

**Supplementary Table 3A**

A list of 1,370 peptides identified using PPS-based approach, sorted by protein name.

MS/MS spectra were searched against the UniProt Human proteomic database; release 09/07.

\* SEQUEST Xcorr observed for an identified peptide; \*\* Human database accession number.

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>14-3-3 protein beta/alpha - Homo sapiens (Human)</i></b>					
	1433B_HUMAN	K.IEAELQDIC#NDVLELLDK.Y	2	2.4548	P31946
	1433B_HUMAN	K.IEAELQDIC#NDVLELLDK.Y	2	3.0147	P31946
	1433B_HUMAN	K.IEAELQDIC#NDVLELLDK.Y	2	4.3961	P31946
	1433B_HUMAN	K.TAFDEAIAELDTLNEESYK.D	2	5.4745	P31946
<b><i>14-3-3 protein epsilon - Homo sapiens (Human)</i></b>					
	1433E_HUMAN	K.AAFDDAIAELDTLSEESYK.D	2	3.9297	P62258
	1433E_HUMAN	K.AAFDDAIAELDTLSEESYK.D	2	2.4262	P62258
<b><i>14-3-3 protein gamma - Homo sapiens (Human)</i></b>					
	1433G_HUMAN	K.NVTELNEPLSNEER.N	2	2.6397	P61981
<b><i>14-3-3 protein zeta/delta - Homo sapiens (Human)</i></b>					
	1433Z_HUMAN	R.DIC#NDVLSLLEK.F	2	2.7905	P63104
	1433Z_HUMAN	K.TAFDEAIAELDTLSEESYK.D	2	2.7409	P63104
	1433Z_HUMAN	K.GIVDQSQQAYQEAFEISK.K	2	3.5216	P63104
	1433Z_HUMAN	K.SVTEQGAELSNEER.N	2	3.9548	P63104
	1433Z_HUMAN	R.DIC#NDVLSLLEK.F	2	2.6152	P63104
	1433Z_HUMAN	K.TAFDEAIAELDTLSEESYK.D	3	4.2294	P63104
	1433Z_HUMAN	K.SVTEQGAELSNEER.N	2	3.677	P63104
	1433Z_HUMAN	K.GIVDQSQQAYQEAFEISK.K	2	4.74	P63104
	1433Z_HUMAN	K.SVTEQGAELSNEER.N	2	3.8013	P63104
	1433Z_HUMAN	R.DIC#NDVLSLLEK.F	2	2.8059	P63104
	1433Z_HUMAN	K.SVTEQGAELSNEER.N	2	3.9712	P63104
	1433Z_HUMAN	R.DIC#NDVLSLLEK.F	2	3.0121	P63104
<b><i>2',3'-cyclic-nucleotide 3'-phosphodiesterase - Homo sapiens (Human)</i></b>					
	CN37_HUMAN	K.APGAE EYAQQDVLK.K	2	3.5784	P09543
	CN37_HUMAN	K.APGAE EYAQQDVLK.K	2	3.2862	P09543
<b><i>40S ribosomal protein S11 - Homo sapiens (Human)</i></b>					
	RS11_HUMAN	R.DVQIGDIVTVGEC#RPLSK.T	2	4.3733	P62280
	RS11_HUMAN	R.DVQIGDIVTVGEC#RPLSK.T	2	2.4041	P62280
<b><i>40S ribosomal protein S14 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	RS14_HUMAN	R.IEDVTPIPSDSTR.R	2	2.801	P62263
	RS14_HUMAN	R.IEDVTPIPSDSTR.R	2	2.8174	P62263
	RS14_HUMAN	K.TPGPGAQSALR.A	2	2.3556	P62263
<b><i>40S ribosomal protein S15 - Homo sapiens (Human)</i></b>					
	RS15_HUMAN	R.GVDLDQLLDMSYEQLMQLYSAR.Q	2	3.7298	P62841
<b><i>40S ribosomal protein S18 - Homo sapiens (Human)</i></b>					
	RS18_HUMAN	K.IPDWFLNR.Q	2	2.7426	P62269
	RS18_HUMAN	K.IPDWFLNR.Q	2	2.329	P62269
	RS18_HUMAN	R.AGELTEDEVER.V	2	2.3231	P62269
<b><i>40S ribosomal protein S20 - Homo sapiens (Human)</i></b>					
	RS20_HUMAN	K.TPVEPEVAIHR.I	3	3.9431	P60866
<b><i>40S ribosomal protein S3 - Homo sapiens (Human)</i></b>					
	RS3_HUMAN	R.ELAEDGYSGVEVR.V	2	2.8917	P23396
	RS3_HUMAN	K.GGKPEPPAMPQPVPPTA.-	2	2.6316	P23396
	RS3_HUMAN	K.GGKPEPPAM*PQPVPPTA.-	2	2.4171	P23396
	RS3_HUMAN	K.DEILPTTPISEQK.G	2	2.4715	P23396
	RS3_HUMAN	K.KPLPDHVSIVEPK.D	3	3.884	P23396
	RS3_HUMAN	K.GGKPEPPAMPQPVPPTA.-	2	2.9793	P23396
	RS3_HUMAN	R.ELAEDGYSGVEVR.V	2	2.9833	P23396
	RS3_HUMAN	R.ELAEDGYSGVEVR.V	2	2.876	P23396
<b><i>40S ribosomal protein S4, X isoform - Homo sapiens (Human)</i></b>					
	RS4X_HUMAN	R.HPGSFDVVHVK.D	3	3.9371	P62701
	RS4X_HUMAN	K.VNDTIQIDLETGK.I	2	2.8603	P62701
	RS4X_HUMAN	R.HPGSFDVVHVK.D	3	4.2664	P62701
	RS4X_HUMAN	R.HPGSFDVVHVK.D	3	4.3937	P62701
	RS4X_HUMAN	K.VNDTIQIDLETGK.I	2	2.5507	P62701
<b><i>40S ribosomal protein S5 - Homo sapiens (Human)</i></b>					
	RS5_HUMAN	K.TIAEC#LADELINAAK.G	2	4.0408	P46782
	RS5_HUMAN	K.TIAEC#LADELINAAK.G	2	5.5156	P46782
	RS5_HUMAN	K.TIAEC#LADELINAAK.G	2	2.8323	P46782
<b><i>4-trimethylaminobutyraldehyde dehydrogenase - Homo sapiens (Human)</i></b>					
	AL9A1_HUMAN	K.SGMERCRIILAAAR.I	2	2.317	P49189
<b><i>5-azacytidine-induced protein 1 - Homo sapiens (Human)</i></b>					
	AZI1_HUMAN	K.ETEKALSRQLQR.Q	1	2.325	Q9UPN4

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>60 kDa heat shock protein, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	CH60_HUMAN	K.TLNDELEIIEGMK.F	2	3.2207	P10809
<b><i>60S ribosomal protein L11 - Homo sapiens (Human)</i></b>					
	RL11_HUMAN	K.VLEQLTGQTPVFSK.A	2	3.2699	P62913
	RL11_HUMAN	K.VLEQLTGQTPVFSK.A	2	2.3965	P62913
	RL11_HUMAN	K.VLEQLTGQTPVFSK.A	2	2.7801	P62913
<b><i>60S ribosomal protein L12 - Homo sapiens (Human)</i></b>					
	RL12_HUMAN	R.C#TGGEVGSALAPK.I	2	3.3402	P30050
<b><i>60S ribosomal protein L18a - Homo sapiens (Human)</i></b>					
	RL18A_HUMAN	R.DLTTAGAVTQC#YR.D	2	2.5021	Q02543
	RL18A_HUMAN	R.DLTTAGAVTQC#YR.D	2	2.7058	Q02543
	RL18A_HUMAN	R.DLTTAGAVTQC#YR.D	2	2.3983	Q02543
<b><i>60S ribosomal protein L35a - Homo sapiens (Human)</i></b>					
	RL35A_HUMAN	K.NNTVTPGGKPNK.T	2	2.3165	P18077
<b><i>60S ribosomal protein L36 - Homo sapiens (Human)</i></b>					
	RL36_HUMAN	R.EELSNVLAAMR.K	2	2.381	Q9Y3U8
<b><i>60S ribosomal protein L6 - Homo sapiens (Human)</i></b>					
	RL6_HUMAN	R.HQEGEIFDTEK.E	2	3.2719	Q02878
	RL6_HUMAN	K.HLTDAYFK.K	2	2.3575	Q02878
	RL6_HUMAN	R.HQEGEIFDTEK.E	2	3.0324	Q02878
	RL6_HUMAN	R.HQEGEIFDTEK.E	2	3.3082	Q02878
	RL6_HUMAN	R.HQEGEIFDTEK.E	2	3.5256	Q02878
<b><i>60S ribosomal protein L7 - Homo sapiens (Human)</i></b>					
	RL7_HUMAN	K.YGIIC#MEDLIHEIYTVGK.R	3	4.0229	P18124
	RL7_HUMAN	K.YGIIC#MEDLIHEIYTVGK.R	3	3.8383	P18124
<b><i>60S ribosomal protein L7a - Homo sapiens (Human)</i></b>					
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	2.3161	P62424
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	3.4862	P62424
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	3.502	P62424
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	3.6929	P62424
<b><i>Acetyl-CoA carboxylase 1 - Homo sapiens (Human)</i></b>					
	COA1_HUMAN	R.ESRGSVLEPEGTVEIK.F	2	2.6704	Q13085
<b><i>Actin, aortic smooth muscle - Homo sapiens (Human)</i></b>					
	ACTA_HUMAN	K.DSYVGDEAQS.K	2	2.5043	P62736

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ACTA_HUMAN		K.LC#YVALDFENEM*ATAASSSSLEK.S	2	2.9462	P62736
ACTA_HUMAN		K.YPIEHGIITNWDDMEK.I	2	2.5817	P62736
ACTA_HUMAN		K.YPIEHGIITNWDDMEK.I	2	3.4953	P62736
ACTA_HUMAN		K.SYELPDGQVITIGNER.F	2	4.1844	P62736
ACTA_HUMAN		K.SYELPDGQVITIGNER.F	2	2.6786	P62736
ACTA_HUMAN		K.SYELPDGQVITIGNER.F	2	4.3354	P62736
ACTA_HUMAN		K.SYELPDGQVITIGNER.F	2	2.5062	P62736
ACTA_HUMAN		R.DLTDYLMK.I	1	2.0545	P62736
ACTA_HUMAN		K.EITALAPSTMK.I	2	2.3757	P62736
ACTA_HUMAN		K.EITALAPSTMK.I	2	2.3554	P62736
ACTA_HUMAN		K.DSYVGDEAQS.K	2	2.4682	P62736
ACTA_HUMAN		K.YPIEHGIITNWDDMEK.I	2	3.3464	P62736
<b><i>Actin, cytoplasmic 1 - Homo sapiens (Human)</i></b>					
ACTB_HUMAN		R.TTGIVM*DSGDGVTHTVPIYEGYALPHAILR.L	3	4.4499	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	4.7415	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	3.5094	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	2.7066	P60709
ACTB_HUMAN		R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	3	5.1798	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	2.3717	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	3.4975	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	5.4905	P60709
ACTB_HUMAN		K.LC#YVALDFEQEMATAASSSSLEK.S	2	4.0558	P60709
ACTB_HUMAN		R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	3	5.087	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	4.9018	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	3.6442	P60709
ACTB_HUMAN		R.C#PEALFQPSFLGMESC#GIHETTFNSIMK.C	3	3.9267	P60709
ACTB_HUMAN		R.VAPEEHPVLLTEAPLNPK.A	2	2.9046	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	5.555	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	2.5227	P60709
<b><i>Adenylyl cyclase-associated protein 1 - Homo sapiens (Human)</i></b>					
CAP1_HUMAN		K.AGAAPYVQAFDSSLAGPVAEYLK.I	2	2.659	Q01518
CAP1_HUMAN		R.VENQENVSNLVIETELK.Q	2	3.8068	Q01518
CAP1_HUMAN		R.VENQENVSNLVIETELK.Q	2	3.1727	Q01518
<b><i>ADP/ATP translocase 2 - Homo sapiens (Human)</i></b>					
ADT2_HUMAN		K.DFLAGGVA AISK.T	2	2.6675	P05141

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	ADT2_HUMAN	K.DFLAGGVAAAISK.T	1	3.0012	P05141
	ADT2_HUMAN	K.DFLAGGVAAAISK.T	2	4.2384	P05141
	ADT2_HUMAN	K.DFLAGGVAAAISK.T	2	2.5456	P05141
	ADT2_HUMAN	K.DFLAGGVAAAISK.T	2	3.8569	P05141
	ADT2_HUMAN	K.DFLAGGVAAAISK.T	2	3.5177	P05141
<b><i>ADP/ATP translocase 3 - Homo sapiens (Human)</i></b>					
	ADT3_HUMAN	K.DFLAGGIAAAISK.T	2	3.2152	P12236
	ADT3_HUMAN	K.DFLAGGIAAAISK.T	2	3.9418	P12236
	ADT3_HUMAN	K.DFLAGGIAAAISK.T	2	3.7957	P12236
	ADT3_HUMAN	K.DFLAGGIAAAISK.T	2	4.1938	P12236
<b><i>ADP-ribosylation factor 1 - Homo sapiens (Human)</i></b>					
	ARF1_HUMAN	K.QDLPNAMNAEITDK.L	2	3.029	P84077
<b><i>A-kinase anchor protein 9 - Homo sapiens (Human)</i></b>					
	AKAP9_HUMAN	K.QMKEKENDLQEK.F	2	2.4166	Q99996
<b><i>Anaphase-promoting complex subunit 4 - Homo sapiens (Human)</i></b>					
	ANC4_HUMAN	R.MENIIDQC#LQKPADVIGK.S	2	2.6898	Q9UJX5
<b><i>Ankyrin-2 - Homo sapiens (Human)</i></b>					
	ANK2_HUMAN	K.M*ADEQGDMDLQISPRK.T	2	2.3087	Q01484
<b><i>Annexin A1 - Homo sapiens (Human)</i></b>					
	ANXA1_HUMAN	K.TPAQFDADEL.R.A	2	3.1619	P04083
	ANXA1_HUMAN	K.GVDEATIIDLTK.R	2	3.5907	P04083
	ANXA1_HUMAN	K.GLGTDEDTLIEILASR.T	2	3.5575	P04083
	ANXA1_HUMAN	K.TPAQFDADEL.R.A	2	3.1115	P04083
	ANXA1_HUMAN	R.SEDFGVNEDLADSDAR.A	2	2.8906	P04083
	ANXA1_HUMAN	K.GVDEATIIDLTK.R	2	2.9551	P04083
	ANXA1_HUMAN	K.GLGTDEDTLIEILASR.T	2	3.6734	P04083
	ANXA1_HUMAN	K.GLGTDEDTLIEILASR.T	2	3.657	P04083
<b><i>Annexin A11 - Homo sapiens (Human)</i></b>					
	ANX11_HUMAN	K.TPVLFDIYEIK.E	2	2.4618	P50995
	ANX11_HUMAN	R.DAQELYAAGENR.L	2	2.6636	P50995
	ANX11_HUMAN	K.GFGTDEQAIIDC#LGSR.S	2	2.9791	P50995
	ANX11_HUMAN	K.GVGTDEAC#LIEILASR.S	2	2.8677	P50995
	ANX11_HUMAN	K.GFGTDEQAIIDC#LGSR.S	2	2.4671	P50995
<b><i>Annexin A2 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.5016	P07355
ANXA2_HUMAN		K.TDLEKDIISDTSGDFR.K	2	4.6437	P07355
ANXA2_HUMAN		R.AEDGSVIDYELIDQDAR.D	2	2.6684	P07355
ANXA2_HUMAN		R.AEDGSVIDYELIDQDAR.D	2	3.6235	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.357	P07355
ANXA2_HUMAN		K.TDLEKDIISDTSGDFR.K	2	3.4621	P07355
ANXA2_HUMAN		R.TNQELQEINR.V	2	2.9638	P07355
ANXA2_HUMAN		K.TDLEKDIISDTSGDFR.K	2	4.1629	P07355
ANXA2_HUMAN		R.RAEDGSVIDYELIDQDAR.D	2	3.9007	P07355
ANXA2_HUMAN		R.RAEDGSVIDYELIDQDAR.D	3	3.9698	P07355
ANXA2_HUMAN		R.DALNIETAIK.T	2	2.6988	P07355
ANXA2_HUMAN		R.RAEDGSVIDYELIDQDAR.D	2	4.6405	P07355
ANXA2_HUMAN		K.GLGTDEDSLIEIIC#SR.T	2	3.3708	P07355
ANXA2_HUMAN		K.GLGTDEDSLIEIIC#SR.T	2	2.8156	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.6321	P07355
ANXA2_HUMAN		K.TDLEKDIISDTSGDFR.K	2	3.6445	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.1602	P07355
ANXA2_HUMAN		R.DALNIETAIK.T	2	2.755	P07355
ANXA2_HUMAN		K.SLYYYIQDQTK.G	2	2.613	P07355
ANXA2_HUMAN		R.RAEDGSVIDYELIDQDAR.D	2	4.0154	P07355
ANXA2_HUMAN		K.TDLEKDIISDTSGDFR.K	2	4.0158	P07355

***Annexin A6 - Homo sapiens (Human)***

ANXA6_HUMAN		K.SLHQAIEGDTSGDFLK.A	2	2.4386	P08133
ANXA6_HUMAN		R.DLEADIIGDTSGHFQK.M	2	2.8435	P08133

***ANXA4 protein - Homo sapiens (Human)***

Q6LES2_HUMA		K.SETSGSFEDALLAIVK.C	2	2.486	Q6LES2
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***Apoptosis regulator BAX, membrane isoform alpha - Homo sapiens (Human)***

BAXA_HUMAN		R.MGGEAPELALDPVPQDASTK.K	2	2.7284	Q07812
BAXA_HUMAN		R.MGGEAPELALDPVPQDASTK.K	2	3.1274	Q07812
BAXA_HUMAN		R.IGDELDSNMELQR.M	2	3.7503	Q07812
BAXA_HUMAN		R.MGGEAPELALDPVPQDASTK.K	2	3.701	Q07812
BAXA_HUMAN		R.MGGEAPELALDPVPQDASTK.K	2	3.2272	Q07812
BAXA_HUMAN		R.IGDELDSNMELQR.M	2	4.0991	Q07812
BAXA_HUMAN		R.MGGEAPELALDPVPQDASTK.K	2	2.9013	Q07812
BAXA_HUMAN		R.IGDELDSNM*ELQR.M	2	3.2652	Q07812

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Apoptosis-inducing factor 3 - Homo sapiens (Human)</i></b>					
AIFM3_HUMAN		R.GRNVVVVGAGFLGMEVAAYLTEK.A	2	2.6413	Q96NN9
AIFM3_HUMAN		R.GRNVVVVGAGFLGMEVAAYLTEK.A	2	2.3939	Q96NN9
<b><i>Astrotactin 2 - Homo sapiens (Human)</i></b>					
A2A2T7_HUMAN		K.M*VSM*ARNTYYLTLISKVSPF.-	2	2.3168	A2A2T7
<b><i>Atlastin-3 - Homo sapiens (Human)</i></b>					
ATLA3_HUMAN		R.YQQELEEEIKELYENFC#K.H	2	4.3985	Q6DD88
ATLA3_HUMAN		R.YQQELEEEIKELYENFC#K.H	2	5.517	Q6DD88
ATLA3_HUMAN		R.YQQELEEEIKELYENFC#K.H	2	4.3234	Q6DD88
ATLA3_HUMAN		R.YQQELEEEIKELYENFC#K.H	3	4.1117	Q6DD88
ATLA3_HUMAN		K.SMLQATAEANNLAAAASAK.D	2	3.7073	Q6DD88
ATLA3_HUMAN		R.YQQELEEEIKELYENFC#K.H	3	4.1868	Q6DD88
<b><i>ATP synthase B chain, mitochondrial precursor - Homo sapiens (Human)</i></b>					
AT5F1_HUMAN		K.HVVQSISTQKEK.E	2	2.7956	P24539
AT5F1_HUMAN		K.TGVTGPYVLGTGLILYALSK.E	2	3.697	P24539
AT5F1_HUMAN		K.TGVTGPYVLGTGLILYALSK.E	2	3.651	P24539
AT5F1_HUMAN		K.TGVTGPYVLGTGLILYALSK.E	2	3.6379	P24539
AT5F1_HUMAN		K.TGVTGPYVLGTGLILYALSK.E	2	4.2992	P24539
AT5F1_HUMAN		K.HVVQSISTQKEK.E	2	3.7883	P24539
AT5F1_HUMAN		K.HVVQSISTQKEK.E	2	3.7649	P24539
AT5F1_HUMAN		K.HVVQSISTQKEK.E	2	3.8684	P24539
<b><i>ATP synthase D chain, mitochondrial - Homo sapiens (Human)</i></b>					
ATP5H_HUMAN		K.YPYWPHQPIENL.-	2	3.0641	O75947
ATP5H_HUMAN		K.YPYWPHQPIENL.-	2	3.4386	O75947
ATP5H_HUMAN		K.YTAQVDAEEKEDVK.S	2	2.5835	O75947
ATP5H_HUMAN		K.YTAQVDAEEKEDVK.S	2	2.6866	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	2.4828	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	5.1207	O75947
ATP5H_HUMAN		K.YPYWPHQPIENL.-	2	3.5086	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	3.1527	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	4.95	O75947
ATP5H_HUMAN		K.YPYWPHQPIENL.-	2	3.7376	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	5.0316	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	2.4485	O75947
ATP5H_HUMAN		K.NLIPFDQM*TIEDLNEAFPETK.L	2	2.7713	O75947

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>ATP synthase f chain, mitochondrial - Homo sapiens (Human)</i></b>					
ATPK_HUMAN		R.DFSPSGIFGAFQR.G	2	2.8601	P56134
ATPK_HUMAN		R.DFSPSGIFGAFQR.G	2	2.4191	P56134
<b><i>ATP synthase subunit alpha, mitochondrial precursor - Homo sapiens (Human)</i></b>					
ATPA_HUMAN		R.TGAIVDVPVGEELLGR.V	2	3.0531	P25705
ATPA_HUMAN		R.TGAIVDVPVGEELLGR.V	2	2.5581	P25705
ATPA_HUMAN		R.ILGADTSVDLEETGR.V	2	3.4327	P25705
ATPA_HUMAN		R.EVAАFAQFGSDLDAATQQLLSR.G	3	4.2238	P25705
ATPA_HUMAN		K.TGTAEMSSILEER.I	2	2.338	P25705
ATPA_HUMAN		R.EVAАFAQFGSDLDAATQQLLSR.G	2	4.2796	P25705
ATPA_HUMAN		R.EVAАFAQFGSDLDAATQQLLSR.G	2	5.7132	P25705
ATPA_HUMAN		R.EVAАFAQFGSDLDAATQQLLSR.G	2	5.2359	P25705
ATPA_HUMAN		K.QGQYSPMAIEEQVAVIYAGVR.G	2	2.7988	P25705
ATPA_HUMAN		K.FENAFLSHVVSQHQALLGTIR.A	3	4.5791	P25705
ATPA_HUMAN		R.ILGADTSVDLEETGR.V	2	3.5817	P25705
ATPA_HUMAN		K.TGTAEMSSILEER.I	2	2.9482	P25705
ATPA_HUMAN		K.TGTAEMSSILEER.I	2	2.6233	P25705
ATPA_HUMAN		K.TGTAEMSSILEER.I	2	2.8654	P25705
<b><i>ATP synthase subunit beta, mitochondrial precursor - Homo sapiens (Human)</i></b>					
ATPB_HUMAN		R.AIAELGIYPAVDPLDSTSR.I	2	2.6942	P06576
ATPB_HUMAN		R.DQEGQDVLLFIDNIFR.F	2	2.7577	P06576
ATPB_HUMAN		R.AIAELGIYPAVDPLDSTSR.I	2	2.8058	P06576
ATPB_HUMAN		R.DQEGQDVLLFIDNIFR.F	2	4.9077	P06576
ATPB_HUMAN		R.AIAELGIYPAVDPLDSTSR.I	2	2.5221	P06576
ATPB_HUMAN		R.AIAELGIYPAVDPLDSTSR.I	2	2.9823	P06576
ATPB_HUMAN		R.DQEGQDVLLFIDNIFR.F	2	4.5136	P06576
ATPB_HUMAN		R.DQEGQDVLLFIDNIFR.F	2	3.5215	P06576
<b><i>ATP synthase-coupling factor 6, mitochondrial precursor - Homo sapiens (Human)</i></b>					
ATP5J_HUMAN		R.QTSGGPVDASSEYQQELER.E	2	2.9976	P18859
<b><i>ATP-binding cassette sub-family D member 1 - Homo sapiens (Human)</i></b>					
ABCD1_HUMAN		R.GLQAPAGEPTQEASGVAAAK.A	2	3.3297	P33897
<b><i>ATP-dependent DNA helicase 2 subunit 1 - Homo sapiens (Human)</i></b>					
KU70_HUMAN		R.DIISIAEDEDLR.V	2	2.5948	P12956
<b><i>ATP-dependent DNA helicase 2 subunit 2 - Homo sapiens (Human)</i></b>					



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
KU86_HUMAN		R.HLM*LPDFDLLEDIESK.I	2	2.7409	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	2	2.3713	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	2	3.0779	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	2	4.1663	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	3	5.0778	P13010
KU86_HUMAN		R.HLM*LPDFDLLEDIESK.I	2	2.7511	P13010
KU86_HUMAN		K.YAPTEAQLNAVDALIDSMFLAK.K	3	4.2559	P13010
KU86_HUMAN		K.YAPTEAQLNAVDALIDSMFLAK.K	2	3.7149	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	2	3.4906	P13010
KU86_HUMAN		R.HLMLPDFDLLEDIESK.I	2	3.6208	P13010
KU86_HUMAN		R.HLM*LPDFDLLEDIESK.I	2	3.4466	P13010
KU86_HUMAN		R.HLM*LPDFDLLEDIESK.I	2	2.7382	P13010
<b><i>ATP-dependent RNA helicase A - Homo sapiens (Human)</i></b>					
DHX9_HUMAN		K.ETPFELIEALLK.Y	2	2.7022	Q08211
DHX9_HUMAN		K.ETPFELIEALLK.Y	2	2.6314	Q08211
DHX9_HUMAN		K.ETPFELIEALLK.Y	2	2.9501	Q08211
DHX9_HUMAN		K.ETPFELIEALLK.Y	2	2.4901	Q08211
<b><i>ATP-dependent RNA helicase DDX3X - Homo sapiens (Human)</i></b>					
DDX3X_HUMAN		K.DLLDLLVEAK.Q	2	2.3684	O00571
<b><i>Bcl-2-like 13 protein - Homo sapiens (Human)</i></b>					
B2L13_HUMAN		K.SSPATSLFVELDEEEVK.A	2	2.5705	Q9BXX5
B2L13_HUMAN		R.EESLVEELSPASEK.K	2	2.9715	Q9BXX5
B2L13_HUMAN		R.EESLVEELSPASEK.K	2	2.8184	Q9BXX5
<b><i>BH3-interacting domain death agonist - Homo sapiens (Human)</i></b>					
BID_HUMAN		R.DLATALEQLLQAYPR.D	3	4.0411	P55957
BID_HUMAN		R.IEADSESQEDIIR.N	2	2.6668	P55957
BID_HUMAN		R.DLATALEQLLQAYPR.D	3	4.1203	P55957
BID_HUMAN		R.HLAQVGDSDMDR.S	2	2.6184	P55957
<b><i>Brain acid soluble protein 1 - Homo sapiens (Human)</i></b>					
BASP_HUMAN		K.AQGPAASAEEPKPVEAPAANSQQTVTVKE.-	3	5.4931	P80723
BASP_HUMAN		K.AQGPAASAEEPKPVEAPAANSQQTVTVKE	3	3.8532	P80723
BASP_HUMAN		K.AQGPAASAEEPKPVEAPAANSQQTVTVKE.-	3	3.8979	P80723
BASP_HUMAN		K.AQGPAASAEEPKPVEAPAANSQQTVTVKE.-	3	4.8189	P80723
<b><i>BRI3-binding protein - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	BRI3B_HUMAN	R.SSPSGPSNPSNPSVEEK.L	2	2.748	Q8WY22
	BRI3B_HUMAN	R.SSPSGPSNPSNPSVEEK.L	2	3.9403	Q8WY22
	BRI3B_HUMAN	R.SSPSGPSNPSNPSVEEK.L	2	4.0793	Q8WY22
<b><i>CAAX prenyl protease 1 homolog - Homo sapiens (Human)</i></b>					
	FACE1_HUMAN	K.DIQEDSGMEPR.N	2	2.6355	O75844
<b><i>Calcium-binding protein p22 - Homo sapiens (Human)</i></b>					
	CHP1_HUMAN	R.IPELAINPLGDR.I	2	2.9043	Q99653
<b><i>Calnexin precursor - Homo sapiens (Human)</i></b>					
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.7897	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.5832	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	3.5884	P27824
	CALX_HUMAN	K.SDAEEDGGTVSQEEEDR.K	2	3.8687	P27824
	CALX_HUMAN	K.TPELNLDQFHDK.T	2	3.8663	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.4758	P27824
	CALX_HUMAN	R.PVIDNPNYK.G	2	2.5081	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	4.2483	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	4.2637	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.6822	P27824
	CALX_HUMAN	R.PVIDNPNYK.G	2	2.5179	P27824
	CALX_HUMAN	K.TPELNLDQFHDK.T	3	3.8899	P27824
	CALX_HUMAN	K.QKSDAEEDGGTVSQEEEDR.K	3	3.8138	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.3884	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	2.9827	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.997	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.4259	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	4.2451	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	4.1753	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	4.4318	P27824
	CALX_HUMAN	R.PVIDNPNYK.G	2	2.306	P27824
<b><i>Carnitine O-palmitoyltransferase I, liver isoform - Homo sapiens (Human)</i></b>					
	CPT1A_HUMAN	R.ILDNTSEPQPGEAR.L	2	2.655	P50416
<b><i>Caspase-1 precursor - Homo sapiens (Human)</i></b>					
	CASP1_HUMAN	R.SMGEGTINGLLDELLQTR.V	2	2.8942	P29466
<b><i>Catechol O-methyltransferase - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	COMT_HUMAN	R.ILNHVLQHAEPGNAQSVLEAIDTYC#EQK.E	3	4.5786	P21964
<b><i>Cathepsin G precursor - Homo sapiens (Human)</i></b>					
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	4.1229	P08311
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	3.0831	P08311
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	3.9247	P08311
<b><i>Cation-independent mannose-6-phosphate receptor precursor - Homo sapiens (Human)</i></b>					
	MPRI_HUMAN	K.VPIDGPPIDIGR.V	2	2.5405	P11717
<b><i>CD302 antigen precursor - Homo sapiens (Human)</i></b>					
	CD302_HUMAN	K.GPDDILLGMFYDTHDASFK.W	2	2.4584	Q8IX05
<b><i>CD44 antigen precursor - Homo sapiens (Human)</i></b>					
	CD44_HUMAN	K.ESSETPDQFMTADETR.N	2	3.4633	P16070
	CD44_HUMAN	K.ESSETPDQFM*TADETR.N	2	2.9127	P16070
	CD44_HUMAN	K.ESSETPDQFMTADETR.N	2	3.7876	P16070
<b><i>CD97 antigen precursor - Homo sapiens (Human)</i></b>					
	CD97_HUMAN	R.HLIATQLLSNLEDIMR.I	2	3.3934	P48960
	CD97_HUMAN	R.HLIATQLLSNLEDIMR.I	2	3.4195	P48960
	CD97_HUMAN	K.LVDELM*EAPGDVEALAPPVR.H	2	2.8974	P48960
<b><i>CDNA FLJ30700 fis, clone FCBBF2000940 - Homo sapiens (Human)</i></b>					
	Q96NK0_HUMA	K.AFLHSLLAFSVAREKTFHF.-	2	2.4159	Q96NK0
<b><i>Cell cycle control protein 50A - Homo sapiens (Human)</i></b>					
	CC50A_HUMAN	R.RPDNTAFK.Q	2	2.6749	Q9NV96
	CC50A_HUMAN	K.DEVDGGPPC#APGGTAK.T	2	2.3191	Q9NV96
<b><i>Cellular tumor antigen p53 - Homo sapiens (Human)</i></b>					
	P53_HUMAN	R.VEYLDDR.N	2	2.4083	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	3.0866	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	2.6459	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	2.803	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	2.8386	P04637
	P53_HUMAN	R.RTEENLR.K	2	2.6067	P04637
<b><i>Centromeric protein E - Homo sapiens (Human)</i></b>					
	CENPE_HUMAN	K.FQESQEEIKSLTK.E	2	2.4465	Q02224
<b><i>Centrosomal protein of 152 kDa - Homo sapiens (Human)</i></b>					
	CE152_HUMAN	R.QMVQDFDHDKQEAVIDR.C	2	2.3151	O94986
<b><i>Clathrin heavy chain 1 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
CLH1_HUMAN		R.LPVVIGLLDVDC#SEDVIK.N	2	3.5112	Q00610
CLH1_HUMAN		R.GQFSTDELVAEVEK.R	2	2.4377	Q00610
CLH1_HUMAN		R.GQFSTDELVAEVEK.R	2	2.9993	Q00610
CLH1_HUMAN		R.GQFSTDELVAEVEK.R	2	2.7256	Q00610
CLH1_HUMAN		K.SVNESLNNLFITEEDYQALR.T	2	2.4704	Q00610
CLH1_HUMAN		K.DTELAEEELLQWFLQEEK.R	2	3.0965	Q00610
CLH1_HUMAN		K.DTELAEEELLQWFLQEEK.R	2	3.9757	Q00610
CLH1_HUMAN		K.AFM*TADLPNELIELLEK.I	2	2.8367	Q00610
CLH1_HUMAN		R.LPVVIGLLDVDC#SEDVIK.N	2	3.9852	Q00610
CLH1_HUMAN		K.SVNESLNNLFITEEDYQALR.T	2	2.5045	Q00610
<b><i>Coatomer subunit gamma - Homo sapiens (Human)</i></b>					
COPG_HUMAN		R.AIVDC#IISIIENSESEK.E	2	2.7527	Q9Y678
<b><i>Cofilin-1 - Homo sapiens (Human)</i></b>					
COF1_HUMAN		K.HELQANC#YEEVKDR.C	3	4.6603	P23528
<b><i>Cohesin subunit SA-2 - Homo sapiens (Human)</i></b>					
STAG2_HUMAN		K.YYNDYGDIKETM*SKTR.Q	2	2.5314	Q8N3U4
<b><i>Coiled-coil domain-containing protein 47 precursor - Homo sapiens (Human)</i></b>					
CCD47_HUMAN		K.YGLPDSLAILSEMGEVTDGMMDTK.M	2	3.3195	Q96A33
<b><i>Core histone macro-H2A.1 - Homo sapiens (Human)</i></b>					
H2AY_HUMAN		K.AASADSTTEGTPADGFTVLSTK.S	2	2.4287	O75367
<b><i>Core histone macro-H2A.2 - Homo sapiens (Human)</i></b>					
H2AW_HUMAN		R.HILLAVANDEELNQLLK.G	2	3.2287	Q9P0M6
H2AW_HUMAN		R.HILLAVANDEELNQLLK.G	2	3.7567	Q9P0M6
H2AW_HUMAN		R.HILLAVANDEELNQLLK.G	2	2.9756	Q9P0M6
<b><i>Coronin-7 - Homo sapiens (Human)</i></b>					
CORO7_HUMAN		R.VPAEGLEEVLTTPETVLTGHTEK.I	3	4.5015	P57737
<b><i>Cytochrome b reductase 1 - Homo sapiens (Human)</i></b>					
Q53TN4_HUMA		R.NLALDEAGQR.S	2	2.4976	Q53TN4
<b><i>Cytochrome b-245 heavy chain - Homo sapiens (Human)</i></b>					
CY24B_HUMAN		R.IVGDWTEGLFNAC#GC#DK.Q	2	2.8523	P04839
CY24B_HUMAN		R.IGVFLC#GPEALAETLSK.Q	2	4.9213	P04839
CY24B_HUMAN		R.IVGDWTEGLFNAC#GC#DK.Q	2	3.2526	P04839
CY24B_HUMAN		R.IGVFLC#GPEALAETLSK.Q	2	2.8403	P04839
CY24B_HUMAN		R.IGVFLC#GPEALAETLSK.Q	2	4.9155	P04839

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	CY24B_HUMAN	R.IVGDWTEGLFNAC#GC#DK.Q	2	2.6739	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	2.4464	P04839
<b><i>Cytochrome b-245 light chain - Homo sapiens (Human)</i></b>					
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	2	4.3861	P13498
	CY24A_HUMAN	K.HMTAVVK.L	2	2.4019	P13498
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	2	4.4304	P13498
	CY24A_HUMAN	K.HMTAVVK.L	2	2.5863	P13498
<b><i>Cytochrome b5 type B precursor - Homo sapiens (Human)</i></b>					
	CYB5B_HUMAN	R.FLNEHPGGEEVLLAQAGVDASESFEDVGHSSDAR.	3	4.6121	O43169
<b><i>Cytochrome c oxidase subunit 4 isoform 1, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.7377	P13073
	COX41_HUMAN	R.DHPLPEVAHVK.H	3	3.7968	P13073
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.8282	P13073
	COX41_HUMAN	R.DHPLPEVAHVK.H	3	4.1384	P13073
	COX41_HUMAN	R.DHPLPEVAHVK.H	3	3.916	P13073
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.8457	P13073
<b><i>Cytochrome c oxidase subunit 5A, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	COX5A_HUMAN	K.GINTLVITYDMVPEPK.I	2	4.1561	P20674
	COX5A_HUMAN	K.GINTLVITYDM*VPEPK.I	2	2.9667	P20674
	COX5A_HUMAN	K.GINTLVITYDM*VPEPK.I	2	2.509	P20674
	COX5A_HUMAN	K.GINTLVITYDM*VPEPK.I	2	2.8511	P20674
	COX5A_HUMAN	K.GINTLVITYDMVPEPK.I	2	4.5986	P20674
<b><i>Cytochrome c oxidase subunit 5B, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	COX5B_HUMAN	R.EDPNLVPSISNK.R	2	2.3091	P10606
<b><i>Cytoskeleton-associated protein 4 - Homo sapiens (Human)</i></b>					
	CKAP4_HUMAN	K.IETNENNLESAG.G	2	2.591	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	3.3494	Q07065
	CKAP4_HUMAN	K.VQEQVHTLLSQDQAQAAR.L	2	4.6148	Q07065
	CKAP4_HUMAN	R.ERDFTSLENTVEER.L	2	3.8402	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	3.2866	Q07065
	CKAP4_HUMAN	K.VASLEESEGNKQDLK.A	2	2.6076	Q07065
	CKAP4_HUMAN	R.DFTSLENTVEER.L	2	2.6626	Q07065
	CKAP4_HUMAN	R.SVGELPSTVESLQK.V	2	3.8083	Q07065
	CKAP4_HUMAN	R.SVGELPSTVESLQK.V	2	2.3053	Q07065

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
CKAP4_HUMAN		R.EELGQGLQGVEQK.V	2	3.048	Q07065
CKAP4_HUMAN		K.SSSSSASAAAAAASSSASC#SR.R	2	2.5436	Q07065
CKAP4_HUMAN		R.SVGELPSTVESLQK.V	2	2.555	Q07065
CKAP4_HUMAN		R.ERDFTSLENTVEER.L	2	3.8979	Q07065
CKAP4_HUMAN		R.LEGLGSSEADQDGLASTVR.S	2	4.0283	Q07065
CKAP4_HUMAN		R.EELGQGLQGVEQK.V	2	3.4478	Q07065
CKAP4_HUMAN		R.SVGELPSTVESLQK.V	2	3.4329	Q07065
CKAP4_HUMAN		R.LEGLGSSEADQDGLASTVR.S	2	4.0553	Q07065
CKAP4_HUMAN		R.SVGELPSTVESLQK.V	2	3.13	Q07065
CKAP4_HUMAN		R.DFTSLENTVEER.L	2	3.0342	Q07065
CKAP4_HUMAN		K.DLSDGIHVVK.D	2	2.3935	Q07065
<b><i>Cytoskeleton-associated protein 5 - Homo sapiens (Human)</i></b>					
CKAP5_HUMAN		K.DQVLAMLEKAK.V	2	2.6213	Q14008
<b><i>DC2 - Homo sapiens (Human)</i></b>					
Q9NRP0_HUMA		R.VPFLVLEC#PNLK.L	2	3.6637	Q9NRP0
Q9NRP0_HUMA		R.VPFLVLEC#PNLK.L	2	3.5747	Q9NRP0
Q9NRP0_HUMA		R.VPFLVLEC#PNLK.L	2	4.0059	Q9NRP0
<b><i>Dehydrogenase/reductase SDR family member 8 precursor - Homo sapiens (Human)</i></b>					
DHRS8_HUMAN		K.NPSTSLGPTLEPEEVNR.L	2	3.6444	Q8NBQ5
DHRS8_HUMAN		K.NPSTSLGPTLEPEEVNR.L	2	4.0758	Q8NBQ5
DHRS8_HUMAN		K.HGLEETAAK.C	2	2.6903	Q8NBQ5
<b><i>Dermatan-sulfate epimerase precursor - Homo sapiens (Human)</i></b>					
DSE_HUMAN		K.HDLAASC#QGRVVAAEEKNGVVFIR.G	2	2.3118	Q9UL01
<b><i>Dihydroorotate dehydrogenase, mitochondrial precursor - Homo sapiens (Human)</i></b>					
PYRD_HUMAN		R.VPIIGVGGVSSGQDALEK.I	2	3.0462	Q02127
<b><i>Dihydropyrimidinase - Homo sapiens (Human)</i></b>					
DPYS_HUMAN		K.VVYEAGVFSVTAGDGK.F	2	2.3684	Q14117
DPYS_HUMAN		K.VVYEAGVFSVTAGDGK.F	2	2.4748	Q14117
<b><i>DNA fragmentation factor subunit alpha - Homo sapiens (Human)</i></b>					
DFFA_HUMAN		K.KTETVQEACERELALR.L	2	2.4816	O00273
<b><i>DNA topoisomerase 2-beta - Homo sapiens (Human)</i></b>					
TOP2B_HUMAN		R.HVDYVVDQVVGK.L	2	2.575	Q02880
<b><i>DNA-dependent protein kinase catalytic subunit - Homo sapiens (Human)</i></b>					
PRKDC_HUMAN		R.MEVQEQEEDISSLIR.S	2	2.4586	P78527

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	PRKDC_HUMAN	K.NLSSNEAISLEEIR.I	2	2.772	P78527
	PRKDC_HUMAN	K.LTLPEDNSMNVDDQGDPSDR.M	2	3.4127	P78527
<b><i>DNA-directed RNA polymerase III subunit RPC3 - Homo sapiens (Human)</i></b>					
	RPC3_HUMAN	K.LDASEIQVDETIFLLESYIEC#TM*KR.Q	2	2.342	Q9BU14
<b><i>DnaJ homolog subfamily C member 13 - Homo sapiens (Human)</i></b>					
	DNJCD_HUMAN	R.LGGYLAEEQATPENPTIR.K	2	2.8644	O75165
<b><i>Dolichol-phosphate mannosyltransferase subunit 3 - Homo sapiens (Human)</i></b>					
	DPM3_HUMAN	R.ELQSQIQEAR.A	2	2.3657	Q9P2X0
<b><i>Dolichyl-diphosphooligosaccharide-protein glycosyltransferase - Homo sapiens (Human)</i></b>					
	Q5VWA5_HUMA	K.LPDVYGVFQFK.V	2	3.1129	Q5VWA5
	Q5VWA5_HUMA	K.LPDVYGVFQFK.V	2	2.8431	Q5VWA5
	Q5VWA5_HUMA	K.LPDVYGVFQFK.V	2	2.6395	Q5VWA5
<b><i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 63 kDa subunit precursor - Ho</i></b>					
	RIB2_HUMAN	R.SIVEEIEDLVAR.L	2	3.037	P04844
	RIB2_HUMAN	K.TGQEVVFAEPDNK.N	2	3.3161	P04844
	RIB2_HUMAN	K.FPEEEAPSTVLSQNLFTPK.Q	2	2.931	P04844
	RIB2_HUMAN	K.TGQEVVFAEPDNK.N	2	3.3313	P04844
<b><i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 67 kDa subunit precursor - Ho</i></b>					
	RIB1_HUMAN	K.NLVEQHIQDIVVHYTFNK.V	3	5.0156	P04843
	RIB1_HUMAN	R.FVDHVFDEQVIDSLTVK.I	2	2.8605	P04843
	RIB1_HUMAN	R.SEDLLDYGPF.R.D	2	2.3191	P04843
	RIB1_HUMAN	K.NIEIDSPYEISR.A	2	2.8502	P04843
	RIB1_HUMAN	K.GEDEEENNLEVR.E	2	2.7637	P04843
	RIB1_HUMAN	K.NIEIDSPYEISR.A	2	2.7412	P04843
	RIB1_HUMAN	R.HFDETVNR.Y	2	2.3938	P04843
	RIB1_HUMAN	K.NIEIDSPYEISR.A	2	2.6436	P04843
	RIB1_HUMAN	K.GEDEEENNLEVR.E	2	3.1634	P04843
	RIB1_HUMAN	R.HFDETVNR.Y	2	2.4144	P04843
	RIB1_HUMAN	R.FVDHVFDEQVIDSLTVK.I	3	3.9712	P04843
	RIB1_HUMAN	K.GEDEEENNLEVR.E	2	2.8297	P04843
	RIB1_HUMAN	K.TEGSDLC#DR.V	2	2.5729	P04843
<b><i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 - Homo sapiens</i></b>					
	DAD1_HUMAN	R.FLEEYLSSTPQR.L	2	2.8627	P61803
<b><i>Dual specificity mitogen-activated protein kinase kinase 1 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	MP2K1_HUMAN	K.LPSGVFSLEFQDFVNK.C	2	4.1568	Q02750
	MP2K1_HUMAN	K.LPSGVFSLEFQDFVNK.C	2	3.6508	Q02750
<b><i>Dual specificity mitogen-activated protein kinase kinase 3 - Homo sapiens (Human)</i></b>					
	MP2K3_HUMAN	K.NMTIPEDILGEIAVSIVR.A	2	2.9707	P46734
	MP2K3_HUMAN	K.NMTIPEDILGEIAVSIVR.A	2	3.172	P46734
	MP2K3_HUMAN	K.QVVEEPSPQLPADR.F	2	2.3218	P46734
<b><i>Dynamain-1-like protein - Homo sapiens (Human)</i></b>					
	DNM1L_HUMAN	R.TLESVDPLGGLNTIDILTAIR.N	2	2.7179	O00429
	DNM1L_HUMAN	R.TLESVDPLGGLNTIDILTAIR.N	2	3.556	O00429
<b><i>Dynamain-2 - Homo sapiens (Human)</i></b>					
	DYN2_HUMAN	K.VPVGDQPPDIEYQIK.D	2	3.1418	P50570
	DYN2_HUMAN	K.VPVGDQPPDIEYQIK.D	2	3.5968	P50570
<b><i>Dynein heavy chain, cytosolic - Homo sapiens (Human)</i></b>					
	DYHC_HUMAN	R.GIFEALRPLETLPVEGLIR.I	2	2.7561	Q14204
<b><i>Dysferlin_v1 - Homo sapiens (Human)</i></b>					
	Q0H395_HUMAN	R.PLYSPLQPDIEQGK.L	2	2.7016	Q0H395
<b><i>Ecotropic viral integration site 5 protein homolog - Homo sapiens (Human)</i></b>					
	EVI5_HUMAN	K.NAMNELQDELMTIRLR.E	2	3.1687	O60447
<b><i>Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial precursor - Homo sap</i></b>					
	ETFD_HUMAN	K.SGILAAESIFNQLTSENLSK.T	2	2.6794	Q16134
<b><i>Elongation factor 1-alpha - Homo sapiens (Human)</i></b>					
	Q6IQ15_HUMAN	R.EHALLAYSQKR.Y	2	2.4758	Q6IQ15
<b><i>Elongation factor 1-alpha 1 - Homo sapiens (Human)</i></b>					
	EF1A1_HUMAN	R.EHALLAYTLGVK.Q	2	2.6922	P68104
	EF1A1_HUMAN	R.LPLQDVYK.I	2	2.3203	P68104
	EF1A1_HUMAN	R.EHALLAYTLGVK.Q	2	2.5266	P68104
<b><i>Elongation factor Tu, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	EFTU_HUMAN	R.TIGTGLVTNTLAMTEEEK.N	2	2.4242	P49411
	EFTU_HUMAN	R.TIGTGLVTNTLAMTEEEK.N	2	2.3956	P49411
	EFTU_HUMAN	R.TIGTGLVTNTLAM*TEEEK.N	2	2.5299	P49411
<b><i>Endoplasmic precursor - Homo sapiens (Human)</i></b>					
	ENPL_HUMAN	R.ELISNASDALDK.I	2	2.5073	P14625
<b><i>Enoyl-CoA hydratase domain-containing protein 1 - Homo sapiens (Human)</i></b>					



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	ECHD1_HUMAN	R.M*NAFSGVMM*LQLLEK.V	2	2.8276	Q9NTX5
	ECHD1_HUMAN	R.M*NAFSGVMM*LQLLEK.V	2	2.7354	Q9NTX5
<b><i>ERGIC-53 protein precursor - Homo sapiens (Human)</i></b>					
	LMAN1_HUMAN	R.LVSGMQHPGSAGGVYETTQHFDIK.E	3	4.0085	P49257
	LMAN1_HUMAN	K.GHPDLQGQPAEEIFESVGDR.E	3	3.8352	P49257
<b><i>Erlin-1 precursor - Homo sapiens (Human)</i></b>					
	ERLN1_HUMAN	K.EALEPSGENVIQNK.E	2	2.8721	O75477
	ERLN1_HUMAN	R.SVQTTLQTDEVK.N	2	2.8525	O75477
	ERLN1_HUMAN	R.SVQTTLQTDEVK.N	2	2.7322	O75477
<b><i>Erlin-2 precursor - Homo sapiens (Human)</i></b>					
	ERLN2_HUMAN	K.LSFGLEDEPLETATK.E	2	3.0559	O94905
	ERLN2_HUMAN	K.LSFGLEDEPLETATK.E	2	2.5243	O94905
<b><i>Erythrocyte band 7 integral membrane protein - Homo sapiens (Human)</i></b>					
	STOM_HUMAN	R.YLQTLTTIAAEK.N	2	2.4892	P27105
	STOM_HUMAN	K.NSTIVFPLPIDMLQGIGAK.H	2	3.1192	P27105
	STOM_HUMAN	K.NLSQILSDREEIAHNMQSTLDDATDAWGIK.V	3	4.1055	P27105
	STOM_HUMAN	R.VQNATLAVANITNADSATR.L	2	2.8131	P27105
	STOM_HUMAN	R.TISFDIPPQEILTK.D	2	2.578	P27105
	STOM_HUMAN	R.VQNATLAVANITNADSATR.L	2	3.8918	P27105
	STOM_HUMAN	R.YLQTLTTIAAEK.N	2	2.6729	P27105
	STOM_HUMAN	R.YLQTLTTIAAEK.N	2	2.4982	P27105
	STOM_HUMAN	K.NSTIVFPLPIDMLQGIGAK.H	2	2.9082	P27105
	STOM_HUMAN	R.TISFDIPPQEILTK.D	2	2.4943	P27105
	STOM_HUMAN	R.TISFDIPPQEILTK.D	2	2.6765	P27105
<b><i>Esophageal cancer associated protein - Homo sapiens (Human)</i></b>					
	Q6PDA0_HUMA	R.TRLEELDDFEESQK.E	2	2.5662	Q6PDA0
<b><i>Estradiol 17-beta-dehydrogenase 12 - Homo sapiens (Human)</i></b>					
	DHB12_HUMAN	K.DKLDQVSSEIK.E	2	2.5506	Q53GQ0
	DHB12_HUMAN	R.TIAVDFASEDIYDK.I	2	2.7648	Q53GQ0
	DHB12_HUMAN	R.TIAVDFASEDIYDK.I	2	3.0326	Q53GQ0
<b><i>Eukaryotic initiation factor 4A-I - Homo sapiens (Human)</i></b>					
	IF4A1_HUMAN	K.GYDVIAQAQSGTGK.T	2	2.4418	P60842
<b><i>Eukaryotic translation initiation factor 3 subunit 5 - Homo sapiens (Human)</i></b>					
	IF35_HUMAN	R.IQDALSTVLQYAEDVLSGK.V	3	5.0048	O00303

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>EVI2B protein precursor - Homo sapiens (Human)</i></b>					
EVI2B_HUMAN		R.SPFADGETPDIC#MDNIR.E	2	4.1159	P34910
EVI2B_HUMAN		K.LFESSENIEDSNNPK.T	2	4.4448	P34910
EVI2B_HUMAN		R.SPFADGETPDIC#M*DNIR.E	2	2.3286	P34910
EVI2B_HUMAN		R.SPFADGETPDIC#MDNIR.E	2	3.5217	P34910
EVI2B_HUMAN		K.LFESSENIEDSNNPK.T	2	3.8501	P34910
EVI2B_HUMAN		R.SPFADGETPDIC#MDNIR.E	2	4.0955	P34910
EVI2B_HUMAN		K.LFESSENIEDSNNPK.T	2	2.6506	P34910
<b><i>Exportin-2 - Homo sapiens (Human)</i></b>					
XPO2_HUMAN		K.LLQTDDEEEAGLLELLK.S	2	3.6692	P55060
XPO2_HUMAN		K.IPGLLGVFQK.L	2	2.6059	P55060
XPO2_HUMAN		K.LLQTDDEEEAGLLELLK.S	2	3.7176	P55060
<b><i>Fatty aldehyde dehydrogenase - Homo sapiens (Human)</i></b>					
AL3A2_HUMAN		K.NVDEAINFINER.E	2	2.3645	P51648
<b><i>Filamin-A - Homo sapiens (Human)</i></b>					
FLNA_HUMAN		K.YTPVQQGPVGVNVVYGGDPIPK.S	2	3.3393	P21333
FLNA_HUMAN		K.VTAQGPGLPSGNIANK.T	2	2.562	P21333
FLNA_HUMAN		R.EAGAGGLAIAVEGPSK.A	2	3.4309	P21333
FLNA_HUMAN		K.VTAQGPGLPSGNIANK.T	2	3.2589	P21333
FLNA_HUMAN		R.SPYTVTVGQAC#NPSAC#R.A	2	2.9662	P21333
<b><i>FLJ00121 protein - Homo sapiens (Human)</i></b>					
Q8TES2_HUMA		K.EGLAVAEELVDHVAELTATPR.D	3	3.9239	Q8TES2
Q8TES2_HUMA		K.EGLAVAEELVDHVAELTATPR.D	3	3.8368	Q8TES2
<b><i>Formin-like protein 1 - Homo sapiens (Human)</i></b>					
FMNL_HUMAN		K.VLQELDMSEDFEEQFK.T	2	3.5916	O95466
FMNL_HUMAN		K.NAVLEHMEELQEQQVALLTER.L	2	4.7587	O95466
FMNL_HUMAN		K.VLQELDM*SEDFEEQFK.T	2	2.4959	O95466
<b><i>Full-length cDNA clone CS0DI085YI08 of Placenta of Homo sapiens - Homo sapiens (Human)</i></b>					
Q86TV4_HUMAN		R.PTC#PSSSGGPARELSSRLR.I	2	2.5759	Q86TV4
<b><i>Gasdermin domain-containing protein 1 - Homo sapiens (Human)</i></b>					
GSDC1_HUMAN		K.DILEPDAAEPDVQR.G	2	2.5047	P57764
<b><i>GCN1-like protein 1 - Homo sapiens (Human)</i></b>					
GCN1L_HUMAN		K.NIVSLLLSMLGHDEDNTR.I	2	3.6647	Q92616
GCN1L_HUMAN		K.NIVSLLLSMLGHDEDNTR.I	2	5.2816	Q92616

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>GEM-interacting protein - Homo sapiens (Human)</i></b>					
	GMIP_HUMAN	R.SQGSPEDESAPQASPGPSK.Q	2	2.4976	Q9P107
<b><i>Glutathione S-transferase kappa 1 - Homo sapiens (Human)</i></b>					
	GSTK1_HUMAN	R.NEDITEPQSILAAAEK.A	2	2.9858	Q9Y2Q3
	GSTK1_HUMAN	R.NEDITEPQSILAAAEK.A	2	3.7887	Q9Y2Q3
<b><i>Golgi-associated plant pathogenesis-related protein 1 - Homo sapiens (Human)</i></b>					
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	3.0191	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	2.8332	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	2.943	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	2.8961	Q9H4G4
<b><i>GPI transamidase component PIG-S - Homo sapiens (Human)</i></b>					
	PIGS_HUMAN	R.SVENLATATTTLSLAQLLQK.I	2	3.2225	Q96S52
<b><i>GTPase IMAP family member 1 - Homo sapiens (Human)</i></b>					
	GIMA1_HUMAN	K.MATDEENVYGLEENAQSR.Q	2	3.8996	Q8WWP7
	GIMA1_HUMAN	R.EQEAQVEQLLGMVEGLVLEHK.G	2	3.1408	Q8WWP7
	GIMA1_HUMAN	K.MATDEENVYGLEENAQSR.Q	2	3.9364	Q8WWP7
	GIMA1_HUMAN	K.MATDEENVYGLEENAQSR.Q	2	2.3763	Q8WWP7
	GIMA1_HUMAN	K.MATDEENVYGLEENAQSR.Q	2	2.5122	Q8WWP7
	GIMA1_HUMAN	R.EQEAQVEQLLGMVEGLVLEHK.G	2	3.8522	Q8WWP7
<b><i>GTPase IMAP family member 4 - Homo sapiens (Human)</i></b>					
	GIMA4_HUMAN	R.EAPEDIQDLMDIFGDR.Y	2	3.8313	Q9NUV9
	GIMA4_HUMAN	R.EAPEDIQDLMDIFGDR.Y	2	2.6685	Q9NUV9
<b><i>Guanine nucleotide-binding protein G(i), alpha-2 subunit - Homo sapiens (Human)</i></b>					
	GNAI2_HUMAN	R.QLFALSC#TAEEQGVLPDDLQSGVIR.R	2	3.83	P04899
	GNAI2_HUMAN	K.YDEAASYIQSK.F	2	2.8459	P04899
	GNAI2_HUMAN	K.YDEAASYIQSK.F	2	2.689	P04899
	GNAI2_HUMAN	R.QLFALSC#TAEEQGVLPDDLQSGVIR.R	2	3.0653	P04899
	GNAI2_HUMAN	K.YDEAASYIQSK.F	2	3.1257	P04899
<b><i>Heat shock protein HSP 90-alpha - Homo sapiens (Human)</i></b>					
	HS90A_HUMAN	R.APFDFENR.K	2	2.9465	P07900
	HS90A_HUMAN	R.NPDDITNEEYGEFYK.S	2	2.8114	P07900
<b><i>Heat shock protein HSP 90-beta - Homo sapiens (Human)</i></b>					
	HS90B_HUMAN	K.HLEINPDHPIVETLR.Q	3	3.8039	P08238
	HS90B_HUMAN	R.GFEVVYMTPEIDEYC#VQQLK.E	2	2.5953	P08238

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	HS90B_HUMAN	R.NPDDITQEEYGEFYK.S	2	4.3726	P08238
	HS90B_HUMAN	R.NPDDITQEEYGEFYK.S	2	3.8581	P08238
<b><i>Heterogeneous nuclear ribonucleoprotein H - Homo sapiens (Human)</i></b>					
	HNRH1_HUMAN	R.STGEAFVQFASQEIAEK.A	2	3.55	P31943
<b><i>Heterogeneous nuclear ribonucleoprotein K - Homo sapiens (Human)</i></b>					
	HNRPK_HUMAN	R.TDYNASVSPDSSGPER.I	2	3.1021	P61978
	HNRPK_HUMAN	R.TDYNASVSPDSSGPER.I	2	3.5951	P61978
	HNRPK_HUMAN	R.TDYNASVSPDSSGPER.I	2	3.6595	P61978
	HNRPK_HUMAN	R.TDYNASVSPDSSGPER.I	2	2.424	P61978
<b><i>Heterogeneous nuclear ribonucleoprotein U-like protein 2 - Homo sapiens (Human)</i></b>					
	HNRL2_HUMAN	R.DLLVQQASQC#LSK.L	2	2.6555	Q1KMD3
	HNRL2_HUMAN	R.DLLVQQASQC#LSK.L	2	3.2113	Q1KMD3
	HNRL2_HUMAN	R.DLLVQQASQC#LSK.L	2	3.0462	Q1KMD3
<b><i>Hexokinase-1 - Homo sapiens (Human)</i></b>					
	HXK1_HUMAN	R.HIDLVEGDEGR.M	2	3.0338	P19367
<b><i>Hexokinase-3 - Homo sapiens (Human)</i></b>					
	HXK3_HUMAN	R.FDASVDQASINPGK.Q	2	3.5046	P52790
<b><i>HFMI protein - Homo sapiens (Human)</i></b>					
	A2RUB2_HUMA	K.ALC#SQRFDWKEK.F	2	2.5498	A2RUB2
<b><i>High affinity immunoglobulin epsilon receptor subunit gamma precursor - Homo sapiens (Hu)</i></b>					
	FCERG_HUMAN	R.NQETYETLK.H	2	2.4784	P30273
	FCERG_HUMAN	K.SDGVYTGLSTR.N	2	2.8191	P30273
	FCERG_HUMAN	K.SDGVYTGLSTR.N	2	2.6658	P30273
	FCERG_HUMAN	K.SDGVYTGLSTR.N	2	2.5197	P30273
<b><i>Histocompatibility - Homo sapiens (Human)</i></b>					
	Q6P189_HUMAN	R.SPLTAASPGELPTEGAGPDVVEDISHLLADVAR.F	3	4.7	Q6P189
<b><i>Histone H2A type 1-A - Homo sapiens (Human)</i></b>					
	H2A1A_HUMAN	R.HLQLAIR.N	2	2.3005	Q96QV6
	H2A1A_HUMAN	R.NDEELNK.L	2	2.4062	Q96QV6
	H2A1A_HUMAN	R.HLQLAIR.N	2	2.3853	Q96QV6
	H2A1A_HUMAN	R.HLQLAIR.N	2	2.3552	Q96QV6
	H2A1A_HUMAN	R.HLQLAIR.N	2	2.3463	Q96QV6
<b><i>Histone H2A type 1-B - Homo sapiens (Human)</i></b>					
	H2A1B_HUMAN	R.VGAGAPVYLAHVLEYLTAEILELAGNAAR.D	2	4.9576	P04908

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
H2A1B_HUMAN		R.VGAGAPVYLAADVLELYLTAEILELAGNAAR.D	4	6.1809	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLELYLTAEILELAGNAAR.D	2	5.2091	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLELYLTAEILELAGNAAR.D	4	5.7893	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLELYLTAEILELAGNAAR.D	4	5.1966	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLELYLTAEILELAGNAAR.D	4	4.7811	P04908
<b><i>Histone H2A type 2-A - Homo sapiens (Human)</i></b>					
H2A2A_HUMAN		R.VGAGAPVYMAADVLELYLTAEILELAGNAAR.D	2	4.9025	Q6F113
H2A2A_HUMAN		R.VGAGAPVYM*AAVLELYLTAEILELAGNAAR.D	2	3.9524	Q6F113
H2A2A_HUMAN		R.VGAGAPVYM*AAVLELYLTAEILELAGNAAR.D	2	4.0714	Q6F113
H2A2A_HUMAN		R.VGAGAPVYMAADVLELYLTAEILELAGNAAR.D	2	4.8799	Q6F113
<b><i>Histone H2AV - Homo sapiens (Human)</i></b>					
H2AV_HUMAN		R.GDEELDSLK.A	2	2.3731	Q71UI9
H2AV_HUMAN		R.GDEELDSLK.A	2	2.605	Q71UI9
<b><i>Histone H3.1 - Homo sapiens (Human)</i></b>					
H31_HUMAN		R.FQSSAVMALQEAC#EAYLVGLFEDTNLC#AIHAK.R	3	4.1636	P68431
<b><i>Histone H4 - Homo sapiens (Human)</i></b>					
H4_HUMAN		R.DAVTYTEHAK.R	2	2.305	P62805
H4_HUMAN		K.TVTAM*DVVYALK.R	2	2.4806	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.2016	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	3.4367	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.2482	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	2.8581	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.0424	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.3493	P62805
H4_HUMAN		K.TVTAM*DVVYALK.R	2	2.4437	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.1408	P62805
<b><i>HLA class I histocompatibility antigen, A-1 alpha chain precursor - Homo sapiens (Human)</i></b>					
1A01_HUMAN		R.DGEDQTQDTELVETR.P	2	4.0365	P30443
1A01_HUMAN		R.DGEDQTQDTELVETRPAGDGTFFQK.W	3	3.7702	P30443
1A01_HUMAN		R.DGEDQTQDTELVETR.P	2	3.5263	P30443
1A01_HUMAN		R.DGEDQTQDTELVETR.P	2	5.0987	P30443
1A01_HUMAN		R.APWIEQEGPEYWDQETR.N	2	3.2609	P30443
<b><i>HLA class I histocompatibility antigen, A-2 alpha chain precursor - Homo sapiens (Human)</i></b>					
1A02_HUMAN		R.FSDAASQR.M	2	2.5787	P01892

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>HLA class I histocompatibility antigen, A-25 alpha chain precursor - Homo sapiens (Human)</i></b>					
1A25_HUMAN		R.APWIEQEGPEYWDR.N	2	3.0361	P18462
1A25_HUMAN		R.APWIEQEGPEYWDR.N	2	3.7456	P18462
1A25_HUMAN		R.APWIEQEGPEYWDR.N	2	3.9579	P18462
1A25_HUMAN		R.APWIEQEGPEYWDR.N	2	3.6625	P18462
<b><i>HLA class II histocompatibility antigen gamma chain - Homo sapiens (Human)</i></b>					
HG2A_HUMAN		R.DLISNNEQLPMLGR.R	2	2.5213	P04233
HG2A_HUMAN		R.DLISNNEQLPMLGR.R	2	3.0894	P04233
HG2A_HUMAN		R.DLISNNEQLPMLGR.R	2	2.7821	P04233
HG2A_HUMAN		R.DLISNNEQLPMLGR.R	2	2.5326	P04233
HG2A_HUMAN		R.DLISNNEQLPMLGR.R	2	2.9254	P04233
<b><i>HLA class II histocompatibility antigen, DR alpha chain precursor - Homo sapiens (Human)</i></b>					
2DRA_HUMAN		R.FASFEAQGALANIAVDK.A	2	4.2577	P01903
2DRA_HUMAN		R.FASFEAQGALANIAVDK.A	2	4.2646	P01903
2DRA_HUMAN		R.FASFEAQGALANIAVDK.A	2	4.3054	P01903
<b><i>Hsp90 co-chaperone Cdc37 - Homo sapiens (Human)</i></b>					
CDC37_HUMAN		R.LGPGGLDPVEVYESLPEELQK.C	2	3.897	Q16543
<b><i>IgG receptor FcRn large subunit p51 precursor - Homo sapiens (Human)</i></b>					
FCGRN_HUMAN		R.GDDTGVLPTPGEAQDADLK.D	2	2.581	P55899
<b><i>Immediate early response 3-interacting protein 1 - Homo sapiens (Human)</i></b>					
IR3IP_HUMAN		K.NIGWGTDQGIGGFGEPEGIK.S	2	3.4964	Q9Y5U9
IR3IP_HUMAN		K.NIGWGTDQGIGGFGEPEGIK.S	2	2.4238	Q9Y5U9
<b><i>Immunoglobulin J chain - Homo sapiens (Human)</i></b>					
IGJ_HUMAN		-.QEDERIVLVDNK.C	2	2.5603	P01591
<b><i>Induced by contact to basement membrane 1 protein - Homo sapiens (Human)</i></b>					
ICB1_HUMAN		K.IPSTLEVDVEDVTASSR.H	2	2.9197	Q5TEJ8
ICB1_HUMAN		K.IPSTLEVDVEDVTASSR.H	2	3.3185	Q5TEJ8
ICB1_HUMAN		K.IPSTLEVDVEDVTASSR.H	2	3.5795	Q5TEJ8
ICB1_HUMAN		K.IPSTLEVDVEDVTASSR.H	2	2.3477	Q5TEJ8
ICB1_HUMAN		K.IPSTLEVDVEDVTASSR.H	2	3.438	Q5TEJ8
<b><i>Inhibitor of nuclear factor kappa-B kinase subunit beta - Homo sapiens (Human)</i></b>					
IKKB_HUMAN		R.EGAILTLLSDIASALR.Y	2	2.4566	O14920
<b><i>Integrin alpha-IIb precursor - Homo sapiens (Human)</i></b>					
ITA2B_HUMAN		R.NRPPLEEDDEEGE.-	2	3.0358	P08514

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	ITA2B_HUMAN	R.DGYNDIAVAAPYGGPSGR.G	2	2.7262	P08514
	ITA2B_HUMAN	R.NRPPEEDDEEGE.-	2	3.4577	P08514
	ITA2B_HUMAN	R.NRPPEEDDEEGE.-	2	3.4194	P08514
	ITA2B_HUMAN	R.NRPPEEDDEEGE.-	2	2.674	P08514
<b><i>Integrin alpha-L precursor - Homo sapiens (Human)</i></b>					
	ITAL_HUMAN	K.VLIITDGEATDSGNIDAAK.D	2	2.8907	P20701
	ITAL_HUMAN	K.ADLQDDTFIGNEPLTPEVR.A	2	2.6961	P20701
<b><i>Integrin alpha-M precursor - Homo sapiens (Human)</i></b>					
	ITAM_HUMAN	R.LPSHSDFLAELR.K	2	2.727	P11215
	ITAM_HUMAN	K.LTDVAIGAPGEEDNR.G	2	3.0424	P11215
	ITAM_HUMAN	R.LPSHSDFLAELR.K	2	2.3048	P11215
	ITAM_HUMAN	K.FGDPLGYEDVIPEADR.E	2	2.9043	P11215
	ITAM_HUMAN	K.FGDPLGYEDVIPEADR.E	2	3.0858	P11215
	ITAM_HUMAN	K.FGDPLGYEDVIPEADR.E	2	3.3551	P11215
	ITAM_HUMAN	K.FGDPLGYEDVIPEADR.E	2	2.7179	P11215
	ITAM_HUMAN	R.LPSHSDFLAELR.K	2	3.0381	P11215
	ITAM_HUMAN	R.LPSHSDFLAELR.K	3	4.0661	P11215
	ITAM_HUMAN	K.LTDVAIGAPGEEDNR.G	2	3.8306	P11215
	ITAM_HUMAN	R.LPSHSDFLAELR.K	2	2.7186	P11215
	ITAM_HUMAN	K.LTDVAIGAPGEEDNR.G	2	3.0653	P11215
<b><i>Integrin beta chain, beta 2 variant - Homo sapiens (Human)</i></b>					
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	3.8836	Q59H50
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	2.9758	Q59H50
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	4.3463	Q59H50
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	3.7253	Q59H50
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	3.8638	Q59H50
	Q59H50_HUMAN	K.SAVGELSESSNVVHLIK.N	2	4.0036	Q59H50
<b><i>Integrin beta-2 precursor - Homo sapiens (Human)</i></b>					
	ITB2_HUMAN	R.TTEGC#LNPR.R	2	2.4899	P05107
<b><i>Integrin beta-3 precursor - Homo sapiens (Human)</i></b>					
	ITB3_HUMAN	K.GSGDSSQVTQVSPQR.I	2	3.4558	P05106
	ITB3_HUMAN	K.GSGDSSQVTQVSPQR.I	2	3.896	P05106
	ITB3_HUMAN	K.GSGDSSQVTQVSPQR.I	2	3.4734	P05106
	ITB3_HUMAN	K.GSGDSSQVTQVSPQR.I	2	2.9107	P05106
	ITB3_HUMAN	R.NDASHLLVFTTDAK.T	2	4.155	P05106

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ITB3_HUMAN		K.IGDTVVSFSIEAK.V	2	2.8902	P05106
ITB3_HUMAN		K.IGDTVVSFSIEAK.V	2	2.7708	P05106
ITB3_HUMAN		K.IGDTVVSFSIEAK.V	2	2.5078	P05106
ITB3_HUMAN		K.GSGDSSQVTQVSPQR.I	2	3.7582	P05106
ITB3_HUMAN		K.GSGDSSQVTQVSPQR.I	2	3.5792	P05106
ITB3_HUMAN		K.IGDTVVSFSIEAK.V	2	2.6218	P05106
<b><i>Intercellular adhesion molecule 2 precursor - Homo sapiens (Human)</i></b>					
ICAM2_HUMAN		R.VPTVEPLDSLTLFLFR.G	2	3.0988	P13598
ICAM2_HUMAN		R.VPTVEPLDSLTLFLFR.G	2	3.0263	P13598
<b><i>Keratin, type I cytoskeletal 10 - Homo sapiens (Human)</i></b>					
K1C10_HUMAN		K.ADLEM*QIESLTEELAYLK.K	2	3.2634	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	2	4.7417	P13645
K1C10_HUMAN		K.ADLEMQIESLTEELAYLK.K	2	3.4991	P13645
K1C10_HUMAN		R.ALEESNYELEGK.I	2	3.3039	P13645
K1C10_HUMAN		K.ELTTEIDNIEQISSYK.S	2	4.2919	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	3	4.0626	P13645
K1C10_HUMAN		K.ELTTEIDNIEQISSYK.S	2	3.4183	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	2	4.2369	P13645
K1C10_HUMAN		K.ELTTEIDNIEQISSYK.S	2	4.0512	P13645
K1C10_HUMAN		R.NVSTGDVNVEMNAAPGVDLTQLLNMR.S	2	3.2528	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	2	5.8869	P13645
K1C10_HUMAN		R.NVSTGDVNVEMNAAPGVDLTQLLNMR.S	2	2.9629	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	2	3.6607	P13645
K1C10_HUMAN		K.YENEVALR.Q	2	2.4535	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	3	5.5405	P13645
K1C10_HUMAN		R.VLDELTLTK.A	2	2.6948	P13645
K1C10_HUMAN		R.VLDELTLTK.A	2	2.4325	P13645
K1C10_HUMAN		K.ADLEMQIESLTEELAYLK.K	2	4.3589	P13645
K1C10_HUMAN		K.ADLEM*QIESLTEELAYLK.K	2	3.8233	P13645
K1C10_HUMAN		R.ALEESNYELEGK.I	2	2.3575	P13645
K1C10_HUMAN		K.NQILNLTTDNANILLQIDNAR.L	2	3.2759	P13645
K1C10_HUMAN		K.ADLEMQIESLTEELAYLK.K	2	2.9345	P13645
K1C10_HUMAN		R.ALEESNYELEGK.I	2	3.3805	P13645
K1C10_HUMAN		K.ADLEM*QIESLTEELAYLK.K	2	3.0669	P13645
K1C10_HUMAN		R.ALEESNYELEGK.I	2	3.4309	P13645



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Keratin, type I cytoskeletal 9 - Homo sapiens (Human)</i></b>					
K1C9_HUMAN		K.SDLEM*QYETLQEELMALK.K	2	2.9683	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELMALK.K	2	5.4804	P35527
K1C9_HUMAN		K.VQALEEANNNDLENK.I	2	4.3705	P35527
K1C9_HUMAN		R.GGSGGSYGGGGSGGGYGGGSGSR.G	2	4.5348	P35527
K1C9_HUMAN		K.EIETYHNLLEGGQEDFESSGAGK.I	2	3.4702	P35527
K1C9_HUMAN		K.EIETYHNLLEGGQEDFESSGAGK.I	2	2.7375	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELM*ALK.K	2	2.5705	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELMALK.K	2	3.2947	P35527
K1C9_HUMAN		R.GGSGGSYGGGGSGGGYGGGSGSR.G	2	2.9533	P35527
K1C9_HUMAN		K.DIENQYETQITQIEHEVSSSQEVQSSAK.E	3	4.7307	P35527
K1C9_HUMAN		K.VQALEEANNNDLENK.I	2	4.6489	P35527
K1C9_HUMAN		K.SDLEM*QYETLQEELMALK.K	2	2.472	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELM*ALK.K	2	2.3079	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELM*ALK.K	2	3.5463	P35527
K1C9_HUMAN		R.HGVQELEIELQSLSK.K	2	4.6278	P35527
K1C9_HUMAN		K.VQALEEANNNDLENK.I	2	4.6055	P35527
K1C9_HUMAN		K.VQALEEANNNDLENK.I	2	4.9163	P35527
K1C9_HUMAN		K.SDLEMQYETLQEELMALK.K	3	4.4094	P35527
K1C9_HUMAN		R.SGGGGGGGLGSGGSIR.S	2	2.6448	P35527
<b><i>Keratin, type II cytoskeletal I - Homo sapiens (Human)</i></b>					
K2C1_HUMAN		K.SLNNQFASFIDK.V	2	3.0592	P04264
K2C1_HUMAN		R.DYQELMNTK.L	2	2.32	P04264
K2C1_HUMAN		K.YEELQITAGR.H	2	3.156	P04264
K2C1_HUMAN		R.TNAENEFVTIK.K	2	2.6725	P04264
K2C1_HUMAN		K.QISNLQQSISDAEQR.G	2	3.4289	P04264
K2C1_HUMAN		K.LNDLEDALQQAK.E	2	3.4804	P04264
K2C1_HUMAN		K.YEELQITAGR.H	2	2.7122	P04264
K2C1_HUMAN		R.TNAENEFVTIK.K	2	2.4518	P04264
K2C1_HUMAN		R.GGGGGYGGSSYGGSSYGGGGGGGR.	2	5.3098	P04264
K2C1_HUMAN		K.LNDLEDALQQAK.E	2	3.2086	P04264
K2C1_HUMAN		K.YEELQITAGR.H	2	3.0369	P04264
K2C1_HUMAN		R.SLDLDSIAEVK.A	2	3.072	P04264
K2C1_HUMAN		R.SLDLDSIAEVK.A	1	2.3109	P04264
K2C1_HUMAN		K.NMQDMVEDYR.N	2	2.3539	P04264

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
K2C1_HUMAN		R.SLDLDSIIAEVK.A	1	2.2855	P04264
K2C1_HUMAN		R.SLDLDSIIAEVK.A	2	2.5203	P04264
K2C1_HUMAN		K.LNDLEDALQQAK.E	2	3.3169	P04264
K2C1_HUMAN		R.TNAENEFVTIK.K	2	2.7825	P04264
K2C1_HUMAN		R.GGGGGGYGSGGSSYGSGGGSYGSGGGGGGGR.	3	4.7775	P04264
K2C1_HUMAN		R.SLDLDSIIAEVK.A	1	2.3322	P04264
K2C1_HUMAN		R.SLDLDSIIAEVK.A	2	3.8614	P04264
K2C1_HUMAN		K.SLNNQFASFIDK.V	2	3.2625	P04264
K2C1_HUMAN		K.LNDLEDALQQAK.E	2	3.9508	P04264
K2C1_HUMAN		K.LNDLEDALQQAK.E	2	2.821	P04264
K2C1_HUMAN		K.YEELQITAGR.H	2	2.7268	P04264
K2C1_HUMAN		R.DYQELMNTK.L	2	2.4075	P04264
K2C1_HUMAN		K.QISNLQQSISDAEQR.G	2	3.2678	P04264
K2C1_HUMAN		K.QISNLQQSISDAEQR.G	2	3.559	P04264
K2C1_HUMAN		R.SLDLDSIIAEVK.A	1	2.1788	P04264
K2C1_HUMAN		R.SLDLDSIIAEVK.A	2	3.9229	P04264
<b><i>Keratin, type II cytoskeletal 2 epidermal - Homo sapiens (Human)</i></b>					
K22E_HUMAN		K.NVQDAIADAEQR.G	2	3.4638	P35908
K22E_HUMAN		R.NLDLDSIIAEVK.A	2	3.2551	P35908
K22E_HUMAN		K.LNDLEEALQQAK.E	2	3.3077	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	4.5191	P35908
K22E_HUMAN		R.NLDLDSIIAEVK.A	2	3.3207	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	4.0113	P35908
K22E_HUMAN		R.NLDLDSIIAEVK.A	2	3.2757	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	3.8334	P35908
K22E_HUMAN		R.NLDLDSIIAEVK.A	2	3.1737	P35908
K22E_HUMAN		K.NVQDAIADAEQR.G	2	3.4187	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	4.375	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	4.0051	P35908
K22E_HUMAN		R.FLEQQNQVLQTK.W	2	4.1679	P35908
<b><i>Keratin, type II cytoskeletal 6B - Homo sapiens (Human)</i></b>					
K2C6B_HUMAN		R.NMQDLVEDLKNK.Y	2	2.5135	P04259
<b><i>KIAA1727 protein - Homo sapiens (Human)</i></b>					
Q9C0D6_HUMA		R.TLTASENESM*RKVM*PITK.S	2	2.3402	Q9C0D6
<b><i>Kinesin-like protein KIF14 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	KIF14_HUMAN	K.LHLKSDM*SECENDDPLLRSAAGK.V	2	2.6812	Q15058
	KIF14_HUMAN	K.LHLKSDM*SECENDDPLLRSAAGK.V	2	2.3698	Q15058
<b><i>Kinesin-like protein KIF9 - Homo sapiens (Human)</i></b>					
	KIF9_HUMAN	R.M*VKNLEKELALLK.Q	2	2.3275	Q9HAQ2
<b><i>Lamin-B receptor - Homo sapiens (Human)</i></b>					
	LBR_HUMAN	R.NDLSPASSGNAVYDFFIGR.E	2	3.8079	Q14739
<b><i>Lamin-B1 - Homo sapiens (Human)</i></b>					
	LMNB1_HUMAN	R.IESLSSQLSNLQK.E	2	3.2662	P20700
	LMNB1_HUMAN	K.SLEGDLEDLKDQIAQLEASLAAAK.K	2	6.3848	P20700
	LMNB1_HUMAN	R.DQMQQQLNDYEQLLDVK.L	2	3.2395	P20700
	LMNB1_HUMAN	R.SLETENSALQLQVTER.E	2	3.1458	P20700
	LMNB1_HUMAN	K.DAALATALGDKK.S	2	2.7247	P20700
	LMNB1_HUMAN	K.SLEGDLEDLKDQIAQLEASLAAAK.K	2	5.5658	P20700
	LMNB1_HUMAN	R.IESLSSQLSNLQK.E	2	3.1075	P20700
	LMNB1_HUMAN	K.SLEGDLEDLKDQIAQLEASLAAAK.K	2	5.3369	P20700
	LMNB1_HUMAN	R.DQMQQQLNDYEQLLDVK.L	2	3.5275	P20700
	LMNB1_HUMAN	R.IESLSSQLSNLQK.E	2	3.3787	P20700
	LMNB1_HUMAN	K.SLEGDLEDLKDQIAQLEASLAAAK.K	2	5.818	P20700
	LMNB1_HUMAN	K.NTSEQDQPMGGWEMIR.K	2	2.8265	P20700
<b><i>LENG4 protein - Homo sapiens (Human)</i></b>					
	Q05DF0_HUMA	R.AGGGPTLQC#PPPSSPEK.A	2	3.6199	Q05DF0
	Q05DF0_HUMA	R.AGGGPTLQC#PPPSSPEK.A	2	3.3679	Q05DF0
<b><i>Leucine-rich repeat serine/threonine-protein kinase 2 - Homo sapiens (Human)</i></b>					
	LRRK2_HUMAN	K.FPNEFDELEIQGK.L	2	3.1722	Q5S007
<b><i>Leukocyte common antigen precursor - Homo sapiens (Human)</i></b>					
	CD45_HUMAN	K.DLQYSTDYTFK.A	2	2.7668	P08575
	CD45_HUMAN	K.HELEMSK.E	2	2.7027	P08575
	CD45_HUMAN	R.SC#NLDEQQELVERDDEK.Q	2	2.5045	P08575
	CD45_HUMAN	R.DPPSEPSPLEAEFQR.L	2	2.605	P08575
	CD45_HUMAN	R.DPPSEPSPLEAEFQR.L	2	2.8741	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	3.8958	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	3.7506	P08575
	CD45_HUMAN	R.DPPSEPSPLEAEFQR.L	2	2.388	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	3.2179	P08575

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Leukotriene B4 receptor 1 - Homo sapiens (Human)</i></b>					
	LT4R1_HUMAN	R.SGPAALEPGPSESLTASSPLK.L	2	2.4265	Q15722
<b><i>Lipase member K precursor - Homo sapiens (Human)</i></b>					
	LIPK_HUMAN	K.YDLPATINFIIIEKTGQK.R	2	2.6949	Q5VXJ0
<b><i>LOC387790 protein - Homo sapiens (Human)</i></b>					
	Q3ZCU0_HUMA	R.ARSADVQGQEKM*DVTAQEAR.T	2	2.4301	Q3ZCU0
<b><i>LOC730092 protein - Homo sapiens (Human)</i></b>					
	Q7L4D9_HUMAN	R.LLTPVIPALWEAEVGGSPSEVRSSR.P	2	2.5307	Q7L4D9
<b><i>Long-chain fatty acid transport protein 3 - Homo sapiens (Human)</i></b>					
	S27A3_HUMAN	R.AAPGAGDAAAGSGAEFAGGDGAAR.G	2	4.0686	Q5K4L6
<b><i>Lysozyme C precursor - Homo sapiens (Human)</i></b>					
	LYSC_HUMAN	R.STDYGIFQINSR.Y	2	3.4726	P61626
	LYSC_HUMAN	R.STDYGIFQINSR.Y	2	3.5298	P61626
	LYSC_HUMAN	R.STDYGIFQINSR.Y	2	3.493	P61626
	LYSC_HUMAN	R.STDYGIFQINSR.Y	2	3.1157	P61626
	LYSC_HUMAN	R.STDYGIFQINSR.Y	2	3.5822	P61626
<b><i>MBC3205 - Homo sapiens (Human)</i></b>					
	Q6UW68_HUMA	R.GLGGEVPGSHQGPDPYR.Q	2	3.2399	Q6UW68
<b><i>Membrane-associated progesterone receptor component 1 - Homo sapiens (Human)</i></b>					
	PGRC1_HUMAN	K.EALKDEYDDLSDLTAAQQETLSDWESQFTFK.Y	3	3.8886	O00264
<b><i>Membrane-associated progesterone receptor component 2 - Homo sapiens (Human)</i></b>					
	PGRC2_HUMAN	R.GLGAGAGAGEESPATSLPR.M	2	2.7734	O15173
<b><i>Methyltransferase-like protein 7A precursor - Homo sapiens (Human)</i></b>					
	MET7A_HUMAN	R.ELFSNLQEFAGPSGK.L	2	4.5344	Q9H8H3
	MET7A_HUMAN	R.ELFSNLQEFAGPSGK.L	2	3.4861	Q9H8H3
	MET7A_HUMAN	R.ELFSNLQEFAGPSGK.L	2	3.895	Q9H8H3
	MET7A_HUMAN	R.ELFSNLQEFAGPSGK.L	2	2.5348	Q9H8H3
	MET7A_HUMAN	R.ELFSNLQEFAGPSGK.L	2	3.3919	Q9H8H3
<b><i>Microspherule protein 1 - Homo sapiens (Human)</i></b>					
	MCRS1_HUMAN	R.HPDAFYLARTAK.A	2	2.5237	Q96EZ8
<b><i>Mitochondrial carrier homolog 2 - Homo sapiens (Human)</i></b>					
	MTCH2_HUMAN	K.VLQHYQESDKGEELGPGNVQK.E	3	5.428	Q9Y6C9
	MTCH2_HUMAN	R.EEGILGFFAGLVPR.L	2	2.6783	Q9Y6C9

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	MTCH2_HUMAN	R.EEGILGFFAGLVPR.L	2	3.2509	Q9Y6C9
	MTCH2_HUMAN	R.EEGILGFFAGLVPR.L	2	3.3758	Q9Y6C9
	MTCH2_HUMAN	K.VLQHYQESDKGEELGPGNVQK.E	3	5.0294	Q9Y6C9
<b><i>Mitochondrial inner membrane protein - Homo sapiens (Human)</i></b>					
	IMMT_HUMAN	R.GIEQAVQSHAVAEER.K	2	2.8406	Q16891
	IMMT_HUMAN	K.TSSAETPTIPLGSAVEAIK.A	2	2.4469	Q16891
	IMMT_HUMAN	R.GIEQAVQSHAVAEER.K	3	3.9266	Q16891
	IMMT_HUMAN	R.GIEQAVQSHAVAEER.K	2	3.5931	Q16891
	IMMT_HUMAN	K.TSSAETPTIPLGSAVEAIK.A	2	3.3464	Q16891
<b><i>Mitochondrial Rho GTPase 1 - Homo sapiens (Human)</i></b>					
	MIRO1_HUMAN	R.FGYDDDLDTPEYLFPLLK.I	2	3.1201	Q8IXI2
	MIRO1_HUMAN	R.FGYDDDLDTPEYLFPLLK.I	2	3.5566	Q8IXI2
	MIRO1_HUMAN	R.FGYDDDLDTPEYLFPLLK.I	2	2.3064	Q8IXI2
	MIRO1_HUMAN	R.FGYDDDLDTPEYLFPLLK.I	2	2.8852	Q8IXI2
<b><i>MMS19-like protein - Homo sapiens (Human)</i></b>					
	MMS19_HUMAN	R.ADAEDLLDSFLSNILQDC#R.H	2	3.7298	Q96T76
<b><i>Moesin - Homo sapiens (Human)</i></b>					
	MOES_HUMAN	K.FYPEDVSEELIQDITQR.L	2	2.4121	P26038
	MOES_HUMAN	K.FYPEDVSEELIQDITQR.L	2	2.4852	P26038
<b><i>Monocyte differentiation antigen CD14 precursor - Homo sapiens (Human)</i></b>					
	CD14_HUMAN	K.ITGTMPPLEATGLALSSLR.L	2	2.4571	P08571
	CD14_HUMAN	K.ITGTMPPLEATGLALSSLR.L	2	2.5064	P08571
	CD14_HUMAN	K.ITGTMPPLEATGLALSSLR.L	2	2.3273	P08571
	CD14_HUMAN	K.ITGTMPPLEATGLALSSLR.L	2	2.4496	P08571
<b><i>Myeloid cell nuclear differentiation antigen - Homo sapiens (Human)</i></b>					
	MNDA_HUMAN	K.NTIYEIQDNTGSM*DVVGSGK.W	2	3.5267	P41218
	MNDA_HUMAN	K.NTIYEIQDNTGSM DVVGSGK.W	2	6.8274	P41218
	MNDA_HUMAN	K.NTIYEIQDNTGSM DVVGSGK.W	2	7.0407	P41218
	MNDA_HUMAN	K.EASSVDFNQNFV PNR.I	2	2.532	P41218
	MNDA_HUMAN	K.EASSVDFNQNFV PNR.I	2	2.9611	P41218
	MNDA_HUMAN	K.NTIYEIQDNTGSM DVVGSGK.W	2	6.5699	P41218
	MNDA_HUMAN	K.NTIYEIQDNTGSM*DVVGSGK.W	2	4.3805	P41218
<b><i>Myeloperoxidase precursor - Homo sapiens (Human)</i></b>					
	PERM_HUMAN	R.NQINALTSFVDASMVY GSEEPLAR.N	2	4.1234	P05164

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	PERM_HUMAN	R.FPTDQLTPDQER.S	2	4.0008	P05164
	PERM_HUMAN	R.FPTDQLTPDQER.S	2	3.3846	P05164
	PERM_HUMAN	R.NQINALTSFVDASMYGSEEPLAR.N	2	3.4304	P05164
<b><i>Myosin-10 - Homo sapiens (Human)</i></b>					
	MYH10_HUMAN	R.EDQSILC#TGESGAGK.T	2	3.3817	P35580
	MYH10_HUMAN	R.TQLEEELEDELQATEDAK.L	2	2.9523	P35580
	MYH10_HUMAN	R.EDQSILC#TGESGAGK.T	2	3.3029	P35580
	MYH10_HUMAN	R.TQLEEELEDELQATEDAK.L	2	3.8174	P35580
	MYH10_HUMAN	R.EDQSILC#TGESGAGK.T	2	2.85	P35580
	MYH10_HUMAN	R.EDQSILC#TGESGAGK.T	2	3.2919	P35580
	MYH10_HUMAN	R.TQLEEELEDELQATEDAK.L	2	3.2573	P35580
	MYH10_HUMAN	R.TQLEEELEDELQATEDAK.L	2	6.5382	P35580
	MYH10_HUMAN	R.EDQSILC#TGESGAGK.T	2	3.2823	P35580
<b><i>Myosin-11 - Homo sapiens (Human)</i></b>					
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.9175	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.5718	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.2302	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.108	P35749
<b><i>Myosin-9 - Homo sapiens (Human)</i></b>					
	MYH9_HUMAN	R.QLEEAEEEEAQR.A	2	2.5582	P35579
	MYH9_HUMAN	K.VEAQLQELQVK.F	2	2.9416	P35579
	MYH9_HUMAN	K.DVLLQVDDERR.N	2	2.7101	P35579
	MYH9_HUMAN	K.HEAMITDLEER.L	2	3.7198	P35579
	MYH9_HUMAN	K.SM*EAEMIQLQEELAAAER.A	2	3.9467	P35579
	MYH9_HUMAN	R.DELADEIANSSGK.G	2	3.0464	P35579
	MYH9_HUMAN	K.DLEAHIDSANK.N	2	3.6401	P35579
	MYH9_HUMAN	K.DMFQETMEAMR.I	2	2.6605	P35579
	MYH9_HUMAN	K.SMEAEMIQLQEELAAAER.A	2	2.4081	P35579
	MYH9_HUMAN	K.DFSALESQQLQDTQELLQEENR.Q	2	5.6836	P35579
	MYH9_HUMAN	K.DFSALESQQLQDTQELLQEENR.Q	2	3.9658	P35579
	MYH9_HUMAN	R.IIGLDQVAGMSETALPGAFK.T	2	2.7328	P35579
	MYH9_HUMAN	R.ELESQISELQEDLESER.A	2	2.9971	P35579
	MYH9_HUMAN	R.IAQLLEEELEEEQGNTELELINDR.L	2	4.7548	P35579
	MYH9_HUMAN	R.IAEFTTNLTTEEEK.S	2	2.5372	P35579
	MYH9_HUMAN	R.DLGEELEALKTELEDTLSTAAQQELR.S	2	4.4676	P35579

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
MYH9_HUMAN		R.KLEGDSTDLSDQIAELQAQIAELK.M	2	5.775	P35579
MYH9_HUMAN		K.DFSALESQIQDTQELLQEENR.Q	2	2.3962	P35579
MYH9_HUMAN		R.LQQELDLLVLDLHQR.Q	2	2.4759	P35579
MYH9_HUMAN		K.M*QQNIQELEEQLLEEEESAR.Q	2	3.419	P35579
MYH9_HUMAN		R.IAEFTTNLTETEEEEK.S	2	3.6025	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTK.R	2	3.9934	P35579
MYH9_HUMAN		K.MQQNIQELEEQLLEEEESAR.Q	2	5.2729	P35579
MYH9_HUMAN		R.VISGVLQLGNIVFK.K	2	3.4905	P35579
MYH9_HUMAN		K.LEEEQIILEDQNC#K.L	2	3.6149	P35579
MYH9_HUMAN		K.IAQLEEQLDNETK.E	2	3.6852	P35579
MYH9_HUMAN		K.KLEEEQIILEDQNC#K.L	2	4.6102	P35579
MYH9_HUMAN		R.ELEDATETADAMNR.E	2	4.5679	P35579
MYH9_HUMAN		R.ELEDATETADAMNR.E	2	3.8141	P35579
MYH9_HUMAN		R.ELEDATETADAM*NR.E	2	3.422	P35579
MYH9_HUMAN		K.TELEDTLSTAAQQELR.S	2	3.5022	P35579
MYH9_HUMAN		R.QLEEAEEEAQR.A	2	2.707	P35579
MYH9_HUMAN		R.ELEDATETADAMNR.E	2	4.0365	P35579
MYH9_HUMAN		K.HEAMITDLEER.L	2	3.2398	P35579
MYH9_HUMAN		K.KLEEEQIILEDQNC#K.L	3	3.7526	P35579
MYH9_HUMAN		K.KLEEEQIILEDQNC#K.L	2	4.4599	P35579
MYH9_HUMAN		K.IAQLEEQLDNETK.E	2	3.0718	P35579
MYH9_HUMAN		R.IAEFTTNLTETEEEEK.S	2	3.1756	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTKR.V	3	5.0598	P35579
MYH9_HUMAN		K.MQQNIQELEEQLLEEEESAR.Q	2	5.3118	P35579
MYH9_HUMAN		R.IIGLDQVAGMSETALPGAFK.T	2	2.9863	P35579
MYH9_HUMAN		K.DFSALESQIQDTQELLQEENR.Q	2	4.2362	P35579
MYH9_HUMAN		K.SMEAEMIQLQEELAAAER.A	2	3.8788	P35579
MYH9_HUMAN		K.SMEAEMIQLQEELAAAER.A	3	4.9751	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTK.R	2	2.8903	P35579
MYH9_HUMAN		R.DLGEELEALKTELEDTLSTAAQQELR.S	2	5.1704	P35579
MYH9_HUMAN		K.DFSALESQIQDTQELLQEENR.Q	2	4.4825	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTK.R	3	3.8165	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTKR.V	3	4.5333	P35579
MYH9_HUMAN		K.LEEEQIILEDQNC#K.L	2	2.9534	P35579
MYH9_HUMAN		K.KLEEEQIILEDQNC#K.L	2	4.8643	P35579
MYH9_HUMAN		K.LEGDSTDLSDQIAELQAQIAELK.M	2	5.3247	P35579

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
MYH9_HUMAN		R.ELEDATETADAMNR.E	2	3.3633	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTK.R	2	2.6898	P35579
MYH9_HUMAN		K.MQQNIQELEEQLLEEEESAR.Q	2	5.3785	P35579
MYH9_HUMAN		K.HSQAVEELAEQLEQTKR.V	3	5.0031	P35579
MYH9_HUMAN		K.LEEEQIILEDQNC#K.L	2	3.5254	P35579
MYH9_HUMAN		K.VEAQLQELQVK.F	2	2.7609	P35579
MYH9_HUMAN		K.HEAMITDLEER.L	2	3.6582	P35579
MYH9_HUMAN		R.ELEDATETADAMNR.E	2	3.5708	P35579
MYH9_HUMAN		K.HEAMITDLEER.L	2	3.5029	P35579
<b><i>Myosin-Ie - Homo sapiens (Human)</i></b>					
MYO1E_HUMAN		K.DIILQSNPLLEAFGNAK.T	2	3.2805	Q12965
MYO1E_HUMAN		K.DIILQSNPLLEAFGNAK.T	2	3.7902	Q12965
<b><i>Myosin-If - Homo sapiens (Human)</i></b>					
MYO1F_HUMAN		K.NPESLFLLEEVR.E	2	2.778	O00160
MYO1F_HUMAN		R.QDDFFILQEDAADSFLSVFK.T	2	3.9469	O00160
MYO1F_HUMAN		R.DVLFSDLIELMQTSEQAFLR.M	3	4.1345	O00160
MYO1F_HUMAN		K.NPESLFLLEEVR.E	2	3.9311	O00160
MYO1F_HUMAN		R.DVLFSDLIELMQTSEQAFLR.M	3	4.367	O00160
<b><i>Myosin-IG - Homo sapiens (Human)</i></b>					
Q6ZNK5_HUMA		R.DYLSSATDNPTASSLFAQR.L	2	3.9285	Q6ZNK5
Q6ZNK5_HUMA		R.DYLSSATDNPTASSLFAQR.L	2	4.1076	Q6ZNK5
<b><i>NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitochondrial precursor -</i></b>					
NDUA9_HUMAN		K.LPHLPGLLEDLGIQATPLELK.A	3	4.4917	Q16795
<b><i>NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 - Homo sapiens (Human)</i></b>					
NDUBA_HUMAN		R.DYKVDQEIIINIMQDR.L	2	3.7328	O96000
NDUBA_HUMAN		K.VDQEIIINIMQDR.L	2	2.5659	O96000
<b><i>NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 - Homo sapiens (Human)</i></b>					
NDUB4_HUMAN		R.TLPETLDPAEYNISPETR.R	2	3.5661	O95168
NDUB4_HUMAN		R.TLPETLDPAEYNISPETR.R	2	2.6037	O95168
<b><i>NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrial precursor - H</i></b>					
NDUB8_HUMAN		R.VEDYEPYPDDGMGYGDYPK.L	2	2.988	O95169
NDUB8_HUMAN		R.VEDYEPYPDDGM*GYGDYPK.L	2	2.3512	O95169
<b><i>NADH-cytochrome b5 reductase 3 - Homo sapiens (Human)</i></b>					
NB5R3_HUMAN		R.APEAWDYGGQGFVNEEMIR.D	2	4.5641	P00387



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEM*IR.D	2	3.6829	P00387
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEMIR.D	2	3.87	P00387
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEM*IR.D	2	3.8219	P00387
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEM*IR.D	2	3.5935	P00387
	NB5R3_HUMAN	K.DILLRPELEELR.N	2	2.5282	P00387
	NB5R3_HUMAN	K.DILLRPELEELR.N	2	2.3275	P00387
	NB5R3_HUMAN	R.GPSGLLVYQ GK.G	2	2.3944	P00387
	NB5R3_HUMAN	R.GPSGLLVYQ GK.G	2	3.1994	P00387
	NB5R3_HUMAN	K.DILLRPELEELR.N	2	2.5662	P00387
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEMIR.D	2	4.6346	P00387
	NB5R3_HUMAN	R.APEAWDYQGQGFVNEEMIR.D	2	4.0611	P00387
	NB5R3_HUMAN	K.DILLRPELEELR.N	2	2.8941	P00387
	NB5R3_HUMAN	K.DILLRPELEELR.N	2	2.3631	P00387
<b><i>Nck-associated protein 1-like - Homo sapiens (Human)</i></b>					
	NCKPL_HUMAN	K.TPEDYADSSIAELLFLEGLR.S	2	2.83	P55160
<b><i>Neurobeachin-like protein 2 - Homo sapiens (Human)</i></b>					
	Q6ZNJ1_HUMAN	R.PGLDSEPGGAEAGK.A	2	2.3633	Q6ZNJ1
<b><i>Neutrophil cytosol factor 2 - Homo sapiens (Human)</i></b>					
	NCF2_HUMAN	R.IHPQQPQEESSPQSDIPAPPSSK.A	3	3.7625	P19878
	NCF2_HUMAN	R.IHPQQPQEESSPQSDIPAPPSSK.A	3	3.9545	P19878
<b><i>Niban protein - Homo sapiens (Human)</i></b>					
	NIBA_HUMAN	K.NDYAVESYENK.E	2	2.5658	Q9BZQ8
<b><i>Nuclear pore membrane glycoprotein 210 precursor - Homo sapiens (Human)</i></b>					
	PO210_HUMAN	R.ELSDEIQVQVFEK.L	2	2.6955	Q8TEM1
	PO210_HUMAN	R.GVAIGQTSLTASVTNK.A	2	4.1943	Q8TEM1
	PO210_HUMAN	R.VFGAPEVLENLEVK.S	2	3.4208	Q8TEM1
	PO210_HUMAN	R.ELSDEIQVQVFEK.L	2	2.3053	Q8TEM1
	PO210_HUMAN	R.GVAIGQTSLTASVTNK.A	2	4.0659	Q8TEM1
	PO210_HUMAN	R.VFGAPEVLENLEVK.S	2	3.2971	Q8TEM1
	PO210_HUMAN	R.LPSQYNFAMNVLGR.V	2	2.8468	Q8TEM1
	PO210_HUMAN	R.GVAIGQTSLTASVTNK.A	2	4.2601	Q8TEM1
	PO210_HUMAN	R.ELYLEDSPLELK.I	2	2.9224	Q8TEM1
<b><i>PDCD6IP protein - Homo sapiens (Human)</i></b>					
	Q6NUS1_HUMA	R.LLDEEEATDNDLR.A	2	2.5938	Q6NUS1

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Peptidyl-prolyl cis-trans isomerase-like 4 - Homo sapiens (Human)</i></b>					
	PPIL4_HUMAN	R.IHVDFSQSVAKVK.W	2	2.433	Q8WUA2
<b><i>Peripherin - Homo sapiens (Human)</i></b>					
	PERI_HUMAN	K.NLQEAEWYK.S	2	2.8437	P41219
	PERI_HUMAN	K.NLQEAEWYK.S	2	2.4827	P41219
<b><i>Peroxiredoxin-6 - Homo sapiens (Human)</i></b>					
	PRDX6_HUMAN	R.ELAILLGMLDPAEKDEK.G	2	2.3487	P30041
	PRDX6_HUMAN	K.LPFPIDDR.N	2	2.5496	P30041
<b><i>Phosphate carrier protein, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	MPCP_HUMAN	R.LPRPPPEMPESLK.K	3	4.1846	Q00325
	MPCP_HUMAN	R.LPRPPPEMPESLKK.K	3	4.0734	Q00325
<b><i>Plasma membrane calcium-transporting ATPase 4 - Homo sapiens (Human)</i></b>					
	AT2B4_HUMAN	R.TPLLDEEEENPDK.A	2	2.7298	P23634
<b><i>Plastin-2 - Homo sapiens (Human)</i></b>					
	PLSL_HUMAN	K.ISTSLPVLDLIDAIQPGSINYDLLK.T	2	3.0177	P13796
	PLSL_HUMAN	R.EITENLMATGDLQDGR.I	2	2.6133	P13796
	PLSL_HUMAN	K.ISTSLPVLDLIDAIQPGSINYDLLK.T	2	3.9544	P13796
	PLSL_HUMAN	K.VNDDIIVNWVNETLR.E	2	3.0046	P13796
	PLSL_HUMAN	R.EITENLMATGDLQDGR.I	2	3.6613	P13796
<b><i>Platelet endothelial cell adhesion molecule precursor - Homo sapiens (Human)</i></b>					
	PECA1_HUMAN	K.SELVTVTESFSTPK.F	2	2.8118	P16284
	PECA1_HUMAN	K.SELVTVTESFSTPK.F	2	2.3593	P16284
	PECA1_HUMAN	R.DQNFVILEFPVEEQDR.V	2	2.9777	P16284
	PECA1_HUMAN	R.DQNFVILEFPVEEQDR.V	2	3.2765	P16284
	PECA1_HUMAN	K.EDTIVSQTQDFTK.I	2	3.7885	P16284
	PECA1_HUMAN	K.EDTIVSQTQDFTK.I	2	2.8659	P16284
	PECA1_HUMAN	K.EDTIVSQTQDFTK.I	2	3.7456	P16284
<b><i>Platelet glycoprotein 4 - Homo sapiens (Human)</i></b>					
	CD36_HUMAN	K.QVVLEEGTIAFK.N	2	2.4969	P16671
<b><i>Plectin-1 - Homo sapiens (Human)</i></b>					
	PLEC1_HUMAN	R.LLDPEDVDVPQPDEK.S	2	2.8764	Q15149
	PLEC1_HUMAN	R.FPSSFEEIEILWSQFLK.F	2	3.5387	Q15149
	PLEC1_HUMAN	R.RPELEDSTLR.Y	2	2.6883	Q15149
	PLEC1_HUMAN	R.SLESLHSFVAAATK.E	2	2.307	Q15149

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Plexin-B2 precursor - Homo sapiens (Human)</i></b>					
	PLXB2_HUMAN	K.VGVSQQPEDSQDLPGER.H	2	2.6563	O15031
	PLXB2_HUMAN	K.VGVSQQPEDSQDLPGER.H	2	3.1723	O15031
<b><i>PNAS-117 - Homo sapiens (Human)</i></b>					
	Q9HB66_HUMA	K.NDDIPEQDSLGLSNLQK.S	2	3.2392	Q9HB66
<b><i>Polypeptide N-acetylgalactosaminyltransferase 2 - Homo sapiens (Human)</i></b>					
	GALT2_HUMAN	K.EIILVDDYSNDPEDGALLGK.I	2	2.6426	Q10471
<b><i>Polypyrimidine tract-binding protein 1 - Homo sapiens (Human)</i></b>					
	PTBP1_HUMAN	K.LPIDVTEGEVISLGLPFGK.V	2	2.8523	P26599
	PTBP1_HUMAN	K.LPIDVTEGEVISLGLPFGK.V	2	3.0954	P26599
<b><i>PRA1 family protein 3 - Homo sapiens (Human)</i></b>					
	PRAF3_HUMAN	R.TPMGIVLDALEQQEEGINR.L	2	4.0004	O75915
	PRAF3_HUMAN	R.TPMGIVLDALEQQEEGINR.L	2	2.9833	O75915
	PRAF3_HUMAN	R.TPMGIVLDALEQQEEGINR.L	2	4.2085	O75915
	PRAF3_HUMAN	R.TPMGIVLDALEQQEEGINR.L	2	3.6452	O75915
<b><i>Probable E3 ubiquitin-protein ligase KIAA1333 - Homo sapiens (Human)</i></b>					
	K1333_HUMAN	K.KHVL PNSNNGITDCLLEESSPK.L	2	2.8615	Q7L622
<b><i>Probable phospholipid-transporting ATPase IIA - Homo sapiens (Human)</i></b>					
	ATP9A_HUMAN	K.EVNSQVYSRLTAR.G	2	2.482	O75110
<b><i>Prohibitin - Homo sapiens (Human)</i></b>					
	PHB_HUMAN	K.DLQNVNITLR.I	2	2.9235	P35232
	PHB_HUMAN	K.DLQNVNITLR.I	2	2.4772	P35232
<b><i>Prolactin - Homo sapiens (Human)</i></b>					
	Q5I0G2_HUMAN	R.GM*QEAPAILSKAVEIEEQTKR.L	2	2.6141	Q5I0G2
<b><i>Prostaglandin G/H synthase 1 precursor - Homo sapiens (Human)</i></b>					
	PGH1_HUMAN	R.VPDASQDDGPAVERPSTEL.-	2	3.1381	P23219
<b><i>Proteasome activator complex subunit 1 - Homo sapiens (Human)</i></b>					
	PSME1_HUMAN	R.IEDGNFVGVAQEK.V	2	3.3336	Q06323
<b><i>Proteasome activator complex subunit 2 - Homo sapiens (Human)</i></b>					
	PSME2_HUMAN	K.IEDGNDFGVAIQEK.V	2	4.0667	Q9UL46
<b><i>Proteasome subunit alpha type 5 - Homo sapiens (Human)</i></b>					
	PSA5_HUMAN	R.PFGVALLFGGVDEK.G	2	3.3313	P28066
	PSA5_HUMAN	R.PFGVALLFGGVDEK.G	2	2.5205	P28066

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Proteasome-associated protein ECM29 homolog - Homo sapiens (Human)</i></b>					
	ECM29_HUMAN	K.DIALVQQLFEALC#K.E	2	2.6854	Q5VYK3
<b><i>Protein FAM49A - Homo sapiens (Human)</i></b>					
	FA49A_HUMAN	R.NPAIQNDFSYYR.R	2	3.2224	Q9H0Q0
	FA49A_HUMAN	R.NPAIQNDFSYYR.R	2	3.1743	Q9H0Q0
<b><i>Protein FAM49B - Homo sapiens (Human)</i></b>					
	FA49B_HUMAN	R.INNVPAEGENEVNNELANR.M	2	3.8214	Q9NUQ9
	FA49B_HUMAN	R.INNVPAEGENEVNNELANR.M	2	2.909	Q9NUQ9
	FA49B_HUMAN	R.INNVPAEGENEVNNELANR.M	2	3.5802	Q9NUQ9
<b><i>Protein flightless-1 homolog - Homo sapiens (Human)</i></b>					
	FLII_HUMAN	R.VPEC#LYTLPSLR.R	2	2.3969	Q13045
<b><i>Protein LYRIC - Homo sapiens (Human)</i></b>					
	LYRIC_HUMAN	R.SWQDELAQQAEEGSAR.L	2	3.8325	Q86UE4
	LYRIC_HUMAN	R.SWQDELAQQAEEGSAR.L	2	3.4109	Q86UE4
	LYRIC_HUMAN	R.SWQDELAQQAEEGSAR.L	2	3.4672	Q86UE4
<b><i>Protein MYSM1 - Homo sapiens (Human)</i></b>					
	MYSM1_HUMAN	K.YLNKTSVRPGLK.N	2	2.3645	Q5VVJ2
	MYSM1_HUMAN	K.YLNKTSVRPGLK.N	2	2.4927	Q5VVJ2
<b><i>Protein S100-A10 - Homo sapiens (Human)</i></b>					
	S10AA_HUMAN	K.EFPGFLENQKDPLAVDK.I	2	2.528	P60903
<b><i>Protein S100-A9 - Homo sapiens (Human)</i></b>					
	S10A9_HUMAN	R.NIETIINTFHQYSVK.L	2	3.0307	P06702
	S10A9_HUMAN	R.NIETIINTFHQYSVK.L	2	2.5724	P06702
	S10A9_HUMAN	K.VIEHIMEDLDTNADK.Q	2	4.075	P06702
	S10A9_HUMAN	R.NIETIINTFHQYSVK.L	2	3.6855	P06702
	S10A9_HUMAN	K.VIEHIMEDLDTNADK.Q	2	2.801	P06702
	S10A9_HUMAN	R.NIETIINTFHQYSVK.L	2	3.2921	P06702
	S10A9_HUMAN	K.VIEHIMEDLDTNADK.Q	2	3.8949	P06702
	S10A9_HUMAN	R.NIETIINTFHQYSVK.L	2	3.3053	P06702
<b><i>Protein-arginine deiminase type-4 - Homo sapiens (Human)</i></b>					
	PADI4_HUMAN	R.ELGLAESDIIDIPQLFK.L	2	3.5022	Q9UM07
	PADI4_HUMAN	R.ELGLAESDIIDIPQLFK.L	2	2.8027	Q9UM07
<b><i>Proteolipid protein 2 - Homo sapiens (Human)</i></b>					
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.3044	Q04941

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.5675	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.5419	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.4839	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.8498	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	3.0994	Q04941
<b><i>Putative eukaryotic translation initiation factor 3 subunit - Homo sapiens (Human)</i></b>					
	IF3X_HUMAN	R.FLENALAVSTKYHGPK.A	2	2.3743	O75153
<b><i>Putative FelC - Homo sapiens (Human)</i></b>					
	Q13718_HUMAN	-.M*AAQSSLYNDDRNLLR.I	2	2.4014	Q13718
<b><i>Putative uncharacterized protein - Homo sapiens (Human)</i></b>					
	A5PLL5_HUMAN	R.MPTTQETDGFQVK.R	2	2.3961	A5PLL5
<b><i>Putative uncharacterized protein DKFZp667I133 - Homo sapiens (Human)</i></b>					
	Q8ND71_HUMA	R.FQDC#VNEAASQEGDKPQGPR.E	3	3.9934	Q8ND71
<b><i>Pyruvate kinase isozymes M1/M2 - Homo sapiens (Human)</i></b>					
	KPYM_HUMAN	R.LAPITSDPTEATAVGAVEASF.K	2	3.5465	P14618
<b><i>Rab11 family-interacting protein 4 - Homo sapiens (Human)</i></b>					
	RFIP4_HUMAN	K.KVTELENDSLTNGDLK.S	2	2.4623	Q86YS3
<b><i>Ras GTPase-activating-like protein IQGAP1 - Homo sapiens (Human)</i></b>					
	IQGA1_HUMAN	R.EEIQSSISGVTAAYNR.E	2	3.3718	P46940
	IQGA1_HUMAN	K.VDFTEEEINNMK.T	2	2.3191	P46940
	IQGA1_HUMAN	K.VNTFSALANIDLALAQGDALALFR.A	2	4.1825	P46940
	IQGA1_HUMAN	K.LPYDVTPEQALAHAEVK.T	3	3.7506	P46940
	IQGA1_HUMAN	K.SWVNQM*ESQTGEASK.L	2	2.4431	P46940
	IQGA1_HUMAN	K.FDVPGDENAEMDAR.T	2	2.8816	P46940
	IQGA1_HUMAN	R.ILAIGLINEALDEGDAQK.T	2	5.6551	P46940
	IQGA1_HUMAN	K.VNTFSALANIDLALAQGDALALFR.A	2	2.5769	P46940
	IQGA1_HUMAN	K.FALGIFAINAEVSGDVGK.T	2	3.4599	P46940
	IQGA1_HUMAN	K.FDVPGDENAEMDAR.T	2	3.167	P46940
	IQGA1_HUMAN	K.NVIFEISPTTEEVGDFEVK.A	2	2.9897	P46940
	IQGA1_HUMAN	K.FALGIFAINAEVSGDVGK.T	2	3.0735	P46940
	IQGA1_HUMAN	R.ILAIGLINEALDEGDAQK.T	2	3.8248	P46940
	IQGA1_HUMAN	K.FALGIFAINAEVSGDVGK.T	2	2.7108	P46940
	IQGA1_HUMAN	K.FALGIFAINAEVSGDVGK.T	2	3.5592	P46940
	IQGA1_HUMAN	R.ILAIGLINEALDEGDAQK.T	2	3.1648	P46940

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	IQGA1_HUMAN	K.NVIFEISPTEEVGDFEVK.A	2	5.3172	P46940
	IQGA1_HUMAN	R.ILAIGLINEALDEGDAQK.T	3	4.7887	P46940
	IQGA1_HUMAN	K.NVIFEISPTEEVGDFEVK.A	2	2.6884	P46940
	IQGA1_HUMAN	K.FDVPGDENAEMDAR.T	2	3.0708	P46940
	IQGA1_HUMAN	K.FALGIFAINAEVESGDVGK.T	2	3.7686	P46940
	IQGA1_HUMAN	K.VNTFSALANIDLALEQQDALALFR.A	2	4.0653	P46940
	IQGA1_HUMAN	K.EELQSGVDAANSAAQQYQR.R	2	4.395	P46940
	IQGA1_HUMAN	K.FDVPGDENAEMDAR.T	2	3.2561	P46940
	IQGA1_HUMAN	K.FDVPGDENAEMDAR.T	2	2.6803	P46940
	IQGA1_HUMAN	K.LPYDVTPEQALAHEEVK.T	3	4.9969	P46940
	IQGA1_HUMAN	K.EELQSGVDAANSAAQQYQR.R	2	5.2487	P46940
<b><i>Ras GTPase-activating-like protein IQGAP2 - Homo sapiens (Human)</i></b>					
	IQGA2_HUMAN	K.LPYDVTTEQALTYPEVK.N	2	4.1092	Q13576
	IQGA2_HUMAN	K.LPYDVTTEQALTYPEVK.N	2	3.9454	Q13576
<b><i>RAS guanyl-releasing protein 4 - Homo sapiens (Human)</i></b>					
	Q8TDF6_HUMA	R.HPEVMHQDPQLEEVIGR.F	2	2.3194	Q8TDF6
<b><i>Ras-related protein Rab-10 - Homo sapiens (Human)</i></b>					
	RAB10_HUMAN	R.NIDEHANEDVER.M	2	2.8393	P61026
<b><i>Ras-related protein Rab-11B - Homo sapiens (Human)</i></b>					
	RB11B_HUMAN	K.NNLSFIETSALDSTNVVEEAFK.N	2	2.656	Q15907
<b><i>Ras-related protein Rab-14 - Homo sapiens (Human)</i></b>					
	RAB14_HUMAN	K.TGENVEDAFLEAAK.K	2	2.5646	P61106
	RAB14_HUMAN	K.TGENVEDAFLEAAK.K	2	3.1756	P61106
<b><i>Ras-related protein Rab-1A - Homo sapiens (Human)</i></b>					
	RAB1A_HUMAN	R.FADDTYTESYISTIGVDFK.I	2	2.8032	P62820
	RAB1A_HUMAN	R.FADDTYTESYISTIGVDFK.I	2	2.7021	P62820
	RAB1A_HUMAN	R.FADDTYTESYISTIGVDFK.I	2	2.4796	P62820
<b><i>Ras-related protein Rab-2A - Homo sapiens (Human)</i></b>					
	RAB2A_HUMAN	K.TASNVEEAFINTAK.E	2	3.1125	P61019
<b><i>Ras-related protein Rab-35 - Homo sapiens (Human)</i></b>					
	RAB35_HUMAN	K.ENVNVEEMFNC#ITELVLR.A	2	3.3701	Q15286
<b><i>Ras-related protein Rab-3A - Homo sapiens (Human)</i></b>					
	RAB3A_HUMAN	R.YADDSFTPAFVSTVGIDFK.V	2	2.3544	P20336
<b><i>Ras-related protein Rab-3D - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	RAB3D_HUMAN	R.DAADQNFDYMFK.L	2	2.3637	Q95716
<b><i>Ras-related protein Rab-5C - Homo sapiens (Human)</i></b>					
	RAB5C_HUMAN	R.GVDLQENNPASR.S	2	2.9106	P51148
<b><i>Ras-related protein Rab-8A - Homo sapiens (Human)</i></b>					
	RAB8A_HUMAN	R.NIEEHASADVEK.M	2	3.5976	P61006
<b><i>Ras-related protein Rab-8B - Homo sapiens (Human)</i></b>					
	RAB8B_HUMAN	R.NIEEHASSDVER.M	2	2.5892	Q92930
<b><i>Receptor expression-enhancing protein 5 - Homo sapiens (Human)</i></b>					
	REEP5_HUMAN	K.HESQMDSVVK.D	2	2.8247	Q00765
	REEP5_HUMAN	K.HESQMDSVVK.D	2	2.3422	Q00765
<b><i>RER1 retention in endoplasmic reticulum 1 homolog - Homo sapiens (Human)</i></b>					
	Q5T091_HUMAN	K.VDPSLMEDSDDGPSLPTK.Q	2	3.6413	Q5T091
	Q5T091_HUMAN	K.VDPSLMEDSDDGPSLPTK.Q	2	4.6029	Q5T091
<b><i>Reticulon-4 - Homo sapiens (Human)</i></b>					
	RTN4_HUMAN	K.NVKDAM*AKIQAK.I	2	2.3027	Q9NQC3
<b><i>Retinal dehydrogenase 1 - Homo sapiens (Human)</i></b>					
	AL1A1_HUMAN	R.DRLLLATM*ESMNGGK.L	2	2.699	P00352
	AL1A1_HUMAN	R.DRLLLATM*ESMNGGK.L	2	3.0847	P00352
<b><i>Rho GDP-dissociation inhibitor 2 - Homo sapiens (Human)</i></b>					
	GDIS_HUMAN	K.APEPHVEEDDDDELDSK.L	3	4.2857	P52566
	GDIS_HUMAN	K.APEPHVEEDDDDELDSK.L	3	3.8527	P52566
	GDIS_HUMAN	K.APEPHVEEDDDDELDSK.L	3	3.9552	P52566
	GDIS_HUMAN	K.TLLGDGPVVTDPK.A	2	2.3597	P52566
<b><i>Rho GTPase-activating protein 4 - Homo sapiens (Human)</i></b>					
	RHG04_HUMAN	R.DLEQQLQDELLEVSELTAK.K	2	2.3196	P98171
<b><i>Rho guanine nucleotide exchange factor 1 - Homo sapiens (Human)</i></b>					
	ARHG1_HUMAN	R.QESGYLIEEIGDVLLAR.F	2	2.3621	Q92888
	ARHG1_HUMAN	R.TQEIQENLLSLEETMK.Q	2	3.8285	Q92888
<b><i>Rho/rac guanine nucleotide exchange factor - Homo sapiens (Human)</i></b>					
	Q5VY92_HUMA	K.ELLSNVDEGIYQLEK.G	2	3.2928	Q5VY92
	Q5VY92_HUMA	K.ELLSNVDEGIYQLEK.G	2	2.5413	Q5VY92
<b><i>Rho-related GTP-binding protein RhoB precursor - Homo sapiens (Human)</i></b>					
	RHOB_HUMAN	K.DEFPEVYVPTVFENYVADIEVDGK.Q	2	3.2169	P62745

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Rho-related GTP-binding protein RhoG precursor - Homo sapiens (Human)</i></b>					
	RHOG_HUMAN	R.LKEQGQAPITPQQGQALAK.Q	3	3.9334	P84095
	RHOG_HUMAN	R.YLEC#SALQQDGVK.E	2	3.3897	P84095
	RHOG_HUMAN	R.YLEC#SALQQDGVK.E	2	3.3702	P84095
<b><i>Ribonuclease inhibitor - Homo sapiens (Human)</i></b>					
	RINI_HUMAN	R.ELC#QGLGQPGSVLR.V	2	2.6696	P13489
	RINI_HUMAN	K.ELSLAGNELGDEGAR.L	2	3.582	P13489
	RINI_HUMAN	R.ELC#QGLGQPGSVLR.V	2	2.5281	P13489
	RINI_HUMAN	K.ELSLAGNELGDEGAR.L	2	3.3381	P13489
<b><i>RING finger protein 13 - Homo sapiens (Human)</i></b>					
	RNF13_HUMAN	R.LDC#NFDIKVLNAQRAGYK.A	2	2.5784	O43567
<b><i>RING finger protein 213 - Homo sapiens (Human)</i></b>					
	RN213_HUMAN	R.VPQFSFLDIFPK.V	2	2.3131	Q63HN8
<b><i>Roundabout homolog 3 precursor - Homo sapiens (Human)</i></b>					
	ROBO3_HUMAN	R.PSFLSRGQGTSTCSTAGSNSSR.G	2	2.4104	Q96MS0
<b><i>RUN and FYVE domain-containing protein 1 - Homo sapiens (Human)</i></b>					
	RUFY1_HUMAN	K.ELRELQDEKAELQK.I	2	2.3532	Q96T51
<b><i>Sad1/unc-84-like protein 2 - Homo sapiens (Human)</i></b>					
	UN84B_HUMAN	R.DDVESQFPAWISQFLAR.G	2	2.573	Q9UH99
	UN84B_HUMAN	R.LEDQLAGLQQELAALALK.Q	2	4.2275	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	2	4.4353	Q9UH99
	UN84B_HUMAN	R.LEDQLAGLQQELAALALK.Q	2	4.9319	Q9UH99
	UN84B_HUMAN	R.LEDQLAGLQQELAALALK.Q	2	3.6537	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	2	4.6475	Q9UH99
	UN84B_HUMAN	R.DDVESQFPAWISQFLAR.G	2	3.5244	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	3	5.0681	Q9UH99
	UN84B_HUMAN	R.LEDQLAGLQQELAALALK.Q	2	2.9619	Q9UH99
	UN84B_HUMAN	R.LEDQLAGLQQELAALALK.Q	2	3.6562	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	3	4.1026	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	2	4.2884	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	3	5.253	Q9UH99
	UN84B_HUMAN	R.RLEDQLAGLQQELAALALK.Q	2	4.3892	Q9UH99
<b><i>SAM domain and HD domain-containing protein 1 - Homo sapiens (Human)</i></b>					
	SAMH1_HUMAN	K.VGNIIDTMITDAFLK.A	2	4.3886	Q9Y3Z3



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	SAMH1_HUMAN	R.FENLGVSSLGER.K	2	2.5668	Q9Y3Z3
	SAMH1_HUMAN	K.EWNDSTSVQNPTR.L	2	3.7049	Q9Y3Z3
	SAMH1_HUMAN	K.VGNIIDTMITDAFLK.A	2	4.1847	Q9Y3Z3
	SAMH1_HUMAN	R.DC#HHLGIQNNFDYK.R	3	3.9917	Q9Y3Z3
	SAMH1_HUMAN	R.FENLGVSSLGER.K	2	2.6105	Q9Y3Z3
	SAMH1_HUMAN	K.VGNIIDTMITDAFLK.A	2	4.1225	Q9Y3Z3
<b><i>Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 - Homo sapiens (Human)</i></b>					
	AT2A2_HUMAN	K.VGEATETALTC#LVEK.M	2	2.5526	P16615
	AT2A2_HUMAN	K.TGTLTTNQMSVC#R.M	2	3.0718	P16615
<b><i>Secretory carrier-associated membrane protein 2 - Homo sapiens (Human)</i></b>					
	SCAM2_HUMAN	R.TGASFQQAQEEFSQGIFSSR.T	2	4.9244	O15127
	SCAM2_HUMAN	R.TGASFQQAQEEFSQGIFSSR.T	2	4.4005	O15127
	SCAM2_HUMAN	R.TGASFQQAQEEFSQGIFSSR.T	2	3.1361	O15127
<b><i>Secretory carrier-associated membrane protein 3 - Homo sapiens (Human)</i></b>					
	SCAM3_HUMAN	K.NYGSYSTQASAAAATAELLK.K	2	3.2372	O14828
<b><i>Septin-9 - Homo sapiens (Human)</i></b>					
	SEPT9_HUMAN	K.APVDFGYVGIDSILEQMR.R	2	2.9614	Q9UHD8
<b><i>Serine palmitoyltransferase 2 - Homo sapiens (Human)</i></b>					
	LCB2_HUMAN	R.LVPLLLDRPFDETTYEETED.-	2	2.5016	O15270
<b><i>Serine/threonine-protein kinase 31 - Homo sapiens (Human)</i></b>					
	STK31_HUMAN	R.MKNLAAKM*EILK.E	2	2.4812	Q9BXU1
<b><i>Sestrin-1 - Homo sapiens (Human)</i></b>					
	SESN1_HUMAN	K.VYIKTVVC#TPEK.V	2	2.3656	Q9Y6P5
<b><i>SH2 domain-containing protein 2A - Homo sapiens (Human)</i></b>					
	SH2A_HUMAN	K.QGQAPVPMQKEGAGEK.E	2	2.4946	Q9NP31
<b><i>Sialic acid-binding Ig-like lectin 9 precursor - Homo sapiens (Human)</i></b>					
	SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A	2	3.0073	Q9Y336
	SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A	2	3.2946	Q9Y336
	SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A	2	3.3121	Q9Y336
	SIGL9_HUMAN	R.PAAGVGDTGIEDANAVR.G	2	4.4161	Q9Y336
<b><i>Sideroflexin-1 - Homo sapiens (Human)</i></b>					
	SFXN1_HUMAN	R.NILLTNEQLESAR.K	2	2.9443	Q9H9B4
<b><i>Sideroflexin-3 - Homo sapiens (Human)</i></b>					
	SFXN3_HUMAN	R.NLLLSGAQLEASR.N	2	2.8707	Q9BWM7

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	SFXN3_HUMAN	R.ELQVGIPVADEAGQR.L	2	2.4752	Q9BWM7
<b><i>Signal peptidase complex subunit 2 - Homo sapiens (Human)</i></b>					
	SPCS2_HUMAN	K.FFDHSGTLVMDAYEPEISR.L	3	4.6988	Q15005
	SPCS2_HUMAN	K.FFDHSGTLVMDAYEPEISR.L	2	3.239	Q15005
	SPCS2_HUMAN	K.DPTGMDPDDIWQLSSSLK.R	2	4.3524	Q15005
	SPCS2_HUMAN	K.FFDHSGTLVMDAYEPEISR.L	3	3.9417	Q15005
	SPCS2_HUMAN	K.DPTGMDPDDIWQLSSSLK.R	2	2.5085	Q15005
<b><i>Signal recognition particle receptor subunit beta - Homo sapiens (Human)</i></b>					
	SRPRB_HUMAN	R.GDVGSADIQDLEK.W	2	2.6232	Q9Y5M8
	SRPRB_HUMAN	K.DVAEFLYQVLIDSMGLK.N	3	3.8801	Q9Y5M8
<b><i>Signal transducer and activator of transcription 1-alpha/beta - Homo sapiens (Human)</i></b>					
	STAT1_HUMAN	K.SLEDLQDEYDFK.C	2	2.7764	P42224
	STAT1_HUMAN	K.SLEDLQDEYDFK.C	2	3.0401	P42224
<b><i>Signal transducer and activator of transcription 3 - Homo sapiens (Human)</i></b>					
	STAT3_HUMAN	R.GLSIEQLTTLAEK.L	2	2.6412	P40763
<b><i>Sister chromatid cohesion protein PDS5 homolog A - Homo sapiens (Human)</i></b>					
	PDS5A_HUMAN	K.SIEGTADDEEEGVSPDTAIR.S	2	3.5516	Q29RF7
	PDS5A_HUMAN	K.SIEGTADDEEEGVSPDTAIR.S	2	3.1146	Q29RF7
<b><i>Sodium/potassium-transporting ATPase subunit alpha-1 precursor - Homo sapiens (Human)</i></b>					
	AT1A1_HUMAN	K.GVGIISEGNETVEDIAAR.L	2	3.0033	P05023
	AT1A1_HUMAN	R.SPFTNENPLETR.N	2	3.5539	P05023
	AT1A1_HUMAN	K.GVGIISEGNETVEDIAAR.L	2	2.5543	P05023
	AT1A1_HUMAN	R.SPFTNENPLETR.N	2	3.5642	P05023
	AT1A1_HUMAN	R.SPFTNENPLETR.N	2	4.1722	P05023
	AT1A1_HUMAN	K.GVGIISEGNETVEDIAAR.L	2	3.8951	P05023
	AT1A1_HUMAN	K.VDNSSLTGESEPQTR.S	2	2.6003	P05023
<b><i>Solute carrier family 25 member 24. - Homo sapiens (Human)</i></b>					
	Q6NUK1_HUMA	K.HEGLGAFYK.G	2	2.5737	Q6NUK1
	Q6NUK1_HUMA	R.DYFLFNPVTDIEIIR.F	2	2.873	Q6NUK1
	Q6NUK1_HUMA	K.HSTGIDIGDSLTPDEFTEDEKK.S	3	3.9435	Q6NUK1
<b><i>Solute carrier family 30 - Homo sapiens (Human)</i></b>					
	Q5SQG7_HUMA	R.TPPLENSLPQC#YQR.V	2	2.4829	Q5SQG7
	Q5SQG7_HUMA	R.TPPLENSLPQC#YQR.V	2	2.5712	Q5SQG7
	Q5SQG7_HUMA	R.TPPLENSLPQC#YQR.V	2	3.0851	Q5SQG7

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Solute carrier family 39 member 11 - Homo sapiens (Human)</i></b>					
	Q8N1S5_HUMA	K.SDPEGPALLFPESELSIR.I	2	3.2091	Q8N1S5
	Q8N1S5_HUMA	K.SDPEGPALLFPESELSIR.I	2	3.0239	Q8N1S5
	Q8N1S5_HUMA	K.SDPEGPALLFPESELSIR.I	2	3.3659	Q8N1S5
<b><i>Sorting and assembly machinery component 50 homolog - Homo sapiens (Human)</i></b>					
	SAM50_HUMAN	K.DDIIC#EIGDVFK.A	2	3.1202	Q9Y512
	SAM50_HUMAN	K.DDIIC#EIGDVFK.A	2	3.1559	Q9Y512
	SAM50_HUMAN	K.DDIIC#EIGDVFK.A	2	3.4957	Q9Y512
<b><i>Spliceosome RNA helicase BAT1 - Homo sapiens (Human)</i></b>					
	UAP56_HUMAN	K.FMQDPMEIFVDETK.L	2	3.1638	Q13838
<b><i>Splicing factor 3B subunit 1 - Homo sapiens (Human)</i></b>					
	SF3B1_HUMAN	R.SLVEIIEHGLVDEQQK.V	2	2.4365	O75533
<b><i>Structural maintenance of chromosomes protein 6 - Homo sapiens (Human)</i></b>					
	SMC6_HUMAN	R.IEELKKSTDQSLEPER.L	2	2.5436	Q96SB8
	SMC6_HUMAN	R.IEELKKSTDQSLEPER.L	2	2.6797	Q96SB8
<b><i>Sulfide:quinone oxidoreductase, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	SQRD_HUMAN	K.VGAENVAIVEPSE.H	2	2.7918	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	2.596	Q9Y6N5
	SQRD_HUMAN	K.VGAENVAIVEPSE.H	2	2.8302	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	3.5034	Q9Y6N5
	SQRD_HUMAN	K.VGAENVAIVEPSE.H	2	3.1753	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	2.7729	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	3.1047	Q9Y6N5
	SQRD_HUMAN	K.VGAENVAIVEPSE.H	2	3.0286	Q9Y6N5
<b><i>Sulfotransferase 1A1 - Homo sapiens (Human)</i></b>					
	ST1A1_HUMAN	K.SGTTWVSQILDMIYQGGDLEK.C	2	3.5656	P50225
<b><i>Surfeit locus protein 4 - Homo sapiens (Human)</i></b>					
	SURF4_HUMAN	R.NLALGGGLLLLLAESR.S	2	3.7927	O15260
	SURF4_HUMAN	R.NLALGGGLLLLLAESR.S	2	2.3637	O15260
<b><i>Syntaxin-7 - Homo sapiens (Human)</i></b>					
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	2.7059	O15400
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	4.1909	O15400
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	3.2468	O15400
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	3.3931	O15400

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	3.1243	O15400
<b><i>Talin-1 - Homo sapiens (Human)</i></b>					
	TLN1_HUMAN	R.GVGAAATAVTQALNELLQHVK.A	2	4.7856	Q9Y490
	TLN1_HUMAN	R.GVAALTSDDPAVQAIVLDTASDVLDK.A	2	3.5189	Q9Y490
	TLN1_HUMAN	K.EVIQEWNLNLIK.R	2	2.8209	Q9Y490
	TLN1_HUMAN	K.LGAASLGAEDPETQVVLINAVK.D	2	2.497	Q9Y490
	TLN1_HUMAN	R.IPEAPAGPPSDFGLFLSDDDPK.K	2	3.3338	Q9Y490
<b><i>TBC1 domain family member 9B - Homo sapiens (Human)</i></b>					
	TBC9B_HUMAN	R.ALLSSSDPPAEVDIFELLK.V	2	2.6501	Q66K14
<b><i>T-complex protein 1 subunit delta - Homo sapiens (Human)</i></b>					
	TCPD_HUMAN	R.DALSDLALHFLNK.M	2	3.7624	P50991
<b><i>T-complex protein 1 subunit eta - Homo sapiens (Human)</i></b>					
	TCPH_HUMAN	K.LPIGDVATQYFADR.D	2	2.5176	Q99832
	TCPH_HUMAN	K.LPIGDVATQYFADR.D	2	2.6608	Q99832
<b><i>T-complex protein 1 subunit gamma - Homo sapiens (Human)</i></b>					
	TCPG_HUMAN	K.IPGGIIEDSC#VLR.G	2	3.0212	P49368
	TCPG_HUMAN	K.IPGGIIEDSC#VLR.G	2	3.3355	P49368
	TCPG_HUMAN	K.IPGGIIEDSC#VLR.G	2	3.3376	P49368
	TCPG_HUMAN	K.IPGGIIEDSC#VLR.G	2	3.3776	P49368
<b><i>Tetracycline transporter-like protein - Homo sapiens (Human)</i></b>					
	Q07706_HUMAN	R.DAADLLSPLALLR.F	2	2.5452	Q07706
<b><i>Tetraspanin-14 - Homo sapiens (Human)</i></b>					
	TSN14_HUMAN	R.DDIDLQNLIDSLQK.A	2	2.7516	Q8NG11
<b><i>Toll-like receptor 2 precursor - Homo sapiens (Human)</i></b>					
	TLR2_HUMAN	R.LFDENNDAAAILLLEPIEK.K	2	3.2337	O60603
	TLR2_HUMAN	R.LFDENNDAAAILLLEPIEK.K	2	3.8901	O60603
	TLR2_HUMAN	R.LFDENNDAAAILLLEPIEK.K	2	2.3502	O60603
<b><i>Toll-like receptor 3 precursor - Homo sapiens (Human)</i></b>					
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	2.7292	O15455
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	3.0688	O15455
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	2.5595	O15455
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	2.932	O15455
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	3.8711	O15455
	TLR3_HUMAN	K.LEPELCQKLPM*LK.V	2	3.0185	O15455

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Toll-like receptor 8 precursor - Homo sapiens (Human)</i></b>					
TLR8_HUMAN		R.DWDPGLAIIDNLMQSIQSK.K	2	2.9945	Q9NR97
<b><i>Torsin family protein C9orf167 - Homo sapiens (Human)</i></b>					
CI167_HUMAN		R.SAEAAAAQAEEDLR.A	2	2.5562	Q9NXH8
CI167_HUMAN		K.TPLLVLDDVELMPR.P	2	3.0429	Q9NXH8
CI167_HUMAN		K.TPLLVLDDVELMPR.P	2	3.253	Q9NXH8
<b><i>Translocon-associated protein subunit alpha precursor - Homo sapiens (Human)</i></b>					
SSRA_HUMAN		K.GEDFPANNIVK.F	2	2.3925	P43307
SSRA_HUMAN		K.GEDFPANNIVK.F	2	2.3094	P43307
<b><i>Transmembrane 9 superfamily protein member 3 precursor - Homo sapiens (Human)</i></b>					
TM9S3_HUMAN		R.NLSGQPNFPC#R.V	2	2.5307	Q9HD45
TM9S3_HUMAN		R.NLSGQPNFPC#R.V	2	2.4986	Q9HD45
<b><i>Transmembrane emp24 domain-containing protein 10 precursor - Homo sapiens (Human)</i></b>					
TMEDA_HUMAN		R.IPDQLVILDMK.H	2	3.0826	P49755
TMEDA_HUMAN		R.IPDQLVILDMK.H	2	2.845	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	2.8184	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	4.0883	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	3.6565	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	4.1025	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	3.4039	P49755
TMEDA_HUMAN		R.IPDQLVILDMK.H	2	3.3469	P49755
TMEDA_HUMAN		R.LEDLSESIVNDFAYMK.K	2	4.2376	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	3.2359	P49755
TMEDA_HUMAN		K.DLLVTGAYEISDQSGGAGGLR.S	2	4.1488	P49755
TMEDA_HUMAN		R.RLEDLSESIVNDFAYMK.K	2	4.2689	P49755
<b><i>Transmembrane emp24 domain-containing protein 2 precursor - Homo sapiens (Human)</i></b>					
TMED2_HUMAN		K.HEQEYMEVR.E	2	2.8977	Q15363
TMED2_HUMAN		K.HEQEYMEVR.E	2	2.8179	Q15363
TMED2_HUMAN		K.HEQEYMEVR.E	2	3.3781	Q15363
TMED2_HUMAN		K.HEQEYMEVR.E	2	3.1338	Q15363
<b><i>Transmembrane emp24 domain-containing protein 5 precursor - Homo sapiens (Human)</i></b>					
TMED5_HUMAN		K.LEDILESINSIK.S	2	3.4724	Q9Y3A6
TMED5_HUMAN		K.LEDILESINSIK.S	2	3.167	Q9Y3A6
TMED5_HUMAN		K.LEDILESINSIK.S	2	3.2693	Q9Y3A6

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	TMED5_HUMAN	K.LEDILESINSIK.S	2	3.3177	Q9Y3A6
	TMED5_HUMAN	K.LEDILESINSIK.S	2	2.9808	Q9Y3A6
<b><i>Transmembrane emp24 domain-containing protein 9 precursor - Homo sapiens (Human)</i></b>					
	TMED9_HUMAN	R.QLVEQVEQIQK.E	2	3.0989	Q9BVK6
<b><i>Transmembrane protease, serine 13 - Homo sapiens (Human)</i></b>					
	TMPD_HUMAN	R.AMTGRIVGGALASDSK.W	2	2.4976	Q9BYE2
<b><i>Transmembrane protein 49 - Homo sapiens (Human)</i></b>					
	TMM49_HUMAN	R.LSGAEPDDEEYQFEEMLEHAESAQDFASR.A	3	4.1371	Q96GC9
<b><i>Trifunctional enzyme subunit alpha, mitochondrial precursor - Homo sapiens (Human)</i></b>					
	ECHA_HUMAN	K.TGIEQGS DAGYLC#ESQK.F	2	3.8824	P40939
	ECHA_HUMAN	K.DLNSDMDSILASLK.L	2	4.3652	P40939
	ECHA_HUMAN	R.DSIFSNLTGQLDYQGFEK.A	2	3.529	P40939
	ECHA_HUMAN	R.FGGNPPELLTQMVSK.G	2	3.2824	P40939
	ECHA_HUMAN	K.TLQEV TQLSQAQR.I	2	3.4555	P40939
	ECHA_HUMAN	K.TLQEV TQLSQAQR.I	2	3.8649	P40939
	ECHA_HUMAN	K.TGIEQGS DAGYLC#ESQK.F	2	4.02	P40939
	ECHA_HUMAN	K.TGIEQGS DAGYLC#ESQK.F	2	3.0244	P40939
	ECHA_HUMAN	K.TGIEQGS DAGYLC#ESQK.F	2	3.8108	P40939
	ECHA_HUMAN	K.DLNSDMDSILASLK.L	2	4.1329	P40939
<b><i>Tubulin alpha-1A chain - Homo sapiens (Human)</i></b>					
	TBA1A_HUMAN	R.FDGALNVDL TEFQTNLVPYPR.I	2	4.0107	Q71U36
	TBA1A_HUMAN	R.FDGALNVDL TEFQTNLVPYPR.I	2	4.0491	Q71U36
	TBA1A_HUMAN	K.EIIDLV LDR.I	2	2.8041	Q71U36
	TBA1A_HUMAN	K.TIGGG DDFSNTFFSETGAGK.H	2	3.7213	Q71U36
	TBA1A_HUMAN	K.VGINYQPPTV VPGDLAK.V	2	2.6558	Q71U36
	TBA1A_HUMAN	R.NLDIERPT YTNLNR.L	2	2.7711	Q71U36
	TBA1A_HUMAN	K.DVNAAIATIK.T	2	2.6526	Q71U36
	TBA1A_HUMAN	R.FDGALNVDL TEFQTNLVPYPR.I	2	3.5541	Q71U36
	TBA1A_HUMAN	R.NLDIERPT YTNLNR.L	2	2.8306	Q71U36
	TBA1A_HUMAN	K.TIGGG DDFSNTFFSETGAGK.H	2	4.2181	Q71U36
	TBA1A_HUMAN	K.TIGGG DDFSNTFFSETGAGK.H	2	4.2982	Q71U36
<b><i>Tubulin beta-1 chain - Homo sapiens (Human)</i></b>					
	TBB1_HUMAN	R.FPGQLNADLR.K	2	2.3437	Q9H4B7
	TBB1_HUMAN	R.FPGQLNADLR.K	2	2.4966	Q9H4B7

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
TBB1_HUMAN		R.FPGQLNADLR.K	2	2.6354	Q9H4B7
TBB1_HUMAN		K.EVDQQLLSVQTR.N	2	2.4552	Q9H4B7
TBB1_HUMAN		R.FPGQLNADLR.K	2	2.4121	Q9H4B7
<b><i>Tubulin beta-2A chain - Homo sapiens (Human)</i></b>					
TBB2A_HUMAN		K.EVDEQM*LNQNK.N	2	3.5741	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.6314	Q13885
TBB2A_HUMAN		K.EVDEQM*LNQNK.N	2	3.5123	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	4.2298	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.3041	Q13885
TBB2A_HUMAN		K.GHYTEGAELVDSVLDVVR.K	2	2.8987	Q13885
TBB2A_HUMAN		K.GHYTEGAELVDSVLDVVR.K	3	4.1245	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	4.1738	Q13885
TBB2A_HUMAN		K.EVDEQM*LNQNK.N	2	3.6404	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.3396	Q13885
TBB2A_HUMAN		K.EVDEQM*LNQNK.N	2	3.4787	Q13885
<b><i>Tudor domain-containing protein 3 - Homo sapiens (Human)</i></b>					
TDRD3_HUMAN		K.TLQVTM*PVKPTNDNDEFEK.Q	2	2.4248	Q9H7E2
<b><i>Tumor necrosis factor, alpha-induced protein 2 - Homo sapiens (Human)</i></b>					
TNAP2_HUMAN		R.ELAAAAAAGGVSEELVR.R	2	2.7882	Q03169
TNAP2_HUMAN		R.ELAAAAAAGGVSEELVR.R	2	2.6805	Q03169
<b><i>Tyrosine-protein phosphatase non-receptor type 12 - Homo sapiens (Human)</i></b>					
PTN12_HUMAN		K.IPEEFNVFNLIQEMR.T	2	3.7411	Q05209
<b><i>Tyrosine-protein phosphatase non-receptor type 6 - Homo sapiens (Human)</i></b>					
PTN6_HUMAN		R.ESLSQPGDFVLSVSDQPK.A	2	2.7832	P29350
PTN6_HUMAN		K.GLDC#DIDIQK.T	2	2.9848	P29350
PTN6_HUMAN		R.ESLSQPGDFVLSVSDQPK.A	2	2.8129	P29350
<b><i>Tyrosine-protein phosphatase non-receptor type substrate 1 precursor - Homo sapiens (Human)</i></b>					
SHPS1_HUMAN		R.EITQDTNDITYADLNLPK.G	2	3.1384	P78324
<b><i>Ubiquinol-cytochrome-c reductase complex core protein 2, mitochondrial precursor - Homo sa</i></b>					
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	2.4214	P22695
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	3.8427	P22695
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	3.4692	P22695
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	3.8101	P22695
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	3.7754	P22695

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Ubiquitin - Homo sapiens (Human)</i></b>					
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	2.9813	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	4.0933	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	4.05	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	4.6406	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	3.4263	P62988
UBIQ_HUMAN		K.ESTLHLVLR.L	2	2.3546	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	4.0995	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	3.7302	P62988
UBIQ_HUMAN		K.ESTLHLVLR.L	2	2.3397	P62988
<b><i>Unc-112-related protein 2 - Homo sapiens (Human)</i></b>					
URP2_HUMAN		R.ARGEELDEDLFLQLTGGEAF.-	2	2.6782	Q86UX7
URP2_HUMAN		K.GC#EVVPDVNVSGQK.F	2	2.5058	Q86UX7
URP2_HUMAN		R.TM*ADSSYTSEVQAILAFLSLQR.T	2	2.7321	Q86UX7
URP2_HUMAN		K.LEGSAPTDVLDLSTTIPELK.D	2	3.2959	Q86UX7
URP2_HUMAN		K.EKEPEEELYDLSK.V	2	2.3284	Q86UX7
URP2_HUMAN		R.TMADSSYTSEVQAILAFLSLQR.T	2	2.8786	Q86UX7
URP2_HUMAN		R.VFVGEEDPEAESVTLR.V	2	3.4512	Q86UX7
URP2_HUMAN		K.LEGSAPTDVLDLSTTIPELK.D	2	3.4989	Q86UX7
URP2_HUMAN		R.GEELDEDLFLQLTGGEAF.-	2	3.0257	Q86UX7
URP2_HUMAN		K.GC#EVVPDVNVSGQK.F	2	2.4427	Q86UX7
<b><i>Unc-13 homolog D - Homo sapiens (Human)</i></b>					
UN13D_HUMAN		R.VGAVLEQQQLQNTLHAQLQSALAGLGHEIR.T	3	5.0997	Q70J99
UN13D_HUMAN		R.TLAEQLEVGIAK.H	2	2.376	Q70J99
UN13D_HUMAN		R.YLQEAHFHVEPEEHQQTLR.V	3	4.4843	Q70J99
<b><i>UNC93 homolog B1 - Homo sapiens (Human)</i></b>					
UN93B_HUMAN		R.RPC#PYEQAQGGDGPEEQ.-	2	2.9289	Q9H1C4
UN93B_HUMAN		R.RPC#PYEQAQGGDGPEEQ.-	2	3.4607	Q9H1C4
UN93B_HUMAN		R.RPC#PYEQAQGGDGPEEQ.-	2	3.701	Q9H1C4
<b><i>Uncharacterized protein KIAA0152 precursor - Homo sapiens (Human)</i></b>					
K0152_HUMAN		K.KEEEEEEEEYDEGSNLKK.Q	3	5.4968	Q14165
K0152_HUMAN		K.KEEEEEEEEYDEGSNLKK.Q	3	6.9483	Q14165
K0152_HUMAN		R.SNPEDQILYQTER.Y	2	3.0228	Q14165
<b><i>Uncharacterized protein KIAA0562 - Homo sapiens (Human)</i></b>					



<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	K0562_HUMAN	K.PM*PSLPQLEER.G	2	2.3743	O60308
<b><i>Uncharacterized protein KIAA0776 - Homo sapiens (Human)</i></b>					
	K0776_HUMAN	K.HIQDAPEEFISELAELYLIKPLNK.T	3	4.1565	O94874
<b><i>Uncharacterized protein KIAA1754 precursor - Homo sapiens (Human)</i></b>					
	K1754_HUMAN	R.EFLEGFVDDLLEALR.S	2	4.7045	Q8IWB1
<b><i>UPF0318 protein FAM120A - Homo sapiens (Human)</i></b>					
	F120A_HUMAN	K.TPLIDLCDGQADQAAK.V	2	2.447	Q9NZB2
<b><i>UPF0404 protein C11orf59 - Homo sapiens (Human)</i></b>					
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	3.5299	Q6IAA8
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	3.7255	Q6IAA8
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	3.0852	Q6IAA8
<b><i>Uveal autoantigen with coiled-coil domains and ankyrin repeats - Homo sapiens (Human)</i></b>					
	UACA_HUMAN	K.KLEMEKLLLENDLSK.D	2	2.5751	Q9BZF9
	UACA_HUMAN	K.KLEMEKLLLENDLSK.D	2	2.7843	Q9BZF9
<b><i>Vacuolar proton translocating ATPase 116 kDa subunit a isoform 3 - Homo sapiens (Human)</i></b>					
	VPP3_HUMAN	R.LGALQQLQQSQELQEVLGETER.F	2	4.376	Q13488
	VPP3_HUMAN	R.LGALQQLQQSQELQEVLGETER.F	2	3.1487	Q13488
	VPP3_HUMAN	R.LGALQQLQQSQELQEVLGETER.F	2	4.1006	Q13488
<b><i>Vam6/Vps39-like protein - Homo sapiens (Human)</i></b>					
	VPS39_HUMAN	K.TPVPAGEEEGELGEYR.Q	2	3.0168	Q96JC1
<b><i>Vesicle transport protein GOT1B - Homo sapiens (Human)</i></b>					
	GOT1B_HUMAN	R.VPVLGSLLNLPGIR.S	2	3.9683	Q9Y3E0
	GOT1B_HUMAN	R.VPVLGSLLNLPGIR.S	2	3.0228	Q9Y3E0
	GOT1B_HUMAN	R.VPVLGSLLNLPGIR.S	2	3.9912	Q9Y3E0
<b><i>Vesicle transport through interaction with t-SNAREs homolog 1A - Homo sapiens (Human)</i></b>					
	VT11A_HUMAN	R.RLEAGYQIAVETEIQIGQEMLENLSHDR.E	3	5.2389	Q96AJ9
<b><i>Vesicle-associated membrane protein 2 - Homo sapiens (Human)</i></b>					
	VAMP2_HUMAN	R.ADALQAGASQFETSAAK.L	2	4.0284	P63027
<b><i>Vesicle-associated membrane protein 8 - Homo sapiens (Human)</i></b>					
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	4.0925	Q9BV40
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	4.4381	Q9BV40
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	3.905	Q9BV40
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	4.0468	Q9BV40

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<b><i>Vesicle-trafficking protein SEC22b - Homo sapiens (Human)</i></b>					
SC22B_HUMAN		R.NLGSINTELQDVQR.I	2	3.5455	O75396
SC22B_HUMAN		R.NLGSINTELQDVQR.I	2	3.7895	O75396
SC22B_HUMAN		R.NLGSINTELQDVQR.I	2	2.8228	O75396
SC22B_HUMAN		R.NLGSINTELQDVQR.I	2	4.3325	O75396
SC22B_HUMAN		R.VADGLPLAASMQEDEQSGR.D	2	2.9689	O75396
<b><i>Vesicular integral-membrane protein VIP36 precursor - Homo sapiens (Human)</i></b>					
LMAN2_HUMAN		R.DHDTFLAVR.Y	2	2.3498	Q12907
LMAN2_HUMAN		K.DNVDDPTGNFR.S	2	2.8489	Q12907
LMAN2_HUMAN		K.DNVDDPTGNFR.S	2	2.7958	Q12907
LMAN2_HUMAN		R.DHDTFLAVR.Y	2	2.6091	Q12907
LMAN2_HUMAN		K.DNVDDPTGNFR.S	2	2.5609	Q12907
<b><i>Vimentin - Homo sapiens (Human)</i></b>					
VIME_HUMAN		R.EMEENFAVEAANYQDTIGR.L	2	4.3925	P08670
VIME_HUMAN		R.EMEENFAVEAANYQDTIGR.L	2	4.8574	P08670
VIME_HUMAN		K.KLHEEEIQELQAQIQEQHVQIDVDVSKPDLTAALR.D	4	5.1316	P08670
VIME_HUMAN		R.QVQSLTC#EVDALK.G	2	2.5869	P08670
VIME_HUMAN		R.EEAENTLQSFR.Q	2	2.3228	P08670
VIME_HUMAN		R.LLQDSVDFSLADAINTEFK.N	2	2.815	P08670
VIME_HUMAN		K.VESLQEEIAFLK.K	2	3.1249	P08670
VIME_HUMAN		K.VESLQEEIAFLK.K	2	3.2063	P08670
VIME_HUMAN		K.LHEEEIQELQAQIQEQHVQIDVDVSKPDLTAALR.D	4	4.8197	P08670
VIME_HUMAN		R.QVQSLTC#EVDALK.G	2	2.3414	P08670
VIME_HUMAN		K.VESLQEEIAFLK.K	2	2.5294	P08670
VIME_HUMAN		R.EEAENTLQSFR.Q	2	2.676	P08670
VIME_HUMAN		R.QVQSLTC#EVDALK.G	2	2.4574	P08670
VIME_HUMAN		K.VESLQEEIAFLK.K	2	2.8092	P08670
VIME_HUMAN		R.LLQDSVDFSLADAINTEFK.N	2	3.6889	P08670
VIME_HUMAN		R.EEAENTLQSFR.Q	2	2.5909	P08670
VIME_HUMAN		K.VESLQEEIAFLK.K	2	3.4336	P08670
VIME_HUMAN		R.EEAENTLQSFR.Q	2	2.4403	P08670
<b><i>Voltage-dependent anion channel 2 - Homo sapiens (Human)</i></b>					
Q5JSD1_HUMA		R.PMC#IPPSYADLGK.A	2	2.4815	Q5JSD1
Q5JSD1_HUMA		R.PMC#IPPSYADLGK.A	2	2.9495	Q5JSD1
<b><i>Voltage-dependent anion-selective channel protein 1 - Homo sapiens (Human)</i></b>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
VDAC1_HUMAN		K.TDEFQLHTNVNDGTEFGGSIYQK.V	3	5.4091	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	2.5859	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	2.5343	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	3.6433	P21796
VDAC1_HUMAN		K.VNNSLIGLGYTQTLKPGIK.L	2	3.468	P21796
VDAC1_HUMAN		K.TDEFQLHTNVNDGTEFGGSIYQK.V	2	4.2172	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	2.5277	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	3.4201	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	3.8125	P21796
VDAC1_HUMAN		K.VNNSLIGLGYTQTLKPGIK.L	2	3.1616	P21796
VDAC1_HUMAN		K.TDEFQLHTNVNDGTEFGGSIYQK.V	2	4.6335	P21796
VDAC1_HUMAN		K.TDEFQLHTNVNDGTEFGGSIYQK.V	2	4.8513	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	3.4499	P21796
VDAC1_HUMAN		K.TDEFQLHTNVNDGTEFGGSIYQK.V	2	4.54	P21796
VDAC1_HUMAN		K.VNNSLIGLGYTQTLKPGIK.L	2	2.7579	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	4.3453	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	3.4818	P21796

***Voltage-dependent anion-selective channel protein 2 - Homo sapiens (Human)***

VDAC2_HUMAN		K.SC#SGVEFSTSGSSNTDTGK.V	2	4.929	P45880
VDAC2_HUMAN		K.WNTDNTLGTEIAIEDQIC#QGLK.L	2	3.0062	P45880
VDAC2_HUMAN		K.VNNSLIGVGYTQTLRPGVK.L	2	3.0908	P45880
VDAC2_HUMAN		R.TGDFQLHTNVNDGTEFGGSIYQK.V	2	5.2069	P45880
VDAC2_HUMAN		K.SC#SGVEFSTSGSSNTDTGK.V	2	4.08	P45880
VDAC2_HUMAN		K.WNTDNTLGTEIAIEDQIC#QGLK.L	2	2.9082	P45880
VDAC2_HUMAN		R.TGDFQLHTNVNDGTEFGGSIYQK.V	2	5.0213	P45880
VDAC2_HUMAN		K.SC#SGVEFSTSGSSNTDTGK.V	2	4.8065	P45880
VDAC2_HUMAN		K.WNTDNTLGTEIAIEDQIC#QGLK.L	2	3.7846	P45880
VDAC2_HUMAN		K.WNTDNTLGTEIAIEDQIC#QGLK.L	2	4.0646	P45880
VDAC2_HUMAN		R.TGDFQLHTNVNDGTEFGGSIYQK.V	3	5.01	P45880
VDAC2_HUMAN		K.VNNSLIGVGYTQTLRPGVK.L	3	4.3737	P45880
VDAC2_HUMAN		K.SC#SGVEFSTSGSSNTDTGK.V	2	4.1038	P45880

***Voltage-dependent anion-selective channel protein 3 - Homo sapiens (Human)***

VDAC3_HUMAN		K.AADFQLHTHVNDGTEFGGSIYQK.V	3	6.1498	Q9Y277
VDAC3_HUMAN		K.WNTDNTLGTEISWENK.L	2	2.5899	Q9Y277
VDAC3_HUMAN		K.AADFQLHTHVNDGTEFGGSIYQK.V	3	5.3321	Q9Y277

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
VDAC3_HUMAN		K.WNTDNTLGTEISWENK.L	2	3.4735	Q9Y277
VDAC3_HUMAN		K.AADFQLHTHVNDGTEFGGSIYQK.V	3	6.1257	Q9Y277
VDAC3_HUMAN		K.AADFQLHTHVNDGTEFGGSIYQK.V	3	5.8775	Q9Y277
VDAC3_HUMAN		K.AADFQLHTHVNDGTEFGGSIYQK.V	3	4.0222	Q9Y277
<b><i>Zinc finger CCHC domain-containing protein 11 - Homo sapiens (Human)</i></b>					
ZCH11_HUMAN		R.M*DDFQLKGIVEEKFK.V	2	2.5007	Q5TAX3
<b><i>Zinc finger protein 12 - Homo sapiens (Human)</i></b>					
ZNF12_HUMAN		K.AFQKDTVFNHM*EEK.P	2	2.3129	P17014
<b><i>Zinc finger protein 44 - Homo sapiens (Human)</i></b>					
ZNF44_HUMAN		R.IHEGHTLEK.P	2	2.3213	P15621
<b><i>Zinc finger protein 667 - Homo sapiens (Human)</i></b>					
ZN667_HUMAN		K.AFNQM*SSLLHKK.I	2	2.3824	Q5HYK9
<b><i>Zinc-binding alcohol dehydrogenase domain-containing protein 1 - Homo sapiens (Human)</i></b>					
ZADH1_HUMAN		K.FEPGILQLSQWFK.E	2	2.3727	Q8N8N7