo sapiens (Human)					
Q9UKE5	3	4.207	R.VSTHSQEMDSGTEYGMGSSTK.A	1	TNIK_HUMAN
TRAF-type zinc finger domain-containing protein 1 - Homo sapiens (Human)					
O14545	2	3.8226	K.LSNSDSQDIQGR.N	3	TRAD1_HUMAN
Transcription elongation factor A protein 1 - Homo sapiens (Human)					
P23193	3	4.1411	R.MTAEEMASDELKEMR.K	2	TCEA1_HUMAN
P23193	2	3.5176	K.NAAGALDLLK.E	1	TCEA1_HUMAN
P23193	2	3.911	K.LLDGPSTEKDLDEK.K	3	TCEA1_HUMAN
P23193	2	3.3521	R.EESTSSGNVSNR.K	5	TCEA1_HUMAN
P23193	2	2.7342	K.NIPM#TLELLQSTR.I	2	TCEA1_HUMAN
Transcription elongation factor A protein 3 - Homo sapiens (Human)					
O75764	2	3.2914	K.MASEIEDHIYQELK.S	1	TCEA3_HUMAN
O75764	2	3.2631	K.MTAEEMASDELRELR.N	1	TCEA3_HUMAN
O75764	2	3.3807	R.NVLSGAISAGLIAK.M	2	TCEA3_HUMAN
Transcription elongation factor A protein-like 3 - Homo sapiens (Human)					
Q969E4	3	4.4628	K.REDEGEPGDEGQLEDEGSQEK.Q	2	TCAL3_HUMAN
Q969E4	2	2.9662	R.EDEGEPGDEGQLEDEGSQEK.Q	2	TCAL3_HUMAN
Transcription elongation factor A protein-like 4 - Homo sapiens (Human)					
Q96EI5	2	3.2119	R.EGESEMEGGSER.E	3	TCAL4_HUMAN
Q96EI5	3	4.2316	R.EGKPEIEGKPESEGEPGSETR.A	1	TCAL4_HUMAN
Q96EI5	2	2.7906	R.EGESEM#EGGSER.E	1	TCAL4_HUMAN
Transcription elongation factor A protein-like 8 - Homo sapiens (Human)					
Q8IYN2	3	6.2997	K.AEEDRPLEDVPQEAEGNPQPSEEGVSQEAE	1	TCAL8_HUMAN
Transcription elongation factor B polypeptide 3 - Homo sapiens (Human)					
Q14241	2	2.9255	K.LSTLPITVDILAETGVGK.T	1	ELOA1_HUMAN
Q14241	3	3.8669	K.SEKPAGADLAK.L	1	ELOA1_HUMAN
Q14241	2	2.7039	K.TSATALGDK.G	1	ELOA1_HUMAN
Q14241	3	5.6707	R.DALQKEEEMEGDYQETWK.A	3	ELOA1_HUMAN
Q14241	2	2.7223	R.HLGEPHGK.G	1	ELOA1_HUMAN
Transcription elongation factor SPT5 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O00267	2	3.8238	R.TPMYGSQTPLQDGSR.T	2	SPT5H_HUMAN
O00267	2	3.0946	R.RPGGMTSTYGR.T	1	SPT5H_HUMAN
O00267	2	3.639	R.TPM#YGSQTPLQDGSR.T	1	SPT5H_HUMAN
Transcription elongation factor SPT6 - Homo sapiens (Human)					
Q7KZ85	2	3.2119	K.MAEQWLQEK.E	2	SPT6H_HUMAN
Transcription elongation regulator 1 - Homo sapiens (Human)					
O14776	3	5.6825	R.RGPPPPPTASEPTRR.S	2	TCRG1_HUMAN
O14776	2	3.2715	R.EALFNEFVAAAR.K	2	TCRG1_HUMAN
Transcription factor 20 - Homo sapiens (Human)					
Q9UGU0	2	2.7759	K.QINLTDYPIPR.K	1	TCF20_HUMAN
Q9UGU0	2	2.8861	R.KSSSTAPEMK.Q	1	TCF20_HUMAN
Transcription factor 4 - Homo sapiens (Human)					
P15884	2	3.532	R.GSGAAGSSQTGDALGK.A	2	ITF2_HUMAN
Transcription factor A, mitochondrial precursor - Homo sapiens (Human)					
Q00059	3	4.0993	R.FKEQLTPSQIM#SLEK.E	2	TFAM_HUMAN
Q00059	2	4.0727	R.SAYNVYVAER.F	1	TFAM_HUMAN
Q00059	2	3.8017	R.FQEAKGDSPQEK.L	4	TFAM_HUMAN
Q00059	3	3.7599	R.FKEQLTPSQIMSLEK.E	1	TFAM_HUMAN
Q00059	2	2.8023	K.SWEEQMIEVGRK.D	1	TFAM_HUMAN
Q00059	2	3.1967	K.SWEEQMIEVGR.K	3	TFAM_HUMAN
Q00059	3	3.8376	K.NLSDSEKELYIQHAKEDETR.Y	1	TFAM_HUMAN
Q00059	3	5.507	R.FKEQLTPSQIMSLEKEIMDK.H	2	TFAM_HUMAN
Transcription factor AP-1 - Homo sapiens (Human)					
P05412	2	4.6625	K.AQNSELASTANMLR.E	2	JUN_HUMAN
P05412	2	3.5778	K.NVTDEQEGFAEGFVR.A	2	JUN_HUMAN
Transcription factor BTF3 - Homo sapiens (Human)					
P20290	4	4.7882	K.LGVNNISGIEEVNMFTNQGTVIHFNNPK.V	1	BTF3_HUMAN
P20290	2	3.5125	R.TGAPAQADSR.G	2	BTF3_HUMAN
P20290	2	4.0784	R.GQEPQMKETIMNQEK.L	1	BTF3_HUMAN
P20290	2	3.6051	R.GQEPQMKETIM#NQEK.L	1	BTF3_HUMAN
P20290	3	4.151	R.GQEPQM#KETIMNQEK.L	2	BTF3_HUMAN
P20290	3	4.1099	K.QLTEMLPSILNQLGADSLTSLR.R	1	BTF3_HUMAN
P20290	2	5.6521	K.VQASLAANTFTITGHAETK.Q	2	BTF3_HUMAN
P20290	3	5.013	K.LGVNNISGIEEVNM#FTNQGTVIHFNNPK.V	1	BTF3_HUMAN
P20290	2	3.2275	R.GGCPGGEATLSQPPPR.G	1	BTF3_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Transcription factor BTF3 homolog 2 - Homo sapiens (Human)					
Q13891	2	2.8295	-.M#KETIM#NQEK.L	1	BT3L2_HUMAN
Q13891	2	3.5621	-.M#KETIMNQEK.L	1	BT3L2_HUMAN
Q13891	2	3.2376	-.MKETIM#NQEK.L	1	BT3L2_HUMAN
Transcription factor E3 - Homo sapiens (Human)					
P19532	2	2.7228	R.TPAMSSSSSSR.V	1	TFE3_HUMAN
Transcription factor EB - Homo sapiens (Human)					
P19484	2	3.3682	K.SRELENHSR.R	4	TFEB_HUMAN
P19484	3	4.6454	R.VHGLPTTSPSGMNMAELAQQVVK.Q	2	TFEB_HUMAN
Transcription factor HES-3 - Homo sapiens (Human)					
Q5TGS1	2	3.0218	R.KLEKADILELSVK.Y	1	HES3_HUMAN
Transcription factor jun-B - Homo sapiens (Human)					
P17275	2	4.4565	R.GGGSGGGAGGAGGGVTEEQEGFADGFVK.A	2	JUNB_HUMAN
Transcription factor jun-D - Homo sapiens (Human)					
P17535	2	4.1196	K.VAASEEQEFAEGFVK.A	2	JUND_HUMAN
P17535	2	4.4036	K.SQNTELASTASLLR.E	2	JUND_HUMAN
Transcription factor MafF - Homo sapiens (Human)					
Q9ULX9	2	4.6253	R.GPATLVAPASVITIVK.S	4	MAFF_HUMAN
Transcription factor PU.1 - Homo sapiens (Human)					
P17947	2	2.8309	K.LTYQFSGEVLGRGGLAER.R	1	SPI1_HUMAN
Transcription factor SOX-17 - Homo sapiens (Human)					
Q9H6I2	3	4.3103	R.LGPEPAGPSIPGLLAPPSALHVYYGAMGSPG	1	SOX17_HUMAN
Transcription factor Sp3 - Homo sapiens (Human)					
Q02447	3	5.7026	R.IKEEEPDPEEWQLSGDSTLNTNDLTHLR.V	1	SP3_HUMAN
Transcription initiation factor IIE subunit beta - Homo sapiens (Human)					
P29084	2	2.9436	K.DYSDITSSK.-	1	T2EB_HUMAN
Transcription initiation factor IIF subunit alpha - Homo sapiens (Human)					
P35269	3	5.3829	R.GNSRPGTPSAEGGSTSSTLR.A	1	T2FA_HUMAN
P35269	2	4.0508	R.TLTAEEAEEEWER.R	1	T2FA_HUMAN
P35269	3	5.0703	R.LKDQDQDEDEEEKEK.R	2	T2FA_HUMAN
P35269	2	2.8133	R.LDTGPQSLSGK.S	1	T2FA_HUMAN
P35269	2	3.8659	K.VNFATWNQAR.L	3	T2FA_HUMAN
P35269	2	3.2656	K.VLNHFSIMQQR.R	1	T2FA_HUMAN
P35269	3	4.5105	K.KTGLSSEQTVNVLAQILK.R	2	T2FA_HUMAN
P35269	2	3.1152	K.TTPNSGDVQVTEDAVR.R	1	T2FA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Transcription initiation factor IIF subunit beta - Homo sapiens (Human)					
P13984	2	3.106	K.NTWELKPEYR.H	2	T2FB_HUMAN
P13984	2	3.2164	K.RLQIEESSKPVR.L	1	T2FB_HUMAN
P13984	2	2.7159	R.LSQQLDK.V	1	T2FB_HUMAN
Transcription initiation factor TFIID subunit 10 - Homo sapiens (Human)					
Q12962	2	4.8712	K.ASPAGTAGGPGAGAAAGGTGPLAAR.A	2	TAF10_HUMAN
Transcription initiation factor TFIID subunit 7 - Homo sapiens (Human)					
Q15545	3	4.3931	K.SKDDAPHELESQFILR.L	1	TAF7_HUMAN
Q15545	3	6.0464	R.EKEQLSSLQEELESLLEK.-	1	TAF7_HUMAN
Transcription initiation factor TFIID subunit 9 - Homo sapiens (Human)					
Q16594	3	4.017	R.LSVGSVTSRPSTPTLGTPTPQTM#SVSTK.V	1	TAF9_HUMAN
Transcription intermediary factor 1-beta - Homo sapiens (Human)					
Q13263	2	3.3813	K.DIVENYFMR.D	1	TIF1B_HUMAN
Transcriptional activator protein Pur-alpha - Homo sapiens (Human)					
Q00577	2	2.8652	K.IAEVGAGGNK.S	1	PURA_HUMAN
Q00577	2	2.7351	R.FFFDVGSNK.Y	1	PURA_HUMAN
Transcriptional adapter 3-like - Homo sapiens (Human)					
O75528	2	2.7782	K.ESGADGASTSPR.N	1	TAD3L_HUMAN
Transcriptional regulator ERG - Homo sapiens (Human)					
P11308	2	3.6344	R.VPQQDWLSQPPAR.V	1	ERG_HUMAN
P11308	2	3.6229	K.TEM#TASSSSDYGQTSK.M	3	ERG_HUMAN
Transcriptional repressor CTCF - Homo sapiens (Human)					
P49711	3	3.8756	R.YTEEGKDVDVSVYDFEEEQQEGLLSEVNAEK	1	CTCF_HUMAN
Transcriptional repressor p66 alpha - Homo sapiens (Human)					
Q86YP4	2	4.2871	R.ALERDPTEDDVESK.K	2	P66A_HUMAN
Q86YP4	2	3.3207	R.ATEATAM#AM#GR.G	1	P66A_HUMAN
Q86YP4	2	3.1819	R.GEGLVGDGPVDMR.T	1	P66A_HUMAN
Q86YP4	2	3.9485	R.GLLASDLNTDGDMR.V	1	P66A_HUMAN
Transcriptional repressor p66 beta - Homo sapiens (Human)					
Q8WXI9	3	3.9985	K.LPSRPGAQGVEPQNLR.T	1	P66B_HUMAN
Q8WXI9	2	3.2689	R.VIAPNPAQLQGQR.G	2	P66B_HUMAN
Transcriptional repressor protein YY1 - Homo sapiens (Human)					
P25490	1	2.8009	K.SHILTHAK.A	1	TYY1_HUMAN
P25490	2	3.0709	R.KIKEDDAPR.T	4	TYY1_HUMAN
P25490	2	4.3198	K.SYLSGGAGAAGGGGADPGNK.K	2	TYY1_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
P25490	3	4.3028	K.KSYLSGGAGAAGGGGADPGNKK.W	2	TYY1_HUMAN
P25490	2	4.1255	K.SYLSGGAGAAGGGGADPGNKK.W	3	TYY1_HUMAN
Transducin-like enhancer protein 1 - Homo sapiens (Human)					
Q04724	2	3.5233	K.DASSSPASTASSASSTSLK.S	1	TLE1_HUMAN
Transducin-like enhancer protein 3 - Homo sapiens (Human)					
Q04726	2	4.6382	K.DAPTSPASVASSSSTPSSK.T	3	TLE3_HUMAN
Q04726	2	3.4277	R.NDAPTPGTSTTPGLR.S	2	TLE3_HUMAN
Transformer-2 protein homolog - Homo sapiens (Human)					
Q13595	2	3.4154	R.YGPLSGVNVVYDQR.T	2	TRA2A_HUMAN
Transforming acidic coiled-coil-containing protein 1 - Homo sapiens (Human)					
O75410	3	7.0383	K.SPPDLKETPGTLSSDTNDSGVELGEESR.S	2	TACC1_HUMAN
O75410	2	3.5472	K.AIGGEFSDTNAAVEGTPLPK.A	2	TACC1_HUMAN
O75410	2	4.1136	R.GGAAGEDEAGGPEGDPEEEDSQAETK.S	3	TACC1_HUMAN
O75410	2	2.751	R.KLGSTLTPK.I	1	TACC1_HUMAN
O75410	2	4.6517	K.ETPGTLSSDTNDSGVELGEESR.S	2	TACC1_HUMAN
O75410	2	5.7422	K.ASYHFSPEELDENTSPLLGDAR.F	1	TACC1_HUMAN
O75410	3	4.3493	K.NFREEPEHDFSK.I	2	TACC1_HUMAN
O75410	2	3.4533	K.DSTDISAVLGTK.A	2	TACC1_HUMAN
Transforming acidic coiled-coil-containing protein 3 - Homo sapiens (Human)					
Q9Y6A5	3	4.6403	K.APQEVEEDDGRSGAGEDPPM#PASR.G	1	TACC3_HUMAN
Q9Y6A5	3	4.7475	K.APQEVEEDDGRSGAGEDPPMPASR.G	1	TACC3_HUMAN
Q9Y6A5	4	4.7399	R.DSPGRPVPVATETSSMHGANETPSGRPR.E	1	TACC3_HUMAN
Transforming growth factor beta-1-induced transcript 1 protein - Homo sapiens (Human)					
O43294	3	4.8303	K.ASATSATLELDRLMASLSDFR.V	2	TGFI1_HUMAN
O43294	3	5.2324	R.VQNHLPASGPTQPPVVSSTNEGSPSPPEPT	2	TGFI1_HUMAN
O43294	2	2.7108	R.RGVPTQAK.G	1	TGFI1_HUMAN
O43294	2	5.6843	K.GSLDTMLGLLQSDLSR.R	10	TGFI1_HUMAN
O43294	2	4.1516	K.VASGEQKEDQSEDK.K	3	TGFI1_HUMAN
Transforming growth factor-beta-induced protein ig-h3 precursor - Homo sapiens (Human)					
Q15582	2	4.5781	R.GDELADSALEIFK.Q	3	BGH3_HUMAN
Transgelin - Homo sapiens (Human)					
Q01995	2	2.7037	K.HVIGLQM#GSNR.G	1	TAGL_HUMAN
Q01995	2	3.053	R.GASQAGMTGYGRPR.Q	1	TAGL_HUMAN
Q01995	2	3.5277	R.GASQAGMTGYGR.P	2	TAGL_HUMAN
Q01995	2	3.8224	K.KYDEELEER.L	3	TAGL_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q01995	2	3.4222	R.GASQAGM#TGYGR.P	1	TAGL_HUMAN
Q01995	2	3.29	R.EFTESQLQEGK.H	5	TAGL_HUMAN
Transgelin-2 - Homo sapiens (Human)					
P37802	2	4.5822	R.GASQAGM#TGYGM#PR.Q	3	TAGL2_HUMAN
P37802	2	3.8573	R.YGINTTDIFQTVDLWEGK.N	1	TAGL2_HUMAN
P37802	1	4.2771	R.TLMNLGGLAVAR.D	5	TAGL2_HUMAN
P37802	2	3.0103	R.TLM#NLGGLAVAR.D	1	TAGL2_HUMAN
P37802	2	3.9846	R.NFSDNQLQEGK.N	4	TAGL2_HUMAN
P37802	2	3.0957	R.GASQAGMTGYGM#PR.Q	2	TAGL2_HUMAN
P37802	2	3.8165	R.DDGLFSGDPNWFPKK.S	3	TAGL2_HUMAN
P37802	2	4.0579	R.DDGLFSGDPNWFPK.K	2	TAGL2_HUMAN
P37802	2	3.2591	K.QM#EQISQFLQAAER.Y	2	TAGL2_HUMAN
P37802	2	4.1534	K.NVIGLQMGTNR.G	8	TAGL2_HUMAN
P37802	2	3.8337	K.NVIGLQM#GTNR.G	3	TAGL2_HUMAN
P37802	2	4.1042	R.GASQAGMTGYGMPR.Q	4	TAGL2_HUMAN
Trans-Golgi network integral membrane protein 2 precursor - Homo sapiens (Human)					
O43493	2	3.9202	K.SGAEDQTPKDVPNK.S	2	TGON2_HUMAN
O43493	2	4.3446	K.SGAEEQGPIDGPSK.S	2	TGON2_HUMAN
O43493	2	3.9433	K.SSAEAQTPEDTPNK.S	3	TGON2_HUMAN
O43493	2	2.7465	K.DVPNKSGADGQTPK.D	2	TGON2_HUMAN
O43493	2	4.1916	K.DGSNKSGAEEQGPIDGPSK.S	2	TGON2_HUMAN
O43493	2	3.3781	K.ADTNQLADKGK.L	2	TGON2_HUMAN
O43493	2	4.0751	K.SGAEEQTSKDSPNK.V	2	TGON2_HUMAN
Transient receptor potential cation channel subfamily V member 4 - Homo sapiens (Human)					
Q9HBA0	2	3.8952	K.GVPNPIDLLESTLYESSVVPGPK.K	2	TRPV4_HUMAN
Q9HBA0	2	2.9328	K.NSNPDEVVVPLDSM#GNPR.C	1	TRPV4_HUMAN
Transitional endoplasmic reticulum ATPase - Homo sapiens (Human)					
P55072	2	3.2474	R.GGNIGDGGGAADR.V	1	TERA_HUMAN
P55072	2	3.7776	R.WALSQSNPSALR.E	7	TERA_HUMAN
P55072	3	4.7546	R.LIVDEAINEDNSVVSLSQPK.M	1	TERA_HUMAN
P55072	3	5.2952	R.KYEMFAQTLQQSR.G	6	TERA_HUMAN
P55072	2	3.2473	K.YEMFAQTLQQSR.G	1	TERA_HUMAN
P55072	2	2.9607	K.NAPAIIFIDELDAIAPK.R	1	TERA_HUMAN
P55072	2	3.2027	K.MDELQLFR.G	1	TERA_HUMAN
P55072	2	3.606	R.LDQLIYIPLPDEK.S	2	TERA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Translocated promoter region - Homo sapiens (Human)					
Q5SWY0	3	5.5495	R.VELLERELQELQDSLNAER.E	1	Q5SWY0_HUMA
Translocation protein SEC62 - Homo sapiens (Human)					
Q99442	2	3.6957	R.IQEVGEPSKEEK.A	2	SEC62_HUMAN
Transmembrane and coiled-coil domains protein 3 - Homo sapiens (Human)					
Q9ULS5	2	4.0007	K.LVNNADKQQAGR.I	2	TMCC3_HUMAN
Q9ULS5	2	4.1822	K.SGM#PGVSLTPPVFVFNK.S	1	TMCC3_HUMAN
Q9ULS5	2	2.8415	K.SGMPGVSLTPPVFVFNK.S	1	TMCC3_HUMAN
Transmembrane emp24 domain-containing protein 10 precursor - Homo sapiens (Human)					
P49755	2	3.4539	R.IPDQLVILDMK.H	1	TMEDA_HUMAN
Transmembrane glycoprotein NMB precursor - Homo sapiens (Human)					
Q14956	2	4.9204	K.DVYVVTDQIPVFVTM#FQK.N	3	GPNMB_HUMAN
Transmembrane protein 126B - Homo sapiens (Human)					
Q8IUX1	2	2.8274	R.DSGVVPVGTEEAPK.V	1	T126B_HUMAN
Transmembrane protein 131 - Homo sapiens (Human)					
Q92545	2	3.6833	K.LSLQTLNADIFLK.Q	2	TM131_HUMAN
Q92545	2	3.4411	R.TSAQAASSQSANK.T	1	TM131_HUMAN
Transmembrane protein 132E precursor - Homo sapiens (Human)					
Q6IEE7	2	3.4701	R.DQAEDPASSPTSK.R	3	T132E_HUMAN
Transmembrane protein 81 precursor - Homo sapiens (Human)					
Q6P7N7	1	2.3566	R.C*DVQLVK.N	2	TMM81_HUMAN
Transthyretin precursor - Homo sapiens (Human)					
P02766	3	4.2695	K.TSESGELHGLTTEEEFVEGIYKVEIDTK.S	3	TTHY_HUMAN
P02766	2	4.2418	R.YTIAALLSPYSYSTTAVVTNPKE.-	8	TTHY_HUMAN
P02766	2	4.4002	R.YTIAALLSPYSYSTTAVVTNPK.E	6	TTHY_HUMAN
P02766	3	5.8443	R.RYTIAALLSPYSYSTTAVVTNPKE.-	6	TTHY_HUMAN
P02766	2	5.0226	R.RYTIAALLSPYSYSTTAVVTNPK.E	1	TTHY_HUMAN
P02766	2	4.62	R.KAADDTWEPFASGK.T	4	TTHY_HUMAN
P02766	2	4.0128	R.GSPAINVAVHVFRK.A	2	TTHY_HUMAN
P02766	2	4.188	R.GSPAINVAVHVFR.K	4	TTHY_HUMAN
P02766	3	5.5826	K.VLDAVRGSPAINVAVHVFR.K	2	TTHY_HUMAN
P02766	3	6.9259	K.TSESGELHGLTTEEEFVEGIYK.V	6	TTHY_HUMAN
P02766	3	5.4644	K.ALGISPFHEHAEVVFTANDSGPR.R	7	TTHY_HUMAN
P02766	2	4.1108	K.AADDTWEPFASGK.T	4	TTHY_HUMAN
P02766	3	4.6646	K.VLDAVRGSPAINVAVHVFRK.A	1	TTHY_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Treacle protein - Homo sapiens (Human)					
Q13428	2	3.6941	K.KNPASLPLTQAALK.V	4	TCOF_HUMAN
Q13428	3	5.0813	R.ISDGKKQEGPATQVSK.K	3	TCOF_HUMAN
Q13428	2	3.7846	K.KQEGPATQVSK.K	5	TCOF_HUMAN
Trichohyalin - Homo sapiens (Human)					
Q07283	2	2.9147	R.KFREEELLHQEQGR.K	1	TRHY_HUMAN
Trinucleotide repeat containing 5 - Homo sapiens (Human)					
Q9BT09	2	2.8554	K.ASPLTHSPPDEL.-	4	Q9BT09_HUMAN
Q9BT09	3	4.3262	K.ELGGLEGDPSPEEDEGIQK.A	3	Q9BT09_HUMAN
Trinucleotide repeat-containing 6B protein - Homo sapiens (Human)					
Q9UPQ9	2	3.5558	K.GMPFGMGLGNTSR.S	2	TNC6B_HUMAN
Q9UPQ9	2	3.0091	R.STDAPSQSTGDR.K	3	TNC6B_HUMAN
Q9UPQ9	2	3.4299	K.GGNNDSWMNPLAK.Q	1	TNC6B_HUMAN
Q9UPQ9	2	3.4667	K.DTESSSENTTDNNSASNPGSEK.S	2	TNC6B_HUMAN
Q9UPQ9	2	3.9842	K.DMGTTDSGPYFEK.G	2	TNC6B_HUMAN
Q9UPQ9	2	3.8319	K.DM#GTTDSGPYFEK.G	3	TNC6B_HUMAN
Q9UPQ9	2	3.3609	R.STDAPSQSTGDRK.T	2	TNC6B_HUMAN
TRIO and F-actin-binding protein - Homo sapiens (Human)					
Q9H2D6	3	4.5925	K.WFEATDSRTPEVPAGEGPR.R	1	TARA_HUMAN
Q9H2D6	2	5.0123	R.QALDYVELSPLTQASPQR.A	2	TARA_HUMAN
Q9H2D6	2	2.8578	R.TPEVPAGEGPR.R	1	TARA_HUMAN
Triosephosphate isomerase - Homo sapiens (Human)					
P60174	3	5.592	K.QSLGELIGTLNAAKVPADTEVVCAPPTAYIDF	1	TPIS_HUMAN
P60174	2	5.1433	K.VPADTEVVCAPPTAYIDFAR.Q	1	TPIS_HUMAN
P60174	4	4.7429	K.VAHALAEGLGVIACIGEKLDEREAGITEK.V	1	TPIS_HUMAN
P60174	3	5.1116	K.VAHALAEGLGVIACIGEKLDER.E	1	TPIS_HUMAN
P60174	2	5.8616	R.KQSLGELIGTLNAAK.V	7	TPIS_HUMAN
P60174	2	5.4677	K.QSLGELIGTLNAAK.V	5	TPIS_HUMAN
P60174	3	3.7051	K.ELASQPDVDGFLVGGASLKPEFVDIINAKQ.-	1	TPIS_HUMAN
P60174	3	3.7866	K.ELASQPDVDGFLVGGASLKPEFVDIINAK.Q	1	TPIS_HUMAN
P60174	2	6.2861	K.VAHALAEGLGVIACIGEK.L	8	TPIS_HUMAN
P60174	2	3.7995	K.TATPQQAQEVHEK.L	7	TPIS_HUMAN
TRIP12 protein - Homo sapiens (Human)					
Q14CA3	3	4.2468	K.ALQHTESPSETNKPHSK.S	4	Q14CA3_HUMA
Q14CA3	2	4.2546	R.NTAGAQPQDDSIGGR.S	2	Q14CA3_HUMA

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Tripartite motif-containing protein 25 - Homo sapiens (Human)					
Q14258	3	5.0921	K.KPPPVPALPSKLPTFGAPEQLVDLK.Q	2	TRI25_HUMAN
Q14258	2	4.3959	K.LPTFGAPEQLVDLK.Q	3	TRI25_HUMAN
Tripartite motif-containing protein 65 - Homo sapiens (Human)					
Q6PJ69	2	3.3944	R.DPGPDPGPGPDPAAR.C	1	TRI65_HUMAN
Tripartite motif-containing protein 68 - Homo sapiens (Human)					
Q6AZZ1	1	2.1052	K.EQEEAWK.L	1	TRI68_HUMAN
Tripeptidyl-peptidase 1 precursor - Homo sapiens (Human)					
O14773	2	3.1097	R.PSYQEEAVTK.F	1	TPP1_HUMAN
Triple functional domain protein - Homo sapiens (Human)					
O75962	3	3.9003	K.SADAGSQKDSDDSAATPQDETVEER.G	2	TRIO_HUMAN
tRNA-dihydrouridine synthase 3-like - Homo sapiens (Human)					
Q96G46	3	3.8612	R.LEDGQTADGQTEEAAEPGEQLQTQKR.A	1	DUS3L_HUMAN
Tropomodulin-2 - Homo sapiens (Human)					
Q9NZR1	2	2.8299	K.FDEETANNK.G	1	TMOD2_HUMAN
Q9NZR1	2	2.9535	K.FGYQFTK.Q	2	TMOD2_HUMAN
Q9NZR1	3	4.2277	K.VKPVFEEPPNPTNVEISLQQMK.A	1	TMOD2_HUMAN
Q9NZR1	2	4.0878	R.SNDPVAIAFADMLK.V	1	TMOD2_HUMAN
Tropomodulin-3 - Homo sapiens (Human)					
Q9NYL9	3	5.7276	K.YKDLDEDELLGNLSETELK.Q	2	TMOD3_HUMAN
Q9NYL9	3	3.7751	K.QLETVLDDLDPENALLPAGFR.Q	1	TMOD3_HUMAN
Q9NYL9	3	4.9801	R.TKENDAHLVEVNLNNIK.N	1	TMOD3_HUMAN
Q9NYL9	2	2.9866	R.SNDPVATAFAEMLK.V	1	TMOD3_HUMAN
Q9NYL9	2	4.1386	K.FGYQFTQQGPR.T	6	TMOD3_HUMAN
Q9NYL9	2	3.3055	K.MLEENTNILK.F	2	TMOD3_HUMAN
Q9NYL9	2	3.0793	K.ILPVFDEPPNPTNVEESLKR.T	2	TMOD3_HUMAN
Q9NYL9	2	3.3758	R.QQLGTAVELEMAK.M	2	TMOD3_HUMAN
Tropomyosin 1 alpha variant 6 - Homo sapiens (Human)					
Q1ZYL5	2	3.0568	R.SLQEQADAAEERAGTLQR.E	1	Q1ZYL5_HUMAN
Q1ZYL5	3	5.4717	K.IRSLQEQADAAEER.A	3	Q1ZYL5_HUMAN
Q1ZYL5	2	5.3891	K.LRETAEADVASLNR.R	6	Q1ZYL5_HUMAN
Q1ZYL5	2	2.9985	K.LRETAEADVASLNRR.I	2	Q1ZYL5_HUMAN
Q1ZYL5	2	4.4036	R.ETAEADVASLNR.R	10	Q1ZYL5_HUMAN
Q1ZYL5	2	4.7687	R.KLRETAEADVASLNR.R	7	Q1ZYL5_HUMAN
Q1ZYL5	2	4.6303	R.SLQEQADAAEER.A	9	Q1ZYL5_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Tropomyosin 2 - Homo sapiens (Human)					
Q5TCU3	3	4.6005	K.TIDDLEETLASAKEENVEIHQTLDQTLLELNNL.	1	Q5TCU3_HUMA
Tropomyosin 3 - Homo sapiens (Human)					
Q5VU72	3	4.2369	K.TIDDLEERLYSQLER.N	1	Q5VU72_HUMA
Tropomyosin alpha-4 chain - Homo sapiens (Human)					
P67936	2	5.4951	K.IQALQQQADEAEDR.A	6	TPM4_HUMAN
P67936	2	4.3497	R.KLVILEGELERAEER.A	5	TPM4_HUMAN
P67936	3	5.2425	R.KIQALQQQADEAEDRAQGLQR.E	9	TPM4_HUMAN
P67936	2	4.8558	K.YSEKEDKYEEEIK.L	2	TPM4_HUMAN
P67936	2	6.16	R.KIQALQQQADEAEDR.A	12	TPM4_HUMAN
P67936	3	3.7239	R.REKAEGDVAALNR.R	1	TPM4_HUMAN
P67936	2	3.6085	R.EKAEGDVAALNR.R	2	TPM4_HUMAN
P67936	1	2.1192	R.AQGLQR.E	2	TPM4_HUMAN
P67936	3	6.0048	R.AEVSELKCGDLEEELKNVTNNLK.S	1	TPM4_HUMAN
P67936	3	4.2417	R.AEERAEVSELK.C	4	TPM4_HUMAN
P67936	2	5.2024	K.TIDDLEEKLAQAK.E	9	TPM4_HUMAN
P67936	1	3.1782	K.TIDDLEEK.L	4	TPM4_HUMAN
P67936	1	3.0588	K.SLEAASEK.Y	19	TPM4_HUMAN
P67936	2	3.8135	K.RKIQALQQQADEAEDR.A	4	TPM4_HUMAN
P67936	2	3.608	K.LVILEGELERAEER.A	6	TPM4_HUMAN
P67936	3	4.492	K.IQALQQQADEAEDRAQGLQR.E	6	TPM4_HUMAN
P67936	2	3.1095	K.EENVGLHQTLDQTLNELNCI.-	1	TPM4_HUMAN
P67936	1	4.1357	K.AEGDVAALNR.R	7	TPM4_HUMAN
P67936	2	4.1076	K.LLSDKLKEAETR.A	4	TPM4_HUMAN
P67936	3	5.5484	K.TIDDLEEKLAQAKEENVGLHQTLDQTLNELNC	3	TPM4_HUMAN
Tropomyosin beta chain - Homo sapiens (Human)					
P07951	2	3.6116	K.TIDDLEDEVYAQK.M	2	TPM2_HUMAN
P07951	2	2.7876	K.YKAISEELDNALNDITSL.-	3	TPM2_HUMAN
Tryptophanyl-tRNA synthetase, cytoplasmic - Homo sapiens (Human)					
P23381	2	4.6536	K.AGNASKDEIDSAVK.M	2	SYWC_HUMAN
TSC22 domain family protein 1 - Homo sapiens (Human)					
Q15714	3	4.4431	R.LDNSSSGASVVAIDNKIEQAMDLVK.S	2	T22D1_HUMAN
TSC22 domain family protein 2 - Homo sapiens (Human)					
O75157	2	3.4393	R.HSSTFDQTAER.D	2	T22D2_HUMAN
O75157	2	3.4024	R.NSLLERENALLK.S	2	T22D2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
TSC22 domain family protein 4 - Homo sapiens (Human)					
Q9Y3Q8	2	3.0825	R.NAALEQENGLLR.A	2	T22D4_HUMAN
Q9Y3Q8	2	3.0484	R.VEAEAGGSGAR.T	2	T22D4_HUMAN
TSC22 domain family, member 3 - Homo sapiens (Human)					
Q5JRI9	2	3.2477	R.GSSGENNNPGSPTVSNFR.Q	1	Q5JRI9_HUMAN
TSPY-like 5 - Homo sapiens (Human)					
Q86VY4	3	4.1822	R.AAGDHGQAAARPGPGK.A	2	Q86VY4_HUMA
TTD non-photosensitive 1 protein - Homo sapiens (Human)					
Q8TAP9	3	6.0314	R.SPAGSQQQFGYSPGQQQTHPQGSPR.T	2	TTDN1_HUMAN
TUBB6 protein - Homo sapiens (Human)					
Q2NKY5	2	2.7615	R.AALVDLEPGTMDSVR.S	2	Q2NKY5_HUMA
Tubulin alpha-4A chain - Homo sapiens (Human)					
P68366	2	3.5618	R.AVFVDLEPTVIDEIR.N	4	TBA4A_HUMAN
Tubulin beta chain - Homo sapiens (Human)					
P07437	3	6.3244	K.FWEVISDEHGIDPTGTYHGDSDLQLDR.I	1	TBB5_HUMAN
Tubulin beta-2C chain - Homo sapiens (Human)					
P68371	2	3.79	R.INVYYNEATGGK.Y	4	TBB2C_HUMAN
Tubulin folding cofactor B - Homo sapiens (Human)					
Q99426	2	3.2734	R.AQQEAEAAQR.L	2	TBCB_HUMAN
Tubulin polymerization-promoting protein - Homo sapiens (Human)					
O94811	3	5.4448	K.AGRVDLVDESGYVSGYK.H	2	TPPP_HUMAN
O94811	2	3.9589	R.TITFEQFQEALEELAK.K	1	TPPP_HUMAN
O94811	2	4.3348	R.VDLVDESGYVSGYK.H	4	TPPP_HUMAN
Tubulin polymerization-promoting protein family member 3 - Homo sapiens (Human)					
Q9BW30	2	5.6594	K.GIAGRQDILDDSGYVSAYK.N	6	TPPP3_HUMAN
Q9BW30	2	2.7648	K.ALEELATKR.F	1	TPPP3_HUMAN
Q9BW30	2	4.1436	K.SVTGTDVDIVFSK.V	5	TPPP3_HUMAN
Q9BW30	2	2.945	K.TGGAVDRLTDTSR.Y	1	TPPP3_HUMAN
Q9BW30	2	3.9765	R.QDILDDSGYVSAYK.N	4	TPPP3_HUMAN
Q9BW30	2	3.3888	R.VINYEEFKK.A	4	TPPP3_HUMAN
Q9BW30	3	6.8721	K.SKEEAFDAICQLVAGKEPANVGVTK.A	11	TPPP3_HUMAN
Tubulin-specific chaperone A - Homo sapiens (Human)					
O75347	2	4.391	K.MRAEDGENYDIKK.Q	10	TBCA_HUMAN
O75347	3	4.6429	R.RLEAAYLDLQR.I	10	TBCA_HUMAN
O75347	1	3.5868	R.LVLDSVKLEA.-	9	TBCA_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O75347	2	4.0991	R.LEAAYLDLQR.I	4	TBCA_HUMAN
O75347	3	6.5155	R.ILENEKDLEEAEEYKEAR.L	9	TBCA_HUMAN
O75347	3	4.4605	R.ILENEKDLEEAEEYK.E	2	TBCA_HUMAN
O75347	2	3.4023	R.AEDGENYDIK.K	1	TBCA_HUMAN
O75347	2	2.9606	K.QAEILQESR.M	2	TBCA_HUMAN
O75347	2	3.3399	K.MRAEDGENYDIK.K	1	TBCA_HUMAN
O75347	3	4.3195	K.M#RAEDGENYDIKK.Q	2	TBCA_HUMAN
O75347	2	3.7482	K.KQAEILQESR.M	3	TBCA_HUMAN
O75347	2	3.0793	K.EAKQQEEKIEK.M	2	TBCA_HUMAN
O75347	2	2.9807	K.VMYEKEAK.Q	1	TBCA_HUMAN
Tubulointerstitial nephritis antigen - Homo sapiens (Human)					
Q9UJW2	2	3.5934	R.HGCNSGSIDR.A	2	TINAG_HUMAN
Tubulointerstitial nephritis antigen-like precursor - Homo sapiens (Human)					
Q9GZM7	2	3.6403	R.IYPVLGTYWDNCNR.C	1	TINAL_HUMAN
Tudor domain-containing protein 3 - Homo sapiens (Human)					
Q9H7E2	2	3.7335	K.HFNVNTDYQNPVR.S	1	TDRD3_HUMAN
Q9H7E2	2	3.3391	R.SNLNM#NAAGNR.N	2	TDRD3_HUMAN
Tuftelin-interacting protein 11 - Homo sapiens (Human)					
Q9UBB9	2	3.1163	K.NAQGIINPIEAK.Q	1	TFP11_HUMAN
Q9UBB9	2	3.1145	K.TGGNFKPSQK.G	2	TFP11_HUMAN
Tumor suppressor candidate 1 - Homo sapiens (Human)					
Q2TAM9	3	3.7721	R.RVPEEASTNRR.A	1	Q2TAM9_HUMA
Tumor suppressor p53-binding protein 1 - Homo sapiens (Human)					
Q12888	2	4.6508	R.ADDPLRLDQELQQPQTQEK.T	3	TP53B_HUMAN
Q12888	3	5.6684	R.KVTEETEEPIVECQECETEVSPSQTGGSSGD	1	TP53B_HUMAN
Q12888	2	3.8784	K.TSNSLTEDSK.M	2	TP53B_HUMAN
Q12888	2	2.9034	K.SPEPEVLSTQEDLFDQSNK.T	1	TP53B_HUMAN
Q12888	2	3.0621	K.QDATVQTER.G	1	TP53B_HUMAN
Q12888	2	3.7941	K.LMLSTSEYSQSPK.M	2	TP53B_HUMAN
Q12888	2	3.471	K.LM#LSTSEYSQSPK.M	2	TP53B_HUMAN
Q12888	2	4.4948	K.EQLSAQELMESGLQIQK.S	2	TP53B_HUMAN
Q12888	2	3.6938	R.VPETVSAATQTIK.N	2	TP53B_HUMAN
Twinfilin-1 - Homo sapiens (Human)					
Q12792	1	2.1924	R.GPAETEATTD.-	2	TWF1_HUMAN
Twisted gastrulation protein homolog 1 precursor - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9GZX9	3	3.7687	K.STVEELHEPIPSLFR.A	1	TWSG1_HUMAN
Tyrosine-protein kinase ABL2 - Homo sapiens (Human)					
P42684	3	6.0797	R.TPSGDLAITEKDPPGVGVAGVAAAPK.G	2	ABL2_HUMAN
Tyrosine-protein kinase receptor - Homo sapiens (Human)					
Q86W07	2	3.364	K.GPGQNVAVTR.A	2	Q86W07_HUMA
Q5VTU6	2	3.0975	K.TGVSIQVAVKM#LKEK.A	2	Q5VTU6_HUMA
Tyrosine-protein kinase SgK269 - Homo sapiens (Human)					
Q9H792	2	2.7613	K.SAPTSPTATNISSK.T	2	SG269_HUMAN
Q9H792	2	2.8398	R.TDQAAVM#EK.G	1	SG269_HUMAN
Q9H792	2	2.8206	R.TDQAAVMEK.G	2	SG269_HUMAN
Q9H792	2	4.978	K.VPIVINPNAYDNLAIYK.S	3	SG269_HUMAN
Q9H792	2	4.3661	K.IEGTQESQM#VGSSSTR.E	3	SG269_HUMAN
Q9H792	3	4.1718	K.ASTDVAGQAVTINLVPTEEQAKPYR.V	1	SG269_HUMAN
Q9H792	2	2.8631	K.GFSNSTEHKR.G	1	SG269_HUMAN
Q9H792	3	4.8042	K.NAIKVPIVINPNAYDNLAIYK.S	1	SG269_HUMAN
Tyrosine-protein phosphatase non-receptor type 11 - Homo sapiens (Human)					
Q06124	2	3.3801	R.VYENVGLMQQQK.S	1	PTN11_HUMAN
Tyrosine-protein phosphatase non-receptor type 12 - Homo sapiens (Human)					
Q05209	3	5.0656	R.EQISENPTEATDIGFGNR.C	3	PTN12_HUMAN
Q05209	3	4.0423	R.TPESFVLASEHNTPVRSEWSELQSQER.S	1	PTN12_HUMAN
Q05209	3	4.2451	R.HNIAGTTHSGAEK.D	1	PTN12_HUMAN
Q05209	2	4.8118	K.STELPGKNESTIEQIDK.K	2	PTN12_HUMAN
Q05209	2	3.1165	K.SFDGNTLLNR.G	1	PTN12_HUMAN
Q05209	2	4.0448	R.SEWSELQSQER.S	5	PTN12_HUMAN
U1 small nuclear ribonucleoprotein 70 kDa - Homo sapiens (Human)					
P08621	3	5.15	K.EELRGGGGDMAEPSEAGDAPPDDGPPGELG	1	RU17_HUMAN
P08621	2	3.4234	R.RLGGGLGGTR.R	3	RU17_HUMAN
P08621	3	3.7949	R.YDERPGPSPLPHR.D	1	RU17_HUMAN
P08621	2	2.7595	R.VNYDTTESK.L	2	RU17_HUMAN
P08621	2	3.2145	R.RQQEVETELK.M	4	RU17_HUMAN
P08621	2	3.3945	R.RGGADVNIR.H	5	RU17_HUMAN
P08621	2	5.1144	R.GGGGGQDNGLEGLGNDSR.D	4	RU17_HUMAN
P08621	3	5.2015	R.GGGGDMAEPSEAGDAPPDDGPPGELGPDG	2	RU17_HUMAN
P08621	3	4.6588	R.GGGGDM#AEPSEAGDAPPDDGPPGELGPD	2	RU17_HUMAN
P08621	3	4.5063	R.GGGGDM#AEPSEAGDAPPDDGPPGELGPD	1	RU17_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
U1 small nuclear ribonucleoprotein A - Homo sapiens (Human)					
P09012	3	4.2837	K.KAVQGGGATPVVGAVQGPVPGM#PPM#TQA	1	SNRPA_HUMAN
P09012	3	5.7952	K.KAVQGGGATPVVGAVQGPVPGMPPMTQAP	3	SNRPA_HUMAN
U2 small nuclear ribonucleoprotein A' - Homo sapiens (Human)					
P09661	3	5.3283	K.GGPSPGDVEAIKNAIANASTLAEVER.L	3	RU2A_HUMAN
U2 small nuclear ribonucleoprotein B'' - Homo sapiens (Human)					
P08579	3	4.3452	R.LVPGRHDIAFVEFENDGQAGAAR.D	2	RU2B_HUMAN
P08579	2	3.5147	K.TVEQTATTTNK.K	4	RU2B_HUMAN
U2-associated protein SR140 - Homo sapiens (Human)					
O15042	3	7.3452	R.SSDVHSSGSSDAHMDASGPSDSDMPSR.T	1	SR140_HUMAN
U3 small nucleolar ribonucleoprotein protein MPP10 - Homo sapiens (Human)					
O00566	2	3.8584	K.TAEEENPEHVEIQK.M	1	MPP10_HUMAN
U3 small nucleolar RNA-associated protein 14 homolog A - Homo sapiens (Human)					
Q9BVJ6	2	3.563	K.DSGSQEVLSELR.V	1	UT14A_HUMAN
Q9BVJ6	2	3.0059	K.LVLADLLEPVK.T	2	UT14A_HUMAN
Q9BVJ6	3	5.7014	R.KVNPAAALEELEKIEK.A	1	UT14A_HUMAN
Q9BVJ6	3	5.1047	R.SELSQDAEPAGSQETKDSGSQEVLSELR.V	1	UT14A_HUMAN
U4/U6 small nuclear ribonucleoprotein Prp31 - Homo sapiens (Human)					
Q8WWY3	2	3.0478	K.YFSSM#AEFLK.V	4	PRP31_HUMAN
Q8WWY3	2	2.7358	K.YFSSMAEFLK.V	1	PRP31_HUMAN
Q8WWY3	2	3.5333	R.VRQTQVNEATK.A	2	PRP31_HUMAN
U4/U6.U5 tri-snRNP-associated protein 1 - Homo sapiens (Human)					
O43290	3	4.6876	R.GEKEAAGTTAAAGTGGATEQPPR.H	2	SNUT1_HUMAN
O43290	2	3.6204	R.RQDLYSAR.D	2	SNUT1_HUMAN
O43290	2	3.0692	R.ADDLLPLGDQTQDGDFGSR.L	1	SNUT1_HUMAN
O43290	2	3.1934	K.TSSGDASSLSIEETNKLR.A	1	SNUT1_HUMAN
O43290	2	4.0483	R.DDGYEAAASSK.T	8	SNUT1_HUMAN
O43290	3	4.1951	K.KPDYLPYAEDESVDDLAQQKPR.S	1	SNUT1_HUMAN
O43290	2	3.3883	R.DSGEKVVEIVK.K	4	SNUT1_HUMAN
O43290	2	3.3639	R.LASEYLTPEEMVTFKK.T	1	SNUT1_HUMAN
O43290	2	3.4209	R.LEQGGTADGLR.E	1	SNUT1_HUMAN
O43290	2	3.1479	R.LEQGGTADGLRER.E	2	SNUT1_HUMAN
O43290	2	3.492	R.LQAQSLSTVGPR.L	2	SNUT1_HUMAN
O43290	2	2.9176	R.QLQQLQQLR.D	2	SNUT1_HUMAN
O43290	2	2.9672	K.TSSGDASSLSIEETNK.L	1	SNUT1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O43290	2	4.5425	R.LRQLQQLQQLR.D	2	SNUT1_HUMAN
O43290	3	4.5346	K.EAGTKEEPVTADVINPM#ALR.Q	1	SNUT1_HUMAN
O43290	3	4.9555	K.LGLKPLEVNAIKK.E	3	SNUT1_HUMAN
O43290	2	5.2918	K.EAAGTTAAAGTGGATEQPPR.H	5	SNUT1_HUMAN
O43290	2	4.9552	K.EAGTKEEPVTADVINPMALR.Q	4	SNUT1_HUMAN
O43290	3	5.5724	K.IKTLGEDDPWLDDTAAWIER.S	3	SNUT1_HUMAN
O43290	2	3.1764	K.KLDEEALLK.K	2	SNUT1_HUMAN
O43290	2	3.946	K.TPYIVLSGSGK.S	2	SNUT1_HUMAN
O43290	3	3.8177	K.LGLKPLEVNAIK.K	1	SNUT1_HUMAN
O43290	2	4.3955	K.RDDGYEAAASSK.T	5	SNUT1_HUMAN
O43290	4	5.7988	K.DKGVLQEEEDVLVNVNLVDKER.A	4	SNUT1_HUMAN
O43290	2	3.0832	K.SM#NANTITK.-	1	SNUT1_HUMAN
O43290	2	2.7696	K.SMNANTITK.-	3	SNUT1_HUMAN
O43290	2	2.8657	K.KLDEEALLKK.M	1	SNUT1_HUMAN
U4/U6.U5 tri-snRNP-associated protein 3 - Homo sapiens (Human)					
Q8WVK2	2	3.5041	K.LMGFASFDSTK.G	3	SNUT3_HUMAN
U5 small nuclear ribonucleoprotein 200 kDa helicase - Homo sapiens (Human)					
O75643	1	2.2765	R.SLQYEYK.A	2	U520_HUMAN
U6 snRNA-associated Sm-like protein LSm3 - Homo sapiens (Human)					
P62310	2	3.8902	R.GDGVVLVAPPLRVG.-	7	LSM3_HUMAN
Ubinuclein - Homo sapiens (Human)					
Q9NPG3	3	4.1529	K.LLTSPSLKPSAVSSVTSSTSLSK.G	1	UBN1_HUMAN
Ubiquilin-2 - Homo sapiens (Human)					
Q9UHD9	2	5.0493	R.GPAAAQGSAAAPAEPK.I	4	UBQL2_HUMAN
Ubiquinol-cytochrome c reductase complex 11 kDa protein, mitochondrial precursor - Homo sapie					
P07919	3	5.4962	R.SHTEEDCTEELFDFLHAR.D	14	UCRH_HUMAN
P07919	1	2.2573	R.EQCEQLEK.C	3	UCRH_HUMAN
P07919	1	2.2005	R.LELCDER.V	1	UCRH_HUMAN
Ubiquinol-cytochrome c reductase complex 14 kDa protein - Homo sapiens (Human)					
P14927	2	3.5094	K.WYYNAAGFNK.L	3	UCR6_HUMAN
P14927	3	6.3945	K.LGLMRDDTIYEDEDVKEAIR.R	2	UCR6_HUMAN
P14927	3	4.5917	K.LGLM#RDDTIYEDEDVKEAIR.R	1	UCR6_HUMAN
Ubiquinol-cytochrome-c reductase complex core protein 2, mitochondrial precursor - Homo sapien					
P22695	2	2.8217	K.KFVSGQKSM#AASGNLGHTPFVDEL.-	1	UQCR2_HUMAN
P22695	2	3.5424	K.SM#AASGNLGHTPFVDEL.-	2	UQCR2_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Ubiquitin carboxyl-terminal hydrolase 14 - Homo sapiens (Human)					
P54578	2	3.5293	K.AQLFALTGVQPAR.Q	1	UBP14_HUMAN
Ubiquitin carboxyl-terminal hydrolase 31 - Homo sapiens (Human)					
Q70CQ4	2	3.4311	K.SQDSVSSPSPQK.Q	2	UBP31_HUMAN
Ubiquitin carboxyl-terminal hydrolase 8 - Homo sapiens (Human)					
P40818	3	4.5816	K.KKQEAEENEITEK.Q	1	UBP8_HUMAN
Ubiquitin carboxyl-terminal hydrolase isozyme L1 - Homo sapiens (Human)					
P09936	3	4.4797	-.MQLKPMEINPEMLNK.V	3	UCHL1_HUMAN
Ubiquitin domain-containing protein UBFD1 - Homo sapiens (Human)					
O14562	2	4.2404	K.IM#VVGSTINDVLAVNTPK.D	3	UBFD1_HUMAN
O14562	3	5.3941	R.GSLQPAPAQPPGDPAAQASVSNGEDAGGGA	2	UBFD1_HUMAN
O14562	3	3.9023	K.HDVKFPLDSTGSELK.Q	1	UBFD1_HUMAN
O14562	2	3.0981	R.LPTVPLSGMYNK.S	1	UBFD1_HUMAN
Ubiquitin fusion degradation protein 1 homolog - Homo sapiens (Human)					
Q92890	2	3.2975	K.KVEEDEAGGR.F	2	UFD1_HUMAN
Q92890	2	4.4093	R.FVAFSGEGQSLR.K	5	UFD1_HUMAN
Ubiquitin interaction motif-containing protein 1 - Homo sapiens (Human)					
Q96RL1	3	5.4517	R.SRPLATGPSSQSHQEK.T	2	UIMC1_HUMAN
Ubiquitin-associated protein 1 - Homo sapiens (Human)					
Q9NZ09	2	5.1113	R.GGSGSVLQDEEVLASLER.A	2	UBAP1_HUMAN
Q9NZ09	3	3.9587	R.NILVGTTGPIMAQLLDNNLPR.G	2	UBAP1_HUMAN
Ubiquitin-associated protein 2-like - Homo sapiens (Human)					
Q14157	1	2.4376	K.DLTQAK.N	5	UBP2L_HUMAN
Q14157	3	4.8978	R.RYPSSISSSPQKDLTQAK.N	1	UBP2L_HUMAN
Q14157	2	4.0474	R.RYPSSISSSPQK.D	4	UBP2L_HUMAN
Q14157	3	4.402	K.STSAPQMSPGSSDNQSSSPQPAQQK.L	6	UBP2L_HUMAN
Q14157	2	4.0326	K.GGSTTGSQFLEQFK.T	4	UBP2L_HUMAN
Ubiquitin-conjugating enzyme E2 L3 - Homo sapiens (Human)					
P68036	1	2.5926	K.NAEEFTK.K	1	UB2L3_HUMAN
P68036	2	2.8682	K.NAEEFTKK.Y	2	UB2L3_HUMAN
Ubiquitin-conjugating enzyme E2 Q1 - Homo sapiens (Human)					
Q7Z7E8	1	2.3348	K.ENLAILEKIK.K	1	UB2Q1_HUMAN
UBX domain-containing protein 1 - Homo sapiens (Human)					
Q9BZV1	2	3.6375	R.QGPTNEAQMAAAAALAR.L	3	UBXD1_HUMAN
UBX domain-containing protein 2 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q92575	2	2.9723	K.TKEEVEAAK.A	1	UBXD2_HUMAN
UBX domain-containing protein 7 - Homo sapiens (Human)					
O94888	2	3.7063	R.FELLTNFPR.R	2	UBXD7_HUMAN
UDP-glucose:glycoprotein glucosyltransferase 1 precursor - Homo sapiens (Human)					
Q9NYU2	2	4.2041	R.ILASPVELALVVMK.D	2	UGGG1_HUMAN
Q9NYU2	3	3.724	R.VEEDVASDLVMKVDALLSAQPK.G	1	UGGG1_HUMAN
UHRF1-binding protein 1 - Homo sapiens (Human)					
Q6BDS2	2	3.7614	K.TDEGVAAPVSGGAAR.L	2	URFB1_HUMAN
UMP-CMP kinase - Homo sapiens (Human)					
P30085	3	3.7752	-.MKPLVVFVLGGPGAGK.G	1	KCY_HUMAN
P30085	3	5.2408	K.RIQTYLQSTKPIIDLYEEMGK.V	2	KCY_HUMAN
P30085	2	2.8813	K.SVDEVFDEVVQIFDKEG.-	2	KCY_HUMAN
P30085	3	5.11	R.IQTYLQSTKPIIDLYEEM#GK.V	2	KCY_HUMAN
P30085	3	5.1391	R.IQTYLQSTKPIIDLYEEMGK.V	1	KCY_HUMAN
P30085	2	2.7968	R.SDDNRESLEK.R	2	KCY_HUMAN
Uncharacterized potential DNA-binding protein C17orf49 - Homo sapiens (Human)					
Q8IXM2	2	4.4587	K.VYEDSGIPLPAESPK.K	2	CQ049_HUMAN
Q8IXM2	2	2.9728	K.VGEIFSAAGAAFTK.L	1	CQ049_HUMAN
Uncharacterized protein C10orf118 - Homo sapiens (Human)					
Q7Z3E2	2	2.7579	K.QLESRIEELNK.E	1	CJ118_HUMAN
Uncharacterized protein C10orf32 - Homo sapiens (Human)					
Q96B45	3	4.5739	K.MAIITTHLQYQQEAIQK.N	2	CJ032_HUMAN
Q96B45	2	3.6581	R.SSELLGQAAR.N	1	CJ032_HUMAN
Uncharacterized protein C11orf52 - Homo sapiens (Human)					
Q96A22	2	3.3749	K.HVHLENATEYATLR.F	1	CK052_HUMAN
Q96A22	2	4.1645	R.VLQQQGSQER.S	4	CK052_HUMAN
Uncharacterized protein C12orf42 - Homo sapiens (Human)					
Q96LP6	2	2.809	-.M#STVIC*M#KQRDEEFLLTIR.P	1	CL042_HUMAN
Uncharacterized protein C12orf43 - Homo sapiens (Human)					
Q96C57	3	3.8998	K.VALEDDGFRLFFTSVPGGR.E	1	CL043_HUMAN
Q96C57	2	4.1813	R.AGAANSQLSTSQPSLR.H	3	CL043_HUMAN
Uncharacterized protein C12orf45 - Homo sapiens (Human)					
Q8N5I9	2	4.3223	R.SPLLDQVQTFLPQMAR.A	1	CL045_HUMAN
Uncharacterized protein C14orf173 - Homo sapiens (Human)					
Q27J81	2	3.4772	R.SQEEVPPDSDDNKTK.K	1	Q27J81_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Uncharacterized protein C14orf179 - Homo sapiens (Human)					
Q96FT9	2	4.0413	K.NSSLTLTGETSSAK.L	1	CN179_HUMAN
Uncharacterized protein C14orf43 - Homo sapiens (Human)					
Q6PJG2	2	3.7676	R.TSAAMLSQQVASVK.W	2	CN043_HUMAN
Uncharacterized protein C15orf38 - Homo sapiens (Human)					
Q7Z6K5	2	3.7944	K.TGASWTDNIMAQK.C	7	CO038_HUMAN
Q7Z6K5	2	4.1717	K.TRGDGPFLDSLAK.L	2	CO038_HUMAN
Q7Z6K5	3	3.813	R.GDGPFLDSLAKLEAGTVTK.C	1	CO038_HUMAN
Q7Z6K5	3	3.7693	R.KVNTGFLM#SSYKVEAK.G	1	CO038_HUMAN
Q7Z6K5	2	4.41	R.KVNTGFLMSSYK.V	1	CO038_HUMAN
Q7Z6K5	3	4.3203	R.KVNTGFLMSSYKVEAK.G	1	CO038_HUMAN
Uncharacterized protein C17orf37 - Homo sapiens (Human)					
Q9BRT3	2	3.024	R.ASNGETLEK.I	1	CQ037_HUMAN
Q9BRT3	2	2.7094	R.RASNGETLEK.I	1	CQ037_HUMAN
Uncharacterized protein C17orf59 - Homo sapiens (Human)					
Q96GS4	2	3.3346	R.ATISSPLELEGTVSR.H	1	CQ059_HUMAN
Uncharacterized protein C18orf25 - Homo sapiens (Human)					
Q96B23	2	3.9186	K.HGSGTQYVSTR.Q	4	CR025_HUMAN
Q96B23	2	3.8642	R.DSSESQLASTESDKPTTGR.V	2	CR025_HUMAN
Q96B23	2	3.0686	R.GGVIQSVSSWK.H	2	CR025_HUMAN
Q96B23	2	2.8087	R.SESETSTMAAK.K	1	CR025_HUMAN
Uncharacterized protein C18orf34 - Homo sapiens (Human)					
Q5BJE1	2	2.7279	K.LTEDNKKLEIDINK.I	1	CR034_HUMAN
Uncharacterized protein C19orf43 - Homo sapiens (Human)					
Q9BQ61	2	3.2405	K.TEDEVLTSK.G	1	CS043_HUMAN
Q9BQ61	2	4.9901	R.KGGPGSTLSFVGK.R	4	CS043_HUMAN
Q9BQ61	2	4.3892	K.TEDEVLTSKGDAWAK.Y	4	CS043_HUMAN
Q9BQ61	2	4.2827	K.KQKTEDEVLTSK.G	2	CS043_HUMAN
Q9BQ61	2	2.8703	K.GGPGSTLSFVGKR.R	2	CS043_HUMAN
Q9BQ61	2	2.9131	K.GGPGSTLSFVGK.R	1	CS043_HUMAN
Q9BQ61	2	2.7546	R.EAPGPAGGGGGGSR.W	1	CS043_HUMAN
Uncharacterized protein C19orf44 - Homo sapiens (Human)					
Q9H6X5	2	2.7931	K.QAPVENISPEAPAGKER.T	1	CS044_HUMAN
Uncharacterized protein C19orf58 - Homo sapiens (Human)					
Q9BW61	2	4.2548	R.DQEQVELEGESSAPPR.K	2	Q9BW61_HUMA

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
Q9BW61	2	3.3125	R.TDSPDMHEDT.-	1	Q9BW61_HUMA
Uncharacterized protein C19orf61 - Homo sapiens (Human)					
Q9H0W8	2	3.059	R.DASEETSTSVM#QK.T	1	CS061_HUMAN
Q9H0W8	2	3.4151	R.DASEETSTSVMQK.T	4	CS061_HUMAN
Uncharacterized protein C19orf62 - Homo sapiens (Human)					
Q9NWV8	3	4.0922	R.SEGEGEAASADDGSLNTSGAGPK.S	1	CS062_HUMAN
Q9NWV8	2	3.4264	R.AVGAQASVGSR.S	2	CS062_HUMAN
Uncharacterized protein C1orf174 - Homo sapiens (Human)					
Q8IYL3	2	2.898	R.AKDDDDDDDDDAEM.-	1	CA174_HUMAN
Q8IYL3	2	5.8666	K.HSAGSGAEESNSSSTVQK.Q	3	CA174_HUMAN
Q8IYL3	2	3.4672	R.AKDDDDDDDDDAEM#.-	3	CA174_HUMAN
Uncharacterized protein C1orf198 - Homo sapiens (Human)					
Q9H425	2	2.8138	R.QEQRPLPNVSTER.E	1	CA198_HUMAN
Q9H425	3	4.5378	R.SRGEGPEAEFQSLTPSQIK.S	2	CA198_HUMAN
Q9H425	2	3.9479	R.LPPAQQDEIIDR.C	3	CA198_HUMAN
Q9H425	3	4.2669	R.FGDEDLTWQDEHSAPFSWETK.S	1	CA198_HUMAN
Q9H425	3	5.0807	R.ERPQPVQAFSSALHEAAPSQLEGK.L	1	CA198_HUMAN
Q9H425	2	3.6513	R.DPGDSEELTR.F	4	CA198_HUMAN
Q9H425	2	2.7806	K.TGFDFLDNW.-	1	CA198_HUMAN
Q9H425	3	3.847	K.LPSPDVRQDDGEDTLFSEPK.F	1	CA198_HUMAN
Q9H425	2	4.2895	K.FAQVSSSNVVLK.T	4	CA198_HUMAN
Uncharacterized protein C1orf31 - Homo sapiens (Human)					
Q5JTJ3	2	3.4635	K.EKFEAGQFEPSETTAKS.-	2	CA031_HUMAN
Q5JTJ3	2	4.8937	K.FEAGQFEPSETTAK.S	5	CA031_HUMAN
Q5JTJ3	2	3.6708	K.FEAGQFEPSETTAKS.-	4	CA031_HUMAN
Q5JTJ3	3	4.7809	K.FKEKFEAGQFEPSETTAK.S	1	CA031_HUMAN
Q5JTJ3	2	3.7451	R.SSFESSCPQQWIK.Y	3	CA031_HUMAN
Q5JTJ3	2	4.8047	K.EKFEAGQFEPSETTAK.S	2	CA031_HUMAN
Uncharacterized protein C1orf50 - Homo sapiens (Human)					
Q9BV19	3	5.2839	R.AGDPLDLVALAEQVQK.A	1	CA050_HUMAN
Q9BV19	2	3.6	R.ESGQQYFSIISPK.E	3	CA050_HUMAN
Uncharacterized protein C1orf55 - Homo sapiens (Human)					
Q6IQ49	2	4.2027	R.VAEVAPEERENVAVAK.L	2	CA055_HUMAN
Q6IQ49	2	3.5866	R.MVTETEETQEK.K	4	CA055_HUMAN
Q6IQ49	2	3.2283	R.M#VTETEETQEK.K	1	CA055_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6IQ49	3	4.639	R.VVNTDHGSPEQLQIPVTDSGR.H	2	CA055_HUMAN
Uncharacterized protein C1orf71 - Homo sapiens (Human)					
Q6PJW8	2	3.4652	K.LVLQSLFSLIR.G	2	CA071_HUMAN
Uncharacterized protein C1orf96 - Homo sapiens (Human)					
Q6IQ19	2	2.7521	R.C*APPSPPPPVEPATQEEAERR.A	1	CA096_HUMAN
Uncharacterized protein C20orf116 precursor - Homo sapiens (Human)					
Q96HY6	4	5.7994	R.TQDTINRIQDLLAEGTITGVIDDRGK.F	1	CT116_HUMAN
Uncharacterized protein C20orf43 - Homo sapiens (Human)					
Q9BY42	2	4.3819	R.SIADSEESEAYK.S	1	CT043_HUMAN
Uncharacterized protein C21orf25 precursor - Homo sapiens (Human)					
Q9Y426	3	4.1388	K.DPGMSQSHNDLVFLEQPEGSR.R	1	CU025_HUMAN
Uncharacterized protein C2orf33 - Homo sapiens (Human)					
Q9GZY8	3	4.4044	K.VAPPNADLEQGFQEGVPNASVIMQVPER.I	1	CB033_HUMAN
Uncharacterized protein C2orf55 - Homo sapiens (Human)					
Q6NV74	2	2.7211	R.EAAEGLGEDSTGK.K	1	CB055_HUMAN
Uncharacterized protein C3orf19 - Homo sapiens (Human)					
Q6PII3	2	2.73	R.QLGVGYFAFAR.D	1	CC019_HUMAN
Uncharacterized protein C4orf8 - Homo sapiens (Human)					
P78312	3	4.9621	R.RVEDLLQFINSSETKPVSSTR.A	2	CD008_HUMAN
Uncharacterized protein C6orf203 - Homo sapiens (Human)					
Q9P0P8	3	5.3288	K.VGDTLDLLIGEDKEAGTETVM#R.I	1	CF203_HUMAN
Q9P0P8	3	5.6429	K.VGDTLDLLIGEDKEAGTETVMR.I	2	CF203_HUMAN
Uncharacterized protein C7orf28 - Homo sapiens (Human)					
O95766	2	3.0046	K.AMEDGGVKLLKER.L	1	CG028_HUMAN
Uncharacterized protein C8orf42 - Homo sapiens (Human)					
Q86YL5	2	3.0364	K.GHLTDSPEEAE.-	2	Q86YL5_HUMAN
Q86YL5	2	3.52	K.KSGFWDNLVLK.Q	2	Q86YL5_HUMAN
Q86YL5	2	2.7987	K.SGFWDNLVLK.Q	1	Q86YL5_HUMAN
Q86YL5	2	3.2668	K.YTSLASSANSSR.W	1	Q86YL5_HUMAN
Q86YL5	3	5.6948	R.VLLDEPPEEEDGLRGGPPPAAAAAAQAQVQ	2	Q86YL5_HUMAN
Uncharacterized protein C8orf45 - Homo sapiens (Human)					
Q4G0Z9	2	2.8605	R.FYM#MQGIVIAM#TTITK.Y	1	CH045_HUMAN
Uncharacterized protein C8orf47 - Homo sapiens (Human)					
Q6P6B1	3	6.92	K.DAGAGTEAESLKGNAEAQPLGPEAK.G	4	CH047_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6P6B1	2	2.8794	R.NVEAGAYVEMIR.N	1	CH047_HUMAN
Q6P6B1	2	2.9491	R.ESTVDGNVQR.E	1	CH047_HUMAN
Q6P6B1	3	6.1315	K.TPEGPGNMEQIQPEGIVGSMEHPAR.N	2	CH047_HUMAN
Uncharacterized protein C9orf142 - Homo sapiens (Human)					
Q9BUH6	2	3.608	K.SKKPAGGVDFDET.-	1	CI142_HUMAN
Q9BUH6	2	3.508	R.FGLSAAEDITPR.F	2	CI142_HUMAN
Uncharacterized protein C9orf40 - Homo sapiens (Human)					
Q8IXQ3	2	3.794	R.LPGGGGDDGAGR.A	13	CI040_HUMAN
Uncharacterized protein C9orf78 - Homo sapiens (Human)					
Q9NZ63	2	3.4762	R.GDSESEEDEQDSEEVR.L	1	CI078_HUMAN
Uncharacterized protein C9orf80 - Homo sapiens (Human)					
Q9NRY2	2	3.8104	R.VAILAELDKEK.R	1	CI080_HUMAN
Uncharacterized protein C9orf82 - Homo sapiens (Human)					
Q9H8G2	2	3.5266	K.SVNEILGLAESSPNEPK.A	1	CI082_HUMAN
Q9H8G2	2	4.3491	R.STDSSSVSGSLQQETK.Y	1	CI082_HUMAN
Uncharacterized protein KIAA0157 - Homo sapiens (Human)					
Q15018	2	3.3689	R.M#PSSGFAAEGR.S	1	K0157_HUMAN
Uncharacterized protein KIAA0256 - Homo sapiens (Human)					
Q93073	3	4.6284	K.NKTPVQLDLGDMLAALEK.Q	1	K0256_HUMAN
Q93073	2	3.2049	K.SLSETTATMLWK.S	2	K0256_HUMAN
Uncharacterized protein KIAA0460 - Homo sapiens (Human)					
Q5VT52	2	3.5159	R.APQFQESVGSFR.S	3	K0460_HUMAN
Q5VT52	3	4.2307	R.DAPTHLPSVDLSNPFTK.E	1	K0460_HUMAN
Q5VT52	2	3.8414	R.LSSSPGLFGAFSVR.G	2	K0460_HUMAN
Uncharacterized protein KIAA0515 - Homo sapiens (Human)					
Q5JSZ5	2	3.2247	K.LQEAPSAASQM#K.R	2	K0515_HUMAN
Q5JSZ5	2	3.737	K.SSQGDSGVDLSAESR.E	1	K0515_HUMAN
Q5JSZ5	2	2.8174	R.AGEQGEAMK.Q	1	K0515_HUMAN
Q5JSZ5	2	3.6136	R.AIGLSPMSFPTADLTLK.M	2	K0515_HUMAN
Q5JSZ5	2	3.5517	R.LLSFSPEEFPTLK.A	4	K0515_HUMAN
Uncharacterized protein KIAA0802 - Homo sapiens (Human)					
Q9Y4B5	2	2.8181	K.AGGGATPVSSPSR.S	1	K0802_HUMAN
Q9Y4B5	2	2.7938	K.APPSPGSLAAPGR.L	2	K0802_HUMAN
Uncharacterized protein KIAA0819 - Homo sapiens (Human)					
O94909	2	3.0161	R.TYTEEELNAK.L	2	K0819_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
O94909	2	3.8483	K.GKSEEELEASK.S	2	K0819_HUMAN
O94909	2	2.7001	K.LMQEWFK.L	1	K0819_HUMAN
O94909	3	3.7895	K.SPQDEGLHLLKPLSIPK.R	1	K0819_HUMAN
O94909	2	3.7232	R.DSLVALLEEQR.L	2	K0819_HUMAN
O94909	3	5.0258	R.EREEDKDLEAAMLSK.G	1	K0819_HUMAN
O94909	2	2.9071	R.GPSQATSPIR.S	2	K0819_HUMAN
O94909	2	3.5523	R.SDLVEEFWMK.S	2	K0819_HUMAN
Uncharacterized protein KIAA1143 - Homo sapiens (Human)					
Q96AT1	2	3.7212	R.NQVSYVRPAEPAFLAR.F	1	K1143_HUMAN
Q96AT1	3	4.5972	K.KPNEDEVNQDSVKK.N	2	K1143_HUMAN
Q96AT1	3	4.5003	K.KKPNEDEVNQDSVK.K	1	K1143_HUMAN
Q96AT1	2	3.5947	K.KGDLSVEEVMK.I	2	K1143_HUMAN
Q96AT1	2	3.7413	K.KGDLSVEEVM#K.I	2	K1143_HUMAN
Q96AT1	2	4.0066	K.AAKADEEPTPADGR.I	2	K1143_HUMAN
Q96AT1	2	3.3412	K.ADEEPTPADGR.I	7	K1143_HUMAN
Uncharacterized protein KIAA1543 - Homo sapiens (Human)					
Q9P1Y5	2	3.7642	R.APAEEEVGPR.K	2	K1543_HUMAN
Q9P1Y5	2	3.2797	K.LSAALSSLQR.D	2	K1543_HUMAN
Q9P1Y5	2	3.6486	K.AVASSPAATNSEVK.M	4	K1543_HUMAN
Q9P1Y5	3	4.5781	K.AEAEAGAGSPTSTPAPPEALSSEMSELSAR.L	1	K1543_HUMAN
Q9P1Y5	3	4.1209	R.LAQEEAPGPAPLVSAVPM#ATPAPAAR.A	1	K1543_HUMAN
Uncharacterized protein KIAA1671 - Homo sapiens (Human)					
Q9BY89	2	3.0407	R.EGDPGPAQVPQPAVR.M	1	K1671_HUMAN
Q9BY89	2	4.0054	R.VETQEVNPGASR.D	2	K1671_HUMAN
Q9BY89	2	2.7512	R.VADGEAAAGGEWASR.R	1	K1671_HUMAN
Q9BY89	2	4.021	R.SPLEDETDNTWMFK.D	1	K1671_HUMAN
Q9BY89	3	3.8616	R.SAPLSQDTKPPVPQEEAGQDHPPSK.A	2	K1671_HUMAN
Q9BY89	2	2.7682	R.RSGPFVDQLK.Q	1	K1671_HUMAN
Q9BY89	2	3.4061	R.MPAFPGMDPAVLK.A	1	K1671_HUMAN
Q9BY89	2	3.3656	R.DLEKEDAPQEK.E	2	K1671_HUMAN
Q9BY89	3	3.8152	K.SPVPSLRPSSTGPSPSGGLSEEPAAK.D	1	K1671_HUMAN
Q9BY89	2	2.8386	K.SPFQPGVLGSR.V	2	K1671_HUMAN
Q9BY89	2	3.4526	K.MSPSGGAPQTTPTLR.S	2	K1671_HUMAN
Q9BY89	3	4.4372	K.ERPLQQVSPVASVPWR.S	2	K1671_HUMAN
Q9BY89	3	5.7429	K.DLDNRMPGLVGQEVGSGEGPR.T	2	K1671_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9BY89	2	3.6961	R.MPGLVGQEVGSGEGPR.T	2	K1671_HUMAN
Uncharacterized protein KIAA1704 - Homo sapiens (Human)					
Q8IXQ4	2	4.2209	R.LAEQVSSYNESK.R	3	K1704_HUMAN
Q8IXQ4	2	4.0383	R.LAEQVSSYNESKR.S	2	K1704_HUMAN
Q8IXQ4	3	5.5688	K.RLAEQVSSYNESKR.S	2	K1704_HUMAN
Q8IXQ4	2	3.5306	R.DLIGPALPPGFK.A	2	K1704_HUMAN
Q8IXQ4	2	3.374	K.KDEEHILSGR.D	2	K1704_HUMAN
Q8IXQ4	3	4.0093	K.GPVNYNVTTEFEKR.A	2	K1704_HUMAN
Q8IXQ4	2	4.406	K.GPVNYNVTTEFEK.R	3	K1704_HUMAN
Q8IXQ4	2	2.8936	K.GDDDSSKPIVR.E	2	K1704_HUMAN
Q8IXQ4	3	3.8574	K.LTKGDDDSSKPIVR.E	1	K1704_HUMAN
Uncharacterized protein KIAA1712 - Homo sapiens (Human)					
Q9C0F1	2	3.2189	K.VMVDENTWTNLLSR.V	1	K1712_HUMAN
Uncharacterized protein RUNDC2B - Homo sapiens (Human)					
Q8IUI4	2	3.0736	K.SIDDEDVDENEDDVYGNSSGR.K	2	RUN2B_HUMAN
UPF0235 protein C15orf40 - Homo sapiens (Human)					
Q8WUR7	2	3.9897	K.LLASTTPEEILEK.L	8	CO040_HUMAN
UPF0318 protein FAM120A - Homo sapiens (Human)					
Q9NZB2	2	4.5377	K.SQGGVQPIPSQGGK.L	2	F120A_HUMAN
Q9NZB2	2	3.8893	K.AEGSSTASSGSQLAEGK.G	2	F120A_HUMAN
Q9NZB2	2	5.9787	K.NQAAIQGRPPYAASAEEVAK.E	3	F120A_HUMAN
UPF0361 protein DC12 - Homo sapiens (Human)					
Q96FZ2	2	3.0327	R.GTAGLLEQWLK.R	1	DC12_HUMAN
UPF0366 protein C11orf67 - Homo sapiens (Human)					
Q9H7C9	2	2.8853	R.ETGTEHSPGVQPADVK.E	1	CK067_HUMAN
Q9H7C9	3	4.1244	R.ETGTEHSPGVQPADVKEVVEK.G	1	CK067_HUMAN
Q9H7C9	2	4.967	R.VLQTEQAVKEYNALVAQGVR.V	3	CK067_HUMAN
Q9H7C9	2	4.0039	K.EYNALVAQGVR.V	3	CK067_HUMAN
UPF0368 protein Cxorf26 - Homo sapiens (Human)					
Q9BVG4	2	3.3821	K.FNGIVEDFNYGTLLR.L	2	CX026_HUMAN
UPF0369 protein C6orf57 precursor - Homo sapiens (Human)					
Q5VUM1	2	3.379	K.TSSSQGGKSELVK.Q	2	CF057_HUMAN
Q5VUM1	3	3.8065	R.FDAPEDSHLEKEPLEKFPDDVNPVTK.E	1	CF057_HUMAN
UPF0378 family protein KIAA0100 precursor - Homo sapiens (Human)					
Q14667	2	2.9704	R.QLIATDDAVPLGPGK.G	1	K0100_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
UPF0384 protein CGI-117 - Homo sapiens (Human)					
Q9Y3C1	3	4.1826	K.RFYPAEWQDFLDSLQKR.K	1	U384_HUMAN
Q9Y3C1	2	3.5522	R.DEKNYYQDTPK.Q	3	U384_HUMAN
UPF0404 protein C11orf59 - Homo sapiens (Human)					
Q6IAA8	2	4.4291	R.TDEQALLSSILAK.T	2	CK059_HUMAN
UPF0418 protein C8orf70 - Homo sapiens (Human)					
Q96GY0	2	3.2307	R.RFNENAADR.H	3	CH070_HUMAN
UPF0428 protein CXorf56 - Homo sapiens (Human)					
Q9H5V9	2	2.8537	R.RLQELAELEAK.K	1	CX056_HUMAN
UPF0430 protein - Homo sapiens (Human)					
Q9NWB6	3	4.8947	K.IRQQEIEEKLIEEETAR.R	2	U430_HUMAN
Q9NWB6	2	3.2999	R.RVEELVAK.R	4	U430_HUMAN
UPF0444 protein C12orf23 - Homo sapiens (Human)					
Q8WUH6	2	5.1868	K.GAVGATIGGVAWIGGK.S	1	CL023_HUMAN
Q8WUH6	2	6.0266	K.GGVSAVAGGVTAVGSAVVNK.V	5	CL023_HUMAN
Q8WUH6	2	6.1657	K.GGVSAVAGGVTAVGSAVVNKVPLTGK.K	1	CL023_HUMAN
Q8WUH6	2	3.6173	K.TAVTTVPSM#GIGLVK.G	1	CL023_HUMAN
Q8WUH6	2	3.7755	K.TAVTTVPSMGIGLVK.G	2	CL023_HUMAN
Q8WUH6	2	3.3867	R.VTGGIFSVTK.G	2	CL023_HUMAN
UPF0446 protein C12orf31 - Homo sapiens (Human)					
Q9BRT6	3	4.3595	K.TLLDQHGQYPIWM#NQR.Q	1	CL031_HUMAN
UPF0448 protein C10orf104 - Homo sapiens (Human)					
Q96DE5	2	3.3024	K.LAGLVEELEADEWR.F	1	CJ104_HUMAN
UPF0449 protein C19orf25 - Homo sapiens (Human)					
Q9UFG5	3	4.4557	R.VLLPTRPAPPTVEQILEDVR.G	2	CS025_HUMAN
Q9UFG5	3	4.3358	R.MMEDAEAPGEQLYQQSR.A	4	CS025_HUMAN
Q9UFG5	2	3.155	R.LQQAGNVLR.Q	1	CS025_HUMAN
Q9UFG5	2	3.2861	R.AGEDLEREVAQMK.Q	3	CS025_HUMAN
Q9UFG5	3	3.9442	R.GAPAEDPVFTILAPEDPPVPFR.M	2	CS025_HUMAN
UPF0451 protein C17orf61 precursor - Homo sapiens (Human)					
Q8N2U0	1	2.7042	K.ELFDKANK.H	2	CQ061_HUMAN
Upstream stimulatory factor 2 - Homo sapiens (Human)					
Q15853	2	3.1942	R.TETNGGQVTYR.V	2	USF2_HUMAN
Uromodulin precursor - Homo sapiens (Human)					
P07911	2	3.1223	K.TALQPMVSALNIR.V	2	UROM_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P07911	2	4.5408	R.DSTIQVVENGESSQGR.F	5	UROM_HUMAN
P07911	1	3.193	R.FVGQGGAR.M	2	UROM_HUMAN
USP6 N-terminal-like protein - Homo sapiens (Human)					
Q92738	2	2.7384	K.LKDEADFQR.K	1	US6NL_HUMAN
Q92738	2	3.0198	R.IEVLPVDTGAGGYSGNSGSPK.N	1	US6NL_HUMAN
UV excision repair protein RAD23 homolog A - Homo sapiens (Human)					
P54725	3	4.128	K.AGQGTSAPPEASPTAAPESSTSFPPAPTSGM	6	RD23A_HUMAN
P54725	2	3.4457	K.IRMEPDETVK.V	1	RD23A_HUMAN
P54725	2	2.8567	R.ASYNNPHR.A	4	RD23A_HUMAN
P54725	3	5.011	R.EDKSPSEESAPTTSPESVSGSVPSSGSSGR.	1	RD23A_HUMAN
P54725	3	8.229	R.QVIQQNPALLPALLQQLGQENPQLLQQISR.H	4	RD23A_HUMAN
P54725	3	4.3269	K.AGQGTSAPPEASPTAAPESSTSFPPAPTSGM	2	RD23A_HUMAN
UV excision repair protein RAD23 homolog B - Homo sapiens (Human)					
P54727	2	2.8911	R.AVEYLLMGIPGDR.E	1	RD23B_HUMAN
P54727	1	3.0716	R.NQPQFQQMR.Q	4	RD23B_HUMAN
P54727	2	4.854	R.QIIQQNPSLLPALLQQIGR.E	1	RD23B_HUMAN
UV radiation resistance-associated gene protein - Homo sapiens (Human)					
Q9P2Y5	2	3.6049	R.IYALNENVSSFR.R	2	UVRAG_HUMAN
Uveal autoantigen with coiled-coil domains and ankyrin repeats - Homo sapiens (Human)					
Q9BZF9	2	3.5774	K.LAQHVKPEEHEQVK.S	1	UACA_HUMAN
Q9BZF9	2	2.8504	R.NLENTQNQIK.A	2	UACA_HUMAN
Q9BZF9	3	4.329	K.SHDAIIDDLNRK.L	2	UACA_HUMAN
Q9BZF9	2	3.8542	K.QTSEILAVQNLLQK.Q	1	UACA_HUMAN
Q9BZF9	3	3.7306	K.KKFEDINQEFVK.I	1	UACA_HUMAN
Q9BZF9	2	4.2805	K.DNKITELLNDVER.L	1	UACA_HUMAN
Q9BZF9	2	3.5849	K.SDLETQISSLNEK.L	1	UACA_HUMAN
Vacuolar ATP synthase catalytic subunit A - Homo sapiens (Human)					
P38606	3	5.5379	K.IKSDYAQLLEDMQNAFR.S	3	VATA_HUMAN
Vacuolar ATP synthase subunit E 1 - Homo sapiens (Human)					
P36543	2	2.8022	R.GALFGANANR.K	1	VATE1_HUMAN
P36543	2	3.5635	R.LDLIAQQMMPEVR.G	2	VATE1_HUMAN
Vacuolar ATP synthase subunit F - Homo sapiens (Human)					
Q16864	3	6.5824	R.HALDAHQQSIPAVLEIPSKEHPYDAAK.D	3	VATF_HUMAN
Q16864	2	3.2208	R.GM#FTAEDLR.-	1	VATF_HUMAN
Q16864	2	3.312	R.GMFTAEDLR.-	2	VATF_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Vacuolar ATP synthase subunit G 1 - Homo sapiens (Human)					
O75348	2	4.7754	K.QAKEEAQAEIEQYR.L	4	VATG1_HUMAN
O75348	3	5.0231	R.LKQAKEEAQAEIEQYR.L	2	VATG1_HUMAN
O75348	2	4.2552	R.GSCSTEVEKETQEK.M	2	VATG1_HUMAN
O75348	2	3.4933	K.M#TILQTYFR.Q	2	VATG1_HUMAN
O75348	2	3.5784	K.AKEAAALGSR.G	3	VATG1_HUMAN
O75348	2	3.6114	K.MTILQTYFR.Q	3	VATG1_HUMAN
Vacuolar protein sorting 37 homolog C - Homo sapiens (Human)					
A5D8V6	3	4.3412	K.FLEGEVPLETFLENFSSM#R.M	1	A5D8V6_HUMAN
Vacuolar protein sorting-associated protein 13C - Homo sapiens (Human)					
Q709C8	2	4.6707	R.GVVGGVTGIITKPVEGAK.K	2	VP13C_HUMAN
Vacuolar protein sorting-associated protein 54 - Homo sapiens (Human)					
Q9P1Q0	2	2.7228	R.KSTSLLGALQSQAIK.F	1	VPS54_HUMAN
Vacuolar protein sorting-associated protein 72 homolog - Homo sapiens (Human)					
Q15906	2	3.5478	R.KVNTPAGSSQK.A	8	VPS72_HUMAN
Vascular cell adhesion protein 1 precursor - Homo sapiens (Human)					
P19320	2	4.7374	K.SLEVTFTPVIEDIGK.V	2	VCAM1_HUMAN
P19320	2	3.536	K.SQEFLEDADRK.S	1	VCAM1_HUMAN
P19320	1	2.9966	K.NTVISVNPSTK.L	3	VCAM1_HUMAN
P19320	3	4.6982	K.EVELIVQEKPFTVEISPGPR.I	1	VCAM1_HUMAN
P19320	2	3.3776	R.QSTQTLYVNVAPR.D	2	VCAM1_HUMAN
Vascular endothelial growth factor receptor 2 precursor - Homo sapiens (Human)					
P35968	2	3.1033	K.TFEDIPLEEPEVK.V	1	VGFR2_HUMAN
Vasodilator-stimulated phosphoprotein - Homo sapiens (Human)					
P50552	2	3.6974	R.SGGGGLM#EEMNAMLAR.R	6	VASP_HUMAN
P50552	2	3.005	K.QEEASGGPTAPK.A	1	VASP_HUMAN
P50552	2	3.2625	R.WLPAGTGPQAFSR.V	2	VASP_HUMAN
P50552	2	2.705	R.VPAQSESVR.R	2	VASP_HUMAN
P50552	3	3.7227	R.VKQELLEEVKK.E	1	VASP_HUMAN
P50552	2	3.4735	R.SGGGGLMEEMNAM#LAR.R	5	VASP_HUMAN
P50552	2	3.3875	R.SGGGGLMEEM#NAM#LAR.R	2	VASP_HUMAN
P50552	2	4.8727	R.KVSKQEEASGGPTAPK.A	2	VASP_HUMAN
P50552	3	5.7431	K.VSKQEEASGGPTAPK.A	6	VASP_HUMAN
P50552	3	5.6284	K.ATQVGEKTPKDESANQEEPEAR.V	2	VASP_HUMAN
P50552	3	4.5728	K.TPKDESANQEEPEAR.V	9	VASP_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P50552	3	4.9247	K.VKEEIIEAFVQELR.K	3	VASP_HUMAN
P50552	2	3.7395	K.DESANQEEPEAR.V	5	VASP_HUMAN
P50552	2	3.6053	R.SGGGGLM#EEMNAM#LAR.R	1	VASP_HUMAN
Vasohibin-1 - Homo sapiens (Human)					
Q7L8A9	2	3.4201	K.VAGGGSSGATPTSAAATAPSGVR.R	1	VASH1_HUMAN
Vasorin precursor - Homo sapiens (Human)					
Q6EMK4	2	2.7073	R.YLQGSSVQLR.S	1	VASN_HUMAN
Versican core protein precursor - Homo sapiens (Human)					
P13611	2	2.7458	R.RTWDAAER.E	1	CSPG2_HUMAN
P13611	3	3.8157	R.SPQETYDVYCYVDHLDGDVFHLTVPSK.F	1	CSPG2_HUMAN
P13611	2	2.8252	R.STDGSFQDR.F	1	CSPG2_HUMAN
P13611	2	3.2135	R.TQEEYEDKK.H	5	CSPG2_HUMAN
P13611	1	2.5671	R.TWDAAER.E	1	CSPG2_HUMAN
P13611	2	4.1177	R.VGHDYQWIGLNDK.M	2	CSPG2_HUMAN
P13611	2	2.9294	R.YEINSLIR.Y	2	CSPG2_HUMAN
P13611	2	3.6995	R.LGEPNYGAEIR.G	4	CSPG2_HUMAN
P13611	2	3.6954	R.YTLNFEAAQK.A	3	CSPG2_HUMAN
P13611	2	3.2699	R.VSVPTHPEAVGDASLTVVK.L	1	CSPG2_HUMAN
P13611	2	2.7053	K.IEVDKNGK.D	1	CSPG2_HUMAN
P13611	3	6.3061	R.LQGAHLTSILSHEEQMFVNR.V	3	CSPG2_HUMAN
P13611	2	2.8262	K.DNSINTSK.H	1	CSPG2_HUMAN
P13611	3	5.3537	R.LQGAHLTSILSHEEQM#FVNR.V	2	CSPG2_HUMAN
P13611	2	2.7926	K.IGQDYKGR.V	2	CSPG2_HUMAN
P13611	2	3.324	K.ITCMNPSAYQR.T	4	CSPG2_HUMAN
P13611	2	4.9334	K.KGTVACGQPPVVENAK.T	2	CSPG2_HUMAN
P13611	2	3.8793	K.LLASDAGLYR.C	7	CSPG2_HUMAN
P13611	2	4.5728	R.AATSRYTLNFEAAQK.A	3	CSPG2_HUMAN
P13611	2	3.6093	R.AQCGGGLLGVR.T	5	CSPG2_HUMAN
P13611	2	4.115	R.LATVGELQAAWR.N	9	CSPG2_HUMAN
Vesicle transport through interaction with t-SNAREs homolog 1B - Homo sapiens (Human)					
Q9UEU0	3	4.0653	K.LIRDFDEKQQEANETLAEMEEELR.Y	1	VTI1B_HUMAN
Q9UEU0	2	3.2904	R.LLGTAGTEEK.K	1	VTI1B_HUMAN
Vesicle-associated membrane protein 2 - Homo sapiens (Human)					
P63027	3	4.9147	R.LQQTQAQVDEVVDIMR.V	1	VAMP2_HUMAN
Vesicle-associated membrane protein 3 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q15836	2	5.1901	R.LQQTQNQVDEVVDIM#R.V	3	VAMP3_HUMAN
Q15836	2	4.0892	R.LQQTQNQVDEVVDIMR.V	5	VAMP3_HUMAN
Q15836	3	4.5014	R.RLQQTQNQVDEVVDIMR.V	5	VAMP3_HUMAN
Vesicle-associated membrane protein 8 - Homo sapiens (Human)					
Q9BV40	3	5.1015	R.NLQSEVEGVKNIMTQNVER.I	3	VAMP8_HUMAN
Q9BV40	2	3.1137	R.NLQSEVEGVK.N	4	VAMP8_HUMAN
Q9BV40	3	3.879	R.NKTEDLEATSEHFK.T	1	VAMP8_HUMAN
Q9BV40	2	2.7064	R.GENLEHLR.N	1	VAMP8_HUMAN
Q9BV40	2	3.2659	K.NIMTQNVER.I	2	VAMP8_HUMAN
Vesicle-associated membrane protein-associated protein A - Homo sapiens (Human)					
Q9P0L0	2	4.6618	K.FKGPFTDVVTTNLK.L	4	VAPA_HUMAN
Q9P0L0	3	3.9019	K.HEQILVLDPPTDLKFK.G	2	VAPA_HUMAN
Q9P0L0	2	4.8815	K.HEQILVLDPPTDLK.F	4	VAPA_HUMAN
Vesicle-associated membrane protein-associated protein B/C - Homo sapiens (Human)					
O95292	2	4.9818	K.FRGPFTDVVTTNLK.L	2	VAPB_HUMAN
O95292	2	4.9873	K.VEQVLSLEPQHELK.F	5	VAPB_HUMAN
Vesicle-trafficking protein SEC22b - Homo sapiens (Human)					
O75396	3	4.8216	R.IMVANIEEVLQRGEALSALDSK.A	2	SC22B_HUMAN
Vesicular integral-membrane protein VIP36 precursor - Homo sapiens (Human)					
Q12907	2	3.2467	K.DNVDDPTGNFR.S	1	LMAN2_HUMAN
Vimentin - Homo sapiens (Human)					
P08670	2	4.1528	R.ISLPLPNFSSLNLR.E	39	VIME_HUMAN
P08670	3	3.9417	R.ISLPLPNFSSLNLRETNLDSLPLVDTHSKR.T	1	VIME_HUMAN
P08670	2	3.7195	R.LGDLYEEEM#R.E	5	VIME_HUMAN
P08670	2	3.8633	R.LGDLYEEEMR.E	5	VIME_HUMAN
P08670	2	3.0715	R.LGDLYEEEMRELR.R	5	VIME_HUMAN
P08670	3	6.0651	R.LLQDSVDFSLADAINTEFK.N	241	VIME_HUMAN
P08670	3	3.9249	R.LLQDSVDFSLADAINTEFKNTR.T	3	VIME_HUMAN
P08670	2	3.5699	R.M#FGGPGTASR.P	3	VIME_HUMAN
P08670	1	3.0482	R.MFGGPGTASR.P	4	VIME_HUMAN
P08670	2	3.2642	R.MFGGPGTASRPSSSR.S	2	VIME_HUMAN
P08670	1	2.1599	R.QVDQLTNDKAR.V	1	VIME_HUMAN
P08670	2	5.6444	R.RQVDQLTNDKAR.V	6	VIME_HUMAN
P08670	2	3.0015	R.RMFGGPGTASRPSSSR.S	1	VIME_HUMAN
P08670	2	4.6721	R.VEVERDNLAEDIMR.L	18	VIME_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P08670	2	3.6188	R.RQVDQLTNDK.A	5	VIME_HUMAN
P08670	1	2.4845	R.SVSSSSYR.R	8	VIME_HUMAN
P08670	4	4.7792	R.TNEKVELQELNDRFANYIDK.V	4	VIME_HUMAN
P08670	2	2.932	R.TYSLGSALRPSTSR.S	6	VIME_HUMAN
P08670	3	5.1779	R.VEVERDNLAEDIM#R.L	6	VIME_HUMAN
P08670	2	3.6559	R.RMFGGPGTASR.P	1	VIME_HUMAN
P08670	2	4.1684	K.VELQELNDRFANYIDK.V	2	VIME_HUMAN
P08670	2	4.5624	R.SLYASSPGGVYATR.S	23	VIME_HUMAN
P08670	2	4.5089	K.ILLAELEQLK.G	10	VIME_HUMAN
P08670	2	5.1085	K.ILLAELEQLKGQGK.S	19	VIME_HUMAN
P08670	2	4.4037	K.SRLGDLYEEEM#R.E	4	VIME_HUMAN
P08670	2	4.3908	K.SRLGDLYEEEMR.E	11	VIME_HUMAN
P08670	3	6.9492	R.FLEQQNKILLAELEQLKGQGK.S	5	VIME_HUMAN
P08670	3	5.6405	K.TVETRDGQVINETSQHHDDLE.-	19	VIME_HUMAN
P08670	2	4.1579	R.DGQVINETSQHHDDLE.-	14	VIME_HUMAN
P08670	2	3.0847	R.DNLAEDIM#R.L	2	VIME_HUMAN
P08670	2	2.7384	R.FANYIDK.V	1	VIME_HUMAN
P08670	2	3.0152	K.SRLGDLYEEEMRELR.R	1	VIME_HUMAN
P08670	2	3.6905	R.FANYIDKVR.F	6	VIME_HUMAN
P08670	2	3.3028	R.DNLAEDIMR.L	3	VIME_HUMAN
P08670	2	3.6814	R.ETNLDSLPLVDTHSKR.T	5	VIME_HUMAN
P08670	2	3.5986	R.ETNLDSLPLVDTHSK.R	6	VIME_HUMAN
P08670	2	4.5237	R.EKLQEEMLQREEAENTLQSFR.Q	4	VIME_HUMAN
P08670	2	3.1379	R.EKLQEEMLQR.E	4	VIME_HUMAN
P08670	3	4.3743	R.EKLQEEM#LQREEAENTLQSFR.Q	1	VIME_HUMAN
P08670	2	2.9464	R.EKLQEEM#LQR.E	1	VIME_HUMAN
P08670	3	4.3204	R.FLEQQNKILLAELEQLK.G	1	VIME_HUMAN
Vimentin-type IF-associated coiled-coil protein - Homo sapiens (Human)					
Q2NL98	3	3.815	R.RPPLAGLLDALAEAER.L	1	Q2NL98_HUMAN
Q2NL98	3	4.2427	R.RRPPLAGLLDALAEAER.L	3	Q2NL98_HUMAN
Vinculin - Homo sapiens (Human)					
P18206	3	4.1783	K.AIPDLTAPVAAVQAAVSNLVR.V	1	VINC_HUMAN
P18206	2	3.9881	R.DPSASPGDAGEQAIR.Q	4	VINC_HUMAN
Vinexin - Homo sapiens (Human)					
O60504	3	4.8803	K.KPLVDDPGEKPSQPIEVLLER.E	1	VINEX_HUMAN

*Accession No**	*Charge*	*XC(Max)***	*Peptide**	*TC****	*Reference No**
O60504	2	2.8544	R.PAHRPGPATSSSGR.S	1	VINEX_HUMAN
O60504	2	5.0382	R.SPADPTDLGGQTSPR.R	5	VINEX_HUMAN
Vitamin D-binding protein variant - Homo sapiens (Human)					
Q53F31	2	3.2422	K.AKLPEATPTELAK.L	2	Q53F31_HUMAN
Q53F31	2	3.6917	K.LPEATPTELAK.L	2	Q53F31_HUMAN
Vitronectin precursor - Homo sapiens (Human)					
P04004	2	3.353	R.SIAQYWLGCPAPGHL.-	1	VTNC_HUMAN
P04004	3	4.9391	K.LIRDVWGIEGPIDAAFTR.I	1	VTNC_HUMAN
P04004	2	4.3322	R.DVWGIEGPIDAAFTR.I	6	VTNC_HUMAN
P04004	3	6.0197	R.DWHGVPGQVDAAM#AGR.I	3	VTNC_HUMAN
P04004	3	4.4682	R.DWHGVPGQVDAAMAGR.I	4	VTNC_HUMAN
P04004	2	3.994	R.FEDGVLDPDYPR.N	4	VTNC_HUMAN
P04004	2	3.2102	R.GQYCYELDEK.A	1	VTNC_HUMAN
P04004	2	3.614	R.IYISGMAPR.P	2	VTNC_HUMAN
Von Willebrand factor A domain containing 1 - Homo sapiens (Human)					
Q6PCB0	2	4.7355	R.EFVGQLVAPLPLGTGALR.A	2	Q6PCB0_HUMA
von Willebrand factor precursor - Homo sapiens (Human)					
P04275	2	2.9915	R.VEDFGNAWK.L	2	VWF_HUMAN
P04275	2	3.2308	R.ILAGPAGDSNVVK.L	2	VWF_HUMAN
VPS10 domain-containing receptor SorCS1 precursor - Homo sapiens (Human)					
Q8WY21	2	2.9222	R.ARGTGASM#AVAAR.S	1	SORC1_HUMAN
WAS/WASL-interacting protein family member 2 - Homo sapiens (Human)					
Q8TF74	2	2.8465	K.KVTNINDR.S	1	WIPF2_HUMAN
Q8TF74	2	4.7721	R.TGPSGQSLAPPPPPYR.Q	3	WIPF2_HUMAN
Q8TF74	2	3.1501	K.GGLFQGGVLK.L	1	WIPF2_HUMAN
Q8TF74	4	5.6925	R.SFLDDFESKYSFHPVEDFPAPEEYKHFQR.I	1	WIPF2_HUMAN
Q8TF74	2	3.0373	R.TPAGPPPPPPPPLR.N	2	WIPF2_HUMAN
Q8TF74	3	4.147	K.HSSSAPPPPPPGRR.A	1	WIPF2_HUMAN
Q8TF74	2	3.0903	K.GSSGGYGSGGAALQPK.G	2	WIPF2_HUMAN
Q8TF74	3	4.5587	K.DGSENLAGKPALQIPSSR.A	5	WIPF2_HUMAN
Q8TF74	2	3.1439	K.HSSSAPPPPPPGR.R	2	WIPF2_HUMAN
WD repeat and HMG-box DNA-binding protein 1 - Homo sapiens (Human)					
O75717	2	3.1693	R.VVDESDETENQEEK.A	2	WDHD1_HUMAN
WD repeat protein 22 - Homo sapiens (Human)					
Q96JK2	2	3.7041	R.EDPTDTPATDSSR.A	1	WDR22_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
WD repeat protein 33 - Homo sapiens (Human)					
Q9C0J8	2	2.7335	K.TPLLGDGPR.A	1	WDR33_HUMAN
Q9C0J8	3	6.1863	R.GRDGFPGPEDFGPEENFDASEEAAR.G	2	WDR33_HUMAN
Q9C0J8	3	4.1777	R.HEQSGGPEHGPER.G	1	WDR33_HUMAN
WD repeat protein 44 - Homo sapiens (Human)					
Q5JSH3	2	3.8753	K.IIESIIEESQK.V	5	WDR44_HUMAN
Q5JSH3	3	6.113	K.VLQLEDDSLDSKGKELSDQATASPIVAR.T	2	WDR44_HUMAN
Q5JSH3	2	3.4226	K.VLQLEDDSLDSK.G	2	WDR44_HUMAN
Q5JSH3	2	5.0723	K.KIIESIIEESQK.V	2	WDR44_HUMAN
Q5JSH3	2	4.8105	K.ELSDQATASPIVAR.T	3	WDR44_HUMAN
Q5JSH3	2	3.0133	K.VGNESPVQELK.Q	2	WDR44_HUMAN
WD repeat protein 91 - Homo sapiens (Human)					
A4D1P6	2	2.7589	K.KESFGGQGTK.G	2	WDR91_HUMAN
A4D1P6	2	4.1036	R.LSPAQGPPQPQSSAK.K	6	WDR91_HUMAN
Wilms tumor 1-associating protein - Homo sapiens (Human)					
Q15007	2	3.6076	R.STMVDPAINLFFLK.M	2	WTAP_HUMAN
Q15007	2	3.9064	R.TTASEPVEQSEATSK.D	2	WTAP_HUMAN
Wings apart-like protein homolog - Homo sapiens (Human)					
Q7Z5K2	3	3.8246	K.RPNFKPDIQEIPK.K	2	WAPL_HUMAN
Q7Z5K2	3	4.434	K.SSQGASNFDKLMDGTSQALAK.A	1	WAPL_HUMAN
Q7Z5K2	2	2.9845	K.WGETTFMAK.L	1	WAPL_HUMAN
Q7Z5K2	2	3.7709	R.IVEDDASISSCNK.L	2	WAPL_HUMAN
Q7Z5K2	2	3.3539	R.SM#DEFTASTPADLGEAGR.L	1	WAPL_HUMAN
Wiskott-Aldrich syndrome protein - Homo sapiens (Human)					
P42768	3	4.916	K.LIYDFIEDQGGLEAVR.Q	1	WASP_HUMAN
P42768	2	4.9624	R.AGISEAQLTDAETSK.L	3	WASP_HUMAN
P42768	3	4.6966	K.HVSHVGWDPQNGFDVNNLDPDLR.S	2	WASP_HUMAN
P42768	3	3.8638	R.AGISEAQLTDAETSKLIYDFIEDQGGLEAVRQ	1	WASP_HUMAN
Wiskott-Aldrich syndrome protein family member 2 - Homo sapiens (Human)					
Q9Y6W5	3	4.1263	K.VTQLDPKEEEVSLQGINTR.K	2	WASF2_HUMAN
Q9Y6W5	2	5.405	R.DVVGNDVATILSR.R	3	WASF2_HUMAN
Q9Y6W5	2	3.2867	R.SDLLSAIR.Q	3	WASF2_HUMAN
Wolframin - Homo sapiens (Human)					
O76024	2	3.3975	R.DAAAPAEPQAQHTR.S	2	WFS1_HUMAN
WW domain-binding protein 11 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q9Y2W2	2	4.0544	K.ATATISAKPQITNPK.A	3	WBP11_HUMAN
Q9Y2W2	2	3.6159	R.AVSILPLLGHGVPR.L	2	WBP11_HUMAN
Q9Y2W2	2	2.8043	R.RRDEDMLYSPELAQR.G	2	WBP11_HUMAN
Q9Y2W2	3	4.2651	R.LYEKENPDIYKELR.K	2	WBP11_HUMAN
Q9Y2W2	2	2.8487	R.KSEDDSAVPLAK.A	1	WBP11_HUMAN
Q9Y2W2	3	5.8852	R.KPPGPPPGPPPPQVVQMYGR.K	2	WBP11_HUMAN
Q9Y2W2	3	3.9047	R.DM#EKLDEMEFNPVQQPQLNEK.V	1	WBP11_HUMAN
Q9Y2W2	2	2.7207	R.DEDM#LYSPELAQR.G	1	WBP11_HUMAN
Q9Y2W2	2	3.6573	K.SEDDSAVPLAK.A	3	WBP11_HUMAN
Q9Y2W2	2	3.6325	K.RAQLSQYFDAVK.N	2	WBP11_HUMAN
Q9Y2W2	3	4.838	K.QIIRDMEKLDEMEFNPVQQPQLNEK.V	1	WBP11_HUMAN
Q9Y2W2	2	3.3535	K.ADDTSAATIEK.K	2	WBP11_HUMAN
Q9Y2W2	2	2.9576	K.ELTPLQAMMLR.M	2	WBP11_HUMAN
Q9Y2W2	4	7.4373	K.NAQHVEVESIPLPDMPHAPSNILIQDIPLPGAQ	1	WBP11_HUMAN
Q9Y2W2	2	3.0979	K.ADDTSAATIEKK.A	2	WBP11_HUMAN
Q9Y2W2	2	3.6228	R.AQLSQYFDAVK.N	3	WBP11_HUMAN
WW domain-binding protein 4 - Homo sapiens (Human)					
O75554	2	2.9677	K.NSDGGSDPETQKEK.S	1	WBP4_HUMAN
O75554	2	3.3107	K.SLDKAKEEEK.A	2	WBP4_HUMAN
O75554	2	4.0613	K.TVTSLGVMADGVAPVFK.K	2	WBP4_HUMAN
WW domain-containing adapter protein with coiled-coil - Homo sapiens (Human)					
Q9BTA9	2	3.4633	R.RGDSQPYQALK.Y	2	WAC_HUMAN
WW domain-containing transcription regulator protein 1 - Homo sapiens (Human)					
Q9GZV5	3	3.864	R.SHSSPASLQLGTGAGAAGSPAQQHAHLR.Q	1	WWTR1_HUMA
XPA-binding protein 1 - Homo sapiens (Human)					
Q9HCN4	2	4.1201	K.DM#GSVALDAGTAK.D	2	XAB1_HUMAN
YLP motif-containing protein 1 - Homo sapiens (Human)					
P49750	3	3.7968	K.AQAVTQPVPLANKPVPAQSTFPSK.T	1	YLPM1_HUMAN
P49750	2	3.0533	R.YEGHPAEGTK.S	2	YLPM1_HUMAN
P49750	2	3.4082	R.VGFQYQGIMQK.H	2	YLPM1_HUMAN
P49750	2	2.7323	R.RFEDLGSR.C	1	YLPM1_HUMAN
P49750	2	3.4663	R.STFETEHAGQR.D	3	YLPM1_HUMAN
P49750	2	4.319	R.DISTNKVEQIPYGER.I	4	YLPM1_HUMAN
P49750	3	4.6324	R.AIGFVVGQTDWEKITDESGHLAEK.A	1	YLPM1_HUMAN
P49750	2	4.4686	K.GPKPAFGQQHQQQPK.S	3	YLPM1_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
P49750	2	5.6858	K.SQAEPLSGNKEPLADTSSNQQK.N	6	YLPM1_HUMAN
P49750	2	3.2037	R.HFDQFWSAAK.T	1	YLPM1_HUMAN
YTH domain family protein 1 - Homo sapiens (Human)					
Q9BYJ9	2	4.9921	K.APGMNSLEQGMVGLK.I	3	YTHD1_HUMAN
Q9BYJ9	3	4.598	K.HNMDIGTWDNKGPVPK.A	2	YTHD1_HUMAN
Q9BYJ9	2	3.0509	K.IGDVSSSAVK.T	1	YTHD1_HUMAN
Q9BYJ9	2	3.0302	K.SGPVMGGGLPPPPIK.H	1	YTHD1_HUMAN
YTH domain family protein 2 - Homo sapiens (Human)					
Q9Y5A9	2	2.8072	K.DGLNDDDFEPYLSPQAR.P	2	YTHD2_HUMAN
Q9Y5A9	2	3.3654	K.LGSTEVASNVPK.V	2	YTHD2_HUMAN
YTH domain family protein 3 - Homo sapiens (Human)					
Q7Z739	3	4.0872	K.HNM#NIGTWDEKGSVVK.A	1	YTHD3_HUMAN
Q7Z739	2	3.0457	K.HNMNIGTWDEK.G	2	YTHD3_HUMAN
Q7Z739	3	4.5669	K.HNMNIGTWDEKGSVVK.A	1	YTHD3_HUMAN
Q7Z739	2	3.2184	K.VPGISSIEQGM#TGLK.I	1	YTHD3_HUMAN
Q7Z739	2	4.6054	K.VPGISSIEQGMTGLK.I	5	YTHD3_HUMAN
Q7Z739	2	3.7061	K.GNVGIGGSAVPPPPIK.H	2	YTHD3_HUMAN
YTH domain-containing protein 1 - Homo sapiens (Human)					
Q96MU7	2	3.633	R.RTQAVVSGR.R	2	YTDC1_HUMAN
Q96MU7	2	4.0305	K.SATEYKNEEYQR.S	3	YTDC1_HUMAN
Q96MU7	2	3.5372	R.QLVSKPLSSSVSNNK.R	1	YTDC1_HUMAN
YY1-associated factor 2 - Homo sapiens (Human)					
Q8IY57	2	3.0156	R.GEASSLNGESH.-	1	YAF2_HUMAN
Q8IY57	3	5.6065	K.SPPASSAASADQHSQSGSSSDNTER.G	3	YAF2_HUMAN
Zinc finger and BTB domain-containing protein 11 - Homo sapiens (Human)					
O95625	3	4.3867	K.DAPSSSSSNSTSNEASGTSSEK.G	1	ZBT11_HUMAN
Zinc finger and SCAN domain-containing protein 16 - Homo sapiens (Human)					
Q9H4T2	2	3.0432	R.TKNEELFQKEDMPK.D	1	ZSC16_HUMAN
Zinc finger and SCAN domain-containing protein 18 - Homo sapiens (Human)					
Q8TBC5	2	3.1588	K.TEEDGPANTEQK.L	2	ZSC18_HUMAN
Zinc finger CCCH domain-containing protein 11A - Homo sapiens (Human)					
O75152	2	2.866	K.AAVAVVPLVSEDK.S	1	ZC11A_HUMAN
O75152	2	2.9573	K.EASGETTGVDITK.I	1	ZC11A_HUMAN
O75152	2	4.4555	K.NLQEGNEVDSQSSIR.T	2	ZC11A_HUMAN
O75152	3	4.1254	R.VQQSSESSTSSPSQHEATPGAR.R	2	ZC11A_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Zinc finger CCCH domain-containing protein 13 - Homo sapiens (Human)					
Q5T200	3	4.2372	K.KKEDDVGIER.G	1	ZC313_HUMAN
Q5T200	2	3.066	R.NTEESSSPVRK.E	2	ZC313_HUMAN
Zinc finger CCCH domain-containing protein 5 - Homo sapiens (Human)					
Q9C0B0	2	3.0568	R.WQETAYVLGNYK.T	1	ZC3H5_HUMAN
Q9C0B0	3	3.959	R.KPPNLEGIVFPGESGLAPGSYK.K	1	ZC3H5_HUMAN
Zinc finger CCCH domain-containing protein C19orf7 - Homo sapiens (Human)					
Q9UPT8	2	3.6899	K.TGSGSPFAGNSPAR.E	3	CS007_HUMAN
Q9UPT8	2	3.3242	R.TVLWNPEDLIPLPIPK.Q	2	CS007_HUMAN
Q9UPT8	3	4.2053	R.QRPGASTDSSTQGANLPDFELLSR.I	1	CS007_HUMAN
Q9UPT8	2	2.9895	R.LQKPTDSTASSR.A	1	CS007_HUMAN
Q9UPT8	3	6.0338	R.LPLPDDREDGELEEGELEDDGAEETQDTSG	1	CS007_HUMAN
Q9UPT8	3	5.3045	R.VLAAGGLGQGGGGGQSSVLSGISLYDPR.T	1	CS007_HUMAN
Q9UPT8	2	4.2512	R.HVEASGGSGPGDSGPSDPR.L	1	CS007_HUMAN
Q9UPT8	3	4.1448	R.AAKPGPAEAPSPTASPSGDASPPATAPYDPR	1	CS007_HUMAN
Q9UPT8	3	4.5535	K.SALEQPETGKAGADGGTPTDR.Y	2	CS007_HUMAN
Q9UPT8	2	4.0392	K.GGMNDDEDFYDEDMGDGGGGSYR.S	1	CS007_HUMAN
Q9UPT8	2	2.818	K.ATEPAADTGAQPK.G	1	CS007_HUMAN
Q9UPT8	2	3.2861	K.AGADGGTPTDR.Y	4	CS007_HUMAN
Q9UPT8	2	2.7731	R.AATAGPPNAR.Q	1	CS007_HUMAN
Q9UPT8	2	3.4787	R.EGEQDAASLKDVFK.G	1	CS007_HUMAN
Zinc finger CCCH type antiviral protein 1 - Homo sapiens (Human)					
Q7Z2W4	2	3.5655	R.ASLEDAPVDDLTR.K	4	ZCC2_HUMAN
Q7Z2W4	2	3.9401	R.SSLGSLQTPEAVTTR.K	2	ZCC2_HUMAN
Q7Z2W4	2	2.7347	R.FFQGSQEFLASASASAER.S	1	ZCC2_HUMAN
Q7Z2W4	2	3.5893	K.SLTSWTNDQGAR.R	5	ZCC2_HUMAN
Q7Z2W4	2	5.2311	K.ATDLGGTSQAGTSQR.F	5	ZCC2_HUMAN
Q7Z2W4	3	4.3261	R.FLENGSQEDLLHGNPGSTYLASNSTSAPNW	1	ZCC2_HUMAN
Zinc finger CCHC domain-containing protein 6 - Homo sapiens (Human)					
Q5VYS8	1	2.4946	R.YPENKEK.R	1	ZCHC6_HUMAN
Zinc finger FYVE domain-containing protein 16 - Homo sapiens (Human)					
Q7Z3T8	3	7.6449	K.DVGLVKEEVDVAVITAAECLKEEGK.T	3	ZFY16_HUMAN
Q7Z3T8	2	2.8466	K.GLTSIQNEK.N	1	ZFY16_HUMAN
Q7Z3T8	2	3.4528	R.EQQNDTSSELQNR.E	2	ZFY16_HUMAN
Zinc finger homeobox protein 4 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q86UP3	2	2.8626	K.DYLAPTTVRQLMAQQELDRIK.K	1	ZFHX4_HUMAN
Zinc finger matrin-type protein 2 - Homo sapiens (Human)					
Q96NC0	1	2.226	R.STLDQVK.K	1	ZMAT2_HUMAN
Q96NC0	2	3.1058	R.STLDQVKK.R	1	ZMAT2_HUMAN
Zinc finger protein 148 - Homo sapiens (Human)					
Q9UQR1	2	2.8795	K.FLQQALDR.T	1	ZN148_HUMAN
Zinc finger protein 161 homolog - Homo sapiens (Human)					
O43829	2	2.9223	R.DVSSPDENNGQSK.S	1	ZF161_HUMAN
Zinc finger protein 185 - Homo sapiens (Human)					
A4FTV3	2	3.7406	K.STGPTQETQAPFIAK.R	2	A4FTV3_HUMAN
O15231	2	4.4248	K.STGSPTQETQAPFIAK.R	4	ZN185_HUMAN
A4FTV3	3	4.0673	R.DLAGEEAFRAPNTDAAR.S	1	A4FTV3_HUMAN
A4FTV3	2	3.0621	R.GSSSATSVSAVPADR.K	2	A4FTV3_HUMAN
A4FTV3	2	3.1893	R.GSSSATSVSAVPADRK.S	2	A4FTV3_HUMAN
A4FTV3	2	4.3176	R.GVFTKPIDSSSQPQQQFPK.A	2	A4FTV3_HUMAN
A4FTV3	3	5.0417	R.SSAQLSDGNVGSGATGSRPEGLAAVDIGSER	1	A4FTV3_HUMAN
A4FTV3	2	4.3765	R.TANAGPPRPSSSGYK.M	2	A4FTV3_HUMAN
Zinc finger protein 22 - Homo sapiens (Human)					
P17026	2	3.591	K.SFFQSSNLIQHR.R	1	ZNF22_HUMAN
P17026	2	3.807	K.SFSQSSTLFQHQK.I	2	ZNF22_HUMAN
Zinc finger protein 318 - Homo sapiens (Human)					
Q5VUA4	2	3.0936	R.GSIPAAQVPVQVSIPSLIR.Y	1	ZN318_HUMAN
Zinc finger protein 330 - Homo sapiens (Human)					
Q9Y3S2	2	3.3118	R.QTGGEEGDGASGYDAYWK.N	1	ZN330_HUMAN
Q9Y3S2	2	3.276	R.TYASGYAHYEEQEN.-	3	ZN330_HUMAN
Zinc finger protein 346 - Homo sapiens (Human)					
Q9UL40	2	3.6913	K.TVASSLGQIPM#QR.Q	2	ZN346_HUMAN
Q9UL40	2	3.8403	K.TVASSLGQIPMQR.Q	3	ZN346_HUMAN
Q9UL40	2	3.3617	R.LADPAVTDFPAGK.G	2	ZN346_HUMAN
Zinc finger protein 37 homolog - Homo sapiens (Human)					
A0AVJ9	2	3.3481	K.DDDQLENIQK.S	2	A0AVJ9_HUMAN
Zinc finger protein 395 - Homo sapiens (Human)					
Q9H8N7	3	5.5216	K.YLGDAFGSPQTDHGFETDPDPFLLDEPAPR.K	1	ZN395_HUMAN
Zinc finger protein 428 - Homo sapiens (Human)					
Q96B54	2	3.3538	R.SPLGEAPPGTPPCR.L	1	Q96B54_HUMAN

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q96B54	2	3.0395	R.ALGEEEEEPPR.A	2	Q96B54_HUMAN
Zinc finger protein 512B - Homo sapiens (Human)					
Q96KM6	3	3.7727	K.KQEGPGPEDARK.K	1	Z512B_HUMAN
Zinc finger protein 513 - Homo sapiens (Human)					
Q86UZ3	3	4.3368	R.DSEGDSLGARPGLPYGLSDDESGGGR.A	2	Q86UZ3_HUMA
Zinc finger protein 518 - Homo sapiens (Human)					
Q6AHZ1	2	2.7989	K.C*VM#PNKTELLKPK.L	1	ZN518_HUMAN
Zinc finger protein 581 - Homo sapiens (Human)					
Q9P0T4	2	3.269	R.DAGELAQHSR.V	2	ZN581_HUMAN
Zinc finger protein 598 - Homo sapiens (Human)					
Q86UK7	3	3.9675	K.KVAQPPLSAQATGSGQPTR.K	1	ZN598_HUMAN
Q86UK7	2	3.8848	R.ASVAAQQQEEAR.R	4	ZN598_HUMAN
Q86UK7	3	3.9837	R.RSEDQEEGGRPK.K	2	ZN598_HUMAN
Zinc finger protein 616 - Homo sapiens (Human)					
Q08AN1	2	2.8507	R.LAELQKVQTEGRLYECNETEK.T	1	ZN616_HUMAN
Zinc finger protein 622 - Homo sapiens (Human)					
Q969S3	3	4.4025	K.GLGVDSVDKDAMNAAIQQAIK.A	1	ZN622_HUMAN
Q969S3	3	5.1723	K.KFASFNAYENHLK.S	1	ZN622_HUMAN
Zinc finger protein 638 - Homo sapiens (Human)					
Q14966	2	4.0359	K.AILQLDSPESAQSMYSFLK.Q	2	ZN638_HUMAN
Q14966	2	2.9636	R.RLPNLPSQSR.N	1	ZN638_HUMAN
Q14966	2	3.5081	K.YGYTEDPLEVR.I	2	ZN638_HUMAN
Q14966	3	4.4732	K.TKLESLSQVGPVNENVMEEDLK.T	1	ZN638_HUMAN
Q14966	2	3.0088	K.QSSVTQVTEQSPK.V	2	ZN638_HUMAN
Q14966	3	6.2584	K.AVVTEPAKGEEAFQMSEVDEESGLKDSEPE	2	ZN638_HUMAN
Q14966	2	2.9284	K.NYQSQADIPIR.S	2	ZN638_HUMAN
Zinc finger protein 691 - Homo sapiens (Human)					
Q5VV50	2	2.9535	K.THLGEQAGKDSS.-	2	Q5VV50_HUMA
Zinc finger protein 703 - Homo sapiens (Human)					
Q9H7S9	2	3.8035	K.DSGSSSVSSTSSSSSSSPGDK.A	1	ZN703_HUMAN
Zinc finger protein 717 - Homo sapiens (Human)					
Q9BY31	2	2.7184	R.NHTLLYIRRTR.A	1	Q9BY31_HUMA
Zinc finger protein 781 - Homo sapiens (Human)					
Q8N8C0	2	2.9384	R.NLMNVRNVKR.L	2	ZN781_HUMAN
Zinc finger protein 787 - Homo sapiens (Human)					

Accession No*	Charge	XC(Max)**	Peptide*	TC***	Reference No*
Q6DD87	2	2.9672	R.LHPELSGPGVAAK.V	2	ZN787_HUMAN
Q6DD87	2	3.8015	R.AGGEEEDDDDEAAGGR.C	4	ZN787_HUMAN
Q6DD87	2	3.8559	K.TFSQSSHLVQHR.R	1	ZN787_HUMAN
Q6DD87	2	3.9974	K.GFGHGAGLLAHQR.A	2	ZN787_HUMAN
Zinc finger protein 8 - Homo sapiens (Human)					
P17098	2	2.7443	R.DSSQAIPITELTK.S	1	ZNF8_HUMAN
Zinc finger protein 91 homolog - Homo sapiens (Human)					
Q96JP5	3	5.4619	K.AAPEEPQQRPPEAVAAAPAGTTSSR.V	5	ZFP91_HUMAN
Zinc finger protein KIAA1802 - Homo sapiens (Human)					
Q96JM3	2	3.3803	K.MDM#TSPEQSR.N	2	K1802_HUMAN
Q96JM3	3	4.4253	K.SSFFIEPQKPVFPETR.K	4	K1802_HUMAN
Q96JM3	2	2.9918	R.KPGPSGPSESPK.A	2	K1802_HUMAN
Zinc finger protein ubi-d4 - Homo sapiens (Human)					
Q92785	2	4.2635	R.GPGLASGQLYSYPAR.R	1	REQU_HUMAN
Q92785	3	3.8596	R.GAPDPRVDDDSLGEFPVTNSR.A	1	REQU_HUMAN
Zinc finger protein with KRAB and SCAN domains 1 - Homo sapiens (Human)					
P17029	2	3.0863	R.SFSLSSNFTTPEEVPTGTK.S	1	ZKSC1_HUMAN
Zinc finger Ran-binding domain-containing protein 2 - Homo sapiens (Human)					
O95218	2	2.8916	R.TGYGGGFNER.E	2	ZRAB2_HUMAN
O95218	1	2.1925	K.AVGPASILK.E	1	ZRAB2_HUMAN
Zinc finger RNA-binding protein - Homo sapiens (Human)					
Q96KR1	2	4.6102	K.GLTTTGNSSLNSTSNTK.V	2	Q96KR1_HUMA
Zinc transporter ZIP10 precursor - Homo sapiens (Human)					
Q9ULF5	2	2.9417	K.LLTNLGLGER.K	1	ZIP10_HUMAN
Zyxin - Homo sapiens (Human)					
Q15942	2	5.4847	R.PSPAISVSVSAPAFYAPQKK.F	3	ZYX_HUMAN
Q15942	2	3.6513	K.FGPVVAPKPK.V	5	ZYX_HUMAN
Q15942	2	3.7649	K.VNPFRPGDSEPPPAPGAQR.A	4	ZYX_HUMAN
Q15942	3	4.8796	R.EKVSSIDLEIDSLSSLLDDM#TKNDPFK.A	1	ZYX_HUMAN
Q15942	2	4.1548	R.PSPAISVSVSAPAFYAPQK.K	4	ZYX_HUMAN