

Table S3. List of HQ peptides

Protein Name	Protein Description	Protein Gene	Peptide Modified Sequence	Peptide Sequence	Precursor Charge	Precursor m/z
sp A5A3E0 POTEF_HUMAN	POTE ankyrin domain family member F OS=Homo sapiens GN=POTEF PE=1 SV=2	POTEF	QEYDESGPSIVHR	QEYDESGPSIVHR	2	758.855
sp Q14791 APOL1_HUMAN	Apolipoprotein L1 OS=Homo sapiens GN=APOL1 PE=1 SV=5	APOL1	VTEPISAESGEQVER	VTEPISAESGEQVER	2	815.8996
sp Q14950 ML12B_HUMAN	Myosin regulatory light chain 12B OS=Homo sapiens GN=MYL12B PE=1 SV=2	MYL12B	FTDEEVDELRYR	FTDEEVDELRYR	2	708.3199
sp Q43866 CD5L_HUMAN	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	CD5L	ELGCG[+57]GAASGTPSGILYEPPAEK	ELGCGAASGTPSGILYEPPAEK	2	1102.531
sp Q60333 KIF1B_HUMAN	Kinesin-like protein KIF1B OS=Homo sapiens GN=KIF1B PE=1 SV=5	KIF1B	IN[+1]DLDLK	INDLDLK	2	416.2266
sp Q75636 FCN3_HUMAN	Ficolin-3 OS=Homo sapiens GN=FCN3 PE=1 SV=2	FCN3	QDGSVDFFR	QDGSVDFFR	2	535.7487
sp Q75636 FCN3_HUMAN	Ficolin-3 OS=Homo sapiens GN=FCN3 PE=1 SV=2	FCN3	LLGEVDHYQLALGK	LLGEVDHYQLALGK	2	778.4276
sp Q75636 FCN3_HUMAN	Ficolin-3 OS=Homo sapiens GN=FCN3 PE=1 SV=2	FCN3	YGIDWASGR	YGIDWASGR	2	512.746
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	LTGSSGFVTDGPGNYK	LTGSSGFVTDGPGNYK	2	800.3861
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	C[+57]TWLIEGQPNR	CTWLIEGQPNR	2	687.3352
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	ISM[+1]SSDTEVC[+57]EC[+57]JENWK	ISNSSDTECECSENWK	2	1023.407
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	GEAC[+57]DIPHC[+57]TDNC[+57]GFPHR	GEACDIPHCNTDCGFPHR	3	714.9561
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	EEYSNLK	EEYSNLK	2	441.7138
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	SVNNVVVR	SVNNVVVR	2	443.7589
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	IDSTGN[+1]VTNELR	IDSTGNVTNELR	2	660.3255
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	C[+57]JIN[+1]QSIQ[+57]EK	CINQSIQEK	2	576.7548
sp Q75882 ATRN_HUMAN	Attractin OS=Homo sapiens GN=ATRN PE=1 SV=2	ATRN	GDEC[+57]QLC[+57]EVENR	GDECQLCEVENR	2	754.8088
sp Q95445 APOM_HUMAN	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1 SV=2	APOM	DGLCL[+57]VPR	DGLCLVPR	2	408.7053
sp Q95445 APOM_HUMAN	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1 SV=2	APOM	AFLLTPR	AFLLTPR	2	409.2502
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	DIASGLIGLIC[+57]K	DIASGLIGLIC	2	735.4234
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	TYC[+57]SEPEK	TYCSEPEK	2	507.2159
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	EYTDASFTRN	EYTDASFTRN	2	602.2675
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	GAYPLSIEPIGVR	GAYPLSIEPIGVR	2	686.3852
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	NNEGTYSPNYPQSR	NNEGTYSPNYPQSR	2	952.4139
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	NNEGTYSPNYP[+1]YNPQSR	NNEGTYSPNYPQSR	2	952.9059
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	EVGPTNADPVC[+57]LAK	EVGPTNADPVC[+57]LAK	2	736.3585
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	EVGPTNADPVC[+57]LAK	EVGPTNADPVC[+57]LAK	2	735.8665
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	EDEDQESNK	EDEDQESNK	2	620.7518
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	YTVNQCI[+57]R	YTVNQCR	2	470.7189
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	ELHHLQEQN[+1]VSN[+1]AFLDK	ELHHLQEQNVSN[+1]AFLDK	2	1012.49
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	ELHHLQEQN[+1]VSN[+1]AFLDK	ELHHLQEQNVSN[+1]AFLDK	2	1011.998
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	GEFYIGSK	GEFYIGSK	2	505.7267
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	QYTDSTFR	QYTDSTFR	2	459.2354
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	SGAGTEDSAC[+57]IPWAYSTVDQVK	SGAGTEDSACIPWAYSTVDQVK	3	835.7128
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	DLYSGLIGPLIVCR	DLYSGLIGPLIVCR	2	788.4318
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	TYSDHPEK	TYSDHPEK	2	488.7222
sp P00450 CERU_HUMAN	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	CP	DDEEFIESNK	DDEEFIESNK	2	613.2646
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	YTAQ[+57]ETAR	YTACETAR	2	486.2162
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	NPDSSTTGPWC[+57]YTDTPTVR	NPDSSTTGPWCYTDTPTVR	2	1077.973
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	GQQYQGR	GQQYQGR	2	418.7041
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	LAVTTHTGLP[+57]JAWASQAQ	LAVTTHTGLPCLAWASQAQ	2	998.0277
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	HQDFNSAVQLVENFCR[+57]R	HQDFNSAVQLVENFCR	2	982.4576
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	TATSEYQTFNPR	TATSEYQTFNPR	3	521.2476
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	TFGSGEAD[+57]GLRPFLEK	TFGSGEADGLRPFLEK	2	942.4515
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	KPVAFSDYHPVC[+57]LPDR	KPVAFSDYHPVCLPDR	3	672.012
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	ETAASLLQAGYK	ETAASLLQAGYK	2	626.3326
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	ETWTANVGK	ETWTANVGK	2	503.2536
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	ETWTANVGK	ETWTANVGK	2	503.7456
sp P00734 THRB_HUMAN	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	F2	GQPSVLQVNLPIVERPVC[+57]K	GQPSVLQVNLPIVERPVC	2	1116.63
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	FC[+57]QGLGSLGNPPGK	FCQGLGSLGNPPGK	2	764.8825
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	NIEGFC[+57]GK	NIEGFCGK	2	462.7158
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	YITEIK	YITEIK	2	434.2447
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	NLPNGDFR	NLPNGDFR	2	466.7329
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	IQYVC[+57]HEPYK	IQYVCHEPYK	2	782.3505
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	ESEQGVYT[+57]TAQGIWK	ESEQGVYTCTAQGIWK	2	928.9278
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	C[+57]LPVC[+57]GKPNPVEQR	CLPVC[+57]GKPNPVEQR	3	584.9676
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	VSVHPDVR	VSVHPDVR	2	486.7485
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	QDAC[+57]QGDSSGVFAVR	QDACQGDSSGVFAVR	2	783.8519
sp P00736 C1R_HUMAN	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	C1R	VLNVYDWIK	VLNVYDWIK	2	575.3188
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	TGDDGVYTLNDK	TGDDGVYTLNDK	2	656.3068
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	LPEC[+57]EAVC[+57]GKPK	LPCEAEVCGKPK	2	694.3391
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	NPANPVQR	NPANPVQR	2	448.2409
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	GSPFWQAK	GSPFWQAK	2	460.7349
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	NLFNLN[+1]HSENAQAK	NLFNLHSENAQAK	2	730.3624
sp P00739 HPTR_HUMAN	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	HPR	QLVEIEK	QLVEIEK	2	429.7502
sp P00740 FA9_HUMAN	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	F9	NC[+57]ELDTVC[+57]JNIK	NCELDVTNLIK	2	683.3105
sp P00740 FA9_HUMAN	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	F9	VVC[+57]JSC[+57]JTEGYR	VVCSTCEGYR	2	615.7657
sp P00740 FA9_HUMAN	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	F9	VDAFC[+57]GGSIVNEK	VDAFCGGSIVNEK	2	698.3323
sp P00742 FA10_HUMAN	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	F10	NC[+57]ELFTR	NCELFTR	2	470.2213
sp P00742 FA10_HUMAN	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	F10	NTEQEEGGEAVHEVEVIK	NTEQEEGGEAVHEVEVIK	3	699.3429
sp P00742 FA10_HUMAN	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	F10	TGIVSGFGR	TGIVSGFGR	2	447.2456
sp P00742 FA10_HUMAN	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	F10	QEDAC[+57]QGDSSGPHVTR	QEDACQGDSSGPHVTR	3	571.913
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	QLGAGSIEEC[+57]AAK	QLGAGSIEECAAK	2	667.3245
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	DVVLFEK	DVVLFEK	2	425.2395
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	VYLSEK[+57]K	VYLSECK	2	449.7206
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	WELC[+57]DIPR	WELCDIPR	2	544.7633
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	NLDENYCR[+57]R	NLDENYCR	2	542.2298
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	WECY[+57]JNLK	WECYCNLK	2	506.7315
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	ATTVTGTGPC[+57]QDWAAQEPHR	ATTVTGTGPCQDWAAQEPHR	2	1063.49
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	LFLEPTR	LFLEPTR	2	438.2529
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	LSSPAVITDK	LSSPAVITDK	2	515.7926
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	VIPAC[+57]LPSPNYVADR	VIPACLPSPNYVADR	3	590.9784
sp P00747 PLMN_HUMAN	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	PLG	EAQLPVNIK	EAQLPVNIK	2	570.8166
sp P00748 FA12_HUMAN	Coagulation factor XII OS=Homo sapiens GN=F12 PE=1 SV=3	F12	GRGPQDPWC[+57]ATTPNFDDQQR	GRGPQDPWCATTPNFDDQQR	3	776.6907
sp P00748 FA12_HUMAN	Coagulation factor XII OS=Homo sapiens GN=F12 PE=1 SV=3	F12	EQPPSLTR	EQPPSLTR	2	464.2483
sp P00748 FA12_HUMAN	Coagulation factor XII OS=Homo sapiens GN=F12 PE=1 SV=3	F12	N[+1]HSC[+57]EPC[+57]QTLAVR	NHSCPECQTLAVR	2	786.8483
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	LLQEGQALEYVC[+57]PSGFFYPYPVQTR	LLQEGQALEYVCPSGFFYPYPVQTR	3	939.1302
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	STGWSWTLK	STGWSWTLK	2	483.7482
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	WSGQTAIC[+57]DN[+1]GAGYC[+57]SNPGPIGTR	WSGQTAICDNAGAGYCSNPGPIGTR	3	885.0638
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	WSGQTAIC[+57]DNAGAGYC[+57]SNPGPIGTR	WSGQTAICDNAGAGYCSNPGPIGTR	3	884.7358
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	C[+57]LVNLIEK	CLVNLIEK	2	494.7784
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	VASYGVKPR	VASYGVKPR	2	488.7824

sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	QLNEINYEDHK	2	701.8335
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	DLLYIGK	2	411.242
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	YGQTRIPC[+57]LPC[+57]TEGTTR	2	1062.022
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	EELLPAQDIK	2	578.3164
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	DAQYAPGYDK	2	564.2538
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	DISEVTPTR	2	508.2746
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	FLC[+57]TGGVSPYADPNTC[+57]R	2	957.9273
sp P00751 CFAB_HUMAN	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	CFB	GDSGGPLVYHK	2	540.2958
sp P00915 CAH1_HUMAN	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2	CA1	NGPEQVWSK	2	473.2249
sp P00915 CAH1_HUMAN	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2	CA1	YSSLAESAAS	2	513.7587
sp P00915 CAH1_HUMAN	Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2	CA1	VLDAQIAIK	2	485.8002
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	IPDATNR	2	400.7167
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	TSQIHFFFAK	2	670.8353
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	LQPLDFK	2	430.7475
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	FSPENTR	2	425.7063
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	FSPENTR	2	426.1983
sp P01008 ANT3_HUMAN	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	SERPINC1	ANRPFLVFIR	2	616.8668
sp P01011 AACT_HUMAN	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	SERPINA3	EQLSLDR	2	487.2693
sp P01011 AACT_HUMAN	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	SERPINA3	LINDVVK	2	432.7449
sp P01011 AACT_HUMAN	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	SERPINA3	EIGELYLPK	2	531.2975
sp P01011 AACT_HUMAN	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	SERPINA3	NLAVSQVVHK	2	547.8195
sp P01019 ANGT_HUMAN	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	AGT	DPTFPAPIQAK	2	649.3612
sp P01019 ANGT_HUMAN	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	AGT	LQAILGVPMWK	2	562.845
sp P01019 ANGT_HUMAN	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	AGT	QPQVQGLALYTPVVLPR	2	949.5486
sp P01019 ANGT_HUMAN	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	AGT	SLDFTELDVAAEK	3	479.9084
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	TIYTPGSVLVYR	3	457.582
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	IFTVNHK	2	429.7452
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	IFTVNHK	2	430.2373
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	FYYIYNK	2	570.274
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	ISLPSLK	2	443.7658
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	VLLDGVQNPQR	2	555.8169
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	QELSEAEQATR	2	631.3046
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	DSC[+57]VGSVLVK	2	532.2762
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	VVLVAVDK	2	421.7709
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	IWDVVEK	2	444.7449
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	ADIGC[+57]TPGSGK	2	531.7479
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	AELQC[+57]PQPAAR	2	620.8088
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	ASHLGLAR	2	412.7405
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	LPYSVVR	2	417.2476
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	NEQVEIR	2	444.2327
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	AVLYNVR	2	449.7427
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	SSLSVPYVIVPLK	2	701.4212
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	TGLQEVVK	2	501.7769
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	QGAELEIK	2	436.266
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	QPSAFAFAVK	2	576.806
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	APSTWLTVAVYK	2	668.369
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	DIC[+57]EEQVNSLPGSITK	2	895.4355
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	VTIKPAPETEK	2	606.8454
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	N[+1]TUIYDK	2	547.3106
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	VHQYFNVELIQGAVK	2	921.4991
sp P01024 CO3_HUMAN	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	C3	ACEPGVDYVYK	2	650.7975
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	FSYSSGHVHLSSENK	3	560.2655
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	FQN[+1]SAILTQPK	2	680.8772
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	FQNSAILTQPK	2	680.3852
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	VYSLNDLKPAAK	2	681.8668
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	EYVLPFHFSVIEPEYFNIGYK	3	844.4213
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	NFEITIK	2	432.7449
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	ELSYSLLEDLNNK	2	794.3805
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	VYLSPPYK	2	435.242
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	LNLVATPLFLKPGIPYPIK	3	698.7624
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	ITHYNYLISK	2	682.8823
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	C[+57]C[+57]YDGC[+57]VNNDET[C+57]EQR	2	1075.885
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	AFTEC[+57]C[+57]JVASQLR	2	770.866
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	TLLPVSKPEIR	2	626.8848
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	GTVYNNR	2	436.7167
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	ESYSGVTLDPK	2	612.2988
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	IPLDLVPK	2	447.7866
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	NADYSYSVWK	2	616.7828
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	LQGTLPVEAR	2	542.3115
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	AFDIC[+57]PLVK	2	531.7863
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	GNPPIYR	2	408.7218
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	DINNVNPKV	2	587.827
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	WLSEQR	2	474.2327
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	TSTSEVC[+57]SFYK	2	775.8558
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	IDTQDIEASHYR	2	724.3442
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	IVAC[+57]JASYKPSR	2	626.3293
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	QTAC[+57]KPEIAYAYK	2	771.8847
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	ATLLDIYK	2	468.7737
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	DSEITFIK	2	476.7529
sp P01031 CO5_HUMAN	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	C5	VTC[+57]TNAELVK	2	567.7948
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	QVAGALNFR	2	502.2878
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	ITYSIVQTN[+1]C[+57]SK	2	707.8478
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	ENFLFLTPDC[+57]JK	2	692.3343
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	IASFSQNC[+57]DIYPGK	2	800.3772
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	IC[+57]JGVC[+57]PR	2	431.2071
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	YFIDFVAR	2	515.7715
sp P01042 KNG1_HUMAN	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	KNG1	ESNEELTESC[+57]ETK	2	778.3251
sp P01344 IGF2_HUMAN	Insulin-like growth factor II OS=Homo sapiens GN=IGF2 PE=1 SV=1	IGF2	GIVEEC[+57]C[+57]JFR	2	585.2575
sp P02042 HBD_HUMAN	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	HBD	LLVYVPWTQR	2	567.8664
sp P02042 HBD_HUMAN	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	HBD	GTFQSLSLHCDK	2	761.3537
sp P02042 HBD_HUMAN	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	HBD	LHVDPENFR	2	563.7856
sp P02042 HBD_HUMAN	Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	HBD	VVAGVANALAHK	2	575.3406
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	WELALGR	2	422.7374
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	SELEEQLTPVAEETR	2	865.9258
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	DADDLQK	2	402.6903
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	LAVYQAGAR	2	474.7667
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	AATVGSLAGQPLQER	2	749.4046
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	AQAWGGER	2	409.2012

sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	EQVAEVR	2	415.722
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	LEEQAAQIR	2	557.7962
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	LQAEAFQAR	2	517.2749
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	QWAGLVEK	2	465.7558
sp P02649 APOE_HUMAN	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	APOE	VQAAVGTSAAVPVPSDNH	2	810.9025
sp P02654 APOC1_HUMAN	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	APOC1	EFGNTLEDK	2	526.7484
sp P02654 APOC1_HUMAN	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	APOC1	EFGN[+1]TLEDK	2	527.2404
sp P02655 APOC2_HUMAN	Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1 SV=1	APOC2	ELSSSYWESAK	2	643.7986
sp P02655 APOC2_HUMAN	Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1 SV=1	APOC2	TYLPAVDEK	2	518.2715
sp P02656 APOC3_HUMAN	Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=1 SV=1	APOC3	DYWSTVK	2	449.7189
sp P02741 CRP_HUMAN	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	CRP	ESDTSYVSLK	2	564.7746
sp P02741 CRP_HUMAN	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	CRP	GYSIFSATK	2	568.7848
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	AYSDLR	2	406.2009
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	AYSLSFYNTQGR	2	703.8386
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	DNELLVYK	2	497.2662
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	VGESLYIGR	2	578.8035
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	QGYFVEAQPK	2	583.7957
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	IVLGEQDQSYGGK	2	697.3515
sp P02743 SAMP_HUMAN	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	APCS	GVIKPLVWV	2	643.8972
sp P02746 C1QB_HUMAN	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	C1QB	TINVPLR	2	406.7531
sp P02747 C1QC_HUMAN	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	C1QC	FQSVFTVTR	2	542.7929
sp P02747 C1QC_HUMAN	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	C1QC	QTHQPPAPNSLIR	2	729.8942
sp P02747 C1QC_HUMAN	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	C1QC	FNAVLNTPQGDDYTSTGK	2	964.4552
sp P02747 C1QC_HUMAN	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	C1QC	TNQVNSGGVLLR	2	629.3491
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	VVEESLAR	2	516.272
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	DGNTLTYYR	2	551.7618
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	TEHYEEQIEAFK	2	762.3543
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	ALPTTYEK	2	461.7477
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	LSPVNLVPVK	2	621.8765
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	AIEDYINEFSVR	3	485.9087
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	C[+57]LJC[+57]JAC[+57]JPFK	2	528.2272
sp P02748 CO9_HUMAN	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	C9	FEGIAECIK	2	577.2815
sp P02749 APOH_HUMAN	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	APOH	VYKPSAGN[+1]NSLYR	2	735.3728
sp P02749 APOH_HUMAN	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	APOH	C[+57]JPFSRPDNGFVNPYAKPTLYYK	3	911.1161
sp P02749 APOH_HUMAN	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	APOH	ATVVYQGER	2	511.7669
sp P02749 APOH_HUMAN	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	APOH	EHSSLAFWK	2	552.7773
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	LQELHLSNGLESLSPEFLRPVPQLR	3	987.2048
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	ALGHLDSLGNR	2	576.8096
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	GPLQLER	2	406.7349
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	DLLLPQPDLR	2	590.3402
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	YFLNGNK	2	484.7636
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	DGFDISGNPWIC[+57]DQNLSDLYR	3	829.0411
sp P02750 A2GL_HUMAN	Leucine-rich alpha-2-glycoprotein OS=Homo sapiens GN=LRG1 PE=1 SV=2	LRG1	GQTLAVAK	2	450.7793
sp P02753 RET4_HUMAN	Retinol-binding protein 4 OS=Homo sapiens GN=RBPA PE=1 SV=3	RBPA	YWGVASFLQK	2	599.8164
sp P02753 RET4_HUMAN	Retinol-binding protein 4 OS=Homo sapiens GN=RBPA PE=1 SV=3	RBPA	QEELC[+57]LAR	2	509.7529
sp P02760 AMBP_HUMAN	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP	ETLLQDFR	2	511.2693
sp P02760 AMBP_HUMAN	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP	GEC[+57]JVPGEQEPELIPR	2	960.4802
sp P02760 AMBP_HUMAN	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP	EC[+57]LQTC[+57]JR	2	483.7102
sp P02760 AMBP_HUMAN	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP	TVAAC[+57]JNLPIVR	2	607.3397
sp P02760 AMBP_HUMAN	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	AMBP	C[+57]JLFPYGGC[+57]QGNGNK	2	835.8743
sp P02765 FETUA_HUMAN	Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1	AHSG	C[+57]JNLIAEK	2	424.2207
sp P02766 TTHY_HUMAN	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	TTR	YTIAALLSPYSYSTAVVTNPK	2	1180.623
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	EFSLGK	2	409.2138
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	ELPEHTVK	2	476.7585
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	DVC[+57]DPGNTK	2	503.2189
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	YTFELSR	2	458.2322
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	THLPEVFLSK	2	585.8295
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	VLEPTLK	2	400.2498
sp P02774 VTDB_HUMAN	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	GC	ELSSFIDK	2	469.7451
sp P02775 CXCL7_HUMAN	Platelet basic protein OS=Homo sapiens GN=PPBP PE=1 SV=3	PPBP	ICLDPDAPR	2	528.7608
sp P02790 HEMO_HUMAN	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	HPX	N[+1]FPSPVDAAFR	2	611.2986
sp P02790 HEMO_HUMAN	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	HPX	GGYTLVSGYVK	2	571.298
sp P02790 HEMO_HUMAN	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	HPX	LWWLDLK	2	487.2789
sp P03951 FA11_HUMAN	Coagulation factor XI OS=Homo sapiens GN=F11 PE=1 SV=1	F11	IVGGTASVR	2	430.2534
sp P03951 FA11_HUMAN	Coagulation factor XI OS=Homo sapiens GN=F11 PE=1 SV=1	F11	VYSGILN[+1]QSEIK	2	676.3588
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	DSVTGTLPK	2	459.2506
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	GVNFN[+1]VSK	2	433.2243
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	YSPGGTPTAIK	2	546.2902
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	VLTPDAFVC[+57]JR	2	589.3053
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	GVNVC[+57]QETC[+57]TK	2	648.2896
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	GVN[+1]VC[+57]QETC[+57]TK	2	648.7814
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	IYSGILNLSITK	2	719.3956
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	DTPFQSK	2	468.2453
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	ELIHHQNYK	2	579.3193
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	LQAPLNLYTEFQKPICLPSK	2	1124.588
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	GEIQNILQK	2	521.7982
sp P03952 KLKB1_HUMAN	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	KLKB1	EQPGVYTK	2	461.2375
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	YTC[+57]LPGYVR	2	564.779
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	GSSVIHC[+57]DADSK	2	638.2853
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	C[+57]HPGYKPTDEPTVIC[+57]QK	3	744.6856
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	WTPYQGC[+57]EALC[+57]C[+57]PEPK	2	998.4235

sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	LNNGEITQHR	LNNGEITQHR	2	591.3047
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	EEIYEC[+57]DK	EEIYECDK	2	599.7685
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	GYILVGQAK	GYILVGQAK	2	474.7793
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	LSC[+57]SYSHWSAPAPQC[+57]K	LSCSYSHWSAPAPQCK	2	939.9167
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	DQYVEPN[+1]VTIQC[+57]DSGYGVVGPGQSITC[+57]SGN[+1]R	DQYVEPNVTIQCDSGYGVVGPGQSITCSG NR	3	1144.505
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	TWYPEVPK	TWYPEVPK	2	510.2635
sp P04003 C4BPA_HUMAN	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	C4BPA	LSLEIEQLELQR	LSLEIEQLELQR	2	735.9118
sp P04004 VTNC_HUMAN	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTN	C[+57]TEGFNVDK	CTEGFNVDK	2	535.2346
sp P04004 VTNC_HUMAN	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTN	AVRPGYPK	AVRPGYPK	2	444.2585
sp P04004 VTNC_HUMAN	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTN	DVWGIEGPIDAAFR	DVWGIEGPIDAAFR	2	823.9123
sp P04004 VTNC_HUMAN	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTN	QPQFISR	QPQFISR	2	438.2403
sp P04004 VTNC_HUMAN	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	VTN	SIAQYWLGC[+57]PAPGHL	SIAQYWLGCAPAGHL	2	835.4114
sp P04070 PROC_HUMAN	Vitamin K-dependent protein C OS=Homo sapiens GN=PROC PE=1 SV=1	PROC	DTEDQEDQVDR	DTEDQEDQVDR	2	723.8026
sp P04070 PROC_HUMAN	Vitamin K-dependent protein C OS=Homo sapiens GN=PROC PE=1 SV=1	PROC	EVFVHPN[+1]YSK	EVFVHPNYSK	2	610.8009
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	YTYNYEAESSSGVPGTADSR	YTYNYEAESSSGVPGTADSR	2	1077.467
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	EVYGFNPEKG	EVYGFNPEKG	2	570.272
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	QVFLYPEK	QVFLYPEK	2	512.2791
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DEPTYLINIK	DEPTYLINIK	2	603.3243
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	GISALLVPPEEAK	GISALLVPPEEAK	2	833.9667
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	GNVATEISTER	GNVATEISTER	2	588.7964
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DLGQC[+57]DR	DLGQCDR	2	432.1874
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TGISPLALIK	TGISPLALIK	2	506.8237
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	EQHLFLPSYK	EQHLFLPSYK	2	704.8666
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LTISEQNIQR	LTISEQNIQR	2	601.3304
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ATLYALSHAVNNYHK	ATLYALSHAVNNYHK	2	851.439
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DQEVLLQTFLDDASPGDK	DQEVLLQTFLDDASPGDK	2	995.9838
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IVQLPWEQNEQVK	IVQLPWEQNEQVK	2	862.4725
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SVSLPSLDPASAK	SVSLPSLDPASAK	2	636.3457
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IEGNLIFDPNNYLPK	IEGNLIFDPNNYLPK	2	873.9567
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ALYVWNGQVPDGVSK	ALYVWNGQVPDGVSK	2	816.9227
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VLVDFHGYTK	VLVDFHGYTK	2	589.8139
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FIIPSPK	FIIPSPK	2	401.2471
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TEVIPPLIENR	TEVIPPLIENR	2	640.8641
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FVTQAEGAK	FVTQAEGAK	2	475.7507
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SEILAHWSPAK	SEILAHWSPAK	2	619.83
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VAWHYDEEK	VAWHYDEEK	2	588.7696
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IEIPLPFGGK	IEIPLPFGGK	2	535.8159
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TPALHFH	TPALHFH	2	407.2345
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	EFQVPTFTIPK	EFQVPTFTIPK	2	653.8557
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FLOSNIK	FLOSNIK	2	418.7293
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LGNNPVSK	LGNNPVSK	2	414.7323
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VSSFYAK	VSSFYAK	2	401.2107
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LNGENSLR	LNGENSLR	2	451.7381
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FN[+1]SSYLQGTNQITGR	FNSSYLQGTNQITGR	2	843.9077
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FN[+1]SSYLQGTN[+1]QITGR	FNSSYLQGTNQITGR	2	844.3997
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	YENVELTK	YENVELTK	2	586.7953
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SEYQADYESLR	SEYQADYESLR	2	680.8044
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IGDQGISSTATNLK	IGDQGISSTATNLK	2	753.3939
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LDNIYSDDK	LDNIYSDDK	2	527.7562
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	QTVNLQLQPSYLVTLTNSDLK	QTVNLQLQPSYLVTLTNSDLK	2	1188.144
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	YNALDITNNKG	YNALDITNNKG	2	611.8068
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	GAYQNNK	GAYQNNK	2	518.7565
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VQGVFEFSHR	VQGVFEFSHR	2	529.7725
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	AEPLAFTFSHDYK	AEPLAFTFSHDYK	2	763.3697
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SISAALEHK	SISAALEHK	2	478.264
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VSAALTPAEQGTGTWK	VSAALTPAEQGTGTWK	2	801.4303
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TQFNNNYSQDLDAYNTK	TQFNNNYSQDLDAYNTK	2	1082.975
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TQFN[+1]JNNYSQDLDAYNTK	TQFNNNYSQDLDAYNTK	2	1083.467
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IGVELTGR	IGVELTGR	2	422.748
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TLADITLDSPIK	TLADITLDSPIK	2	700.4058
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DAVEKPKQFTIVAFVK	DAVEKPKQFTIVAFVK	2	910.9933
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ITENDIQALDDAK	ITENDIQALDDAK	2	779.9016
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DSYDLHDLK	DSYDLHDLK	2	553.2617
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TIHDLHLFIENIDFNK	TIHDLHLFIENIDFNK	2	985.0125
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SGSSTASWQNVDTK	SGSSTASWQNVDTK	2	790.8812
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	QHIEAIDVR	QHIEAIDVR	2	540.7935
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LSNLVQLQVK	LSNLVQLQVK	2	514.8086
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SFDYHQFVDETNDK	SFDYHQFVDETNDK	2	872.8761
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LFLEETK	LFLEETK	2	440.2447
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ETLEDTR	ETLEDTR	2	432.2089
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NLTDFAEQYSIQDWAK	NLTDFAEQYSIQDWAK	2	964.9549
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ALVEQGFTVPEIK	ALVEQGFTVPEIK	2	715.8981
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ATQPTDPDFVPLTDLR	ATQPTDPDFVPLTDLR	2	917.4909
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IPSVQINFK	IPSVQINFK	2	523.3057
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VEDIPLAR	VEDIPLAR	2	456.7611
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ITLPDFR	ITLPDFR	2	431.2451
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SNTVASLHTEK	SNTVASLHTEK	2	593.8068
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NTELSN[+1]GVIVK	NTELSNGVIVK	2	644.3614
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	INNQLTLDSENTK	INNQLTLDSENTK	2	680.857
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	AGHIAWTSSEK	AGHIAWTSSEK	2	557.7856
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LPYTIITPPKL	LPYTIITPPKL	2	678.9105
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	DFSLWEK	DFSLWEK	2	462.7267
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	QSFDSLWK	QSFDSLWK	2	462.2453
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NNALDFVTK	NNALDFVTK	2	511.2693
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LSPDFK	LSPDFK	2	410.2342
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	AQIPLR	AQIPLR	2	405.7634
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	EYSGTIASEANTYLNK	EYSGTIASEANTYLNK	3	616.6268
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IDDIWNLVWK	IDDIWNLVWK	2	622.8297
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ENFAGEATLQR	ENFAGEATLQR	2	618.3044
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NHQLQGLFFTTNGEHTSK	NHQLQGLFFTTNGEHTSK	3	691.3447
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NIILPVYDK	NIILPVYDK	2	537.8133

sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	SLWDFLK	2	454.7475
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LDVTTISGR	2	481.2693
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	VSTAFVYTK	2	508.2766
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	NPNGYSFSIPVK	2	661.8406
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	TSSFALNLPFLPEVK	2	808.9483
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FPEVDVLTG	2	524.2897
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FSPVAGIVIPSFQALTAR	2	937.5304
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IEDGTASK	2	467.248
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	YEGLOQEWEGK	2	619.788
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ESDEETQIK	2	539.7486
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	ATGVLYDYVYNK	2	621.8219
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	QIDDIDIVR	2	487.2511
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	AASGTTGTGYQEWK	2	700.3281
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	AQNLVQELLTQEQQASFQGLK	2	1183.603
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	FQFPGKPGIYTR	2	705.8801
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	EVGTVLVSQVYSK	2	655.3535
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	IISDYHQQFR	2	653.8306
sp P04114 APOB_HUMAN	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	APOB	LQDQFSDQLSDYYEK	2	875.894
sp P04180 LCAT_HUMAN	Phosphatidylcholine-sterol acyltransferase OS=Homo sapiens GN=LCAT PE=1 SV=1	LCAT	STELC[+57]GLWQGR	2	653.8141
sp P04196 HRG_HUMAN	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG	VIDFM[+1]C[+57]TTSSVSSALANTK	2	1008.483
sp P04196 HRG_HUMAN	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG	EENDDFASFR	2	615.2571
sp P04196 HRG_HUMAN	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG	QIGSVYR	2	411.7271
sp P04196 HRG_HUMAN	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG	GEVLP[PEANFSPFLPHHK	2	1113.589
sp P04196 HRG_HUMAN	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	HRG	HPLKPDNQPFQPSVSESC[+57]PGK	3	783.7146
sp P04217 A1BG_HUMAN	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4	A1BG	NGVAQEPVHLDSPAIK	2	837.9441
sp P04217 A1BG_HUMAN	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4	A1BG	LLELTGPK	2	435.7684
sp P04217 A1BG_HUMAN	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4	A1BG	GVTFLLR	2	403.2502
sp P04217 A1BG_HUMAN	Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4	A1BG	C[+57]EGPIPDVTFELLR	2	823.4163
sp P04275 VWF_HUMAN	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	VWF	IGEADFNR[+1]R	2	461.7169
sp P04275 VWF_HUMAN	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	VWF	YDAAQLR	2	418.7167
sp P04275 VWF_HUMAN	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	VWF	ILAGAPGDSNVVK	2	620.8484
sp P04275 VWF_HUMAN	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	VWF	GLRPS[C+57]PNSQSPVK	3	509.5963
sp P04278 SHBG_HUMAN	Sex hormone-binding globulin OS=Homo sapiens GN=SHBG PE=1 SV=2	SHBG	TSSSFEVR	2	456.7247
sp P04406 G3P_HUMAN	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	GAPDH	GALQNIIPASTGAAK	2	706.3988
sp P05062 ALDOB_HUMAN	Fructose-bisphosphate aldolase B OS=Homo sapiens GN=ALDOB PE=1 SV=2	ALDOB	LDQGGAPLAGTNK	2	621.3279
sp P05090 APOD_HUMAN	Apolipoprotein B OS=Homo sapiens GN=APOD PE=1 SV=1	APOD	ADGTNVNQEIGEATPVN[+1]LTEPAK	2	1128.056
sp P05154 IPSP_HUMAN	Plasma serine protease inhibitor OS=Homo sapiens GN=SERPINA5 PE=1 SV=3	SERPINA5	GFQQLLQELNQPR	2	785.9204
sp P05154 IPSP_HUMAN	Plasma serine protease inhibitor OS=Homo sapiens GN=SERPINA5 PE=1 SV=3	SERPINA5	TLYLADTFPTNFR	2	779.8986
sp P05154 IPSP_HUMAN	Plasma serine protease inhibitor OS=Homo sapiens GN=SERPINA5 PE=1 SV=3	SERPINA5	VVGVPYQGN[+1]ATALFILPSEK	2	1081.081
sp P05155 IC1_HUMAN	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	SERPING1	DFTC[+57]VHQALK	2	609.8004
sp P05155 IC1_HUMAN	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	SERPING1	TLYSSSPR	2	455.7351
sp P05155 IC1_HUMAN	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	SERPING1	FQPTLLTLPR	2	593.3531
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	VITYSQEDLVEK	2	706.3512
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	C[+57]JEGTC[+57]VC[+57]K	2	563.7381
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	SFPTYC[+57]QQK	2	579.766
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	SLECH[+57]LHPGTK	2	571.2871
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	FLNN[+1]JGTC[+57]TAEKG	2	656.7955
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	AC[+57]DGINDC[+57]GDQSDCLC[+57]C[+57]K	2	1058.887
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	AQLGDLVPQWAIK	2	719.9063
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	IVIEYVDR	2	503.782
sp P05156 CFAI_HUMAN	Complement factor I OS=Homo sapiens GN=CFI PE=1 SV=2	CFI	VFSLQWGEVK	2	596.8217
sp P05452 TETN_HUMAN	Tetranectin OS=Homo sapiens GN=CLEC3B PE=1 SV=3	CLEC3B	EQQALQTVLC[+57]LK	2	659.3452
sp P05452 TETN_HUMAN	Tetranectin OS=Homo sapiens GN=CLEC3B PE=1 SV=3	CLEC3B	TFHEASEDC[+57]ISR	2	726.3146
sp P05452 TETN_HUMAN	Tetranectin OS=Homo sapiens GN=CLEC3B PE=1 SV=3	CLEC3B	NWETEITAQPDGGK	2	773.3626
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	AQWANFPDPK	2	630.804
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	TEDSSSFLDK	2	621.2984
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	NALALFVLPK	2	543.3995
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	GWVDLFPVK	2	530.7949
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	AVLHIGEK	2	433.7584
sp P05543 THBG_HUMAN	Thyroxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	SERPINA7	GTEAAVPEVELSDQEPNTFLHPIQIDR	3	1063.874
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	GGETAQSDAPQWEQLNNK	2	986.9534
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	FAFNLYR	2	465.7452
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	YEITTHNLFR	2	703.875
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	NFGYTLLR	2	435.7271
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	QFPILLDFK	2	560.8237
sp P05546 HEP2_HUMAN	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	SERPIND1	NYNLVESLK	2	540.7822
sp P06276 CHLE_HUMAN	Cholinesterase OS=Homo sapiens GN=BCHE PE=1 SV=1	BCHE	DEGTAFLVVGAPGFSK	2	829.9067
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	HVVPNEVVQR	2	638.362
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	SEDC[+57]FILDHKG	2	660.7981
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	TASDFITK	2	441.732
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	QTQVSVLPPEGGETPLFK	2	915.4858
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	DPDQTDGLGLSYLSHIANVER	2	1194.077
sp P06396 GELS_HUMAN	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	GSN	AGALNSNDAFVLK	2	660.3513
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	C[+57]SSNLVLTGSSER	2	705.3381
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	EC[+57]QGN[+1]GVWSVGTEPIC[+57]R	2	617.5981
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	EC[+57]QGN[+1]GVWSVGTEPIC[+57]R	2	925.4014
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	HAILITDGK	2	540.8242
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	ELININQK	2	486.2796
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	ELNELGSK	2	445.2349
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	ELN[+1]ELGSK	2	445.7269
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	TPWHVTIKPK	2	603.8533
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	DGNDHSLWR	2	550.2494
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	EVVTDQFLC[+57]SGTQDESPC[+57]K	2	1165.002
sp P06681 CO2_HUMAN	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	C2	GESGGAFLER	2	561.2829
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	SELTQNALFQDK	2	817.9229
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LGEVNTYAGDLQK	2	704.3594
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LGEVN[+1]TYAGDLQK	2	704.8514

sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LVPFATELHER	LVPFATELHER	3	437.9067
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LLPHANEVSQK	LLPHANEVSQK	2	618.3408
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LLPHANJ+1IEVSQK	LLPHANEVSQK	2	618.8328
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LEPYADQLR	LEPYADQLR	2	552.7878
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	ENADSLQASLRPHADELK	ENADSLQASLRPHADELK	2	997.5005
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	IDQNVEELK	IDQNVEELK	2	544.2851
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	IDQTVEELR	IDQTVEELR	2	551.7906
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	SLAPYAQDTQEK	SLAPYAQDTQEK	2	675.8304
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	ISASAEELR	ISASAEELR	2	488.2589
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	LAPLAEDVR	LAPLAEDVR	2	492.2796
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	GNTGELQK	GNTGELQK	2	423.7194
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	VEPYGENFNK	VEPYGENFNK	2	598.7828
sp P06727 APOA4_HUMAN	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	APOA4	VNSFFSTFK	VNSFFSTFK	2	538.7742
sp P06753 TPM3_HUMAN	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=1 SV=2	TPM3	IQLVEEELDR	IQLVEEELDR	2	622.3301
sp P07225 PROS_HUMAN	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	PROS1	ASFTC[+57]TC[+57]KPGWQGEK	ASFTCTCKPGWQGEK	2	878.8927
sp P07225 PROS_HUMAN	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	PROS1	DVDEC[+57]SLKPSIC[+57]GTAVC[+57]K	DVDECSLKPSICGTAVCK	2	1019.966
sp P07225 PROS_HUMAN	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	PROS1	IETISHEDLQR	IETISHEDLQR	2	670.8439
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	FGGTIC[+57]SGDIWDQASC[+57]JSSSTTC[+57]VVR	FGGTICSGDIWDQASCSSSTTCVVR	3	884.709
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	HLVC[+57]NGDQDC[+57]LDGSEDDDC[+57]EDV	HLVCNGDQDCLDGSEDDCEDVVR	3	908.3465
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	AIDEDC[+57]SQYEIPGSQK	AIDEDCSQYEIPIGSQK	2	968.9333
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	INVGGGLSGDHC[+57]K	INVGGGLSGDHCK	2	657.317
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	GGSSGWSGGLAQN[+1]R	GGSSGWSGGLAQNR	2	667.8078
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	HTSLGLEAK	HTSLGLEAK	2	526.7904
sp P07357 C08A_HUMAN	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	C8A	LGSLGAAC[+57]EQTQTEGAK	LGSLGAACECQTQTEGAK	2	860.9122
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	YAYLLQPSQFHGEPCE[+57]NFS DK	YAYLLQPSQFHGEPGCFNFS DK	3	801.0354
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	C[+57]EGFVC[+57]JAQTGR	CEGFVCAQTGR	2	642.7766
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	LLC[+57]NGDNDC[+57]GDQSEANCC[+57]R	LLCNGDNDCGDQSEANCR	2	1106.918
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	SGFSFGFK	SGFSFGFK	2	438.7162
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	SVFLHAR	SVFLHAR	2	415.2376
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	SDLEVAHYK	SDLEVAHYK	2	531.2667
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	GDYTLNNVHAC[+57]AK	GDYTLNNVHACAK	2	731.8408
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	QALEEFQK	QALEEFQK	2	496.756
sp P07358 C08B_HUMAN	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	C8B	C[+57]DC[+57]IC[+57]PVGSQLAC[+57]EVS YR	CDICIPVGSQGLACEVS YR	2	1115.971
sp P07359 GP1BA_HUMAN	Platelet glycoprotein Ib alpha chain OS=Homo sapiens GN=GP1BA PE=1 SV=2	GP1BA	LTSPLPLGALR	LTSPLPLGALR	2	520.8268
sp P07359 GP1BA_HUMAN	Platelet glycoprotein Ib alpha chain OS=Homo sapiens GN=GP1BA PE=1 SV=2	GP1BA	GLGELQELYLK	GLGELQELYLK	2	631.8532
sp P07360 C08G_HUMAN	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	C8G	RPASPISTIQPK	RPASPISTIQPK	2	647.8775
sp P07360 C08G_HUMAN	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	C8G	FLQEQGHR	FLQEQGHR	2	507.7594
sp P07360 C08G_HUMAN	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	C8G	QLYGDGTVLGR	QLYGDGTVLGR	2	589.8118
sp P07360 C08G_HUMAN	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	C8G	SLPVSDSVLSGFEQR	SLPVSDSVLSGFEQR	2	810.915
sp P07360 C08G_HUMAN	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	C8G	VQEAHLTDQIFYFPK	VQEAHLTDQIFYFPK	2	982.9913
sp P07737 PROF1_HUMAN	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	PFN1	EGVHGGGLNK	EGVHGGGLNK	2	512.2827
sp P07996 TSP1_HUMAN	Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2	THBS1	GTLALLER	GTLALLER	2	436.7636
sp P07996 TSP1_HUMAN	Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2	THBS1	GGVNDNFQGVQLQNR	GGVNDNFQGVQLQNR	2	808.9106
sp P07996 TSP1_HUMAN	Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2	THBS1	AQYSGLSVK	AQYSGLSVK	2	505.2693
sp P08185 CBG_HUMAN	Corticosteroid-binding globulin OS=Homo sapiens GN=SERPINA6 PE=1 SV=1	SERPINA6	HLVALSPK	HLVALSPK	2	432.7687
sp P08185 CBG_HUMAN	Corticosteroid-binding globulin OS=Homo sapiens GN=SERPINA6 PE=1 SV=1	SERPINA6	QINSYVK	QINSYVK	2	426.2347
sp P08571 CD14_HUMAN	Monocyte differentiation antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	CD14	VLSIAQAHSAPFSC[+57]EQVR	VLSIAQAHSAPFSCQVR	3	667.3388
sp P08571 CD14_HUMAN	Monocyte differentiation antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	CD14	AFPALTSLDLSDNPNGLGER	AFPALTSLDLSDNPNGLGER	2	987.0024
sp P08571 CD14_HUMAN	Monocyte differentiation antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	CD14	ATVNPSAPR	ATVNPSAPR	2	456.7485
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	NTEILTGWSWSDQTYPEGTQAIYK	NTEILTGWSWSDQTYPEGTQAIYK	3	868.0818
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	GEWWALNPLR	GEWWALNPLR	2	577.8195
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	EC[+57]DTDGTWNTDIPIC[+57]EVVK	ECTDGTWNTDIPICEVVK	2	1075.972
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]LPVTAPENGK	CLPVTAPENGK	2	593.3003
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]LPVTAPEN[+1]GK	CLPVTAPENGK	2	593.7923
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]VEISC[+57]JR	CVEISCK	2	448.2042
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	SPDIVNGSPISQK	SPDIVNGSPISQK	2	671.3541
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	GDAVC[+57]JTESGWRPLPSC[+57]EEK	GDAVCTESGWRPLPSCEEK	2	1089.483
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	TGDEITYQC[+57]JR	TGDEITYQCR	2	621.7746
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]JTSGWIPAPR	CTSGTWIPAPR	2	623.3059
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]JLKPC[+57]DYDPDK	CTLKPCDYDPDK	2	755.3574
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	SIDVAC[+57]HPGYALPK	SIDVACHPGYALPK	2	764.3849
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	NDFTWFK	NDFTWFK	2	479.2269
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	IDVHLVPDR	IDVHLVPDR	2	532.2984
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	EEYGHSEVVEYYC[+57]NPR	EEYGHSEVVEYYCNP	2	1015.931
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	SSNLIUEHLK	SSNLIUEHLK	2	698.3957
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	EGWIHTVC[+57]JIN[+1]GR	EGWIHTVCINGR	2	721.8459
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	EGWIHTVC[+57]JINGR	EGWIHTVCINGR	2	721.3539

sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	WQSIPLC[+57]VEK	WQSIPLCVEK	2	630.3263
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	IPC[+57]SQPPQIEHGTIN[+1]SSR	IPCSQPPQIEHGTINSSR	3	674.6618
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	LSVTC[+57]JEGGFR	LSVTCGGGFR	2	595.269
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	WSSPPQC[+57]EGLPC[+57]K	WSSPPQCEGLPCK	2	773.3449
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	DTSC[+57]VNPPTVQNAVIVSR	DTSCVNPPTVQNAVIVSR	2	1010.991
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	C[+57]LHPC[+57]VISR	CLHPCVISR	2	571.2839
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	TGESVEFVC[+57]K	TGESVEFVCK	2	578.2712
sp P08603 CFAH_HUMAN	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	CFH	TTC[+57]JWDGK	TTCWJWDGK	2	434.1869
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	QLTSGPNQEQVSPLTLLK	QLTSGPNQEQVSPLTLLK	2	977.0362
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	LGNOEPGGQTALK	LGNOEPGGQTALK	2	656.8464
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	LGN[+1]QEPGGQTALK	LGNOEPGGQTALK	2	657.3384
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	SPPGV[C][+57]SR	SPPGVCSR	2	430.2082
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	LQQLVHAGSGPCLPHLLSR	LQQLVHAGSGPCLPHLLSR	3	695.0458
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	EDFLEQSEQLFGAKPVSLTGK	EDFLEQSEQLFGAKPVSLTGK	3	775.0654
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	QEDDLANINQWVK	QEDDLANINQWVK	2	786.8863
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	FDPSLTQR	FDPSLTQR	2	482.2483
sp P08697 A2AP_HUMAN	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	SERPINF2	DFLQSLK	DFLQSLK	2	425.7371
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	SDFSNEER	SDFSNEER	2	492.2069
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	SDFS[N][+1]EER	SDFSNEER	2	492.6989
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	QFGPYC[+57]GHGFGPLNIETK	QFGPYCGHFGPGLNIETK	3	707.007
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	EDTPNSVWPEAK	EDTPNSVWPEAK	2	686.8226
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	VGATSFYSTC[+57]QSNKG	VGATSFYSTCQSNKG	2	803.9671
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	C[+57]QPVC[+57]GIPESIEGK	CQPVDCGIPESIEGK	2	901.906
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	VEDPESTLFGSVIR	VEDPESTLFGSVIR	3	516.935
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	C[+57]JVPVC[+57]GVPR	CPVPCGVPR	2	522.2599
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	IIGGSDADIK	IIGGSDADIK	2	494.7691
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	LLEVPGR	LLEVPGR	2	456.7611
sp P09871 C1S_HUMAN	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	C1S	GDGGGAFVQDPNDK	GDGGGAFVQDPNDK	2	739.3313
sp P0C0L4 CO4A_HUMAN	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	C4A	NNVPC[+57]SPK	NNVPCSPK	2	458.2213
sp P0C0L4 CO4A_HUMAN	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	C4A	N[+1]NNVPC[+57]SPK	NNVPCSPK	2	458.7133
sp P0C0L4 CO4A_HUMAN	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	C4A	VDFTLSSER	VDFTLSSER	2	527.2642
sp P0C0L5 CO4B_HUMAN	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	C4B	C[+57]SVFYGAPSK	CSVFYGAPSK	2	558.2631
sp P0C0L5 CO4B_HUMAN	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	C4B	FAC[+57]YYPR	FACYPR	2	488.7209
sp P0C0L5 CO4B_HUMAN	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	C4B	VEYGVQK	VEYGVQK	2	485.2556
sp P0C0L5 CO4B_HUMAN	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	C4B	ITQVLHFTK	ITQVLHFTK	2	543.8189
sp P0C0L5 CO4B_HUMAN	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	C4B	AAC[+57]AQLNDFLQEYGTQGC[+57]QV	AACAQLNDFLQEYGTQGCQV	2	1137.002
sp P0CG47 UBB_HUMAN	Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1	UBB	TITLEVEPSDITENVK	TITLEVEPSDITENVK	2	894.4673
sp P0DJI8 SAA1_HUMAN	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	SAA1	EANYIGSDK	EANYIGSDK	2	498.7353
sp P0DJI8 SAA1_HUMAN	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	SAA1	EAN[+1]YIGSDK	EANYIGSDK	2	499.2273
sp P0DJI8 SAA1_HUMAN	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	SAA1	FFGHGAEDSLADQAANEWGR	FFGHGAEDSLADQAANEWGR	2	1089.485
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	GC[+57]PTEEGC[+57]GER	GCPTEECCGER	2	626.2401
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	SLVC[+57]JNGDSDC[+57]DEDSADEDR	SLVCNGDSDCEDSDADEDR	2	1079.892
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	SLVC[+57]JN[+1]GDSDC[+57]DEDSADEDR	SLVCNGDSDCEDSDADEDR	2	1080.384
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	SYTSHNEIHK	SYTSHNEIHK	2	685.8151
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	ELSHLPSLYDSAYR	ELSHLPSLYDSAYR	3	605.2967
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	QNDFNSVEEK	QNDFNSVEEK	2	605.2727
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	SSGWHFVVK	SSGWHFVVK	2	523.7745
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	ELENALK	ELENALK	2	408.7267
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	AASGTQNNVLR	AASGTQNNVLR	2	565.7993
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	YSAWAESVTLNPQVIK	YSAWAESVTLNPQVIK	2	903.4753
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	NVVYTC[+57]JNEGYSLIGNPVAR	NVVYTCNEGYSLIGNPVAR	2	1063.52
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	N[+1]YTLTGR	NYTLTGR	2	413.2087
sp P10643 C07_HUMAN	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	C7	DS[C][+57]JTLPASAEK	DSCTLPASAEK	2	589.7715
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	EIQNAVGVK	EIQNAVGVK	2	536.2933
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	TLLSNLEAK	TLLSNLEAK	2	559.3086
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	EDALN[+1]JETR	EDALNETR	2	474.7171
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	IDSLLENDR	IDSLLENDR	2	537.7749
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	ASSIDELFQDR	ASSIDELFQDR	2	697.3515
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	EILSVDC[+57]STNNPSQAK	EILSVDCSTNNPSQAK	2	881.9174
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	LANL[+1]LTQGEDQYLR	LANLTLQGEDQYLR	2	842.9125
sp P10909 CLUS_HUMAN	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	CLU	LFDSDPITVTPPVEVSR	LFDSDPITVTPPVEVSR	2	937.4989
sp P11226 MBL2_HUMAN	Mannose-binding protein C OS=Homo sapiens GN=MBL2 PE=1 SV=2	MBL2	EEAFLGITDEK	EEAFLGITDEK	2	626.3088
sp P12111 CO6A3_HUMAN	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=5	COL6A3	VAVVQYSR	VAVVQYSR	2	518.7747
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	LEC[+57]JNGENDC[+57]GDNSDER	LECNCGENDCGDNSDER	2	942.342
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	VPANLENVGFEVQTAEDDLK	VPANLENVGFEVQTAEDDLK	2	1094.542
				DLTSLGHNENQQGSSFGSSFVPIYSSK		
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	DLTSLGHNENQQGSSFGSSFVPIYSSK	SSK	3	1131.527
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	SEINHNNSAFK	SEINHNNSAFK	2	630.802
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	ALNHLPLEYNSALYSR	ALNHLPLEYNSALYSR	2	930.9838
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	SEYGAALAWEK	SEYGAALAWEK	2	612.7984
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	C[+57]LPDGTWR	CLPDGTWR	2	502.7346
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	IGESIELTC[+57]JK	IGESIELTCPK	2	623.821
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	YTC[+57]JQGNSWTPPISNSLT[C][+57]JEK	YTCQGNSWTPPISNSLTCEK	2	1172.023
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	QLEWGLER	QLEWGLER	2	515.7694
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	ESC[+57]JGVDTC[+57]JYDWEK	ESCGVDTCYDWEK	2	856.8138
sp P13671 CO6_HUMAN	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	C6	GGNQLYC[+57]JK	GGNQLYCVK	2	519.7555
	General transcription factor IIIF subunit 2 OS=Homo sapiens GN=GTTF2F2 PE=1 SV=2	GTTF2F2	DLVDITK	DLVDITK	2	402.2291
sp P13984 T2FB_HUMAN	L-selectin OS=Homo sapiens GN=SELL PE=1 SV=2	SELL	DN[+1]YTDLVAIQNK	DNYTDLVAIQNK	2	697.8435
sp P14151 LYAM1_HUMAN	L-selectin OS=Homo sapiens GN=SELL PE=1 SV=2	SELL	AEIEYLEK	AEIEYLEK	2	497.7582
sp P15169 CBPN_HUMAN	Carboxypeptidase N catalytic chain OS=Homo sapiens GN=CPN1 PE=1 SV=1	CPN1	IVQLQDTR	IVQLQDTR	2	543.3193
sp P15169 CBPN_HUMAN	Carboxypeptidase N catalytic chain OS=Homo sapiens GN=CPN1 PE=1 SV=1	CPN1	SIPQVSPVR	SIPQVSPVR	2	491.7876
sp P15924 DESP_HUMAN	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	DSP	QQIYAAEK	QQIYAAEK	2	475.7507
sp P18428 LBP_HUMAN	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	LBP	LAEGFPLPLK	LAEGFPLPLK	2	599.3657
sp P19823 ITI2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	ITI2	VQSTITSR	VQSTITSR	2	446.2483
sp P19823 ITI2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	ITI2	VVNSPQPNVVFDVQIPK	VVNSPQPNVVFDVQIPK	2	1061.568
sp P19823 ITI2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	ITI2	VQFELHYQEVK	VQFELHYQEVK	2	710.367
sp P19823 ITI2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	ITI2	IYLPQGR	IYLPQGR	2	423.7452
sp P19823 ITI2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	ITI2	TILDDLRL	TILDDLRL	2	423.24

sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	AEDHFSVIDFNQNI	AEDHFSVIDFNQNI	2	902.9343
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	NDLISATK	NDLISATK	2	431.2375
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	IQPSGGTNINEALLR	IQPSGGTNINEALLR	2	791.931
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	LSNENHGIAQR	LSNENHGIAQR	2	619.8155
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	IYGNQDTSSQLK	IYGNQDTSSQLK	2	677.3359
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	FYNQVSTPLLR	FYNQVSTPLLR	2	669.3642
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	HADPDFTR	HADPDFTR	2	479.7225
sp P19823 ITIH2_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	ITIH2	SLAPTAAAK	SLAPTAAAK	2	415.2425
sp P19827 ITIH1_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1	FAHYVVTSQVNTANEAR	FAHYVVTSQVNTANEAR	2	1003.508
sp P19827 ITIH1_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1	EVAFDLEIPK	EVAFDLEIPK	2	580.8135
sp P19827 ITIH1_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1	AAISGENAGLVR	AAISGENAGLVR	2	579.3173
sp P19827 ITIH1_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1	LDAQASFLPK	LDAQASFLPK	2	545.3006
sp P19827 ITIH1_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	ITIH1	VTYDVS	VTYDVS	2	420.2165
sp P20742 PZP_HUMAN	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	PZP	ISEITNIVSK	ISEITNIVSK	2	552.319
sp P20742 PZP_HUMAN	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	PZP	ATVLNVLPK	ATVLNVLPK	2	509.8002
sp P20742 PZP_HUMAN	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	PZP	YGAATFTR	YGAATFTR	2	443.7245
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	EWDN[+1]TTTEC[+57]R	EWDNTTTECR	2	656.7591
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	LGHC[+57]PDPVLVNGEFSSSGPVN[+1]VSDK	LGHC PDPVLVNGEFSSSGPVNVSDK	3	871.4148
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	LGHC[+57]PDPVLVN[+1]GEFSSSGPVN[+1]VSDK	LGHC PDPVLVNGEFSSSGPVNVSDK	3	871.7428
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	SQC[+57]LEDHTWAPFPIC[+57]K	SQCLEDHTWAPFPICK	3	695.9889
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	LIQEAQPECK[+57]EK	LIQEAQPECEK	2	721.3714
sp P20851 C4BPB_HUMAN	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	C4BPB	ALLAFQESK	ALLAFQESK	2	503.782
sp P21333 FLNA_HUMAN	Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4	FLNA	FGGEHVPNSPFQVTALAGDQPSVQPLR	FGGEHVPNSPFQVTALAGDQPSVQPLR	3	982.5017
sp P21333 FLNA_HUMAN	Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4	FLNA	ATC[+57]APQHGAPGPGPADASK	ATCAPQHGAPGPGPADASK	3	597.2811
sp P22352 GPX3_HUMAN	Glutathione peroxidase 3 OS=Homo sapiens GN=GPX3 PE=1 SV=2	GPX3	QEPGENSEILPTLK	QEPGENSEILPTLK	2	777.9041
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	AFGSNPN[+1]LTK	AFGSNPNLTK	2	525.2667
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	LSNNALSLPQGVFGK	LSNNALSLPQGVFGK	2	801.4359
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	DLEELVK	DLEELVK	2	423.2344
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	LYLGSNN[+1]LTALHPALFQN[+1]LSK	LYLGSNNLTALHPALFQNLK	3	772.746
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	LELLLSK	LELLLSK	2	451.7815
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	LLNIQTYC[+57]AGPAYLK	LLNIQTYCAGPAYLK	2	862.9556
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	QLVC[+57]PVTR	QLVCPVTR	2	486.7684
sp P22792 CPN2_HUMAN	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	CPN2	SQC[+57]TYSNPEGTVVLC[+57]DQAQC[+57]R	SQCTYSNPEGTVVLCDDQAQCR	3	848.7019
sp P23142 FBLN1_HUMAN	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	FBLN1	SQETGDDLVDGGLQETDK	SQETGDDLVDGGLQETDK	2	896.4158
sp P23142 FBLN1_HUMAN	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	FBLN1	IIIEVEEQEDPYLND	IIIEVEEQEDPYLND	2	995.9656
sp P23142 FBLN1_HUMAN	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	FBLN1	DSSC[+57]GTGYELTEDNSC[+57]K	DSSCGTGYELTEDNSCK	2	961.8726
sp P25311 ZA2G_HUMAN	Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	AZGP1	DYIEFNK	DYIEFNK	2	464.7242
sp P25311 ZA2G_HUMAN	Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	AZGP1	EIPAWVPFDPAQAQITK	EIPAWVPFDPAQAQITK	2	891.9749
sp P25311 ZA2G_HUMAN	Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	AZGP1	AGEVQEPELR	AGEVQEPELR	2	564.2882
sp P26927 HGFL_HUMAN	Hepatocyte growth factor-like protein OS=Homo sapiens GN=MST1 PE=1 SV=2	MST1	EFK[+57]DLPR	EFCDLPR	2	468.7158
sp P26927 HGFL_HUMAN	Hepatocyte growth factor-like protein OS=Homo sapiens GN=MST1 PE=1 SV=2	MST1	GTAN[+1]TTTAGVPC[+57]QR	GTANTTTAGVPCQR	2	717.8357
sp P27169 PON1_HUMAN	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	PON1	EVQPVELPNC[+57]NLVK	EVQPVELPNCNLVK	2	819.9296
sp P27169 PON1_HUMAN	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	PON1	SFNPNSPGK	SFNPNSPGK	2	474.2327
sp P27169 PON1_HUMAN	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	PON1	VYVIAELLAHK	VYVIAELLAHK	2	660.3715
sp P27169 PON1_HUMAN	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	PON1	IFFYDSENPPASEVLR	IFFYDSENPPASEVLR	2	942.4623
sp P27169 PON1_HUMAN	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	PON1	IQNILTEEPK	IQNILTEEPK	2	592.8297
sp P27348 1433T_HUMAN	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1	YWHAQ	NLLSVAYK	NLLSVAYK	2	454.266
sp P27918 PROP_HUMAN	Properdin OS=Homo sapiens GN=CFP PE=1 SV=2	CFP	SISC[+57]QIEIPGQQSR	SISCQIEIPGQQSR	2	745.3568
sp P27918 PROP_HUMAN	Properdin OS=Homo sapiens GN=CFP PE=1 SV=2	CFP	LC[+57]TPLLPK	LCTPLLPK	2	471.2781
sp P29622 KAIN_HUMAN	Kallistatin OS=Homo sapiens GN=SERPINA4 PE=1 SV=3	SERPINA4	IAPANADFAFR	IAPANADFAFR	2	596.8091
sp P29622 KAIN_HUMAN	Kallistatin OS=Homo sapiens GN=SERPINA4 PE=1 SV=3	SERPINA4	ALWEKPFISSR	ALWEKPFISSR	2	667.3668
sp P29622 KAIN_HUMAN	Kallistatin OS=Homo sapiens GN=SERPINA4 PE=1 SV=3	SERPINA4	LGFTDLFSK	LGFTDLFSK	2	514.2766
sp P29622 KAIN_HUMAN	Kallistatin OS=Homo sapiens GN=SERPINA4 PE=1 SV=3	SERPINA4	WADLSGITK	WADLSGITK	2	495.7664
sp P29622 KAIN_HUMAN	Kallistatin OS=Homo sapiens GN=SERPINA4 PE=1 SV=3	SERPINA4	FFSAQTNR	FFSAQTNR	2	485.7407
sp P35542 SAA4_HUMAN	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2	SAA4	GNYDAAQR	GNYDAAQR	2	447.7068
sp P35858 ALS_HUMAN	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	IGFALS	ELVLAGNR	ELVLAGNR	2	436.2534
sp P35858 ALS_HUMAN	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	IGFALS	ANVFVQLPR	ANVFVQLPR	2	522.3035
sp P35858 ALS_HUMAN	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	IGFALS	LEYLLSR	LEYLLSR	2	503.8002
sp P35858 ALS_HUMAN	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	IGFALS	DFALQNPSAVPR	DFALQNPSAVPR	2	657.8437
sp P35858 ALS_HUMAN	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	IGFALS	DLSEAHFAPC[+57]	DLSEAHFAPC	2	573.7479
sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	TESIHR	TESIHR	2	428.2378
sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	ALYDILSSPDIHGT	ALYDILSSPDIHGT	2	978.4911

sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	ELLDTVTAPQK	ELLDTVTAPQK	2	607.835
sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	SSFVAPLEK	SSFVAPLEK	2	489.2688
sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	TSLEDFYLDEER	TSLEDFYLDEER	2	758.8437
sp P36955 PEDF_HUMAN	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	SERPINF1	TVQAVLTVPK	TVQAVLTVPK	2	528.3266
sp P36980 FHR2_HUMAN	Complement factor H-related protein 2 OS=Homo sapiens GN=CFHR2 PE=1 SV=1	CFHR2	INHGLYDEEK	INHGLYDEEK	2	665.8355
sp P41222 PTGDS_HUMAN	Prostaglandin-H2 D-isomerase OS=Homo sapiens GN=PTGDS PE=1 SV=1	PTGDS	AQGFTEDTIVFLPQTDK	AQGFTEDTIVFLPQTDK	2	955.4807
sp P43251 BTD_HUMAN	Biotinidase OS=Homo sapiens GN=BTD PE=1 SV=2	BTD	SHLIAQVAK	SHLIAQVAK	2	540.3322
sp P43251 BTD_HUMAN	Biotinidase OS=Homo sapiens GN=BTD PE=1 SV=2	BTD	NPVGLIGAEN[+1]ATGETDPSHSK	NPVGLIGAENATGETDPSHSK	2	1048.001
sp P43251 BTD_HUMAN	Biotinidase OS=Homo sapiens GN=BTD PE=1 SV=2	BTD	LSSGLVTAALYGR	LSSGLVTAALYGR	2	654.3695
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	SDVGLFPFPTLDPEEK	SDVGLFPFPTLDPEEK	3	629.984
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	ESLLNHFLYEVAR	ESLLNHFLYEVAR	2	795.9174
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	AIPVTQYLK	AIPVTQYLK	2	516.808
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	AFSSYQK	AFSSYQK	2	415.7058
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	HVC[+57]GALLK	HVCGALLK	2	449.2524
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	GQC[+57]JNSNK	GQCINSNK	2	517.2584
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	FTDSENVC[+57]QER	FTDSENVCCQER	2	692.7935
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	DADPDTFFAK	DADPDTFFAK	2	563.7562
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	FTFEYSR	FTFEYSR	2	475.2243
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	N[+1]C[+57]JNTENPPG[C+57]YR	NCCNTENPPGCYR	2	821.8058
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	HFQNLGK	HFQNLGK	2	422.2272
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	IAPQLSTEELVSLGEK	IAPQLSTEELVSLGEK	2	857.4671
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	TINPAVDHC[+57]C[+57]K	TINPAVDHCCK	2	657.8001
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	FLVNLVK	FLVNLVK	2	416.7682
sp P43652 AFAM_HUMAN	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	AFM	AESPEVC[+57]FNEESPK	AESPEVCFNEESPK	2	811.8538
sp P49747 COMP_HUMAN	Cartilage oligomeric matrix protein OS=Homo sapiens GN=COMP PE=1 SV=2	COMP	LVPNPGQEDADR	LVPNPGQEDADR	2	655.8204
sp P49908 SEPP1_HUMAN	Selenoprotein P OS=Homo sapiens GN=SEPP1 PE=1 SV=3	SEPP1	EGYSNI[+1]SYIVVNHQGISR	EGYSNI[+1]SYIVVNHQGISR	3	708.6834
sp P49908 SEPP1_HUMAN	Selenoprotein P OS=Homo sapiens GN=SEPP1 PE=1 SV=3	SEPP1	DDFLYDR	DDFLYDR	2	528.7535
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	NNQIDHIDEK	NNQIDHIDEK	2	613.294
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	LHINHNN[+1]LTVESVGLPK	LHINHNNLTVESVGLPK	3	628.6706
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	SLEDLQLTHNK	SLEDLQLTHNK	2	649.341
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	ISNIPEYFK	ISNIPEYFK	2	613.3086
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	FNALQYLR	FNALQYLR	2	512.7824
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	NIPTVNIENLNYLEVNLQLEK	NIPTVNIENLNYLEVNLQLEK	3	846.0904
sp P51884 LUM_HUMAN	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	LUM	ILGLPSYSK	ILGLPSYSK	2	489.2869
sp P55056 APOC4_HUMAN	Apolipoprotein C-IV OS=Homo sapiens GN=APOC4 PE=1 SV=1	APOC4	ELLETVVNR	ELLETVVNR	2	536.8035
sp P55056 APOC4_HUMAN	Apolipoprotein C-IV OS=Homo sapiens GN=APOC4 PE=1 SV=1	APOC4	ELLETVVN[+1]R	ELLETVVNR	2	537.2955
sp P55056 APOC4_HUMAN	Apolipoprotein C-IV OS=Homo sapiens GN=APOC4 PE=1 SV=1	APOC4	AWFLESK	AWFLESK	2	440.7318
sp P55058 PLTP_HUMAN	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	PLTP	VSN[+1]VSC[+57]QASVSR	VSNVSCQASVSR	2	647.8064
sp P60174 TPIS_HUMAN	Triosephosphate isomerase OS=Homo sapiens GN=TP1 PE=1 SV=3	TP1	TATPQQAEQVHEK	TATPQQAEQVHEK	3	489.5793
sp P61626 LYSC_HUMAN	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1	LYZ	WESGYNTR	WESGYNTR	2	506.7278
sp P62328 TYB4_HUMAN	Thymosin beta-4 OS=Homo sapiens GN=TMSB4X PE=1 SV=2	TMSB4X	ETIEQEK	ETIEQEK	2	438.7191
sp P62937 PPIA_HUMAN	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	PPIA	FEDENFILK	FEDENFILK	2	577.79
sp P63104 14332_HUMAN	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	YWHAZ	FLIPNASQAESK	FLIPNASQAESK	2	652.8459
sp P67936 TPM4_HUMAN	Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4 PE=1 SV=3	TPM4	AEGDVAALNR	AEGDVAALNR	2	508.262
sp P68431 H3I_HUMAN	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2	HIST1H3A	STELLIR	STELLIR	2	416.2504
sp P68871 HBB_HUMAN	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2	HBB	SAYTALWKG	SAYTALWKG	2	466.7636
sp P68871 HBB_HUMAN	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2	HBB	GTFTATSELHC[+57]DK	GTFTATSELHCDK	2	739.8508
sp P68871 HBB_HUMAN	Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2	HBB	EFTFPVQAAYQK	EFTFPVQAAYQK	2	689.8537
sp P69905 HBA_HUMAN	Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2	HBA1	TYFPHFDLSHGSAQVK	TYFPHFDLSHGSAQVK	2	917.4496
sp P80108 PHLD_HUMAN	Phosphatidylinositol-glycan-specific phospholipase D OS=Homo sapiens GN=GPLD1 PE=1 SV=3	GPLD1	N[+1]LTSLTESVDR	NLTSLTESVDR	2	668.8332
sp P80108 PHLD_HUMAN	Phosphatidylinositol-glycan-specific phospholipase D OS=Homo sapiens GN=GPLD1 PE=1 SV=3	GPLD1	QVLLVGAPTYDDVSK	QVLLVGAPTYDDVSK	2	802.9301
sp P80108 PHLD_HUMAN	Phosphatidylinositol-glycan-specific phospholipase D OS=Homo sapiens GN=GPLD1 PE=1 SV=3	GPLD1	SWITPC[+57]PEEK	SWITPCPEEK	2	623.7923
sp Q02153 GCB1_HUMAN	Guanylate cyclase soluble subunit beta-1 OS=Homo sapiens GN=GUCY1B3 PE=1 SV=1	GUCY1B3	FDTLTDSR	FDTLTDSR	2	477.73
sp Q03591 FHR1_HUMAN	Complement factor H-related protein 1 OS=Homo sapiens GN=CFHR1 PE=1 SV=2	CFHR1	ITC[+57]TEEGWSPTPK	ITCTEEGWSPTPK	2	753.3507
sp Q03591 FHR1_HUMAN	Complement factor H-related protein 1 OS=Homo sapiens GN=CFHR1 PE=1 SV=2	CFHR1	TGESAEFVC[+57]K	TGESAEFVCK	2	564.2555
sp Q04756 HGFA_HUMAN	Hepatocyte growth factor activator OS=Homo sapiens GN=HGFA PE=1 SV=1	HGFAC	YEYLEGGDR	YEYLEGGDR	2	551.246
sp Q04756 HGFA_HUMAN	Hepatocyte growth factor activator OS=Homo sapiens GN=HGFA PE=1 SV=1	HGFAC	TWC[+57]EGTR	TWCEGTR	2	455.1978
sp Q04756 HGFA_HUMAN	Hepatocyte growth factor activator OS=Homo sapiens GN=HGFA PE=1 SV=1	HGFAC	DSALSWEYC[+57]R	DSALSWEYCR	2	643.7771
sp Q04756 HGFA_HUMAN	Hepatocyte growth factor activator OS=Homo sapiens GN=HGFA PE=1 SV=1	HGFAC	SQFVQPIC[+57]LPEPGSTFPAGHK	SQFVQPICLPEPGSTFPAGHK	3	766.3843
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	SLPEGVANGIEVYSTK	SLPEGVANGIEVYSTK	2	832.4305
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	SLPEGVAN[+1]GIEVYSTK	SLPEGVANGIEVYSTK	2	832.9225
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	EVSFDOVELPK	EVSFDOVELPK	2	581.8032
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	VTFELTYEELLK	VTFELTYEELLK	2	742.8978
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	GHVSFKPSLDQQR	GHVSFKPSLDQQR	2	749.8917
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	EHLVQATPENLQEAR	EHLVQATPENLQEAR	2	867.9421
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	DYIFGNVIER	DYIFGNVIER	2	645.3117
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	VSDIRPGSDPTKPDATLVVK	VSDIRPGSDPTKPDATLVVK	3	699.0513
sp Q06033 ITI3_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI3 PE=1 SV=2	ITI3	NHQLIVTR	NHQLIVTR	2	490.7854
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	DAGVVC[+57]JTN[+1]ETR	DAGVVCJTNETR	2	611.7721
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	IDITLSSVK	IDITLSSVK	2	488.2897
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	AVDTWSWGER	AVDTWSWGER	2	603.7805
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	ASHEEVEGLVEK	ASHEEVEGLVEK	2	663.8304

sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	GLN[+1]LTEDTYKPR	GLNLTEDTYKPR	2	704.3594
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	IYTSPTWSAFVTDSSWSAR	IYTSPTWSAFVTDSSWSAR	2	1081.513
sp Q08380 LG3BP_HUMAN	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	LGALS3BP	AAIPSAALDTN[+1]SSK	AAIPSAALDTNSSK	2	638.325
sp Q12805 FBLN3_HUMAN	EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 PE=1 SV=2	EFEMP1	ADQVC[+57]INLR	ADQVCINLR	2	544.7795
sp Q12805 FBLN3_HUMAN	EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 PE=1 SV=2	EFEMP1	GSFAC[+57]QC[+57]PPGYQK	GSFACQCPPGYQK	2	750.3239
sp Q13201 MMRN1_HUMAN	Multimerin-1 OS=Homo sapiens GN=MMRN1 PE=1 SV=3	MMRN1	FNPGAESVLSN[+1]STLK	FNPGAESVLSNSTLK	2	832.4305
sp Q13790 APOF_HUMAN	Apolipoprotein F OS=Homo sapiens GN=APOF PE=1 SV=2	APOF	SLPTEDC[+57]ENEK	SLPTEDCENEK	2	661.2825
sp Q14520 HABP2_HUMAN	Hyaluronan-binding protein 2 OS=Homo sapiens GN=HABP2 PE=1 SV=1	HABP2	FC[+57]EIGSDDC[+57]YVGDGYSYR	FCEIGSDDCYVGDGYSYR	2	1081.925
sp Q14520 HABP2_HUMAN	Hyaluronan-binding protein 2 OS=Homo sapiens GN=HABP2 PE=1 SV=1	HABP2	VVLGDQDLK	VVLGDQDLK	2	493.7795
sp Q14520 HABP2_HUMAN	Hyaluronan-binding protein 2 OS=Homo sapiens GN=HABP2 PE=1 SV=1	HABP2	DEIPHNDIALLK	DEIPHNDIALLK	2	689.3723
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	NGIDIYSLTVDSR	NGIDIYSLTVDSR	2	726.8701
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	FAHTVVTSR	FAHTVVTSR	2	509.2774
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	LGVYELLK	LGVYELLK	2	524.3261
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	VRPQQLVK	VRPQQLVK	2	484.306
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	FKPTLSQQQK	FKPTLSQQQK	2	602.8379
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	NVVFVIDK	NVVFVIDK	2	467.2738
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	ILDDLSPR	ILDDLSPR	2	464.7585
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	SIQNNVR	SIQNNVR	2	415.7276
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	LALDNGGLAR	LALDNGGLAR	2	500.2827
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	GPDLVTATVSGK	GPDLVTATVSGK	2	572.8141
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	NVHSGSTFFK	NVHSGSTFFK	2	562.2802
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	YYLQGAK	YYLQGAK	2	421.724
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	QGPVNLLSDPEQGVETGQYER	QGPVNLLSDPEQGVETGQYER	3	805.7311
sp Q14624 ITIH4_HUMAN	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	ITIH4	TGLLLLSDPK	TGLLLLSDPK	2	586.3321
sp Q15848 ADIPO_HUMAN	Adiponectin OS=Homo sapiens GN=ADIPOQ PE=1 SV=1	ADIPOQ	GDIGETGVPGAEGPR	GDIGETGVPGAEGPR	2	706.3442
sp Q15848 ADIPO_HUMAN	Adiponectin OS=Homo sapiens GN=ADIPOQ PE=1 SV=1	ADIPOQ	IFYNQNHVDGSGTGK	IFYNQNHVDGSGTGK	3	591.2727
sp Q16610 ECM1_HUMAN	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	ECM1	LLPAQLPAEK	LLPAQLPAEK	2	540.3266
sp Q16610 ECM1_HUMAN	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	ECM1	EVGPPLPQEAIVPLQK	EVGPPLPQEAIVPLQK	2	801.4485
sp Q16610 ECM1_HUMAN	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	ECM1	ELPSLQHPNEQK	ELPSLQHPNEQK	2	710.365
sp Q16610 ECM1_HUMAN	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	ECM1	FSC[+57]FQEEAPQPHYQLR	FSCFQEEAPQPHYQLR	3	679.6476
sp Q16610 ECM1_HUMAN	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	ECM1	NVALVSGDTENAK	NVALVSGDTENAK	2	659.3359
sp Q16635 TAZ_HUMAN	Tafazzin OS=Homo sapiens GN=TAZ PE=1 SV=1	TAZ	EVLYELIEK	EVLYELIEK	2	568.3159
sp Q571H1 EYS_HUMAN	Protein eyes shut homolog OS=Homo sapiens GN=EYS PE=1 SV=5	EYS	FLN[+1]FGIR	FLNFGIR	2	434.2398
sp Q577N2 LITD1_HUMAN	LINE-1 type transposase domain-containing protein 1 OS=Homo sapiens GN=LITD1 PE=1 SV=1	LITD1	DIAPVLDLK	DIAPVLDLK	2	492.2922
sp Q57CS8 KAD9_HUMAN	Adenylate kinase 9 OS=Homo sapiens GN=AK9 PE=1 SV=2	AK9	ITSEYGLK	ITSEYGLK	2	455.7477
sp Q8N3Z3 GTPB8_HUMAN	GTP-binding protein 8 OS=Homo sapiens GN=GTPB8 PE=2 SV=1	GTPB8	C[+57]FIASVTGSLD	CFIASVTGSLD	2	585.279
sp Q8NF91 SYNE1_HUMAN	Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=1 SV=4	SYNE1	LSEFSLK	LSEFSLK	2	468.7737
sp Q8TD26 CHD6_HUMAN	Chromodomain-helicase-DNA-binding protein 6 OS=Homo sapiens GN=CHD6 PE=1 SV=4	CHD6	AVLQDINR	AVLQDINR	2	464.7642
sp Q8WU4 ZN296_HUMAN	Zinc finger protein 296 OS=Homo sapiens GN=ZNF296 PE=2 SV=1	ZNF296	TLSSFSN[+1]LK	TLSSFSNLK	2	499.2637
sp Q92820 GGH_HUMAN	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2	GGH	SINGILFPGGSVDLR	SINGILFPGGSVDLR	2	772.9252
sp Q92820 GGH_HUMAN	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2	GGH	FFNLTTNTDGGK	FFNLTTNTDGGK	2	678.8433
sp Q92820 GGH_HUMAN	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2	GGH	NLDGISHAPNAVK	NLDGISHAPNAVK	2	668.3544
sp Q92954 PRG4_HUMAN	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	PRG4	C[+57]FESFER	CFESFER	2	487.7055
sp Q96PD5 PGRP2_HUMAN	N-acetylmuramoyl-L-alanine amidase OS=Homo sapiens GN=PGLYRP2 PE=1 SV=1	PGLYRP2	GC[+57]PDVQASLPDAK	GCPDVQASLPDAK	2	679.3245
sp Q96PD5 PGRP2_HUMAN	N-acetylmuramoyl-L-alanine amidase OS=Homo sapiens GN=PGLYRP2 PE=1 SV=1	PGLYRP2	TFTLLDPK	TFTLLDPK	2	467.7658
sp Q96PD5 PGRP2_HUMAN	N-acetylmuramoyl-L-alanine amidase OS=Homo sapiens GN=PGLYRP2 PE=1 SV=1	PGLYRP2	DTLPSC[+57]AVR	DTLPSCAVR	2	509.7529
sp Q98QI4 CCDC3_HUMAN	Coiled-coil domain-containing protein 3 OS=Homo sapiens GN=CCDC3 PE=2 SV=1	CCDC3	HLELANQK	HLELANQK	2	476.7642
sp Q98VG8 KIFC3_HUMAN	Kinesin-like protein KIFC3 OS=Homo sapiens GN=KIFC3 PE=1 SV=4	KIFC3	GVDC[+57]STGLR	GVDCSTGLR	2	482.7295
sp Q9C0F0 ASXL3_HUMAN	Putative Polycomb group protein ASXL3 OS=Homo sapiens GN=ASXL3 PE=2 SV=3	ASXL3	IEDDQSTR	IEDDQSTR	2	482.2225
sp Q9NZP8 C1RL_HUMAN	Complement C1r subcomponent-like protein OS=Homo sapiens GN=C1RL PE=1 SV=2	C1RL	GSEAINAPGDNPAK	GSEAINAPGDNPAK	2	670.8257
sp Q9UGM5 FETUB_HUMAN	Fetuin-B OS=Homo sapiens GN=FETUB PE=1 SV=2	FETUB	GSVQYLPDLDDK	GSVQYLPDLDDK	2	675.3328
sp Q9UK55 ZPI_HUMAN	Protein Z-dependent protease inhibitor OS=Homo sapiens GN=SERPINA10 PE=1 SV=1	SERPINA10	ETFFN[+1]LSK	ETFFNLSK	2	493.7451
sp Q9UK55 ZPI_HUMAN	Protein Z-dependent protease inhibitor OS=Homo sapiens GN=SERPINA10 PE=1 SV=1	SERPINA10	TVIEVDER	TVIEVDER	2	480.7535
sp Q9ULV4 COR1C_HUMAN	Coronin-1C OS=Homo sapiens GN=COR1C PE=1 SV=1	COR1C	NDQC[+57]YDDIR	NDQCYDDIR	2	599.7433
sp Q9Y2M2 SSUH2_HUMAN	Protein SSUH2 homolog OS=Homo sapiens GN=SSUH2 PE=1 SV=1	SSUH2	DISLASQR	DISLASQR	2	445.2405
sp Q9Y5Y7 LYVE1_HUMAN	Lymphatic vessel endothelial hyaluronin acid receptor 1 OS=Homo sapiens GN=LYVE1 PE=1 SV=2	LYVE1	DQVETALK	DQVETALK	2	452.2427