

Supplementary Table 2A

A list of 1,403 peptides identified using methanol/PPS combination, 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>
<i>40S ribosomal protein S18 - Homo sapiens (Human)</i>					
	RS18_HUMAN	R.VITIMQNPR.Q	2	2.3746	P62269
<i>40S ribosomal protein S2 - Homo sapiens (Human)</i>					
	RS2_HUMAN	K.AFVAIGDYNGHVGLGVK.C	2	2.8167	P15880
	RS2_HUMAN	K.AFVAIGDYNGHVGLGVK.C	2	3.7086	P15880
<i>40S ribosomal protein S24 - Homo sapiens (Human)</i>					
	RS24_HUMAN	K.TTGFGMIYDSL DYAK.K	2	2.9052	P62847
	RS24_HUMAN	K.TTPDVIFVFGFR.T	2	3.0943	P62847
	RS24_HUMAN	K.TTGFGMIYDSL DYAK.K	2	2.796	P62847
	RS24_HUMAN	K.TTPDVIFVFGFR.T	2	2.3489	P62847
	RS24_HUMAN	K.TTGFGMIYDSL DYAK.K	2	2.6718	P62847
	RS24_HUMAN	K.TTPDVIFVFGFR.T	2	2.599	P62847
	RS24_HUMAN	K.TTGFGMIYDSL DYAK.K	2	2.9064	P62847
<i>40S ribosomal protein S27-like protein - Homo sapiens (Human)</i>					
	RS27L_HUMAN	R.LVQSPNSYFMDVK.C	2	2.4384	Q71UM5
<i>40S ribosomal protein S3 - Homo sapiens (Human)</i>					
	RS3_HUMAN	K.GGKPEPPAMPQPVPTA.-	2	3.2637	P23396
	RS3_HUMAN	K.GGKPEPPAMPQPVPTA.-	2	3.1759	P23396
	RS3_HUMAN	K.GGKPEPPAMPQPVPTA.-	2	2.9073	P23396
	RS3_HUMAN	K.GGKPEPPAMPQPVPTA.-	2	2.5791	P23396
<i>40S ribosomal protein S4, X isoform - Homo sapiens (Human)</i>					
	RS4X_HUMAN	R.TDITYPAGFMDVISIDK.T	2	2.7898	P62701
	RS4X_HUMAN	R.HPGSFDVVHVK.D	3	4.2251	P62701
<i>40S ribosomal protein S5 - Homo sapiens (Human)</i>					
	RS5_HUMAN	K.TIAEC#LADELINAAK.G	3	4.0208	P46782
<i>40S ribosomal protein S7 - Homo sapiens (Human)</i>					
	RS7_HUMAN	K.AIIIFVPVQLK.S	2	2.4173	P62081
<i>60S acidic ribosomal protein P0 - Homo sapiens (Human)</i>					
	RLA0_HUMAN	R.GHLENNPALEK.L	2	2.6263	P05388
	RLA0_HUMAN	R.GHLENNPALEK.L	2	3.1326	P05388

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	RLA0_HUMAN	K.IIQLDDYPK.C	2	2.621	P05388
	RLA0_HUMAN	R.GHLENNPALEK.L	2	2.5929	P05388
<i>60S ribosomal protein L10a - Homo sapiens (Human)</i>					
	RL10A_HUMAN	K.KYDAFLASESLIK.Q	2	2.6385	P62906
<i>60S ribosomal protein L10-like - Homo sapiens (Human)</i>					
	RL10L_HUMAN	R.VHIGQVIMSIR.T	2	2.4459	Q96L21
<i>60S ribosomal protein L11 - Homo sapiens (Human)</i>					
	RL11_HUMAN	K.VLEQLTGQTPVFSK.A	2	3.6064	P62913
<i>60S ribosomal protein L13 - Homo sapiens (Human)</i>					
	RL13_HUMAN	K.STESLQANVQR.L	2	2.652	P26373
	RL13_HUMAN	K.STESLQANVQR.L	2	2.4198	P26373
<i>60S ribosomal protein L14 - Homo sapiens (Human)</i>					
	RL14_HUMAN	R.ALVDGPC#TQVR.R	2	2.581	P50914
	RL14_HUMAN	K.C#MQLTDFILK.F	2	2.4894	P50914
	RL14_HUMAN	R.ALVDGPC#TQVR.R	2	2.636	P50914
	RL14_HUMAN	R.ALVDGPC#TQVR.R	2	3.0109	P50914
	RL14_HUMAN	R.ALVDGPC#TQVR.R	2	2.798	P50914
	RL14_HUMAN	K.C#MQLTDFILK.F	2	2.6589	P50914
	RL14_HUMAN	R.ALVDGPC#TQVR.R	2	2.4865	P50914
	RL14_HUMAN	K.LVAIVDVIDQNR.A	2	3.4674	P50914
<i>60S ribosomal protein L18 - Homo sapiens (Human)</i>					
	RL18_HUMAN	K.ILTFDQLALDSPK.G	2	3.9814	Q07020
	RL18_HUMAN	K.ILTFDQLALDSPK.G	2	3.5481	Q07020
	RL18_HUMAN	K.GC#GTVLLSGPR.K	2	2.4692	Q07020
	RL18_HUMAN	K.ILTFDQLALDSPK.G	2	3.367	Q07020
	RL18_HUMAN	K.ILTFDQLALDSPK.G	2	2.8506	Q07020
<i>60S ribosomal protein L18a - Homo sapiens (Human)</i>					
	RL18A_HUMAN	R.DLTTAGAVTQC#YR.D	2	2.824	Q02543
	RL18A_HUMAN	K.SSGEIVYC#GQVFEK.S	2	3.2284	Q02543
	RL18A_HUMAN	R.DLTTAGAVTQC#YR.D	2	2.7283	Q02543
	RL18A_HUMAN	K.SSGEIVYC#GQVFEK.S	2	2.9374	Q02543
<i>60S ribosomal protein L23 - Homo sapiens (Human)</i>					
	RL23_HUMAN	R.ISLGLPVGAVINC#ADNTGAK.N	2	2.4817	P62829
<i>60S ribosomal protein L24 - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	RL24_HUMAN	K.VFQFLNAKC#ESAFLSKR.N	2	2.4107	P83731
	RL24_HUMAN	K.VFQFLNAKC#ESAFLSKR.N	2	2.3442	P83731
<i>60S ribosomal protein L6 - Homo sapiens (Human)</i>					
	RL6_HUMAN	K.HLTDAYFK.K	2	2.4537	Q02878
	RL6_HUMAN	R.ASITPGTILIILTGR.H	2	3.2609	Q02878
	RL6_HUMAN	R.ASITPGTILIILTGR.H	2	2.7782	Q02878
	RL6_HUMAN	R.ASITPGTILIILTGR.H	2	3.0492	Q02878
<i>60S ribosomal protein L7 - Homo sapiens (Human)</i>					
	RL7_HUMAN	R.IALTDNALIAR.S	2	2.9696	P18124
	RL7_HUMAN	R.IVEPYIAWGYPNLK.S	2	2.9473	P18124
	RL7_HUMAN	R.IVEPYIAWGYPNLK.S	2	2.8266	P18124
	RL7_HUMAN	R.IVEPYIAWGYPNLK.S	2	2.7256	P18124
	RL7_HUMAN	R.IALTDNALIAR.S	2	2.6854	P18124
	RL7_HUMAN	R.IVEPYIAWGYPNLK.S	2	2.5953	P18124
<i>60S ribosomal protein L7a - Homo sapiens (Human)</i>					
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	2.4564	P62424
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	2.9275	P62424
	RL7A_HUMAN	K.VPPAINQFTQALDR.Q	2	3.113	P62424
	RL7A_HUMAN	R.KTCTTVAFQVNSQEDK.G	2	2.3604	P62424
<i>60S ribosomal protein L9 - Homo sapiens (Human)</i>					
	RL9_HUMAN	R.TIC#SHVQNMIIK.G	2	2.6089	P32969
	RL9_HUMAN	R.TIC#SHVQNMIIK.G	2	2.8809	P32969
<i>Actin, aortic smooth muscle - Homo sapiens (Human)</i>					
	ACTA_HUMAN	K.DSYVGDEAQS.K	2	2.6591	P62736
	ACTA_HUMAN	K.AGFAGDDAPR.A	2	2.7112	P62736
	ACTA_HUMAN	R.HQGVVMVGMGQK.D	2	2.4777	P62736
	ACTA_HUMAN	R.LDLAGRDLTDYLMK.I	2	2.4095	P62736
	ACTA_HUMAN	R.LDLAGRDLTDYLMKILTER.G	3	3.7781	P62736
	ACTA_HUMAN	R.LDLAGRDLTDYLMKILTER.G	3	4.287	P62736
	ACTA_HUMAN	K.DSYVGDEAQS.K	2	2.4821	P62736
	ACTA_HUMAN	K.AGFAGDDAPR.A	2	2.5252	P62736
	ACTA_HUMAN	R.LDLAGRDLTDYLMK.I	2	3.3284	P62736
	ACTA_HUMAN	R.LDLAGRDLTDYLMKILTER.G	3	4.4045	P62736
	ACTA_HUMAN	K.DSYVGDEAQS.K	2	2.523	P62736
	ACTA_HUMAN	K.AGFAGDDAPR.A	2	2.4662	P62736

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ACTA_HUMAN		R.HQGVMVGMGQK.D	2	2.8007	P62736
ACTA_HUMAN		K.EITALAPSTM*K.I	2	2.4897	P62736
ACTA_HUMAN		R.LDLAGRDLTDYLM*K.I	2	2.4253	P62736
ACTA_HUMAN		R.LDLAGRDLTDYLMK.I	2	2.5905	P62736
ACTA_HUMAN		K.DSYVGDEAQSQR.G	2	2.3643	P62736
ACTA_HUMAN		K.AGFAGDDAPR.A	2	2.6045	P62736
ACTA_HUMAN		K.EITALAPSTM*K.I	2	2.4469	P62736
ACTA_HUMAN		K.EITALAPSTMK.I	2	2.428	P62736
ACTA_HUMAN		R.LDLAGRDLTDYLMK.I	2	2.6957	P62736
<i>Actin, cytoplasmic 1 - Homo sapiens (Human)</i>					
ACTB_HUMAN		R.GYSFTTTAER.E	2	2.6212	P60709
ACTB_HUMAN		R.GYSFTTTAER.E	2	2.3589	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	3.04	P60709
ACTB_HUMAN		R.GYSFTTTAER.E	2	2.4565	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	3.5705	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	4.3093	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	3.5558	P60709
ACTB_HUMAN		R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	3	5.522	P60709
ACTB_HUMAN		R.C#PEALFQPSFLGMESC#GIHETTFNSIMK.C	3	4.2996	P60709
ACTB_HUMAN		K.YPIEHGIVTNWDDM*EKIWHHTFYNELR.V	4	4.5903	P60709
ACTB_HUMAN		R.GYSFTTTAER.E	2	2.9371	P60709
ACTB_HUMAN		K.DLYANTVLSGGTTMYPGIADR.M	2	4.429	P60709
<i>Adipocyte plasma membrane-associated protein - Homo sapiens (Human)</i>					
APMAP_HUMAN		R.RPLRPQVVTDGQQAPEAK.D	3	4.4374	Q9HDC9
<i>ADP/ATP translocase 1 - Homo sapiens (Human)</i>					
ADT1_HUMAN		R.YFPTQALNFAFK.D	2	2.804	P12235
ADT1_HUMAN		K.LLLQVQHASK.Q	2	2.4117	P12235
ADT1_HUMAN		K.GADIMYGTVDC#WR.K	2	2.391	P12235
ADT1_HUMAN		R.YFPTQALNFAFK.D	2	2.6301	P12235
ADT1_HUMAN		K.LLLQVQHASK.Q	2	2.6144	P12235
ADT1_HUMAN		R.YFPTQALNFAFK.D	2	2.3005	P12235
ADT1_HUMAN		R.YFPTQALNFAFK.D	2	3.7556	P12235
ADT1_HUMAN		R.YFPTQALNFAFK.D	2	3.1878	P12235
ADT1_HUMAN		K.GADIMYGTVDC#WR.K	2	2.4745	P12235
ADT1_HUMAN		R.YFPTQALNFAFKDK.Y	2	2.4669	P12235

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>ADP/ATP translocase 2 - Homo sapiens (Human)</i>					
ADT2_HUMAN		R.GMGGAFVLVLYDEIKKYT.-	2	2.8304	P05141
ADT2_HUMAN		K.DFLAGGVAAAISK.T	2	3.6781	P05141
ADT2_HUMAN		K.DFLAGGVAAAISK.T	2	2.9874	P05141
ADT2_HUMAN		K.DFLAGGVAAAISK.T	2	3.2	P05141
ADT2_HUMAN		R.GMGGAFVLVLYDEIKKYT.-	2	3.2393	P05141
ADT2_HUMAN		K.DFLAGGVAAAISK.T	2	4.5688	P05141
ADT2_HUMAN		K.DFLAGGVAAAISK.T	1	2.178	P05141
ADT2_HUMAN		R.GMGGAFVLVLYDEIKKYT.-	2	3.0375	P05141
<i>ADP/ATP translocase 3 - Homo sapiens (Human)</i>					
ADT3_HUMAN		K.DFLAGGIAAAISK.T	2	3.8179	P12236
ADT3_HUMAN		K.DFLAGGIAAAISK.T	2	3.7338	P12236
ADT3_HUMAN		K.DFLAGGIAAAISK.T	2	3.5625	P12236
ADT3_HUMAN		K.DFLAGGIAAAISK.T	2	3.2756	P12236
ADT3_HUMAN		K.DFLAGGIAAAISK.T	2	2.7364	P12236
ADT3_HUMAN		K.DFLAGGIAAAISK.T	1	2.5381	P12236
<i>ADP-ribosyl cyclase 2 precursor - Homo sapiens (Human)</i>					
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	3.1148	Q10588
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	2.8019	Q10588
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	2.6575	Q10588
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	2.8897	Q10588
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	2.5833	Q10588
BST1_HUMAN		K.GFFADYEIPNLQK.E	2	2.8181	Q10588
<i>Aldehyde dehydrogenase 3B1 - Homo sapiens (Human)</i>					
AL3B1_HUMAN		K.NLATQLDSAFIR.K	2	2.4486	P43353
AL3B1_HUMAN		R.FYGDDPQSSPNLGR.I	2	3.4846	P43353
<i>Annexin A1 - Homo sapiens (Human)</i>					
ANXA1_HUMAN		K.ALTGHLEEVVLLALK.T	2	3.1095	P04083
<i>Annexin A2 - Homo sapiens (Human)</i>					
ANXA2_HUMAN		R.TNQELQEINR.V	2	2.9352	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.1626	P07355
ANXA2_HUMAN		K.SYSPYDMLESIR.K	2	2.5416	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	2.9794	P07355
ANXA2_HUMAN		K.SALSGHLETVILGLLK.T	3	3.9586	P07355
ANXA2_HUMAN		R.TNQELQEINR.V	2	3.0887	P07355

<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	3.25	P07355
ANXA2_HUMAN		K.SLYYYIQDQTK.G	2	3.4341	P07355
ANXA2_HUMAN		K.AYTNFDAERDALNIETAIK.T	2	2.3249	P07355
ANXA2_HUMAN		K.SYSPYDMLESIRK.E	2	2.6198	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	2.577	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	3.3456	P07355
ANXA2_HUMAN		K.SALSGHLETIVILGLLK.T	2	4.3126	P07355
ANXA2_HUMAN		R.DLYDAGVKR.K	2	2.4015	P07355
ANXA2_HUMAN		R.TNQELQEINR.V	2	3.1814	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.8152	P07355
ANXA2_HUMAN		K.LSLEGDHSTPPSAYGSVK.A	2	3.1983	P07355
ANXA2_HUMAN		R.AEDGSVIDYELIDQDAR.D	2	3.4092	P07355
ANXA2_HUMAN		R.AEDGSVIDYELIDQDAR.D	2	2.4255	P07355
ANXA2_HUMAN		K.SYSPYDMLESIR.K	2	2.3697	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	2.7963	P07355
ANXA2_HUMAN		K.SALSGHLETIVILGLLK.T	2	2.4054	P07355
ANXA2_HUMAN		R.TNQELQEINR.V	2	3.194	P07355
ANXA2_HUMAN		K.TPAQYDASELK.A	2	3.6265	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	3.3728	P07355
ANXA2_HUMAN		K.GVDEVTIVNILTNR.S	2	3.0007	P07355
ANXA2_HUMAN		K.SALSGHLETIVILGLLK.T	2	4.4282	P07355
ANXA2_HUMAN		K.SLYYYIQDQTKGDYQK.A	2	3.7215	P07355
ANXA2_HUMAN		R.AEDGSVIDYELIDQDAR.D	2	2.8201	P07355
ANXA2_HUMAN		K.GDLENAFLNLVQC#IQNKPLYFADR.L	2	3.3201	P07355

Apoptosis regulator BAX, membrane isoform alpha - Homo sapiens (Human)

BAXA_HUMAN		R.MIAAVDTDSPR.E	2	2.5613	Q07812
BAXA_HUMAN		R.MIAAVDTDSPR.E	2	2.5276	Q07812
BAXA_HUMAN		K.TGALLLQGFQDR.A	2	3.154	Q07812
BAXA_HUMAN		K.TGALLLQGFQDR.A	2	3.441	Q07812
BAXA_HUMAN		K.TGALLLQGFQDR.A	2	3.1732	Q07812
BAXA_HUMAN		R.IGDELDSNMELQR.M	2	3.1251	Q07812

Atlastin-3 - Homo sapiens (Human)

ATLA3_HUMAN		K.SMLQATAEANNLAAAASAK.D	2	3.4138	Q6DD88
ATLA3_HUMAN		R.LAMDEIFQKPFQTLMLVLR.D	3	4.2907	Q6DD88

ATP synthase a chain - Homo sapiens (Human)

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ATP6_HUMAN		R.LITTQQWLIK.L	2	2.6533	P00846
ATP6_HUMAN		R.LITTQQWLIK.L	2	2.5048	P00846
ATP6_HUMAN		R.LITTQQWLIK.L	2	2.6973	P00846
<i>ATP synthase B chain, mitochondrial precursor - Homo sapiens (Human)</i>					
AT5F1_HUMAN		K.HVVQSISTQQEK.E	2	2.686	P24539
AT5F1_HUMAN		R.YGLIPEEFFQFLYPK.T	2	2.6633	P24539
AT5F1_HUMAN		K.SQQALVQK.R	2	2.3885	P24539
AT5F1_HUMAN		R.YGLIPEEFFQFLYPK.T	2	2.5564	P24539
AT5F1_HUMAN		R.YGLIPEEFFQFLYPK.T	2	3.1315	P24539
AT5F1_HUMAN		K.SQQALVQK.R	2	2.3146	P24539
AT5F1_HUMAN		K.HVVQSISTQQEK.E	2	2.8591	P24539
AT5F1_HUMAN		K.QASIQHIQNAIDTEK.S	2	2.6913	P24539
<i>ATP synthase D chain, mitochondrial - Homo sapiens (Human)</i>					
ATP5H_HUMAN		K.NLIPFDQMTIEDLNEAFPETK.L	2	2.624	O75947
ATP5H_HUMAN		K.YPYWPHQPIENL.-	2	3.1325	O75947
<i>ATP synthase e chain, mitochondrial - Homo sapiens (Human)</i>					
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	3.5039	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	2.6429	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	2.9455	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	3.1184	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	2.8861	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	3.3263	P56385
ATP5I_HUMAN		R.YSALFLGVAYGATR.Y	2	3.1449	P56385
ATP5I_HUMAN		R.ELAEDDSILK.-	1	2.2819	P56385
<i>ATP synthase f chain, mitochondrial - Homo sapiens (Human)</i>					
ATPK_HUMAN		K.LGELPSWILM*R.D	2	2.3061	P56134
ATPK_HUMAN		K.LGELPSWILM*R.D	2	2.8183	P56134
ATPK_HUMAN		R.DFSPSGIFGAFQR.G	2	2.4436	P56134
ATPK_HUMAN		K.LGELPSWILMR.D	2	3.2747	P56134
ATPK_HUMAN		K.LGELPSWILMR.D	2	2.7755	P56134
ATPK_HUMAN		R.DFSPSGIFGAFQR.G	2	2.5802	P56134
ATPK_HUMAN		K.LGELPSWILM*R.D	2	2.3272	P56134
ATPK_HUMAN		R.DFSPSGIFGAFQR.G	2	2.4921	P56134
<i>ATP synthase gamma chain, mitochondrial precursor - Homo sapiens (Human)</i>					
ATPG_HUMAN		K.HLLIGVSSDR.G	2	2.3432	P36542

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	ATPG_HUMAN	K.HLLIGVSSDR.G	2	2.5936	P36542
	ATPG_HUMAN	K.HLLIGVSSDR.G	2	2.468	P36542
<i>ATP synthase subunit alpha, mitochondrial precursor - Homo sapiens (Human)</i>					
	ATPA_HUMAN	K.HALIIYDDLK.Q	2	2.511	P25705
	ATPA_HUMAN	K.HALIIYDDLK.Q	2	3.0634	P25705
<i>ATP synthase subunit beta, mitochondrial precursor - Homo sapiens (Human)</i>					
	ATPB_HUMAN	K.TVLIMELINNVAK.A	2	2.7392	P06576
	ATPB_HUMAN	R.LVLEVAQHLGESTVR.T	2	2.541	P06576
<i>ATP synthase subunit g, mitochondrial - Homo sapiens (Human)</i>					
	ATP5L_HUMAN	K.IVNSAQTGSFK.Q	2	2.5007	O75964
<i>ATWD578 - Homo sapiens (Human)</i>					
	Q6UW11_HUMAN	R.AAAPDVAPAPGPAPR.A	2	2.7871	Q6UW11
	Q6UW11_HUMAN	R.AAAPDVAPAPGPAPR.A	2	2.4093	Q6UW11
	Q6UW11_HUMAN	R.AAAPDVAPAPGPAPR.A	2	2.8106	Q6UW11
	Q6UW11_HUMAN	R.IINGIIISVK.T	2	2.536	Q6UW11
	Q6UW11_HUMAN	R.AAAPDVAPAPGPAPR.A	2	3.1134	Q6UW11
	Q6UW11_HUMAN	K.QMNVQLAAK.I	2	2.4142	Q6UW11
<i>B-cell receptor-associated protein 31 - Homo sapiens (Human)</i>					
	BAP31_HUMAN	K.AENQVLAMR.K	2	2.6534	P51572
	BAP31_HUMAN	K.AENQVLAMR.K	2	2.4014	P51572
	BAP31_HUMAN	K.YMEENDQLKK.G	2	2.3373	P51572
	BAP31_HUMAN	K.AENQVLAM*R.K	2	2.4955	P51572
	BAP31_HUMAN	K.AENQVLAMR.K	2	2.4808	P51572
	BAP31_HUMAN	K.LDVGNAEVKLEENR.S	2	3.2891	P51572
<i>BH3-interacting domain death agonist - Homo sapiens (Human)</i>					
	BID_HUMAN	R.HLAQVGDSDMR.S	2	2.7367	P55957
	BID_HUMAN	R.HLAQVGDSDMR.S	2	2.3472	P55957
	BID_HUMAN	R.DVFHTTVNFINQNL.R.T	2	2.3792	P55957
	BID_HUMAN	R.HLAQVGDSDMR.S	2	2.5405	P55957
	BID_HUMAN	R.DVFHTTVNFINQNL.R.T	2	2.8099	P55957
<i>Brain acid soluble protein 1 - Homo sapiens (Human)</i>					
	BASP_HUMAN	K.AQGPAASAEEPKEPAANSQDQTVTVKE.-	3	4.4936	P80723
<i>BTB/POZ domain-containing protein KCTD12 - Homo sapiens (Human)</i>					
	KCD12_HUMAN	R.MFTQQQPQELAR.D	2	3.1906	Q96CX2

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	KCD12_HUMAN	R.MFTQQQPQELAR.D	2	3.4497	Q96CX2
<i>C1orf121 protein - Homo sapiens (Human)</i>					
	Q8WUE8_HUMAN	R.C#FRPGC#RGPGCTLSAR.L	2	2.4735	Q8WUE8
<i>Calnexin precursor - Homo sapiens (Human)</i>					
	CALX_HUMAN	K.DKGDEEEEGEEKLEEK.Q	3	4.2063	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.7832	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.4471	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.6478	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	2.913	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.6295	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	2.7463	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	2.9321	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.5692	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.5594	P27824
	CALX_HUMAN	K.AEEDEILNR.S	2	2.8388	P27824
	CALX_HUMAN	R.IVDDWANDGWGLK.K	2	3.3322	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	2.9671	P27824
	CALX_HUMAN	K.APVPTGEVYFADSFDR.G	2	2.8812	P27824
	CALX_HUMAN	R.IVDDWANDGWGLK.K	2	3.5818	P27824
	CALX_HUMAN	R.IVDDWANDGWGLK.K	2	3.0348	P27824
	CALX_HUMAN	R.IVDDWANDGWGLK.K	2	2.3239	P27824
	CALX_HUMAN	K.IPNPDFFEDLEPFR.M	2	3.4322	P27824
	CALX_HUMAN	K.IPDPEAVKPDWDEDAPAK.I	2	3.0715	P27824
	CALX_HUMAN	K.IPDPEAVKPDWDEDAPAK.I	2	3.0033	P27824
<i>Carnitine O-palmitoyltransferase I, liver isoform - Homo sapiens (Human)</i>					
	CPT1A_HUMAN	R.ILDNTSEPQPGEAR.L	2	3.4956	P50416
	CPT1A_HUMAN	R.ILDNTSEPQPGEAR.L	2	3.9612	P50416
<i>Cathepsin G precursor - Homo sapiens (Human)</i>					
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	3.2873	P08311
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	4.8768	P08311
	CATG_HUMAN	R.TIQNDIM*LLQLSR.R	2	2.6596	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	2.93	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	3.175	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	4.3398	P08311
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	4.2502	P08311

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	CATG_HUMAN	R.TIQNDIM*LLQLSR.R	2	2.6286	P08311
	CATG_HUMAN	R.TIQNDIM*LLQLSR.R	2	2.5648	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	3.1208	P08311
	CATG_HUMAN	R.PYMAYLQIQSPAGQSR.C	2	3.3312	P08311
	CATG_HUMAN	R.TIQNDIM*LLQLSR.R	2	3.4374	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	3.9129	P08311
	CATG_HUMAN	R.TIQNDIMLLQLSR.R	2	3.2747	P08311
<i>Cation-dependent mannose-6-phosphate receptor precursor - Homo sapiens (Human)</i>					
	MPRD_HUMAN	R.HTLADNFNPVSEER.G	2	3.7658	P20645
	MPRD_HUMAN	R.HTLADNFNPVSEER.G	2	3.1211	P20645
<i>CD44 antigen precursor - Homo sapiens (Human)</i>					
	CD44_HUMAN	K.ALSIGFETC#R.Y	2	2.3237	P16070
<i>CD48 antigen precursor - Homo sapiens (Human)</i>					
	CD48_HUMAN	R.C#YTC#QVSNVSSK.N	2	2.7445	P09326
<i>CD9 antigen - Homo sapiens (Human)</i>					
	CD9_HUMAN	K.EVQEFYKDTYNK.L	2	2.8692	P21926
<i>CD97 antigen precursor - Homo sapiens (Human)</i>					
	CD97_HUMAN	R.HLIATQLLSNLEDIMR.I	2	4.3944	P48960
	CD97_HUMAN	R.HLIATQLLSNLEDIMR.I	2	3.6254	P48960
<i>CDNA FLJ46362 fis, clone TESTI4050293 - Homo sapiens (Human)</i>					
	Q6ZRH1_HUMAN	K.MLEAHIKTFRM*R.M	2	2.3204	Q6ZRH1
<i>CDP-diacylglycerol--inositol 3-phosphatidyltransferase - Homo sapiens (Human)</i>					
	CDIPT_HUMAN	R.FGAMLDMILTDR.C	2	2.3921	O14735
	CDIPT_HUMAN	K.SLISVIHLITAAR.N	2	2.8701	O14735
<i>Cellular tumor antigen p53 - Homo sapiens (Human)</i>					
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	2.9382	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	3.0498	P04637
	P53_HUMAN	R.RTEEENLR.K	2	2.4847	P04637
	P53_HUMAN	K.SVTC#TYSPALNK.M	2	3.083	P04637
	P53_HUMAN	K.TC#PVQLWVDSTPPPGTR.V	2	2.5429	P04637
<i>CKLF-like MARVEL transmembrane domain-containing protein 6 - Homo sapiens (Human)</i>					
	CKLF6_HUMAN	R.SGLAAYFFMGR.L	2	2.776	Q9NX76
	CKLF6_HUMAN	R.SGLAAYFFMGR.L	2	3.2259	Q9NX76
	CKLF6_HUMAN	R.SGLAAYFFMGR.L	2	2.8355	Q9NX76

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Clathrin coat assembly protein AP180 - Homo sapiens (Human)</i>					
	AP180_HUMAN	R.NTLFNLSNFLDK.S	2	2.6558	O60641
<i>Clathrin heavy chain 1 - Homo sapiens (Human)</i>					
	CLH1_HUMAN	R.NNLAGAEELFAR.K	2	2.7536	Q00610
<i>Coatomer subunit alpha - Homo sapiens (Human)</i>					
	COPA_HUMAN	K.LVGQSIIAYLQK.K	2	3.0906	P53621
	COPA_HUMAN	K.LVGQSIIAYLQK.K	2	3.0917	P53621
	COPA_HUMAN	K.LVGQSIIAYLQK.K	2	2.5141	P53621
	COPA_HUMAN	K.LVGQSIIAYLQK.K	2	2.742	P53621
<i>Coatomer subunit beta - Homo sapiens (Human)</i>					
	COPB_HUMAN	K.LPGLLMTIIR.F	2	2.6946	P53618
	COPB_HUMAN	K.LPGLLMTIIR.F	2	3.0215	P53618
<i>Core histone macro-H2A.2 - Homo sapiens (Human)</i>					
	H2AW_HUMAN	R.HILLAVANDEELNQLLK.G	2	2.4584	Q9P0M6
<i>Cytochrome b-245 heavy chain - Homo sapiens (Human)</i>					
	CY24B_HUMAN	R.KLLGSALALAR.A	2	2.5857	P04839
	CY24B_HUMAN	K.LLGSALALAR.A	2	2.8734	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.5062	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	3.138	P04839
	CY24B_HUMAN	R.KLLGSALALAR.A	2	2.5359	P04839
	CY24B_HUMAN	K.LLGSALALAR.A	2	2.8769	P04839
	CY24B_HUMAN	K.LLGSALALAR.A	2	3.0439	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.5818	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.4602	P04839
	CY24B_HUMAN	R.IVGDWTEGLFNAC#GC#DK.Q	2	3.0179	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	2.8402	P04839
	CY24B_HUMAN	R.KLLGSALALAR.A	2	3.0123	P04839
	CY24B_HUMAN	K.LLGSALALAR.A	2	2.9982	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.6795	P04839
	CY24B_HUMAN	R.IVGDWTEGLFNAC#GC#DK.Q	2	2.7196	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	3.2208	P04839
	CY24B_HUMAN	K.LEWHPFTLTSAPPEEDFFSIHIR.I	3	3.7916	P04839
	CY24B_HUMAN	K.LLGSALALAR.A	2	3.2775	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.6385	P04839
	CY24B_HUMAN	K.MEVGQYIFVK.C	2	2.8065	P04839

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	2.3923	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	3.8563	P04839
	CY24B_HUMAN	R.IGVFLC#GPEALAETLSK.Q	2	3.7194	P04839
	CY24B_HUMAN	R.IVGDWTEGLFNAC#GC#DK.Q	2	2.7578	P04839
<i>Cytochrome b-245 light chain - Homo sapiens (Human)</i>					
	CY24A_HUMAN	K.HMTAVVK.L	2	2.6397	P13498
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	3	3.9445	P13498
	CY24A_HUMAN	K.HMTAVVK.L	2	2.4776	P13498
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	3	4.0812	P13498
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	3	4.0268	P13498
	CY24A_HUMAN	R.KKPSEEEEEAAAAGGPPGGPQVNPPIVTDEVV.-	2	4.7791	P13498
<i>Cytochrome c oxidase polypeptide VIc precursor - Homo sapiens (Human)</i>					
	COX6C_HUMAN	K.AGIFQSVK.-	2	2.555	P09669
	COX6C_HUMAN	K.AGIFQSVK.-	2	2.5486	P09669
	COX6C_HUMAN	K.AGIFQSVK.-	2	2.5423	P09669
<i>Cytochrome c oxidase polypeptide VIIa-liver/heart, mitochondrial precursor - Homo sapiens (H</i>					
	CX7A2_HUMAN	K.LFQEDDEIPLYLK.G	2	2.4474	P14406
	CX7A2_HUMAN	K.LFQEDDEIPLYLK.G	2	2.4163	P14406
	CX7A2_HUMAN	K.LFQEDDEIPLYLK.G	2	2.5025	P14406
<i>Cytochrome c oxidase subunit 2 - Homo sapiens (Human)</i>					
	COX2_HUMAN	R.ILYMTDEVNDPSLTIK.S	2	2.6602	P00403
	COX2_HUMAN	R.LLDVDNRVVLPIEAPIR.M	3	3.8103	P00403
	COX2_HUMAN	R.LLDVDNRVVLPIEAPIR.M	2	2.4544	P00403
	COX2_HUMAN	R.MMITSQDVLHSHWAVPTLGLK.T	2	4.073	P00403
	COX2_HUMAN	R.ILYM*TDEVNDPSLTIK.S	2	2.4854	P00403
	COX2_HUMAN	R.ILYM*TDEVNDPSLTIK.S	2	3.558	P00403
	COX2_HUMAN	R.ILYMTDEVNDPSLTIK.S	2	3.9082	P00403
	COX2_HUMAN	R.MMITSQDVLHSHWAVPTLGLK.T	2	2.6969	P00403
<i>Cytochrome c oxidase subunit 4 isoform 1, mitochondrial precursor - Homo sapiens (Human)</i>					
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.7423	P13073
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.4543	P13073
	COX41_HUMAN	R.DHPLPEVAHVK.H	3	3.8231	P13073
	COX41_HUMAN	K.SEDFSLPAYMDR.R	2	2.7968	P13073
<i>Cytochrome c oxidase subunit 5A, mitochondrial precursor - Homo sapiens (Human)</i>					
	COX5A_HUMAN	K.EIYPYVIQELRPTLNELGISTPEELGLDKV.-	3	4.6806	P20674

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Cytochrome c oxidase subunit 5B, mitochondrial precursor - Homo sapiens (Human)</i>					
	COX5B_HUMAN	K.KGLDPYNVLAPK.G	2	2.5215	P10606
<i>Cytoplasmic FMRI-interacting protein 1 - Homo sapiens (Human)</i>					
	CYFP1_HUMAN	K.SLLQGTLQYVK.T	2	2.3823	Q7L576
<i>Cytoskeleton-associated protein 4 - Homo sapiens (Human)</i>					
	CKAP4_HUMAN	K.IETNENNLESAK.G	2	3.116	Q07065
	CKAP4_HUMAN	R.RLEEELR.Q	2	2.3737	Q07065
	CKAP4_HUMAN	K.SSSSSASAAAAAASSSASC#SR.R	2	2.4566	Q07065
	CKAP4_HUMAN	R.STLQTMESDIYEV.R	2	3.0013	Q07065
	CKAP4_HUMAN	R.RLEEELR.Q	2	2.3995	Q07065
	CKAP4_HUMAN	R.STLQTMESDIYEV.R	2	3.1423	Q07065
	CKAP4_HUMAN	R.RLEEELR.Q	2	2.6395	Q07065
	CKAP4_HUMAN	K.IETNENNLESAK.G	2	2.6181	Q07065
	CKAP4_HUMAN	K.VQEQVHTLLSQDQAQAAR.L	3	4.7923	Q07065
	CKAP4_HUMAN	K.VQEQVHTLLSQDQAQAAR.L	3	5.7991	Q07065
	CKAP4_HUMAN	R.DFTSLENTVEER.L	2	3.4853	Q07065
	CKAP4_HUMAN	R.QTESLESLLSK.S	2	2.3543	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	3.8805	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	4.9141	Q07065
	CKAP4_HUMAN	R.ERDFTSLENTVEER.L	2	2.7099	Q07065
	CKAP4_HUMAN	K.SSSSSASAAAAAASSSASC#SR.R	2	3.6758	Q07065
	CKAP4_HUMAN	K.VQEQVHTLLSQDQAQAAR.L	2	3.8214	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	3.1186	Q07065
	CKAP4_HUMAN	R.LEGLGSSEADQDGLASTVR.S	2	4.1936	Q07065
<i>DC2 - Homo sapiens (Human)</i>					
	Q9NRP0_HUMAN	R.VPFLVLEC#PNLK.L	2	3.5117	Q9NRP0
	Q9NRP0_HUMAN	R.VPFLVLEC#PNLK.L	2	3.1145	Q9NRP0
	Q9NRP0_HUMAN	R.VPFLVLEC#PNLK.L	2	3.3595	Q9NRP0
	Q9NRP0_HUMAN	R.VPFLVLEC#PNLK.L	2	2.8493	Q9NRP0
<i>Dehydrogenase/reductase SDR family member 8 precursor - Homo sapiens (Human)</i>					
	DHRS8_HUMAN	K.NPSTSLGPTLEPEEVNR.L	2	3.3771	Q8NBQ5
	DHRS8_HUMAN	K.NPSTSLGPTLEPEEVNR.L	2	3.3846	Q8NBQ5
<i>Derlin-1 - Homo sapiens (Human)</i>					
	DERL1_HUMAN	K.LGLISPAYLFLWPEAFLYR.F	2	3.5409	Q9BUN8
<i>Diacylglycerol O-acyltransferase 1 - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	DGAT1_HUMAN	R.DAAAGPDVGAAGDAPAPAPNK.D	2	3.232	O75907
<i>DNA damage-binding protein 1 - Homo sapiens (Human)</i>					
	DDB1_HUMAN	K.PTAVNGC#VTGHFTSAEDLNLLIAK.N	2	2.5771	Q16531
<i>DNA topoisomerase 3-alpha - Homo sapiens (Human)</i>					
	TOP3A_HUMAN	K.LLTEADLIALM*EK.H	2	2.3945	Q13472
<i>DNA-dependent protein kinase catalytic subunit - Homo sapiens (Human)</i>					
	PRKDC_HUMAN	R.LGASLAFNNIYR.E	2	2.4674	P78527
<i>DnaJ homolog subfamily C member 13 - Homo sapiens (Human)</i>					
	DNJCD_HUMAN	R.LGGYLAEEQATPENPTIR.K	2	2.8369	O75165
<i>Dolichyl-diphosphooligosaccharide-protein glycosyltransferase - Homo sapiens (Human)</i>					
	Q5VWA5_HUMAN	K.LPDVYGVFQFK.V	2	2.9471	Q5VWA5
	Q5VWA5_HUMAN	K.LPDVYGVFQFK.V	2	2.3308	Q5VWA5
	Q5VWA5_HUMAN	K.LPDVYGVFQFK.V	2	2.4627	Q5VWA5
	Q5VWA5_HUMAN	K.LPDVYGVFQFK.V	2	2.3388	Q5VWA5
	Q5VWA5_HUMAN	R.YSQTGNYELAVLSR.W	2	4.0576	Q5VWA5
<i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 67 kDa subunit precursor - Ho</i>					
	RIB1_HUMAN	R.HFDETVNR.Y	2	2.4849	P04843
	RIB1_HUMAN	K.ALTSEIALLQSR.L	2	2.5259	P04843
	RIB1_HUMAN	R.HFDETVNR.Y	2	2.6126	P04843
	RIB1_HUMAN	R.HFDETVNR.Y	2	2.4905	P04843
	RIB1_HUMAN	K.ALTSEIALLQSR.L	2	2.7355	P04843
<i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 - Homo sapiens</i>					
	DAD1_HUMAN	R.FLEEYLSSTPQR.L	2	2.4675	P61803
	DAD1_HUMAN	R.FLEEYLSSTPQR.L	2	2.7359	P61803
	DAD1_HUMAN	K.ADFQGISPER.A	2	2.3418	P61803
<i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit STT3A - Homo sapiens</i>					
	STT3A_HUMAN	R.FYSLLDPSYAK.N	2	2.3611	P46977
	STT3A_HUMAN	R.FYSLLDPSYAK.N	2	3.033	P46977
<i>Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit STT3B - Homo sapiens</i>					
	STT3B_HUMAN	R.NILDDFREAYFWLR.Q	2	2.5045	Q8TCJ2
	STT3B_HUMAN	R.NILDDFREAYFWLR.Q	2	3.0089	Q8TCJ2
<i>Elongation factor 1-alpha 1 - Homo sapiens (Human)</i>					
	EF1A1_HUMAN	K.STTTGHLIYK.C	2	2.4464	P68104
	EF1A1_HUMAN	K.IGGIGTVPVGR.V	2	2.3508	P68104

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	EF1A1_HUMAN	K.IGGIGTVPVGR.V	2	2.53	P68104
	EF1A1_HUMAN	K.STTTGHLIYK.C	2	2.92	P68104
	EF1A1_HUMAN	K.IGGIGTVPVGR.V	2	2.6054	P68104
	EF1A1_HUMAN	R.EHALLAYTLGVK.Q	2	2.3042	P68104
<i>Elongation of very long chain fatty acids protein 1 - Homo sapiens (Human)</i>					
	ELOV1_HUMAN	R.ALQQNGAPGIAK.V	2	2.7113	Q9BW60
	ELOV1_HUMAN	R.ALQQNGAPGIAK.V	2	2.9035	Q9BW60
<i>Endoplasmic precursor - Homo sapiens (Human)</i>					
	ENPL_HUMAN	K.GVVDSDDLPLNVSR.E	2	2.6557	P14625
	ENPL_HUMAN	K.GVVDSDDLPLNVSR.E	2	2.3719	P14625
<i>Eosinophil cationic protein precursor - Homo sapiens (Human)</i>					
	ECP_HUMAN	R.YPVVPHLDTTI.-	2	2.8613	P12724
<i>ER lumen protein retaining receptor 2 - Homo sapiens (Human)</i>					
	ERD22_HUMAN	R.LTGDLSHLAAIVILLK.I	3	3.9103	P33947
	ERD22_HUMAN	R.LTGDLSHLAAIVILLK.I	2	3.913	P33947
	ERD22_HUMAN	R.LTGDLSHLAAIVILLK.I	2	3.8056	P33947
<i>Erlin-1 precursor - Homo sapiens (Human)</i>					
	ERLN1_HUMAN	R.SVQTTLQTDEVK.N	2	2.7331	O75477
	ERLN1_HUMAN	R.SVQTTLQTDEVK.N	2	2.6926	O75477
	ERLN1_HUMAN	K.EALEPSGENVIQNK.E	2	2.8835	O75477
<i>Erlin-2 precursor - Homo sapiens (Human)</i>					
	ERLN2_HUMAN	K.VAQVAEITYGQK.V	2	2.8652	O94905
<i>Erythrocyte band 7 integral membrane protein - Homo sapiens (Human)</i>					
	STOM_HUMAN	R.AMAAEAEASR.E	2	2.6783	P27105
	STOM_HUMAN	K.NSTIVFPLPIDMLQGIGAK.H	2	3.2485	P27105
	STOM_HUMAN	K.NSTIVFPLPIDMLQGIGAK.H	2	3.8223	P27105
	STOM_HUMAN	R.AMAAEAEASR.E	2	2.7011	P27105
	STOM_HUMAN	R.TISFDIPPQEILTK.D	2	2.499	P27105
	STOM_HUMAN	R.TISFDIPPQEILTK.D	2	2.4574	P27105
	STOM_HUMAN	R.AMAAEAEASR.E	2	2.7773	P27105
	STOM_HUMAN	K.EASMVITESPAALQLR.Y	2	3.2488	P27105
	STOM_HUMAN	K.NSTIVFPLPIDMLQGIGAK.H	2	2.6316	P27105
	STOM_HUMAN	R.AMAAEAEASR.E	2	2.6246	P27105
	STOM_HUMAN	K.VIAAEGEM*NASR.A	2	2.8428	P27105
	STOM_HUMAN	K.VIAAEGEMNASR.A	2	3.3748	P27105

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	STOM_HUMAN	R.VQNATLAVANITNADSATR.L	2	2.9789	P27105
	STOM_HUMAN	K.EASMVITESPAALQLR.Y	2	4.3809	P27105
<i>Eukaryotic initiation factor 4A-I - Homo sapiens (Human)</i>					
	IF4A1_HUMAN	K.GYDVIAQAQSGTGK.T	2	3.82	P60842
	IF4A1_HUMAN	K.MFVLDEADEMLSR.G	2	3.9323	P60842
	IF4A1_HUMAN	K.MFVLDEADEMLSR.G	2	3.8905	P60842
	IF4A1_HUMAN	K.MFVLDEADEMLSR.G	2	4.2433	P60842
	IF4A1_HUMAN	K.GYDVIAQAQSGTGK.T	2	3.945	P60842
	IF4A1_HUMAN	K.MFVLDEADEMLSR.G	2	4.1378	P60842
<i>Eukaryotic translation initiation factor 3 subunit 10 - Homo sapiens (Human)</i>					
	IF3A_HUMAN	R.FNVLQYVVPEVK.D	2	2.6427	Q14152
	IF3A_HUMAN	R.FNVLQYVVPEVK.D	2	2.7388	Q14152
<i>Eukaryotic translation initiation factor 3 subunit 6-interacting protein - Homo sapiens (Human)</i>					
	IF36I_HUMAN	K.MLGYFSLVGLLR.L	2	2.6107	Q9Y262
<i>Exportin-1 - Homo sapiens (Human)</i>					
	XPO1_HUMAN	K.YYGLQILENVIK.T	2	2.5155	O14980
<i>Ferritin light chain - Homo sapiens (Human)</i>					
	FRIL_HUMAN	R.LGGPEAGLGEYLFER.L	2	2.3823	P02792
<i>Filamin-A - Homo sapiens (Human)</i>					
	FLNA_HUMAN	R.SPYTVTVGQAC#NPSAC#R.A	2	2.7651	P21333
	FLNA_HUMAN	K.VTAQGPGLPSGNIANK.T	2	2.9905	P21333
	FLNA_HUMAN	K.VTAQGPGLPSGNIANK.T	2	2.6292	P21333
	FLNA_HUMAN	R.SPYTVTVGQAC#NPSAC#R.A	2	3.1685	P21333
	FLNA_HUMAN	R.AYGPGIEPTGNMVK.K	2	2.3232	P21333
	FLNA_HUMAN	R.VTYC#PTEPGNYIINIK.F	2	2.368	P21333
	FLNA_HUMAN	R.AGQSAAGAAPGGGVDR.D	2	2.5456	P21333
	FLNA_HUMAN	R.SPYTVTVGQAC#NPSAC#R.A	2	3.2514	P21333
	FLNA_HUMAN	K.VDINTEDLEDGTC#R.V	2	2.7771	P21333
	FLNA_HUMAN	K.YTPVQQGPVGVNVTYGGDPIPK.S	2	4.0491	P21333
	FLNA_HUMAN	K.YTPVQQGPVGVNVTYGGDPIPK.S	2	3.2362	P21333
	FLNA_HUMAN	K.YTPVQQGPVGVNVTYGGDPIPK.S	2	2.9129	P21333
<i>Formin-like protein 3 - Homo sapiens (Human)</i>					
	FMNL3_HUMAN	R.NFLSTNEGKLDKLR.D	2	2.5592	Q81VF7

Full-length cDNA clone CS0DI085YI08 of Placenta of Homo sapiens - Homo sapiens (Human)

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	Q86TV4_HUMAN	R.PTC#PSSSGGPARELSSRLR.I	2	2.7222	Q86TV4
	Q86TV4_HUMAN	R.PTC#PSSSGGPARELSSRLR.I	2	2.8843	Q86TV4
<i>Glia maturation factor gamma - Homo sapiens (Human)</i>					
	GMFG_HUMAN	K.ETDNAAIIMKVDK.D	2	2.5641	O60234
<i>Glyceraldehyde-3-phosphate dehydrogenase - Homo sapiens (Human)</i>					
	G3P_HUMAN	K.LVINGNPITIFQER.D	2	2.7138	P04406
<i>Glycoprotein Ib beta - Homo sapiens (Human)</i>					
	Q14422_HUMAN	R.LSLTDPLVAER.A	2	2.4589	Q14422
<i>Golgi-associated plant pathogenesis-related protein 1 - Homo sapiens (Human)</i>					
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	3.1934	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	3.03	Q9H4G4
	GAPR1_HUMAN	K.ASASDGSSFVVAR.Y	2	2.6071	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	3.2899	Q9H4G4
	GAPR1_HUMAN	K.HGVPPLK.L	2	2.584	Q9H4G4
	GAPR1_HUMAN	R.EAQQYSEALASTR.I	2	3.1447	Q9H4G4
	GAPR1_HUMAN	K.ASASDGSSFVVAR.Y	2	2.6673	Q9H4G4
<i>G-protein coupled receptor 98 precursor - Homo sapiens (Human)</i>					
	GPR98_HUMAN	R.NIIVSEDTQM*IR.L	2	2.458	Q8WXC9
<i>Growth hormone-inducible transmembrane protein - Homo sapiens (Human)</i>					
	GHITM_HUMAN	R.VATMLATGGNR.K	2	2.6046	Q9H3K2
	GHITM_HUMAN	K.YDPINSMLSIYMDTLNIFMR.V	2	4.6025	Q9H3K2
	GHITM_HUMAN	K.YDPINSMLSIYMDTLNIFMR.V	2	4.4829	Q9H3K2
	GHITM_HUMAN	R.VATMLATGGNR.K	2	2.6975	Q9H3K2
	GHITM_HUMAN	R.VATMLATGGNR.K	2	2.523	Q9H3K2
	GHITM_HUMAN	K.YDPINSMLSIYMDTLNIFMR.V	2	3.9739	Q9H3K2
	GHITM_HUMAN	R.TPVL MNFMMR.G	2	2.3116	Q9H3K2
<i>Growth-arrest-specific protein 2 - Homo sapiens (Human)</i>					
	GAS2_HUMAN	-.M*C#TALSPKVR.S	2	2.4491	O43903
<i>GTPase IMAP family member 1 - Homo sapiens (Human)</i>					
	GIMA1_HUMAN	R.FTAQDQQAVR.Q	2	3.2476	Q8WWP7
	GIMA1_HUMAN	R.FTAQDQQAVR.Q	2	3.4339	Q8WWP7
	GIMA1_HUMAN	R.FTAQDQQAVR.Q	2	2.3331	Q8WWP7
	GIMA1_HUMAN	R.FTAQDQQAVR.Q	2	3.2506	Q8WWP7
	GIMA1_HUMAN	R.FTAQDQQAVR.Q	2	3.9412	Q8WWP7

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Guanine nucleotide binding protein, alpha transducing 3 - Homo sapiens (Human)</i>					
A4D1B2_HUMAN		K.AM*TTLGIDYVNPR.S	2	2.4254	A4D1B2
<i>Guanine nucleotide-binding protein G(i), alpha-2 subunit - Homo sapiens (Human)</i>					
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.7287	P04899
GNAI2_HUMAN		R.EYQLNDSAAYYLNDLER.I	2	3.3445	P04899
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.1274	P04899
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.4711	P04899
GNAI2_HUMAN		R.AVVYSNTIQSIMAIVK.A	2	3.8353	P04899
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.306	P04899
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.3203	P04899
GNAI2_HUMAN		R.EYQLNDSAAYYLNDLER.I	2	2.4384	P04899
GNAI2_HUMAN		K.YDEAASYIQSK.F	2	2.4786	P04899
GNAI2_HUMAN		R.IAQSDYIPTQQDVL.R.T	2	4.048	P04899
GNAI2_HUMAN		K.AM*GNLQIDFADPSR.A	2	2.8106	P04899
GNAI2_HUMAN		K.AMGNLQIDFADPSR.A	2	3.2318	P04899
GNAI2_HUMAN		R.SREYQLNDSAAYYLNDLER.I	2	3.251	P04899
<i>Heat shock protein HSP 90-alpha - Homo sapiens (Human)</i>					
HS90A_HUMAN		K.YIDQEELNK.T	2	2.4172	P07900
<i>Heat shock protein HSP 90-beta - Homo sapiens (Human)</i>					
HS90B_HUMAN		R.NPDDITQEEYGEFYK.S	2	2.9143	P08238
<i>Heavy chain Fab - Homo sapiens (Human)</i>					
A2NYU6_HUMAN		K.SRVTMVSDTSKNQVSLR.L	2	2.3397	A2NYU6
<i>Heterogeneous nuclear ribonucleoprotein H - Homo sapiens (Human)</i>					
HNRH1_HUMAN		R.ATENDIYNFFSPLNPVR.V	2	2.4681	P31943
<i>Heterogeneous nuclear ribonucleoprotein K - Homo sapiens (Human)</i>					
HNRPK_HUMAN		K.IILDLISESPIK.G	2	2.6645	P61978
HNRPK_HUMAN		K.IILDLISESPIK.G	2	2.3501	P61978
<i>Heterogeneous nuclear ribonucleoprotein U - Homo sapiens (Human)</i>					
HNRPU_HUMAN		R.NFILDQTNVSAQAQR.R	2	2.9633	Q00839
<i>Hexokinase-1 - Homo sapiens (Human)</i>					
HXK1_HUMAN		K.MVSGMYLGELVR.L	2	2.5251	P19367
HXK1_HUMAN		K.MISGMYLGEIVR.N	2	2.6037	P19367
HXK1_HUMAN		R.LSDETLIDIMTR.F	2	2.6818	P19367
<i>Hexokinase-3 - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	3.2554	P52790
HXK3_HUMAN		R.QLGLDQGILLNWTG.G	2	2.9612	P52790
HXK3_HUMAN		K.QGLSGQSLPLGFTFSFPC#R.Q	2	3.086	P52790
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	3.1645	P52790
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	2.9111	P52790
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	3.0962	P52790
HXK3_HUMAN		K.ASDC#EGQDVVSLLR.E	2	2.3637	P52790
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	3.0211	P52790
HXK3_HUMAN		K.FLSEIESDSLALR.Q	2	2.5559	P52790

High affinity immunoglobulin epsilon receptor subunit gamma precursor - Homo sapiens (Hum)

FCERG_HUMAN		K.SDGVYTGLSTR.N	2	2.4618	P30273
FCERG_HUMAN		K.SDGVYTGLSTR.N	2	2.937	P30273
FCERG_HUMAN		K.SDGVYTGLSTR.N	2	2.834	P30273
FCERG_HUMAN		K.SDGVYTGLSTR.N	2	2.5583	P30273

Histone H1.1 - Homo sapiens (Human)

H11_HUMAN		K.ALAAAGYDVEK.N	2	2.5601	Q02539
H11_HUMAN		K.ALAAAGYDVEK.N	2	2.647	Q02539
H11_HUMAN		K.ALAAAGYDVEK.N	2	2.4942	Q02539
H11_HUMAN		K.ALAAAGYDVEK.N	2	2.3085	Q02539
H11_HUMAN		K.ALAAAGYDVEKNNSR.I	2	2.8677	Q02539

Histone H1.5 - Homo sapiens (Human)

H15_HUMAN		K.ALAAGGYDVEK.N	2	2.4425	P16401
H15_HUMAN		K.ALAAGGYDVEK.N	2	2.6342	P16401
H15_HUMAN		K.ALAAGGYDVEK.N	2	2.6702	P16401
H15_HUMAN		K.ALAAGGYDVEK.N	2	2.6067	P16401

Histone H2A type 1-A - Homo sapiens (Human)

H2A1A_HUMAN		R.HLQLAIR.N	2	2.4864	Q96QV6
H2A1A_HUMAN		R.AGLQFPVGR.I	2	2.5754	Q96QV6
H2A1A_HUMAN		R.AGLQFPVGR.I	2	2.6349	Q96QV6
H2A1A_HUMAN		R.HLQLAIR.N	2	2.6381	Q96QV6
H2A1A_HUMAN		R.HLQLAIR.N	2	2.3771	Q96QV6
H2A1A_HUMAN		R.AGLQFPVGR.I	2	2.5374	Q96QV6
H2A1A_HUMAN		R.HLQLAIR.N	2	2.3884	Q96QV6
H2A1A_HUMAN		R.AGLQFPVGR.I	2	2.4881	Q96QV6
H2A1A_HUMAN		R.HLQLAIR.N	2	2.575	Q96QV6

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
H2A1A_HUMAN		R.HLQLAIR.N	2	2.8487	Q96QV6
H2A1A_HUMAN		R.AGLQFPVGR.I	2	2.3302	Q96QV6
<i>Histone H2A type 1-B - Homo sapiens (Human)</i>					
H2A1B_HUMAN		R.VGAGAPVYLAADVLEYLTAIEILELAGNAAR.D	2	4.8376	P04908
H2A1B_HUMAN		R.HLQLAIRNDEELNKLLGR.V	4	4.6656	P04908
H2A1B_HUMAN		R.VTIAQGGVLPNIQAVLLPK.K	2	2.3523	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLEYLTAIEILELAGNAAR.D	2	4.4231	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLEYLTAIEILELAGNAAR.D	2	5.1416	P04908
H2A1B_HUMAN		R.VGAGAPVYLAADVLEYLTAIEILELAGNAAR.D	2	4.9606	P04908
H2A1B_HUMAN		R.HLQLAIRNDEELNKLLGR.V	2	4.0312	P04908
<i>Histone H2A type 1-D - Homo sapiens (Human)</i>					
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.2431	P20671
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.3089	P20671
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.5046	P20671
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.0467	P20671
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.3013	P20671
H2A1D_HUMAN		R.NDEELNKLLGK.V	2	3.3202	P20671
H2A1D_HUMAN		R.HLQLAIRNDEELNKLLGK.V	3	3.8348	P20671
H2A1D_HUMAN		R.HLQLAIRNDEELNKLLGK.V	2	3.5893	P20671
<i>Histone H2A type 2-A - Homo sapiens (Human)</i>					
H2A2A_HUMAN		R.VGAGAPVYMAADVLEYLTAIEILELAGNAAR.D	4	5.1835	Q6FI13
H2A2A_HUMAN		R.VGAGAPVYMAADVLEYLTAIEILELAGNAAR.D	2	4.0911	Q6FI13
H2A2A_HUMAN		R.VGAGAPVYMAADVLEYLTAIEILELAGNAAR.D	2	4.3754	Q6FI13
H2A2A_HUMAN		R.VGAGAPVYMAADVLEYLTAIEILELAGNAAR.D	2	4.9873	Q6FI13
H2A2A_HUMAN		R.VGAGAPVYMAADVLEYLTAIEILELAGNAAR.D	2	3.7722	Q6FI13
<i>Histone H2AV - Homo sapiens (Human)</i>					
H2AV_HUMAN		R.VGATAAVYSAAILEYLTAEVLELAGNASK.D	4	6.3814	Q71UI9
<i>Histone H2B type 1-B - Homo sapiens (Human)</i>					
H2B1B_HUMAN		R.KESYSIYVYK.V	2	2.3926	P33778
H2B1B_HUMAN		R.KESYSIYVYK.V	2	2.8637	P33778
H2B1B_HUMAN		K.AM*GIM*NSFVNDIFER.I	2	3.7546	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	3.9302	P33778
H2B1B_HUMAN		K.AM*GIMNSFVNDIFER.I	2	5.0029	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.7184	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	3.8548	P33778

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.0424	P33778
H2B1B_HUMAN		R.KESYSIYVYK.V	2	2.3577	P33778
H2B1B_HUMAN		K.AM*GIM*NSFVNDIFER.I	2	3.5253	P33778
H2B1B_HUMAN		K.AM*GIMNSFVNDIFER.I	2	3.9563	P33778
H2B1B_HUMAN		K.AM*GIMNSFVNDIFER.I	2	4.4609	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.8814	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	5.0904	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFERIAGEASR.L	2	3.2958	P33778
H2B1B_HUMAN		R.KESYSIYVYK.V	2	2.3729	P33778
H2B1B_HUMAN		R.KESYSIYVYK.V	2	3.0414	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	2.9906	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	4.691	P33778
H2B1B_HUMAN		K.AM*GIMNSFVNDIFER.I	2	4.619	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.3402	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.639	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	3.0942	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	3.1586	P33778
H2B1B_HUMAN		R.KESYSIYVYK.V	2	3.5188	P33778
H2B1B_HUMAN		K.AM*GIM*NSFVNDIFER.I	2	3.2665	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	2.5672	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	2.9967	P33778
H2B1B_HUMAN		K.AMGIM*NSFVNDIFER.I	2	2.6764	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	3.6348	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	4.4809	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	2	2.5541	P33778
H2B1B_HUMAN		K.AMGIMNSFVNDIFER.I	1	3.0751	P33778

Histone H2B type 1-C/E/F/G/I - Homo sapiens (Human)

H2B1C_HUMAN		R.KESYSVYVYK.V	2	2.949	P62807
H2B1C_HUMAN		R.KESYSVYVYK.V	2	2.4358	P62807
H2B1C_HUMAN		R.KESYSVYVYK.V	2	3.2853	P62807
H2B1C_HUMAN		R.KESYSVYVYK.V	2	2.9715	P62807
H2B1C_HUMAN		R.KESYSVYVYK.V	2	3.1679	P62807

Histone H2B type 2-C - Homo sapiens (Human)

H2B2C_HUMAN		K.AMGIM*NSFLNDIFER.I	2	2.3462	Q6DN03
H2B2C_HUMAN		K.AM*GIMNSFLNDIFER.I	2	2.6029	Q6DN03
H2B2C_HUMAN		K.AM*GIMNSFLNDIFER.I	2	2.6266	Q6DN03

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
H2B2C_HUMAN		K.AM*GIMNSFLNDIFER.I	2	2.3807	Q6DN03
H2B2C_HUMAN		K.AMGIMNSFLNDIFER.I	2	3.6176	Q6DN03
H2B2C_HUMAN		K.AM*GIMNSFLNDIFER.I	2	2.3895	Q6DN03
H2B2C_HUMAN		K.AMGIMNSFLNDIFER.I	2	3.3873	Q6DN03
H2B2C_HUMAN		K.AM*GIM*NSFLNDIFER.I	2	2.8391	Q6DN03
H2B2C_HUMAN		K.AM*GIMNSFLNDIFER.I	2	2.5185	Q6DN03

Histone H3.It - Homo sapiens (Human)

H31T_HUMAN		R.EIAQDFKTDLR.F	2	2.4713	Q16695
H31T_HUMAN		K.STELLIR.K	2	2.4172	Q16695
H31T_HUMAN		R.EIAQDFK.T	1	2.0046	Q16695
H31T_HUMAN		R.EIAQDFKTDLR.F	2	2.3565	Q16695
H31T_HUMAN		R.EIAQDFKTDLR.F	2	2.7251	Q16695

Histone H4 - Homo sapiens (Human)

H4_HUMAN		R.DAVTYTEHAK.R	2	3.1602	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.3861	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	2.9938	P62805
H4_HUMAN		R.ISGLIYEETR.G	2	2.9208	P62805
H4_HUMAN		K.TVTAMDVVYALKR.Q	2	2.8223	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	3.2857	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	2.585	P62805
H4_HUMAN		K.VFLENVIRDAVTYTEHAK.R	3	4.9742	P62805
H4_HUMAN		R.DAVTYTEHAK.R	2	3.1041	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.4877	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.1422	P62805
H4_HUMAN		R.ISGLIYEETR.G	2	3.1062	P62805
H4_HUMAN		R.KTVTAMDVVYALKR.Q	3	5.1125	P62805
H4_HUMAN		K.TVTAMDVVYALKR.Q	2	2.987	P62805
H4_HUMAN		K.TVTAMDVVYALKR.Q	2	3.1027	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	2.4941	P62805
H4_HUMAN		R.ISGLIYEETRGLVKVFLENVIR.D	3	5.0535	P62805
H4_HUMAN		R.DAVTYTEHAK.R	2	3.4207	P62805
H4_HUMAN		R.DAVTYTEHAK.R	2	2.5897	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.4872	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.1633	P62805
H4_HUMAN		R.ISGLIYEETR.G	2	3.0256	P62805
H4_HUMAN		K.VFLENVIR.D	2	2.3686	P62805

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
H4_HUMAN		K.TVTAMDVVYALK.R	2	2.7565	P62805
H4_HUMAN		R.DAVTYTEHAK.R	2	3.1298	P62805
H4_HUMAN		R.DNIQGITKPAIR.R	2	3.3804	P62805
H4_HUMAN		R.ISGLIYEETR.G	2	3.1359	P62805
H4_HUMAN		K.TVTAM*DVVYALK.R	2	2.3458	P62805
H4_HUMAN		K.TVTAMDVVYALKR.Q	2	2.8216	P62805
H4_HUMAN		K.TVTAMDVVYALKR.Q	2	3.1439	P62805
H4_HUMAN		K.TVTAMDVVYALK.R	2	2.45	P62805
H4_HUMAN		R.ISGLIYEETRGLK.V	2	3.2378	P62805
H4_HUMAN		R.ISGLIYEETRGLK.V	2	2.3915	P62805
H4_HUMAN		R.ISGLIYEETRGLKVFLENVIR.D	3	4.8495	P62805

HLA class I histocompatibility antigen, A-1 alpha chain precursor - Homo sapiens (Human)

1A01_HUMAN		K.WAAVVVPSGEEQR.Y	2	2.3943	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	3.418	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	2.5477	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	3.4356	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	2.7491	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	3.2646	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	2.7254	P30443
1A01_HUMAN		R.FIAVGYVDDTQFVR.F	2	2.4514	P30443

HLA class I histocompatibility antigen, A-2 alpha chain precursor - Homo sapiens (Human)

1A02_HUMAN		R.FSDAASQR.M	2	2.3133	P01892
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HLA class I histocompatibility antigen, A-25 alpha chain precursor - Homo sapiens (Human)

1A25_HUMAN		R.APWIEQEGPEYWR.N	2	3.4924	P18462
1A25_HUMAN		R.APWIEQEGPEYWR.N	2	3.7141	P18462
1A25_HUMAN		R.APWIEQEGPEYWR.N	2	3.6917	P18462
1A25_HUMAN		R.APWIEQEGPEYWR.N	2	2.354	P18462
1A25_HUMAN		R.APWIEQEGPEYWR.N	2	3.0113	P18462

HLA class I histocompatibility antigen, B-7 alpha chain precursor - Homo sapiens (Human)

1B07_HUMAN		R.FSDAASPR.E	2	2.5507	P01889
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HLA class II histocompatibility antigen gamma chain - Homo sapiens (Human)

HG2A_HUMAN		K.LTVTSQNLQLENLR.M	2	2.5326	P04233
HG2A_HUMAN		K.LTVTSQNLQLENLR.M	2	3.5097	P04233
HG2A_HUMAN		K.LTVTSQNLQLENLR.M	2	3.1517	P04233

HLA class II histocompatibility antigen, DR alpha chain precursor - Homo sapiens (Human)

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	4.0398	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	4.1845	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	3.8184	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	3.8187	P01903
2DRA_HUMAN		R.NGKPVTTGVSETVFLPR.E	2	3.2359	P01903
2DRA_HUMAN		R.NGKPVTTGVSETVFLPR.E	2	2.6158	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	3.7092	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	5.6319	P01903
2DRA_HUMAN		R.FASFEAQQGALANIAVDK.A	2	2.3309	P01903
<i>HLA class II histocompatibility antigen, DRB1-1 beta chain precursor - Homo sapiens (Human)</i>					
2B11_HUMAN		R.FDSVDGEYR.A	2	3.2815	P04229
2B11_HUMAN		R.FDSVDGEYR.A	2	2.5855	P04229
2B11_HUMAN		R.FDSVDGEYR.A	2	2.3673	P04229
2B11_HUMAN		R.HNYGVGESFTVQR.R	2	2.9717	P04229
<i>HLA class II histocompatibility antigen, DRB1-8 beta chain precursor - Homo sapiens (Human)</i>					
2B18_HUMAN		R.YFYNQEEYVR.F	2	2.6442	Q30134
2B18_HUMAN		R.FLDRYFYNQEEYVR.F	2	3.4517	Q30134
2B18_HUMAN		R.FLDRYFYNQEEYVR.F	2	2.4914	Q30134
2B18_HUMAN		R.YFYNQEEYVR.F	2	2.6096	Q30134
2B18_HUMAN		R.FLDRYFYNQEEYVR.F	2	2.4321	Q30134
<i>HLA class II histocompatibility antigen, DR-W53 beta chain precursor - Homo sapiens (Human)</i>					
HB2K_HUMAN		R.YIYNQEEYAR.Y	2	2.7151	P13762
HB2K_HUMAN		R.YIYNQEEYAR.Y	2	2.5839	P13762
HB2K_HUMAN		R.YIYNQEEYAR.Y	2	2.5283	P13762
HB2K_HUMAN		R.YIYNQEEYAR.Y	2	2.6057	P13762
<i>IGHM protein - Homo sapiens (Human)</i>					
Q6GMX5_HUMAN		K.QVGSVTTDQVQAEAK.E	2	3.1113	Q6GMX5
<i>Immediate early response 3-interacting protein 1 - Homo sapiens (Human)</i>					
IR3IP_HUMAN		K.NIGWGTDQGIGGFGEPEGIK.S	2	3.2637	Q9Y5U9
<i>Inhibitor of nuclear factor kappa-B kinase subunit beta - Homo sapiens (Human)</i>					
IKKB_HUMAN		R.EGAILTLLSDIASALR.Y	2	2.5131	O14920
<i>Integrin alpha-IIb precursor - Homo sapiens (Human)</i>					
ITA2B_HUMAN		R.NRPPLEEDDEEGE.-	2	2.8583	P08514
ITA2B_HUMAN		R.ALSNVEGFER.L	2	2.3612	P08514
ITA2B_HUMAN		R.GEAQVWTQLLR.A	2	2.7571	P08514

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ITA2B_HUMAN		R.SRPSQVLDSPFPTGSAFGFSLR.G	3	3.8215	P08514
ITA2B_HUMAN		R.ALSNVEGFER.L	2	2.4009	P08514
ITA2B_HUMAN		R.GEAQVWTQLLR.A	2	2.7388	P08514
ITA2B_HUMAN		R.GEAQVWTQLLR.A	2	2.7835	P08514
ITA2B_HUMAN		R.NRPPEEDDEEGE.-	2	2.671	P08514
ITA2B_HUMAN		R.ALSNVEGFER.L	2	2.4183	P08514
ITA2B_HUMAN		R.GEAQVWTQLLR.A	2	2.5278	P08514
ITA2B_HUMAN		R.ALSNVEGFER.L	2	2.3724	P08514
ITA2B_HUMAN		R.NRPPEEDDEEGE.-	2	2.6832	P08514
<i>Integrin alpha-M precursor - Homo sapiens (Human)</i>					
ITAM_HUMAN		R.VVVGAPQEIVAANQR.G	2	3.4419	P11215
ITAM_HUMAN		R.GFGQSVVQLQGSR.V	2	2.8158	P11215
ITAM_HUMAN		R.LFTALFPFEK.N	2	2.3217	P11215
ITAM_HUMAN		R.VMQHQYQVSNLGQR.S	3	3.7874	P11215
ITAM_HUMAN		R.GFGQSVVQLQGSR.V	2	3.2979	P11215
ITAM_HUMAN		R.VMQHQYQVSNLGQR.S	3	4.3306	P11215
ITAM_HUMAN		R.VVVGAPQEIVAANQR.G	2	3.2634	P11215
ITAM_HUMAN		R.LFTALFPFEK.N	2	2.3074	P11215
ITAM_HUMAN		R.LFTALFPFEK.N	2	2.49	P11215
ITAM_HUMAN		R.VMQHQYQVSNLGQR.S	3	4.3896	P11215
ITAM_HUMAN		R.GFGQSVVQLQGSR.V	2	3.1732	P11215
ITAM_HUMAN		R.MKEFVSTVMEQLKK.S	3	3.8276	P11215
ITAM_HUMAN		R.VMQHQYQVSNLGQR.S	2	3.8065	P11215
ITAM_HUMAN		R.VMQHQYQVSNLGQR.S	2	3.2777	P11215
ITAM_HUMAN		R.QYKDMMSEGGPPGAEPQ.-	2	2.7639	P11215
ITAM_HUMAN		R.VVVGAPQEIVAANQR.G	2	2.6602	P11215
ITAM_HUMAN		K.YAVYMVVTSHGVSTK.Y	2	2.8447	P11215
<i>Integrin beta chain, beta 2 variant - Homo sapiens (Human)</i>					
Q59H50_HUMAN		K.SAVGELSESSNVVHLIK.N	2	4.0551	Q59H50
Q59H50_HUMAN		K.SAVGELSESSNVVHLIK.N	2	3.7355	Q59H50
Q59H50_HUMAN		K.SAVGELSESSNVVHLIK.N	2	3.5128	Q59H50
<i>Integrin beta-2 precursor - Homo sapiens (Human)</i>					
ITB2_HUMAN		R.ALNEITESGR.I	2	2.8005	P05107
ITB2_HUMAN		R.VFLDHNALPDTLK.V	2	3.761	P05107
ITB2_HUMAN		K.VTATEC#IQEQSFVIR.A	2	3.3421	P05107

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	3.3084	P05107
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	2.3064	P05107
ITB2_HUMAN		R.LLVFATDDGFHFAGDGK.L	2	3.1178	P05107
ITB2_HUMAN		R.YNGQVC#GGPGR.G	2	2.6449	P05107
ITB2_HUMAN		R.ALNEITESGR.I	2	2.696	P05107
ITB2_HUMAN		K.LGAILTPNDGR.C	2	2.5318	P05107
ITB2_HUMAN		R.VFLDHNALPDTLK.V	2	3.7362	P05107
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	3.0422	P05107
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	3.7888	P05107
ITB2_HUMAN		R.LLVFATDDGFHFAGDGK.L	2	3.6021	P05107
ITB2_HUMAN		K.GYPIDLYYLDLSYSMLDDL.R.N	2	3.16	P05107
ITB2_HUMAN		R.YNGQVC#GGPGR.G	2	2.7069	P05107
ITB2_HUMAN		R.ALNEITESGR.I	2	2.8597	P05107
ITB2_HUMAN		K.LGGDLLR.A	2	2.3238	P05107
ITB2_HUMAN		K.VTATEC#IQEQSFVIR.A	2	2.5082	P05107
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	3.1983	P05107
ITB2_HUMAN		K.LAENNIQPIFAVTSR.M	2	3.0969	P05107
ITB2_HUMAN		R.LLVFATDDGFHFAGDGK.L	2	2.4688	P05107
ITB2_HUMAN		K.GYPIDLYYLDLSYSMLDDL.R.N	2	3.9499	P05107
ITB2_HUMAN		R.YNGQVC#GGPGR.G	2	2.5124	P05107
ITB2_HUMAN		R.TTEGC#LNPR.R	2	2.772	P05107
ITB2_HUMAN		R.C#HPGFEGSAC#QC#ER.T	3	4.1009	P05107
ITB2_HUMAN		R.ALNEITESGR.I	2	2.9642	P05107
ITB2_HUMAN		R.SNEFDYPSVQLAHLK.L	2	3.3725	P05107
ITB2_HUMAN		R.VFLDHNALPDTLK.V	2	2.4559	P05107
ITB2_HUMAN		K.VTATEC#IQEQSFVIR.A	2	3.1386	P05107
ITB2_HUMAN		R.LLVFATDDGFHFAGDGK.L	2	3.0177	P05107
<i>Integrin beta-3 precursor - Homo sapiens (Human)</i>					
ITB3_HUMAN		K.GSGDSSQVTQVSPQR.I	2	3.7417	P05106
ITB3_HUMAN		K.HVLTLTDQVTR.F	2	2.7605	P05106
ITB3_HUMAN		K.GSGDSSQVTQVSPQR.I	2	3.1899	P05106
ITB3_HUMAN		K.GSGDSSQVTQVSPQR.I	2	3.4351	P05106
<i>Interferon-induced transmembrane protein 1 - Homo sapiens (Human)</i>					
IFM1_HUMAN		K.MVGDVTGAQAYASTAK.C	2	3.7608	P13164
IFM1_HUMAN		K.MVGDVTGAQAYASTAK.C	2	3.8018	P13164
IFM1_HUMAN		K.MVGDVTGAQAYASTAK.C	2	3.8894	P13164

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>IQ motif and Sec7 domain-containing protein 3 - Homo sapiens (Human)</i>					
	IQEC3_HUMAN	R.QMIGEFLGNSKKQFNR.D	2	2.3412	Q9UAPP2
<i>Jagged-1 precursor - Homo sapiens (Human)</i>					
	JAG1_HUMAN	K.IIDLVSKR.D	2	2.3394	P78504
<i>Lamin-B receptor - Homo sapiens (Human)</i>					
	LBR_HUMAN	K.VVEGTPLIDGR.R	2	2.4417	Q14739
	LBR_HUMAN	K.VVEGTPLIDGR.R	2	2.4716	Q14739
	LBR_HUMAN	K.VVEGTPLIDGR.R	2	2.4056	Q14739
	LBR_HUMAN	R.NDLSPASSGNAVYDFFIGR.E	2	3.2656	Q14739
	LBR_HUMAN	K.VVEGTPLIDGR.R	2	2.3457	Q14739
<i>Lamin-B1 - Homo sapiens (Human)</i>					
	LMNB1_HUMAN	R.LSSEMNTSTVNSAR.E	2	2.8377	P20700
	LMNB1_HUMAN	R.LSSEMNTSTVNSAR.E	2	3.3151	P20700
<i>Leukocyte common antigen precursor - Homo sapiens (Human)</i>					
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	4.3277	P08575
	CD45_HUMAN	R.LFLAEFQSIPR.V	2	2.6115	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	4.9279	P08575
	CD45_HUMAN	R.LFLAEFQSIPR.V	2	2.8764	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	4.7381	P08575
	CD45_HUMAN	R.LFLAEFQSIPR.V	2	3.0768	P08575
	CD45_HUMAN	R.SC#NLDEQQELVER.D	2	3.8775	P08575
	CD45_HUMAN	R.VKDLQYSTDYTFK.A	2	3.197	P08575
<i>Leukocyte elastase inhibitor - Homo sapiens (Human)</i>					
	ILEU_HUMAN	R.LGVQDLFNSSK.A	2	3.2356	P30740
	ILEU_HUMAN	K.LVLVNAIFYK.G	2	2.6079	P30740
	ILEU_HUMAN	K.LVLVNAIFYK.G	2	2.8507	P30740
	ILEU_HUMAN	R.FKLEESYTLNSDLAR.L	2	2.6394	P30740
	ILEU_HUMAN	K.IPELLASGMVDNMTK.L	2	3.5882	P30740
<i>Leukocyte elastase precursor - Homo sapiens (Human)</i>					
	ELNE_HUMAN	R.VVLGAHNLSR.R	2	2.5746	P08246
	ELNE_HUMAN	R.VVLGAHNLSR.R	2	2.5131	P08246
	ELNE_HUMAN	R.LGNGVQC#LAMGWLLGR.N	2	2.6474	P08246
	ELNE_HUMAN	R.LGNGVQC#LAMGWLLGR.N	2	2.5082	P08246
	ELNE_HUMAN	R.LGNGVQC#LAMGWLLGR.N	2	3.1195	P08246

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Leukocyte immunoglobulin-like receptor subfamily B member 1 precursor - Homo sapiens (Hu)</i>					
LIRB1_HUMAN		R.KADFQHPAGAVGPEPTDR.G	3	4.0669	Q8NHL6
LIRB1_HUMAN		R.SSPAADAQEENLYAAVK.H	2	3.4819	Q8NHL6
<i>Leukosialin precursor - Homo sapiens (Human)</i>					
LEUK_HUMAN		R.TGALVLSR.G	2	2.6072	P16150
<i>Lysozyme C precursor - Homo sapiens (Human)</i>					
LYSC_HUMAN		R.ATNYNAGDR.S	2	2.398	P61626
LYSC_HUMAN		K.RLGMDGYR.G	2	2.4653	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	1	2.3904	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	2.391	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	3.11	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	3.3342	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	3.6745	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	3.305	P61626
LYSC_HUMAN		R.ATNYNAGDR.S	2	2.6106	P61626
LYSC_HUMAN		K.RLGMDGYR.G	2	2.4501	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	2	2.9349	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	1	2.6812	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	2.9514	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	2.8318	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	3.3712	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	1	2.1476	P61626
LYSC_HUMAN		R.ATNYNAGDR.S	2	2.4955	P61626
LYSC_HUMAN		R.ATNYNAGDR.S	2	2.3713	P61626
LYSC_HUMAN		R.ATNYNAGDR.S	2	2.6471	P61626
LYSC_HUMAN		K.RLGMDGYR.G	2	2.7299	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	2	3.8134	P61626
LYSC_HUMAN		K.TPGAVNACHLSCSALLQDNIADAVAC#AK.R	3	4.3733	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	2.4242	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	2.7271	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	1	2.1776	P61626
LYSC_HUMAN		R.VVRDPQGIR.A	2	2.5284	P61626
LYSC_HUMAN		K.RLGMDGYR.G	2	2.7245	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	2	3.9565	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	2	4.1827	P61626

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
LYSC_HUMAN		R.STDYGIFQINSR.Y	2	2.9782	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	2.5283	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	2.8494	P61626
LYSC_HUMAN		R.GISLANWM*C#LAK.W	2	3.2983	P61626
LYSC_HUMAN		R.GISLANWMC#LAK.W	2	2.5992	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	1	3.026	P61626
LYSC_HUMAN		R.STDYGIFQINSR.Y	1	2.893	P61626
LYSC_HUMAN		K.TPGAVNAC#HLSC#SALLQDNIADAVAC#AK.R	2	4.0675	P61626
<i>Mannose-P-dolichol utilization defect 1 protein - Homo sapiens (Human)</i>					
MPU1_HUMAN		K.GLGLGIVAGSLLVK.L	2	2.9635	O75352
MPU1_HUMAN		K.GLGLGIVAGSLLVK.L	2	2.9374	O75352
MPU1_HUMAN		K.GLGLGIVAGSLLVK.L	2	2.7362	O75352
<i>Mannosyl-oligosaccharide glucosidase - Homo sapiens (Human)</i>					
GCS1_HUMAN		R.LGPLLDILADSR.H	2	2.6688	Q13724
GCS1_HUMAN		R.VTVEPQDSGTSALPLVSLFFYVVDGKEVLLPEVG	3	3.864	Q13724
<i>Mast cell-expressed membrane protein 1 - Homo sapiens (Human)</i>					
MCEM1_HUMAN		R.SKIDRLETTLAGIK.N	2	3.1752	Q8IX19
MCEM1_HUMAN		R.GWDSVQQSITM*VR.S	2	2.8646	Q8IX19
MCEM1_HUMAN		R.GWDSVQQSITMVR.S	2	2.8342	Q8IX19
MCEM1_HUMAN		R.GWDSVQQSITMVR.S	2	2.715	Q8IX19
MCEM1_HUMAN		R.GWDSVQQSITMVR.S	2	3.1165	Q8IX19
<i>MBC3205 - Homo sapiens (Human)</i>					
Q6UW68_HUMAN		R.GLGGEVPGSHQGPDPYR.Q	2	2.8264	Q6UW68
<i>Membrane-associated progesterone receptor component 2 - Homo sapiens (Human)</i>					
PGRC2_HUMAN		K.FYGPAGPYGIFAGR.D	2	2.7791	O15173
PGRC2_HUMAN		K.FYGPAGPYGIFAGR.D	2	2.4289	O15173
PGRC2_HUMAN		R.GLGAGAGAGEESPATSLPR.M	2	3.1144	O15173
<i>Methyltransferase-like protein 7A precursor - Homo sapiens (Human)</i>					
MET7A_HUMAN		R.VTC#IDPNPNFEK.F	2	2.6695	Q9H8H3
MET7A_HUMAN		K.LSLLEVGC#GTGANFK.F	2	3.5125	Q9H8H3
MET7A_HUMAN		R.ELFSNLQEFAGPSGK.L	2	3.3522	Q9H8H3
MET7A_HUMAN		R.VTC#IDPNPNFEK.F	2	2.9438	Q9H8H3
MET7A_HUMAN		R.VTC#IDPNPNFEK.F	2	3.1708	Q9H8H3
MET7A_HUMAN		K.LSLLEVGC#GTGANFK.F	2	3.3575	Q9H8H3
MET7A_HUMAN		R.ELFSNLQEFAGPSGK.L	2	2.8285	Q9H8H3

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Microsomal glutathione S-transferase 1 - Homo sapiens (Human)</i>					
MGST1_HUMAN		R.KVFANPEDC#VAFGK.G	2	3.5222	P10620
MGST1_HUMAN		K.MMLMSTATAFYR.L	2	3.2697	P10620
MGST1_HUMAN		K.MMLMSTATAFYR.L	2	2.6095	P10620
MGST1_HUMAN		R.KVFANPEDC#VAFGK.G	2	2.9192	P10620
MGST1_HUMAN		K.MMLMSTATAFYR.L	2	2.6726	P10620
MGST1_HUMAN		K.VFANPEDC#VAFGK.G	2	2.5656	P10620
MGST1_HUMAN		K.MMLMSTATAFYR.L	2	2.5862	P10620
MGST1_HUMAN		K.MMLMSTATAFYR.L	2	3.1998	P10620
MGST1_HUMAN		R.IYHTIAYLTPLPQPNR.A	2	3.4624	P10620
<i>Microsomal glutathione S-transferase 3 - Homo sapiens (Human)</i>					
MGST3_HUMAN		R.VLYAYGYTGEPSKR.S	2	3.1094	O14880
MGST3_HUMAN		K.YKVEYPIMYSTDPENGHIFNC#IQR.A	3	4.2044	O14880
MGST3_HUMAN		R.IASGLGLAWIVGR.V	2	3.4553	O14880
MGST3_HUMAN		R.IASGLGLAWIVGR.V	2	2.9675	O14880
MGST3_HUMAN		R.IASGLGLAWIVGR.V	2	2.8869	O14880
MGST3_HUMAN		R.VLYAYGYTGEPSKR.S	2	3.4215	O14880
MGST3_HUMAN		R.VLYAYGYTGEPSKR.S	2	3.4692	O14880
MGST3_HUMAN		R.IASGLGLAWIVGR.V	2	3.2573	O14880
MGST3_HUMAN		R.IASGLGLAWIVGR.V	2	3.2765	O14880
MGST3_HUMAN		R.VLYAYGYTGEPSKR.S	2	2.8933	O14880
<i>Mitochondrial 2-oxoglutarate/malate carrier protein - Homo sapiens (Human)</i>					
M2OM_HUMAN		R.LGIYTVLFR.L	2	2.6188	Q02978
<i>Mitochondrial carrier homolog 2 - Homo sapiens (Human)</i>					
MTCH2_HUMAN		K.VLIQVGYEPLPPTIGR.N	2	2.3481	Q9Y6C9
MTCH2_HUMAN		K.VLIQVGYEPLPPTIGR.N	2	2.3074	Q9Y6C9
<i>Mitochondrial import receptor subunit TOM22 homolog - Homo sapiens (Human)</i>					
TOM22_HUMAN		K.LQMEQQQLQQR.Q	2	4.001	Q9NS69
<i>Monocyte differentiation antigen CD14 precursor - Homo sapiens (Human)</i>					
CD14_HUMAN		R.LTVGAAQVPAQLLVGALR.V	2	2.4556	P08571
CD14_HUMAN		K.FPAIQNLALR.N	2	2.7192	P08571
CD14_HUMAN		K.FPAIQNLALR.N	2	2.6048	P08571
CD14_HUMAN		R.AFPALTSLDLSDNPGLGER.G	2	3.6198	P08571
CD14_HUMAN		R.LTVGAAQVPAQLLVGALR.V	2	3.3594	P08571
CD14_HUMAN		R.LTVGAAQVPAQLLVGALR.V	2	2.9632	P08571

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	CD14_HUMAN	R.AFPALTSLDLSDNPGLGER.G	2	5.5742	P08571
	CD14_HUMAN	R.SWLAELQQWLKPKLK.V	2	2.8424	P08571
	CD14_HUMAN	R.AFPALTSLDLSDNPGLGER.G	2	3.6414	P08571
<i>Myeloblastin precursor - Homo sapiens (Human)</i>					
	PRTN3_HUMAN	R.LVNVVLGAHNVR.T	2	2.7012	P24158
	PRTN3_HUMAN	R.LVNVVLGAHNVR.T	2	2.8274	P24158
	PRTN3_HUMAN	R.LVNVVLGAHNVR.T	2	2.8236	P24158
	PRTN3_HUMAN	R.LVNVVLGAHNVR.T	2	3.1136	P24158
<i>Myeloid cell nuclear differentiation antigen - Homo sapiens (Human)</i>					
	MNDA_HUMAN	K.LVC#GSHSFIK.V	2	2.5536	P41218
	MNDA_HUMAN	K.VITISDYSEC#K.G	2	2.3314	P41218
	MNDA_HUMAN	K.QASGTMVYGLFMLQK.K	2	2.332	P41218
	MNDA_HUMAN	K.VITISDYSEC#K.G	2	2.572	P41218
	MNDA_HUMAN	K.NTIYEIQDNTGSMDVVGSGK.W	2	4.5772	P41218
<i>Myeloid-associated differentiation marker - Homo sapiens (Human)</i>					
	MYADM_HUMAN	R.TTITTTTTSSSGLSPMIVGSPR.A	2	4.8381	Q96S97
<i>Myeloperoxidase precursor - Homo sapiens (Human)</i>					
	PERM_HUMAN	R.FPTDQLTPDQER.S	2	3.5995	P05164
	PERM_HUMAN	K.NNIFM*SNSYPR.D	2	2.6143	P05164
	PERM_HUMAN	R.VVLEGGIDPILR.G	2	2.8049	P05164
	PERM_HUMAN	R.VVLEGGIDPILR.G	2	2.7576	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.4528	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.7467	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.5637	P05164
	PERM_HUMAN	R.WLPAEYEDGFSLPYGWTPGVK.R	2	2.537	P05164
	PERM_HUMAN	R.FPTDQLTPDQER.S	2	3.6082	P05164
	PERM_HUMAN	R.IANVFTNAFR.Y	2	3.1063	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.7387	P05164
	PERM_HUMAN	K.NNIFM*SNSYPR.D	2	2.6979	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.6975	P05164
	PERM_HUMAN	R.WLPAEYEDGFSLPYGWTPGVK.R	2	3.3398	P05164
	PERM_HUMAN	K.NNIFMSNSYPR.D	2	2.949	P05164
	PERM_HUMAN	R.VVLEGGIDPILR.G	2	2.3521	P05164
	PERM_HUMAN	R.VVLEGGIDPILR.G	2	2.3203	P05164
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	2.5673	P05164

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	PERM_HUMAN	R.IGLDLPALNMQR.S	2	3.2533	P05164
<i>Myosin-10 - Homo sapiens (Human)</i>					
	MYH10_HUMAN	K.KFDQLLAEEK.S	2	2.4609	P35580
	MYH10_HUMAN	K.KFDQLLAEEK.S	2	2.6724	P35580
	MYH10_HUMAN	K.KFDQLLAEEK.S	2	2.4628	P35580
	MYH10_HUMAN	R.AGVLAHLEEEER.D	2	3.0529	P35580
<i>Myosin-11 - Homo sapiens (Human)</i>					
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	2.7626	P35749
	MYH11_HUMAN	K.QLLQANPILEAFGNAK.T	2	3.1103	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.8828	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.5119	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	3.0553	P35749
	MYH11_HUMAN	R.NTDQASMPDNTAAQK.V	2	2.6505	P35749
<i>Myosin-9 - Homo sapiens (Human)</i>					
	MYH9_HUMAN	R.ELEDATETADAMNR.E	2	3.1606	P35579
	MYH9_HUMAN	K.IAQLEEQLDNETK.E	2	4.6601	P35579
	MYH9_HUMAN	K.IAQLEEQLDNETK.E	2	4.4803	P35579
	MYH9_HUMAN	R.IMGIPEEEQMGLLR.V	2	2.6806	P35579
	MYH9_HUMAN	R.IMGIPEEEQMGLLR.V	2	2.3042	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFKK.E	2	2.4947	P35579
	MYH9_HUMAN	R.TFHIFYLLSGAGEHLK.T	3	4.6782	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFK.K	2	3.1529	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFK.K	2	4.0607	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFK.K	2	2.759	P35579
	MYH9_HUMAN	K.IAQLEEQLDNETK.E	2	2.9088	P35579
	MYH9_HUMAN	R.IAEFTTNLTETEEEEK.S	2	3.4293	P35579
	MYH9_HUMAN	K.ANLQIDQINTDLNLER.S	2	3.9662	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFK.K	2	2.6211	P35579
	MYH9_HUMAN	K.IAQLEEQLDNETK.E	2	4.5863	P35579
	MYH9_HUMAN	K.IAQLEEQLDNETK.E	2	4.1911	P35579
	MYH9_HUMAN	K.ALELDSNLYR.I	2	2.4031	P35579
	MYH9_HUMAN	K.ANLQIDQINTDLNLER.S	2	2.5897	P35579
	MYH9_HUMAN	R.IMGIPEEEQMGLLR.V	2	2.5255	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFKK.E	2	2.9152	P35579
	MYH9_HUMAN	R.VISGVLQLGNIVFK.K	2	4.0932	P35579

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
MYH9_HUMAN		R.TVGQLYKEQLAK.L	2	2.8696	P35579
MYH9_HUMAN		R.VISGVLQLGNIVFK.K	2	2.4986	P35579
MYH9_HUMAN		K.KLEEEQIILEDQNC#K.L	2	4.392	P35579
MYH9_HUMAN		K.IAQLEEQLDNETK.E	2	3.1768	P35579
MYH9_HUMAN		R.IAEFTTNLTETEEEK.S	2	2.946	P35579
MYH9_HUMAN		R.IAEFTTNLTETEEEK.S	2	4.4916	P35579
MYH9_HUMAN		K.TDLLLEPYNKYR.F	2	2.3255	P35579
MYH9_HUMAN		R.IMGIPEEEQMGLLR.V	2	2.4661	P35579
MYH9_HUMAN		R.VISGVLQLGNIVFKK.E	2	3.0503	P35579
MYH9_HUMAN		R.VISGVLQLGNIVFK.K	2	3.4446	P35579
MYH9_HUMAN		R.VISGVLQLGNIVFK.K	2	4.0141	P35579

Myosin-IG - Homo sapiens (Human)

Q6ZNK5_HUMAN		R.LLYNSTDPTLR.A	2	2.4388	Q6ZNK5
Q6ZNK5_HUMAN		R.DYLSSATDNPTASSLFAQR.L	2	4.2567	Q6ZNK5
Q6ZNK5_HUMAN		R.LIPIVLLLQK.A	2	2.7839	Q6ZNK5
Q6ZNK5_HUMAN		R.LIPIVLLLQK.A	2	2.9049	Q6ZNK5
Q6ZNK5_HUMAN		R.LIPIVLLLQK.A	2	2.5603	Q6ZNK5
Q6ZNK5_HUMAN		R.DYLSSATDNPTASSLFAQR.L	2	3.3048	Q6ZNK5
Q6ZNK5_HUMAN		R.LIPIVLLLQK.A	2	2.7466	Q6ZNK5
Q6ZNK5_HUMAN		K.VAAMGALQGLR.Q	2	2.5136	Q6ZNK5

NACHT, LRR and PYD domains-containing protein 8 - Homo sapiens (Human)

NALP8_HUMAN		R.IGNNKEVQVSFAFCLKR.C	2	2.632	Q86W28
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NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 - Homo sapiens (Human)

NDUA4_HUMAN		R.LALFNPVC#WDR.N	2	2.5406	O00483
NDUA4_HUMAN		R.LALFNPVC#WDR.N	2	2.8725	O00483
NDUA4_HUMAN		K.FYSVNVDSK.L	2	2.314	O00483
NDUA4_HUMAN		R.LALFNPVC#WDR.N	2	2.5737	O00483

NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitochondrial precursor - H

NDUA9_HUMAN		R.LFLPFPLPFAYR.W	2	2.7798	Q16795
NDUA9_HUMAN		R.LFLPFPLPFAYR.W	2	2.5351	Q16795

NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 - Homo sapiens (Human)

NDUBA_HUMAN		R.YQDLGAYSSAR.K	2	2.557	O96000
NDUBA_HUMAN		R.YQDLGAYSSAR.K	2	3.0515	O96000

NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrial precursor - Ho

NDUB8_HUMAN		R.VEDYEPYPDDGMGYDYPK.L	2	2.3137	O95169
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<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>NADH-cytochrome b5 reductase 3 - Homo sapiens (Human)</i>					
NB5R3_HUMAN		K.SVGMIAGGTGITPMLQVIR.A	2	2.3104	P00387
NB5R3_HUMAN		R.GPSGLLVYQGK.G	2	2.3452	P00387
NB5R3_HUMAN		K.SVGMIAGGTGITPMLQVIR.A	2	2.4224	P00387
NB5R3_HUMAN		R.STPAITLESPIKYLPLR.L	2	2.9897	P00387
<i>NADH-ubiquinone oxidoreductase chain 4 - Homo sapiens (Human)</i>					
NU4M_HUMAN		K.LIVPTIMLLPLTWLSK.K	2	3.2721	P03905
NU4M_HUMAN		K.MPLYGLHLWLPK.A	2	2.5434	P03905
<i>NADH-ubiquinone oxidoreductase chain 5 - Homo sapiens (Human)</i>					
NU5M_HUMAN		R.FPTLTNINENNPTLLNPIK.R	2	2.8521	P03915
<i>NEDD9-interacting protein with calponin homology and LIM domains - Homo sapiens (Human)</i>					
MICA1_HUMAN		R.IYNQSFQSLK.A	2	2.8762	Q8TDZ2
<i>Neurolysin, mitochondrial precursor - Homo sapiens (Human)</i>					
NEUL_HUMAN		R.FDIEMSM*RGDIFER.I	2	2.4227	Q9BYT8
<i>Neutrophil cytosol factor 2 - Homo sapiens (Human)</i>					
NCF2_HUMAN		R.IC#FNIGC#MYTILK.N	2	2.6582	P19878
NCF2_HUMAN		R.IHPQQPQEESPPQSDIPAPSSK.A	3	4.7232	P19878
<i>Nuclear factor NF-kappa-B p105 subunit - Homo sapiens (Human)</i>					
NFKB1_HUMAN		R.RLEPVVSDAIYDSK.A	2	2.5701	P19838
<i>Nuclear pore membrane glycoprotein 210 precursor - Homo sapiens (Human)</i>					
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.1591	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	2.678	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.2966	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.326	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	2.4113	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	2.6465	Q8TEM1
PO210_HUMAN		R.VFGAPEVLENLEVK.S	2	2.7087	Q8TEM1
PO210_HUMAN		K.TGAYQYTIR.A	2	2.4443	Q8TEM1
PO210_HUMAN		K.AVDPTSGQLYGLAR.E	2	2.8121	Q8TEM1
PO210_HUMAN		R.GVAIGQTSLTASVTNK.A	2	3.7122	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.133	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.0247	Q8TEM1
PO210_HUMAN		R.LPSQYNFAMNVLGR.V	2	3.3645	Q8TEM1
<i>Peptidyl-prolyl cis-trans isomerase A - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	PPIA_HUMAN	K.SIYGKFEFENFILK.H	2	2.8562	P62937
<i>Peripherin - Homo sapiens (Human)</i>					
	PERI_HUMAN	K.NLQEAEWYK.S	2	2.3227	P41219
<i>Peroxiredoxin-6 - Homo sapiens (Human)</i>					
	PRDX6_HUMAN	K.DINAYNC#EEPTK.L	2	2.5337	P30041
<i>Phosphate carrier protein, mitochondrial precursor - Homo sapiens (Human)</i>					
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.8696	Q00325
	MPCP_HUMAN	K.VLYSNMLGEENTYLWR.T	2	3.3114	Q00325
	MPCP_HUMAN	K.GWAPTFLGYSMQGLC#K.F	2	2.6215	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.4403	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.9696	Q00325
	MPCP_HUMAN	K.GWAPTFLGYSMQGLC#K.F	2	2.4814	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.9319	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	3.0071	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.6159	Q00325
	MPCP_HUMAN	R.IQTQPGYANTLR.D	2	2.6014	Q00325
<i>Plastin-2 - Homo sapiens (Human)</i>					
	PLSL_HUMAN	R.YTLNILEEIGGGQK.V	2	2.7998	P13796
	PLSL_HUMAN	K.LNLAFIANLFNR.Y	2	3.7844	P13796
	PLSL_HUMAN	K.LNLAFIANLFNR.Y	2	3.0525	P13796
	PLSL_HUMAN	R.YTLNILEEIGGGQK.V	2	3.36	P13796
	PLSL_HUMAN	K.LNLAFIANLFNR.Y	2	3.8418	P13796
	PLSL_HUMAN	K.LNLAFIANLFNR.Y	2	3.8814	P13796
	PLSL_HUMAN	K.LNLAFIANLFNR.Y	2	3.4597	P13796
	PLSL_HUMAN	R.VYALPEDLVEVNP.K.M	2	2.396	P13796
	PLSL_HUMAN	R.YTLNILEEIGGGQK.V	2	3.2524	P13796
<i>Platelet glycoprotein 4 - Homo sapiens (Human)</i>					
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	2.9212	P16671
	CD36_HUMAN	R.TYLDIEPITGFTLQFAK.R	2	2.455	P16671
	CD36_HUMAN	R.TYLDIEPITGFTLQFAK.R	2	2.5732	P16671
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	3.1792	P16671
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	2.8482	P16671
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	2.8256	P16671
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	2.6568	P16671
	CD36_HUMAN	R.SIYAVFESDVNLK.G	2	2.3905	P16671

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Poly [ADP-ribose] polymerase 4 - Homo sapiens (Human)</i>					
	PARP4_HUMAN	K.YIIKFSM*PGDQIK.D	2	2.3141	Q9UUKK3
<i>Poly(rC)-binding protein 1 - Homo sapiens (Human)</i>					
	PCBP1_HUMAN	R.INISEGNC#PER.I	2	3.0262	Q15365
	PCBP1_HUMAN	R.IITLTGPTNAIFK.A	2	2.5242	Q15365
	PCBP1_HUMAN	R.IITLTGPTNAIFK.A	2	2.5772	Q15365
	PCBP1_HUMAN	R.INISEGNC#PER.I	2	3.0041	Q15365
	PCBP1_HUMAN	R.IITLTGPTNAIFK.A	2	2.6541	Q15365
	PCBP1_HUMAN	R.IITLTGPTNAIFK.A	2	3.0212	Q15365
	PCBP1_HUMAN	R.INISEGNC#PER.I	2	3.1477	Q15365
	PCBP1_HUMAN	R.ESTGAQVQVAGDMLPNSTER.A	2	3.9682	Q15365
<i>Poly(rC)-binding protein 2 - Homo sapiens (Human)</i>					
	PCBP2_HUMAN	R.IITLAGPTNAIFK.A	2	2.8649	Q15366
	PCBP2_HUMAN	R.IITLAGPTNAIFK.A	2	2.6625	Q15366
	PCBP2_HUMAN	R.IITLAGPTNAIFK.A	2	2.5305	Q15366
	PCBP2_HUMAN	R.IITLAGPTNAIFK.A	2	2.6848	Q15366
<i>Polypyrimidine tract-binding protein 1 - Homo sapiens (Human)</i>					
	PTBP1_HUMAN	K.LSLDGQNIYNAC#C#TLR.I	2	3.7542	P26599
<i>PRA1 family protein 3 - Homo sapiens (Human)</i>					
	PRAF3_HUMAN	R.AWDDFFPGSDR.F	2	2.4809	O75915
	PRAF3_HUMAN	R.AWDDFFPGSDR.F	2	2.5708	O75915
<i>Probable ATP-dependent RNA helicase DDX17 - Homo sapiens (Human)</i>					
	DDX17_HUMAN	R.MLDMGFEPQIR.K	2	2.8503	Q92841
<i>Prohibitin - Homo sapiens (Human)</i>					
	PHB_HUMAN	R.FDAGELITQR.E	2	2.6648	P35232
	PHB_HUMAN	K.DLQNVNITLR.I	2	2.8585	P35232
	PHB_HUMAN	R.IFTSIGEDYDER.V	2	2.4339	P35232
	PHB_HUMAN	R.FDAGELITQR.E	2	2.6686	P35232
	PHB_HUMAN	R.ILFRPVASQLPR.I	2	2.4915	P35232
	PHB_HUMAN	R.KLEAAEDIAYQLSR.S	2	4.0971	P35232
<i>Prohibitin-2 - Homo sapiens (Human)</i>					
	PHB2_HUMAN	K.IVQAEGEAEAAK.M	2	2.5851	Q99623
	PHB2_HUMAN	R.IPWFQYPIIDIR.A	2	2.6232	Q99623
	PHB2_HUMAN	K.IVQAEGEAEAAK.M	2	3.1053	Q99623

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
PHB2_HUMAN		K.FNASQLITQR.A	2	2.9279	Q99623
PHB2_HUMAN		R.IYLTADNLVLNLQDESFTR.G	2	2.5971	Q99623
PHB2_HUMAN		K.IVQAEGEAEAAK.M	2	3.2251	Q99623
PHB2_HUMAN		R.LGLDYEER.V	2	2.3268	Q99623
PHB2_HUMAN		R.IGGVQQDTILAEGLHFR.I	3	4.1001	Q99623
PHB2_HUMAN		K.IVQAEGEAEAAK.M	2	2.7091	Q99623
PHB2_HUMAN		K.FNASQLITQR.A	2	2.7923	Q99623
PHB2_HUMAN		K.FNASQLITQR.A	2	2.9658	Q99623
PHB2_HUMAN		K.LLLGAGAVAYGVR.E	2	2.4611	Q99623
PHB2_HUMAN		R.VLSRPNAQELPSMYQR.L	2	3.2936	Q99623
<i>Proteasome activator complex subunit 2 - Homo sapiens (Human)</i>					
PSME2_HUMAN		R.AFYAELYHISSNLEK.I	2	3.077	Q9UL46
PSME2_HUMAN		R.AFYAELYHISSNLEK.I	2	4.0342	Q9UL46
PSME2_HUMAN		R.AFYAELYHISSNLEK.I	2	2.6221	Q9UL46
<i>Protein FAM62A - Homo sapiens (Human)</i>					
FA62A_HUMAN		R.LLVPLVPDLQDVAQLR.S	2	3.4462	Q9BSJ8
<i>Protein kinase C-binding protein 1 - Homo sapiens (Human)</i>					
PKCB1_HUMAN		R.LIAEVKKQLELEK.Q	2	2.3067	Q9ULU4
<i>Protein LYRIC - Homo sapiens (Human)</i>					
LYRIC_HUMAN		R.TELGLDLGLEPK.R	2	2.6699	Q86UE4
LYRIC_HUMAN		R.SWQDELAQQAEEGSAR.L	2	3.4474	Q86UE4
<i>Protein S100-A8 - Homo sapiens (Human)</i>					
S10A8_HUMAN		K.LLETEC#PQYIR.K	2	2.7924	P05109
S10A8_HUMAN		K.LLETEC#PQYIR.K	2	2.5517	P05109
S10A8_HUMAN		K.LLETEC#PQYIR.K	2	2.8931	P05109
S10A8_HUMAN		K.LLETEC#PQYIR.K	2	2.4833	P05109
S10A8_HUMAN		K.LLETEC#PQYIR.K	2	2.6589	P05109
<i>Protein S100-A9 - Homo sapiens (Human)</i>					
S10A9_HUMAN		R.NIETIINTFHQYSVK.L	2	3.1467	P06702
S10A9_HUMAN		R.NIETIINTFHQYSVK.L	2	3.428	P06702
S10A9_HUMAN		R.NIETIINTFHQYSVK.L	2	2.9079	P06702
<i>Protein transport protein Sec61 subunit alpha isoform 1 - Homo sapiens (Human)</i>					
S61A1_HUMAN		R.GMEFEGAIILFHLLATR.T	3	4.0181	P61619
S61A1_HUMAN		R.GMEFEGAIILFHLLATR.T	2	3.9445	P61619

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Proteolipid protein 2 - Homo sapiens (Human)</i>					
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.6251	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.3578	Q04941
	PLP2_HUMAN	R.HTAAPTDPADGPV.-	2	2.5996	Q04941
<i>Putative uncharacterized protein - Homo sapiens (Human)</i>					
	A5D8X1_HUMAN	R.FIAVGYVDDTEFVR.F	2	4.0161	A5D8X1
	A5D8X1_HUMAN	R.FIAVGYVDDTEFVR.F	2	2.8839	A5D8X1
	A5D8X1_HUMAN	R.FIAVGYVDDTEFVR.F	2	2.6639	A5D8X1
<i>Pyruvate kinase isozymes M1/M2 - Homo sapiens (Human)</i>					
	KPYM_HUMAN	K.GADFLVTEVENGGSLGSK.K	2	2.5362	P14618
<i>Rab11 family-interacting protein 4 - Homo sapiens (Human)</i>					
	RFIP4_HUMAN	K.KVTELENDSLTNGDLK.S	2	2.7758	Q86YS3
<i>Ras GTPase-activating-like protein IQGAP1 - Homo sapiens (Human)</i>					
	IQGA1_HUMAN	R.SNQQLENDLNLMDIK.I	2	2.8679	P46940
	IQGA1_HUMAN	K.LTELGTVDPK.N	2	2.3317	P46940
	IQGA1_HUMAN	R.SNQQLENDLNLMDIK.I	2	2.9638	P46940
	IQGA1_HUMAN	K.VNVNLLIFLLNK.K	2	3.103	P46940
	IQGA1_HUMAN	K.SWVNQMESQTGEASK.L	2	3.1883	P46940
	IQGA1_HUMAN	K.SKVDQIQEIVTGNPTVIK.M	2	3.0469	P46940
<i>Ras GTPase-activating-like protein IQGAP2 - Homo sapiens (Human)</i>					
	IQGA2_HUMAN	K.LPYDVTTEQUALTYPEVK.N	2	3.4655	Q13576
<i>Ras-related C3 botulinum toxin substrate 1 precursor - Homo sapiens (Human)</i>					
	RAC1_HUMAN	K.YLEC#SALTQR.G	2	2.4444	P63000
	RAC1_HUMAN	K.YLEC#SALTQR.G	2	2.5506	P63000
	RAC1_HUMAN	K.YLEC#SALTQR.G	2	2.4905	P63000
	RAC1_HUMAN	K.YLEC#SALTQR.G	2	2.3411	P63000
<i>Ras-related protein Rab-10 - Homo sapiens (Human)</i>					
	RAB10_HUMAN	K.AFLTLAEDILR.K	2	2.3268	P61026
	RAB10_HUMAN	K.AFLTLAEDILR.K	2	2.7012	P61026
<i>Ras-related protein Rab-3D - Homo sapiens (Human)</i>					
	RAB3D_HUMAN	R.LADDLGFEFFFEASAK.E	2	3.2682	O95716
	RAB3D_HUMAN	R.LADDLGFEFFFEASAK.E	2	3.7809	O95716
	RAB3D_HUMAN	R.LADDLGFEFFFEASAK.E	2	2.802	O95716
	RAB3D_HUMAN	R.LADDLGFEFFFEASAK.E	2	3.5789	O95716

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Ras-related protein Rab-5C - Homo sapiens (Human)</i>					
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	2.7961	P51148
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	3.2409	P51148
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	3.1379	P51148
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	3.3823	P51148
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	2.9479	P51148
RAB5C_HUMAN		R.GVDLQENNPASR.S	2	3.2858	P51148
<i>Ras-related protein Rap-1A precursor - Homo sapiens (Human)</i>					
RAP1A_HUMAN		K.INVNEIFYDLVR.Q	2	4.1354	P62834
RAP1A_HUMAN		K.LVVLGSGGVGK.S	2	2.5519	P62834
RAP1A_HUMAN		K.INVNEIFYDLVR.Q	2	3.7014	P62834
RAP1A_HUMAN		K.SKINVNEIFYDLVR.Q	2	2.7059	P62834
RAP1A_HUMAN		K.INVNEIFYDLVR.Q	2	4.238	P62834
<i>Receptor expression-enhancing protein 5 - Homo sapiens (Human)</i>					
REEP5_HUMAN		K.HESQMDSVVK.D	2	2.9746	Q00765
REEP5_HUMAN		K.HESQMDSVVK.D	2	2.5697	Q00765
REEP5_HUMAN		K.HESQMDSVVK.D	2	3.0614	Q00765
<i>Renin receptor precursor - Homo sapiens (Human)</i>					
RENH_HUMAN		K.NPASPYNLAYK.Y	2	2.4263	O75787
<i>RER1 retention in endoplasmic reticulum 1 homolog - Homo sapiens (Human)</i>					
Q5T091_HUMAN		K.VDPSLMEDSDDGPSLPTK.Q	2	3.5476	Q5T091
Q5T091_HUMAN		K.VDPSLMEDSDDGPSLPTK.Q	2	3.6236	Q5T091
Q5T091_HUMAN		K.VDPSLM*EDSDDGPSLPTK.Q	2	2.9471	Q5T091
Q5T091_HUMAN		K.VDPSLM*EDSDDGPSLPTK.Q	2	2.4101	Q5T091
Q5T091_HUMAN		K.VDPSLMEDSDDGPSLPTK.Q	2	3.2652	Q5T091
<i>Reticulon-3 - Homo sapiens (Human)</i>					
RTN3_HUMAN		K.TQIDHYVGIAR.D	2	3.1943	O95197
RTN3_HUMAN		K.TQIDHYVGIAR.D	2	3.2557	O95197
<i>Reticulon-4 - Homo sapiens (Human)</i>					
RTN4_HUMAN		R.HQAQIDHYLGLANK.N	3	4.0591	Q9NQC3
RTN4_HUMAN		R.HQAQIDHYLGLANK.N	2	3.877	Q9NQC3
RTN4_HUMAN		R.HQAQIDHYLGLANK.N	3	4.0036	Q9NQC3
RTN4_HUMAN		R.HQAQIDHYLGLANK.N	3	4.0548	Q9NQC3
RTN4_HUMAN		R.HQAQIDHYLGLANK.N	2	3.4418	Q9NQC3
RTN4_HUMAN		K.YSNSALGHVNC#TIK.E	2	3.7896	Q9NQC3

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	RTN4_HUMAN	R.HQAQIDHYLGLANK.N	2	3.6937	Q9NQC3
<i>Rho GTPase-activating protein 1 - Homo sapiens (Human)</i>					
	RHG01_HUMAN	K.FLLDHQGELEFPSPDPSGL.-	2	2.7511	Q07960
<i>Rho guanine nucleotide exchange factor 1 - Homo sapiens (Human)</i>					
	ARHG1_HUMAN	R.LRPLLSQLGGNSVPQPGC#T.-	2	2.4585	Q92888
<i>Rho-related GTP-binding protein RhoG precursor - Homo sapiens (Human)</i>					
	RHOG_HUMAN	R.LKEQGQAPITPQQGQALAK.Q	3	4.1111	P84095
	RHOG_HUMAN	K.EQGQAPITPQQGQALAK.Q	2	2.4269	P84095
	RHOG_HUMAN	K.EQGQAPITPQQGQALAK.Q	2	2.5243	P84095
<i>Ribonuclease inhibitor - Homo sapiens (Human)</i>					
	RINI_HUMAN	R.ELC#QGLGQPGSVLR.V	2	2.5383	P13489
	RINI_HUMAN	R.ELC#QGLGQPGSVLR.V	2	2.7461	P13489
<i>RNA adenosine deaminase - Homo sapiens (Human)</i>					
	Q86XN4_HUMAN	R.FLYAQLELYLNNK.E	2	2.3876	Q86XN4
	Q86XN4_HUMAN	R.FLYAQLELYLNNK.E	2	2.3589	Q86XN4
<i>RRP5 protein homolog - Homo sapiens (Human)</i>					
	RRP5_HUMAN	K.KGKAINIGQLVDVK.V	2	2.4508	Q14690
<i>SAC domain-containing protein 3 - Homo sapiens (Human)</i>					
	SAC3_HUMAN	-.M*PTAAAPIISSVQK.L	2	2.3636	Q92562
<i>SAM domain and HD domain-containing protein 1 - Homo sapiens (Human)</i>					
	SAMH1_HUMAN	K.EWNDSTSVQNPTR.L	2	3.0779	Q9Y3Z3
	SAMH1_HUMAN	R.NFTKPQDGDVIAPLITPQK.K	2	2.4901	Q9Y3Z3
	SAMH1_HUMAN	K.VGNIIDTMITDAFLK.A	2	3.2595	Q9Y3Z3
	SAMH1_HUMAN	K.EWNDSTSVQNPTR.L	2	2.9805	Q9Y3Z3
	SAMH1_HUMAN	K.VGNIIDTMITDAFLK.A	2	3.6995	Q9Y3Z3
	SAMH1_HUMAN	K.YVGETQPTGQIK.I	2	2.4376	Q9Y3Z3
	SAMH1_HUMAN	K.EWNDSTSVQNPTR.L	2	3.1863	Q9Y3Z3
	SAMH1_HUMAN	K.EWNDSTSVQNPTR.L	2	2.8884	Q9Y3Z3
<i>Secretory carrier-associated membrane protein 3 - Homo sapiens (Human)</i>					
	SCAM3_HUMAN	K.AQQEFAAGVFSNPAVR.T	2	2.7414	O14828
<i>Sialic acid-binding Ig-like lectin 7 precursor - Homo sapiens (Human)</i>					
	SIGL7_HUMAN	K.APVATNPAWAVQEETR.D	2	4.0756	Q9Y286
<i>Sialic acid-binding Ig-like lectin 9 precursor - Homo sapiens (Human)</i>					
	SIGL9_HUMAN	R.PAAGVGDTGIEDANAVR.G	2	3.5569	Q9Y336

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A		2	3.324	Q9Y336
SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A		2	2.5592	Q9Y336
SIGL9_HUMAN	R.EGANTDQDAPVATNNPAR.A		2	2.7113	Q9Y336
SIGL9_HUMAN	R.PAAGVGDGTGIEDANAVR.G		2	3.495	Q9Y336
<i>Sideroflexin-1 - Homo sapiens (Human)</i>					
SFXN1_HUMAN	R.NILLTNEQLESAR.K		2	3.0843	Q9H9B4
<i>Signal peptidase complex catalytic subunit SEC11A - Homo sapiens (Human)</i>					
SC11A_HUMAN	R.GFVPYIGIVTILMNDYPK.F		2	2.7355	P67812
<i>Signal recognition particle receptor subunit beta - Homo sapiens (Human)</i>					
SRPRB_HUMAN	R.SAAPSTLDSSTAPAQLGK.K		2	2.5044	Q9Y5M8
SRPRB_HUMAN	R.SAAPSTLDSSTAPAQLGK.K		2	3.0953	Q9Y5M8
SRPRB_HUMAN	R.AIVFVVDAAAFQR.E		2	2.401	Q9Y5M8
SRPRB_HUMAN	R.SAAPSTLDSSTAPAQLGK.K		2	4.2563	Q9Y5M8
<i>Similar to ribosomal protein L18; 60S ribosomal protein L18 - Homo sapiens (Human)</i>					
A4D0W6_HUMAN	K.NLTFDQLALDSPK.V		2	3.069	A4D0W6
<i>Sodium/potassium-transporting ATPase subunit alpha-1 precursor - Homo sapiens (Human)</i>					
AT1A1_HUMAN	K.VDNSSLTGESEPQTR.S		2	2.8133	P05023
AT1A1_HUMAN	K.GVGIISEGNETVEDIAAR.L		2	3.2532	P05023
<i>Sodium/potassium-transporting ATPase subunit beta-3 - Homo sapiens (Human)</i>					
AT1B3_HUMAN	K.LFIYNPTTGEFLGR.T		2	3.4942	P54709
AT1B3_HUMAN	K.LFIYNPTTGEFLGR.T		2	2.8453	P54709
AT1B3_HUMAN	K.LFIYNPTTGEFLGR.T		2	3.0578	P54709
AT1B3_HUMAN	K.LFIYNPTTGEFLGR.T		2	2.5496	P54709
<i>Solute carrier family 22 member 11 - Homo sapiens (Human)</i>					
OAT4_HUMAN	R.INGHKEAKNLTIEVLMSSVK.E		2	2.383	Q9NSA0
<i>Solute carrier family 25 member 24. - Homo sapiens (Human)</i>					
Q6NUK1_HUMAN	R.QLLAGGIAGAVSR.T		2	3.8477	Q6NUK1
Q6NUK1_HUMAN	R.QLLAGGIAGAVSR.T		2	2.6163	Q6NUK1
Q6NUK1_HUMAN	R.QLLAGGIAGAVSR.T		2	3.0839	Q6NUK1
<i>Sorting nexin-7 - Homo sapiens (Human)</i>					
SNX7_HUMAN	R.MSGLSEALLPVVHEYVLYSEM*LMGVMKR.R		3	3.7526	Q9UNH6
<i>Structural maintenance of chromosomes protein 3 - Homo sapiens (Human)</i>					
SMC3_HUMAN	K.KM*VTKNDVMNLLSAGFSR.S		2	2.4367	Q9UQE7
<i>Sulfide:quinone oxidoreductase, mitochondrial precursor - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	SQRD_HUMAN	K.TAAAVAAQSGILDR.T	2	3.8223	Q9Y6N5
	SQRD_HUMAN	K.ANIIFNTSLGAIFGVK.K	2	2.4241	Q9Y6N5
	SQRD_HUMAN	K.ADLMPFLYWNMLLR.G	2	3.2642	Q9Y6N5
	SQRD_HUMAN	K.TAAAVAAQSGILDR.T	2	3.3695	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	2.959	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	2.5298	Q9Y6N5
	SQRD_HUMAN	K.IMYLSEAYFR.K	2	2.4918	Q9Y6N5
	SQRD_HUMAN	K.YADALQEIIQER.N	2	3.3993	Q9Y6N5
	SQRD_HUMAN	K.VGAENVAIVEPSE.H	2	2.9343	Q9Y6N5
	SQRD_HUMAN	K.TAAAVAAQSGILDR.T	2	3.0484	Q9Y6N5
	SQRD_HUMAN	K.YADALQEIIQER.N	2	3.6877	Q9Y6N5
	SQRD_HUMAN	K.YADALQEIIQER.N	2	2.4687	Q9Y6N5
	SQRD_HUMAN	K.YADALQEIIQER.N	2	3.3043	Q9Y6N5
	SQRD_HUMAN	R.KVGAENVAIVEPSE.H	2	2.3123	Q9Y6N5
	SQRD_HUMAN	R.KVGAENVAIVEPSE.H	2	2.3346	Q9Y6N5
	SQRD_HUMAN	K.EGNAIFTFPNTPVK.C	2	2.7891	Q9Y6N5
	SQRD_HUMAN	R.HFYQPIWTLVGAGAK.Q	2	2.8874	Q9Y6N5
	SQRD_HUMAN	R.HFYQPIWTLVGAGAK.Q	2	2.9018	Q9Y6N5
<i>Surfeit locus protein 1 - Homo sapiens (Human)</i>					
	SURF1_HUMAN	K.LNLIAELESR.V	2	2.5107	Q15526
<i>Synaptogyrin-1 - Homo sapiens (Human)</i>					
	SNG1_HUMAN	K.AGGAFDPYTLVR.Q	2	2.3263	O43759
<i>Synaptogyrin-2 - Homo sapiens (Human)</i>					
	SNG2_HUMAN	K.DVLVGADSVR.A	2	2.364	O43760
<i>Synaptophysin-like protein 1 - Homo sapiens (Human)</i>					
	SYPL1_HUMAN	K.TVTATFGYPFR.L	2	2.3664	Q16563
<i>Syntaxin-7 - Homo sapiens (Human)</i>					
	STX7_HUMAN	K.ITQC#SVEIQR.T	2	2.4811	O15400
	STX7_HUMAN	R.LVAEFTTSLTNFQK.V	2	2.6768	O15400
	STX7_HUMAN	K.ITQC#SVEIQR.T	2	2.7161	O15400
	STX7_HUMAN	R.TLNQLGTPQDSPELR.Q	2	2.3593	O15400
	STX7_HUMAN	R.LVAEFTTSLTNFQK.V	2	2.508	O15400
	STX7_HUMAN	R.LVAEFTTSLTNFQK.V	2	3.1107	O15400
<i>Tapasin precursor - Homo sapiens (Human)</i>					
	TPSN_HUMAN	K.WASGLTPAQNC#PR.A	2	2.8969	O15533

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	TPSN_HUMAN	K.WASGLTPAQNC#PR.A	2	2.9556	O15533
	TPSN_HUMAN	K.WASGLTPAQNC#PR.A	2	3.3483	O15533
	TPSN_HUMAN	K.WASGLTPAQNC#PR.A	2	2.8735	O15533
<i>T-cell surface glycoprotein CD1a precursor - Homo sapiens (Human)</i>					
	CD1A_HUMAN	R.FILGLLDAGKAHLQR.Q	2	2.9632	P06126
<i>T-complex protein 1 subunit delta - Homo sapiens (Human)</i>					
	TCPD_HUMAN	K.IGLIQFC#LSAPK.T	2	2.7222	P50991
	TCPD_HUMAN	R.DALSDLALHFLNK.M	2	3.3397	P50991
<i>Thromboxane A synthase 1 - Homo sapiens (Human)</i>					
	Q53F23_HUMAN	R.FQAC#PETQVPLQLESK.S	2	2.8436	Q53F23
<i>Thromboxane synthase - Homo sapiens (Human)</i>					
	Q16843_HUMAN	R.YAESGDAFDIQR.C	2	3.5721	Q16843
	Q16843_HUMAN	R.YAESGDAFDIQR.C	2	2.3701	Q16843
	Q16843_HUMAN	K.PSPFIGNLTFFR.Q	2	2.6019	Q16843
	Q16843_HUMAN	R.MFIVISEPDMIK.Q	2	2.3918	Q16843
<i>Translocator protein - Homo sapiens (Human)</i>					
	TSPO_HUMAN	R.FVHGEGLR.W	2	2.4388	P30536
	TSPO_HUMAN	K.AVVPLGLYTGQLALNWAWPPIFFGAR.Q	2	2.9049	P30536
<i>Translocon-associated protein subunit alpha precursor - Homo sapiens (Human)</i>					
	SSRA_HUMAN	K.GEDFPANNIVK.F	2	2.5849	P43307
<i>Translocon-associated protein subunit delta precursor - Homo sapiens (Human)</i>					
	SSRD_HUMAN	R.VQNMALYADVGGK.Q	2	2.7726	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	2.9253	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	3.0536	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	3.0703	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	3.1083	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	3.0943	P51571
	SSRD_HUMAN	R.VQNMALYADVGGK.Q	2	3.3553	P51571
	SSRD_HUMAN	R.FFDEESYSLLR.K	2	2.4746	P51571
<i>Translocon-associated protein subunit gamma - Homo sapiens (Human)</i>					
	SSRG_HUMAN	K.QQSEEDLLLQDFSR.N	2	2.5379	Q9UNL2
	SSRG_HUMAN	K.QQSEEDLLLQDFSR.N	2	2.9348	Q9UNL2
	SSRG_HUMAN	K.QQSEEDLLLQDFSR.N	2	2.41	Q9UNL2
<i>Transmembrane 9 superfamily protein member 3 precursor - Homo sapiens (Human)</i>					

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	TM9S3_HUMAN	R.NLSGQPNFPC#R.V	2	2.7794	Q9HD45
	TM9S3_HUMAN	R.NLSGQPNFPC#R.V	2	2.7978	Q9HD45
	TM9S3_HUMAN	R.NLSGQPNFPC#R.V	2	2.558	Q9HD45
<i>Transmembrane and coiled-coil domain-containing protein 1 - Homo sapiens (Human)</i>					
	TMCO1_HUMAN	K.LPFTPLSYIQGLSHR.N	3	4.2222	Q9UM00
<i>Transmembrane emp24 domain-containing protein 1 precursor - Homo sapiens (Human)</i>					
	TMED1_HUMAN	R.SIQMLTLLR.A	2	2.3077	Q13445
<i>Transmembrane emp24 domain-containing protein 10 precursor - Homo sapiens (Human)</i>					
	TMEDA_HUMAN	R.IPDQLVILDMK.H	2	2.9262	P49755
	TMEDA_HUMAN	R.IPDQLVILDMK.H	2	2.8589	P49755
	TMEDA_HUMAN	R.IPDQLVILDMK.H	2	2.5427	P49755
	TMEDA_HUMAN	R.RLEDLSESIVNDFAYMK.K	2	2.4268	P49755
	TMEDA_HUMAN	R.IPDQLVILDMK.H	2	2.4224	P49755
	TMEDA_HUMAN	R.RLEDLSESIVNDFAYMK.K	2	4.2595	P49755
	TMEDA_HUMAN	R.RLEDLSESIVNDFAYMK.K	2	3.7101	P49755
<i>Transmembrane emp24 domain-containing protein 2 precursor - Homo sapiens (Human)</i>					
	TMED2_HUMAN	K.IVMFTIDIGEAPK.G	2	2.733	Q15363
	TMED2_HUMAN	K.IVMFTIDIGEAPK.G	2	2.5482	Q15363
<i>Transmembrane emp24 domain-containing protein 9 precursor - Homo sapiens (Human)</i>					
	TMED9_HUMAN	K.DKLSSELQLR.V	2	2.315	Q9BVK6
	TMED9_HUMAN	R.QLVEQVEIQK.E	2	2.8735	Q9BVK6
	TMED9_HUMAN	R.QLVEQVEIQK.E	2	3.0789	Q9BVK6
<i>Transmembrane protein 109 precursor - Homo sapiens (Human)</i>					
	TM109_HUMAN	R.EAPVDVLTQIGR.S	2	2.505	Q9BVC6
	TM109_HUMAN	R.EAPVDVLTQIGR.S	2	2.8243	Q9BVC6
<i>Transmembrane protein 167 precursor - Homo sapiens (Human)</i>					
	TM167_HUMAN	K.TGLLGIFWK.C	2	2.3385	Q8TBQ9
<i>Transmembrane protein 16E - Homo sapiens (Human)</i>					
	TM16E_HUMAN	K.FLLAWMIPDVPK.D	2	2.6298	Q75V66
	TM16E_HUMAN	K.FLLAWMIPDVPK.D	2	2.3513	Q75V66
<i>Transmembrane protein 33 - Homo sapiens (Human)</i>					
	TMM33_HUMAN	R.ALLANALTSALR.L	2	3.5783	P57088
	TMM33_HUMAN	R.ALLANALTSALR.L	2	3.8143	P57088
	TMM33_HUMAN	R.ALLANALTSALR.L	2	3.1314	P57088

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	TMM33_HUMAN	R.SVLDKLSANQQNILK.F	2	4.1993	P57088
	TMM33_HUMAN	R.SVLDKLSANQQNILK.F	2	2.6638	P57088
<i>Transmembrane protein 43 - Homo sapiens (Human)</i>					
	TMM43_HUMAN	R.FFLSSGLIDKVDNFK.S	2	3.3431	Q9BTV4
<i>Transporter 1, ATP-binding cassette, sub-family B - Homo sapiens (Human)</i>					
	Q96CP4_HUMAN	R.IFSLLVPTALPLLR.V	2	2.34	Q96CP4
<i>Trifunctional enzyme subunit alpha, mitochondrial precursor - Homo sapiens (Human)</i>					
	ECHA_HUMAN	K.TGIEQGS DAGYLC#ESQK.F	2	2.6909	P40939
	ECHA_HUMAN	K.TLQEVTQLS QEAQR.I	2	2.4831	P40939
	ECHA_HUMAN	R.FGGGNPELLTQM VSK.G	2	2.7753	P40939
	ECHA_HUMAN	R.FGGGNPELLTQM VSK.G	2	2.5008	P40939
<i>Tubulin alpha-1A chain - Homo sapiens (Human)</i>					
	TBA1A_HUMAN	K.DVNAAIATIK.T	2	2.8774	Q71U36
	TBA1A_HUMAN	K.TIGGGD DSFN TFFSETGAGK.H	2	3.0804	Q71U36
	TBA1A_HUMAN	K.TIGGGD DSFN TFFSETGAGK.H	2	4.2575	Q71U36
	TBA1A_HUMAN	R.GHYTIGKEIIDLVLDR.I	2	3.0361	Q71U36
	TBA1A_HUMAN	K.LADQC#TGLQGFLVFH SFGGGTGSGFTSLLMER.L	3	4.2029	Q71U36
	TBA1A_HUMAN	R.GHYTIGKEIIDLVLDR.I	2	3.6793	Q71U36
	TBA1A_HUMAN	K.DVNAAIATIK.T	2	2.5342	Q71U36
	TBA1A_HUMAN	R.NLDIERPTYTNLNR.L	2	2.5339	Q71U36
	TBA1A_HUMAN	R.LDHKFDL MYAK.R	3	4.1124	Q71U36
	TBA1A_HUMAN	K.EIIDLVLDR.I	1	2.1598	Q71U36
	TBA1A_HUMAN	R.NLDIERPTYTNLNR.L	2	2.5406	Q71U36
	TBA1A_HUMAN	R.NLDIERPTYTNLNR.L	2	2.6403	Q71U36
	TBA1A_HUMAN	R.GHYTIGKEIIDLVLDR.I	2	2.4964	Q71U36
	TBA1A_HUMAN	K.LADQC#TGLQGFLVFH SFGGGTGSGFTSLLMER.L	3	5.7302	Q71U36
	TBA1A_HUMAN	K.LADQC#TGLQGFLVFH SFGGGTGSGFTSLLMER.L	3	5.7818	Q71U36
<i>Tubulin alpha-1B chain - Homo sapiens (Human)</i>					
	TBA1B_HUMAN	R.SIQFVDWC#PTGFK.V	2	3.2419	P68363
	TBA1B_HUMAN	R.SIQFVDWC#PTGFK.V	2	2.5547	P68363
	TBA1B_HUMAN	R.LISQIVSSITASLR.F	2	2.9277	P68363
	TBA1B_HUMAN	R.LISQIVSSITASLR.F	2	3.1178	P68363
	TBA1B_HUMAN	R.LISQIVSSITASLR.F	2	2.4457	P68363
	TBA1B_HUMAN	R.SIQFVDWC#PTGFK.V	2	2.8995	P68363
	TBA1B_HUMAN	R.LISQIVSSITASLR.F	2	3.6917	P68363

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>Tubulin beta chain - Homo sapiens (Human)</i>					
TBB5_HUMAN		K.MAVTFIGNSTAIQELFKR.I	2	3.9085	P07437
TBB5_HUMAN		R.ALTVPELTQQVFDAK.N	2	2.3794	P07437
<i>Tubulin beta-1 chain - Homo sapiens (Human)</i>					
TBB1_HUMAN		K.LAVNMVPPFR.L	2	3.5196	Q9H4B7
TBB1_HUMAN		K.LAVNMVPPFR.L	2	3.2563	Q9H4B7
TBB1_HUMAN		R.FPGQLNADLR.K	2	2.6215	Q9H4B7
TBB1_HUMAN		K.LAVNM*VPPFR.L	2	2.3953	Q9H4B7
TBB1_HUMAN		K.LAVNMVPPFR.L	2	3.0207	Q9H4B7
TBB1_HUMAN		K.LAVNMVPPFR.L	2	3.38	Q9H4B7
<i>Tubulin beta-2A chain - Homo sapiens (Human)</i>					
TBB2A_HUMAN		K.NMMAAC#DPR.H	2	2.5448	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.5294	Q13885
TBB2A_HUMAN		R.ISEQFTAM*FR.R	2	2.9367	Q13885
TBB2A_HUMAN		R.ISEQFTAMFR.R	2	2.9926	Q13885
TBB2A_HUMAN		R.LHFFM*PGFAPLTSR.G	2	2.9558	Q13885
TBB2A_HUMAN		K.NSSYFVEWIPNNVK.T	2	2.7399	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	3	3.8886	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	3.0999	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.3251	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	2.8856	Q13885
TBB2A_HUMAN		R.ISEQFTAM*FR.R	2	2.7492	Q13885
TBB2A_HUMAN		R.ISEQFTAMFR.R	2	2.7948	Q13885
TBB2A_HUMAN		K.NSSYFVEWIPNNVK.T	2	4.0956	Q13885
TBB2A_HUMAN		K.NSSYFVEWIPNNVK.T	2	2.5243	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	2.7723	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.4385	Q13885
TBB2A_HUMAN		R.ISEQFTAM*FR.R	2	2.7082	Q13885
TBB2A_HUMAN		R.ISEQFTAMFR.R	2	3.2069	Q13885
TBB2A_HUMAN		K.NSSYFVEWIPNNVK.T	2	3.3443	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	2.8836	Q13885
TBB2A_HUMAN		K.NMMAAC#DPR.H	2	2.3165	Q13885
TBB2A_HUMAN		K.NMMAAC#DPR.H	2	2.5119	Q13885
TBB2A_HUMAN		K.EVDEQMLNVQNK.N	2	3.1691	Q13885
TBB2A_HUMAN		R.ISEQFTAM*FR.R	2	2.7247	Q13885

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
TBB2A_HUMAN		R.ISEQFTAMFR.R	2	2.4334	Q13885
TBB2A_HUMAN		K.NSSYFVEWIPNNVK.T	2	3.5737	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	2.5683	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	4.0923	Q13885
TBB2A_HUMAN		R.LHFFMPGFAPLTSR.G	2	2.6894	Q13885
<i>Tubulin beta-2C chain - Homo sapiens (Human)</i>					
TBB2C_HUMAN		R.IMNTFSVVPSPK.V	2	2.8355	P68371
TBB2C_HUMAN		R.YLTVAAVFR.G	2	2.3067	P68371
TBB2C_HUMAN		R.YLTVAAVFR.G	2	2.346	P68371
TBB2C_HUMAN		R.ALTVPELTQQMFDAK.N	2	2.38	P68371
TBB2C_HUMAN		R.YLTVAAVFR.G	2	2.9921	P68371
<i>Tubulin beta-3 chain - Homo sapiens (Human)</i>					
TBB3_HUMAN		K.ECENCDC#LQGFQLTHSLGGGTGSGMGTLISK.V	3	4.5146	Q13509
<i>Type-1 angiotensin II receptor-associated protein - Homo sapiens (Human)</i>					
ATRAP_HUMAN		R.SAYQTIDSAEAPADPFVPEGR.S	2	2.7816	Q6RW13
<i>Tyrosine-protein kinase HCK - Homo sapiens (Human)</i>					
HCK_HUMAN		R.VIEDNEYTAR.E	2	2.3339	P08631
<i>Tyrosine-protein kinase SYK - Homo sapiens (Human)</i>					
KSYK_HUMAN		K.KFDTLWQLVEHYSYK.A	3	3.941	P43405
<i>Tyrosine-protein phosphatase non-receptor type 6 - Homo sapiens (Human)</i>					
PTN6_HUMAN		K.GLDC#DIDIQK.T	2	2.6299	P29350
<i>Ubiquinol-cytochrome-c reductase complex core protein 2, mitochondrial precursor - Homo sap</i>					
UQCR2_HUMAN		K.TIAQGNLSNTDVQAAK.N	2	3.0025	P22695
<i>Ubiquitin - Homo sapiens (Human)</i>					
UBIQ_HUMAN		K.ESTLHLVLR.L	2	2.7939	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	3.565	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	3.854	P62988
UBIQ_HUMAN		K.TITLEVEPSDTIENVK.A	2	2.9731	P62988
<i>Unc-112-related protein 2 - Homo sapiens (Human)</i>					
URP2_HUMAN		R.TMADSSYTSEVQAILAFLSLQR.T	2	3.4665	Q86UX7
URP2_HUMAN		R.TGSGGPGNHPHPDASAEGLNPYGLVAPR.F	3	3.9501	Q86UX7
URP2_HUMAN		R.LTQLYEQAR.W	2	2.5207	Q86UX7
URP2_HUMAN		K.SQDEAPGDPIQQLNLK.G	2	2.8915	Q86UX7
URP2_HUMAN		R.TMADSSYTSEVQAILAFLSLQR.T	2	2.7158	Q86UX7

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
<i>UNC93 homolog B1 - Homo sapiens (Human)</i>					
	UN93B_HUMAN	R.SVGWGNIFQLPFK.H	2	2.3946	Q9H1C4
	UN93B_HUMAN	R.RPC#PYEQAQGGDGPEEQ.-	2	3.0312	Q9H1C4
<i>Uncharacterized protein C12orf51 - Homo sapiens (Human)</i>					
	CL051_HUMAN	K.LVLTEGERNSGLSQLR.D	2	2.3799	Q3MJD5
<i>UPF0404 protein C11orf59 - Homo sapiens (Human)</i>					
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	3.8165	Q6IAA8
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	2.3937	Q6IAA8
	CK059_HUMAN	R.TDEQALLSSILAK.T	2	2.9718	Q6IAA8
<i>Vacuolar ATP synthase 16 kDa proteolipid subunit - Homo sapiens (Human)</i>					
	VATL_HUMAN	K.SGTGIAAMSVMRPEQIMK.S	2	3.6451	P27449
	VATL_HUMAN	K.SGTGIAAMSVMRPEQIMK.S	2	3.0026	P27449
	VATL_HUMAN	K.SGTGIAAMSVMRPEQIMK.S	2	2.7081	P27449
<i>Vacuolar proton translocating ATPase 116 kDa subunit a isoform 1 - Homo sapiens (Human)</i>					
	VPP1_HUMAN	R.ASLYPC#PETPQER.K	2	2.3075	Q93050
<i>Vesicle-associated membrane protein 8 - Homo sapiens (Human)</i>					
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	3.7696	Q9BV40
	VAMP8_HUMAN	R.NKTEDLEATSEHFK.T	3	4.1858	Q9BV40
<i>Vesicle-fusing ATPase - Homo sapiens (Human)</i>					
	NSF_HUMAN	R.FSNLVLQALLVLLK.K	2	3.6152	P46459
	NSF_HUMAN	R.FSNLVLQALLVLLK.K	3	3.8939	P46459
	NSF_HUMAN	R.FSNLVLQALLVLLK.K	2	3.6699	P46459
<i>Vesicle-trafficking protein SEC22b - Homo sapiens (Human)</i>					
	SC22B_HUMAN	K.ANNLSSLK.K	2	2.3345	O75396
	SC22B_HUMAN	R.GEALSALDSK.A	2	2.7533	O75396
	SC22B_HUMAN	R.NLGSINTELQDVQR.I	2	3.0841	O75396
<i>Vesicular integral-membrane protein VIP36 precursor - Homo sapiens (Human)</i>					
	LMAN2_HUMAN	K.DNVDDPTGNFR.S	2	3.0461	Q12907
	LMAN2_HUMAN	K.DNVDDPTGNFR.S	2	2.7499	Q12907
	LMAN2_HUMAN	K.DNVDDPTGNFR.S	2	3.0687	Q12907
	LMAN2_HUMAN	K.DNVDDPTGNFR.S	2	2.8584	Q12907
<i>Vimentin - Homo sapiens (Human)</i>					
	VIME_HUMAN	K.FADLSEAANR.N	2	3.5557	P08670
	VIME_HUMAN	R.SLYASSPGGVYATR.S	2	3.2086	P08670

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
VIME_HUMAN		K.FADLSEAANR.N	2	2.6552	P08670
VIME_HUMAN		R.SLYASSPGGVYATR.S	2	2.9691	P08670
VIME_HUMAN		R.QVQSLTC#EVDALK.G	2	2.8193	P08670
VIME_HUMAN		R.QDVDNASLAR.L	2	2.5796	P08670
VIME_HUMAN		K.FADLSEAANR.N	2	3.5152	P08670
VIME_HUMAN		R.SLYASSPGGVYATR.S	2	3.1671	P08670
VIME_HUMAN		R.SLYASSPGGVYATR.S	2	2.6251	P08670
VIME_HUMAN		R.LGDLYEEEMR.E	2	2.4666	P08670
VIME_HUMAN		K.FADLSEAANR.N	2	2.739	P08670
VIME_HUMAN		R.SLYASSPGGVYATR.S	2	3.1368	P08670
VIME_HUMAN		R.LGDLYEEEMR.E	2	2.3076	P08670
VIME_HUMAN		R.QVQSLTC#EVDALK.G	2	2.9191	P08670
VIME_HUMAN		R.TYSLGSALRPSTSR.S	2	2.3714	P08670
VIME_HUMAN		R.KVESLQEEIAFLK.K	2	3.3233	P08670

Voltage-dependent anion channel 2 - Homo sapiens (Human)

Q5JSD1_HUMAN		R.PMC#IPPSYADLGK.A	2	2.3078	Q5JSD1
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Voltage-dependent anion-selective channel protein 1 - Homo sapiens (Human)

VDAC1_HUMAN		R.VTQSNFAVGYK.T	2	2.4296	P21796
VDAC1_HUMAN		R.VTQSNFAVGYK.T	2	2.6489	P21796
VDAC1_HUMAN		K.YQIDPDAC#FSAK.V	2	2.68	P21796
VDAC1_HUMAN		K.VNNSLIGLGYTQTLKPGIK.L	2	2.3844	P21796
VDAC1_HUMAN		K.YQIDPDAC#FSAK.V	2	3.0888	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	2.454	P21796
VDAC1_HUMAN		R.VTQSNFAVGYK.T	2	2.419	P21796
VDAC1_HUMAN		K.LTFDSSFSPNTGK.K	2	3.7633	P21796
VDAC1_HUMAN		K.LTFDSSFSPNTGK.K	2	2.6129	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	3.327	P21796
VDAC1_HUMAN		K.SENGLEFTSSGSANTETTK.V	2	3.1667	P21796
VDAC1_HUMAN		K.VNNSLIGLGYTQTLKPGIK.L	2	2.4867	P21796
VDAC1_HUMAN		K.WNTDNTLGTEITVEDQLAR.G	2	4.2515	P21796

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

VDAC2_HUMAN		K.VNNSLIGVGYTQTLRPGVK.L	2	2.6529	P45880
VDAC2_HUMAN		K.VNNSLIGVGYTQTLRPGVK.L	2	3.0911	P45880
VDAC2_HUMAN		K.VNNSLIGVGYTQTLRPGVK.L	3	4.0779	P45880
VDAC2_HUMAN		R.NNFAVGYR.T	2	2.3789	P45880

<i>Protein</i>	<i>Reference</i>	<i>Peptide sequence</i>	<i>Charge</i>	<i>XCorr*</i>	<i>Accession**</i>
	VDAC2_HUMAN	R.TGDFQLHTNVNDGTEFGGSIYQK.V	3	4.82	P45880
	VDAC2_HUMAN	K.VNNSLIGVGYTQTLRPGVK.L	2	2.6781	P45880
<i>Voltage-dependent anion-selective channel protein 3 - Homo sapiens (Human)</i>					
	VDAC3_HUMAN	K.LSQNNFALGYK.A	2	2.9668	Q9Y277
	VDAC3_HUMAN	K.LTLDTIFVPNTGK.K	2	2.5733	Q9Y277
	VDAC3_HUMAN	K.VNNASLIGLGYTQTLRPGVK.L	2	5.0243	Q9Y277
	VDAC3_HUMAN	K.LTLDTIFVPNTGK.K	2	2.4628	Q9Y277
	VDAC3_HUMAN	K.LTLDTIFVPNTGK.K	2	2.4452	Q9Y277
	VDAC3_HUMAN	K.VC#NYGLTFTQK.W	2	2.7216	Q9Y277
	VDAC3_HUMAN	K.VC#NYGLTFTQK.W	2	2.6627	Q9Y277
	VDAC3_HUMAN	K.AADFQLHTHVNDGTEFGGSIYQK.V	3	4.9122	Q9Y277
	VDAC3_HUMAN	K.VNNASLIGLGYTQTLRPGVK.L	2	4.6703	Q9Y277
<i>WD repeat and FYVE domain-containing protein 3 - Homo sapiens (Human)</i>					
	WDFY3_HUMAN	K.GM*LKNSTAALYIDGQLVNTVK.L	2	2.5015	Q8IZQ1
	WDFY3_HUMAN	K.GM*LKNSTAALYIDGQLVNTVK.L	2	2.3626	Q8IZQ1
	WDFY3_HUMAN	K.GM*LKNSTAALYIDGQLVNTVK.L	2	2.4356	Q8IZQ1
<i>Zinc finger, CDGSH-type domain 2 - Homo sapiens (Human)</i>					
	Q8N5K1_HUMAN	K.VVNEINIEDLC#LTK.A	2	3.7081	Q8N5K1
	Q8N5K1_HUMAN	K.VVNEINIEDLC#LTK.A	2	2.491	Q8N5K1
	Q8N5K1_HUMAN	K.VVNEINIEDLC#LTK.A	2	2.8317	Q8N5K1
	Q8N5K1_HUMAN	K.VVNEINIEDLC#LTK.A	2	2.5035	Q8N5K1