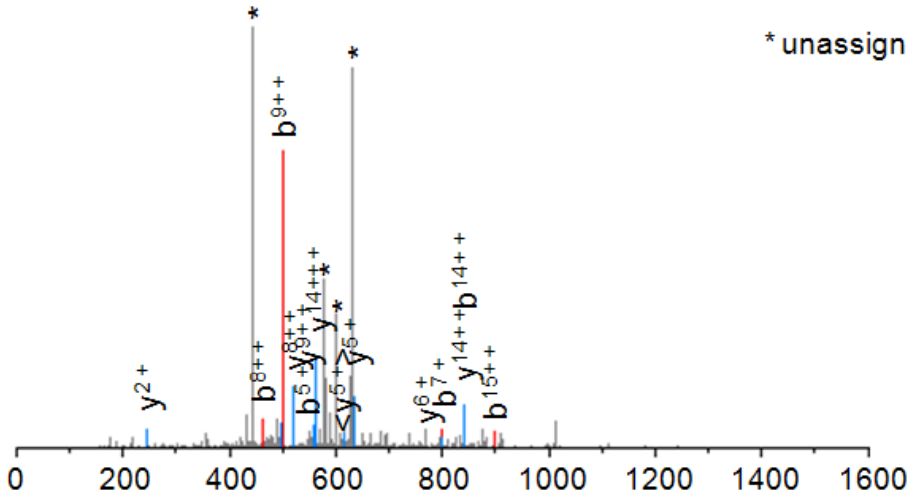
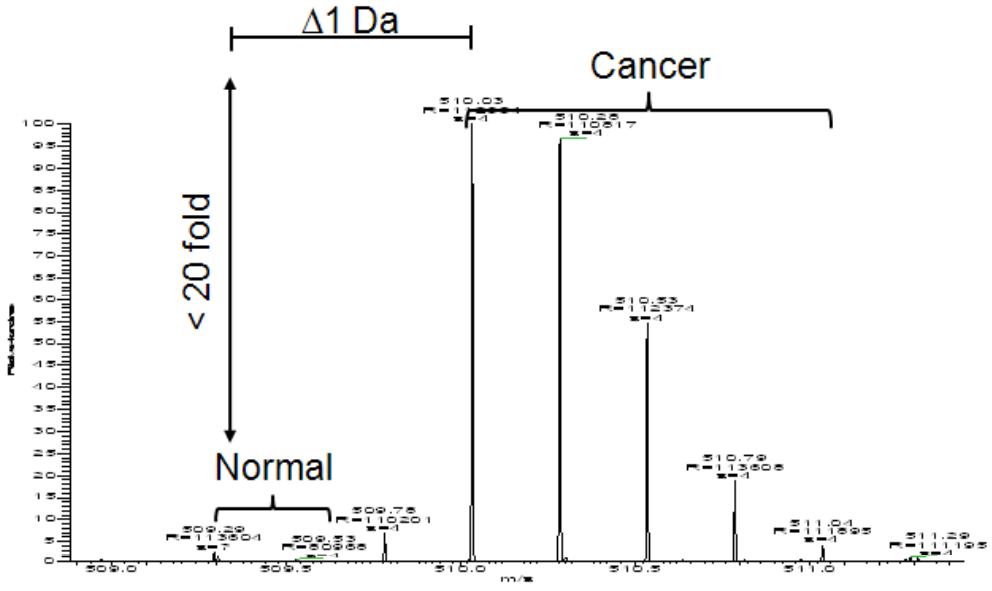
A

Probability	Precursor	z	MH+	ΔCn	Xcorr	Peptide
0.9815	51.03	4	2037.11	0.257	3.307	M.AAAPTQIEAELYYLIAR.F



B



Supplemental Figure 1. Identification and Quantification of the BRWD3. (A) MS analysis output identifying BRWD3 protein. MH+, theoretical mass for the 4^+ charged molecular ion; ΔCn, delta correlation; Xcorr, cross-correlation score. Tandem mass spectrum for the selected precursor ion is shown below. Spectrum was generated using the Trans Proteomic Pipeline. (B) Quantification of BRW1 through the mTRAQ-labeled parent ion signal of peptide * AAAPTQIEAELYYLIAR.

Supplemental Table 1. List of peptides identified by mTRAQ and LC-MSMS.

entry no.	accession number[a]	probability	coverage	protein ratio (L/H)	SD[b]	XNP[c]	USP[b]	protein name	peptide sequence	z[c]	m/z	Xcorr	deltCn
1	IPI00001610	1	11.1	0.67	0	2	2	IGF1 Insulin-like growth factor IA precursor	GFYFNKPTGYGSSSR	3	649.9953	3.596	0.345
1	IPI00001610	1	11.1	0.67	0	2	2	IGF1 Insulin-like growth factor IA precursor	n[145]GFYFNK[272]PTGYGSSSR	3	652.6667	3.664	0.316
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	LTFINDLCGPR	2	717.8652	3.775	0.428
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	n[145]LTFINDLCGPR	2	719.8688	2.648	0.339
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	LLPAQLPAEK	2	680.4216	2.905	0.344
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	n[145]RAPYPNYDR	3	432.5586	1.928	0.11
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	VTPNLM[147]GH	2	512.7659	1.909	0.056
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	n[145]VTPNLM[147]GH	2	514.7694	2.066	0.244
2	IPI00003351	1	9.2	1.07	0.11	11	6	ECM1 Extracellular matrix protein 1 precursor	AQLPAEK	2	518.8112	1.728	0.078
3	IPI00003590	1	6.6	0.86	0.1	4	4	QSOX1 Isoform 1 of Sulfhydryl oxidase 1 precursor	n[145]AHFSPSNIILDFPAAGSAAR	3	729.3873	2.878	0.298
3	IPI00003590	1	6.6	0.86	0.1	4	4	QSOX1 Isoform 1 of Sulfhydryl oxidase 1 precursor	n[145]SFYTAYLQR	2	646.8414	2.619	0.297
3	IPI00003590	1	6.6	0.86	0.1	4	4	QSOX1 Isoform 1 of Sulfhydryl oxidase 1 precursor	n[145]AAPGQEPPEHM[147]AELQR	3	640.9833	2.989	0.28
3	IPI00003590	1	6.6	0.86	0.1	4	4	QSOX1 Isoform 1 of Sulfhydryl oxidase 1 precursor	SFYTAYLQR	2	644.8379	2.715	0.067
3	IPI00003590	1	6.6	0.86	0.1	4	4	QSOX1 Isoform 1 of Sulfhydryl oxidase 1 precursor	YILR	2	352.7263	1.622	0.012
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	LQSLFDSPDFSK	2	832.4382	4.518	0.5
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]AAVSNFGYDLYR	2	760.3867	4.243	0.373
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]LAAAVSNFGYDLYR	2	852.4473	3.584	0.392
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]LQSLFDSPDFSK[272]	2	836.4453	3.93	0.263
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	ALYYDLISSPDIHGTYK	3	746.0598	5.58	0.49
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]ALYYDLISSPDIHGTYK[272]	3	748.7312	5.187	0.494
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]EIPDEISILLLGVAHFK[272]	3	728.0957	4.175	0.288
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]ELLDTVTAPQK[272]	2	751.9371	3.631	0.386
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]YGLDSDLSCK[272]	2	717.846	3.359	0.407
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	LSYEGEVTK	2	653.3561	2.722	0.283
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]TESIIHR	2	500.2888	2.847	0.214
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]ITGK[272]PIK[272]	2	594.9057	2.731	0.22
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]SSFVAPLEK[272]	2	633.3708	2.712	0.258
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]VLTGNPR	2	450.7728	2.513	0.173
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	KTSLEDFYLDEER	3	642.3266	2.918	0.173
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	YGLDSDLSCK	2	713.8389	2.687	0.295
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]LTQVEHR	2	513.7943	2.447	0.133
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	ITGKPIK	2	588.895	2.338	0.096
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]LSYEGEVTK[272]	2	657.3632	1.801	0.185
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	ELLDTVTAPQK	2	747.93	2.164	0.129
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]ITGK[272]PIK[272]	3	396.9395	1.769	0.072
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	n[145]SLQEM[147]K[272]	2	520.2885	1.641	0.043
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	ALYYDLISSPDIHGTYK	4	559.7967	1.84	0.044
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	AAVSNFGYDLYR	2	758.3832	2.749	0.206
4	IPI00006114	1	34.9	1.04	0.27	35	23	SERPINF1 Pigment epithelium-derived factor precursor	EM[147]K	1	703.3808	1.121	0.072
5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	n[145]SFFSFLGEAFDGAR	2	847.9184	4.657	0.449
5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	n[145]EANYIGSDK[272]	2	642.8374	2.886	0.26
5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	EANYIGSDK	2	638.8303	2.4	0.303
5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	n[145]YFHAR	2	419.228	2.15	0.188

5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	n[145]DPNHFRPAGLPEK[272]Y	3	643.6789	2.891	0.231
5	IPI00006146	1	34.4	0.7	0.12	6	6	SAA1;SAA2 serum amyloid A2	DPNHFRPAGLPEKY	3	641.0075	2.372	0.113
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	CPNPPVQENFDVNK	2	963.9637	3.832	0.473
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	NILTSNNIDVK	2	755.9331	4.21	0.429
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]NILTSNNIDVK[272]	2	759.9402	4.216	0.4
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	KM[147]TVTDQVNCPK	3	615.9843	3.913	0.297
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	M[147]TVTDQVNCPK	2	789.3779	3.903	0.392
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]NPNLPPETVDSLK[272]	2	856.4771	3.284	0.307
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]CPNPPVQENFDVNK[272]	2	967.9708	3.259	0.347
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	VLNQELR	2	506.3009	2.496	0.115
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]WYEIEK[272]	2	578.3181	2.223	0.1
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]VLNQELR	2	508.3045	2.44	0.041
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	NPNLPPETVDSLK	2	852.47	1.97	0.038
6	IPI00006662	1	33.3	0.88	0.06	18	12	APOD Apolipoprotein D precursor	n[145]NILTSNNIDVK[272]	3	506.9625	2.031	0.127
7	IPI00007047	1	20.4	1.6	0.13	5	5	S100A8 Protein S100-A8	ALNSIIDVYHK	2	776.946	3.934	0.421
7	IPI00007047	1	20.4	1.6	0.13	5	5	S100A8 Protein S100-A8	ALNSIIDVYHK	3	518.2997	4.032	0.386
7	IPI00007047	1	20.4	1.6	0.13	5	5	S100A8 Protein S100-A8	n[145]ALNSIIDVYHK[272]	3	520.9711	4.011	0.374
7	IPI00007047	1	20.4	1.6	0.13	5	5	S100A8 Protein S100-A8	GNFHAVYR	3	368.5297	2.206	0.166
7	IPI00007047	1	20.4	1.6	0.13	5	5	S100A8 Protein S100-A8	GNFHAVYR	2	552.2909	2.483	0.134
8	IPI00007199	1	9.3	0.65	0.17	6	4	SERPINA10 Protein Z-dependent protease inhibitor precursor	n[145]LILVDYILFK[272]	2	762.9858	3.212	0.369
8	IPI00007199	1	9.3	0.65	0.17	6	4	SERPINA10 Protein Z-dependent protease inhibitor precursor	n[145]IFSPFADLSELSATGR	2	927.9896	2.585	0.199
8	IPI00007199	1	9.3	0.65	0.17	6	4	SERPINA10 Protein Z-dependent protease inhibitor precursor	n[145]ETSNFGFSLLR	2	707.876	2.497	0.147
8	IPI00007199	1	9.3	0.65	0.17	6	4	SERPINA10 Protein Z-dependent protease inhibitor precursor	n[145]TVIEVDER	2	552.8045	1.25	0.029
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	GFQQLLQELNQPR	2	855.9679	5.238	0.414
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	n[145]GFQQLLQELNQPR	2	857.9715	4.931	0.381
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	RVEDLHVGATVAPSSR	3	612.0025	2.933	0.329
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	AVVEVDESGTR	2	651.3384	3.449	0.453
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	n[145]AVVEVDESGTR	2	653.342	3.374	0.37
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	AAAATGTIFTFR	2	683.8775	3.196	0.219
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	n[145]AAAATGTIFTFR	2	685.8811	3.447	0.221
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	n[145]TLYLADTFPTNFR	2	851.9497	3.18	0.281
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	n[145]GFQQLLQELNQPR	3	572.3167	2.136	0.002
9	IPI00007221	1	16.7	0.96	0.09	11	9	SERPINA5 Plasma serine protease inhibitor precursor	EM[147]K	1	703.3808	1.121	0.072
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	IAQYYYTFK	2	738.898	3.914	0.425
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	VLHGDLIDFVCK	2	842.9493	4.249	0.525
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]CFDHHFLEGSR	2	769.3435	3.646	0.516
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]IAQYYYTFK[272]	2	742.9051	4.056	0.465
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]VLHGDLIDFVCK[272]	2	846.9564	4.002	0.537
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]SGYLLHGSNEITCNR	3	618.63	4.277	0.344
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	CFDHHFLEGSR	4	384.1736	2.815	0.434
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	LIENGYFHPVK	3	532.9681	2.81	0.19
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	VLHGDLIDFVCK	3	562.302	3.503	0.331
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]LIENGYFHPVK[272]	3	535.6395	3.69	0.277
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]VACEEPPFIENGAANLHSK[272]	3	787.3921	4.793	0.301
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]CFDHHFLEGSR	4	385.1754	2.441	0.332
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]CNEYYLLR	2	632.3028	3.248	0.326
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	IQTHSTTYR	2	623.8306	2.287	0.23
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	CFDHHFLEGSR	3	511.8957	2.276	0.161

10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]IQTHSTTYR	2	625.8341	2.323	0.167
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]IQTHSTTYR	3	417.5585	1.855	0.226
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	CNEYYLLR	2	630.2992	2.155	0.069
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	n[145]CFDHHFLEGSR	3	513.2314	1.658	0.109
10	IPI00007240	1	14.7	0.96	0.07	21	19	F13B Coagulation factor XIII B chain precursor	EM[147]K	1	703.3808	1.121	0.072
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	n[145]GGISGYTLR	2	534.3019	3.265	0.252
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	QFPSLEHR	2	577.3093	2.464	0.098
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	EIIIHDQYK	3	480.2733	3.268	0.161
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	n[145]DIAILK[272]	2	480.8203	1.803	1
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	n[145]CQFFTY	1	998.4233	1.511	0.238
11	IPI00008556	1	6.7	0.48	0.22	3	3	F11 Isoform 1 of Coagulation factor XI precursor	SVAK	2	342.7238	0.841	0.121
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	EQQALQTVCLK	2	793.9233	5.152	0.437
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	n[145]EQQALQTVCLK[272]	2	797.9304	4.53	0.378
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	n[145]TFHEASEDCISR	3	528.9016	4.097	0.422
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	n[145]SRLDTLAQEVALLK[272]	3	615.7065	4.04	0.384
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	n[145]LDTLAQEVALLK[272]	2	801.4895	4.139	0.302
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	SRLDTLAQEVALLK	3	613.0351	3.464	0.296
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	n[145]CFLAFTQTK[272]	2	696.8666	3.449	0.254
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	CFLAFTQTK	2	692.8595	3.532	0.262
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	LDTLAQEVALLK	2	797.4824	3.314	0.254
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	KDVVNTK	2	612.3772	2.602	0.214
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	KIVNAK	3	364.9133	2.582	0.08
12	IPI00009028	1	29.2	1.08	0.19	13	12	CLEC3B Tetranectin precursor	KIVNAK	2	546.8663	2.146	0.057
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	DLHLSDVFLK	2	733.922	4.234	0.494
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	HEGSFIQGAEK	2	741.8887	4.098	0.429
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	YYQENFCEQICSK	2	1013.9301	4.17	0.549
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]DLHLSDVFLK[272]	2	737.9291	4.211	0.414
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ENPAVIDFELAPIVDLVR	2	1077.6	5.652	0.409
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]HEGSFIQGAEK[272]	2	745.8958	4.347	0.466
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]SENINHNSAFK[272]	2	774.9041	4.999	0.412
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]YYQENFCEQICSK[272]	2	1017.9372	4.316	0.449
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	DLHLSDVFLK	3	489.6171	3.611	0.331
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ALNHLPLEYNSALYSR	3	669.0256	4.9	0.401
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]DLHLSDVFLK[272]	3	492.2885	3.882	0.296
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]VPANLENVGFEVQTAEDDLK[272]TDFYK[272]	3	1092.233	5.806	0.559
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	HEGSFIQGAEK	3	494.9282	3.895	0.226
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	SENINHNSAFK	3	514.2671	2.983	0.325
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]AK[272]DLHLSDVFLK[272]	3	606.6999	3.397	0.319
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]LSEK[272]HEGSFIQGAEK[272]	3	698.0515	3.777	0.372
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]SENINHNSAFK[272]	3	516.9385	2.729	0.367
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]CLNNQQLHFLHIGSCQDGR	4	605.5307	3.913	0.39
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ALNHLPLEYNSALYSR	2	1003.0348	3.729	0.379
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]TLNICEVGTIR	2	704.8741	3.751	0.337
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	NSGLTEEEAK	2	679.3516	3.814	0.382
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ECNNPAPQR	2	609.7774	2.691	0.29
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]GFVVAGPSR	2	517.2992	3.228	0.287
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]NSGLTEEEAK[272]	2	683.3587	2.814	0.359
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]QAIQASHK[272]K[272]	3	481.6317	2.91	0.197

13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	IEEADCK	2	567.2701	2.4	0.291
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]RSENINHNSAFK[272]	3	568.9722	2.665	0.176
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]GEVLDNSFTGGICK[272]	2	887.4413	2.486	0.254
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	KM[147]EILHPGK	2	744.9397	2.213	0.218
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	NIPCAVTK	2	586.32	2.138	0.186
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ALQEYAAK[272]	2	591.3421	2.897	0.054
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	ALQEYAAK	2	587.335	2.578	0.071
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	QAIQASHKK	3	477.6246	1.886	0.19
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	AKDLHLSDVFLK	3	602.6928	3.507	0.132
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]HEGSFIQGAEK[272]	3	497.5996	2.954	0.081
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]TFSEWLESVK[272]	2	757.4107	2.535	0.277
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]FLAEK[272]	2	448.2782	1.72	0.009
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]ENPAVIDFELAPIVDLVR	3	718.7357	2.892	0.109
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	KM[147]EILHPGK	4	372.9735	1.877	0.018
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]IEEADCK[272]	2	571.2772	2.243	0.216
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	KALQEYAAK	3	481.2891	1.659	0.022
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	M[147]EILHPGK	3	407.5655	1.668	0.003
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	n[145]CLNNQQLH	2	580.2771	1.81	0.17
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	SVK[272]	1	617.4104	0.88	0.378
13	IPI00009920	1	26	0.7	0.19	54	43	C6 Complement component C6 precursor	EM[147]K	1	703.3808	1.121	0.072
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	EALIQFLEQVHQGIK	3	678.3896	4.126	0.359
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]EALIQFLEQVHQGIK[272]	3	681.061	5.373	0.494
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]IHILPSM[147]NPDGYEVAAAQGPNK[272]PGYLV	3	1123.5936	4.63	0.469
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]IHILPSM[147]NPDGYEVAAAQGPNK[272]PGYLV	4	842.9471	4.7	0.381
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]YVGNMHGNEALGR	3	521.2602	3.553	0.383
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]YVGNM[147]HGNEALGR	3	526.5919	2.084	0.437
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	IHILPSM[147]NPDGYEVAAAQGPNKPGYLVGR	4	840.9435	3.511	0.213
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]IVQLIQDTR	2	615.3704	2.994	0.316
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]VQNECPGITR	2	653.8218	2.757	0.282
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]TASTPTPDDK[272]	2	660.8479	2.902	0.187
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	TASTPTPDDK	2	656.8408	2.108	0.109
14	IPI00010295	1	20.1	0.73	0.08	18	12	CPN1 Carboxypeptidase N catalytic chain precursor	n[145]VYSIGR	2	419.7488	1.533	0.016
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	AIDEDCSQYEPIPGSQK	2	1103.5114	4.78	0.542
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	INVGGGLSGDHCK	2	791.8951	4.267	0.545
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]AIDEDCSQYEPIPGSQK[272]	2	1107.5185	4.919	0.605
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]ALDQYLMEFNACR	2	882.4054	4.425	0.511
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]INVGGGLSGDHCK[272]	2	795.9022	3.689	0.511
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	INVGGGLSGDHCK	3	528.2658	4.191	0.402
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]INVGGGLSGDHCK[272]	3	530.9372	5.132	0.511
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]YNPVVIDFEMQPIHEVLR	3	781.7478	5.587	0.509
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]YNPVVIDFEM[147]QPIHEVLR	3	787.0794	6.193	0.541
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	HTSLGPLEAK	2	666.8854	3.59	0.411
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]HTSLGPLEAK[272]	2	670.8925	3.337	0.362
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]AMAVEDIISR	2	624.8406	2.86	0.308
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	KPYNFLK	3	443.9396	2.368	0.206
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]STITYR	2	442.7516	2.224	0.154
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	LYYGDDEK	2	641.8194	2.752	0.123
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]YDSTCER	2	532.2189	1.766	0.201

15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	KFGGGK	3	338.5467	1.807	0.139
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]DITTCFGGSLGIQYEDKINVGGGLSGDHCK	5	720.9473	2.521	0.037
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]AM[147]AVEDIISR	2	632.838	2.073	0.053
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	KVQTQAC	2	552.2887	1.203	0.192
15	IPI00011252	1	24.7	0.56	0.21	23	21	C8A Complement component C8 alpha chain precursor	n[145]FIFTR	2	414.2484	1.442	0.099
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]SLPVSDSVLSGFEQR	2	882.9661	4.493	0.474
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]AEATTLHVAPQGTAM[147]AVSTFR	3	773.7343	4.599	0.433
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	VQEAHLTEDQIFYFPK	3	749.0599	3.76	0.345
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]FLQEQGHR	2	579.8105	2.707	0.286
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]FLQEQGHR	3	386.8761	3.312	0.151
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]QLYGDTGVLGR	2	661.8629	3.441	0.272
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]AGQLSVK[272]	2	495.813	2.803	0.206
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	AGQLSVK	2	491.8059	2.498	0.189
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]FLLQAR	2	446.2803	2.19	0.063
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	n[145]VQEAHLTEDQIFYFPK[272]	3	751.7313	2.73	0.064
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	SLPVSDSVLSGFEQR	2	880.9625	1.789	0.019
16	IPI00011261	1	41.6	0.78	0.04	16	11	C8G Complement component C8 gamma chain precursor	SVK[272]	1	617.4104	0.88	0.378
17	IPI00012792	1	4.5	0.98	0.08	3	3	CDH5 Cadherin-5 precursor	n[145]ELDSTGTPTGK[272]	2	697.3743	3.478	0.294
17	IPI00012792	1	4.5	0.98	0.08	3	3	CDH5 Cadherin-5 precursor	KGDIYNEK	2	693.8907	2.091	0.142
17	IPI00012792	1	4.5	0.98	0.08	3	3	CDH5 Cadherin-5 precursor	n[145]VHFLPVVISDNGM[147]PSR	3	643.3442	2.225	0.11
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	EYTDASFTNR	1	1343.6227	2.585	0.425
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]EYTDASFTNR	1	1347.6298	2.886	0.373
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	AGLQAFFQVQECNK	2	954.9766	5.013	0.516
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIASGLIGPLIICK	2	870.0016	4.769	0.492
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIFTGLIGPMK	2	736.4207	4.43	0.413
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DLYSGLIGPLIVCR	2	852.9624	3.337	0.49
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	EVGPTNADPVCLAK	2	870.4446	3.919	0.435
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	MYYSAVDPTK	2	727.8711	3.877	0.443
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VDKDNEDFQESNR	2	938.4452	4.66	0.522
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VNKDDEEFIESNK	2	994.0102	5.162	0.467
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AGLQAFFQVQECNK[272]	2	958.9837	5.545	0.541
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]ALYLQYTDETFR	2	832.4261	3.721	0.512
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIASGLIGPLIICK[272]	2	874.0087	4.485	0.49
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIFTGLIGPMK[272]	2	740.4278	4.336	0.409
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DLYSGLIGPLIVCR	2	854.966	3.029	0.355
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]MYYSAVDPTK[272]	2	731.8782	4.224	0.465
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NNEGTYYSPNYNPQSR	2	1024.465	5.688	0.516
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]RDTANLFPQTSLTLH	2	929.5006	3.92	0.46
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VDK[272]DNEDFQESNR	2	942.4523	3.751	0.406
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VNK[272]DDEEFIESNK[272]	2	1000.0209	4.14	0.332
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	AEEEHLGILGPQLHADVGDK	3	803.4238	5.476	0.478
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	AGLQAFFQVQECNK	3	636.9868	3.863	0.358
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KAEEEHLGILGPQLHADVGDK	3	892.8205	4.517	0.442
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KALYLQYTDETFR	3	643.3474	3.777	0.396
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KLISVDTEHSNIYLQNGPDR	3	860.4574	6.174	0.526
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	MFTTAPDQVDKEDEDFQESNK	3	965.4553	5.166	0.49
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	M[147]FTTAPDQVDKEDEDFQESNK	3	970.7869	5.616	0.516
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	NLASRPYTFHSHGITYYK	3	812.4255	4.636	0.472

18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	SVPPSASHVAPTETFTYEWTVPK	3	937.8166	6.12	0.662
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYCSEPEKVDKDNEDFQESNR	3	1000.4552	4.419	0.491
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VNKDDEEFIESNK	3	663.0092	4.565	0.45
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AEEEHLGILGPQLHADVGDK[272]	3	806.0952	6.769	0.495
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AGLQAFFQVQECNK[272]	3	639.6582	3.28	0.395
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIASGLIGPLIICK[272]	3	583.0082	2.904	0.173
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]MFTTAPDQVDK[272]EDEDFQESNK[272]	3	969.4624	5.008	0.526
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NLASRPYTFHSHGITYYK[272]	3	815.0969	4.896	0.459
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]SVPPSASHVAPTETFTYEWTVPK[272]	3	940.488	6.565	0.634
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TYCSEPEK[272]VDK[272]DNEDFQESNR	3	1004.4623	6.265	0.495
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VNK[272]DDEEFIESNK[272]	3	667.0163	3.676	0.397
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VTFHNK[272]GAYPLSIEPIGVR	3	796.121	4.715	0.564
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KAEEEHLGILGPQLHADVGDK	4	669.8672	3.542	0.342
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AEEEHLGILGPQLHADVGDK[272]	4	604.8232	4.761	0.438
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NLASRPYTFHSHGITYYK[272]	4	611.5745	3.725	0.358
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]GAYPLSIEPIGVR	2	758.4362	3.656	0.34
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	AETGDKVYVHLK	3	594.0087	3.513	0.355
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIASGLIGPLIICK	3	580.3368	2.684	0.121
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	SVPPSASHVAPTETFTYEWTVPK[272]	3	939.1523	4.512	0.349
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VTFHNKGAYPLSIEPIGVR	3	793.4496	3.82	0.393
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AETGDK[272]VYVHLK[272]	3	598.0158	3.296	0.334
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]GAYPLSIEPIGVR	3	505.9599	2.863	0.282
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]GPEEEHLGILGPVIWAEVGDTIR	3	877.8015	3.002	0.332
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]IYHSHIDAPK[272]	3	490.2766	3.862	0.02
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VDK[272]DNEDFQESNR	3	628.6373	4.32	0.331
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYCSEPEKVDKDNEDFQESNR	4	750.5932	4.365	0.399
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]AETGDK[272]VYVHLK[272]	4	448.7637	3.575	0.358
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIFTGLIGPM[147]K	2	744.4182	3.409	0.336
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KALYLQYTDETFR	2	964.5175	4.059	0.383
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIFTGLIGPM[147]K[272]	2	748.4253	4.166	0.363
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]EYTDASFTNR	2	674.3185	2.764	0.382
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	IYHSHIDAPK	3	487.6052	3.34	0.076
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]GVYSSDVFDIFPGTYQTLEM[147]FPR	3	943.7903	3.045	0.315
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	AETGDKVYVHLK	4	445.7584	2.523	0.293
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	M[147]FTTAPDQVDKEDEDFQESNK	4	728.342	2.396	0.097
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	GEFYIGSK	2	590.8217	3.179	0.315
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	M[147]YYSAVDPTK	2	735.8685	2.995	0.247
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]EVGPTNADPVCLAK[272]	2	874.4517	4.546	0.324
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]GEFYIGSK[272]	2	594.8288	2.933	0.28
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TYSDHPEK[272]	2	632.8243	3.239	0.338
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]YTVNQCR	2	537.2531	1.911	0.255
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYSDHPEK	2	628.8172	2.564	0.326
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]IYHSHIDAPK[272]	2	734.9112	3.692	0.185
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIFTGLIGPM[147]K	3	496.6145	3.064	0.044
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	MYYSAVDPTKDIFTGLIGPM[147]K	4	696.6183	3.055	0.23
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TYCSEPEK[272]VDK[272]DNEDFQESNR	4	753.5986	1.897	0.164
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]YTVNQCR	1	1073.4989	2.636	0.169
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	IYHSHIDAPK	2	730.9041	3.861	0.22

18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]QYTDSTFR	2	581.2865	1.983	0.297
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KDSLDKEK	3	508.3033	2.53	0.259
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	M[147]YYSAVDPTKDIFTGLIGPM[147]K	3	933.8202	4.006	0.312
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]EVGPTNADPVCLAK[272]	3	583.3036	2.902	0.181
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VTFHNK[272]	2	517.3053	2.42	0.186
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NM[147]ATRPYSIHAH	3	519.9286	2.509	0.25
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	YTVNQCR	2	535.2495	2.191	0.256
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIFTGLIGPMK[272]	3	493.9543	2.965	0.117
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VYVHLK[272]	2	523.8337	2.172	0.259
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYCSEPEK	2	641.794	2.418	0.209
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYSDHPEK	3	419.5472	1.804	0.068
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]QSEDSTFYLGER	2	788.374	2.66	0.386
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VDKDNEDFQESNR	3	625.9659	2.345	0.272
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	VTFHNK	2	513.2982	2.163	0.125
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KAEEEHLGILGPQLHADVGDK	5	536.0952	3.021	0.217
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DIFTGLIGPM[147]K[272]	3	499.2859	2.657	0.131
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]RPYLK[272]	2	482.8128	1.743	0.175
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KDSLDKEK	4	381.4793	1.977	0.197
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	QYTDSTFR	2	579.2829	2.208	0.098
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DIFTGLIGPMK	3	491.2829	2.325	0.133
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VTFHNK[272]GAYPLSIEPIGVR	4	597.3426	2.535	0.188
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KDSLDK	3	375.8924	2.164	0.083
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	EYTDASFTNRK	3	537.9424	1.753	0.172
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VTFHNK[272]	3	345.206	2.146	0.092
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TYYSPNYNPQSR	2	817.39	3.69	0.377
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]YAPSGIDIFTK[272]	2	750.4211	3.793	0.35
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DSLDKEK	2	627.8563	2.613	0.143
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]IGGSYK[272]K[272]	2	592.8718	1.805	0.055
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	IGGSYKK	2	586.8612	1.853	0.042
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KDSLDKEK	2	761.9513	2.708	0.09
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DSLDK[272]EK[272]	2	633.867	2.197	0.176
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]YPLSIEPIGVR	2	694.4069	2.681	0.385
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DLYSGLIGPLIVCR	3	568.9774	2.662	0.164
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KVVYR	3	315.5395	1.253	0.178
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DLYSGLIGPLIVCR	3	570.3131	2.357	0.122
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	DTANLFPQTSLTLH	2	849.4465	2.771	0.209
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TPGIWLLH	2	538.8162	2.982	0.218
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TTIEK[272]PVWLGF	2	789.9604	2.26	0.38
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]IYHSHIDAPK[272]	4	367.9593	2.114	0.061
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]RPYLK[272]	3	322.2109	1.754	0.042
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DTANLFPQTSLTLH	2	851.4501	4.259	0.297
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]YSTVDQVK[272]	2	614.3448	2.481	0.288
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]RDTANLFPQTSLTLH	3	620.0028	2.689	0.239
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TTIEKPVWLGFLGPIIK	3	779.4791	2.458	0.247
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TPGIWLLH	2	540.8198	2.729	0.186
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NM[147]ATRPYSIH	2	675.3413	2.465	0.308
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	IGGSYK	2	452.7662	1.373	0.093
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VFNPR	2	388.7304	1.311	0.014

18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	KLVYR	2	479.8135	1.207	0.093
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NM[147]ATRPYSIHAH	4	390.1983	1.665	0.134
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]DSLDK[272]EK[272]	3	422.9137	1.715	0.092
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TYCSEPEK[272]	2	645.8011	1.292	0.035
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	LTTDHYTGGM[147]K	3	507.2573	2.462	0.131
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	IGGSYKK	3	391.5765	1.815	0.013
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]TTIEKPVWLGFLGPIIK	4	584.8612	1.848	0.262
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NLASRPY	2	482.7703	2.083	0.1
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]VYVHLK[272]	3	349.5582	1.444	0.039
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]SIEPIGVR	2	507.8069	2.16	0.063
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NLASRPYTF	2	606.8283	2.341	0.088
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	TYCSEPEKVDK	3	588.9602	1.941	0.014
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	HYYIGIIETTWDYASDHGEK[272]	3	894.7701	4.253	0.364
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]NM[147]ATRPYSIHAH	2	779.3893	2.038	0.101
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]LSIEPIGVR	2	564.3489	1.999	0.092
18	IPI00017601	1	48.8	0.73	0.08	265	##	CP Ceruloplasmin precursor	n[145]IYHSHID	2	514.7677	1.811	0.06
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	CGHGFPGPLNIETK	2	898.4528	4.081	0.484
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	GDSGGAFAVQDPNDK	2	879.4263	4.674	0.462
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]CGHGFPGPLNIETK[272]	2	902.4599	4.602	0.495
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]GDSGGAFAVQDPNDK[272]	2	883.4334	4.826	0.506
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	QFGPYCGHGFPGPLNIETK	3	796.7258	4.935	0.466
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	SSNNPHSPIVEEFQVPYNK	3	822.7519	4.668	0.437
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]SSNNPHSPIVEEFQVPYNK[272]	3	825.4233	4.99	0.578
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]EDTPNSVWEPAK[272]	2	830.9247	4.049	0.319
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]VEDPESTLFGSVIR	2	846.9499	3.114	0.385
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]CGHGFPGPLNIETK[272]	3	601.9757	4.218	0.335
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	EDTPNSVWEPAK	2	826.9176	3.557	0.189
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	YHGDPM[147]PCPK	2	743.8357	2.595	0.285
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	CGHGFPGPLNIETK	3	599.3043	3.804	0.323
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]TNFDNDIALVR	2	711.3789	3.148	0.303
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]CVPVCGVPR	2	583.2772	2.125	0.138
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]LLEVPEGR	2	528.8121	2.604	0.227
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]SSNNPHSPIVEEFQVPYNK[272]	4	619.3193	3.106	0.157
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]QFGPYCGHGFPGPLNIETK[272]	3	799.3972	2.473	0.203
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]CEYQIR	2	501.2369	2.091	0.139
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]SDFSNEER	2	564.2579	2.197	0.145
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	EPFEEK	2	529.7795	2.167	0.12
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	YHGDPM[147]PCPK	3	496.2262	2.244	0.127
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]YVFR	2	364.7142	1.594	0.143
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	M[147]ICAGGEK	2	575.7745	2.347	0.252
19	IPI00017696	1	24.2	0.7	0.07	29	23	C1S Complement C1s subcomponent precursor	n[145]VEK[272]PTADAEAYVFTPN	2	1020.5301	3.074	0.272
20	IPI00018219	1	6.6	1.09	0.06	3	3	TGFBI Transforming growth factor-beta-induced protein ig-h3 precur	n[145]DGTPPIDAHTR	3	441.8973	4.142	0.38
20	IPI00018219	1	6.6	1.09	0.06	3	3	TGFBI Transforming growth factor-beta-induced protein ig-h3 precur	DGTPPIDAHTR	3	440.5616	3.226	0.303
20	IPI00018219	1	6.6	1.09	0.06	3	3	TGFBI Transforming growth factor-beta-induced protein ig-h3 precur	VFVYR	2	412.2449	1.039	0.052
21	IPI00018305	1	17.9	0.7	0.06	6	5	IGFBP3 Insulin-like growth factor-binding protein 3 precursor	SAGSVESPSVSSTHR	3	543.2727	4.484	0.388
21	IPI00018305	1	17.9	0.7	0.06	6	5	IGFBP3 Insulin-like growth factor-binding protein 3 precursor	n[145]SAGSVESPSVSSTHR	3	544.6084	3.287	0.387
21	IPI00018305	1	17.9	0.7	0.06	6	5	IGFBP3 Insulin-like growth factor-binding protein 3 precursor	n[145]ALAQCAPPPAVCAELVR	2	972.986	3.825	0.279
21	IPI00018305	1	17.9	0.7	0.06	6	5	IGFBP3 Insulin-like growth factor-binding protein 3 precursor	n[145]FLNVLSPR	2	545.3305	2.891	0.078

21	IPI00018305	1	17.9	0.7	0.06	6	5	IGFBP3 Insulin-like growth factor-binding protein 3 precursor	n[145]GK[272]EDVHCYSM[147]QS	2	867.3804	2.714	0.372
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]AYWDIM[147]ISNHQNSNR	3	670.3187	3.965	0.382
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]EALQGVGDM[147]GR	2	646.8229	2.928	0.349
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	EALQGVGDM[147]GR	2	644.8194	2.735	0.234
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]EALQGVGDMGR	2	638.8255	2.476	0.257
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	FRPDGLPK	2	605.3588	2.124	0.109
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]FRPDGLPK[272]	2	609.3659	2.303	0.06
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	AYWDIM[147]ISNHQNSNR	3	668.983	3.097	0.105
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]YIYAR	2	415.2381	1.89	0
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	n[145]FRPDGLPK[272]	3	406.5797	1.691	0.021
22	IPI00019399	1	33.1	0.85	0.08	18	10	SAA4 Serum amyloid A-4 protein precursor	GPGGVWAAK[272]	2	563.8281	1.915	0.021
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ETAASLLQAGYK	2	766.4276	5.437	0.444
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KPVAFSDYIHPVCLPDR	2	1142.0925	5.416	0.544
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KSPQELLCGASLISDR	2	1022.0399	5.176	0.545
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	TFGSGEADCGLRPLFEK	2	1077.0296	5.728	0.43
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ETAASLLQAGYK[272]	2	770.4347	4.973	0.489
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GQPSVLQVVNLPIVER	2	946.5579	4.493	0.489
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]NPDSSTTGPWCYTTDPTVR	2	1144.5076	4.599	0.488
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SEGSSVNLSPPLEQCVPDR	2	1102.5258	4.629	0.347
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]TATSEYQTFFNPR	2	853.4188	3.755	0.481
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]TFGSGEADCGLRPLFEK[272]	2	1081.0367	5.865	0.635
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]VAFSDYIHPVCLPDR	2	961.4747	5.5	0.479
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]YGFYTHVFR	2	667.3441	2.488	0.388
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	GQPSVLQVVNLPIVERPVCK	3	834.4747	4.247	0.503
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KPVAFSDYIHPVCLPDR	3	761.7308	4.547	0.48
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KSPQELLCGASLISDR	3	681.6957	3.268	0.395
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	LKKPVAFSDYIHPVCLPDR	3	888.8221	3.644	0.319
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GQPSVLQVVNLPIVERPVCK[272]	3	837.1461	4.214	0.461
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]RQECSIPVCGQDQVTVAM[147]TPR	3	857.3973	5.584	0.447
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]TFGSGEADCGLRPLFEK[272]	3	721.0269	4.301	0.393
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]VAFSDYIHPVCLPDR	3	641.3189	4.5	0.419
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]YGFYTHVFR	3	445.2318	3.486	0.286
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ITDNM[147]FCAGYKPDEGKR	4	607.5459	3.279	0.548
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]LK[272]K[272]PVAFSDYIHPVCLPDR	4	669.8737	5.2	0.466
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	LKKPVAFSDYIHPVCLPDR	5	533.6962	2.474	0.098
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]HQDFNSAVQLVENFCR	3	699.6636	4.038	0.315
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]LK[272]K[272]PVAFSDYIHPVCLPDR	3	892.8292	3.245	0.385
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ITDNM[147]FCAGYK[272]PDEGK[272]R	4	610.5513	2.385	0.313
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]LK[272]K[272]PVAFSDYIHPVCLPDR	5	536.1004	2.871	0.217
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GDACEGDSGGPFVMK[272]	2	902.4014	2.22	0.34
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ENLDRDIALMK[272]	3	535.9673	3.338	0.298
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ETWTANVGK	2	643.3486	3.311	0.229
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GDACEGDSGGPFVM[147]K[272]	2	910.3988	3.965	0.368
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ETWTANVGK[272]	2	647.3557	3.278	0.262
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]YTACETAR	2	552.7504	2.34	0.314
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ITDNM[147]FCAGYKPDEGKR	3	809.7255	2.9	0.229
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ELLESYIDGR	2	669.8547	1.631	0.042
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	YTACETAR	2	550.7468	2.338	0.288

23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SPQELLCGASLISDR	2	889.9485	3.12	0.325
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	LKKPVAFSDYIHPVCLPDR	4	666.8684	3.335	0.296
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	DIALM[147]K	2	493.7888	2.4	0.145
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]DIALMK[272]	2	489.7985	2.418	0.149
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	DIALMK	2	485.7914	2.365	0.12
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]DIALM[147]K[272]	2	497.7959	2.562	0.149
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SGIECQLWR	2	641.3137	2.873	0.164
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SLEDK[272]TER	3	422.5695	2.977	0.189
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	GDACEGDSGGPFVM[147]K	2	906.3917	4.275	0.229
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ISMLEK[272]	2	504.8038	2.293	0.129
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]IYIHPR	2	471.7857	2.014	0.13
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KPVAFSDYIHPVCL	3	639.0038	3.006	0.41
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	SGIECQLWR	2	639.3101	2.537	0.125
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ISM[147]LEK[272]	2	512.8012	1.741	0.105
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]PVAFSDYIHPVCLPDR	3	673.6698	2.689	0.304
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]FSDYIHPVCLPDR	3	584.617	2.792	0.263
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ETAASLLQAGYK[272]	3	513.9589	2.684	0.048
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ISMLEK	2	500.7967	1.889	0.011
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	KPVAFSDYIHPVCLPD	3	709.697	2.751	0.329
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SLEDK[272]TER	2	633.3506	2.366	0.154
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]YGFYTH	2	466.2252	2.103	0.443
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SDYIHPVCLPDR	2	802.8877	3.275	0.389
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ENLDRDIALM[147]K[272]	2	811.4448	2.771	0.198
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]SAVQLVENFCR	2	728.3639	3.065	0.321
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GFYTHVFR	2	585.8125	3.115	0.251
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GFYTHVFR	3	390.8774	2.382	0.227
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ISM[147]LEK	2	508.7941	1.548	0.084
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	SLEDKTER	3	419.8981	2.528	0.082
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]NIEK[272]ISM[147]LEK[272]	3	551.6588	1.922	0.153
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	SLEDKTER	2	629.3435	2.415	0.136
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	TFGSGEADCGLRPLFEK	3	718.3555	1.898	0.096
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	GDACEGDSGGPFVMK	2	898.3943	1.447	0.058
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]GQPSVLQVVNLPIVER	3	631.3744	3.447	0.116
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]TFGSGEADCGLR	2	701.8142	2.324	0.278
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	ANTFLEEVRK	3	496.2841	2.015	0.079
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]LAVTTHGLPC	2	601.3131	1.628	0.173
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	n[145]ENLDRDIALM[147]K[272]	3	541.2989	1.531	0.058
23	IPI00019568	1	48.9	0.74	0.07	147	72	F2 Prothrombin precursor (Fragment)	LAVTTH	2	391.232	1.78	0.087
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	n[145]QEDACQGDSGGPHVTR	3	616.2691	3.737	0.47
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	n[145]ETYDFDIAVLR	2	743.3889	3.564	0.428
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	n[145]YGIYTK[272]	2	516.8021	2.207	0.24
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	GKYGIYTK	3	450.6029	2.629	0.155
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	n[145]TGIVSGFGR	2	519.2967	1.973	0.138
24	IPI00019576	1	9.6	0.73	0.04	4	5	F10 Coagulation factor X precursor	EM[147]K	1	703.3808	1.121	0.072
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	EPLDDYVNTQGASLFSVTK	2	1182.6078	6.482	0.615
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	QLGAGSIEECAAK	2	801.9026	4.271	0.439
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]DGDVGGPWCYTTNPR	2	914.3992	3.426	0.536
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EPLDDYVNTQGASLFSVTK[272]	2	1186.6149	6.028	0.549

25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]FSPATHPSEGLEENYCR	2	1063.9732	4.26	0.536
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]GNVAVTVSGH	2	542.797	3.286	0.454
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]QLGAGSIEECAAK[272]	2	805.9097	4.574	0.517
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VIPACLPSPNYVVADR	2	952.4982	3.933	0.42
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]CTTPPPSSGPTYQCLK[272]	3	687.3236	4.117	0.366
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]FSPATHPSEGLEENYCR	3	709.6512	5.495	0.55
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]LYDYCDVPQCAAPSFDCGK[272]PQVEPK[272]	3	1115.1715	4.72	0.407
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]NPDNDPQGPWCYTTDPEK[272]R	3	856.3893	3.356	0.328
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	KLYDYCDVPQCAAPSFDCGKPQVEPK	4	900.6726	3.8	0.48
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TM[147]SGLECQAWDSQSPHAHGYIPSK	3	992.7902	4.314	0.39
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VCNRYEFLNGR	3	520.9226	3.463	0.338
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TM[147]SGLECQAWDSQSPHAHGYIPSK[272]	4	745.8462	3.953	0.354
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	KLYDYCDVPQCAAPSFDCGKPQVEPK	5	720.7396	3.542	0.196
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	LSSPAVITDK	2	655.8876	2.727	0.349
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EQQCVIM[147]AENR	2	763.8477	3.89	0.399
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]HSIFTPETNPR	2	721.8791	1.922	0.095
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]LSSPAVITDK[272]	2	659.8947	2.546	0.064
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TPENYPNAGLTM[147]NYCR	3	683.9698	3.026	0.26
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VVGGCVAHPH	3	389.1957	3.281	0.42
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EQQCVIMAENR	2	755.8503	3.631	0.334
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TPENFPCK[272]	2	635.3142	3.104	0.313
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	TPENFPCK	2	631.3071	3.134	0.267
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TPENYPNAGLTM[147]NYCR	2	1025.4511	3.847	0.342
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]CEEDEEFTCR	2	748.7678	2.978	0.419
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]YEFLNGR	2	521.7756	1.828	0.305
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	TKNGITCQK	3	486.9355	2.644	0.289
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]NLDENYCR	2	608.764	2.509	0.352
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]NPDGDVGGPWCYTTNPR	2	1019.947	3.831	0.457
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]WEYCNLK[272]	2	645.3167	2.16	0.322
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VYLSECK[272]	2	588.3058	2.679	0.158
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	AFQYHSK	2	580.8142	2.769	0.162
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	WEYCNLK	2	641.3096	1.897	0.105
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TK[272]NGITCQK[272]	3	490.9426	3.224	0.172
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VILGAHQEVNLEPHVQEIEVSR	3	880.8124	3.66	0.412
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]WELCDIPR	2	611.2975	2.998	0.275
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	VYLSECK	2	584.2987	2.322	0.182
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]LFLEPTR	2	510.304	2.079	0.067
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]GTSSTTTTGK[272]K[272]	3	500.958	2.09	0.267
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	KVYLSECK	3	479.2649	2.571	0.105
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EAQLPVIENK[272]	2	714.9187	3.311	0.217
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]AHGYIPSK[272]FPNK[272]	3	597.6808	3.515	0.32
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VQSTELCAGHLAGGTDSCQGDSGGPLVCFEK[2	4	873.8873	2.89	0.065
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]WEYCNLK	2	643.3131	2.184	0.276
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]DIAILK[272]	2	480.8203	1.803	1
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	EAQLPVIENK	2	710.9116	2.212	0.091
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]SPRPSSYK[272]	2	605.3452	2.191	0.157
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EPLDDYVNTQGASLFSVTK[272]	3	791.4123	3.265	0.261
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VIPACLPSPNYVVADR	3	635.3345	2.716	0.074

25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	KDIALLK	3	407.6078	2.624	0.098
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]SPRPSSYK[272]	3	403.8992	2.045	0.098
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	DVVLFEK	2	565.3345	1.682	0.107
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	NPDADKGPWCFTTDPSVR	3	778.0332	2.372	0.17
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	PNKPGVYVR	2	655.3906	2.58	0.368
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	LFLEPTR	2	508.3004	1.75	0.089
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]PNK[272]PGVYVR	2	659.3977	2.737	0.306
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	PNKPGVYVR	3	437.2628	1.999	0.27
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	CTTPPPSSGPTYQCLK	3	684.6522	2.593	0.12
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]ELRPWCFTTDPNK[272]	3	647.6582	2.55	0.04
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	VIPACLPSPNYVVADR	3	633.9988	2.497	0.068
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VVGGCVAHPH	2	583.29	2.157	0.41
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VQSTELCAGH	2	617.7875	2.583	0.236
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	AHGYIPSKFPNK	3	593.6737	2.909	0.218
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]GNVAVTVSGHTCQH	3	533.9211	2.745	0.145
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]AFQYH	2	405.2068	1.626	0.251
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]EQQCVIM[147]AENRK[272]	3	600.2999	2.204	0.027
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VIPACLPSPN	2	600.8155	2.216	0.223
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	GNVAVTVSGH	1	1080.5797	2.09	0.475
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]GNVAVTVSGH	1	1084.5868	2.507	0.419
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]ENFPCK[272]	2	536.2639	2.022	0.113
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]ELRPWCF	2	570.7762	1.497	0.296
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]DSQSPHAHGYIPSK[272]	3	604.6475	2.687	0.122
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	AFQYH	2	403.2032	1.51	0.184
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	ENFPCK	2	532.2568	2.001	0.074
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]YVVADR	2	433.7463	1.935	0.05
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TPENYPNAGLTM[147]N	2	791.3704	2.224	0.147
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]TPENYPNAGLTMN	2	783.373	2.21	0.081
25	IPI00019580	1	57.2	0.77	0.08	131	78	PLG Plasminogen precursor	n[145]VQSTELCAGHLAG	2	738.3588	2.052	0.074
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	LCHCPVGYTGPFCDVDTK	3	791.6724	5.393	0.442
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]LCHCPVGYTGPFCDVDTK[272]	3	794.3438	5.863	0.526
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]CLEVEGHR	3	378.1835	3.062	0.221
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]VVGGLVALR	2	514.3409	3.009	0.245
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]CFEPQLLR	2	598.3078	1.955	0.034
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]CLEVEGHR	2	566.7716	2.351	0.251
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	CFEPQLLR	2	596.3043	1.864	0.179
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]EQPPSLTR	2	536.2994	2.009	0.015
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	EQPPSLTR	2	534.2958	2.073	0.014
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	LCHCPVGY	2	562.2314	1.622	0.248
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	n[145]VK[272]DHCSK[272]	3	432.2373	2.096	0.005
26	IPI00019581	1	10.9	0.81	0.05	17	11	F12 Coagulation factor XII precursor	NWGLGGHAF	2	549.7776	1.757	0.064
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	ALFVSEEEKK	2	800.4589	4.551	0.357
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	EAGIPEFYDYDVALIK	2	1062.0566	4.294	0.49
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KDNEQHVFK	2	782.9334	3.645	0.464
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	PQGSCSLEGVEIK	2	836.9235	3.875	0.471
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VKDISEVVTPR	2	761.9513	3.852	0.447
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VSEADSSNADWVTK	2	894.9418	5.915	0.558
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	YGLVTYATYPK	2	778.4296	4.491	0.497

27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]FLCTGGVSPYADPNTCR	2	1018.9446	5.63	0.507
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]GDSGGPLIVHK[272]	2	684.3979	4.088	0.421
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]LEDSVTYHCSR	2	750.3406	3.611	0.489
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]LLQEGQALEYVCPSGFYPYPVQTR	2	1474.7258	5.354	0.5
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VSEADSSNADWVTK[272]	2	898.9489	4.828	0.438
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGLVTYATYPK[272]	2	782.4367	4.317	0.525
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]ALFVSEEEK[272]K[272]	3	537.9821	3.504	0.438
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]LEDSVTYHCSR	3	500.5628	2.567	0.217
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]QLNEINYEDHK[272]	3	564.2928	5.06	0.251
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VASYGVK[272]PR	3	422.2587	2.307	0.258
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	ALFVSEEEKK	3	533.975	3.362	0.326
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	GDSGGPLIVHK	3	453.9296	3.176	0.227
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KDNEQHVFK	3	522.2914	3.057	0.33
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KVGSQYR	3	373.2208	2.915	0.311
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VASYGVKPR	3	419.5873	1.888	0.174
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VKDISEVVTPR	3	508.3033	3.841	0.336
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DLEIEVVLFHPNYNINGK[272]	3	801.4368	3.363	0.449
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]GDSGGPLIVHK[272]	3	456.601	3.455	0.199
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VK[272]DISEVVTPR	3	510.9747	3.609	0.394
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGQTIRPICLPCTEGTTR	3	749.0288	3.399	0.381
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KDNEQHVFK	4	391.9704	2.48	0.085
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	GDSGGPLIVHK	2	680.3908	3.954	0.395
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VASYGVKPR	2	628.8774	1.818	0.172
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]LPPTTTCQQQK[272]	2	789.9148	2.879	0.403
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VASYGVK[272]PR	2	632.8845	1.907	0.083
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	CLVNLIEK	2	629.3565	3.292	0.285
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]CLVNLIEK[272]	2	633.3636	3.337	0.344
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DAQYAPGYDK[272]	2	708.3559	3.583	0.376
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DNEQHVFK[272]	2	652.8455	3.485	0.224
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]QLNEINYEDHK[272]	2	845.9356	3.583	0.301
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]FLCTGGVSPYADPNTCR	3	679.6321	2.432	0.166
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	QLNEINYEDHK	2	841.9285	2.822	0.382
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	DAQYAPGYDK	2	704.3488	3.168	0.36
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]ALFVSEEEK[272]K[272]	2	806.4696	3.206	0.232
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	DLLYIGK	2	551.337	1.512	0.033
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	DNEQHVFK	2	648.8384	2.896	0.199
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DLLYIGK[272]	2	555.3441	2.465	0.035
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]HVIILM[147]TDGLHNM[147]GGD	3	633.6455	3.274	0.373
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]ALFVSEEEK[272]	2	670.371	2.849	0.187
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DISEVVTPR	2	580.3256	1.919	0.082
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	ALFVSEEEK	2	666.3639	2.749	0.194
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VK[272]DISEVVTPR	2	765.9584	3.288	0.38
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]EELLPAQDIK[272]	2	722.4185	2.755	0.175
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]EAGIPEFYDYDVALIK[272]	2	1066.0637	2.8	0.17
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	DNEQHVFK	3	432.8947	2.194	0.159
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KEVYIK	2	600.3792	2.533	0.052
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	EELLPAQDIK	2	718.4114	2.507	0.098
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]TVDDK[272]EHSIK[272]	4	401.7309	2.847	0.335

27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]EVYIK[272]	2	470.2913	2.249	0.168
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]PQGSCSLEGVEIK[272]	2	840.9306	4.34	0.247
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]HVIILMTDGLHNM[147]GGDPITVIDEIR	3	973.8445	2.187	0.251
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DNEQHVFK[272]	3	435.5661	2.423	0.051
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VSEADSSNADWVTK[272]	2	896.9454	4.116	0.383
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	EVYIK	2	466.2842	1.949	0.19
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VSVGGEK[272]	2	482.2893	2.001	0.021
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	EELLPAQDIK	3	479.2767	2.322	0.114
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KGTDYHK	2	634.8592	1.947	0.068
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KGTDYHK	4	317.9332	2.048	0.121
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]PICLPCTEGTTR	2	763.8514	2.583	0.253
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	VASYGVK	2	502.3004	2.579	0.188
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VCPSGFYPYPVQTR	2	902.4376	3.349	0.354
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGQTIRPICLPCTEG	2	943.9413	2.364	0.345
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]CYDGYTLR	2	590.766	2.347	0.127
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]TAAHCFTVDDK[272]EHSIK[272]	4	570.7924	2.592	0.188
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	QLNEINYEDHK	3	561.6214	1.824	0.064
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VASYGVK[272]	2	506.3075	2.249	0.105
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]FVSEEEK[272]K[272]	3	476.6084	3.176	0.034
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	TVDDKEHSIK	3	531.2983	2.179	0.218
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	KALQAVYSM[147]	2	653.8631	2.086	0.089
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]VCPSGFYPYPVQTR	3	601.9608	2.589	0.099
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGLVTY	1	859.4682	1.718	0.25
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGQTIRPICLPC	2	800.3854	1.681	0.244
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]DFHINLF	2	525.2805	1.727	0.093
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	DVCK	2	395.7012	1.198	0.182
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	SYGVKPR	2	543.8246	1.635	0.055
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	YGLVTY	1	855.4611	1.367	0.328
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]YGQTIR	2	441.2517	1.544	0.033
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]LEDSVTYHCS	2	672.29	1.46	0.112
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	n[145]TVDDK[272]EHSIK[272]	3	535.3054	2.085	0.109
27	IPI00019591	1	45.4	0.68	0.07	154	77	CFB Isoform 1 of Complement factor B precursor (Fragment)	SK[272]K[272]	2	395.7731	0.913	0.136
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	AESPEVCFNEESPK	2	946.4319	5.385	0.584
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	ESLLNHFLYEVAR	2	865.9649	4.374	0.462
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	IAPQLSTEELVSLGEK	2	997.5621	4.736	0.526
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	TINPAVDHCCK	2	786.8613	3.975	0.529
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]AESPEVCFNEESPK[272]	2	950.439	4.709	0.429
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]ESLLNHFLYEVAR	2	867.9684	4.372	0.403
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]TINPAVDHCCK[272]	2	790.8684	4.275	0.453
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	TINPAVDHCCK	3	524.91	3.799	0.483
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]LK[272]HELTDEELQSLFTNFANVVDK[272]	3	1041.5627	6.172	0.567
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]TINPAVDHCCK[272]	3	527.5814	4.51	0.527
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	GQCIINSNKDDRPK	3	685.3593	4.607	0.273
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	KSDVGFLPPFPTLDPEEK	3	812.7773	3.693	0.346
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]GQCIINSNK[272]DDRPK[272]	3	689.3664	4.831	0.312
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]RHPDLSIPELLR	3	530.6463	2.493	0.158
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]SCCEEQNK[272]	2	660.7684	2.59	0.45
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]CQAYESNR	2	580.7509	2.994	0.381

28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	AIPVTQYLK	2	656.903	2.924	0.351
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	MVQQECK	2	596.2878	2.607	0.322
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]LCFFYNK[272]	2	634.8242	2.852	0.32
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	HELTDEELQSLFTNFANVVDK	3	910.4642	2.473	0.293
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	LPNNVLQEK	2	667.8932	2.769	0.217
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	SCCEEQNK	2	656.7613	2.295	0.358
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	LCFFYNK	2	630.8171	2.676	0.249
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]LPNNVLQEK[272]	2	671.9003	2.113	0.109
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]FTFEYSR	2	547.2754	1.853	0.145
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]HELTDEELQSLFTNFANVVDK[272]	4	685.1036	3.208	0.167
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]YHYLIR	2	504.791	2.465	0.177
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	FLVNLVK	2	556.8632	2.763	0.164
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]FTDSENVCQER	2	759.3277	2.719	0.272
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	FTFEYSR	2	545.2718	2.135	0.059
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]RHPDLSIPELLR	2	795.4658	2.973	0.334
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]FLVNLVK[272]	2	560.8703	2.173	0.256
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	YHYLIR	2	502.7875	2.45	0.142
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]TLPECSK[272]	2	556.2901	2.36	0.164
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]AIPVTQYLK[272]	2	660.9101	2.464	0.162
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	HFQNLGK	2	562.3222	2.348	0.164
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]TVAVHFEEVAK[272]	3	506.6237	3.45	0.283
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	ICAM[147]EGLPQK	2	716.3615	2.145	0.231
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	TVAVHFEEVAK	3	503.9523	3.188	0.294
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]HFQNLGK[272]	2	566.3293	2.255	0.117
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]DMVEYK[272]	2	536.783	1.838	0.165
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	IVQIYK	2	522.3343	2.332	0.158
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]ECCEK[272]	2	496.2016	1.788	0.169
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	TLPECSK	2	552.283	1.76	0.03
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]MVQQECK[272]	2	600.2949	1.912	0.122
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	ECCEK	2	492.1945	1.776	0.147
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	DADPDTFFAK	2	703.8512	2.196	0.127
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]IVQIYK[272]	2	526.3414	2.207	0.169
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	RHPDLSIPELLR	3	529.3106	1.921	0.177
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	LCFFYNKK	3	510.2771	2.443	0.026
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]VNCLQTR	2	512.2634	1.626	0.027
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	NNVLQEK	2	562.8248	2.925	0.184
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]IK[272]ECCEK[272]	3	459.5639	2.803	0.162
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]NNVLQEK[272]	2	566.8319	2.342	0.068
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	IKECCEK	2	682.8315	2.35	0.226
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]YHYLIR	3	336.8631	1.699	0.004
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	LKHELTDEELQSLFTNFANVVDK	4	778.4185	2.494	0.064
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]ESLLNHFLYEVAR	3	578.9814	2.162	0.007
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	IKECCEK	3	455.5568	2.094	0.106
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]CFESLK[272]	2	530.7741	2.122	0.15
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	CFESLK	2	526.767	1.8	0.141
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]ICAM[147]EGLPQK[272]	2	720.3686	2.41	0.03
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	n[145]RNPFVFAPTLL	2	709.9175	2.155	0.131
28	IPI00019943	1	48.1	0.81	0.1	86	59	AFM Afamin precursor	RNPFVFAPTLL	2	707.9139	2.065	0.187

29	IPI00020019	1	12.3	1	0.01	5	4	ADIPOQ Adiponectin precursor	IFYNQQNHYDGSTGK	3	684.6693	3.839	0.393
29	IPI00020019	1	12.3	1	0.01	5	4	ADIPOQ Adiponectin precursor	n[145]IFYNQQNHYDGSTGK[272]	3	687.3407	4.434	0.402
29	IPI00020019	1	12.3	1	0.01	5	4	ADIPOQ Adiponectin precursor	GDIGETGVPGAEGPR	2	776.3917	3.632	0.442
29	IPI00020019	1	12.3	1	0.01	5	4	ADIPOQ Adiponectin precursor	n[145]GDIGETGVPGAEGPR	2	778.3953	2.447	0.273
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]EHVAHLLFLR	2	689.9074	3.635	0.38
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]TLMFGSYLDDEK[272]	2	853.9311	4.492	0.516
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]EHVAHLLFLR	3	460.274	4.407	0.351
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]YEGGREHVAHLLFLR	3	647.6907	3.933	0.423
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	TLM[147]FGSYLDDEKNWGLSFYADKPETTK	4	933.971	4.955	0.514
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	TLMFGSYLDDEK	2	849.924	4.243	0.385
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]EHVAHLLFLR	4	345.4574	1.76	0.054
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]YEGGREHVAHLLFLR	4	486.0199	3.803	0.29
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]TLM[147]FGSYLDDEK[272]	2	861.9286	3.89	0.352
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]DK[272]CEPLEK[272]	3	480.5915	3.228	0.147
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	SDVMYTDWK	2	712.8476	3.179	0.348
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	DKCEPLEK	3	476.5844	2.348	0.263
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]YEGGREHVAHLLFLR	5	389.0173	2.84	0.137
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]TEDTIFLR	1	1138.6225	2.6	0.235
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]TEDTIFLR	2	569.8149	2.252	0.02
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]EQLGEFYEALDCLCIPR	3	745.674	2.237	0.292
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]NWGLSFYADKPETTK	3	727.7202	3.22	0.281
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]SDVMYTDWK[272]	2	716.8547	2.857	0.213
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	SDVM[147]YTDWK	2	720.8451	2.809	0.209
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]CEPLEK[272]	2	526.7716	1.507	0.229
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]CEPLEK[272]	1	1052.5359	1.61	0.168
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	DKCEPLEK	2	714.3729	2.542	0.263
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]DK[272]CEPLEK[272]	2	720.3836	2.88	0.121
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]FYIASAFR	2	559.8094	2.117	0.267
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	CEPLEK	2	522.7645	1.847	0.016
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]SSYLNVQR	2	555.8048	2.701	0.185
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]YIASAFR	2	486.2752	1.866	0.286
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]IASAFR	2	404.7435	1.822	0.055
30	IPI00020091	1	49.8	0.98	0.29	27	24	ORM2 Alpha-1-acid glycoprotein 2 precursor	n[145]YLNVQR	2	468.7728	1.886	0.018
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	LKEDAVSAAFK	2	799.9669	4.588	0.382
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]LK[272]EDAVSAAFK[272]	2	805.9776	4.661	0.497
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]SLEDLQLTHNK[272]	2	793.4431	4.842	0.399
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	SLEDLQLTHNK	3	526.6264	3.645	0.42
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]NNQIDHIDEK[272]	3	505.2665	3.852	0.373
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]SLEDLQLTHNK[272]	3	529.2978	4.121	0.458
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	LKEDAVSAAFK	3	533.647	3.077	0.295
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	SLEDLQLTHNK	2	789.436	4.199	0.362
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	NNQIDHIDEK	2	753.389	3.586	0.323
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]FNALQYLR	2	584.8334	3.234	0.325
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	NNQIDHIDEK	3	502.5951	2.953	0.272
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]SLEYLDLSFNQIAR	2	906.9843	3.303	0.286
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	FNALQYLR	2	582.8299	2.62	0.071
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	n[145]ILGPLSYSK[272]	2	633.389	2.668	0.209
31	IPI00020986	1	18.6	0.93	0.05	18	15	LUM Lumican precursor	ILGPLSYSK	2	629.3819	1.91	0.078

32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]DLHFLEELQLGHNR	2	932.993	4.709	0.445
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]DLHFLEELQLGHNR	3	622.3311	6.299	0.316
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LHSLHLEGSCLGR	3	537.9454	3.868	0.357
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]SFEGLGQLEVLTLDHNQLQEVK[272]	3	929.1707	4.858	0.414
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]DLHFLEELQLGHNR	4	467.0001	4.527	0.354
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	NLIAAVAPGAFLGLK	3	579.0258	2.922	0.312
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]AFWLDVSHNR	3	463.5779	3.819	0.273
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]DFALQNPSAVPR	2	729.8947	4.408	0.361
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LAYLQPALFSGLAELR	2	953.5496	3.345	0.447
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LEALPNSLLAPLGR	2	804.4837	3.712	0.285
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]WLDLSHNR	3	395.5479	3.225	0.193
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]ANVFVQLPR	2	594.3545	3.363	0.248
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LEYLLLSR	2	575.8513	3.028	0.275
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LAELPADALGPLQR	2	804.4655	2.728	0.308
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LHSLHLEGSCLGR	2	806.4144	3.409	0.269
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]NLIAAVAPGAFLGLK[272]	2	872.0422	2.878	0.351
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]LHSLHLEGSCLGR	4	403.7109	2.24	0.147
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]TFTPQPPGLER	2	693.8785	2.682	0.233
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]NLPEQVFR	2	573.823	2.481	0.182
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	NLIAAVAPGAFLGLK	2	868.0351	2.572	0.334
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]WLDLSHNR	2	592.8183	2.479	0.129
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]YLSLR	2	398.2459	1.721	0.003
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]SLALGTFAH	2	530.799	2.34	0.326
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	SFEGLGQLEVLTLDHNQLQEVK	3	926.4993	2.735	0.094
32	IPI00020996	1	33.4	0.76	0.07	38	25	IGFALS Insulin-like growth factor-binding protein complex acid labile	n[145]GLLEDTFPGLLGLR	2	822.9757	2.346	0.313
33	IPI00021364	1	8.7	0.67	0.03	4	4	CFP Properdin precursor	n[145]TCNHPVPQHGGPFCAGDATR	4	576.003	4.95	0.553
33	IPI00021364	1	8.7	0.67	0.03	4	4	CFP Properdin precursor	LCTPLLPK	2	605.8562	2.091	0.172
33	IPI00021364	1	8.7	0.67	0.03	4	4	CFP Properdin precursor	n[145]CSAPEPSQK[272]PPGK[272]	3	601.9868	3.122	0.286
33	IPI00021364	1	8.7	0.67	0.03	4	4	CFP Properdin precursor	n[145]LCTPLLPK[272]	2	609.8633	1.873	0.162
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	EEIIYECDK	2	734.3466	3.528	0.413
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	GSSVIHCDADSK	2	772.8635	4.45	0.38
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	TPSCGDICNFPPK	2	875.8928	3.534	0.503
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GSSVIHCDADSK[272]	2	776.8706	4.139	0.345
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GVGWSHPLPQCEIVK[272]	2	992.523	4.248	0.438
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GYILVGQAK[272]	2	618.8814	4.147	0.441
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]SCSEGFFLIGSTTSR	2	891.4196	5.423	0.549
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	CHPGYKPTTDEPTTVICQK	3	877.4248	4.368	0.434
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	DVSHGEMVSGFGPIYNYK	3	760.7088	5.668	0.58
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	DVSHGEM[147]VSGFGPIYNYK	3	766.0404	5.439	0.44
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	GSSVIHCDADSK	3	515.5781	3.891	0.489
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEMVSGFGPIYNYK	3	882.4564	6.119	0.564
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEM[147]VSGFGPIYNYK	3	887.788	5.936	0.604
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]CHPGYK[272]PTTDEPTTVICQK[272]	3	881.4319	4.489	0.448
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]DVSHGEM[147]VSGFGPIYNYK[272]	3	768.7118	6.325	0.505
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GSSVIHCDADSK[272]	3	518.2495	2.712	0.434
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GVGWSHPLPQCEIVK[272]	3	662.0177	4.488	0.444
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEM[147]VSGFGPIYNYKDTIVFK	4	876.9642	3.731	0.449
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]CHPGYK[272]PTTDEPTTVICQK[272]	4	661.3258	4.007	0.417

34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	GVGWSHPLPQCEIVK	3	659.3463	3.691	0.318
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]FK[272]TGTTLK[272]	3	443.2819	2.704	0.39
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]LNNGEITQHR	3	442.5729	2.472	0.041
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	CHPGYKPTTDEPTTVICQK	4	658.3204	3.312	0.32
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEMVSGFGPIYNYK	4	662.0941	3.151	0.302
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEM[147]VSGFGPIYNYK	4	666.0928	3.009	0.138
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]FSAICQGDGTWSPR	2	857.8935	4.031	0.374
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]LNNGEITQHR	2	663.3558	3.975	0.341
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]LSLEIEQLELQR	2	807.9628	4.093	0.373
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]TPSCGDICNFPPK[272]	2	879.8999	3.093	0.439
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]YTCLPGYVR	2	631.3131	2.08	0.232
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPDVSHGEM[147]VSGFGPIYNYK	5	533.0757	2.532	0.223
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	GYILVGQAK	2	614.8743	2.969	0.267
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]EDVYVVGTVLR	2	697.394	3.25	0.32
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPELVNGR	3	398.2436	2.364	0.217
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]EEIIYECDK[272]	2	738.3537	3.194	0.272
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]DTIVFK[272]	2	505.8099	2.417	0.175
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]DVSHGEMVSGFGPIYNYK[272]	3	763.3802	4.533	0.293
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]VSHGEM[147]VSGFGPIYNYK[272]	3	730.3695	5.011	0.367
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	DTIVFK	2	501.8028	2.182	0.177
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	KPELVNGR	2	596.8617	2.151	0.176
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	IAHGHYK	2	553.3169	2.015	0.165
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	MALEVYK	2	567.323	2.388	0.041
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GVGWSHPLPQCEIVK	2	990.5194	4.213	0.362
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	NGQVEIK	2	534.3141	2.045	0.085
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]GVGWSHPLPQCEIVK	3	660.682	2.732	0.193
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]NGQVEIK[272]	2	538.3212	2.27	0.031
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]SCGDICNFPPK[272]	2	780.8497	2.781	0.303
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	LNNGEITQHR	3	441.2372	2.546	0.005
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]CEVQDR	2	470.2109	1.73	0.037
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	FKTGTTLK	3	439.2748	1.673	0.15
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]WTPYQGCEALCCPEPK[272]	3	750.9857	2.182	0.007
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	CKPPPDIR	3	417.8949	1.608	0.012
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]TCLPGYVR	2	549.7815	2.102	0.175
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]TPSCGDICNFPPK[272]	3	586.9357	1.955	0.044
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	CCPEPK	2	524.726	1.717	0.184
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]PSPPTCEK[272]	2	596.8009	2.08	0.1
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]FSLLGH	2	409.2381	2.096	0.235
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]LSCSYSH	2	493.7132	1.811	0.056
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]NCGPPPTLSFAAPM[147]DITLTETR	3	846.7435	3.429	0.046
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	LSCSYSH	2	491.7097	1.663	0.063
34	IPI00021727	1	47.6	0.68	0.11	95	56	C4BPA C4b-binding protein alpha chain precursor	n[145]CCPEPK[272]	2	528.7331	1.576	0.064
35	IPI00021817	1	5.9	1.03	0.07	4	4	PROC Vitamin K-dependent protein C precursor	n[145]LGDDLLQCHPAVK[272]	3	581.6439	3.744	0.41
35	IPI00021817	1	5.9	1.03	0.07	4	4	PROC Vitamin K-dependent protein C precursor	LGDDLLQCHPAVK	2	867.9551	3.95	0.423
35	IPI00021817	1	5.9	1.03	0.07	4	4	PROC Vitamin K-dependent protein C precursor	n[145]TFVLNFIK[272]	2	635.3941	3.399	0.303
35	IPI00021817	1	5.9	1.03	0.07	4	4	PROC Vitamin K-dependent protein C precursor	TFVLNFIK	2	631.387	2.215	0.123
35	IPI00021817	1	5.9	1.03	0.07	4	4	PROC Vitamin K-dependent protein C precursor	SK[272]K[272]	2	395.7731	0.913	0.136
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLATVYVDVLK	1	1515.8782	1.274	0.431

36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDELR	1	1441.7435	2.987	0.377
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTK	1	1666.9051	3.441	0.516
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPYSDELR	1	1445.7506	3.308	0.337
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VQPYLDDFQK[272]	1	1540.8251	2.874	0.341
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSALEEYTK[272]	1	1674.9193	0.942	0.228
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ATEHLSTLSEK	2	748.4094	3.301	0.318
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLATVYVDVLK	2	758.4427	3.576	0.51
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DSGRDYVSQFEGSALGK	2	1048.524	5.284	0.412
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DYVSQFEGSALGK	2	840.9333	5.288	0.467
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEK	2	1107.0655	5.359	0.407
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEK[272]	2	1109.0691	4.67	0.428
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LLDNWDSVTSTFSK	2	946.9913	3.275	0.313
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QGLLPVLESFK	2	755.9533	2.621	0.377
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VKDLATVYVDVLK	2	942.0719	3.956	0.415
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VQPYLDDFQK	2	766.9091	1.92	0.127
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VQPYLDDFQKK	2	901.004	4.246	0.478
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTK	2	833.9562	3.643	0.412
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTKK	2	968.0512	5.008	0.392
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ATEHLSTLSEK[272]	2	752.4165	3.43	0.306
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLATVYVDVLK[272]	2	762.4498	4.392	0.474
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DSGRDYVSQFEGSALGK[272]	2	1052.5311	5.166	0.429
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DYVSQFEGSALGK[272]	2	844.9404	4.879	0.471
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]EQLGPVTQEFWDNLEK[272]	2	1111.0726	4.041	0.488
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LLDNWDSVTSTFSK[272]	2	950.9984	3.962	0.421
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VK[272]DLATVYVDVLK[272]	2	948.0826	4.264	0.41
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VQPYLDDFQK[272]	2	770.9162	2.954	0.163
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VQPYLDDFQK[272]K[272]	2	907.0147	4.123	0.432
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSALEEYTK[272]	2	837.9633	4.141	0.497
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSALEEYTK[272]K[272]	2	974.0618	5.4	0.437
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]YVSQFEGSALGK[272]	2	787.4269	4.445	0.469
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLATVYVDVLKDSGR	3	644.3581	4.205	0.418
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DSGRDYVSQFEGSALGK	3	699.3518	4.579	0.467
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEK	3	738.3794	5.921	0.543
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEKETEGLR	3	966.8259	5.659	0.603
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KWQEEMELYR	3	564.624	3.375	0.342
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LAEYHAKATEHLSTLSEK	3	816.7797	4.611	0.501
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LREQLGPVTQEFWDNLEK	3	828.1078	5.508	0.437
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VEPLRAELQEGAR	3	536.6336	3.635	0.533
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VKDLATVYVDVLK	3	628.3837	3.692	0.436
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VKDLATVYVDVLKDSGR	3	766.7775	4.964	0.611
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTKK	3	645.7032	3.854	0.345
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLATVYVDVLK[272]DSGR	3	647.0295	4.004	0.436
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DSGRDYVSQFEGSALGK[272]	3	702.0232	6.193	0.501
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]EQLGPVTQEFWDNLEK[272]	3	741.0508	5.75	0.5
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]EQLGPVTQEFWDNLEK[272]ETEGLR	3	969.4973	5.738	0.591
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LREQLGPVTQEFWDNLEK[272]	3	830.7792	5.473	0.427
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VK[272]DLATVYVDVLK[272]	3	632.3908	2.321	0.248
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VK[272]DLATVYVDVLK[272]DSGR	3	770.7846	5.374	0.51

36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VQPYLDDFQK[272]K[272]	3	605.0122	2.011	0.131
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSALEEYTK[272]K[272]	3	649.7103	5.497	0.498
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLATVYVDVLKDSGRDYVSQFEGSALGK	4	863.957	4.404	0.523
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LREQLGPVTQEFWDNLEKETEGLR	4	792.6676	5.487	0.606
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VKDLATVYVDVLKDSGR	4	575.335	3.217	0.321
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LREQLGPVTQEFWDNLEK[272]ETEGLR	4	794.6711	5.988	0.534
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AHVDALR	3	307.847	2.743	0.277
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KWQEEM[147]ELYR	3	569.9556	2.683	0.335
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LAEYHAK	3	371.2135	1.727	0.142
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDELR	3	481.2527	3.152	0.3
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDELRQR	3	575.9726	3.04	0.353
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VQPYLDDFQKK	3	601.0051	3.022	0.151
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AHVDALR	3	309.1827	2.638	0.24
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]VQPYLDDFQK[272]K[272]	3	719.4236	3.511	0.367
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LAEYHAK[272]	3	373.8849	2.497	0.262
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QGLLPVLESFK[272]	3	506.976	3.11	0.29
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]LHELQEK[272]	3	528.986	3.602	0.301
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTKK	4	484.5292	3.755	0.313
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]VQPYLDDFQK[272]K[272]	4	539.8195	3.226	0.411
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLATVYVDVLK[272]DSGRDYVSQFEGSALGK[27	4	866.9623	4.971	0.37
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]EQLGPVTQEFWDNLEK[272]ETEGLRQEMSK[27	4	914.2175	4.184	0.425
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VK[272]DLATVYVDVLK[272]DSGR	4	578.3403	2.054	0.087
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDELR	2	721.3754	3.292	0.287
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QGLLPVLESFK[272]	2	759.9604	2.635	0.255
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALEDLR	3	431.5944	1.77	0.125
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKLHELQEK	3	524.9789	3.038	0.158
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSALEEYTK[272]	3	558.9779	2.041	0.043
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKVQPYLDDFQKK	4	535.8124	2.143	0.289
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AELQEGAR	2	507.2724	3.094	0.227
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALEDLR	2	646.8879	3.054	0.362
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LAEYHAK	2	556.3166	1.809	0.12
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LSPLGEEMR	2	588.3142	2.003	0.024
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKVQPYLDDFQKK	3	714.0808	3.742	0.331
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLRAELQEGAR	4	501.7892	2.259	0.321
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ATEHLSTLSEK	1	1495.8116	2.81	0.32
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VQPYLDDFQK	1	1532.8109	2.565	0.33
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LSPLGEEMR	1	1175.6211	2.859	0.282
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEMR	2	586.3106	3.08	0.347
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	WQEEMELYR	2	712.3374	2.209	0.448
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AELQEGAR	2	509.2759	3.27	0.26
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LAEYHAK[272]	2	560.3237	2.878	0.242
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ATEHLSTLSEK	3	499.2754	2.434	0.092
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QGLLPVLESFK	3	504.3046	3.134	0.078
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LHELQEK[272]	3	395.5674	2.438	0.083
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEMR	1	1171.614	2.885	0.254
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]PALEDLR	1	1300.7828	2.561	0.303
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ATEHLSTLSEK[272]	1	1503.8258	2.904	0.31
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLATVYVDVLK[272]	1	1523.8924	2.304	0.492

36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LHELQEK[272]	1	1184.6878	2.564	0.142
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEM[147]R	2	594.3081	2.484	0.379
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	WQEEM[147]ELYR	2	720.3348	2.763	0.24
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LSPLGEEM[147]R	2	596.3116	2.31	0.16
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]PALEDLR	3	434.2658	2.16	0.177
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLRAELQEGAR	4	503.7928	2.728	0.109
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALEDLR	1	1292.7686	2.347	0.245
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LAEYHAK	1	1111.626	2.577	0.272
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AHVDALR	2	461.2669	2.058	0.129
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLRAELQEGAR	3	668.7165	2.687	0.31
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ATEHLSTLSEK[272]	3	501.9468	2.952	0.163
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LHELQEK	1	1176.6736	2.468	0.133
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPYSDELR	2	723.3789	3.166	0.158
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPYSDELR	3	482.5884	1.603	0.091
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AHVDALR	1	921.5265	2.217	0.148
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]PALEDLR	2	650.895	2.245	0.281
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLRAELQEGAR	3	671.3879	3.136	0.242
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEKETEGLRQEMSK	4	911.2122	4.249	0.34
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AHVDALR	1	925.5336	2.202	0.174
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LHELQEK[272]	2	592.8475	2.486	0.1
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEMSK[272]	2	798.4187	3.919	0.371
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEMSK	2	794.4116	3.701	0.38
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AHVDALR	2	463.2704	2.024	0.276
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LEALK[272]ENGGAR	2	723.4194	3.685	0.396
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LREQLGPVTQEFWDNLEK[272]ETEGLR	4	793.6694	3.696	0.254
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LHELQEK	2	588.8404	1.894	0.105
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDELRQR	2	863.4552	3.986	0.349
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DIEEVK[272]	1	1020.5816	2.261	0.111
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LREQLGPVTQEFWDNLEK[272]	3	829.4435	3.576	0.327
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DIEEVK	1	1012.5674	2.152	0.113
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]WQEEM[147]ELYR	2	722.3384	2.269	0.325
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LEALK[272]ENGGAR	3	482.6153	3.44	0.17
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEM[147]SKDLEEVK	4	615.0718	2.183	0.253
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLRAELQEGAR	2	1002.5711	3.772	0.365
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LAEYHAK[272]ATEHLSTLSEK[272]	4	615.842	2.706	0.173
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLR	1	1149.7104	2.291	0.152
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLR	1	1157.7246	2.187	0.156
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LLDNWDSVTSTFSKLR	3	722.7274	2.924	0.22
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEM[147]SK[272]	2	806.4162	3.031	0.377
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LEALKENGGAR	3	479.9439	2.421	0.22
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EQLGPVTQEFWDNLEK[272]	3	739.7151	3.81	0.246
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QEMSK[272]DLEEVK[272]	3	589.989	3.249	0.194
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALEEYTK	3	556.3065	2.172	0.2
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LEALK[272]	1	861.5649	1.823	0.151
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KVSFLSALEEYTK	3	645.7032	4.074	0.308
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LHELQEK	3	392.896	2.576	0.082
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]LHELQEK[272]	2	792.9754	3.148	0.282
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEM[147]SK	2	802.4091	2.592	0.347

36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LEALKENGGAR	2	719.4123	3.431	0.272
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DIEEVK[272]	2	510.7944	2.478	0.158
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DIEEVK	2	506.7873	2.563	0.181
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKLHELQEK	2	786.9647	3.284	0.202
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLEEVK[272]AK[272]	3	455.2768	2.885	0.126
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]TEHLSTLSEK[272]	3	478.2677	3.044	0.249
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VEPIR	2	377.2345	2.11	0.076
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLEEVKAK	3	451.2697	2.799	0.126
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LEALK	2	427.279	1.738	0.19
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLR	2	422.7298	2.282	0.116
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLRAELQEGAR	2	1006.5782	4.187	0.107
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLR	2	575.3588	2.5	0.155
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	WQEEM[147]ELYR	1	1439.6624	1.437	0.255
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LLPVLESFK[272]	2	667.4203	2.757	0.162
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	YVSQFEGSALGK	2	783.4198	3.971	0.38
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LSALEEYTK[272]	2	671.3789	3.697	0.354
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDEL	2	643.3248	3.089	0.476
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEM[147]SK[272]	3	537.9465	1.52	0.076
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLR	2	579.3659	2.646	0.088
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPYSDEL	2	645.3284	2.967	0.361
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSALEEYTK	2	667.3718	3.718	0.319
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DSVTSTFSK	2	626.3326	3.228	0.341
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LEALK[272]	2	431.2861	1.844	0.126
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]WQEEMELYR	2	714.3409	2.082	0.092
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	QKVEPLR	3	383.9083	2.254	0.117
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DEPPQSPWDR	2	683.823	3.125	0.427
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLR	2	424.7334	1.906	0.007
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DSVTSTFSK[272]	2	630.3397	2.529	0.081
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEMSK[272]	3	532.6149	2.374	0.165
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LPVLESFK[272]	2	610.8783	2.046	0.12
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]GLLPVLESFK[272]	2	695.9311	2.456	0.171
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	GLLPVLESFK	2	691.924	2.82	0.212
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LLPVLESFK	2	663.4132	2.942	0.249
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LLDNWDSVTSTFSK	2	948.9949	3.547	0.251
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]SVTSTFSK[272]	2	572.8263	2.64	0.307
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DEPPQSPWDR	2	685.8265	2.638	0.309
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QEEM[147]ELYR	2	629.2987	2.691	0.314
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]TEHLSTLSEK[272]	2	716.898	2.903	0.272
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEM[147]SKDLEEVK	3	819.76	2.643	0.173
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	SALEEYTK	2	610.8297	3.212	0.396
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEMSK	3	529.9435	2.051	0.167
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]TVYVDVLK[272]	2	612.8758	2.487	0.255
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]HLAPYSDELR	2	672.8551	2.754	0.285
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	ETEGLRQEM[147]SK	3	535.2751	1.635	0.186
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEMRDR	3	481.5855	2.082	0.143
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KPALEDLR	2	611.3694	2.466	0.244
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSD	2	522.2615	1.807	0.386
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QK[272]VEPLR	3	386.5797	2.097	0.064

36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VEPLRAELQEGAR	3	537.9693	2.355	0.106
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]SK[272]DLEEVK[272]	3	460.6084	2.858	0.238
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	TEHLSTLSEK	2	712.8909	3.184	0.293
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEM[147]RDR	3	486.9172	2.024	0.189
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKVQPYLDDFQK	3	624.6841	2.197	0.129
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]GLLPVLESFK[272]	3	464.2898	2.765	0.129
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LDDFQK	2	523.2875	2.231	0.163
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALE	2	454.7819	1.84	0.244
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	SKDLEEVK	3	456.6013	2.669	0.175
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DEPPQSPWDR	1	1366.6387	2.018	0.394
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]WQEEM[147]ELYR	1	1443.6695	1.308	0.092
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]PALE	2	458.789	1.994	0.179
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALEDLRQGLLPVLESFK	4	661.8942	2.714	0.063
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]AK[272]VQPYLDDFQK[272]	3	628.6912	1.764	0.083
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QEM[147]SK[272]DLEEVK[272]	3	595.3206	1.951	0.081
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]HLAPYSDELR	3	448.9058	2.095	0.173
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	TVYVDVLK	2	608.8687	2.742	0.047
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	EGSALGK	2	471.2744	1.882	0.071
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEM[147]	2	626.8016	2.584	0.301
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DEPPQSPWDR	1	1370.6458	1.973	0.295
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	AKPALED	2	512.2953	1.405	0.289
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	SKDLEEVK	2	684.3983	2.913	0.182
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]DLATVYVD	1	1039.5429	1.695	0.362
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LSPLGEEM[147]RDR	2	729.8721	1.639	0.139
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KDLATVYVDVLK	3	595.3609	2.732	0.207
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPYS	2	466.7516	1.76	0.126
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]SK[272]DLEEVK[272]	2	690.409	2.625	0.203
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	THLAPYSDE	2	586.7828	1.963	0.113
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]SPLGEEM[147]R	2	539.7696	1.674	0.147
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	LEEYTK	2	531.7952	1.754	0.107
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KLHELQEK	3	482.2927	2.63	0.094
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	VSFLSALE	1	1005.5616	1.54	0.329
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEM	2	618.8042	2.213	0.227
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	DLATVYVD	1	1035.5358	1.562	0.287
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]LDDFQK[272]	2	527.2946	1.759	0.071
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]THLAPY	2	423.2355	1.421	0.194
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KATEHLSTLSEK	3	588.672	2.522	0.1
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]VSFLSA	1	767.442	1.741	0.1
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	KVEPLR	3	341.2221	2.207	0.069
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]ETEGLRQEM[147]S	2	670.3177	2.368	0.187
36	IPI00021841	1	85.4	0.95	0.11	976	##	APOA1 Apolipoprotein A-I precursor	n[145]QGLLPVLESF	2	623.8618	1.662	0.053
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LAVYQAGAR	1	1092.6283	2.812	0.341
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	AKLEEQAQQIR	2	797.4572	4.398	0.486
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	KVEQAVETEPEPELR	2	1017.547	5.073	0.492
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]AATVGSLAGQPLQER	2	821.4557	4.307	0.437
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]GEVQAM[147]LGQSTEELR	2	904.4525	3.55	0.366
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LGPLVEQGR	2	556.8309	2.055	0.212
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]SELEEQLTPVAEETR	2	937.9768	5.135	0.412

37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]VQAAVGTSAAPVPSDNH	2	882.9535	4.623	0.506
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	AKLEEQAQQIR	3	531.9739	2.411	0.344
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]AK[272]LEEQAQQIR	3	534.6453	4.031	0.364
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	ALMDETMK	2	609.8147	3.388	0.352
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]ALM[147]DETMK[272]	2	621.8193	3.069	0.374
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LGADMEDVCGR	2	678.2973	2.831	0.405
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	ALM[147]DETMK	2	617.8122	2.732	0.323
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]ALMDETM[147]K[272]	2	621.8193	2.689	0.396
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]GEVQAMLGQSTEELR	2	896.455	3.765	0.343
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LAVYQAGAR	2	546.8178	3.341	0.265
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]ALM[147]DETM[147]K[272]	2	629.8167	2.638	0.34
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	ALM[147]DETM[147]K	2	625.8096	2.613	0.292
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	LAVYQAGAR	2	544.8142	2.678	0.276
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]ALMDETMK[272]	2	613.8218	2.894	0.262
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	ELQAAQAR	2	513.7882	2.203	0.105
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]WVQTLSEQVQEELLSSQVTQELR	3	958.837	3.084	0.241
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LQAEAFQAR	2	589.3259	1.768	0.141
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]AK[272]LEEQAQQIR	2	801.4643	4.342	0.317
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]ELQAAQAR	2	515.7917	2.198	0.043
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	WELALGR	2	492.7849	2.431	0.182
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]SWFEPLVEDMQR	2	840.9123	2.905	0.28
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]WELALGR	2	494.7885	2.791	0.103
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	DADDLQK	2	542.7853	2.779	0.118
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	KVEQAVETEPEPELR	3	678.7004	4.018	0.344
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LEEQAQQIR	2	629.8472	2.813	0.057
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	ALMDETM[147]K	2	617.8122	2.274	0.171
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]SWFEPLVEDM[147]QR	2	848.9097	2.658	0.105
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	EQVAEVR	2	485.7695	2.075	0.069
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]DADDLQK[272]	2	546.7924	2.261	0.039
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]GLSAIR	2	380.7435	1.949	0.026
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LGADM[147]EDVCGR	2	686.2948	1.632	0.057
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	DRLDEVK	3	385.5558	2.658	0.039
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]DRLDEVK[272]	3	388.2272	2.396	0.016
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	AVYQAGAR	2	488.2722	2.557	0.169
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]LGQSTEELR	2	588.8207	2.436	0.151
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]DRLDEVK[272]	2	581.8372	1.314	0.002
37	IPI00021842	1	69.1	0.75	0.15	74	44	APOE Apolipoprotein E precursor	n[145]QWAGLVEK	2	607.8544	2.028	0.104
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SPELQAEAK	1	1252.6897	3.017	0.39
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SPELQAEAK[272]	1	1260.7039	2.708	0.335
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	EPCVESLVSQYFQTVTDYGK	2	1310.1271	5.335	0.587
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	VKSPELQAEAK	2	810.4777	3.495	0.275
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]CVESLVSQYFQTVTDYGK[272]	2	1201.0866	4.843	0.578
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]EPCVESLVSQYFQTVTDYGK[272]	2	1314.1342	5.208	0.613
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]VK[272]SPELQAEAK[272]	2	816.4883	3.489	0.26
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	CVESLVSQYFQTVTDYGK	3	798.3887	4.312	0.487
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	EPCVESLVSQYFQTVTDYGK	3	873.7538	2.587	0.013
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	VKSPELQAEAK	3	540.6542	2.492	0.276
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]EPCVESLVSQYFQTVTDYGK[272]	3	876.4252	3.575	0.517

38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]VK[272]SPELQAEAK[272]	3	544.6613	3.99	0.194
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]EPCVESLVSQYFQTVTDYGK[272]DLM[147]EK[2	4	851.6673	3.327	0.335
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	EPCVESLVSQYFQTVTDYGKDLM[147]EK	4	848.662	2.821	0.172
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SKEQLTPLIKK	4	462.0476	2.364	0.367
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SYFEK[272]SK[272]EQLTPLIK[272]	4	597.6063	2.22	0.342
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SPELQAEAK	2	626.8485	3.422	0.349
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SPELQAEAK[272]	2	630.8556	3.38	0.403
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SYFEKSKEQLTPLIK	3	791.1298	3.237	0.314
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SKEQLTPLIK	3	526.331	2.414	0.121
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SK[272]EQLTPLIK[272]	3	530.3381	2.207	0.093
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SYFEK[272]SK[272]EQLTPLIK[272]	3	796.4726	2.536	0.168
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]EQLTPLIK[272]	2	615.389	2.237	0.161
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	EQLTPLIK	2	611.3819	2.709	0.214
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SYFEK[272]	1	961.5234	1.955	0.134
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SYFEK[272]	2	481.2653	1.786	0.118
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]SK[272]EQLTPLIK[272]	2	795.0036	1.197	0.014
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SYFEK	1	953.5092	1.787	0.138
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SYFEK	2	477.2582	1.966	0.167
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SKEQLTPLIK	2	788.9929	2.585	0.192
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	EQLTPLIKK	3	497.3204	2.605	0.137
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]EPCVESLVSQYFQTVTDYGK[272]	4	657.5708	2.734	0.074
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	SYFEKSKEQLTPLIK	4	593.5992	1.998	0.196
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]DLMEK[272]	2	462.2592	2.054	0.2
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	DLMEK	2	458.2521	2.001	0.039
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	FQTVTDYGK	2	669.8563	3.252	0.291
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]FQTVTDYGK[272]	2	673.8634	3.108	0.278
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	KSPELQAEAK	3	507.6314	2.878	0.227
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]ELQAEAK[272]	2	538.8132	2.361	0.127
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	ELQAEAK	2	534.8061	2.434	0.128
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]AGTELVNFLSY	1	1357.7121	2.372	0.357
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	KAGTELVNFLSY	2	811.4511	2.106	0.285
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	KSYFEK	3	407.9046	2.615	0.134
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	KEPCVESLVSQYFQTVTDYGK	4	722.6147	2.888	0.208
38	IPI00021854	1	64	0.79	0.06	228	43	APOA2 Apolipoprotein A-II precursor	n[145]YFEK[272]	2	437.7493	1.194	0.17
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	DVSSALDK	1	1114.6104	2.93	0.353
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]DVSSALDK[272]	1	1122.6246	3.053	0.38
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]EFGNTLEDK[272]	1	1340.6937	2.987	0.306
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]EFGNTLEDK[272]	2	670.8505	2.254	0.199
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]IK[272]QSELSAK[272]	2	718.4459	3.761	0.407
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	LKEFGNTLEDK	3	571.9893	4.016	0.378
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]LK[272]EFGNTLEDK[272]	3	575.9964	4.34	0.381
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]IK[272]QSELSAK[272]	3	479.2997	3.675	0.317
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	EWFSETFQK	2	741.3748	3.703	0.381
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	IKQSELSAK	3	475.2926	2.182	0.085
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	EFGNTLEDK	1	1332.6795	3.134	0.222
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	EFGNTLEDK	2	666.8434	3.172	0.225
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]EWFSETFQK[272]	2	745.3819	2.846	0.258
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	LKEFGNTLEDK	2	857.4804	3.527	0.358

39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]QSELSAK[272]	2	525.8053	2.44	0.134
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	QSELSAK	2	521.7982	2.067	0.199
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]LK[272]EFGNTLEDK[272]	2	863.491	3.457	0.34
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	IKQSELSAK	2	712.4353	3.639	0.28
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]EWFSETFQK	2	743.3784	3.058	0.404
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]DVSSALDK[272]	2	561.8159	3.467	0.295
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]DVSSALDK[272]LK[272]	3	503.3067	2.906	0.249
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	DVSSALDK	2	557.8088	2.635	0.304
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]TPDVSSALDK[272]	2	660.8661	2.525	0.239
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	TPDVSSALDK	2	656.859	2.774	0.186
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	IKQSELSAKM[147]R	3	576.3381	1.532	0.217
39	IPI00021855	1	49.4	0.92	0.1	46	26	APOC1 Apolipoprotein C-I precursor	n[145]IK[272]QSELSAK[272]M[147]R	3	580.3452	1.904	0.078
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]ESLSSYWESAK[272]	2	787.9007	4.37	0.445
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]TAAQNLYEK[272]	2	663.3688	3.9	0.32
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	TQQPQQDEM[147]PSPTFLTQVK	3	833.7565	4.577	0.476
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]TQQPQQDEM[147]PSPTFLTQVK[272]	3	836.4279	4.912	0.533
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	STAAM[147]STYTGIFTDQVLSVLK	3	843.7805	3.616	0.425
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	TAAQNLYEK	2	659.3617	3.883	0.335
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]TYLPAVDEK[272]	2	662.3736	2.762	0.293
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	ESLSSYWESAK	2	783.8936	3.342	0.343
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	TYLPAVDEKLRDLYSK	3	777.7739	3.507	0.266
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]STAAM[147]STYTGIFTDQVLSVLK[272]	3	846.4519	3.428	0.256
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	TYLPAVDEK	2	658.3665	2.778	0.275
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]STAAMSTYTGIFTDQVLSVLK[272]GEE	3	946.1558	2.713	0.247
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	TYLPAVDEKLRDLYSK	4	583.5823	2.164	0.138
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	STAAMSTYTGIFTDQVLSVLKGEE	3	943.4844	2.396	0.258
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]DLYSK[272]	2	457.2653	1.878	0.198
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	DLYSK	2	453.2582	1.635	0.17
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	n[145]TGIFTDQVLSVLK[272]	2	855.008	3.865	0.23
40	IPI00021856	1	78.2	0.77	0.03	29	18	APOC2 Apolipoprotein C-II precursor	ESLSSYWESAK[272]	2	785.8971	2.63	0.123
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	DALSSVQESQVAQQAR	2	928.9767	5.545	0.567
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	GWVTDGFSSLK	2	738.8959	3.842	0.414
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	SEAEDASLLSFM[147]QGYM[147]K	2	1110.0216	5.86	0.569
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]DALSSVQESQVAQQAR	2	930.9802	5.13	0.515
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]GWVTDGFSSLK[272]	2	742.903	3.446	0.411
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]SEAEDASLLSFMQGYMK[272]	2	1098.0338	5.303	0.563
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]SEAEDASLLSFM[147]QGYM[147]K[272]	2	1114.0287	5.944	0.527
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	DALSSVQESQVAQQAR	3	619.6535	3.248	0.239
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]DYWSTVK[272]	2	593.821	1.62	0.294
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	DYWSTVK	2	589.8139	2.745	0.271
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]GWVTDGFSSLK	2	740.8995	3.534	0.261
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]GWVTDGFSSLK	3	494.2687	2.215	0.183
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]SEAEDASLLSFMQGYM[147]K[272]	2	1106.0313	5.589	0.551
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	SEAEDASLLSFMQGYM[147]K	2	1102.0242	4.768	0.56
41	IPI00021857	1	51.5	0.72	0.2	24	14	APOC3 Apolipoprotein C-III precursor	n[145]SEAEDASLLSFM[147]QGYM[147]K[272]	3	743.0216	2.643	0.134
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TVIGPDGHK	1	1203.6845	2.721	0.374
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDGHK[272]	1	1211.6987	2.629	0.394
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIMEILR	2	885.4644	4.646	0.388

42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIM[147]EILR	2	893.4618	1.722	0.064
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ESSSHHPGIAEFPSR	2	889.4345	4.859	0.444
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	EVDLKDYEDQQK	2	965.4995	4.798	0.37
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GGSTSYGTGSETESPR	2	856.8898	3.42	0.497
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GLIDEVNQDFTNR	2	830.9181	4.681	0.443
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GSESGIFTNTK	2	710.8752	4.617	0.49
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	HPDEAAFFDTASTGK	2	937.4576	5.841	0.538
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	HRHPDEAAFFDTASTGK	2	1084.0376	6.287	0.555
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	LGEFVSETESR	2	697.3515	3.229	0.535
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MADEAGSEADHEG	2	729.7937	3.66	0.545
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MADEAGSEADHEGTHSTK	2	1076.9819	6.706	0.666
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MKPVPDLVPGNFK	2	931.5397	4.21	0.473
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]ADEAGSEADHEGTHSTK	2	1084.9793	5.447	0.518
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VQHIQLLQK	2	693.9327	3.969	0.414
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ADSGEGDFLAEGGGVR	2	840.9009	3.405	0.373
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DEAAFFDTASTGK[272]	2	824.4089	3.8	0.461
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSGEGDFLAEGGGVR	2	805.3824	2.892	0.259
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSHSLTTNIM[147]EILR	2	895.4654	3.824	0.47
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]EVDLK[272]DYEDQQK[272]	2	971.5102	4.691	0.481
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GGSTSYGTGSETESPR	2	858.8933	2.609	0.458
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GLIDEVNQDFTNR	2	832.9217	4.267	0.538
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GSESGIFTNTK[272]	2	714.8823	4.374	0.476
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]HPDEAAFFDTASTGK[272]	2	941.4647	4.934	0.614
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]HRHPDEAAFFDTASTGK[272]	2	1088.0447	6.28	0.539
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MADEAGSEADHEGTHSTK[272]	2	1080.989	6.697	0.673
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NSLFEYQK[272]	2	658.8581	3.41	0.234
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPMLGEFVSETESR	2	1205.0813	6.174	0.504
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPM[147]LGEFVSETESR	2	1213.0788	5.059	0.521
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VQHIQLLQK[272]	2	697.9398	3.95	0.417
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIMEILR	3	590.6453	3.079	0.134
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIM[147]EILR	3	595.977	4.824	0.509
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ESSSHHPGIAEFPSR	3	593.2921	3.306	0.324
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GDFSSANNRDNTYNR	3	624.2857	2.933	0.391
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	HPDEAAFFDTASTGK	3	625.3075	4.688	0.385
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	HRHPDEAAFFDTASTGK	3	723.0275	5.378	0.557
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MADEAGSEADHEGTHSTK	3	718.3237	3.999	0.512
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]ADEAGSEADHEGTHSTK	3	723.6553	3.74	0.61
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	NNKDSHSLTTNIM[147]EILR	3	761.4022	4.451	0.476
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TVIGPDGHK	3	401.8997	1.822	0.141
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSHSLTTNIMEILR	3	591.981	4.051	0.381
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSHSLTTNIM[147]EILR	3	597.3127	2.419	0.288
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ESSSHHPGIAEFPSR	3	594.6278	3.825	0.452
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]HRHPDEAAFFDTASTGK[272]	3	725.6989	5.635	0.476
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MADEAGSEADHEGTHSTK[272]	3	720.9951	3.81	0.46
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NNK[272]DSHSLTTNIM[147]EILR	3	764.0736	4.523	0.473
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPM[147]LGEFVSETESR	3	809.0549	2.525	0.226
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDGHK[272]	3	404.5711	1.445	0.175
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VIEK[272]VQHIQLLQK[272]	3	670.093	4.904	0.569

42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MADEAGSEADHEGTHSTK	4	538.9946	2.614	0.201
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]ADEAGSEADHEGTHSTK	4	542.9933	4.717	0.451
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GSESGIFTNTK[272]ESSSHHPGIAEFPSR	4	762.632	4.499	0.483
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MADEAGSEADHEGTHSTK[272]	4	540.9981	5.594	0.469
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MADEAGSEADHEGTHSTK[272]R	4	580.0234	3.286	0.385
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VIEK[272]VQHIQLLQK[272]	4	502.8216	2.975	0.454
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GSESGIFTNTK[272]ESSSHHPGIAEFPSR	5	610.3071	3.584	0.327
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ESSSHHPGIAEFPSR	2	891.438	3.128	0.44
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	EKVTSGSTTTTR	3	516.6178	2.956	0.347
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VIEKVQHIQLLQK	3	666.0859	3.023	0.427
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VQHIQLLQK	3	462.9575	3.29	0.152
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]EVDLK[272]DYEDQQK[272]	3	648.0092	3.613	0.285
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MELERPGGNEITR	3	549.2865	2.813	0.277
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NNK[272]DSHSLTTNIMEILR	3	758.742	4.175	0.393
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDGHK[272]EVTK[272]	3	605.023	3.581	0.406
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VQHIQLLQK[272]	3	465.6289	4.049	0.272
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ESSSHHPGIAEFPSR	4	445.2209	1.855	0.021
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VIEKVQHIQLLQK	4	499.8162	2.837	0.379
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDGHK[272]EVTK[272]	4	454.0191	3.071	0.352
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]KPVPDLVPGNFK	2	939.5372	2.909	0.232
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	NSLFEYQK	2	654.851	2.693	0.245
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSHSLTTNIMEILR	2	887.4679	4.245	0.379
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MK[272]PVPDLVPGNFK[272]	2	937.5504	3.595	0.454
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDGHK[272]	2	606.353	1.416	0.161
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MELERPGGNEITR	3	547.9508	2.604	0.115
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	NNKDSHSLTTNIMEILR	3	756.0706	3.682	0.303
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]EK[272]VTSGSTTTTR	3	519.2892	2.722	0.352
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	AQLVDMK	2	542.8128	2.492	0.234
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GDSTFESK	2	575.7906	2.129	0.189
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MELERPGGNEITR	2	821.4225	3.414	0.324
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	QFTSSTSYNR	2	665.823	2.473	0.14
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]AQLVDMK[272]	2	546.8199	2.625	0.339
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GDSTFESK[272]	2	579.7977	2.411	0.225
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GGSTSYGTGSETESPR	3	572.9313	2.69	0.284
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MK[272]PVPDLVPGNFK[272]	3	625.3693	2.773	0.214
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	NSLFEYQK	1	1308.6948	2.743	0.277
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DYEDQQK	2	603.2935	2.813	0.314
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DYEDQQK[272]	2	607.3006	2.769	0.287
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MELERPGGNEITR	2	823.4261	2.05	0.114
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GSESGIFTNTKESSSHHPGIAEFPSR	4	760.6285	3.832	0.36
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NNK[272]DSHSLTTNIM[147]EILR	4	573.3071	3.491	0.24
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GSESGIFTNTKESSSHHPGIAEFPSR	5	608.7042	2.917	0.147
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GDSTFESK	1	1150.574	2.472	0.337
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NSLFEYQK[272]	1	1316.709	2.425	0.305
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GDFSSANNR	2	554.2625	1.9	0.267
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TVIGPDGHK	2	602.3459	3.099	0.243
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GDSTFESK[272]	1	1158.5882	2.485	0.314
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GDFSSANNRDNTYNR	3	625.6214	1.924	0.203

42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ALTDMPQMR	2	601.8047	2.677	0.31
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]QFTSSTSYNR	2	667.8265	2.121	0.161
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	EVDLKDYEDQQK	3	644.0021	2.183	0.12
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TVIGPDGHKEVTK	3	601.0159	2.863	0.286
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ALTDMPQMR	2	603.8082	2.551	0.317
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GDFSSANNR	2	556.2661	1.501	0.093
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ESSSHHPGIAEFPSR	4	446.2227	2.822	0.102
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ALTDM[147]PQM[147]R	2	617.7996	1.729	0.064
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	QLEQVIAK	2	604.8717	2.808	0.125
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]QLEQVIAK[272]	2	608.8788	3.106	0.121
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	AQLVDM[147]K	2	550.8103	1.919	0.058
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]RLEVDIDIK[272]	3	463.6166	2.281	0.007
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ALTDM[147]PQM[147]R	2	619.8031	1.926	0.236
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]AQLVDM[147]K[272]	2	554.8174	1.326	0.094
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SSHHPGIAEFPSR	3	522.6029	3.548	0.317
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VTSGSTTTTR	2	577.8103	2.359	0.298
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	IRPLVTQ	2	483.8084	1.853	0.268
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]QHLPLIK[272]	2	568.8734	2.069	0.211
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	NSLFEYQK	3	436.9031	3.095	0.073
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VTSGSTTTTR	2	575.8068	2.182	0.231
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	RLEVDIDIK	1	1380.821	1.971	0.129
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]ELERPGGNEITR	3	553.2824	1.462	0.094
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SHHPGIAEFPSR	3	493.5922	2.65	0.355
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ALTDM[147]PQMR	2	611.8057	2.417	0.185
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VPPEWK[272]	2	522.3101	2.094	0.184
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	SSHHPGIAEFPSR	3	521.2672	2.949	0.317
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	AQLVDM[147]KR	2	628.8609	2.307	0.12
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	RLEVDIDIK	3	460.9452	2.451	0.115
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]MADEAGSEADHEGTH	3	567.5695	2.603	0.331
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]KPVPDLVPGNFK	3	626.6939	2.923	0.092
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]ALTDMPQM[147]R	2	611.8057	2.534	0.334
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DILPSR	2	422.7541	1.772	0.006
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SHSLTTNIM[147]EILR	3	558.9704	3.306	0.261
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	RLEVDIDIK	2	690.9141	2.224	0.05
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]FTSSTSYNR	2	603.7972	2.966	0.478
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]LVTSK[272]GDK[272]	2	640.401	2.432	0.265
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VPPEWK	2	518.303	1.867	0.106
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]PGSTGTWNPGSSER	2	788.8773	4.012	0.388
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]RLEVDIDIK[272]	2	694.9212	2.295	0.066
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]AQLVDM[147]K[272]R	3	422.2477	1.688	0.167
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]RK[272]VIEK[272]	4	301.958	2.259	0.086
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ALTDMPQM[147]R	2	609.8021	2.388	0.302
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPMLGEFVSETESR	3	803.7233	2.611	0.087
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SLTTNIM[147]EILR	2	725.9064	3.513	0.247
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]LK[272]NSLFEYQK[272]	3	568.0015	2.602	0.122
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	MKPVPDLVPGNFK	3	621.3622	2.441	0.041
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DEAAFFDTASTGK	2	820.4018	3.728	0.412
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VSEDLR	2	431.7412	1.701	0.064

42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DLIPSR	2	420.7505	1.92	0.016
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]NSLFEYQK[272]	3	439.5745	1.956	0.106
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SLTTNIMEILR	2	717.909	3.462	0.215
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	LVTSKGDKELR	3	556.0052	2.41	0.125
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]STSYGTGSETESPR	2	801.8718	2.758	0.465
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VSEDLR	2	429.7376	1.918	0.04
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	GSESGIFTNTK	3	474.2526	2.521	0.048
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	SLTTNIM[147]EILR	2	723.9029	3.574	0.338
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	LVTSKGDK	2	634.3903	2.249	0.183
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GDFSSANNRDNTYNR	2	937.9285	2.699	0.392
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]RLEVDIDIK[272]	1	1388.8352	1.694	0.22
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	EVDLK	2	442.266	1.828	0.087
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DFLAEGGGVR	2	582.8101	3.228	0.313
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	KVIEK	2	518.8476	2.301	0.075
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ALTDM[147]PQMR	2	609.8021	1.557	0.064
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]LVTSK[272]	2	418.2782	1.601	0.179
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	QHLPLIK	2	564.8663	1.509	0.028
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]LGEFVSETESR	2	699.3551	2.83	0.283
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SSHHPGIAEFPSR	2	783.4007	3.391	0.353
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SGEGDFLAEGGGVR	2	747.8689	2.9	0.255
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	M[147]ELERPGGNEITR	2	829.42	1.239	0.009
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GEFVSETESR	2	642.8131	2.521	0.302
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	LKDYEDQQK	3	529.6228	2.765	0.227
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	LKNSLFEYQK	3	563.9944	2.016	0.102
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPM[147]LG	2	680.84	2.051	0.363
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIM[147]	2	637.8059	2.491	0.224
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]DSHSLTTNIMEIL	2	809.4174	2.186	0.278
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	SLFEYQK	2	597.8295	2.571	0.123
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VSEDLRSR	2	551.3042	1.826	0.061
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SLFEYQK[272]	2	601.8366	2.419	0.131
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SSSHHPGIAEFPSR	3	551.6136	2.751	0.153
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]RK[272]VIEK[272]	3	402.2749	1.869	0.022
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TVIGPDG	1	798.4357	2.272	0.392
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]HPDEAAFFDTASTG	2	805.3662	2.208	0.207
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	SSHHPGIAEFPSR	2	781.3972	2.803	0.299
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	RKVIEK	2	596.8981	1.565	0.018
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GSESGIFTNTK[272]	3	476.924	1.869	0.017
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]EVNQDFTNR	2	633.8134	1.923	0.25
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	DSHSLTTNIM	2	629.8085	2.306	0.189
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPM[147]	2	595.7873	1.734	0.181
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]SHSLTTNIM[147]EILR	2	837.9519	2.738	0.133
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	ALTDM[147]PQM[147]R	3	412.2021	1.446	0.02
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]IMEILR	2	459.7818	2.002	0.079
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]IM[147]EILR	2	467.7793	2.077	0.061
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	VQHIQLL	2	495.8084	1.504	0.11
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	TFPGFFSPM[147]	2	593.7837	1.534	0.136
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]VIGPDGHK[272]	2	555.8292	1.859	0.07
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]GSAGHWTSESSVSG	2	746.8429	2.084	0.054

42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TFPGFFSPM	2	587.7898	1.813	0.004
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]TVIGPDG	1	802.4428	1.825	0.067
42	IPI00021885	1	42.1	0.91	0.17	521	##	FGA Isoform 1 of Fibrinogen alpha chain precursor	n[145]LK[272]DYEDQQK[272]	3	533.6299	2.223	0.088
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DNCCILDER	1	1316.5224	2.197	0.387
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	AIQLTYNPDESSKPNMIDAATLK	2	1470.7787	5.094	0.532
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	DTVQIHDITGK	2	753.9174	2.746	0.228
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	EGFGHLSPTGTTEFWLGNEK	2	1244.1188	6.265	0.484
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	EGFGHLSPTGTTEFWLGNEK[272]	2	1246.1224	4.795	0.417
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	GIADFLSTYQTK	2	812.4407	3.051	0.398
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLGGAK	2	913.5052	6.014	0.506
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYFIKPLK	2	857.5244	1.858	0.052
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	TSTADYAMFK	2	707.8554	4.457	0.417
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	TSTADYAM[147]FK	2	715.8529	3.863	0.469
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YLQEIYNSNNQK	2	897.4627	4.713	0.457
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PN	2	983.012	4.094	0.355
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PNMIDAATLK[272]	2	1476.7893	4.506	0.499
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DTVQIHDITGK[272]	2	757.9245	2.184	0.264
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]EGFGHLSPTGTTEF	2	812.3922	4.052	0.387
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]EGFGHLSPTGTTEFWLGNEK[272]	2	1248.1259	5.966	0.53
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]GIADFLSTYQTK[272]	2	816.4478	4.5	0.481
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGGAK[272]	2	917.5123	5.644	0.542
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYFIK[272]PLK[272]	2	863.5351	4.383	0.491
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TCGIADFLSTYQTK[272]	2	941.4701	4.266	0.52
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TSTADYAMFK[272]	2	711.8625	4.331	0.491
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TSTADYAM[147]FK[272]	2	719.86	3.398	0.494
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TVQIHDITGK[272]	2	700.411	4.305	0.445
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VAQLEAQCQEPCK[272]	2	913.9292	3.131	0.474
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YEASILTHDSSIR	2	818.4266	2.532	0.345
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YLQEIYNSNNQK[272]	2	901.4698	4.827	0.399
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	AIQLTYNPDESSKPNMIDAATLK	3	980.8549	5.053	0.466
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	AIQLTYNPDESSKPNM[147]IDAATLK	3	986.1865	4.989	0.469
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	CHAGHLNGVYYQGGTYSK	3	761.0262	6.364	0.582
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	DTVQIHDITGK	3	502.9474	3.135	0.345
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	EGFGHLSPTGTTEFWLGNEK	3	829.7483	4.738	0.529
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	FAGGDAGDAFDGFDFGDDPSDK	3	835.0277	4.955	0.548
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KMLEEIMKYEASILTHDSSIR	3	972.1892	4.613	0.483
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLGGAK	3	609.3392	2.59	0.105
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VAQLEAQCQEPCKDTVQIHDITGK	3	1056.1887	5.282	0.512
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YLQEIYNSNNQK	3	598.6442	4.373	0.402
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PNMIDAATLK[272]	3	984.862	3.571	0.279
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	3	990.1936	4.764	0.525
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]CHAGHLNGVYYQGGTYSK[272]	3	763.6976	6.804	0.526
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DTVQIHDITGK[272]	3	505.6188	2.543	0.318
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FAGGDAGDAFDGFDFGDDPSDK[272]	3	837.6991	5.273	0.599
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IHLISTQSAIPYALR	3	609.6918	3.428	0.234
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLG	3	478.5921	4.393	0.484
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGGAK[272]	3	612.0106	3.661	0.316
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYFIK[272]PLK[272]	3	576.0258	3.663	0.352

43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VAQLEAQCQEPCK[272]DTVQIHDITGK[272]	3	1060.1958	6.647	0.576
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YEASILTHDSSIR	3	545.9535	1.619	0.018
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KMLEEIMKYEASILTHDSSIR	4	729.3937	5.105	0.517
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLGGAK	4	457.2563	3.972	0.446
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYFIKPLK	4	429.2659	1.97	0.306
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VAQLEAQCQEPCKDTVQIHDITGK	4	792.3934	4.279	0.383
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGGAK[272]	4	459.2598	2.972	0.326
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYFIK[272]PLK[272]	4	432.2712	2.567	0.337
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VAQLEAQCQEPCK[272]DTVQIHDITGK[272]	4	795.3987	5.295	0.4
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KMLEEIMKYEASILTHDSSIR	5	583.7164	4.728	0.502
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYFIKPLK	3	572.0187	2.77	0.129
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	RLDGSVDFK	3	439.5822	4.1	0.236
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VGPEADKYR	3	438.9104	3.208	0.4
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YLQEIYNSNNQK[272]	3	601.3156	3.845	0.254
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	4	742.897	3.57	0.303
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]RLDGSVDFK[272]K[272]	4	399.9913	1.638	0.129
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LDGSVDFKK	2	714.9142	2.941	0.408
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YEASILTHDSSIR	2	816.423	3.879	0.4
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IHLISTQSAIPYALR	2	914.0341	1.996	0.145
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIMK	3	486.6122	1.855	0.026
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]RLDGSVDFK[272]	3	442.2536	2.947	0.247
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	DCQDIANK	2	616.7918	2.496	0.194
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VAQLEAQCQEPCK	2	909.9221	2.805	0.445
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DCQDIANK[272]	2	620.7989	2.82	0.28
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DNCCILDER	2	658.7648	2.981	0.348
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]RLDGSVDFK[272]K[272]	2	798.9754	3.28	0.353
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	EGFGHLSPTGTTEFWLGNEK[272]	3	831.084	4.4	0.389
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	AIQLTYNPDESSKPNMIDAATLK	4	735.893	3.439	0.235
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIM[147]KYEASILTHDSSIR	4	737.3912	3.365	0.256
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DCQDIANK[272]	1	1240.5905	2.453	0.308
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	IHLISTQSAIPYALR	3	608.3561	2.856	0.191
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	DCQDIANK	1	1232.5763	2.406	0.352
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LDGSVDFK[272]	1	1168.6453	2.342	0.262
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	RLDGSVDFKK	2	792.9647	3.362	0.24
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VELEDWNGR	2	629.3148	2.308	0.088
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YEASILTHDSSIR	3	544.6178	2.921	0.077
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]EGFGHLSPTGTTEFWLGNEK[272]	3	832.4197	3.443	0.259
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VGPEADK[272]YR	3	441.5818	2.251	0.103
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VELEDWNGR	1	1257.6223	2.928	0.147
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VELEDWNGR	1	1261.6294	2.673	0.18
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VELEDWNGR	2	631.3183	3.031	0.313
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LDGSVDFK	1	1160.6311	2.327	0.233
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLG	3	477.2564	3.871	0.321
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGGA	3	521.2783	3.566	0.346
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LDGSVDFK[272]	2	584.8263	2.144	0.151
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VGPEADK[272]	2	502.2868	2.179	0.28
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TVQIHDITGK[272]	3	467.2764	2.782	0.376
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LDGSVDFK	2	580.8192	1.873	0.032

43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHL	3	459.5849	3.542	0.315
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]MLEEIMK[272]	2	591.3293	2.962	0.134
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	RLDGSVDFK	2	658.8697	2.103	0.095
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VGPEADKYR	2	657.8619	2.808	0.375
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VGPEADK[272]YR	1	1322.7308	2.449	0.2
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KMLEEIMK	2	721.4172	2.78	0.129
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]RLDGSVDFK[272]	2	662.8768	2.291	0.16
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]VGPEADK[272]YR	2	661.869	2.769	0.276
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]RLDGSVDFK[272]K[272]	3	532.986	1.735	0.026
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIM[147]K	2	737.4121	2.285	0.204
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	VGPEADK	2	498.2797	2.189	0.169
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]SSK[272]PNM[147]IDAATLK[272]	3	608.6802	2.767	0.347
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	2	1484.7868	3.126	0.486
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	RLDGSVDFKK	4	396.986	2.457	0.278
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	MLEEIMK	2	587.3222	2.417	0.101
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	M[147]LEEIM[147]K	2	603.3171	2.223	0.107
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHH	3	421.8903	2.967	0.282
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TIGEGQQHHLGGAK[272]	3	574.316	3.612	0.27
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LDGSVDFKK	3	476.9452	1.837	0.188
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]MLEEIM[147]K[272]	1	1197.6462	1.765	0.072
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	IVNLKEK	2	632.4111	2.819	0.118
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	MLEEIM[147]K	1	1189.632	1.934	0.082
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]MLEEIM[147]K[272]YEASILTHDSSIR	4	668.3485	2.549	0.25
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LDGSVDFK[272]K[272]	2	720.9248	2.054	0.087
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LDGSVDFK[272]K[272]	3	480.9523	1.955	0.135
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	MLEEIM[147]K	2	595.3196	2.315	0.031
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]TSTADYAM[147]FK[272]	3	480.2424	1.865	0.191
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLGG	3	496.2636	3.146	0.254
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LISTQSAIPYALR	2	788.9626	4.069	0.373
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	TVQIHDITGK	2	696.4039	3.838	0.335
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	TSTADYAM[147]FK	3	477.571	2.856	0.018
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]GIADFLSTYQTK[272]	3	544.6343	2.582	0.283
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LYFIKPLK	2	721.4684	3.112	0.352
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGG	2	745.8953	3.614	0.358
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQHHLG	2	715.381	3.461	0.335
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IVNLK[272]EK[272]	2	638.4217	2.263	0.11
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLG	2	717.3845	3.463	0.384
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	GLYFIKPLK	2	749.9791	2.89	0.437
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYFIK[272]	2	622.3681	2.709	0.342
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]SGLYFIK[272]PLK[272]	3	533.3396	3.119	0.258
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	AIQLTYNPDESSKPN	2	979.0049	4.288	0.405
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	FFTSHNGM[147]QF	2	686.3112	2.747	0.326
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	SGLYFIKPLK	2	793.4951	3.781	0.265
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYFIK	2	618.361	2.567	0.286
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]GLYFIK[272]PLK[272]	2	755.9898	3.012	0.296
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LYFIK[272]PLK[272]	2	727.479	3.091	0.256
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]DTVQIHDITG	2	621.826	2.288	0.391
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LYFIKPLK	3	481.3147	1.885	0.271

43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]SGLYFIK[272]PLK[272]	2	799.5058	3.404	0.239
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	DTVQIHDITG	2	619.8224	2.759	0.286
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHLGG	3	497.5993	2.822	0.184
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FFTSHNGM[147]QF	2	688.3147	2.877	0.265
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	SGLYFIKPLK	3	529.3325	2.977	0.136
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]ISTQSAIPYALR	2	732.4206	3.391	0.321
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQHHL	2	688.8738	2.869	0.216
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YEASILTHDSSIR	4	409.7169	1.687	0.015
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YEASILTHDS	2	638.3144	1.862	0.352
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIMKYEASILTHDSSIR	4	733.3924	3.879	0.427
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IHLISTQ	2	478.2883	2.054	0.266
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KMLEEIM[147]KYEASILTHDSSIR	4	733.3924	3.519	0.472
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FFTSHN	2	448.7228	2.161	0.188
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYFIK[272]PL	2	727.4365	2.447	0.395
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]EGFGHLSPTGTT	2	674.3367	2.533	0.223
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YEASILTHDS	2	640.318	2.22	0.234
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FFTSH	2	391.7013	1.92	0.227
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIM[147]KYEASILTHDSSIR	3	982.8525	2.812	0.121
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IHLISTQSAIPY	2	743.923	2.414	0.205
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	KM[147]LEEIMKYEASILTHDSSIR	3	977.5208	3.306	0.46
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYFIKPL	2	723.4294	1.875	0.23
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YFIK[272]PLK[272]	2	670.937	2.033	0.13
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YVATR	2	375.2189	1.994	0.107
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	GLYFIKPLK	3	500.3218	2.139	0.144
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]HLISTQSAIPYALR	2	857.4921	2.743	0.248
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	M[147]IDAATLK	2	579.8312	2.801	0.147
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	YFIKPLK	2	664.9263	2.041	0.169
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]SILTHDSSIR	3	424.9058	2.689	0.065
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	FFTSHN	2	446.7192	1.767	0.192
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]IHLISTQSAIPYAL	2	835.9835	1.722	0.273
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	IHLISTQSAIPY	2	741.9194	2.17	0.169
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	MIDAATLK	2	571.8338	2.63	0.131
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	FFTSH	2	389.6978	1.74	0.137
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]GLYFIK[272]PLK[272]	3	504.3289	2.159	0.114
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	IHLISTQ	2	476.2848	1.79	0.173
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FFTSH	1	782.3954	1.633	0.216
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]MIDAATLK[272]	2	575.8409	2.197	0.172
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	TCGIADFLSTYQTK	2	937.463	4.226	0.314
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	SAIPYALR	2	515.8059	2.267	0.008
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LYFIK[272]PLK[272]	3	485.3218	2.426	0.045
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]YEASILTHD	2	596.802	1.94	0.09
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTIGEGQQH	2	561.7988	1.874	0.079
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	QSGLYF	1	854.4407	1.393	0.317
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]FIK[272]PLK[272]	3	393.2726	2.208	0.024
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	IHLISTQSAIPYAL	2	833.98	1.798	0.058
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	FFTSH	1	778.3883	1.635	0.134
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]QSGLYF	1	858.4478	1.427	0.281
43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	n[145]LTIGEGQQH	2	563.8023	1.31	0.145

43	IPI00021891	1	63.4	N_A	N_A	N_A	99	FGG Isoform Gamma-B of Fibrinogen gamma chain precursor	LTYAYF	1	917.4768	1.334	0.289
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ATFQTPDFIVPLTDLR	2	987.5384	4.37	0.434
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ATGVLYDYVNK	2	761.9169	3.637	0.507
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EVGTVLSQVYSK	2	795.4485	4.291	0.334
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	GNVATEISTER	2	658.8439	3.649	0.443
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KGNVATEISTER	2	792.9389	4.231	0.48
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LLSGGNTLHLVSTTK	2	911.0333	5.16	0.562
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ITENDIQIALDDAK	2	919.9966	4.539	0.436
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NIQEYLSILTDPDGK	2	993.5308	3.292	0.533
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NNALDFVTK	2	651.3643	3.685	0.421
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NSEEFAAAMSR	2	676.8168	3.568	0.463
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVSDGIAALDLNAVANK	2	969.5364	5.665	0.398
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TKNSEEFAAAM[147]SR	2	869.4331	3.669	0.494
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TIEMSAFGYVFPK	2	885.4684	4.031	0.436
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TSSFALNLPTLPEVK	2	949.0433	4.945	0.456
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VLVDHFGYTK	2	729.9089	4.134	0.517
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VNWEEEAASGLLTSLK	2	1014.0441	6.504	0.545
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ATGVLYDYVNK[272]	2	765.924	4.846	0.475
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AVSM[147]PSFSILGSDVR	2	863.4517	3.584	0.339
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EVGTVLSQVYSK[272]	2	799.4556	4.257	0.335
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]GFEPTLEALFGK[272]	2	798.9475	4.344	0.474
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IADFELPTIIVPEQTIEIPSIK[272]	2	1377.7928	4.212	0.516
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LLSGGNTLHLVSTTK[272]	2	915.0404	5.559	0.572
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LPQQANDYLNSFNWER	2	1070.0224	4.332	0.478
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ITENDIQIALDDAK[272]	2	924.0037	4.895	0.481
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NNALDFVTK[272]	2	655.3714	3.303	0.434
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NSEEFAAAM[147]SR	2	686.8178	3.694	0.466
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SVSDGIAALDLNAVANK[272]	2	973.5435	5.499	0.399
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TK[272]NSEEFAAAM[147]SR	2	873.4402	3.187	0.546
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TIEMSAFGYVFPK[272]	2	889.4755	4.636	0.49
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TSSFALNLPTLPEVK[272]	2	953.0504	4.66	0.52
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VLVDHFGYTK[272]	2	733.916	3.578	0.49
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VNDESTEGK[272]TSYR	2	887.4465	3.724	0.462
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VNWEEEAASGLLTSLK[272]	2	1018.0512	5.146	0.468
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YTYNYEAESSSGVPGTADSR	2	1149.5176	6.378	0.589
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AEPLAFTFSHDYK	3	602.6456	3.843	0.359
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AHLDIAGSLEGHLR	3	543.6338	5.202	0.398
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ASGSLPYTQTLQDHLNSLK	3	785.0885	4.697	0.4
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DKDQEVLLQTFLDDASPGDKR	3	937.4943	3.571	0.507
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	GISTSAASPAVGTVGM[147]DM[147]DEDDDFSK	3	972.4356	5.865	0.574
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KGNVATEISTER	3	528.9617	4.645	0.413
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KIISDYHQQFR	3	572.3178	3.809	0.436
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KLTISEQNIQR	3	537.3177	2.091	0.229
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KM[147]GLAFESTK	3	516.6253	3.103	0.375
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KM[147]TSNFPVDLSDYPK	3	726.7137	4.329	0.474
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KYTYNYEAESSSGVPGTADSR	3	854.7418	4.207	0.361
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IHSGSFQSQVELSNDQEK	3	771.7205	4.571	0.373
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ILGEELGFASLHDLQLLGK	3	778.4458	6.146	0.581

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LLLQM[147]DSSATAYGSTVSKR	3	775.4142	3.194	0.336
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NHLQLEGLFFTNGEHTSK	3	784.7414	4.731	0.438
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QVFLYPEKDEPTYILNIK	3	877.4913	5.05	0.47
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QVFLYPEKDEPTYILNIKR	3	929.525	4.681	0.468
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SGVQM[147]NTNFFHESGLEAHVALK	3	904.7903	5.934	0.605
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SKPTVSSSM[147]EFK	3	588.6502	4.932	0.407
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLDEHYHIR	3	437.2264	3.603	0.432
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SNTVASLHTEK	3	489.6036	3.558	0.331
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TILGTM[147]PAFEVSLQALQK	3	748.4205	3.687	0.371
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TSQCTLKEVYGFNPEGK	3	789.7327	5.047	0.45
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VLVDHFGYTK	3	486.9417	2.617	0.145
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AEPLAFTFSHDYK[272]	3	605.317	4.63	0.559
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AHLDIAGSLEGHLR	3	544.9695	3.241	0.4
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ASGSLPYTQTLQDHLNSLK[272]	3	787.7599	4.884	0.419
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ATGVLYDYVNK[272]	3	510.9517	3.669	0.357
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LLLQM[147]DSSATAYGSTVSK[272]R	3	778.0856	4.345	0.501
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NK[272]YGM[147]VAQVTQTLK[272]	3	677.0541	4.763	0.466
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QVFLYPEK[272]DEPTYILNIK[272]	3	881.4984	4.386	0.416
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SGVQM[147]NTNFFHESGLEAHVALK[272]	3	907.4617	5.909	0.498
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SK[272]PTVSSSM[147]EFK[272]	3	592.6573	4.031	0.343
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SPAFTDLHLR	3	434.2425	3.995	0.361
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TK[272]NSEEFAAAM[147]SR	3	582.6292	4.825	0.439
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TIHDLHLFIENIDFNK[272]	3	753.0788	4.273	0.407
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TILGTM[147]PAFEVSLQALQK[272]	3	751.0919	2.357	0.29
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AHLDIAGSLEGHLR	4	407.9772	4.49	0.466
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DKDQEVLLQTFLDDASPGDKR	4	703.3726	4.134	0.508
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QVFLYPEKDEPTYILNIKR	4	697.3956	3.337	0.305
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LEIQSQVDSQHVGHSVLTAK[272]	4	616.8404	3.634	0.154
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NHLQLEGLFFTNGEHTSK[272]	4	590.8114	4.386	0.364
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QVFLYPEK[272]DEPTYILNIK[272]R	4	700.4009	4.749	0.499
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SGVQM[147]NTNFFHESGLEAHVALK[272]	4	680.8481	3.865	0.505
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	GFEPTLEALFGK	2	794.9404	3.631	0.434
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QGFFPDSVNK[272]	2	713.8821	3.574	0.335
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ATGVLYDYVNK	3	508.2803	2.897	0.282
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DKAQNLYQELLTQEGQASFQGLK	3	1010.54	3.642	0.361
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EFNLQNM[147]GLPDFHIPENLFLK	3	938.1546	4.22	0.383
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IISDYHQQFR	3	482.9211	2.667	0.32
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LTLDIQNKK	3	498.3119	2.758	0.296
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SPAFTDLHLR	3	432.9068	3.439	0.272
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	STSPPKQAEAVLK	3	592.6841	3.349	0.342
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TKNSEEFAAAM[147]SR	3	579.9578	3.608	0.358
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FFGEGTK[272]K[272]	3	449.2662	2.616	0.357
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]HINIDQFVR	3	429.2426	2.816	0.349
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ILGEELGFASLHDLQLLGK[272]	3	781.1172	3.059	0.318
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IISDYHQQFR	3	484.2568	2.529	0.174
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LLSGGNTLHLVSTTK[272]	3	610.3627	3.021	0.284
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]INDVLEHVK[272]	3	452.2693	3.178	0.238
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NSLK[272]IEIPLPFGGK[272]	3	649.0664	3.974	0.365

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QVFLYPEK[272]DEPTYILNIK[272]R	3	933.5321	4.364	0.342
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SISAALEHK[272]	3	415.2465	4.079	0.226
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SNTVASLHTEK[272]	3	492.275	3.299	0.329
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]STSPPK[272]QAEAVLK[272]	3	596.6912	2.72	0.044
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TSSFALNLPTLPEVK[272]	3	635.7027	2.342	0.022
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VLVDHFGYTK[272]	3	489.6131	1.98	0.076
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VPLLLSEPINIIDALEM[147]R	3	732.7525	4.117	0.369
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VPSYTLILPSLELPVLHVPR	3	796.4776	4.309	0.355
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]WNFYYSPQSSPDKK	3	724.3729	4.17	0.387
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AHLDIAGSLEGHLR	4	408.9789	2.928	0.262
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YHWEHTGLTLR	4	389.9577	3.031	0.267
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ALVEQGFTVPEIK	2	855.9931	3.85	0.403
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KIISDYHQQFR	2	857.973	4.028	0.386
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KLTISEQNIQR	2	805.4729	3.461	0.406
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IAELSATAQEIIK	2	833.9906	4.314	0.336
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	MGLAFESTK	2	632.342	3	0.397
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NPNGYSFSIPVK	2	801.9356	4.038	0.346
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SISAALEHK	2	618.359	3.466	0.35
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVSLPSLDPASAK	2	776.4407	3.38	0.189
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VSTAFVYTK	2	648.3716	2.945	0.438
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LAAYLM[147]LM[147]R	2	629.3444	2.395	0.372
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IAELSATAQEIIK[272]	2	837.9977	4.454	0.264
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IISDYHQQFR	2	725.8816	3.257	0.419
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NIQEYLSILTDPDGK[272]	2	997.5379	3.362	0.424
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NPNGYSFSIPVK[272]	2	805.9427	3.851	0.379
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SNTVASLHTEK[272]	2	737.9089	3.581	0.382
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SVSLPSLDPASAK[272]	2	780.4478	3.103	0.09
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TLQGIPQM[147]IGEVIR	2	857.9858	2.598	0.184
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VNDESTEGK[272]	2	633.8245	3.1	0.423
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VSTAFVYTK[272]	2	652.3787	2.904	0.421
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FQFPGKPGIYTR	3	564.3191	2.803	0.271
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SISAALEHK	3	412.5751	2.925	0.229
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]CVQSTK[272]PSLM[147]IQK[272]	3	653.0264	2.76	0.273
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EFNLQNM[147]GLPDFHIPENLFLK[272]	3	940.826	4.717	0.354
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VQGVEFSHR	3	401.5515	3.168	0.193
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ESDEETQIK	2	679.8436	3.687	0.322
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	HVAEAICK	2	598.8176	3.014	0.372
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LDVTTSIGR	2	551.3168	2.453	0.295
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QGFFPDSVNK	2	709.875	3.353	0.37
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLHMYANR	2	566.29	3.121	0.307
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TGISPLALIK	2	646.9187	3.34	0.295
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VNDESTEGK	2	629.8174	3.137	0.4
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	YHWEHTGLTLR	2	776.9046	3.123	0.416
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AVSMPSFSILGSDVR	2	855.4543	2.991	0.371
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EFQVPTFTIPK[272]	2	797.9578	3.291	0.325
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FVTQAEGAK[272]	2	619.8528	2.533	0.238
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NSEEFAAAMSR	2	678.8204	3.733	0.309
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SPSQADINK[272]	2	624.3454	3.905	0.214

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TGISPLALIK[272]	2	650.9258	3.718	0.363
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EKLTALTK	3	441.9502	3.479	0.218
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KSISAALEHK	3	501.9717	3.268	0.182
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VKYTLNK	3	429.2712	2.513	0.303
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VRESDEETQIK	3	538.6213	3.359	0.269
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FQFPGKPGIYTR	2	845.9751	3.449	0.31
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IADFELPTIIVPEQTIEIPSIK	2	1373.7857	2.845	0.434
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NSEEFAAAM[147]SR	2	684.8143	3.185	0.395
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVGFHLPSR	2	570.3197	2.942	0.305
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TLQGIPQM[147]IGEVIR	2	855.9822	2.488	0.308
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]HVAEAICK[272]	2	602.8247	2.907	0.329
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IEGNLIFDPNNYLPK[272]	2	1018.0588	2.363	0.27
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IGVELTGR	2	494.799	2.981	0.281
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NKYGM[147]VAQVTQTLK	3	673.047	4.326	0.415
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLHMYANR	3	377.8624	2.816	0.193
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ANLFNK[272]LVTELR	3	569.345	2.717	0.309
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]GISTSAASPAVGTVGM[147]DM[147]DEDDDFSK	3	975.107	2.933	0.307
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YHWEHTGLTLR	3	519.6079	2.801	0.223
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EFQVPTFTIPK	2	793.9507	2.655	0.248
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IGVELTGR	2	492.7955	2.292	0.134
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLWDFLK	2	594.8425	2.856	0.236
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ESDEETQIK[272]	2	683.8507	2.96	0.312
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FQFPGK[272]PGIYTR	2	849.9822	3.234	0.292
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LTISEQNIQR	2	673.3815	3.436	0.212
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SISAALEHK[272]	2	622.3661	3.364	0.328
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SPAFTDLHLR	2	650.8602	2.884	0.277
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VQGVEFSHR	2	601.8236	2.672	0.301
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	YHWEHTGLTLR	4	388.956	2.448	0.224
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IEIPLPFGGK	2	675.9109	2.714	0.274
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LGNNPVSK[272]	2	558.8344	2.735	0.313
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LPYTIITTPPLK[272]	2	823.0126	2.883	0.368
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]MGLAFESTK[272]	2	636.3491	1.653	0.178
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YHWEHTGLTLR	2	778.9082	3.378	0.313
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DDK[272]HEQDMVNGIM[147]LSVEK[272]	3	846.0965	3.136	0.28
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VK[272]YTLNK[272]	3	433.2783	3.225	0.194
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FVTQAEGAK	2	615.8457	2.672	0.179
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IEDGTLASK	2	607.343	2.769	0.172
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LPYTIITTPPLK	2	819.0055	3.404	0.307
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	M[147]GLAFESTK	2	640.3394	2.501	0.282
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FPEVDVLTK[272]	2	668.3918	3.055	0.279
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]GVISIPR	2	443.2856	1.667	0.088
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IPSVQINFK[272]	2	667.4078	2.767	0.277
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLHMYANR	2	568.2936	2.746	0.232
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VNDESTEGKTSYR	3	589.2954	3.315	0.378
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLDEHYHIR	3	438.5621	2.574	0.199
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VRESDEETQIK[272]	3	541.2927	3.43	0.257
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QAEAVLK	2	519.819	2.581	0.242
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LDVTTSIGR	2	553.3203	2.298	0.206

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLDEHYHIR	2	657.3396	3.075	0.241
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SQAIATK[272]	2	503.8104	2.567	0.324
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LELELRPTGEIEQYSVSATYELQR	3	990.1843	3.154	0.208
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NIILPVYDK	2	677.9083	3.256	0.209
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]GNVATEISTER	2	660.8474	3.534	0.321
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TEVIPPLIENR	2	712.9151	2.503	0.183
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LTLDIQNK[272]K[272]	3	502.319	3.12	0.148
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	KSISAALEHK	2	752.454	3.108	0.164
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	INPLALK	2	524.8475	2.41	0.225
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LTLDIQNK	2	612.8692	2.985	0.144
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QAEAVLK[272]	2	523.8261	2.561	0.223
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SQAIATK	2	499.8033	2.591	0.175
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QTEATM[147]TFK[272]	2	680.8547	2.912	0.188
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EAQEVFK	2	565.8139	2.426	0.223
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FFGEGTK[272]	2	537.2972	2.293	0.228
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DLKVEDIPLAR	3	516.9751	3.488	0.235
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EAQEVFK[272]	2	569.821	2.525	0.219
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLHM[147]YANR	3	384.5298	2.157	0.204
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LTISEQNIQR	2	671.3779	3.166	0.137
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LTLDIQNK[272]	2	616.8763	2.844	0.178
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLHM[147]YANR	2	576.2911	1.883	0.131
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	GVISIPR	2	441.282	2.482	0.204
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DK[272]IGVELTGR	3	459.2764	2.78	0.224
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EIFNMAR	2	512.772	2.305	0.191
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EIQIYK[272]	2	541.3285	2.169	0.106
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QTEATMTFK[272]	2	672.8572	2.659	0.181
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLWDFLK[272]	2	598.8496	2.609	0.19
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EVYGFNPEGK	2	710.367	2.774	0.194
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VRESDEETQIK	2	807.4283	3.489	0.347
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NYQLYK[272]	2	558.8183	2.13	0.218
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LAIPEGK[272]	2	508.3232	2.195	0.235
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SEILAHWSPAK	3	506.9524	2.449	0.172
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ESQLPTVM[147]DFR	2	741.8726	2.63	0.182
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ITLPDFR	2	503.2961	1.667	0.038
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EVYGFNPEGK[272]	2	714.3741	2.975	0.14
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]HINIDQFVR	2	643.3603	2.99	0.073
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NIILPVYDK[272]	2	681.9154	2.35	0.181
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EIQIYK	2	537.3214	1.998	0.122
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FFGEGTK	2	533.2901	2.14	0.191
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LIDVISM[147]YR	2	633.3498	1.726	0.141
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	QTEATM[147]TFK	2	676.8476	2.194	0.205
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLDEHYHIR	4	328.1717	1.6	0.111
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LAIPEGK	2	504.3161	2.626	0.1
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IEDGTLASK[272]	2	611.3501	2.485	0.176
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VK[272]YTLNK[272]NSLK[272]	3	628.7304	2.569	0.26
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ALVEQGFTVPEIK[272]	2	860.0002	2.643	0.2
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SVGFHLPSR	2	572.3232	2.882	0.24
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SVGFHLPSR	3	381.8846	2.716	0.171

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VKYTLNK	2	643.4032	3.081	0.258
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NIFNFK[272]	2	535.8155	2.25	0.13
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	YTLNKNSLK	2	750.9485	3.11	0.3
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ITLPDFR	2	501.2926	1.912	0.165
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DLGQCDR	2	498.7216	2.179	0.167
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VRESDEETQIK[272]	2	811.4354	2.682	0.341
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLDEHYHIR	4	329.1734	1.961	0.073
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ENFAGEATLQR	2	690.3554	2.62	0.124
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	YTLNKNSLK	3	500.9681	2.071	0.232
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IEIPLPFGGK[272]	2	679.918	2.433	0.151
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LIDVISM[147]YR	2	635.3533	2.701	0.108
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EK[272]LTALTK[272]	3	445.9573	2.738	0.138
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QIDDIDVR	2	559.3021	2.638	0.056
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]FLDSNIK[272]	2	562.8314	2.194	0.11
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TIHDLHLFIENIDFNK	3	750.4074	3.313	0.059
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FPEVDVLTK	2	664.3847	1.392	0.111
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SQAIATK[272]K[272]	2	639.909	1.943	0.122
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VK[272]YTLNK[272]	2	649.4139	2.775	0.229
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LKETIQK	3	427.2748	2.6	0.142
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TLQELK[272]K[272]	2	646.4192	2.349	0.01
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VKYTLNKNSLK	3	623.3876	2.154	0.235
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AIVDTLK	2	520.3292	2.299	0.231
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SK[272]EVPEAR	3	401.9027	2.821	0.112
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DLGQCDR	2	496.7181	1.967	0.133
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SQAIATKK	3	422.9346	2.069	0.124
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ITLPDFR	1	1001.5779	1.712	0.061
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LLLM[147]GAR	2	465.2837	2.395	0.012
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LIDVISMYR	2	627.3559	2.424	0.039
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FSHVEK	2	513.7902	1.938	0.072
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NYQLYK	2	554.8112	1.753	0.125
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IEGNLIFDPNNYLPK	2	1014.0517	2.112	0.102
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TLADLTLLDSPIK[272]	2	844.5079	2.719	0.196
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LEDTPK[272]	2	495.7892	2.092	0.064
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TEVIPPLIENR	2	710.9116	2.12	0.122
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LAPGELTIIL	1	1179.7348	1.531	0.161
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SLHM[147]YANR	3	383.1941	1.96	0.113
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SK[272]EVPEAR	2	602.3504	2.183	0.282
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LVTELR	2	435.774	2.059	0.15
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SPSQADINK	2	620.3383	3.167	0.238
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]MLETVR	2	446.7558	1.961	0.008
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IHSGSFQSQVELSNDQEK[272]	3	774.3919	2.102	0.124
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EEEMLENVSLVCPK[272]	2	977.4823	4.208	0.351
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IPSVQINFK	2	663.4007	2.077	0.056
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DK[272]IGVELTGR	2	688.411	2.773	0.156
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VNDESTEGKTSYR	2	883.4394	2.87	0.405
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IPSVQINFK[272]DLK[272]	3	612.0436	2.639	0.124
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VNDESTEGK[272]TSYR	3	591.9668	2.299	0.155
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	CVQSTKPSLM[147]IQK	3	649.0193	2.691	0.076

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LLLM[147]GAR	2	467.2872	1.734	0.046
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TSQCTLK	2	555.792	1.863	0.032
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LAAYLM[147]LM[147]R	2	627.3409	1.722	0.048
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]IEIPLPFGGK[272]	3	453.6144	1.632	0.149
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DDKHEQDM[147]VNGIM[147]LSVEK	3	847.421	2.89	0.106
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ALYWVNGQVPDGVSK[272]	3	641.0189	3.123	0.043
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YTLNK[272]NSLK[272]	2	756.9592	2.231	0.194
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]INFNEK[272]	2	526.8026	1.632	0.07
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NIFNFK	2	531.8084	1.615	0.029
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	VLADK	2	413.2633	1.802	0.059
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EEEMLENVSLVCPK	2	973.4752	3.33	0.434
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]TNFFHESGLEAHVALK[272]	4	522.7834	3.056	0.228
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	HIPSFTIDFVEM[147]K	3	620.6623	3.107	0.197
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ATLYALSH	2	510.2858	2.74	0.27
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DLK[272]VEDIPLAR	2	778.9662	2.601	0.245
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	ATLYALSH	2	508.2822	3.065	0.218
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	FFGEGTKK	3	445.2591	1.762	0.109
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EVDVLTK[272]	2	546.3312	2.839	0.233
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SKEVPEAR	2	598.3433	1.844	0.155
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	NSLKIEIPLPFGGK	3	645.0593	2.821	0.125
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SKEVPEAR	3	399.2313	2.434	0.047
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SLHMYANR	3	379.1981	1.516	0.095
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]EEEM[147]LENVSLVCPK[272]	2	985.4798	3.986	0.414
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AFTDLHLR	2	556.8142	3.113	0.121
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AYLM[147]LM[147]R	2	537.2838	2.319	0.279
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AFTDLHLR	2	558.8178	3.265	0.083
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	DKIGVELTGR	3	456.605	2.268	0.076
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EVDVLTK	2	542.3241	2.389	0.233
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DNVFDGLVR	2	589.818	1.418	0.054
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EEEM[147]LENVSLVCPK	2	981.4727	2.402	0.241
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DLK[272]VEDIPLAR	3	519.6465	2.16	0.08
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LK[272]ETIQK[272]	3	431.2819	2.051	0.044
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ETLEDTRDR	3	426.8851	2.033	0.13
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVSLPSLDPASAK	3	517.9629	2.004	0.003
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	IPSVQINFKDLK	3	608.0365	1.902	0.096
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QVFLYPEK[272]DEPTYILNIK[272]	4	661.3756	2.866	0.043
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AVSM[147]PSFSILGSDVR	3	575.9702	1.823	0.077
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]SQADINK[272]	2	532.303	1.904	0.19
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	TKNSEEFAAAMSR	3	574.6262	2.263	0.018
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LK[272]QHIEAIDVR	3	537.3258	1.894	0.041
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AYLM[147]LM[147]R	2	535.2803	2.007	0.139
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]AFTDLHLR	3	372.8809	2.445	0.08
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]NFVASHIAN	2	558.7996	2.175	0.234
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]QQANDYLNSFNWER	2	964.954	4.36	0.322
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]PGIYTR	2	425.7488	1.958	0.019
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	EM[147]K	1	703.3808	1.121	0.072
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	LKETIQK[272]LSNVLQQVK	3	811.8395	1.313	0.13
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	AFTDLHLR	3	371.5452	2.238	0.035

44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LLSGGNTLH	2	528.3019	2.099	0.025
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]LAPGELTIIL	1	1183.7419	1.374	0.103
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]HIPSFTIDFVEM[147]K[272]	3	623.3337	2.665	0.027
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]ETLEDTRDR	2	639.824	2.049	0.025
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]VLVDHFG	2	465.7619	1.4	0.109
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]YFEK[272]	2	437.7493	1.194	0.17
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SK[272]K[272]	2	395.7731	0.913	0.136
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVAK	2	342.7238	0.841	0.121
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	SVK[272]	1	617.4104	0.88	0.378
44	IPI00022229	1	34.7	0.84	0.11	378	##	APOB Apolipoprotein B-100 precursor	n[145]DFLK[272]	1	810.4964	1.211	0.02
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	ADLFYDVEALDLESPK	2	1053.0437	3.977	0.497
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	GGEGTGYFVDFSVR	2	815.8967	3.978	0.488
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	RPSEIVIGQCK	2	778.426	4.109	0.479
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	YKEENDDFASFR	2	900.9312	4.683	0.478
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]VSPTDCSAVEPEAEK[272]	2	948.4521	4.871	0.558
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]YK[272]EENDDFASFR	2	904.9383	4.256	0.41
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	DHHHPHKPHEHGPPPPPDER	3	877.4339	2.98	0.362
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	GEVLPLPEANFPSFPLPHHK	3	836.125	5.369	0.572
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	HPLKPDNQPFPQSVSESCPGK	3	920.1317	6.47	0.553
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	KGEVLPLPEANFPSFPLPH	3	790.4387	3.98	0.384
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	SSTTKPPFKPHGSR	3	649.7023	2.494	0.09
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]DHHHPHK[272]PHEHGPPPPPDER	3	880.1053	4.218	0.51
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]GEVLPLPEANFPSFPLPHHK[272]	3	838.7964	4.035	0.499
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]HPLK[272]PDNQPFPQSVSESCPGK[272]	3	924.1388	5.605	0.497
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]YK[272]EENDDFASFR	3	603.628	4.033	0.478
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	DHHHPHKPHEHGPPPPPDER	4	658.3273	3.96	0.429
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	GEVLPLPEANFPSFPLPHHK	4	627.3456	3.088	0.395
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	KGEVLPLPEANFPSFPLPHHK	4	694.3931	3.275	0.45
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]GEVLPLPEANFPSFPLPHHK[272]	4	629.3492	2.738	0.134
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]RPSEIVIGQCK[272]	2	782.4331	3.662	0.416
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	RDGYLFQLLR	3	474.2735	4.517	0.204
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]RPSEIVIGQCK[272]	3	521.9578	3.738	0.225
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]DHHHPHK[272]PHEHGPPPPPDER	4	660.3308	3.369	0.31
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]HPLK[272]PDNQPFPQSVSESCPGK[272]	4	693.3559	2.11	0.086
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]SSTTK[272]PPFK[272]PHGSR	4	490.5339	2.544	0.253
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	DGYLFQLLR	2	632.8561	3.328	0.356
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]IADAHLDR	2	527.7917	2.975	0.396
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]ADLFYDVEALDLESPK[272]	2	1057.0508	3.627	0.316
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	YKEENDDFASFR	3	600.9566	3.187	0.271
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]RDGYLFQLLR	3	475.6092	3.385	0.154
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]DGYLFQLLR	2	634.8596	2.841	0.304
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	QIGSVYR	2	481.7746	2.695	0.249
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	DHHHPHKPHEHGPPPPPDER	5	526.8633	2.002	0.05
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	SSTTKPPFKPHGSR	4	487.5286	2.443	0.12
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	IADAHLDR	2	525.7882	2.795	0.202
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]SGFPQVSM[147]FFTHTFPK[272]	3	721.371	2.823	0.22
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]ALDLINK[272]	2	537.8417	2.807	0.083
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]SSTTK[272]PPFK[272]PHGSR	3	653.7094	2.884	0.179

45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	ALDLINK	2	533.8346	2.319	0.079
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]IADAHLDR	3	352.1969	1.617	0.178
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	SGFPQVSM[147]FFTH	2	770.8663	2.798	0.484
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	IADAHLDR	3	350.8612	1.995	0.093
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]RDGYLFQLLR	2	712.9102	2.422	0.159
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	DHSHGPPLPQGPPPLLPM[147]	2	1022.0277	3.459	0.415
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	GEVLPLPEANFPSFPLPH	2	1051.0595	3.888	0.381
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]DHHHPHK[272]PHEHGPPPPPDER	5	528.4661	2.339	0.092
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]DHSHGPPLPQGPPPLLPM[147]SC	3	761.6995	2.298	0.309
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]NVFGFCR	2	516.7474	2.44	0.281
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	RPSEIVIGQCK	3	519.2864	2.414	0.11
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	NVFGFCR	2	514.7439	2.128	0.251
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]HPNVFGF	2	481.2543	2.261	0.17
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	n[145]SGFPQVSM[147]FFTH	2	772.8699	2.345	0.264
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	HPNVFGF	2	479.2507	2.07	0.14
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	HPLKPDNQPFPQSVSESCPGK	4	690.3506	2.174	0.047
45	IPI00022371	1	42.1	0.89	0.17	105	55	HRG Histidine-rich glycoprotein precursor	SEIVIGQCK	2	651.8491	3.142	0.302
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	IVLGQEQDSYGGK	2	837.4465	4.304	0.466
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	QGYFVEAQPK	2	723.8907	3.229	0.32
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]AYSLFSYNTQGR	2	775.8896	3.894	0.453
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]QGYFVEAQPK[272]	2	727.8978	3.285	0.375
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]AYSLFSYNTQGRDNELLVYK[272]ER	4	739.3867	5.231	0.462
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]VGEYSLYIGR	2	650.8545	2.634	0.313
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	AYSLFSYNTQGR	2	773.8861	3.328	0.419
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]AYSLFSYNTQGRDNELLVYK[272]ER	3	985.5132	4.461	0.352
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]IVLGQEQDSYGGK[272]	2	841.4536	3.168	0.168
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]VFVFPR	2	454.7774	1.961	0.161
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	VFVFPR	2	452.7738	1.971	0.015
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	HTDLSGKVFVFPR	3	595.0053	2.82	0.366
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]HTDLSGK[272]VFVFPR	4	448.5094	2.601	0.24
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	n[145]HTDLSGK[272]VFVFPR	3	597.6767	2.85	0.169
46	IPI00022391	1	32.7	0.79	0.09	20	14	APCS Serum amyloid P-component precursor	KVIEK	2	518.8476	2.301	0.075
47	IPI00022392	1	10	0.72	0.05	13	3	C1QA Complement C1q subcomponent subunit A precursor	SLGFCDTTNK	2	706.3391	3.532	0.284
47	IPI00022392	1	10	0.72	0.05	13	3	C1QA Complement C1q subcomponent subunit A precursor	n[145]SLGFCDTTNK[272]	2	710.3462	3.549	0.411
47	IPI00022392	1	10	0.72	0.05	13	3	C1QA Complement C1q subcomponent subunit A precursor	n[145]PAFSAIR	2	453.2699	1.82	0.144
47	IPI00022392	1	10	0.72	0.05	13	3	C1QA Complement C1q subcomponent subunit A precursor	n[145]EDLCR	2	413.1894	1.977	0.143
48	IPI00022394	1	21.2	0.73	0.02	6	5	C1QC Complement C1q subcomponent subunit C precursor	FNAVLTNPQGDYDTSTGK	2	1104.5502	5.983	0.537
48	IPI00022394	1	21.2	0.73	0.02	6	5	C1QC Complement C1q subcomponent subunit C precursor	n[145]FNAVLTNPQGDYDTSTGK[272]	2	1108.5573	5.753	0.569
48	IPI00022394	1	21.2	0.73	0.02	6	5	C1QC Complement C1q subcomponent subunit C precursor	n[145]QTHQPPAPNSLIR	3	534.9659	3.174	0.295
48	IPI00022394	1	21.2	0.73	0.02	6	5	C1QC Complement C1q subcomponent subunit C precursor	n[145]TNQVNSGGVLLR	2	701.4002	4.007	0.27
48	IPI00022394	1	21.2	0.73	0.02	6	5	C1QC Complement C1q subcomponent subunit C precursor	n[145]FQSVFTVTR	2	614.844	2.88	0.231
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	TEHYEEQIEAFK	2	902.4493	4.478	0.514
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]TEHYEEQIEAFK[272]	2	906.4564	4.547	0.39
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	AEQCCEETASSISLHGK	3	722.3213	3.739	0.478
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]AEQCCEETASSISLHGK[272]	3	724.9927	3.491	0.351
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	KYAFELK	3	440.2676	2.921	0.228
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]CTDAVGDRR	3	394.854	2.266	0.243
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	LSPIYNLVPVK	2	761.9715	3.578	0.387

49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]CLCACPFK[272]	2	655.7786	2.871	0.415
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]LSPIYNLVPVK[272]	2	765.9786	3.126	0.414
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]DRDGNTLTYYR	3	506.587	3.672	0.281
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]GEIHLGR	2	463.2704	3.044	0.3
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]VVEESELAR	2	588.3231	1.851	0.158
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]DGNTLTYYR	2	623.8129	2.737	0.325
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	ALPTTYEK	2	601.8426	2.32	0.058
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	DVVLTTTFVDDIK	2	873.4879	3.999	0.292
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]RPWNVASLIYETK	3	621.0199	3.95	0.257
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	KGVELK	3	365.2413	2.606	0.144
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]ALPTTYEK[272]	2	605.8497	2.334	0.009
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]CTDAVGDR	2	513.7269	2.855	0.219
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]FTPTETNK[272]	2	613.337	2.303	0.247
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	FTPTETNK	2	609.3299	2.325	0.135
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]CTDAVGDRR	2	591.7774	2.523	0.116
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]TEHYEEQIEAFK[272]	3	604.64	3.499	0.062
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	CLCACPFK	2	651.7715	2.017	0.193
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	KYAFELK	2	659.8978	2.21	0.072
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]GEIHLGR	3	309.1827	2.781	0.032
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]DVVLTTTFVDDIK[272]	2	877.495	3.212	0.368
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	TEHYEEQIEAFK	3	601.9686	2.268	0.114
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	n[145]FEGIACEISK[272]	2	715.8667	2.006	0.078
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	YAFELK	2	525.8028	1.521	0.06
49	IPI00022395	1	27.2	0.79	0.23	40	30	C9 Complement component C9 precursor	SK[272]K[272]	2	395.7731	0.913	0.136
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]ALGHLDLSGNR	2	648.8607	4.039	0.456
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]VAAGAFQGLR	2	567.331	3.382	0.366
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]ALGHLDLSGNR	3	432.9095	2.867	0.259
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]LHLEGNK[272]	3	366.8885	3.098	0.164
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	LHLEGNK	3	364.2171	2.502	0.201
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]DCQVFR	2	479.2238	1.892	0.092
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	n[145]CAGPEAVK[272]	2	554.7903	2.226	0.186
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	ENQLEVLEVSWLHGLK[272]	3	726.7395	3.587	0.141
50	IPI00022417	1	19	0.72	0.05	11	8	LRG1 Leucine-rich alpha-2-glycoprotein precursor	GIQPHVSR	2	517.2987	1.545	0.092
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]LIVHNGYCDGR	3	479.568	3.379	0.296
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]LIVHNGYCDGR	2	718.8484	2.47	0.207
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]QRQEELCLAR	3	479.2471	1.998	0.09
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]DPNGLPPEAQK[272]	2	727.3981	3.441	0.35
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]QEELCLAR	2	576.2871	2.754	0.212
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]GNDDHWIVDTDYDTYAVQYSCR	3	942.7346	2.649	0.195
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]YWGVASFLQK	2	741.9149	3.255	0.306
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]LLNLDGTCADSYSFVFSR	3	733.3508	2.407	0.146
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]QRQEELCLAR	2	718.367	3.156	0.237
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]VK[272]ENFDK[272]AR	2	769.9544	2.765	0.199
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	VKENFDKAR	2	763.9438	2.684	0.259
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]VK[272]ENFDK[272]AR	3	513.6387	2.345	0.171
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	VKENFDKAR	3	509.6316	2.413	0.153
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	VKENFDK	2	650.3747	2.544	0.136
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]YWGVASFLQK	3	494.9457	3.199	0.142

51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]VK[272]ENFDK[272]	3	437.926	2.189	0.139
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]VSSFR	2	370.2146	1.747	0.054
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]VK[272]ENFDK[272]	2	656.3853	2.279	0.086
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	VKENFDKAR	4	382.4755	1.896	0.019
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	n[145]AVQYSCR	2	508.7423	1.981	0.177
51	IPI00022420	1	48.2	0.82	0.14	27	21	RBP4 Plasma retinol-binding protein precursor	VKENFDK	3	433.9189	1.83	0.019
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	AFIQLWAFDAVK	2	844.9798	3.989	0.529
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	GVCEETSGAYEK	2	799.8631	3.869	0.434
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]GECVPGEQEPEPILIPR	2	1027.0144	3.962	0.523
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]GVCEETSGAYEK[272]	2	803.8702	3.535	0.524
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]TVAACNLPIVR	2	673.8739	3.769	0.362
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]EYCGVPGDGDEELLR	2	921.412	3.083	0.455
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]HHGPTITAK[272]	3	417.2467	3.147	0.188
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]CVLFPYGGCQGNGNK[272]	2	968.9427	2.692	0.443
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]ECLQTCR	2	544.7275	2.137	0.259
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	HHGPTITAK	3	414.5753	3.062	0.06
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]HHGPTITAK[272]	2	625.3664	1.879	0.195
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]SAGPCM[147]GM[147]TSR	2	660.2703	2.582	0.49
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]FYSEK[272]	2	481.2653	1.408	0.025
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	CVLFPYGGCQGNGNK	2	964.9356	2.373	0.312
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	n[145]GNNFVTEK[272]	2	598.8293	2.61	0.116
52	IPI00022426	1	34.7	0.73	0.04	23	16	AMBP AMBP protein precursor	GNNFVTEK	2	594.8222	2.222	0.061
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	TYMLAFDVNDEK	2	863.4295	3.986	0.413
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	TYM[147]LAFDVNDEK	2	871.4269	4.347	0.519
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]EQLGEFYEALDCLR	2	938.4405	4.97	0.591
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TYMLAFDVNDEK[272]	2	867.4366	4.147	0.435
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]VGGQEHFAHLLILR	2	867.5002	4.714	0.417
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHLLILR	2	949.0319	6.59	0.566
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAH	3	430.2112	3.285	0.363
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHLLILR	3	633.0237	2.878	0.172
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	NWGLSVYADKPETTK	3	710.3845	3.081	0.345
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]GLSVYADK[272]PETTK[272]	3	614.3509	4.239	0.408
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHLLILR	4	475.0196	4.339	0.257
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	SDVVYTDWK	2	696.8616	2.874	0.296
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]DK[272]CEPLEK[272]	3	480.5915	3.228	0.147
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]EQLGEFYEALDCLR	3	625.9628	2.196	0.072
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]NWGLSVYADKPETTK	3	711.7202	3.502	0.185
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]NWGLSVYADK[272]PETTK[272]	3	714.3916	3.078	0.321
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]QEHFAHLLILR	3	507.6321	3.456	0.426
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	DKCEPLEK	3	476.5844	2.348	0.263
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]SDVVYTDWK[272]	2	700.8687	2.514	0.13
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TYM[147]LAFDVNDEK[272]	2	875.434	2.952	0.308
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]VGGQEHFAHLLILR	3	578.6692	4.04	0.37
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TEDTIFLR	1	1138.6225	2.6	0.235
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TEDTIFLR	2	569.8149	2.252	0.02
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]SDVVYTDWK[272]	1	1400.7301	2.869	0.288
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	GLSVYADKPETTK	3	610.3438	3.206	0.344
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]QEHFAHLLILR	4	380.9759	3.196	0.306

53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]CEPLEK[272]	2	526.7716	1.507	0.229
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHL	3	467.9059	3.525	0.255
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]CEPLEK[272]	1	1052.5359	1.61	0.168
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	DKCEPLEK	2	714.3729	2.542	0.263
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]GQEHFAHLLILR	3	526.6393	3.518	0.244
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	SDVVYTDWK[272]	2	698.8651	3.278	0.34
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHLL	3	505.6006	2.284	0.339
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]DK[272]CEPLEK[272]	2	720.3836	2.88	0.121
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]GQEHFAHLLILR	2	789.4553	4.07	0.396
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]VGGQEHFAHLLILR	4	434.2538	2.997	0.197
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TTYLNVQR	2	569.8205	2.844	0.349
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]FYIASAFR	2	559.8094	2.117	0.267
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAHLL	2	757.8973	3.011	0.398
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	CEPLEK	2	522.7645	1.847	0.016
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YVGGQEHFAH	2	644.8132	2.62	0.197
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]GQEHFAHLLILR	4	395.2313	2.892	0.181
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	SDVVYTDWK[272]	3	466.2458	2.664	0.146
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YIASAFR	2	486.2752	1.866	0.286
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]IASAFR	2	404.7435	1.822	0.055
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]YLNVQR	2	468.7728	1.886	0.018
53	IPI00022429	1	48.3	0.6	0.09	67	41	ORM1 Alpha-1-acid glycoprotein 1 precursor	n[145]TYLNVQR	2	519.2967	1.922	0.012
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	EHAVEGDCDFQLLK	2	965.4635	4.277	0.485
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	M[147]GVVSLGSPSGEVSHPR	2	926.4727	4.851	0.415
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	TVSGTDCVAK	2	653.8283	3.974	0.464
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]EHAVEGDCDFQLLK[272]	2	969.4706	5.091	0.549
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFM[147]GVVSLGSPSGEVSHPR	2	1121.0638	5.508	0.577
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTLNQIDEVK[272]	2	742.9192	2.036	0.198
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]TVSGTDCVAK[272]	2	657.8354	3.912	0.414
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]TVVQPSVGAAAGPVVPPCPGR	2	1075.0726	6.269	0.477
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	AQLVPLPPSTYVEFTVSGTDCVAK	3	950.159	1.978	0.127
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	LDGKFSVVYAK	3	549.6592	3.841	0.436
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	M[147]GVVSLGSPSGEVSHPR	3	617.9842	5.123	0.466
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFMGVVSLGSPSGEVSHPR	3	742.38	5.258	0.465
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFM[147]GVVSLGSPSGEVSHPR	3	747.7116	2.633	0.355
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]LDGK[272]FSVVYAK[272]	3	553.6663	3.696	0.499
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]TVVQPSVGAAAGPVVPPCPGR	3	717.0508	3.5	0.291
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]VVSLGSPSGEVSHPR	3	551.301	5.01	0.394
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]AQLVPLPPSTYVEFTVSGTDCVAK[272]	3	952.8304	2.929	0.105
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]CNLLAEK[272]QYGFCK[272]	3	681.0083	3.997	0.341
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTLNQIDEVK[272]	3	495.6152	3.645	0.292
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	HTLNQIDEVK	2	738.9121	1.795	0.188
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]CNLLAEK[272]	2	562.806	2.924	0.187
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFMGVVSLGSPSGEVSHPR	4	557.0368	3.315	0.185
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFM[147]GVVSLGSPSGEVSHPR	4	561.0355	2.998	0.286
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	CNLLAEK	2	558.7989	3.119	0.262
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	FSVVYAK	2	547.3239	2.961	0.289
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]FSVVYAK[272]	2	551.331	2.405	0.182
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]VLLAAPPGHQLHR	3	518.3113	3.455	0.348

54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	CNLLAEK	1	1116.5905	2.196	0.161
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	HTLNQIDEVK	3	492.9438	3.067	0.076
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]CNLLAEK[272]	1	1124.6047	2.005	0.178
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]GSPSGEVSHPR	3	418.55	3.038	0.372
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	ATLSEK	2	464.7768	2.031	0.033
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]ATLSEK[272]	2	468.7839	1.996	0.076
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HGPGLIYR	3	352.8739	2.986	0.206
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]AHYDLR	3	306.8353	2.483	0.037
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	AHYDLR	3	305.4996	2.584	0.004
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HTFM[147]GVVSLGSPSG	2	768.3859	3.744	0.345
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	QYGFCK	2	536.2594	1.219	0.14
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]QYGFCK[272]	2	540.2665	1.241	0.104
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]SLGSPSGEVSHPR	3	485.2554	2.961	0.234
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]HGPGLIYR	2	528.8072	2.294	0.174
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]QIDEVK[272]	2	510.3024	2.611	0.112
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	VLDPTPVAR	2	554.3297	1.829	0.23
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	SPSGEVSHPR	3	398.2072	2.538	0.084
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]GPGLIYR	2	460.2777	2.404	0.051
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	QIDEVK	2	506.2953	2.15	0.014
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]VLDPTPVAR	2	556.3332	1.973	0.034
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	HGPGLIYR	1	1052.6	1.82	0.185
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]AALAAF	1	707.4209	1.548	0.22
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	AALAAF	1	703.4138	1.443	0.26
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]CDFQLLK[272]	2	600.8216	1.9	0.239
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	HGPGLIYR	3	351.5382	1.976	0.065
54	IPI00022431	1	45.5	0.76	0.1	124	45	AHSG Alpha-2-HS-glycoprotein precursor	n[145]SPPDSHVLLAAPPGHQLHR	4	544.0491	2.259	0.007
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	AADDTWEPFASGK	2	837.9098	4.1	0.563
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GPTGTGESKCPLM[147]VK	2	994.0124	3.582	0.537
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	KAADDTWEPFASGK	2	972.0048	4.465	0.533
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]AADDTWEPFASGK[272]	2	841.9169	3.735	0.392
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]GPTGTGESK[272]CPLM[147]VK[272]	2	1000.023	4.571	0.45
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]GSPAINVAVHVFR	2	755.9342	4.07	0.497
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	ALGISPFHEHAEVVFTANDSGPRR	3	916.4719	5.031	0.503
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	TSESGELHGLTTEEEFVEGIYKVEIDTK	3	1187.6051	6.317	0.613
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]GSPAINVAVHVFR	3	504.2919	3.978	0.206
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	TSESGELHGLTTEEEFVEGIYKVEIDTK	4	890.9557	3.164	0.419
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]TSESGELHGLTTEEEFVEGIYK[272]VEIDTK[272]	4	893.961	7.253	0.607
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	YTIAALLSPYSYSTTAVVTNPKE	3	923.8285	3.401	0.389
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	ALGISPFHEHAEVVFTANDSGPR	3	864.4382	3.595	0.283
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]TSESGELHGLTTEEEFVEGIYK[272]	3	915.1233	3.498	0.269
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GSPAINVAVHVFR	3	502.9562	2.045	0.178
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GPTGTGESKCPLM[147]VK	3	663.0107	2.091	0.22
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	VIDAVR	2	406.7531	1.998	0.073
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GSPAINVAVH	2	552.8117	3.385	0.393
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]VEIDTK[272]	2	496.797	2.05	0.094
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	VEIDTK	2	492.7899	1.911	0.107
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	KAADDTWEPFASGK	3	648.3389	1.931	0.091
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]ALGISPFH	2	493.283	2.17	0.243

55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]GPTGTGESK[272]	2	561.3057	2.193	0.216
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GPTGTGESK	2	557.2986	1.992	0.123
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	GPTGTGESKCPLM[147]VK	4	497.5098	2.624	0.114
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]GPTGTGESK[272]CPLM[147]VK[272]	3	667.0178	2.054	0.218
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]KAADDTWEPFASGK	2	974.0083	3.349	0.309
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	TSESGELHGLTTEEE	2	879.9051	2.206	0.128
55	IPI00022432	1	79.6	0.86	0.21	52	25	TTR Transthyretin precursor	n[145]YTIAALLSPYSYSTTAVVTNPK[272]E	3	926.4999	2.011	0.058
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AVMDDFAAFVEK	1	1622.8248	3.042	0.474
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCTESLVNR	1	1256.5255	2.625	0.448
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVNEVTEFAK	1	1429.805	3.092	0.39
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NECFLQHK	1	1344.6552	2.519	0.376
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QTALVELVK	1	1280.7937	3.011	0.382
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AVMDDFAAFVEK[272]	1	1630.839	3.256	0.514
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCTESLVNR	1	1260.5326	3.271	0.418
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DLGEENFK[272]	1	1239.646	2.184	0.264
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVNEVTEFAK[272]	1	1437.8192	3.013	0.335
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AAFTECCQAADK	2	815.3483	2.069	0.216
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AEFAEVSKLVTDLTK	2	1036.0936	5.03	0.415
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AQYLQQCPFEDHVK	2	1016.4926	3.237	0.478
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AVMDDFAAFVEK	2	811.916	4.232	0.482
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AVM[147]DDFAAFVEK	2	819.9135	3.653	0.402
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCAAADPHECYAK	2	900.3469	4.382	0.533
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCTESLVNR	2	628.7664	2.692	0.374
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CTAFHDNEETFLK	2	940.9371	4.315	0.575
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVFLGMFLYEYAR	2	882.4449	4.873	0.517
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVFLGM[147]FLYEYAR	2	890.4424	3.239	0.403
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ECCEKPLLEK	2	852.4212	3.598	0.364
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHADICTLSEK	2	1265.0931	5.669	0.553
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ETYGEMADCCAK	2	846.8316	3.399	0.657
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ETYGEM[147]ADCCAK	2	854.829	1.631	0.208
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FKDLGEENFK	2	823.9487	4.224	0.364
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FTECCQAADK	2	744.3111	4.566	0.548
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FYAPELLFFAK	2	813.4582	3.499	0.446
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPELLFFAK	2	1012.0457	4.247	0.379
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KQTALVELVK	2	774.9955	1.955	0.065
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KVPQVSTPTLVEVSR	2	960.5675	3.012	0.399
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKECCEKPLLEK	2	1043.0583	4.177	0.369
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVNEVTEFAK	2	715.4061	4.868	0.392
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVMCTAFHDNEETFLK	2	1460.2137	5.861	0.564
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QEPERNECFLQHK	2	992.48	4.497	0.446
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QNCELFEQLGEYK	2	963.9581	5.547	0.508
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QTALVELVK	2	640.9005	3.245	0.333
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPDYSVVLLLR	2	804.4727	2.486	0.257
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RPCFSALEVDETYVPK	2	1090.5476	2.432	0.155
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEMPADLPSLAADFVESK	2	1622.254	3.891	0.671
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEM[147]PADLPSLAADFVESK	2	1630.2515	5.814	0.585
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TCVADESAENCDK	2	878.8541	4.785	0.6
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VADESAENCDK	2	753.8318	3.339	0.524

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VFDEFKPLVEEPQNLIK	2	1233.1938	4.425	0.416
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VMDDFAAFVEK	2	776.3975	3.866	0.535
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VPQVSTPTLVEVSR	2	826.4725	2.141	0.242
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YAPELLFFAK	2	739.924	4.198	0.416
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YFYAPELLFFAK	2	894.9898	4.368	0.422
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YICENQDSISSK	2	856.9028	2.601	0.254
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AAFTECCQAADK[272]	2	819.3554	3.908	0.54
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AQYLQQCPFEDHVK[272]	2	1020.4997	5.163	0.471
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AVMDDFAAFVEK[272]	2	815.9231	5.077	0.562
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AVM[147]DDFAAFVEK[272]	2	823.9206	2.519	0.241
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCAAADPHECYAK[272]	2	904.354	3.962	0.608
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CFSALEVDETYVPK[272]	2	967.9778	3.131	0.413
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CTAFHDNEETFLK[272]	2	944.9442	4.593	0.483
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DESAENCDK[272]	2	672.7862	2.878	0.52
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DVFLGM[147]FLYEYAR	2	892.4459	4.101	0.406
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EFNAETFTFHADICTLSEK[272]	2	1269.1002	5.117	0.512
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ETYGEMADCCAK[272]	2	850.8387	4.145	0.676
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ETYGEM[147]ADCCAK[272]	2	858.8361	2.203	0.38
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FK[272]DLGEENFK[272]	2	829.9594	3.295	0.302
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FTECCQAADK[272]	2	748.3182	3.483	0.5
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FYAPELLFFAK[272]	2	817.4653	3.968	0.405
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]HPYFYAPELLFFAK[272]	2	1016.0528	3.845	0.487
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LK[272]ECCEK[272]PLLEK[272]	2	1051.0725	3.48	0.423
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVNEVTEFAK[272]	2	719.4132	4.454	0.4
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QNCELFEQLGEYK[272]	2	967.9652	6.123	0.494
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RPCFSALEVDETYVPK[272]	2	1094.5547	3.134	0.28
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SALEVDETYVPK[272]	2	819.9451	4.898	0.516
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEMPADLPSLAADFVESK[272]	2	1626.2611	5.918	0.625
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEM[147]PADLPSLAADFVESK[272]	2	1634.2586	5.819	0.545
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TCVADESAENCDK[272]	2	882.8612	4.845	0.584
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TFHADICTLSEK[272]	2	849.9253	4.682	0.466
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VFDEFK[272]PLVEEPQNLIK[272]	2	1239.2045	4.628	0.615
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDR	2	1099.4228	4.473	0.477
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VMDDFAAFVEK[272]	2	780.4046	3.863	0.448
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YAPELLFFAK[272]	2	743.9311	4.088	0.354
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YFYAPELLFFAK[272]	2	898.9969	5.206	0.557
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YICENQDSISSK[272]	2	860.9099	4.147	0.419
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AAFTECCQAADK	3	543.9013	1.207	0.066
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ALVLIAFAQYLQQCPFEDHVK	3	920.4852	5.27	0.568
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AQYLQQCPFEDHVK	3	677.9975	2.605	0.11
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCAAADPHECYAK	3	600.567	1.828	0.544
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVFLGM[147]FLYEYAR	3	593.964	4.771	0.383
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ECCEKPLLEK	3	568.6166	2.411	0.022
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHADICTLSEK	3	843.7311	6.119	0.531
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EQLKAVMDDFAAFVEK	3	754.4048	4.63	0.535
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ETYGEM[147]ADCCAK	3	570.2218	1.468	0.213
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPDYSVVLLLR	3	484.6172	4.46	0.223
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KQTALVELVK	3	516.9994	2.132	0.056

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KVPQVSTPTLVEVSR	3	640.7141	2.558	0.274
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KYLYEIAR	3	445.931	2.961	0.374
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	IAFAQYLQQCPFEDHVK	3	788.394	5.082	0.457
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVMCTAFHDNEETFLK	3	973.8116	6.495	0.557
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVM[147]CTAFHDNEETFLK	3	979.1432	4.963	0.53
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	M[147]PCAEDYLSVVLNQLCVLHEK	3	931.4486	4.146	0.464
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NYAEAKDVFLGM[147]FLYEYAR	3	866.1017	3.73	0.399
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QNCELFEQLGEYK	3	642.9745	4.592	0.454
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QNCELFEQLGEYKFQNALLVR	3	956.8227	6.205	0.436
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPDYSVVLLLR	3	536.6509	3.13	0.426
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPYFYAPELLFFAK	3	727.0666	4.331	0.5
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RPCFSALEVDETYVPK	3	727.3675	2.281	0.089
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEMPADLPSLAADFVESK	3	1081.8384	2.275	0.297
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEM[147]PADLPSLAADFVESK	3	1087.1701	5.633	0.535
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TCVADESAENCDK	3	586.2385	4.238	0.53
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VFDEFKPLVEEPQNLIK	3	822.465	3.958	0.183
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGDLLECADDRADLAK	3	944.7403	5.455	0.574
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YKAAFTECCQAADK	3	687.6524	3.168	0.481
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YTKKVPQVSTPTLVEVSR	3	818.1477	3.654	0.486
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AAFTECCQAADK[272]	3	546.5727	1.799	0.174
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AQYLQQCPFEDHVK[272]	3	680.6689	2.602	0.207
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CTAFHDNEETFLK[272]K[272]	3	721.0309	4.607	0.327
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ECCEK[272]PLLEK[272]	3	572.6237	3.05	0.401
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EFNAETFTFHADICTLSEK[272]	3	846.4025	4.348	0.45
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVMCTAFHDNEETFLK[272]	3	976.483	7.422	0.535
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVM[147]CTAFHDNEETFLK[272]	3	981.8146	4.687	0.311
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NYAEAK[272]DVFLGM[147]FLYEYAR	3	868.7731	4.042	0.489
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QEPERNECFLQHK[272]	3	664.6605	4.934	0.356
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QNCELFEQLGEYK[272]FQNALLVR	3	959.4941	6.254	0.475
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RHPDYSVVLLLR	3	537.9866	3.686	0.399
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RHPYFYAPELLFFAK[272]	3	729.738	5.019	0.381
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RPCFSALEVDETYVPK[272]	3	730.0389	3.259	0.461
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEMPADLPSLAADFVESK[272]	3	1084.5098	7.996	0.658
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEM[147]PADLPSLAADFVESK[272]	3	1089.8415	5.796	0.4
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TCVADESAENCDK[272]	3	588.9099	4.151	0.356
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VFDEFK[272]PLVEEPQNLIK[272]	3	826.4721	3.616	0.354
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDR	3	733.2843	4.329	0.503
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDRADLAK[272]	3	947.4117	6.649	0.593
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YK[272]AAFTECCQAADK[272]	3	691.6595	3.515	0.316
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ALVLIAFAQYLQQCPFEDHVK	4	690.6158	2.951	0.38
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHADICTLSEKER	4	704.3361	4.472	0.499
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVMCTAFHDNEETFLK	4	730.6105	5.399	0.465
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVM[147]CTAFHDNEETFLK	4	734.6093	1.991	0.144
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVM[147]CTAFHDNEETFLKK	4	801.6567	4.966	0.439
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QEPERNECFLQHK	4	496.7437	2.783	0.271
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QNCELFEQLGEYKFQNALLVR	4	717.8689	3.507	0.452
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEMPADLPSLAADFVESK	4	811.6306	3.096	0.301
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SHCIAEVENDEM[147]PADLPSLAADFVESK	4	815.6294	4.946	0.539

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGDLLECADDRADLAK	4	708.8071	3.73	0.44
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVMCTAFHDNEETFLK[272]K[272]	4	800.6633	4.932	0.272
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDR	4	550.2151	3.004	0.429
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDRADLAK[272]	4	710.8106	5.761	0.541
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVMCTAFHDNEETFLKK	5	638.3279	3.736	0.354
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVMCTAFHDNEETFLK[272]K[272]	5	640.7321	5.021	0.523
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVM[147]CTAFHDNEETFLK[272]K[272]	5	643.9311	2.574	0.336
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLECADDRADLAK[272]	5	568.85	3.839	0.306
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LAKTYETTLEK	3	573.0014	2.364	0.105
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKCASLQKFGER	3	616.0112	3.933	0.302
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NECFLQHK	3	448.8899	3.032	0.182
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QEPERNECFLQHK	3	661.9891	4.448	0.354
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AVM[147]DDFAAFVEK[272]	3	549.6161	3.658	0.258
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ETYGEM[147]ADCCAK[272]	3	572.8932	3.334	0.451
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]HPYFYAPELLFFAK[272]	3	677.7043	4.079	0.347
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LDELRDEGK[272]ASSAK[272]	3	651.0322	3.303	0.443
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LK[272]CASLQK[272]FGER	3	620.0183	3.054	0.254
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NECFLQHK[272]	3	451.5613	2.729	0.117
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHADICTLSEK	4	633.0502	3.09	0.345
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVMCTAFHDNEETFLKK	4	797.658	3.994	0.248
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGDLLECADDR	4	549.2133	2.574	0.361
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EFNAETFTFHADICTLSEK[272]ER	4	706.3397	4.762	0.415
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LK[272]ECCEK[272]PLLEK[272]	4	526.0399	1.968	0.02
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVMCTAFHDNEETFLK[272]	4	732.6141	4.143	0.3
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVM[147]CTAFHDNEETFLK[272]	4	736.6128	3.297	0.092
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QNCELFEQLGEYK[272]FQNALLVR	4	719.8724	3.54	0.338
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DLGEENFK	2	616.3195	3.99	0.311
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NECFLQHK	2	672.8312	3.2	0.346
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCTESLVNR	2	630.7699	3.019	0.386
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DVFLGMFLYEYAR	2	884.4485	4.101	0.345
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ECCEK[272]PLLEK[272]	2	858.4319	3.685	0.382
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NECFLQHK[272]	2	676.8383	3.339	0.377
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QTALVELVK[272]	2	644.9076	2.689	0.235
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CTAFHDNEETFLKK	3	717.0238	5.044	0.342
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FKDLGEENFK	3	549.6349	2.533	0.039
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKECCEKPLLEK	3	695.7079	2.544	0.045
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AVMDDFAAFVEK[272]	3	544.2845	2.876	0.262
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCAAADPHECYAK[272]	3	603.2384	1.305	0.465
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FK[272]DLGEENFK[272]	3	553.642	3.066	0.319
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKCASLQKFGER	4	462.2602	2.724	0.294
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RPCFSALEVDETYVPK	4	545.7774	2.102	0.112
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEM[147]PADLPSLAADFVESK[272]	4	817.6329	3.175	0.251
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VFDEFK[272]PLVEEPQNLIK[272]	4	620.1059	1.885	0.106
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LCTVATLR	2	531.7936	2.833	0.272
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TYETTLEK	2	632.8428	3.339	0.314
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DLGEENFK[272]	2	620.3266	3.623	0.314
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DVCK[272]NYAEAK[272]	3	540.2813	2.755	0.117
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LK[272]ECCEK[272]PLLEK[272]	3	701.0507	2.656	0.014

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YICENQDSISSK[272]	3	574.2757	1.853	0.09
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QEPERNECFLQHK[272]	4	498.7472	3.49	0.26
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DLGEENFK	1	1231.6318	2.658	0.335
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TYETTLEK	1	1264.6784	2.28	0.288
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TYETTLEK[272]	1	1272.6926	2.407	0.304
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DDNPNLPR	2	540.7753	2.883	0.316
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KYLYEIAR	2	668.3928	2.621	0.313
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LCTVATLR	2	533.7971	2.812	0.346
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AVM[147]DDFAAFVEK	3	546.9447	2.642	0.096
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPDYSVVLLLR	2	726.4221	2.285	0.424
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TYETTLEK[272]	2	636.8499	2.648	0.227
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YICENQDSISSK	3	571.6043	1.796	0.035
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AEFAEVSK	1	1160.6311	2.481	0.066
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AEFAEVSK[272]	1	1168.6453	2.905	0.048
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DDNPNLPR	1	1084.5504	2.2	0.299
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YLYEIAR	2	534.2979	2.505	0.117
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DDNPNLPR	1	1080.5433	1.91	0.311
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YLYEIAR	1	1067.5885	2.485	0.127
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YLYEIAR	1	1071.5956	2.423	0.119
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKECCEKPLLEK	4	522.0328	2.602	0.226
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LK[272]CASLQK[272]FGER	4	465.2656	2.234	0.231
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SIHTLFGDK	1	1297.7264	2.868	0.274
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LDELRDEGK[272]	1	1362.7468	2.185	0.271
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SLHTLFGDK[272]	1	1305.7406	3.102	0.266
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DDNPNLPR	2	542.7788	2.281	0.19
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AEFAEVSK	2	580.8192	2.656	0.032
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVTDLTK[272]	2	539.3416	1.491	0.104
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LDELRDEGK	1	1354.7326	2.427	0.225
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVTDLTK	1	1069.6617	2.119	0.085
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVTDLTK	2	535.3345	2.415	0.102
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AEFAEVSK[272]	2	584.8263	3.664	0.07
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LAKTYETTLEK	2	858.9984	3.58	0.386
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVTDLTK[272]	1	1077.6759	1.923	0.042
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AACLLPK[272]	2	525.3081	2.062	0.084
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AACLLPK	2	521.301	2.231	0.104
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QEPERNECFLQHK[272]	2	996.4871	4.303	0.369
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHADICTLSEKER	3	938.779	3.567	0.34
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CTAFHDNEETFLK[272]K[272]	4	541.025	3.211	0.334
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FQNALIVR	2	550.8324	3.137	0.218
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EPERNECFLQHK[272]	3	621.9743	3.874	0.365
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NECFLQHKDDNPNLPR	4	567.2769	3.025	0.197
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KDLGEENFK	3	500.6121	3.852	0.332
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YLYEIAR	2	536.3014	1.823	0.031
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LAK[272]TYETTLEK[272]	2	865.0091	3.883	0.323
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LCTVATLR	1	1066.587	1.901	0.165
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SIHTLFGDK[272]	2	653.3739	1.949	0.197
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YTKKVPQVSTPTLVEVSR	4	613.8626	2.907	0.272
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SIHTLFGDK	2	649.3668	3.473	0.235

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGD	3	411.482	2.073	0.398
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AEFAEVSKLVTDLTK	3	691.0648	2.985	0.265
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LDELRDEGK	2	677.8699	2.878	0.242
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FQNALIVR	1	1104.6647	2.248	0.187
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FQNALLVR	1	1100.6576	2.297	0.179
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPELLFFAK	3	675.0329	2.863	0.13
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVNEVTEFAK	3	477.2732	2.425	0.134
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QLCVLHEK[272]	3	435.2417	3.108	0.271
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AACLLPK[272]	1	1049.609	1.764	0.11
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TFHADICTLSEK[272]	3	566.9526	3.063	0.282
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKECCEKPLL	3	563.2971	2.261	0.257
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NYAEAK	2	488.2666	1.936	0.042
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ADDK[272]ETCFAEEGQK[272]	3	683.6592	3.192	0.306
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NYAEAK[272]	2	492.2737	1.795	0.108
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LDELRDEGK[272]	3	454.9204	2.139	0.239
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LDELRDEGK[272]	2	681.877	2.741	0.17
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVDVM[147]CTAFHDNEETFLKK	5	641.5268	2.87	0.041
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LDELRDEGK	3	452.249	2.395	0.195
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VFDEFKPLVEEPQNLIK	4	617.1006	1.996	0.152
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKCASLQK	2	678.8964	2.777	0.262
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TPVSDRVTK	2	641.8776	2.508	0.263
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AACLLPK	1	1041.5948	1.714	0.088
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ATKEQLK	3	413.2591	2.161	0.198
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SIHTLFGDK[272]	3	435.9184	3.187	0.259
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CASIQK[272]	2	492.2665	1.47	0.143
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ADDKETCFAEEGQK	3	679.6521	3.029	0.354
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SIHTLFGDK	3	433.247	2.485	0.15
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LCTVATLR	1	1062.5799	1.972	0.064
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YFYAPELLFFAK[272]	3	599.667	3.015	0.25
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]AEFAEVSK[272]LVTDLTK[272]	3	695.0719	2.43	0.243
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ATK[272]EQLK[272]	3	417.2662	2.466	0.13
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ADIAK	2	399.2477	1.927	0.203
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CASIQK	2	488.2594	1.741	0.13
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TFHADICTLSEK	2	845.9182	3.833	0.399
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGD	2	616.7194	2.566	0.487
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DESAENCDK	2	668.7791	2.983	0.443
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	PHECYAK	3	391.8563	2.641	0.143
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TALVELVK[272]	2	580.8783	2.643	0.308
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVCKNYAEAK	3	536.2742	2.8	0.105
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DFAAFVEK[272]	2	607.8366	3.607	0.344
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HADICTLSEK	2	721.8602	3.244	0.401
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NLGK[272]VGSK[272]	2	617.8959	1.852	0.313
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ESAENCDK	2	611.2656	2.693	0.445
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CTESLVNR	2	556.2715	1.944	0.168
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]APELLFFAK[272]	2	662.3994	3.664	0.315
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NLGKVGSK	2	611.8852	2.128	0.266
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]HADICTLSEK[272]	2	725.8673	3.061	0.439
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LAK[272]TYETTLEK[272]	3	577.0085	2.385	0.148

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CFSALEVDETYVPK	2	963.9707	3.729	0.387
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SEVAHR	2	419.7301	1.482	0.138
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LKCASLQK	3	452.9333	3.023	0.057
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DYSVVLLLR	2	609.3663	2.639	0.269
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TALVELVK	2	576.8712	3.131	0.223
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	PHECYAK	2	587.2808	2.555	0.389
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QCPFEDHVK[272]	3	479.5683	2.755	0.208
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FKPLVEEPQNLIK	3	659.0628	3.046	0.217
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FSALEVDETYVPK	2	889.4722	3.683	0.293
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LGM[147]FLYEYAR	2	709.8605	2.963	0.34
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	APELLFFAK	2	658.3923	3.288	0.274
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]PELLFFAK[272]	2	626.8809	3.433	0.253
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EFNAETFTFHADICTLSEK[272]	4	635.0537	2.398	0.122
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VHTECCHGDLLECADDR	3	731.9486	2.424	0.104
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QVSTPTLVEVSR	2	728.4119	2.45	0.147
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGD	2	618.723	1.91	0.513
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FYAPELLFFAK[272]	3	545.3126	2.639	0.238
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVCK	2	395.7012	1.198	0.182
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DDFAAFVEK	2	661.343	2.685	0.32
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EPERNECFLQHK	4	464.729	2.543	0.267
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DYSVVLLLR	2	611.3698	1.786	0.081
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ADICTLSEK[272]	2	657.3378	2.997	0.306
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]QVSTPTLVEVSR	2	730.4155	2.58	0.171
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPELLF	2	768.8979	2.819	0.314
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ECFLQHK	3	410.8756	2.092	0.246
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ALEVDETYVPK	2	772.422	2.913	0.312
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ATKEQLK	2	619.385	2.366	0.054
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FLYEYAR	2	551.29	2.182	0.236
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPYFYAPELL	2	773.4143	2.366	0.06
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	PELLFFAK	2	622.8738	3.169	0.353
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ESAENCDK[272]	2	615.2727	2.34	0.339
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ELLFFAK	2	574.3474	2.61	0.24
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	SALEVDETYVPK	2	815.938	3.903	0.418
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YLQQCPFEDHVK	3	611.6323	4.971	0.456
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VNEVTEFAK[272]	2	662.8712	2.654	0.29
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CVLHEK[272]	2	531.7876	2.428	0.209
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CTVATLR	2	477.2551	2.278	0.255
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QLCVLHEK	2	648.3518	2.737	0.21
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RM[147]PCAEDYLSVVLNQL	2	1029.003	2.39	0.322
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FSALEVDETYVPK[272]	2	893.4793	3.017	0.328
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EPERNECFLQHK	3	619.3029	2.903	0.25
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LQQCPFEDHVK	3	557.2778	2.56	0.229
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPELL	2	695.3637	2.509	0.257
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EFAEVSK[272]	2	549.3077	3.16	0.135
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TECCQAADK	2	670.7769	2.543	0.256
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPEL	2	638.8217	1.934	0.114
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EVDETYVPK[272]	2	684.3685	2.87	0.194
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	HPYFYAPELLFF	2	842.4321	2.294	0.321

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DDFAAFVEK[272]	2	665.3501	2.633	0.281
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCKHPEAK	3	476.5641	2.283	0.077
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCKHPEAK	2	714.3426	2.083	0.016
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LGEENFK	2	558.8061	2.741	0.142
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHAD	2	784.8545	2.207	0.325
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CVLHEK	2	527.7805	2.238	0.165
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DVFLGM[147]FLYEYAR	3	595.2997	2.059	0.068
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFAEVSK	2	545.3006	2.94	0.106
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	YLQQCPFEDHVK	2	916.9448	3.681	0.465
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]PHECYAK[272]	2	591.2879	2.285	0.18
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FYAPELLFFAK	3	542.6412	2.783	0.06
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KVPQVSTPTLV	2	724.9454	2.074	0.255
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LGMFLYEYAR	2	701.863	2.55	0.312
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]HADICTLSEK[272]	3	484.2473	2.679	0.136
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EK[272]PLLEK[272]	3	430.2782	2.425	0.093
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EKPLLEK	3	426.2711	2.747	0.078
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LYEYAR	2	479.7594	1.845	0.11
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KVPQVSTPTLVEV	2	839.0009	2.257	0.265
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TPVSDRVTK	3	428.2541	2.177	0.051
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]EQLGEYK[272]	2	577.8184	2.498	0.129
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TESLVNR	2	479.7695	2.317	0.127
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ETYVPK	2	508.7924	2.07	0.15
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ADFVESK	2	538.2928	2.292	0.131
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TAFHDNEETFLKK	4	500.7705	3.421	0.134
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ALVLIAFAQ	1	1085.6718	1.794	0.523
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EQLGEYK	2	573.8113	2.337	0.108
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LVRPEVD	2	484.2822	1.528	0.263
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ECFLQHK	2	615.8098	1.989	0.216
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LYEYAR	2	477.7558	1.6	0.041
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVDVMCTAFH	3	602.9656	2.912	0.11
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPDYSVVL	2	613.338	2.009	0.202
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EVDETYVPK	2	680.3614	2.361	0.258
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RPCFSALEVDE	2	726.3365	1.298	0.387
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KDLGEENFK	2	750.4145	3.506	0.213
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DVFLGMFL	1	1081.5751	1.912	0.346
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	AVMDDFAAFVEK	3	541.6131	1.843	0.04
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LLECADDR	2	562.7635	2.137	0.129
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FAAFVEK[272]	2	550.3232	2.005	0.139
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YLQQCPFEDHVK[272]	3	614.3037	3.621	0.34
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ELLFFAK[272]	2	578.3545	2.092	0.101
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DNPNLPR	2	483.2618	1.957	0.159
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	VSTPTLVEVSR	2	664.3827	2.392	0.141
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ECCEK[272]	2	496.2016	1.788	0.169
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RPCFSALEVD	2	663.8188	1.951	0.164
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DVFLGMFL	1	1085.5822	1.985	0.326
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCAAAD	1	789.252	1.321	0.57
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	EFNAETFTFHA	2	727.341	2.286	0.289
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ECFLQHK[272]	3	413.547	2.315	0.111

56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPDYSVVLL	2	669.8801	2.053	0.236
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]NECFLQH	2	540.7398	1.917	0.122
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RHPYFYAPEL	2	718.8758	2.125	0.089
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LLFFAK[272]	2	513.8332	1.704	0.126
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ECCEK	2	492.1945	1.776	0.147
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	PLLEK	2	440.2868	1.519	0.055
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	LDELRDEGKASSAK	4	485.5207	2.256	0.031
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FK[272]PLVEEPQNLIK[272]	3	663.0699	2.152	0.129
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NLGKVGSK	3	408.2592	1.713	0.064
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RHPYFYAPEL	2	716.8723	2.058	0.158
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LGEENFK[272]	2	562.8132	2.122	0.075
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RHPDYSVVLLL	2	728.4257	1.706	0.222
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KPLLEK	3	383.2569	2.599	0.063
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	QNALLVR	2	477.2982	2.25	0.1
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VTDLTK[272]	2	482.7995	2.013	0.049
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VFDEFK[272]	2	536.7995	2.015	0.046
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]YLQQCPFEDHVK[272]	2	920.9519	4.393	0.335
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	ADICTLSEK	2	653.3307	2.644	0.243
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]IYEIAR	2	454.7697	2.144	0.122
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	KVPQVSTPTLVEVS	2	882.517	2.651	0.424
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLLE	2	796.3283	1.656	0.239
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CTAFHDNEETFLKK	4	538.0197	2.233	0.031
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CCAAAD	1	785.2449	1.34	0.407
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGD	3	412.8177	1.451	0.196
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]PLLEK[272]	2	444.2939	1.942	0
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]TECCQAADK[272]	2	674.784	1.931	0.105
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]VHTECCHGDLL	2	731.807	2.095	0.12
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]LVRPEVD	2	486.2858	1.559	0.126
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RHPDYSVV	2	558.7996	1.829	0.13
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	RPCFSAL	2	490.2462	1.689	0.117
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RPCFSALEVDET	2	778.8639	1.97	0.136
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]FK[272]DLGEEN	2	620.3266	1.912	0.262
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]RM[147]PCAEDYLSVVLNQL	3	686.3377	2.426	0.084
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TFHADICTLSEK	3	564.2812	1.912	0.135
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]SHCIAEVENDEM[147]PAD	2	933.3773	1.687	0.39
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]DELRDEGK[272]	3	417.2257	2.222	0.116
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	NECFLQH	2	538.7362	1.533	0.053
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	CQAADK	2	481.2334	1.528	0.049
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	TAFHDNEETFLK	2	866.4387	3.075	0.309
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	DELRDEGK	2	621.3279	2.296	0.107
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	FKDLGEEN	2	616.3195	1.758	0.25
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]ADDK[272]ETCFAEEG	2	824.8573	2.224	0.215
56	IPI00022434	1	86.3	1.24	0.15	###	##	ALB Uncharacterized protein ALB	n[145]CCAAADPH	2	512.1855	1.346	0.093
57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	n[145]GK[272]EESLDSDLYAELR	2	1007.0226	4.78	0.433
57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	n[145]GK[272]EESLDSDLYAELR	3	671.6842	2.928	0.458
57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	n[145]GTHCNQVEVIATLK[272]	3	616.3301	3.688	0.316
57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	GTHCNQVEVIATLK	3	613.6587	3.214	0.281
57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	GKEESLDSDLYAELR	3	669.0128	2.893	0.275

57	IPI00022445	1	29.7	0.26	0.01	7	6	PPBP Platelet basic protein precursor	n[145]ICLDPDAPR	2	595.2949	2.146	0.12
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	NFPSPVDAAFR	1	1360.7009	3.116	0.478
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]NFPSPVDAAFR	1	1364.708	3.508	0.47
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VDGALCMEK[272]	1	1299.635	3.075	0.431
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GGYTLVSGYPK	2	711.393	3.986	0.38
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	QGHNSVFLIK	2	711.9145	4.115	0.389
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	SGAQATWTELPWPHEK[272]	2	1061.5456	4.395	0.446
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILDSVDAAFICPGSSR	2	1316.155	5.944	0.555
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EWFWDLATGTMK[272]	2	886.9497	3.371	0.491
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GECQAEGVLFFQGDR	2	923.4227	4.843	0.417
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GGYTLVSGYPK[272]	2	715.4001	3.313	0.256
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GIILDSVDAAFICPGSSR	2	1006.0091	5.108	0.488
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]LLQDEFPGIPSPLDAAVECHR	2	1249.1204	5.348	0.507
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VDGALCMEK[272]	2	650.3211	3.552	0.383
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YYCFQGNQFLR	2	814.8772	2.478	0.415
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GNVAEGETKPDPDVTER	3	698.6869	3.347	0.362
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GPGLYLIHGPNLYCYSDVEK	3	855.4274	5.81	0.522
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILDSVDAAFICPGSSR	3	877.7724	4.156	0.397
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GLYLIHGPNLYCYSDVEK[272]	3	806.7407	5.147	0.474
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GPGLYLIHGPNLYCYSDVEK[272]	3	858.0988	5.214	0.495
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]LLQDEFPGIPSPLDAAVECHR	3	833.0827	4.065	0.318
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]SANGPGLYLIHGPNLYCYSDVEK[272]	3	948.8028	6.009	0.467
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]DYFMPCPGR	2	638.2757	2.784	0.294
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]NFPSPVDAAFR	2	682.8576	2.164	0.282
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]QGHNSVFLIK[272]	2	715.9216	4.121	0.31
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSDNHG	3	614.2983	2.723	0.223
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	VDGALCMEK	2	646.314	3.024	0.47
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]ALPQPQNVTSLLGCTH	2	934.9776	3.422	0.389
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YYCFQGNQFLR	3	543.5872	3.076	0.218
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	NFPSPVDAAFR	2	680.8541	2.986	0.331
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VDGALCM[147]EK[272]	2	658.3186	2.46	0.302
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]DYFM[147]PCPGR	2	646.2731	2.374	0.238
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	VDGALCM[147]EK	2	654.3115	2.67	0.306
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GLYLIHGPNLYCYSDVEK	3	804.0693	4.34	0.385
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]DYFMPCPGR	1	1275.5441	1.85	0.214
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]SLGPNSCSANGPGLYLIHGPN	3	753.3653	4.356	0.331
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]FDPVRGEVPPR	3	471.5971	1.555	0.23
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSDNHGA	3	637.9773	3.871	0.321
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GNVAEGETK[272]PDPDVTER	3	701.3583	3.82	0.196
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]FDPVRGEVPPR	2	706.892	2.391	0.104
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSDNHGATY	3	726.0143	3.028	0.353
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]LLQDEFPGIPSPLDAAVECHR	4	625.0639	2.273	0.136
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	EWFWDLATGTM[147]K	2	890.94	2.121	0.222
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]FDPVR	2	389.2224	1.747	0.08
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GEFVWK	2	523.2951	2.264	0.092
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	EVGTPHGIILDSVDAAFICPGSSR	3	876.4367	2.756	0.253
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GEFVWK[272]	2	527.3022	2.517	0.052
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSDNHGATYA	3	749.6933	2.823	0.389

58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	EIISER	1	886.4993	1.844	0.112
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]ALPQPQNVTSLLGCTH	3	623.6541	2.409	0.115
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	VAEGETKPDPDVTER	3	641.6655	2.398	0.384
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YCFQGNQFLR	2	733.3455	3.305	0.424
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EK[272]GYPK[272]	3	385.2362	2.004	0.145
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YYCFQGNQFL	2	736.8266	2.404	0.451
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VTSLLGCTH	2	560.7842	2.93	0.331
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	YYCFQGNQFLR	2	812.8736	1.748	0.121
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YYCFQGNQF	2	680.2846	2.816	0.374
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	EKGYPK	2	571.3401	2.166	0.125
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VWVYPPEK	2	651.3724	2.518	0.194
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]TSDNHGATYAFSGTHYWR	3	739.0073	3.59	0.355
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]IHLM[147]AGR	3	319.8522	2.272	0.149
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GALCM[147]EK	2	547.2638	2.368	0.26
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]DYFM[147]PCPGR	3	431.1845	1.815	0.025
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	NVAEGETKPDPDVTER	3	679.6798	2.954	0.153
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]ETK[272]PDPDVTER	2	787.9169	3.212	0.196
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GECQAEGVLFFQGDR	3	615.9509	2.307	0.01
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]SVFLIK[272]	2	497.8306	2.606	0.137
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]AAVECHR	2	488.2347	2.751	0.135
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILDSVDAA	2	869.4606	3.292	0.383
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	SVFLIK	2	493.8235	2.285	0.18
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSDNHGATYAF	3	798.7161	2.473	0.296
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]VWVYPPEKK	3	523.9807	2.337	0.131
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GEVPPRYPR	2	607.8418	2.393	0.046
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]GALCM[147]EK[272]	2	551.2709	2.206	0.187
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTSD	2	766.8821	2.561	0.358
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]AAFICPGSSR	2	599.7951	2.741	0.123
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EK[272]GYPK[272]	2	577.3507	1.922	0.131
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]YFM[147]PCPGR	2	588.7597	1.594	0.327
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]LYCYSDVEK[272]	2	727.3509	2.425	0.18
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	ETKPDPDVTER	2	783.9098	2.186	0.184
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CYSDVEK[272]	2	589.2772	1.64	0.208
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]ELISER	1	890.5064	1.565	0.089
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILDSVD	2	798.4235	2.709	0.263
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	GEFVWK[272]	2	525.2987	1.713	0.028
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	CYSDVEK	2	585.2701	1.265	0.277
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]WKNFPSPVDAAFR	3	606.9972	3.04	0.134
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSAL	2	615.3288	2.134	0.149
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]ICPGSSR	2	455.2238	1.721	0.109
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	LYCYSDVEK	2	723.3438	1.995	0.057
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]CSPHLVLSALTS	2	709.3686	1.835	0.126
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILD	2	647.8598	2.123	0.072
58	IPI00022488	1	62.8	0.8	0.14	222	86	HPX Hemopexin precursor	n[145]EVGTPHGIILDSVDA	2	833.9421	2.183	0.065
59	IPI00022731	1	16.5	0.67	0.03	4	3	APOC4 Apolipoprotein C-IV precursor	n[145]GFMQTYYDDHLR	3	563.9319	4.57	0.498
59	IPI00022731	1	16.5	0.67	0.03	4	3	APOC4 Apolipoprotein C-IV precursor	n[145]GFM[147]QTYYDDHLR	3	569.2635	2.275	0.177
59	IPI00022731	1	16.5	0.67	0.03	4	3	APOC4 Apolipoprotein C-IV precursor	n[145]ELLETVVNR	2	608.8545	2.64	0.1
60	IPI00022733	1	4.9	1.01	0.01	4	3	PLTP 45 kDa protein	n[145]GAFFPLTER	2	591.3254	3.024	0.282

60	IPI00022733	1	4.9	1.01	0.01	4	3	PLTP 45 kDa protein	GAFFPLTER	2	589.3219	2.632	0.206
60	IPI00022733	1	4.9	1.01	0.01	4	3	PLTP 45 kDa protein	n[145]AVEPQLQEEER	2	736.3791	2.503	0.273
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LLELTGPK[272]	1	1158.7337	2.948	0.271
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]CEGPIPDVTFELLR	2	889.9505	3.979	0.454
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]HQFLLTGDTQGR	2	758.9031	3.89	0.429
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]TPGAAANLELIFVGPQHAGNYR	2	1220.6463	5.434	0.557
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	NGVAQEPVHLDSPAIK	3	652.3618	4.991	0.514
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]AVALGDGGHYTCR	3	503.9068	4.698	0.402
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]GAAANLELIFVGPQHAGNYR	3	748.0665	5.854	0.468
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]HQFLLTGDTQGR	3	506.2711	4.132	0.368
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]NGVAQEPVHLDSPAIK[272]	3	655.0332	4.818	0.506
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]TPGAAANLELIFVGPQHAGNYR	3	814.1	6.481	0.503
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LELHVDGPPPRPQLR	3	623.3589	2.405	0.144
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LELHVDGPPPRPQLR	4	467.771	1.971	0.267
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]IFFHLNAVALGDGGHYTCR	4	571.0336	3.344	0.279
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LETPDFQLFK[272]	2	763.4289	1.763	0.171
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LELHVDGPPPRPQLR	3	622.0232	2.029	0.11
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LLELTGPK	2	575.8634	2.677	0.23
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]CLAPLEGAR	2	560.2922	2.449	0.213
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LLELTGPK[272]	2	579.8705	1.435	0.094
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LETPDFQLFK	2	759.4218	2.683	0.228
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	HHGESSQVLHPGNK	4	452.4894	2.85	0.375
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]GVTFLLR	2	475.3012	3.137	0.177
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]SWVPHTFESELSDPVELLVAES	2	1308.1535	2.511	0.453
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LELHVDGPPPRPQLR	4	466.7693	2.489	0.017
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]HHGESSQVLHPGNK[272]	4	454.493	3.924	0.32
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LELHVDGPPPRPQLR	2	932.5312	2.423	0.4
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	ELLVPR	1	866.5459	1.803	0.173
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LELHVDGPPPRPQLR	2	934.5348	2.605	0.135
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]ELLVPR	1	870.553	1.852	0.013
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]ELLVPR	2	435.7801	2.007	0.014
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]LELIFVGPQHAGNYR	3	620.0079	3.106	0.309
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	GVTFLLR	2	473.2977	2.06	0.015
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]AVALGDGGHYTCR	2	755.3566	4.12	0.344
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]VAPLSGVDFQLR	2	723.4153	3.275	0.287
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	n[145]VTFELLR	2	511.3118	2.532	0.179
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	HHGESSQVLHPGNK	3	602.9835	2.011	0.171
61	IPI00022895	1	37.6	0.77	0.12	98	35	A1BG Alpha-1B-glycoprotein precursor	LELHVD	2	433.2426	1.325	0.139
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	SQHLDNFSNQIGK	3	589.9761	4.232	0.342
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	n[145]SQHLDNFSNQIGK[272]	3	592.6475	4.141	0.326
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	DIHSGLIGPLLICQK	3	645.0302	4.734	0.255
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	n[145]DIHSGLIGPLLICQK[272]	3	647.7016	4.209	0.343
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	n[145]FCENPDEVK[272]R	3	524.2581	2.569	0.321
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	n[145]YLDSTFTK[272]	2	631.8472	1.968	0.112
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	SK[272]K[272]	2	395.7731	0.913	0.136
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	n[145]NLASRPY	2	482.7703	2.083	0.1
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	LLLLK	2	440.3232	1.8	1
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	FEGNTNTK	2	595.8119	2.135	0.003

62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	SPLR	1	612.3828	1.332	0.125
62	IPI00022937	1	3.6	1.07	0.46	10	6	F5 Coagulation factor V	EM[147]K	1	703.3808	1.121	0.072
63	IPI00023019	1	10	0.89	0.04	6	5	SHBG Isoform 1 of Sex hormone-binding globulin precursor	n[145]IALGGLLFPASNLR	2	793.481	4.124	0.423
63	IPI00023019	1	10	0.89	0.04	6	5	SHBG Isoform 1 of Sex hormone-binding globulin precursor	n[145]LPLVPALDGCLR	2	728.9105	2.182	0.076
63	IPI00023019	1	10	0.89	0.04	6	5	SHBG Isoform 1 of Sex hormone-binding globulin precursor	LVPALDGCLR	2	621.8385	2.093	0.245
63	IPI00023019	1	10	0.89	0.04	6	5	SHBG Isoform 1 of Sex hormone-binding globulin precursor	LPLVPALDGCLR	2	726.9069	1.79	0.019
63	IPI00023019	1	10	0.89	0.04	6	5	SHBG Isoform 1 of Sex hormone-binding globulin precursor	n[145]LVPALDGCLR	2	623.8421	2.205	0.147
64	IPI00023673	1	4.7	0.88	0.09	4	4	LGALS3BP Galectin-3-binding protein precursor	KSQLVYQSR	3	463.6052	3.169	0.22
64	IPI00023673	1	4.7	0.88	0.09	4	4	LGALS3BP Galectin-3-binding protein precursor	STHTLDLSR	2	585.3173	2.974	0.341
64	IPI00023673	1	4.7	0.88	0.09	4	4	LGALS3BP Galectin-3-binding protein precursor	n[145]SQLVYQSR	2	562.8127	3.275	0.239
64	IPI00023673	1	4.7	0.88	0.09	4	4	LGALS3BP Galectin-3-binding protein precursor	n[145]YFYSR	2	440.2277	1.748	0.039
64	IPI00023673	1	4.7	0.88	0.09	4	4	LGALS3BP Galectin-3-binding protein precursor	SVK[272]	1	617.4104	0.88	0.378
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	GVLHNEVK	2	588.3484	2.938	0.309
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	n[145]GVLHNEVK[272]	3	395.2394	2.816	0.202
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	n[145]IQYSPAR	2	489.7781	1.866	0.082
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	n[145]GVLHNEVK[272]	2	592.3555	2.51	0.248
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	GVLHNEVK	3	392.568	2.153	0.083
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	ETQSIEK	2	557.8088	1.812	0.012
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	n[145]ETQSIEK[272]	2	561.8159	1.833	0.035
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	VYFFK	2	492.2893	1.735	0.167
65	IPI00024825	1	2.5	0.55	0.15	9	7	PRG4 Isoform A of Proteoglycan-4 precursor	PSLPP	2	325.6972	1.303	0.198
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	LADGPGHCK	3	408.539	2.191	0.268
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	n[145]LADGPGHCK[272]	3	411.2104	2.399	0.252
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	n[145]CYGPGVGR	2	499.7371	2.904	0.24
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	n[145]LADGPGHCK[272]	2	616.3119	2.67	0.295
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	LEVLHK	2	509.8241	2.284	0.123
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	n[145]LEVLHK[272]	2	513.8312	2.263	0.141
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	n[145]LEVLHK[272]	3	342.8899	2.108	0.032
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	LADGPGHCK	2	612.3048	1.972	0.094
66	IPI00025204	1	6.6	0.87	0.03	9	9	CD5L CD5 antigen-like precursor	LEVLHK	3	340.2185	1.847	0.01
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	AVGYLITGYQR	2	690.8854	3.71	0.44
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	GSFALSFPVESDVAPIAR	2	1002.0335	4.66	0.456
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	M[147]VSGFIPLKPTVK	2	927.0578	2.248	0.234
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]ATVLNYLPK[272]	2	653.9023	1.422	0.077
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]MVSGFIPLK[272]PTVK[272]	2	925.0709	3.962	0.447
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	MVSGFIPLKPTVK	3	613.0426	2.28	0.11
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	SGTHTLPVESGDM[147]K	3	585.6296	2.796	0.425
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	TEPQYMVLVPSLLHTEAPK	3	811.7785	5.251	0.482
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	VQTVPQTCDGHK	3	546.9412	4.632	0.516
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AGAFCLSEDAGLGISSTASLR	3	739.3651	2.89	0.13
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]MVSGFIPLK[272]PTVK[272]	3	617.0497	2.357	0.273
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]SGTHTLPVESGDM[147]K[272]	3	588.301	4.082	0.469
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VQTVPQTCDGHK[272]	3	549.6126	2.739	0.408
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	ATVLNYLPK	2	649.8952	2.663	0.344
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	M[147]VSGFIPLKPTVK	3	618.3743	1.861	0.035
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]HQDGSYSTFGER	3	509.903	2.865	0.341
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]IREEGTDLEVTANR	3	582.9749	4.454	0.312
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	SSGSLLNNAIK	2	692.4014	3.62	0.366

67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AVGYLITGYQR	2	692.8889	3.767	0.41
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]SSGSLLNNAIK[272]	2	696.4085	3.957	0.357
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]YGAATFTR	2	515.7756	2.604	0.238
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	EGTHGSHVYTK	3	499.2599	3.063	0.23
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]TEPQYM[147]VLVPSLLHTEAPK[272]	3	819.7816	5.043	0.379
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]DNGCFR	2	451.1925	2.125	0.346
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]ISEITNIVSK[272]	2	696.4211	3.295	0.336
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AFQPFFVELTM[147]PYSVIR	3	735.7227	3.847	0.203
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]TEPQYMVLVPSLLHTEAPK[272]	3	814.4499	5.168	0.322
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	YGAATFTR	1	1026.5368	2.424	0.24
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VQTVPQTCDGHK[272]	2	823.9153	2.91	0.422
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	DNGCFR	2	449.1889	2.036	0.331
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	SLFTDLVAEK	2	701.9007	2.844	0.253
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	TEPQYM[147]VLVPSLLHTEAPK	3	817.1102	4.958	0.349
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]YGAATFTR	1	1030.5439	1.848	0.355
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	VQTVPQTCDGHK	2	819.9082	2.677	0.245
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VVSVDENFRPR	3	487.9359	2.23	0.228
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]HQDGSYSTFGER	2	764.3509	2.988	0.194
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	GVPIPNK	2	502.8162	2.324	0.225
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AFQPFFVELTM[147]PYSVIR	2	1103.0804	4.222	0.318
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]GVPIPNK[272]	2	506.8233	2.066	0.197
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	VSGFIPLKPTVK	3	569.3625	2.601	0.35
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	VDSHFR	1	900.4687	1.824	0.075
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VDSHFR	2	452.7415	2.06	0.125
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]YNILPEK[272]EDSPFALK[272]	3	732.7474	2.727	0.309
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	VDSHFR	2	450.738	1.968	0.107
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]YNILPEK[272]	2	582.847	2.486	0.138
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VDSHFR	1	904.4758	1.669	0.064
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]FVELTM[147]PYSVIR	2	807.9377	3.594	0.266
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]NQGNTWLTAFVLK	2	888.4999	3.914	0.225
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]EGTHGSHVYTK[272]	3	501.9313	1.986	0.154
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	QLNYK	2	473.2795	1.725	0.097
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	DLKPAIVK	3	435.2869	2.515	0.232
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]FVELTMPYSVIR	2	799.9403	3.516	0.329
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VSGFIPLK[272]PTVK[272]	2	859.5507	2.916	0.469
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	M[147]VSGFIPLKPTVK	4	464.0325	1.742	0.022
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	HQDGSYSTFGER	2	762.3473	2.011	0.005
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FVELTM[147]PYSVIR	2	805.9342	2.746	0.284
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FVELTMPYSVIR	2	797.9367	3.457	0.208
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]QLNYK[272]	2	477.2866	1.602	0.005
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AGLGISSTASLR	2	638.8707	2.778	0.268
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	DLKPAIVK	2	652.4267	2.425	0.2
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FIPLKPTVK	2	731.9791	1.951	0.3
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]LK[272]PAIVK[272]	2	600.9239	2.228	0.263
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	YNILPEKEDSPFALK	3	728.7403	2.48	0.081
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	ALLAYAF	2	454.7657	1.983	0.287
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]VSGFIPLK[272]PTVK[272]	3	573.3696	2.821	0.167
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]NQGNTWLTAFVLK	3	592.669	2.126	0.115

67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	DLFHCVSF	2	577.2621	1.853	0.36
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]DLK[272]PAIVK[272]	2	658.4374	2.086	0.106
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	M[147]VSGFIPLK	2	644.3783	2.288	0.258
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]TM[147]PYSVIR	2	563.806	2.066	0.244
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]GFIPLK[272]PTVK[272]	3	511.3361	1.821	0.239
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	IPLKPTVK	2	658.4449	1.879	0.24
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FIPLKPTVK	3	488.3218	1.843	0.206
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	LKPAIVK	3	396.9446	2.98	0.033
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]TMPYSVIR	2	555.8085	2.107	0.14
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]FIPLK[272]PTVK[272]	2	737.9898	2.539	0.209
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]IPLK[272]PTVK[272]	2	664.4555	1.94	0.175
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	LKPAIVK	2	594.9132	1.839	0.153
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	RKDTVIK	2	640.4141	1.645	0.035
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	SDVAPIAR	2	484.7798	2.266	0.022
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AFQPFFVEL	2	621.338	2.101	0.048
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]NQGNTWLTAF	2	648.3287	2.02	0.284
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FVELTMPYSVIR	3	532.2936	2.128	0.093
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]ALLAYAF	2	456.7692	1.824	0.061
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	ALLAYAF	1	908.5241	1.559	0.282
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]AFQPF	1	753.4052	1.648	0.137
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]DLK[272]PAIVK[272]	3	439.294	2.155	0.009
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	AFQPF	1	749.3981	1.518	0.155
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]ALLAYAF	1	912.5312	1.571	0.199
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	n[145]FIPLK[272]PTVK[272]	3	492.3289	2.151	0.005
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	FCLESAWNVAK[272]	3	533.2711	2.194	0.063
67	IPI00025426	1	21.4	1.02	0.04	32	27	PZP Isoform 1 of Pregnancy zone protein precursor	EM[147]K	1	703.3808	1.121	0.072
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]SQCLEDHTWAPPFPICK[272]	3	784.7011	3.402	0.391
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	GYHLVGKK	3	441.275	3.332	0.239
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]ALLAFQESK[272]	2	647.8841	3.514	0.394
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	LIQEAPKPECEK	3	617.6671	3.296	0.242
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	ALLAFQESK	2	643.877	3.289	0.292
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]ESGM[147]TM[147]EELK[272]	2	737.854	2.628	0.222
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	SQCLEDHTWAPPFPICK	3	782.0297	3.017	0.275
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	YSLELK	2	516.8081	1.931	0.059
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]YSLELK[272]	2	520.8152	1.497	0.033
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]EVEGQILGTYVCIK[272]	2	943.5039	2.247	0.036
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	n[145]EHCPELPPVDNSIFVAK[272]	3	743.7144	3.012	0.12
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	M[147]CNDHYILK	3	493.57	2.031	0.092
68	IPI00025862	1	40.6	0.65	0.14	16	11	C4BPB Isoform 1 of C4b-binding protein beta chain precursor	LELK	2	391.7604	1.578	1
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	IFFPGVSEFGK	2	754.4191	3.579	0.424
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	n[145]IFFPGVSEFGK[272]	2	758.4262	3.604	0.367
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	n[145]VIVVSM[147]NYR	2	620.8457	2.606	0.271
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	n[145]FK[272]K[272]PQSLTK[272]	3	551.6897	1.9	0.181
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	FKKPQSLTK	3	546.3469	2.354	0.183
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	n[145]FYYFEHR	3	402.5335	2.391	0.166
69	IPI00025864	1	6.1	2.79	2.34	7	6	BCHE Cholinesterase precursor	SK[272]K[272]	2	395.7731	0.913	0.136
70	IPI00026199	1	5.3	0.93	0.07	11	4	GPX3 Glutathione peroxidase 3 precursor	FLVGPDGIPIM[147]R	2	735.9105	3.208	0.268
70	IPI00026199	1	5.3	0.93	0.07	11	4	GPX3 Glutathione peroxidase 3 precursor	n[145]FLVGPDGIPIMR	2	729.9166	3.195	0.304

70	IPI00026199	1	5.3	0.93	0.07	11	4	GPX3 Glutathione peroxidase 3 precursor	n[145]FLVGPDGIPIM[147]R	2	737.9141	3.486	0.204
70	IPI00026199	1	5.3	0.93	0.07	11	4	GPX3 Glutathione peroxidase 3 precursor	FLVGPDGIPIMR	2	727.9131	3.351	0.079
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	AGALNSNDAFVLK	2	800.4463	4.846	0.47
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EVQGFESATFLGYFK	2	1002.0173	5.108	0.473
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]AGALNSNDAFVLK[272]	2	804.4534	5.173	0.481
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLM[147]SLFGGK[272]	2	795.9313	4.063	0.497
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EVQGFESATFLGYFK[272]	2	1006.0244	5.073	0.498
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLMSLFGGK	3	522.9536	3.805	0.443
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLM[147]SLFGGK	3	528.2852	3.505	0.403
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	KGGVASGFK	3	424.2592	1.97	0.296
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	SEDCFILDHGK	3	530.5865	3.849	0.492
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLM[147]SLFGGK[272]	3	530.9566	3.807	0.518
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]SEDCFILDHGK[272]	3	533.2579	2.111	0.205
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	DSQEEEKTEALTSAK	3	696.027	3.513	0.226
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]DSQEEEK[272]TEALTSAK[272]	3	700.0341	3.967	0.262
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	HVVPNEVVVQR	2	708.4095	3.159	0.45
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLM[147]SLFGGK[272]PM[147]	3	612.3193	3.996	0.386
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	KGGVASGFK	2	635.8852	1.24	0.003
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]TASDFITK[272]	2	585.8341	2.781	0.341
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	TASDFITK	2	581.827	2.502	0.229
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLM[147]SLFGGKPM[147]	3	609.6479	3.553	0.391
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	AVEVLPK	2	518.3317	2.711	0.212
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLM[147]SLFGGK[272]PM	3	606.9877	3.488	0.35
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]TGAQELLR	2	516.3019	3.149	0.145
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]HVVPNEVVVQR	2	710.4131	1.985	0.351
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	YIETDPANR	2	609.8093	1.7	0.251
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLMSLFGGK[272]	3	525.625	3.039	0.34
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]AVEVLPK[272]	2	522.3388	1.54	0.03
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]SEDCFILDHGK[272]	2	799.3833	2.976	0.299
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EVQGFESATFLGYFK	3	668.3473	3.25	0.246
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	AQPVQVAEGSEPDGFWEALGGK	3	851.4309	3.428	0.129
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	VHVSEEGTEPEAM[147]LQVLGPK	3	816.0931	3.687	0.318
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	TGAQELLR	2	514.2984	2.657	0.031
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]AMAELAA	1	820.4356	1.383	0.222
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	LFACSNK	2	554.7858	2.193	0.018
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLM[147]SLFGGK	2	791.9242	3.639	0.401
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]YIETDPANR	2	611.8129	1.367	0.231
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	LFQVK	2	457.7948	1.766	0.025
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]SM[147]VVEHPEFLK[272]	3	540.6284	2.472	0.256
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]EPAHLMSLFGGK[272]PM[147]	3	606.9877	2.254	0.156
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]PALPAGTEDTAK[272]	2	729.9058	3.326	0.317
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]PNSM[147]VVEHPEFLK[272]	3	610.9936	2.625	0.242
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	VPFDAATLHTSTAMAAQHGM[147]DDDGTGQK	4	793.1224	3.66	0.266
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]PNSMVVEHPEFLK[272]	3	605.662	2.821	0.17
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	PALPAGTEDTAK	2	725.8987	2.493	0.196
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLMSLFGGKPM[147]	3	604.3163	3.355	0.116
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	GGVASGFK	2	501.7902	1.81	0.093
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	EPAHLM[147]SLFGGKPM	3	604.3163	2.476	0.058

71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	SMVVEHPEFLK	3	532.6253	2.562	0.012
71	IPI00026314	1	29.9	0.91	0.08	88	45	GSN Isoform 1 of Gelsolin precursor	n[145]VHVSEEGTEPEAM[147]LQVLGPK[272]	3	818.7645	2.157	0.137
72	IPI00026625	1	0.5	1.05	0.14	19	4	NUP155 Isoform 1 of Nuclear pore complex protein Nup155	LHELEEK	2	589.3324	2.338	0.133
72	IPI00026625	1	0.5	1.05	0.14	19	4	NUP155 Isoform 1 of Nuclear pore complex protein Nup155	n[145]LHELEEK[272]	2	593.3395	1.707	0.024
72	IPI00026625	1	0.5	1.05	0.14	19	4	NUP155 Isoform 1 of Nuclear pore complex protein Nup155	LHELEEK	3	393.224	2.958	0.079
72	IPI00026625	1	0.5	1.05	0.14	19	4	NUP155 Isoform 1 of Nuclear pore complex protein Nup155	n[145]LHELEEK[272]	3	395.8954	2.267	0.158
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]AAAAAAVSGSAAAEAK[272]	2	802.946	5.029	0.554
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]SEAACLAAGPGIR	2	703.3561	4.408	0.45
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]SVNNVVVR	2	515.8099	3.067	0.166
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	EEYSNLK	2	581.8088	2.617	0.227
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]LADDLYR	2	505.2754	2.765	0.117
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]AK[272]EQYAVVGH	3	463.9292	3.015	0.395
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]EEYSNLK[272]	2	585.8159	2.203	0.107
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]IYM[147]YGGK[272]	2	568.3067	1.928	0.123
73	IPI00027235	1	5.9	1.1	0.54	10	9	ATRN Isoform 1 of Attractin precursor	n[145]YQGNPLR	2	496.2757	2.08	0.044
74	IPI00027350	1	17.7	0.69	0.06	5	4	PRDX2 Peroxiredoxin-2	IGKPAPDFK	3	464.9503	2.537	0.391
74	IPI00027350	1	17.7	0.69	0.06	5	4	PRDX2 Peroxiredoxin-2	n[145]IGK[272]PAPDFK[272]	3	468.9574	2.617	0.293
74	IPI00027350	1	17.7	0.69	0.06	5	4	PRDX2 Peroxiredoxin-2	KEGGLGPLNIPLLADVTR	3	715.0914	2.393	0.073
74	IPI00027350	1	17.7	0.69	0.06	5	4	PRDX2 Peroxiredoxin-2	GLFIIDGK	2	571.8503	1.601	0.042
74	IPI00027350	1	17.7	0.69	0.06	5	4	PRDX2 Peroxiredoxin-2	n[145]GLFIIDGK[272]	2	575.8574	1.524	0.048
75	IPI00027462	1	13.2	1.57	0	2	2	S100A9 Protein S100-A9	NIETIINTFHQYSVK	3	696.381	4.608	0.53
75	IPI00027462	1	13.2	1.57	0	2	2	S100A9 Protein S100-A9	n[145]NIETIINTFHQYSVK[272]	3	699.0524	4.16	0.481
75	IPI00027462	1	13.2	1.57	0	2	2	S100A9 Protein S100-A9	SVK[272]	1	617.4104	0.88	0.378
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	VLVNYIFFK	2	711.9291	3.836	0.382
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	n[145]EENFYVDETTVVK[272]	2	930.9771	4.33	0.399
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	SETEIHQGFQHLHQLFAK	3	810.7608	5.103	0.446
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	n[145]SETEIHQGFQHLHQLFAK[272]	3	813.4322	5.445	0.402
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	SETEIHQGFQHLHQLFAK	4	608.3224	3.644	0.385
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	n[145]SETEIHQGFQHLHQLFAK[272]	4	610.326	4.728	0.444
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	ITQDAQLK	2	598.8535	1.913	0.033
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	n[145]ITQDAQLK[272]	2	602.8606	2.375	0.064
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	M[147]DPNAAYVNM[147]SNHHR	3	643.6212	2.547	0.318
76	IPI00027482	1	15.6	0.78	0.02	14	10	SERPINA6 Corticosteroid-binding globulin precursor	n[145]VLVNYIFFK[272]	2	715.9362	3.813	0.333
77	IPI00027843	1	4.8	0.84	0	3	3	PROZ Isoform 1 of Vitamin K-dependent protein Z precursor	DFAEHLLIPR	3	450.9225	3.097	0.225
77	IPI00027843	1	4.8	0.84	0	3	3	PROZ Isoform 1 of Vitamin K-dependent protein Z precursor	n[145]DFAEHLLIPR	3	452.2582	2.967	0.202
77	IPI00027843	1	4.8	0.84	0	3	3	PROZ Isoform 1 of Vitamin K-dependent protein Z precursor	n[145]DFCGGVIIR	2	585.3	1.859	0.195
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]LVDEDM[147]NSFK[272]	2	751.376	3.581	0.481
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]VTFELTYEELLK[272]R	3	643.7027	3.361	0.467
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	VTFELTYEELLKR	3	641.0313	3.598	0.291
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	KGHVSFKPSLDQQR	4	512.5445	3.019	0.322
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]DFLGFYVVDSHR	3	533.6076	2.7	0.28
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]GMTNINDGLLR	2	674.3622	3.257	0.333
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	KGHVSFKPSLDQQR	3	683.0569	2.863	0.293
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]GM[147]TNINDGLLR	2	682.3597	2.243	0.077
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]LVDEDMNSFK[272]	2	743.3785	3.097	0.378
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]GK[272]YEMYLK[272]	3	488.6146	2.905	0.185
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]EVSFDVELPK[272]	2	725.9053	2.241	0.118
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]FAHNVVTM[147]R	2	617.8278	1.85	0.022

78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	LVDEDM[147]NSFK	2	747.3689	1.899	0.204
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	ALDISIK	2	520.3292	1.666	0.038
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]DYIFGNYIER	2	717.3627	1.75	0.184
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	GKYEM[147]YLK	2	734.4051	2.308	0.144
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]GK[272]YEM[147]YLK[272]	2	740.4158	2.063	0.176
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	KGHVSFK	3	408.2522	2.715	0.109
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	n[145]GK[272]YEM[147]YLK[272]	3	493.9463	1.322	0.096
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	GKYEM[147]YLK	3	489.9392	1.246	0.034
78	IPI00028413	1	11.6	0.67	0.03	28	19	ITIH3 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H3 precu	KSFSGK	2	537.327	1.865	0.132
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	n[145]VSLATVDK[272]	2	560.8445	2.338	0.232
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	DDFLIYDR	2	598.801	2.826	0.12
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	n[145]DDFLIYDR	2	600.8045	2.289	0.136
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	VSLATVDK	2	556.8374	2.236	0.072
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	HLIFEK	2	533.8241	1.869	0.135
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	n[145]HLIFEK[272]	2	537.8312	1.572	0.087
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	IAYCEK	2	526.767	1.726	0.039
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	n[145]ESQDQSSLCK[272]	2	729.844	2.99	0.219
79	IPI00029061	1	10	0.75	0.04	9	9	SEPP1 Selenoprotein P precursor	FLIYDR	2	483.774	2.406	0.019
80	IPI00029168	1	7.7	0.45	0.12	4	3	LPA Apolipoprotein	n[145]EAQLLVIENEVCNHYK[272]	3	746.0497	5.711	0.54
80	IPI00029168	1	7.7	0.45	0.12	4	3	LPA Apolipoprotein	n[145]GTYSTTVTGR	2	593.8129	3.406	0.279
80	IPI00029168	1	7.7	0.45	0.12	4	3	LPA Apolipoprotein	n[145]GSFSTTVTGR	2	578.8076	3.01	0.239
80	IPI00029168	1	7.7	0.45	0.12	4	3	LPA Apolipoprotein	n[145]DIAILK[272]	2	480.8203	1.803	1
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]EALVPLVADHK[272]	3	493.964	3.413	0.408
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]LHK[272]PGVYTR	3	453.6098	2.553	0.314
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]SQFVQPICLPEPGSTFPAGHK[272]	3	858.7744	3.52	0.221
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]TLYSVFNPSDHDLVLIR	3	711.7202	4.508	0.324
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]SVFNPSDHDLVLIR	3	585.9885	2.817	0.238
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	LHKPGVYTR	3	450.9384	2.279	0.038
81	IPI00029193	1	11	0.98	0.28	7	7	HGFAC Hepatocyte growth factor activator precursor	n[145]ACQGDSGGPLACEK	2	856.3733	2.917	0.288
82	IPI00029260	1	5.3	0.62	0.07	4	4	CD14 Monocyte differentiation antigen CD14 precursor	LKELTLEDLK	3	541.3345	2.517	0.159
82	IPI00029260	1	5.3	0.62	0.07	4	4	CD14 Monocyte differentiation antigen CD14 precursor	n[145]LK[272]ELTLEDLK[272]	3	545.3416	2.996	0.127
82	IPI00029260	1	5.3	0.62	0.07	4	4	CD14 Monocyte differentiation antigen CD14 precursor	n[145]SLDLSHNSLR	3	429.2375	3.584	0.111
82	IPI00029260	1	5.3	0.62	0.07	4	4	CD14 Monocyte differentiation antigen CD14 precursor	n[145]SLDLSHNSLR	2	643.3527	3.035	0.071
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	CFEGFGIDGPAIAK	2	875.9364	4.479	0.526
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	EQVQSCGPPPELLNGNVK	2	1118.0667	5.864	0.538
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SIDVACHPGYALPK	2	898.963	5.021	0.501
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SPDVINGSPISQK	2	811.4491	4.574	0.513
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	TLTGGNVFEYGVK	2	832.954	4.725	0.525
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]AGEQVTYTCATYYK[272]	2	966.4597	5.364	0.439
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CYFPYLENGYNQNYGR	2	1095.9783	4.818	0.586
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]DTSCVNPPTVQNAYIVSR	2	1077.5256	5.388	0.467
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EIM[147]ENYNIALR	2	763.3937	3.642	0.438
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]HFGLSPDLPICK[272]	2	830.9433	4.081	0.506
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SIDVACHPGYALPK[272]	2	902.9701	5.03	0.477
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SPDVINGSPISQK[272]	2	815.4562	4.398	0.476
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SSQESYAHGTK[272]	2	741.875	4.427	0.49
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TGESVEFVCK[272]	2	716.8564	4.015	0.487
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WSSPPQCEGLPCK[272]	2	906.4132	3.606	0.548

83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	CTLKPCDYPDIK	3	636.6465	3.275	0.443
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	GDAVCTESGWRPLPSCEEK	3	812.6987	5.132	0.511
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IEGDEEM[147]HCSDDGFWSK	3	776.3186	3.737	0.464
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SIDVACHPGYALPK	3	599.6444	2.462	0.159
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SSNLIILEEHLK	3	559.3296	1.758	0.019
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SSQESYAHGTK	3	492.2477	4.097	0.404
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	TKEEYGHSEVVEYYCNPR	3	843.7228	4.68	0.549
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CTLK[272]PCDYPDIK[272]	3	640.6536	3.659	0.362
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CYFPYLENGYNQNYGR	3	730.988	3.345	0.335
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]DTSCVNPPTVQNAYIVSR	3	718.6862	3.799	0.441
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EYHFGQAVR	3	417.5515	3.862	0.284
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]GDAVCTESGWRPLPSCEEK[272]	3	815.3701	5.03	0.527
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]HGGLYHENMR	3	453.2232	3.656	0.376
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAMEPDREYHFGQAVR	3	779.3906	4.058	0.286
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SIDVACHPGYALPK[272]	3	602.3158	4.48	0.433
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SSIDIENGFISESQYTYALK[272]	3	851.7703	4.726	0.393
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SSNLIILEEHLK[272]	3	562.001	4.649	0.328
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SSQESYAHGTK[272]	3	494.9191	3.761	0.441
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SVQCYHFGLSPDLPICK[272]	3	763.0377	5.332	0.417
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TK[272]EEYGHSEVVEYYCNPR	3	846.3942	4.189	0.417
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	HGGLYHENMR	4	339.1675	3.917	0.431
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAMEPDREYHFGQAVR	4	584.7948	3.125	0.213
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]RPCGHPGDTPFGTFTLTGGNVFEYGVK[272]	4	797.8939	4.402	0.35
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SPPEISHGVVAHMSDSYQYGEEVTYK[272]	4	800.3878	4.281	0.357
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	RPYFPVAVGK	3	471.9504	3.43	0.212
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EFDHNSNIR	3	425.8781	2.969	0.345
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IEGDEEMHCSDDGFWSK[272]	3	773.6584	3.818	0.315
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAM[147]EPDREYHFGQAVR	3	784.7223	3.334	0.397
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]RPYFPVAVGK[272]	3	474.6218	3.399	0.296
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TK[272]EEYGHSEVVEYYCNPR	4	635.0475	3.676	0.419
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	CTLKPCDYPDIK	2	954.4662	3.475	0.416
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CFEGFGIDGPAIAK[272]	2	879.9435	3.483	0.466
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CLPVTAPENGK[272]	2	731.8855	3.271	0.306
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CNMGYEYSER	2	721.287	2.997	0.465
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SSNLIILEEHLK[272]	2	842.4978	4.21	0.306
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TDCLSLPSFENAIPMGEK[272]	2	1143.5566	3.515	0.49
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAM[147]EPDREYHFGQAVR	4	588.7936	3.161	0.279
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SSNLIILEEHLK	2	838.4907	4.048	0.291
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SSQESYAHGTK	2	737.8679	3.363	0.452
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CTLK[272]PCDYPDIK[272]	2	960.4768	3.3	0.341
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EIMENYNIALR	2	755.3962	3.934	0.346
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]LSYTCEGGFR	2	661.8031	1.437	0.133
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SVQCYHFGLSPDLPICK	3	760.3663	4.251	0.388
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]AVYTCNEGYQLLGEINYR	3	766.0335	3.315	0.237
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CLHPCVISR	3	421.8699	2.184	0.298
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IIYK[272]ENERFQYK[272]	3	688.3932	3.792	0.257
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]NK[272]K[272]EFDHNSNIR	3	645.3571	2.512	0.353
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EYHFGQAVR	2	625.8236	2.745	0.335

83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SCDIPVFMNAR	2	721.8392	3.047	0.264
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SCDNPYIPNGDYSPLR	2	1000.9518	3.455	0.284
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	CLPVTAPENGK	2	727.8784	3.624	0.237
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	CVEISCK	2	577.2655	2.689	0.255
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	LEYPTCAK	2	625.8173	2.06	0.264
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CNM[147]GYEYSER	2	729.2844	2.525	0.376
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TDCLSLPSFENAIPM[147]GEK[272]	2	1151.554	2.275	0.123
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	RPYFPVAVGK	2	707.4219	3.215	0.194
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAM[147]EPDR	2	632.8198	2.99	0.259
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SCDIPVFM[147]NAR	2	729.8366	2.618	0.325
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SLGNVIM[147]VCR	2	647.3274	3.172	0.214
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]CVEISCK[272]	2	581.2726	2.338	0.314
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]LEYPTCAK[272]	2	629.8244	2.457	0.162
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SLGNVIM[147]VCR	2	649.331	2.73	0.346
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]NK[272]K[272]EFDHNSNIR	5	387.6172	2.197	0.121
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IEGDEEM[147]HCSDDGFWSK[272]	3	778.99	3.212	0.275
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IIYK[272]ENERFQYK[272]	4	516.5467	2.44	0.229
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]NGFYPATR	2	535.281	2.199	0.248
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TGDEITYQCR	2	688.3088	2.529	0.358
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	TGESVEFVCK	2	712.8493	2.499	0.303
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IIYKENERFQYK	3	684.3861	3.334	0.237
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EFDHNSNIR	2	638.3136	2.518	0.268
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TGESVEFVCK[272]R	3	530.2737	3.5	0.093
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	TGESVEFVCKR	3	527.6023	3.169	0.111
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]RPYFPVAVGK[272]	2	711.429	3.304	0.147
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IDVHLVPDR	2	604.3494	2.726	0.218
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TDCLSLPSFENAIPM[147]GEK[272]	3	768.0384	2.566	0.256
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]NDFTWFK[272]	2	623.329	2.202	0.137
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IEGDEEM[147]HCSDDGFWSK[272]	3	777.6543	2.55	0.338
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IIYKENER	2	672.8854	3.198	0.23
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]ECELPK[272]	2	526.7716	2.214	0.141
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	NDFTWFK	2	619.3219	2.296	0.114
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SPPEISHGVVAHM[147]SDSYQYGEEVTYK[272]	4	804.3865	2.679	0.2
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	ECELPK	2	522.7645	1.818	0.199
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EDCNELPPRR	3	473.5594	2.608	0.296
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SPPEISHGVVAH	3	458.5813	3.436	0.244
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IIYK[272]ENER	2	676.8925	2.726	0.237
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]HGGLYHENMR	2	679.3312	2.793	0.417
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WSHPPSCIK	2	692.853	3.003	0.25
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IVSSAMEPDR	2	624.8224	1.829	0.146
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TTCWDGK[272]	2	572.7721	1.69	0.116
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	KDVYK	3	358.222	1.863	0.172
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]IIYK[272]ENER	3	451.5974	2.69	0.09
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WSSPPQCEGLPCK	2	904.4097	3.321	0.404
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WQSIPLCVEK	2	766.9079	3.718	0.198
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]TLTGGNVFEYGVK[272]	2	836.9611	3.283	0.428
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	NKKEFDHNSNIR	3	641.35	2.77	0.098
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IIYKENER	3	448.926	2.332	0.112

83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]RPCGHPGDTPFGTF	3	560.2577	2.769	0.25
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EDCNELPPR	2	631.7849	2.621	0.229
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	KFVQGK	2	563.8585	2.067	0.123
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SIDVACHPGY	2	626.2846	2.861	0.397
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	KEFDHNSNIR	3	513.9391	2.22	0.031
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WQSIPLCVEK[272]	3	512.9434	1.936	0.09
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SPPQCEGLPCK	2	765.8504	2.897	0.241
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	FSCKPGFTIVGPN	2	846.9337	2.992	0.321
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	AKYQCK	2	603.828	1.817	0.123
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]GFYPATR	2	478.2595	2.28	0.254
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]WSHPPSCIK	3	462.2377	2.295	0.165
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]SCDIPVFM[147]NAR	3	486.8935	2.162	0.017
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	GLSPDLPICK	2	684.8726	2.607	0.252
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]KGEWVALNPLR	3	522.9794	2.812	0.033
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]AK[272]YQCK[272]	3	406.8948	2.253	0.025
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	GDAVCTESGWRPLPSCEEK[272]	3	814.0344	2.489	0.171
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]KGEWVALNPLRK	3	612.3761	3.069	0.094
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	HFGLSPDLPICK	3	551.6266	2.987	0.107
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	IVSSAMEPDR	2	622.8188	1.521	0.053
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	NKKEFDHNSNIR	4	481.2644	2.107	0.206
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]DGWSAQPTCIK	2	768.377	2.816	0.122
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]FSCK[272]PGFTIVGPN	2	850.9408	2.711	0.275
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	K[272]GEWVALNPLR	2	783.9655	2.459	0.008
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	RPCGHPGDTPFGTF	3	558.922	2.027	0.133
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]AYIVSR	2	426.7566	2.161	0.139
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]LEYPTCA	1	986.4444	1.966	0.147
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]DVHLVPDR	2	547.8074	2.118	0.132
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	n[145]EDCNELPPRR	2	709.8355	2.092	0.106
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	DTSCVNPPTVQ	2	673.8076	1.634	0.169
83	IPI00029739	1	52.1	0.7	0.16	204	##	CFH Isoform 1 of Complement factor H precursor	SPLR	1	612.3828	1.332	0.125
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LCQDLGPGAFR	2	683.84	3.6	0.486
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LGNQEPGGQTALK[272]	2	800.9485	4.823	0.358
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	WFLLEQPEIQVAHFPFK	3	803.7755	4.301	0.405
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]GDK[272]LFGPDLK[272]	3	507.9716	3.87	0.324
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]GFPIK[272]EDFLEQSEQLFGAK[272]	3	872.4742	4.638	0.5
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]WFLLEQPEIQVAHFPFK	3	805.1112	4.931	0.423
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	LGNQEPGGQTALK	2	796.9414	4.529	0.366
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	ELKEQQDSPGNK	3	598.3233	3.95	0.386
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	GDKLFGPDLK	3	503.9645	3.407	0.317
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LQQVLHAGSGPCLPHLLSR	4	554.8033	3.644	0.354
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]DSFHLDEQFTVPVEMMQAR	3	808.7197	3.473	0.245
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]ELK[272]EQQDSPGNK[272]	3	602.3304	3.886	0.322
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]DFLQSLK[272]	2	569.8392	2.613	0.155
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LQQVLHAGSGPCLPH	3	582.9682	2.81	0.399
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	DFLQSLK	2	565.8321	1.699	0.012
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LFGPDLK[272]	2	539.331	2.089	0.004
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]NK[272]FDPSLTQR	2	747.4194	2.535	0.309
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	NKFDPSLTQR	2	743.4123	2.556	0.293

84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]FDPSLTQR	2	554.2994	2.014	0.076
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	NQEQVSPLTLLK	2	825.4829	3.878	0.19
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]NQEQVSPLTLLK[272]	2	829.49	4.108	0.243
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	LQQVLHAGSGPCLPH	2	871.9451	3.83	0.38
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]MYLQK[272]	2	485.7854	1.618	0.073
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LQQVLHAGSGPCLPH	2	873.9486	4.281	0.382
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	EQQDSPGNKDFLQSLK	3	752.0658	2.39	0.172
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	NPNPSAPR	2	496.7673	1.251	0.16
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	GFPIKEDFLEQSEQLFGAK	3	868.4671	2.34	0.187
84	IPI00029863	1	33	0.77	0.17	53	26	SERPINF2 SERPINF2 protein	n[145]LQQVLH	2	441.2699	1.88	0.036
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	SLTSCLDSK	2	640.3229	1.671	0.188
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	WIYHLTEGSTDLR	2	865.9467	3.849	0.377
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]SLTSCLDSK[272]	2	644.33	1.95	0.1
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]AFLLTPR	2	481.3012	2.34	0.184
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]DGLCVPR	2	475.2394	1.709	0.143
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]FLLYNR	1	969.5639	2.045	0.182
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]FLLYNR	2	485.2856	1.891	0.066
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	DGLCVPR	2	473.2359	2.149	0.155
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	CVEEFK	2	540.7645	2.104	0.163
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	SPHPPEK	2	536.301	1.833	0.201
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]CVEEFK[272]	2	544.7716	2.065	0.183
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	TEGRPDMK	2	607.3216	1.594	0.181
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	FLLYNR	1	965.5568	2.016	0.028
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]WIYHLTEGSTDLR	3	578.9692	2.567	0.033
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	n[145]TEGRPDMK[272]	2	611.3287	1.506	0.143
85	IPI00030739	1	54.3	0.85	0.06	29	16	APOM Apolipoprotein M	TEGRPDMK	3	405.2168	1.696	0.035
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	ADGESCSASM[147]M[147]YQEGK	2	1026.4125	5.048	0.689
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	DDLYVSDAFHK	2	795.4016	4.085	0.401
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	EQLQDMGLVDLFSPEK	2	1065.051	4.496	0.448
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	EQLQDM[147]GLVDLFSPEK	2	1073.0485	5.292	0.473
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	FATTFYQHLADSK	2	904.9702	5.586	0.442
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	NDNDNIFLSPLSISTAFAM[147]TK	2	1298.1598	5.61	0.604
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	TSDQIHFFFAK	2	810.9303	4.615	0.494
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	VAEGTQVLELPFK	2	855.9931	4.639	0.522
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]ADGESCSASM[147]M[147]YQEGK[272]	2	1030.4196	3.79	0.606
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]AFLEVNEEGSEAAASTAVVIAGR	2	1218.1304	6.161	0.519
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EQLQDMGLVDLFSPEK[272]	2	1069.0581	4.387	0.478
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EQLQDM[147]GLVDLFSPEK[272]	2	1077.0556	4.587	0.437
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FATTFYQHLADSK[272]	2	908.9773	4.136	0.498
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]NDNDNIFLSPLSISTAFAM[147]TK[272]	2	1302.1669	6.123	0.525
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]TSDQIHFFFAK[272]	2	814.9374	4.303	0.452
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]VAEGTQVLELPFK[272]	2	860.0002	4.917	0.491
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	ADGESCSASM[147]M[147]YQEGK	3	684.6108	3.675	0.422
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	DDLYVSDAFHK	3	530.6035	3.977	0.407
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	FATTFYQHLADSK	3	603.6492	3.653	0.463
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	TSDQIHFFFAK	3	540.956	3.174	0.359
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DDLYVSDAFHK[272]	3	533.2749	3.125	0.302
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FATTFYQHLADSK[272]	3	606.3206	4.512	0.378

86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FRIEDGFSLK[272]	3	500.6202	4.566	0.38
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LPGIVAEGRDDLYVSDAFHK[272]	4	623.3363	4.673	0.422
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]LPGIVAEGRDDLYVSDAFHK[272]	4	713.1435	2.854	0.028
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]LPGIVAEGRDDLYVSDAFHK[272]	5	570.7163	3.9	0.385
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	GDDITM[147]VLILPKPEK	3	702.4062	2.975	0.345
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	KATEDEGSEQK	3	547.953	3.234	0.319
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	SKLPGIVAEGR	3	469.6208	3.94	0.374
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]NDNDNIFLSPLSISTAFAM[147]TK[272]	3	868.447	2.584	0.132
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]LPGIVAEGR	3	472.2922	4.034	0.317
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]TSDQIHFFFAK[272]	3	543.6274	2.621	0.225
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DDLYVSDAFHK[272]	2	799.4087	3.921	0.377
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EVPLNTIIFMGR	2	767.4326	2.112	0.039
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	FRIEDGFSLK	3	497.9488	3.969	0.202
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	EVPLNTIIFMGR	2	765.4291	3.288	0.34
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EVPLNTIIFM[147]GR	2	775.4301	2.835	0.306
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]FSPENTRK[272]	3	542.6494	2.838	0.308
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	GDDITMVLILPKPEK	3	697.0745	3.069	0.282
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]GDDITM[147]VLILPK[272]PEK[272]	3	706.4133	2.642	0.28
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	ATEDEGSEQK	2	687.3308	2.896	0.377
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DIPMNPMCIYR	2	771.8565	3.245	0.246
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]GDDITMVLILPK[272]PEK[272]	2	1051.1188	4.079	0.333
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]GDDITMVLILPK[272]PEK[272]	3	701.0816	2.82	0.231
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]IEDGFSLK[272]	1	1196.6766	2.55	0.155
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LQPLDFK[272]	2	574.8496	2.385	0.263
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	IEDGFSLK	1	1188.6624	2.533	0.124
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]ELFYK[272]	2	494.2913	2.138	0.245
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DIPM[147]NPM[147]CIYR	2	787.8514	1.997	0.196
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	LQPLDFKENAEQSR	3	652.3497	2.527	0.296
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	KELFYK	2	624.3792	2.463	0.157
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	LQPLDFK	2	570.8425	1.888	0.001
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]ATEDEGSEQK[272]	2	691.3379	2.563	0.266
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DIPMNPM[147]CIYR	2	779.854	3.33	0.249
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FSPENTR	2	497.7574	2.225	0.202
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LQPLDFK[272]ENAEQSR	3	655.0211	2.483	0.275
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	IEDGFSLK	2	594.8348	2.38	0.214
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	SKLPGIVAEGR	2	703.9276	3.29	0.386
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	ELFYK	2	490.2842	2.057	0.22
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LQPLDFK[272]ENAEQSR	2	982.028	3.013	0.439
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LPGIVAEGR	2	528.3201	1.918	0.164
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	EVPLNTIIFM[147]GR	2	773.4265	2.727	0.081
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]LPGIVAEGRDDLYVSDAFHK[272]	3	950.5223	3.35	0.234
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]LPGIVAEGR	2	707.9347	3.002	0.327
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	FDTISEK	2	560.3059	2.545	0.082
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]IEDGFSLK[272]	2	598.8419	1.879	0.247
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EVPLNTIIFM[147]GR	3	517.2891	3.049	0.102
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	LQPLDFKENAEQSR	2	978.0209	2.322	0.408
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]VANPCVK[272]	2	532.7954	1.904	0.15
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]DIPM[147]NPMCIYR	2	779.854	2.327	0.297

86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	DIPM[147]NPMCIYR	2	777.8504	2.326	0.363
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FRIEDGFSLK[272]	2	750.4267	3.163	0.217
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	FSPENTR	2	495.7538	1.747	0.143
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	KELFYK	3	416.5886	1.95	0.064
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]FATTFYQH	2	579.7887	3.023	0.328
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]VWELSK[272]	2	525.3153	1.647	0.042
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	SKLPGIVAEGRDDLYVSDAFHK	4	710.1382	2.704	0.136
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]TSDQIHFFF	2	643.3203	2.113	0.38
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]TSDQIHFFFA	2	678.8389	2.73	0.412
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]FSPENTR	3	451.917	1.967	0.112
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	KATEDEGSEQK	2	821.4258	1.385	0.211
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]YQHLADSK[272]	2	625.3426	2.422	0.254
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]SK[272]FSPENTRK[272]	4	407.2389	1.835	0.001
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]EVPLNTIIFMGR	3	511.9575	1.933	0.035
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]YQHLADSK[272]	3	417.2308	2.906	0.067
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	ATTFYQHLADSK	3	554.6264	2.574	0.145
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]LYVSDAFHK[272]	3	456.5902	2.185	0.148
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	YQHLADSK	2	621.3355	1.942	0.138
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	n[145]ATTFYQHLADSK[272]	3	557.2978	2.985	0.043
86	IPI00032179	1	52.3	0.77	0.08	80	89	SERPINC1 Antithrombin III variant	NDNDNIFLSPLSISTAFAM[147]TK	3	865.7756	1.6	0.062
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	ALQDQLVLVAAK	2	774.9773	4.754	0.463
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	QPFVQGLALYTPVVLPR	2	1019.5961	5.403	0.576
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	SLDFTELDVAAEK	2	859.454	4.49	0.461
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	VQGLALYTPVVLPR	2	833.5062	3.779	0.521
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]ALQDQLVLVAAK[272]	2	778.9844	4.453	0.369
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]DPTFIPAPIQAK[272]	2	793.4633	3.234	0.307
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]FVQGLALYTPVVLPR	2	909.0439	4.119	0.469
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]SLDFTELDVAAEK[272]	2	863.4611	4.457	0.393
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	AVYDQSATALHFLGR	3	596.9846	4.672	0.414
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	QPFVQGLALYTPVVLPR	3	680.0665	3.196	0.22
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]AVYDQSATALHFLGR	3	598.3203	4.941	0.518
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]LFAVYDQSATALHFLGR	3	685.0378	5.518	0.464
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]QPFVQGLALYTPVVLPR	3	681.4022	4	0.344
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	DPTFIPAPIQAK	2	789.4562	2.597	0.387
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	ALQDQLVLVAAK	3	516.9873	2.242	0.256
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]LDTEDK[272]LR	3	426.5816	2.457	0.263
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]QPFVQGLALYTPVVLPR	2	1021.5996	3.823	0.404
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	LDTEDKLR	3	423.9102	2.691	0.215
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	LDTEDK	2	500.7692	2.069	0.116
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	VANPLSTA	1	912.515	1.727	0.121
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]LDTEDK[272]	2	504.7763	2.325	0.041
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]VANPLSTA	1	916.5221	1.397	0.211
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]LDTEDK[272]LR	2	639.3688	2.137	0.233
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]ALYTPVVLPR	2	636.8935	2.897	0.311
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]ALQDQLVLVAAK[272]	3	519.6587	1.922	0.141
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	ALYTPVVLPR	2	634.8899	3.035	0.319
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	LQAILGVPWK[272]	2	704.9435	2.652	0.27
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	TSPVDEK	2	528.2902	1.463	0.014

87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	YTPVVLPR	2	542.8293	2.04	0.212
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	YASDLDK	2	546.2902	1.912	0.083
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	VYIHPF	2	458.258	1.307	0.197
87	IPI00032220	1	24.1	0.89	0.06	53	29	AGT Angiotensinogen precursor	n[145]YASDLDK[272]	2	550.2973	1.745	0.102
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	ALVEGVDQLFTDYQIK	2	1060.0754	5.533	0.543
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	DINYVNPVIK	2	727.922	4.324	0.374
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	FSYSSGHVHLSSENK	2	979.9896	5.173	0.507
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VSITSITVENVFVK	2	908.5326	5.053	0.481
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]AFTECCVVASQLR	2	831.8833	3.708	0.459
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]ALVEGVDQLFTDYQIK[272]	2	1064.0825	3.892	0.488
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSYSSGHVHLSSENK[272]	2	983.9967	5.712	0.522
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VSITSITVENVFVK[272]	2	912.5397	5.258	0.542
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	ALLVGEHLNIIVTPK	3	633.0593	3.667	0.379
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	EESSSGSSHAVM[147]DISLPTGISANEEDLK	3	1062.8418	4.756	0.428
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	FSYSSGHVHLSSENK	3	653.6622	3.98	0.416
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	KAFDICPLVK	3	533.9753	2.95	0.31
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSDASYQSINIPVTQNM[147]VPSSR	3	867.7627	4.376	0.401
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSYSSGHVHLSSENK[272]	3	656.3336	4.426	0.399
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]TGEAVAEK[272]DSEITFIK[272]	3	724.0704	5.172	0.437
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSYSSGHVHLSSENK[272]	4	492.502	4.388	0.446
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	IVACASYKPSR	3	507.6074	2.805	0.172
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	TGEAVAEKDSEITFIK	3	720.0633	4.387	0.382
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VSITSITVENVFVK	3	606.0242	3.254	0.293
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]ALLVGEHLNIIVTPK[272]	3	635.7307	3.613	0.404
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IDTQDIEASHYR	3	531.2659	3.668	0.261
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IVACASYK[272]PSR	3	510.2788	3.3	0.322
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VSITSITVENVFVK[272]	3	608.6956	3.251	0.324
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	MPITYDNGFLFIHTDKPVYTPDQSVK	4	862.4509	3.582	0.298
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	M[147]PITYDNGFLFIHTDKPVYTPDQSVK	4	866.4497	4.336	0.382
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	KAFDICPLVK	2	800.4594	3.411	0.352
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]DINYVNPVIK[272]	2	731.9291	3.688	0.345
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FQNSAILTIQPK[272]	2	824.4873	3.566	0.367
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IVACASYK[272]PSR	2	764.9146	3.574	0.365
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]TDAPDLPEENQAR	2	800.3902	3.74	0.229
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	KVTCTNAELVK	3	557.981	3.181	0.224
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]QLPGGQNPVSYVYLEVVSK[272]	3	789.1051	3.245	0.17
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VYSLNDDLK[272]PAK[272]R	4	488.5389	2.033	0.253
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]ATLLDIYK[272]	2	612.8758	1.346	0.147
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]DINYVNPVIK[272]	3	488.2885	3.227	0.203
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	AFDICPLVK	2	666.3644	3.211	0.211
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]AFDICPLVK[272]	2	670.3715	3.407	0.201
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]DSEITFIK[272]	2	620.855	3.192	0.288
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]ENSQYQPIK[272]	2	697.8796	3.238	0.281
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FLK[272]PGIPYPIK[272]	3	569.0308	3.114	0.394
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YK[272]HSVVK[272]K[272]	3	522.3389	2.756	0.286
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	KIEEIAAK	2	661.4138	2.958	0.288
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	ITHYNYLILSK	2	822.9773	4.871	0.311
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VCEGAACK	2	576.7553	2.523	0.295

88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]GTVYNYR	2	508.7677	2.892	0.224
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	GTVYNYR	2	506.7642	2.475	0.266
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]GYGNSDYK[272]	2	596.2979	2.078	0.079
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	ATLLDIYK	2	608.8687	2.784	0.179
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	GEQIQLK	2	548.3297	2.459	0.174
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]GIYGTISR	2	505.7912	2.832	0.182
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]EGM[147]LSIM[147]SYR	2	681.8293	2.255	0.313
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YNFSFR	2	489.2517	2.389	0.207
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]GEQIQLK[272]	2	552.3368	2.358	0.08
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]LK[272]EGM[147]LSIM[147]SYR	3	583.3157	3.09	0.236
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	DINYVNPVIK	3	485.6171	2.613	0.21
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VYSLNDDLK[272]PAK[272]R	3	651.0494	2.512	0.206
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]NFEITIK[272]	2	576.847	2.216	0.244
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	YKHSVVKK	3	516.9961	2.118	0.263
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]GNPPIYR	2	480.7728	2.386	0.127
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VTCTNAELVK	2	702.3729	2.869	0.139
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]LQGTLPVEAR	2	614.3625	3.227	0.079
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VYSLNDDLKPAK	3	595.0086	1.822	0.249
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	YVLSPYK	2	575.337	2.228	0.142
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]SYPDK[272]K[272]	2	585.3482	1.962	0.18
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VFQFIEK[272]	2	599.8574	2.681	0.167
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VCEGAACK[272]	2	580.7624	2.02	0.13
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IDTALIK[272]	2	531.3441	2.494	0.105
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VFQFIEK	2	595.8503	2.574	0.142
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]EALQIK[272]	2	495.3153	2.408	0.032
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	EALQIK	2	491.3082	2.489	0.037
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	GYGNSDYK	2	592.2908	2.083	0.234
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VVPEGVK[272]	2	508.3232	1.748	0.016
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]EDLK[272]DDQK[272]	3	474.9316	2.416	0.142
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]EGM[147]LSIMSYR	2	673.8319	2.334	0.042
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	VVPEGVK	2	504.3161	1.765	0.034
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]TSTSEEVCSFYLK[272]	2	914.441	3.025	0.369
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]DSSVPNTGTAR	2	624.8187	2.276	0.158
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YK[272]EDFSTTGTAYFEVK[272]	3	773.4021	2.925	0.188
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YFYNK[272]	2	511.7811	1.779	0.027
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	TSTSEEVCSFYLK	2	910.4339	1.744	0.106
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]WLSEEQR	2	546.2837	2.128	0.013
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	IPLDLVPK	2	587.8816	1.673	0.005
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	FLKPGIPYPIK	3	565.0237	3.004	0.194
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	ITHYNYLILSK	3	548.9873	2.73	0.065
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	EDLKDDQK	2	705.8831	2.102	0.151
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	FSYSSGHVHLSSENK	4	490.4985	2.317	0.042
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	LKEGM[147]LSIM[147]SYR	3	580.6443	1.712	0.083
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YVLSPYK[272]	2	579.3441	1.562	0.018
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSYSSGH	2	464.7177	1.906	0.312
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	EDLKDDQK	3	470.9245	1.956	0.098
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]FSDASYQSINIPVTQNMVPSSR	3	862.4311	2.393	0.021
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]VYSLNDDLK[272]PAK[272]	3	599.0157	1.774	0.056

88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]SIPSSDFLCVR	2	707.353	2.609	0.147
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]EFPYR	2	428.2277	1.337	0
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	SYPDKK	3	386.5608	1.257	0.027
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]DGHVILQLN	2	576.8283	2.078	0.168
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]ALSLGDK[272]	2	496.305	2.226	0.066
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IPLDLVPK[272]	2	591.8887	1.298	0.009
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YSVVR	2	384.2303	1.803	0.049
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]YSVVR	1	767.4533	1.967	0.106
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	n[145]IPYSVVR	2	489.2987	1.877	0.088
88	IPI00032291	1	33.1	0.71	0.07	151	98	C5 Complement C5 precursor	SVK[272]	1	617.4104	0.88	0.378
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]SPVTLLAAVMSLPEEHNK[272]	3	742.0798	4.696	0.459
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]VQLYDLGLQIHK[272]	3	572.3413	4.674	0.374
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	SPVTLLAAVMSLPEEHNK	3	739.4084	4.141	0.318
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	VQLYDLGLQIHK	3	569.6699	3.157	0.359
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	LAEGFPLPLLK	2	739.4607	3.824	0.352
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]LAEGFPLPLLK[272]	2	743.4678	3.613	0.305
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	ITGFLKPGK	2	690.94	3.025	0.281
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]ITGFLK[272]PGK[272]	2	696.9506	2.978	0.307
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	ITGFLKPGK	3	460.9624	1.994	0.087
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]VK[272]VELK[272]	3	383.264	1.608	0.102
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	n[145]ANPGLVAR	2	471.2861	2.432	0.139
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	GDLR	1	600.3464	1.375	0.017
89	IPI00032311	1	14.8	0.65	0.39	13	11	LBP Lipopolysaccharide-binding protein precursor	SVK[272]	1	617.4104	0.88	0.378
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	AATGECTATVGK	2	717.8576	3.932	0.488
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	DIPTNSPELEETLTHTITK	2	1210.1394	6.156	0.619
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	IASFSQNCDIYPGK	2	934.9553	4.661	0.541
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	TVGSDTFYSFK	2	766.3932	3.797	0.308
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]AATGECTATVGK[272]	2	721.8647	4.023	0.39
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]AVDAALK[272]K[272]	2	624.4061	3.265	0.429
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]DIPTNSPELEETLTHTITK[272]	2	1214.1465	6.661	0.621
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]	2	938.9624	4.537	0.539
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]TVGSDTFYSFK[272]	2	770.4003	3.883	0.422
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]YNSQNQSNNQFVLYR	2	1009.9937	6.31	0.461
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	DIPTNSPELEETLTHTITK	3	807.0954	3.436	0.161
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	FKLDDDLEHQGGHVLDHGHK	3	906.4684	6.016	0.538
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	GHGLGHGHEQQHGLGHGHK	3	754.0496	5.892	0.517
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	KYFIDFVAR	3	480.2784	4.364	0.354
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	KYNSQNQSNNQFVLYR	3	761.7258	4.334	0.45
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]DIPTNSPELEETLTHTITK[272]	3	809.7668	2.435	0.122
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]GHGLGHGHEQQHGLGHGHK[272]	3	756.721	6.481	0.565
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	FKLDDDLEHQGGHVLDHGHK	4	680.1032	4.229	0.474
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	GHGLGHGHEQQHGLGHGHK	4	565.7891	5.789	0.571
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]GHGLGHGHEQQHGLGHGHK[272]	4	567.7926	3.962	0.304
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]GHGLGHGHEQQHGLGHGHK[272]	5	454.4355	3.332	0.322
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	AVDAALKK	3	412.5994	2.374	0.342
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]AVDAALK[272]K[272]	3	416.6065	2.264	0.345
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]DFVQPPTK[272]	3	978.5016	4.255	0.387
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]VQVVAGK[272]K[272]	3	420.9503	2.485	0.327

90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]DFVQPPTK[272]	4	734.128	3.901	0.397
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]VQVVAGK[272]K[272]	2	630.9219	2.949	0.454
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	TVGSDTFYSFKYEIK	3	735.7246	3.593	0.356
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	VQVVAGKK	3	416.9432	2.217	0.31
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]YNSQNQSNNQFVLYR	3	673.6649	3.213	0.311
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	TWQDCEYK	2	699.8127	2.887	0.436
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]EGDCPVQSGK[272]	2	677.3227	2.709	0.264
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]ENFLFLTPDCK[272]	2	830.9195	3.004	0.385
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	EGDCPVQSGK	2	673.3156	2.229	0.245
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	KYFIDFVAR	2	719.9139	3.42	0.289
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	VQVVAGKK	2	624.9112	2.668	0.303
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]YFIDFVAR	2	587.8225	2.207	0.392
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	DFVQPPTK	2	606.3428	2.294	0.202
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]SVSEINPTTQM[147]K[272]	2	819.9342	3.043	0.333
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	VQVVAGK	2	490.8162	3.032	0.186
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]TWQDCEYK[272]	2	703.8198	2.659	0.315
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]TVGSDTFYSFK[272]YEIK[272]	3	739.7317	3.68	0.229
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]VQVVAGK[272]	2	494.8233	2.593	0.178
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]AVDAALK[272]	2	488.3075	2.198	0.188
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]ICVGCPR	2	492.2244	1.902	0.206
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	AVDAALK	2	484.3004	2.449	0.188
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]DFVQPPTK[272]	2	610.3499	2.114	0.137
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]QVVAGLNFR	2	574.3389	2.605	0.145
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	ITEATK	2	471.7846	1.741	0.035
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]ITEATK[272]	2	475.7917	2.142	0.073
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	AVDAALKK	2	618.3954	1.972	0.143
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]ETTCSK[272]	2	501.7456	1.891	0.168
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	IDNVKK	2	568.8612	2.026	0.076
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	IDNVK	2	434.7662	1.944	0.066
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	YEIKEGDCPVQSGK	3	673.6728	2.449	0.181
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]SDTFYSFK[272]	2	641.8316	3.152	0.331
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]FIDFVAR	2	506.2908	2.415	0.089
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	TVGSDTFYSFK	3	511.2646	2.527	0.079
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]CQPLGM[147]ISLM[147]K[272]	2	793.9045	3.027	0.286
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]IYPTVNCQPLGM[147]	2	771.3678	2.928	0.33
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	CQPLGM[147]ISLM[147]K	2	789.8974	2.869	0.234
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]IYPTVNCQPLGM	2	763.3703	2.776	0.138
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	IYPTVN	1	846.472	1.7	0.123
90	IPI00032328	1	38.2	0.82	0.03	9	52	KNG1 Isoform HMW of Kininogen-1 precursor	n[145]RPPGFSPF	2	524.7885	1.56	0.098
91	IPI00044369	1	3.1	1.08	0	2	2	PLXDC2 Isoform 1 of Plexin domain-containing protein 2 precursor	n[145]VGLSDAFVVVHR	3	481.6127	4.162	0.411
91	IPI00044369	1	3.1	1.08	0	2	2	PLXDC2 Isoform 1 of Plexin domain-containing protein 2 precursor	VGLSDAFVVVHR	3	480.277	3.096	0.242
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	AGLLRPDYALLGHR	2	846.4888	4.1	0.498
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	GCPDVQASLPDAK	2	813.9026	4.395	0.535
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]GCPDVQASLPDAK[272]	2	817.9097	4.428	0.491
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]AGLLRPDYALLGHR	3	565.9973	3.185	0.288
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]EFTEAFLGCPAIHPR	3	626.6435	5.278	0.46
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]AGLLRPDYALLGHR	4	424.7498	2.43	0.061
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]AGLLRPDYALLGHR	2	848.4924	4.395	0.389

92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]TDCPGDALFDLLR	2	813.3928	3.958	0.371
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]PSLSHLLSQYYGAGVAR	3	655.0221	3.485	0.346
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	SLPLLM[147]DSVIQALAELEQK	3	798.782	3.795	0.276
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]DTLPSCAVR	2	576.2871	2.298	0.049
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]SLPLLM[147]DSVIQALAELEQK[272]	3	801.4534	3.228	0.267
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	AGLLRPDYALLGHR	4	423.7481	1.934	0.019
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]EFTEAFLGCPAIHPR	2	939.4616	2.569	0.252
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	AGLLRPDYALLG	2	699.9088	2.12	0.303
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]WGAAPYR	2	482.7597	2.027	0.055
92	IPI00163207	1	18.6	0.68	0.22	21	15	PGLYRP2 Isoform 1 of N-acetylmuramoyl-L-alanine amidase precur	n[145]AGLLRPDYALLG	2	701.9124	1.677	0.164
93	IPI00165972	1	11.2	1.49	0.64	6	4	CFD Complement factor D preproprotein	RPDSLQHVLLPVLDR	3	633.3705	3.989	0.522
93	IPI00165972	1	11.2	1.49	0.64	6	4	CFD Complement factor D preproprotein	n[145]RPDSLQHVLLPVLDR	3	634.7062	3.654	0.376
93	IPI00165972	1	11.2	1.49	0.64	6	4	CFD Complement factor D preproprotein	n[145]RPDSLQHVLLPVLDR	4	476.2815	4.244	0.448
93	IPI00165972	1	11.2	1.49	0.64	6	4	CFD Complement factor D preproprotein	VQVLLGAH	2	488.8006	2.162	0.393
93	IPI00165972	1	11.2	1.49	0.64	6	4	CFD Complement factor D preproprotein	n[145]PGIYTR	2	425.7488	1.958	0.019
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	INHGILYDEEK	2	805.9305	4.335	0.363
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	STDTSCVNPPTVQNAH	2	928.9169	5.071	0.517
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]EIM[147]ENYNIALR	2	763.3937	3.642	0.438
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]INHGILYDEEK[272]	2	809.9376	4.647	0.433
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]STDTSCVNPPTVQNAH	2	930.9205	4.271	0.508
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]INHGILYDEEK[272]	3	540.2942	3.53	0.38
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]EIMENYNIALR	2	755.3962	3.934	0.346
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]CLHPCVISR	3	421.8699	2.184	0.298
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	TGESAEFVCK	2	698.8336	2.45	0.405
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	LEYPTCAK	2	625.8173	2.06	0.264
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]LEYPTCAK[272]	2	629.8244	2.457	0.162
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	INHGILYDEEK	3	537.6228	3.311	0.167
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]YECR	2	380.6656	1.459	0.182
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]TTCWDGK[272]	2	572.7721	1.69	0.116
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]TGESAEFVCK[272]	2	702.8407	1.331	0.144
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	n[145]LEYPTCA	1	986.4444	1.966	0.147
94	IPI00167093	1	23	1.58	0.4	8	8	CFHR1 complement factor H-related 1	DTSCVNPPTVQ	2	673.8076	1.634	0.169
95	IPI00178375	1	0.7	0.94	0.05	12	4	HISPPD1 Isoform 2 of Inositol hexakisphosphate and diphosphoinos	n[145]ARLHEILQK[272]	3	465.9607	3.761	0.163
95	IPI00178375	1	0.7	0.94	0.05	12	4	HISPPD1 Isoform 2 of Inositol hexakisphosphate and diphosphoinos	ARLHEILQK	2	694.4303	2.584	0.104
95	IPI00178375	1	0.7	0.94	0.05	12	4	HISPPD1 Isoform 2 of Inositol hexakisphosphate and diphosphoinos	ARLHEILQK	3	463.2893	3.384	0.077
95	IPI00178375	1	0.7	0.94	0.05	12	4	HISPPD1 Isoform 2 of Inositol hexakisphosphate and diphosphoinos	n[145]ARLHEILQK[272]	2	698.4374	2.662	0.149
96	IPI00178926	1	8.8	0.78	0.05	4	4	IGJ immunoglobulin J chain	n[145]FVYHLSDLCK[272]	2	779.9037	3.912	0.329
96	IPI00178926	1	8.8	0.78	0.05	4	4	IGJ immunoglobulin J chain	FVYHLSDLCK	2	775.8966	3.861	0.233
96	IPI00178926	1	8.8	0.78	0.05	4	4	IGJ immunoglobulin J chain	n[145]FVYHLSDLCK[272]	3	520.2715	2.668	0.122
96	IPI00178926	1	8.8	0.78	0.05	4	4	IGJ immunoglobulin J chain	FVYHLSDLCK	3	517.6001	2.582	0.064
96	IPI00178926	1	8.8	0.78	0.05	4	4	IGJ immunoglobulin J chain	SPLR	1	612.3828	1.332	0.125
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]VTEPISAESGEQVER	2	887.9506	4.184	0.443
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	SETAEELKK	3	485.612	3.051	0.243
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]SETAEELK[272]K[272]	3	489.6191	3.763	0.286
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]SETAEELK[272]K[272]	2	733.925	3.045	0.307
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	ALADGVQK	2	541.3219	2.811	0.141
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	NEADELRK	2	627.8437	2.881	0.158
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	SETAEELKK	2	727.9144	3.304	0.313

97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	VAQELEEK	2	613.343	3.54	0.141
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	ALDNLAR	2	456.7667	2.587	0.126
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]NEADELRK[272]	2	631.8508	2.643	0.181
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	LNILNNNYK	2	693.3987	3.274	0.058
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]ANLQSVPHASASR	3	494.6029	2.856	0.282
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]NEADELRK[272]	3	421.5696	2.588	0.136
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]ALADGVQK[272]	2	545.329	2.15	0.085
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	SKLEDNIR	2	627.8619	3.082	0.119
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	SKLEDNIR	3	418.9104	2.659	0.109
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]SK[272]LEDNIR	2	631.869	2.772	0.066
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	n[145]VNEPSILEM[147]SR	2	717.8726	2.198	0.026
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	NEADELRK	3	418.8982	1.801	0.041
97	IPI00186903	1	25.4	1.04	0.34	29	19	APOL1 Isoform 2 of Apolipoprotein-L1 precursor	AELPR	1	725.4305	1.443	0.051
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	AATGECTATVGK	2	717.8576	3.932	0.488
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	DIPTNSPELEETLTHTITK	2	1210.1394	6.156	0.619
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IASFSQNCDIYPGK	2	934.9553	4.661	0.541
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IGEIKEETTSHLR	2	896.9995	4.222	0.466
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	TVGSDTFYSFK	2	766.3932	3.797	0.308
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]AATGECTATVGK[272]	2	721.8647	4.023	0.39
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]AVDAALK[272]K[272]	2	624.4061	3.265	0.429
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]DIPTNSPELEETLTHTITK[272]	2	1214.1465	6.661	0.621
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]	2	938.9624	4.537	0.539
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IGEIK[272]EETTSHLR	2	901.0066	4.58	0.405
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]TVGSDTFYSFK[272]	2	770.4003	3.883	0.422
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]YNSQNQSNNQFVLYR	2	1009.9937	6.31	0.461
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	DIPTNSPELEETLTHTITK	3	807.0954	3.436	0.161
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	KYFIDFVAR	3	480.2784	4.364	0.354
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	KYNSQNQSNNQFVLYR	3	761.7258	4.334	0.45
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]DIPTNSPELEETLTHTITK[272]	3	809.7668	2.435	0.122
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	AVDAALKK	3	412.5994	2.374	0.342
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IGEIKEETTSHLR	3	598.3354	3.577	0.374
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]AVDAALK[272]K[272]	3	416.6065	2.264	0.345
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]DFVQPPTK[272]	3	978.5016	4.255	0.387
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]VQVVAGK[272]K[272]	3	420.9503	2.485	0.327
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IASFSQNCDIYPGK[272]DFVQPPTK[272]	4	734.128	3.901	0.397
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]VQVVAGK[272]K[272]	2	630.9219	2.949	0.454
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	TVGSDTFYSFKYEIK	3	735.7246	3.593	0.356
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	VQVVAGKK	3	416.9432	2.217	0.31
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]YNSQNQSNNQFVLYR	3	673.6649	3.213	0.311
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	TWQDCEYK	2	699.8127	2.887	0.436
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]EGDCPVQSGK[272]	2	677.3227	2.709	0.264
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]ENFLFLTPDCK[272]	2	830.9195	3.004	0.385
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	EGDCPVQSGK	2	673.3156	2.229	0.245
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	KYFIDFVAR	2	719.9139	3.42	0.289
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IGEIK[272]EETTSHLR	4	451.0069	2.554	0.266
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	VQVVAGKK	2	624.9112	2.668	0.303
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]YFIDFVAR	2	587.8225	2.207	0.392
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	DFVQPPTK	2	606.3428	2.294	0.202

98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	VQVVAGK	2	490.8162	3.032	0.186
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]TWQDCEYK[272]	2	703.8198	2.659	0.315
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]TVGSDTFYSFK[272]YEIK[272]	3	739.7317	3.68	0.229
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]VQVVAGK[272]	2	494.8233	2.593	0.178
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]AVDAALK[272]	2	488.3075	2.198	0.188
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]ICVGCPR	2	492.2244	1.902	0.206
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	AVDAALK	2	484.3004	2.449	0.188
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	SCEYK	2	478.2225	1.952	0.216
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]DFVQPPTK[272]	2	610.3499	2.114	0.137
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]QVVAGLNFR	2	574.3389	2.605	0.145
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	ITEATK	2	471.7846	1.741	0.035
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]ITEATK[272]	2	475.7917	2.142	0.073
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	AVDAALKK	2	618.3954	1.972	0.143
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]ETTCSK[272]	2	501.7456	1.891	0.168
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]SCEYK[272]	2	482.2296	1.745	0.166
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IDNVKK	2	568.8612	2.026	0.076
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IGEIK[272]EETTSHLR	3	601.0068	3.42	0.122
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IDNVK	2	434.7662	1.944	0.066
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	YEIKEGDCPVQSGK	3	673.6728	2.449	0.181
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]SDTFYSFK[272]	2	641.8316	3.152	0.331
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]FIDFVAR	2	506.2908	2.415	0.089
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	TVGSDTFYSFK	3	511.2646	2.527	0.079
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]CQPLGM[147]ISLM[147]K[272]	2	793.9045	3.027	0.286
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IYPTVNCQPLGM[147]	2	771.3678	2.928	0.33
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	CQPLGM[147]ISLM[147]K	2	789.8974	2.869	0.234
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]IYPTVNCQPLGM	2	763.3703	2.776	0.138
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	IYPTVN	1	846.472	1.7	0.123
98	IPI00215894	1	49.9	0.75	0.05	11	9	KNG1 Isoform LMW of Kininogen-1 precursor	n[145]RPPGFSPF	2	524.7885	1.56	0.098
99	IPI00215983	1	27.7	0.9	0.04	5	4	CA1 Carbonic anhydrase 1	SLLSNVEGDNAVPM[147]QHNNRPTQPLK	4	764.649	2.651	0.295
99	IPI00215983	1	27.7	0.9	0.04	5	4	CA1 Carbonic anhydrase 1	VLDALQAIK	2	625.8952	3.281	0.217
99	IPI00215983	1	27.7	0.9	0.04	5	4	CA1 Carbonic anhydrase 1	n[145]VGEANPK[272]	2	501.7948	2.24	0.09
99	IPI00215983	1	27.7	0.9	0.04	5	4	CA1 Carbonic anhydrase 1	n[145]VLDALQAIK[272]	2	629.9023	1.944	0.055
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	n[145]LSSGLVTAALYGR	2	726.4206	4.57	0.512
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	DAQEVHCDEATK	3	557.9204	3.077	0.325
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	n[145]WNPCLEPHR	3	447.8819	3.386	0.237
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	n[145]ILSGDPYCEK[272]	2	729.8642	2.495	0.223
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	n[145]VDLITFDTPFAGR	2	798.4311	2.894	0.335
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	n[145]DAQEVHCDEATK[272]	3	560.5918	2.377	0.008
100	IPI00218413	1	12.3	1.27	0.52	7	7	BTD biotinidase precursor	MAHAHIQGGR	3	406.5489	1.688	0.04
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	IFFYDSENPPASEVLR	2	1012.5098	5.019	0.548
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	YVYIAELLAHK	2	800.4665	4.607	0.39
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]IFFYDSENPPASEVLR	2	1014.5134	4.572	0.542
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]YVYIAELLAHK[272]	2	804.4736	4.643	0.398
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	YVYIAELLAHK	3	533.9801	4.57	0.321
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]YVYIAELLAHK[272]	3	536.6515	2.808	0.224
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	FDVSSFNPHGISTFTDEDNAM[147]YLLVVNHPDAK	4	969.9689	7.797	0.629
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]FDVSSFNPHGISTFTDEDNAMYLLVVNHPDAK[27	4	967.9737	5.491	0.433
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	SFNPNSPGK	2	614.3277	2.94	0.338

101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]SFNPNSPGK[272]	2	618.3348	2.559	0.269
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]EVQPVELPNCNLVK[272]	2	958.5148	2.628	0.246
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]IHVYEK[272]	2	538.8208	1.819	0.208
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]IHVYEK[272]	3	359.5496	2.144	0.117
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	LLIGTVFHK	3	436.6115	3.294	0.264
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	IQNILTEEPK	2	732.9247	1.463	0.273
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]STVEIFK[272]	2	556.3337	2.331	0.232
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	STVEIFK	2	552.3266	1.676	0.141
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	ALYCEL	1	897.4209	1.455	0.132
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	EVQPVELPNCNLVK	2	954.5077	2.232	0.061
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	IHVYEK	2	534.8137	1.466	0.063
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]YVYIAELLAH	2	668.3751	3.166	0.437
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	n[145]ALYCEL	1	901.428	1.189	0.139
101	IPI00218732	1	31.8	0.84	0.05	24	22	PON1 Serum paraoxonase/arylesterase 1	FDVSSFNPH	2	595.2855	1.588	0.145
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]GSQAVSYTR	2	556.7945	2.971	0.356
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]LTYNHGGITGSR	3	473.9202	2.237	0.196
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]AYSHSVSLTR	2	632.842	2.832	0.235
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	GSQAVSYTR	2	554.7909	2.618	0.214
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]GNPAVSYVR	2	553.8074	2.663	0.118
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]VLVENEHR	2	570.3181	2.034	0.078
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]VLVENEHR	3	380.5478	1.419	0.194
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	n[145]YHGAVCGLCGNM[147]DR	3	583.2369	2.129	0.086
102	IPI00242956	1	2.4	0.6	0.22	9	7	FCGBP IgGFc-binding protein precursor	SK[272]K[272]	2	395.7731	0.913	0.136
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	DQTVSDNELQEMSNQGSK	2	1145.5345	5.66	0.51
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	DQTVSDNELQEM[147]SNQGSK	2	1153.5319	5.142	0.501
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	EILSVDCSTNNPSQAK	2	1016.4956	5.604	0.538
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KTLLSNLEEAK	2	833.4986	4.029	0.392
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	VTTVASHTSDSDVPSGVTEVVVK	2	1297.6873	6.806	0.599
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]DQTVSDNELQEM[147]SNQGSK[272]	2	1157.539	5.163	0.588
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]EILSVDCSTNNPSQAK[272]	2	1020.5027	4.618	0.508
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]TLLSNLEEAK[272]K[272]	2	839.5092	4.364	0.385
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]VTTVASHTSDSD	2	682.3265	3.669	0.467
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	DQTVSDNELQEM[147]SNQGSK	3	769.357	4.767	0.467
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KTLLSNLEEAKK	3	645.3981	4.193	0.422
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	VTTVASHTSDSDVPSGVTEVVVK	3	865.4606	4.914	0.479
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHMLDVMQDHFSR	3	672.9882	4.59	0.433
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHM[147]LDVMQDHFSR	3	678.3198	4.939	0.275
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHM[147]LDVM[147]QDHFSR	3	683.6515	3.709	0.414
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]VTTVASHTSDSDVPSGVTEVVVK[272]	3	868.132	4.574	0.488
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KTLLSNLEEAKK	4	484.3004	3.151	0.347
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHMLDVMQDHFSR	4	504.993	4.292	0.358
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHMLDVM[147]QDHFSR	4	508.9917	3.247	0.361
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHM[147]LDVMQDHFSR	4	508.9917	3.394	0.32
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHM[147]LDVM[147]QDHFSR	4	512.9904	2.705	0.213
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KTLLSNLEEAK	3	556.0015	1.993	0.243
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]TLLSNLEEAK[272]K[272]	3	560.0086	2.889	0.104
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	EIQNAVNGVK	2	676.3883	3.816	0.318
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	TLLSNLEEAKK	2	833.4986	3.873	0.388

103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]EIQNAVNGVK[272]	2	680.3954	3.429	0.234
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	TLLSNLEEAKK	3	556.0015	3.344	0.197
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]RELDESLQVAER	3	530.2852	4.271	0.165
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]ASSIIDELFQDR	2	769.4026	2.99	0.286
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	TLLSNLEEAK	2	699.4036	1.73	0.087
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	FM[147]ETVAEK	2	625.8261	3.004	0.185
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]ELDESLQVAER	2	716.8737	2.761	0.276
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]LFDSDPITVTVPVEVSR	2	1009.55	2.98	0.357
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KYNELLK	2	664.4085	3.059	0.12
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]FMETVAEK[272]	2	621.8358	2.821	0.182
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]MLDVM[147]QDHFSR	3	513.5786	3.362	0.322
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]TLLSNLEEAK[272]	2	703.4107	2.472	0.242
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KYNELLK	3	443.2748	2.697	0.107
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	ALQEYR	2	460.2535	2.063	0.115
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]FM[147]ETVAEK[272]	2	629.8332	2.556	0.102
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]TLIEK[272]	2	446.2913	2.044	0.245
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]QQTHMLDVM[147]QDHFSR	3	678.3198	4.882	0.485
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]MLDVMQDHFSR	2	761.8668	4.239	0.315
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]ALQEYR	2	462.257	1.969	0.008
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]ALQEYR	1	923.5068	1.651	0.067
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]PITVTVPVEVSR	2	720.9308	2.192	0.207
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]IDSLLENDR	2	609.826	3.181	0.208
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]MK[272]DQCDK[272]	3	449.2261	1.988	0.135
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]MLDVM[147]QDHFSR	2	769.8643	3.471	0.319
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	EPQDTYHYLPF	2	775.3697	2.307	0.132
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	KKEDALNETR	3	541.9774	2.083	0.035
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	MLDVM[147]QDHFSR	3	512.2429	3.112	0.393
103	IPI00291262	1	44.2	0.69	0.12	79	51	CLU Clusterin precursor	n[145]EPQDTYHYLPF	2	777.3733	2.156	0.162
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	DFTCVHQALK	2	744.3785	4.451	0.391
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	GVTSVSQIFHSPDLAIR	2	984.0391	5.33	0.519
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KYPVAHFIDQTLK	2	990.5752	4.488	0.446
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LEDM[147]EQALSPSVFK	2	945.4875	3.779	0.493
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	TNLESILSYPK	2	772.9378	4.173	0.404
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	VTTSQDM[147]LSIM[147]EK	2	897.9525	3.247	0.521
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]DFTCVHQALK[272]	2	748.3856	4.016	0.452
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]GVTSVSQIFHSPDLAIR	2	986.0426	6.57	0.474
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]HRLEDMEQALSPSVFK[272]	2	1088.0772	4.343	0.434
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LEDM[147]EQALSPSVFK[272]	2	949.4946	4.166	0.458
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LVLLNAIYLSAK[272]	2	803.5128	4.498	0.371
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]TNLESILSYPK[272]	2	776.9449	3.977	0.396
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VTTSQDMLSIMEK[272]	2	885.9647	4.802	0.339
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VTTSQDM[147]LSIM[147]EK[272]	2	901.9596	4.542	0.464
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]YPVAHFIDQTLK[272]	2	860.4873	4.26	0.388
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	DFTCVHQALK	3	496.5881	3.464	0.354
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	HRLEDM[147]EQALSPSVFK	3	728.3808	3.961	0.403
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KYPVAHFIDQTLK	3	660.7192	4.96	0.417
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]DFTCVHQALK[272]	3	499.2595	3.832	0.318
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]GVTSVSQIFHSPDLAIR	3	657.6975	4.517	0.396

104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]HRLEDMEQALSPSVFK[272]	3	725.7205	4.444	0.464
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	YPVAHFIDQTLK	3	571.3225	3.311	0.292
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]HRLEDM[147]EQALSPSVFK[272]	3	731.0522	5.308	0.298
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]YPVAHFIDQTLK[272]	3	573.9939	3.345	0.241
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	VTTSQDMLSIMEK	2	881.9576	3.847	0.333
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LEDMEQALSPSVFK[272]	2	941.4972	3.48	0.44
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]FQPTLLTLPR	2	665.4042	1.471	0.113
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	HRLEDMEQALSPSVFK	3	723.0491	3.727	0.27
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LFVLWDQQHK[272]	3	534.6396	4.289	0.325
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VTTSQDMLSIM[147]EK[272]	3	596.3105	3.34	0.226
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LEDMEQALSPSVFK	2	937.4901	3.863	0.278
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LYHAFSAM[147]K	2	682.3632	3.307	0.261
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LYHAFSAM[147]K[272]	2	686.3703	2.981	0.278
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VATTVISK[272]	2	553.8548	1.538	0.156
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LYHAFSAM[147]K	3	455.2446	3.067	0.167
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	NSVIKVPM[147]M[147]NSK	3	600.6672	2.678	0.308
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LVLLNAIYLSAK	2	799.5057	2.988	0.437
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	M[147]EPFHFK	2	616.3183	2.48	0.207
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	YPVAHFIDQTLK	2	856.4802	3.807	0.405
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LYHAFSAM[147]K[272]	3	457.916	3.176	0.133
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KYPVAHFIDQTLK	4	495.7912	3.276	0.152
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	VATTVISK	2	549.8477	2.268	0.218
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]TNLESILSYPK[272]	3	518.299	2.049	0.048
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LLDSLPSDTR	2	630.8495	2.643	0.215
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KYPVAH	3	332.1993	2.413	0.342
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VTTSQDM[147]LSIMEK[272]	2	893.9621	4.029	0.396
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]TLYSSSPR	2	527.7861	2.089	0.113
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]TLYSSSPR	1	1054.565	1.959	0.115
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	VTTSQDMLSIM[147]EK	2	889.955	4.332	0.505
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	GVTSVSQIFHSPDLAIR	3	656.3618	4.028	0.178
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]AIMEK[272]	2	440.2643	2.159	0.176
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]MEPFHFK[272]	3	408.5544	2.373	0.093
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]VTTSQDMLSIM[147]EK[272]	2	893.9621	2.62	0.229
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LEMSK	2	444.2546	1.926	0.03
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]IEMSK[272]	2	448.2617	1.625	0.148
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	FPVFM[147]GR	2	505.268	1.428	0.161
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LFVLWDQQHK	3	531.9682	2.858	0.197
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]LFVLWDQQHK[272]	2	801.4558	3.243	0.343
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LYHAFSAM[147]KK	4	408.7328	1.82	0.03
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LFVLWDQQHK	2	797.4487	3.289	0.249
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]FIDQTLK[272]	2	576.847	2.294	0.067
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]GVTSVSQIFHS	2	653.3496	2.902	0.34
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	TNLESILSYPK	3	515.6276	2.03	0.056
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]GVTSVSQIFH	2	609.8336	2.072	0.273
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KVETNMAFSPF	2	775.9055	2.371	0.181
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	n[145]DFTCVH	2	456.1972	1.459	0.216
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	KVETNM[147]AFSPF	2	783.9029	1.999	0.054
104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	LYHAF	2	395.716	1.426	0.03

104	IPI00291866	1	38.8	0.72	0.04	121	66	SERPING1 Plasma protease C1 inhibitor precursor	SK[272]K[272]	2	395.7731	0.913	0.136
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	KVTYTSQEDLVEK	2	980.5412	5.266	0.459
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]AQLGDLPWQVAIK[272]	2	864.0084	4.489	0.444
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]GLETSLAECTFTK[272]	2	867.4383	4.463	0.483
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]SLECLHPGTK[272]	2	709.8724	3.632	0.45
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	IIFHENYNAGTYQNDIALIEM[147]K	3	965.1566	4.62	0.399
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]IIFHENYNAGTYQNDIALIEM[147]K[272]	3	967.828	5.605	0.443
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	AQLGDLPWQVAIK	2	860.0013	4.024	0.42
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]YTHLSCDK[272]	3	434.2138	2.743	0.369
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	SLECLHPGTK	2	705.8653	3.556	0.421
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]HGNTDSEGIVEVK[272]	2	836.9409	3.344	0.409
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	SLECLHPGTK	3	470.9126	2.729	0.256
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	YTHLSCDK	2	646.81	2.769	0.3
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]YTHLSCDK[272]	2	650.8171	2.394	0.316
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]CIEGTCVCK[272]	2	691.2896	2.71	0.235
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]IVIEYVDR	2	575.8331	2.441	0.23
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	HGNTDSEGIVEVK	2	832.9338	2.506	0.253
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]VTYTSQEDLVEK[272]K[272]	3	658.037	2.408	0.15
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]SLECLHPGTK[272]	3	473.584	2.318	0.151
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]FYGNR	2	400.7122	1.609	0.095
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	YTHLSCDK	3	431.5424	2.09	0.067
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	YISQM[147]K	2	533.2917	1.648	0.084
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	LVDQDK	2	499.2875	1.528	0.074
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]TM[147]GYQDFADVVCYTQK[272]	3	740.3402	1.983	0.056
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	n[145]SLLPK[272]	2	423.2886	0.884	0.016
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	KVTYTSQEDLVEK	3	654.0299	2.231	0.184
105	IPI00291867	1	25.4	0.64	0.19	26	24	CFI Complement factor I precursor	EM[147]K	1	703.3808	1.121	0.072
106	IPI00292218	1	4.6	0.68	0.19	4	4	MST1 Hepatocyte growth factor-like protein precursor	n[145]TPFDYCALR	2	638.3028	2.497	0.191
106	IPI00292218	1	4.6	0.68	0.19	4	4	MST1 Hepatocyte growth factor-like protein precursor	n[145]WDAQIPHQHR	3	477.9169	3.261	0.143
106	IPI00292218	1	4.6	0.68	0.19	4	4	MST1 Hepatocyte growth factor-like protein precursor	n[145]AAFCYQIR	2	581.2869	2.189	0.085
106	IPI00292218	1	4.6	0.68	0.19	4	4	MST1 Hepatocyte growth factor-like protein precursor	n[145]YTPTLR	2	447.7619	1.739	0.019
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	IYEDHDATQQLQGFYSQVAK	2	1311.1534	6.542	0.614
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]FAHYVVTSQVVNTANEAR	2	1075.5592	7.304	0.573
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]FPLYNLGFGH	2	654.8465	4.26	0.47
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GHMLENHVER	2	683.3443	3.156	0.407
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GHM[147]LENHVER	2	691.3418	2.972	0.512
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GSLVQASEANLQAAQDFVR	2	1074.5619	7.387	0.577
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]ILGDM[147]QPGDYFDLVLFGTR	2	1159.0864	5.617	0.545
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]IYEDHDATQQLQGFYSQVAK[272]	2	1315.1605	6.647	0.584
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]MSLDYGFVTPLTSM[147]SIR	2	1039.5246	4.58	0.471
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GHMLENHVER	3	454.5629	3.551	0.402
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GHM[147]LENHVER	3	459.8946	4.241	0.437
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	IYEDHDATQQLQGFYSQVAK	3	874.438	2.876	0.069
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]FAHYVVTSQVVNTANEAR	3	717.3752	5.352	0.456
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GHMLENHVER	3	455.8986	4.606	0.307
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GHM[147]LENHVER	3	461.2303	3.763	0.419
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]IYEDHDATQQLQGFYSQVAK[272]	3	877.1094	6.5	0.612
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]TMEQFTIHLTVNPQSK[272]	3	721.3888	4.965	0.357

107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]TM[147]EQFTIHLTVNPQSK[272]	3	726.7204	4.952	0.487
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GMADQDGLK[272]PTIDK[272]PSEDSPPLEMLGR	4	857.4457	4.067	0.404
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]AAISGENAGLVR	2	651.3683	4.086	0.399
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GM[147]ADQDGLKPTIDKPSEDSPPLEM[147]LGPR	3	1149.5814	3.836	0.329
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	TMEQFTIHLTVNPQSK	3	718.7174	4.547	0.338
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GSLVQASEANLQAAQDFVR	3	716.7104	3.004	0.364
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]NHM[147]QYEIVIK[272]	3	526.9566	3.568	0.24
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]TFVLSALQPSPTHSSSNTQR	3	768.0684	4.17	0.321
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GMADQDGLKPTIDKPSEDSPPLEMLGPR	4	854.4404	4.231	0.363
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GHMLENHVER	2	681.3408	2.986	0.457
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GHM[147]LENHVER	2	689.3383	2.833	0.49
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GFSLDEATNLNGGLLR	2	910.9848	4.733	0.326
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]NHM[147]QYEIVIK[272]	2	789.9313	3.986	0.343
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]ELAAQTIK[272]K[272]	3	478.6399	3.198	0.203
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	LDAQASFLPK	2	685.3956	3.075	0.185
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	NHM[147]QYEIVIK	2	785.9242	3.656	0.294
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]LDAQASFLPK[272]	2	689.4027	2.021	0.171
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]QYYEGSEIVVAGR	2	807.9159	3.931	0.361
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]QAVDTAVDGVFIR	2	767.9209	3.809	0.437
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]VTFQLTYEEVLK[272]R	3	638.7028	2.968	0.281
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	TM[147]EQFTIHLTVNPQSK	3	724.049	4.571	0.321
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	ELAAQTIK	2	577.3506	2.818	0.035
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	KAAISGENAGLVR	2	783.4598	3.62	0.2
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]VTYDVSR	2	492.2676	2.337	0.206
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GM[147]ADQDGLKPTIDKPSEDSPPLEM[147]LGPR	4	862.4379	2.82	0.243
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]ELAAQTIK[272]	2	581.3577	2.707	0.087
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GM[147]ADQDGLKPTIDKPSEDSPPLEMLGPR	4	858.4391	2.775	0.243
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	EVAFDLEIPK	2	720.9085	2.603	0.179
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]PLLVDVDLQYPQDAVLALTQNHHK[272]	4	754.6681	2.891	0.307
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GHM[147]LENHVER	4	346.1746	2.393	0.105
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	VTYDVSR	2	490.264	2.049	0.169
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GMADQDGLK[272]PTIDK[272]PSEDSPPLEM[147	4	861.4444	2.201	0.005
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]EALLK[272]	2	431.2861	1.702	0.145
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]HYVVTSQVVNTANEAR	3	644.6734	3.343	0.248
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GM[147]ADQDGLK[272]PTIDK[272]PSED	3	789.0688	2.481	0.336
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]RNHM[147]QYEIVIK[272]	3	578.9903	2.29	0.089
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	SLKVNCK	3	419.9105	2.054	0.168
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	EAIIK	2	427.279	1.917	0.224
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GRFPLYNLGFGH	3	507.941	2.648	0.342
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]FAHYVVTSQVVN	2	754.4049	2.882	0.374
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]SLK[272]VNCK[272]	2	635.3728	2.446	0.168
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]EVAFDLEIPK[272]	2	724.9156	1.966	0.039
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]ILGDM[147]QPGDYFDLVLFGTR	3	773.06	2.375	0.135
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]LTIQELLAK[272]	2	658.9232	3.968	0.144
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]TAFISDFAVTADGNAFIGDIK[272]	3	821.1	2.443	0.225
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	KSFSGK	2	537.327	1.865	0.132
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GHM[147]LENHVER	4	345.1728	1.736	0.008
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]AISGENAGLVR	2	615.8498	3.773	0.194

107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	LTIQELLAK	2	654.9161	3.647	0.145
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	SLKVNCK	2	629.3622	1.905	0.145
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	TAFISDFAVTADGNAFIGDIK	3	818.4286	2.507	0.092
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]ADVQAHGEGQEF	2	716.3347	3.063	0.134
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]SLK[272]VNCK[272]	3	423.9176	1.829	0.035
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]GRFPLYNLGF	2	664.3676	1.824	0.225
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GRFPLYNLGF	2	662.3641	2.252	0.288
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GRFPLYNLGFGH	2	759.4043	2.838	0.158
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	n[145]MSLDYGFVTPLTSM[147]SIR	3	693.3521	3.067	0.033
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	GM[147]ADQDGLKPTIDKPSED	3	785.0617	2.279	0.083
107	IPI00292530	1	41.6	0.73	0.1	119	69	ITIH1 Inter-alpha-trypsin inhibitor heavy chain H1 precursor	EM[147]K	1	703.3808	1.121	0.072
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]QEINSHVEM[147]QTK[272]	3	583.3011	3.72	0.388
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	QEINSHVEMQTK	3	575.2981	2.724	0.339
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	QEINSHVEM[147]QTK	3	580.6297	3.384	0.346
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]TEDSSSFLIDK[272]	2	765.4005	3.761	0.364
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]NALALFVLPK[272]	2	687.4416	3.476	0.339
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]SFMLLILER	2	633.3741	2.982	0.341
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	NALALFVLPK	2	683.4345	3.582	0.395
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	TEDSSSFLIDK	2	761.3934	2.863	0.309
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	FTVETPDK	2	608.8323	2.837	0.151
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]HLK[272]PLAK[272]	2	619.9191	2.035	0.25
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]GWVDLFVPK[272]	2	674.897	2.585	0.201
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	HLKPLAK	2	613.9085	2.113	0.151
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	GWVDLFVPK	2	670.8899	2.067	0.103
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	HLKPLAK	3	409.6081	1.887	0.13
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]FAFNLYR	2	537.7963	2.516	0.3
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]FLNDVK[272]	2	512.3075	2	0.019
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]AFNLYR	2	464.2621	2.458	0.176
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]HLK[272]PLAK[272]	3	413.6152	1.893	0.322
108	IPI00292946	1	19	0.96	0.2	18	18	SERPINA7 Thyroxine-binding globulin precursor	n[145]FNLYR	2	428.7435	1.775	0.037
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	SVNDLYIQK	2	680.3852	3.856	0.37
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]EYYFAEAQIADFSDPAFISK[272]	2	1300.6436	5.857	0.494
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]SVNDLYIQK[272]	2	684.3923	4.111	0.364
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]VEM[147]THNHNFR	2	722.8473	3.766	0.426
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]YEITTIHNLFR	2	775.926	3.894	0.392
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	EYYFAEAQIADFSDPAFISK	3	864.7601	5.707	0.359
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	FTVDRPFLFLIYEHR	3	698.3826	3.989	0.349
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]EYYFAEAQIADFSDPAFISK[272]	3	867.4315	4.653	0.488
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEM[147]THNHNFR	4	422.9576	3.152	0.401
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]GETHEQVHSILHFK[272]	4	488.2664	3.594	0.395
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEM[147]THNHNFR	3	563.6076	2.849	0.348
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FTVDRPFLFLIYEHR	3	699.7183	3.732	0.412
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]YEITTIHNLFR	3	517.6198	3.163	0.318
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FAFNLYR	2	537.7963	2.516	0.3
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]NYNLVESLK[272]	2	684.3923	2.796	0.275
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	NYNLVESLK	2	680.3852	2.636	0.281
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	QFPILLDFK	2	700.9187	2.595	0.251
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]QFPILLDFK[272]	2	704.9258	1.982	0.204

109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEMTHNHNFR	3	558.276	2.569	0.138
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEMTHNHNFR	2	836.9104	3.539	0.243
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]TLEAQLTPR	2	586.8414	2.739	0.21
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]NFGYTLR	2	507.7781	2.523	0.132
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]VEMTHNHNFR	3	476.9023	3.348	0.353
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]TSCLLFM[147]GR	2	617.2992	2.542	0.236
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEM[147]THNHNFR	2	844.9078	3.413	0.358
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]VEM[147]THNHNFR	3	482.2339	2.378	0.13
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPVEMTHNHNFR	4	418.9588	2.249	0.098
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]LFLIYEHR	3	412.2403	2.97	0.252
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]IAIDIFK[272]	2	554.3545	2.03	0.154
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	EVIIPK	2	489.821	1.843	0.095
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	IAIDIFK	2	550.3474	2.258	0.054
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]EVIIPK[272]	2	493.8281	1.968	0.036
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]TVDRPFLFLIYEHR	4	488.2734	3.198	0.306
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]LIYEHR	3	325.5228	2.513	0.179
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]LFLIYEHR	2	617.8569	2.999	0.223
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]ILNCIYFK[272]	2	674.374	2.873	0.28
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	FPILLDFK	2	636.8894	2.684	0.183
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]AFNLYR	2	464.2621	2.458	0.176
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]GETHEQVHSILH	3	510.9309	2.49	0.198
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FTVDRPFLF	2	643.3567	1.992	0.078
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]LIYEHR	2	487.7806	2.345	0.132
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]VEM[147]THNHNFR	4	361.9273	2.337	0.234
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]QFPILLDFK[272]	3	470.2863	1.84	0.032
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FPILLDFK[272]	2	640.8965	2.149	0.08
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]FNLYR	2	428.7435	1.775	0.037
109	IPI00292950	1	29.3	0.7	0.07	84	44	SERPIND1 Serpin peptidase inhibitor	n[145]CIYFK[272]	2	504.2685	1.655	0.213
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	FSEGTAGDSLSLHSGR	2	880.9318	4.897	0.477
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	YAVSEAAAHK	2	663.8619	3.957	0.537
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]YAVSEAAAHK[272]	2	667.869	4.352	0.56
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	LLGEVDHYQLALGK	3	612.6841	3.226	0.436
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	YAVSEAAAHK	3	442.9104	4.704	0.49
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]FSEGTAGDSLSLHSGR	3	588.9593	4.884	0.457
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]LLGEVDHYQLALGK[272]	3	615.3555	2.607	0.166
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]YAVSEAAAHK[272]	3	445.5818	4.274	0.459
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]TFAHYATFR	3	419.8881	2.767	0.249
110	IPI00293925	1	17	0.67	0.03	15	10	FCN3 Isoform 1 of Ficolin-3 precursor	n[145]TFAHYATFR	2	629.3285	2.074	0.239
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	DVDECSLKPSICGTAVCK	3	809.3742	4.651	0.465
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]DVDECSLK[272]PSICGTAVCK[272]	3	813.3813	4.508	0.492
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]IQALSLCSDQQSHLEFR	3	722.3582	4.144	0.518
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]DCK[272]DVDECSLK[272]PSICGTAVCK[272]	4	744.343	4.123	0.548
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	KVESELIKPINPR	3	648.3982	3.909	0.305
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	KVESELIKPINPR	4	486.5505	3.146	0.344
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]HCLVTVEK[272]	2	631.8456	2.705	0.426
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	NNLELSTPLK	2	704.9116	3.52	0.277
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]IETISHEDLQR	2	742.8949	3.353	0.304
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]SFQTGLFTAAR	2	671.8654	2.864	0.369

111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]VYFAGFPR	2	550.8041	2.675	0.361
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]SQDILLSVENTVIYR	2	947.5238	3.368	0.219
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	HCLVTVEK	2	627.8385	2.785	0.307
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]NNLELSTPLK[272]	2	708.9187	3.354	0.157
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]QLAVLDK[272]	2	537.8417	2.031	0.131
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]NIPGDFECECPEGYR	2	982.892	1.661	0.225
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	SFQTGLFTAAR	2	669.8619	2.279	0.124
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]IEVQLK[272]	2	509.331	2.039	0.067
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]SCEVVSVCLPLNLDTK[272]	3	700.6866	2.34	0.039
111	IPI00294004	1	23.4	0.73	0.19	25	19	PROS1 Vitamin K-dependent protein S precursor	n[145]SELIK[272]	2	439.2835	2.103	0.159
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	AEAQAQYSAAVAK	2	794.4281	4.944	0.367
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ANTVQEATFQM[147]ELPK	2	1002.017	5.354	0.526
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ETLFSVMPGLK	2	751.426	4.357	0.421
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ETLFSVM[147]PGLK	2	759.4235	3.992	0.419
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ITFELVYEELLK	2	889.011	3.679	0.459
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]AEAQAQYSAAVAK[272]	2	798.4352	4.983	0.513
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ETLFSVMPGLK[272]	2	755.4331	3.501	0.447
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ETLFSVM[147]PGLK[272]	2	763.4306	3.517	0.46
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	DQFNLIVFSTEATQWRPSLVPASAENVNK	3	1181.2861	5.92	0.573
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	EKAEAQAQYSAAVAK	3	662.3654	4.51	0.529
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ITFELVYEELLKR	3	645.0434	3.128	0.455
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]EK[272]AEAQAQYSAAVAK[272]	3	666.3725	4.565	0.38
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]EK[272]NGIDIYSLTVDSR	3	666.6964	4.116	0.439
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]FK[272]PTLSQQQK[272]	3	546.3298	2.089	0.021
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ITFELVYEELLK[272]R	3	647.7148	3.584	0.375
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]NPLVWVHASPEHVVVTR	3	695.3889	3.303	0.383
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]VPDHAAYHPFR	3	485.2534	4.443	0.388
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]YIFHNFMER	3	467.5682	3.928	0.357
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	NPLVWVHASPEHVVVTR	4	520.7917	4.709	0.506
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	FKPTLSQQQK	3	542.3227	1.987	0.155
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]NPLVWVHASPEHVVVTR	4	521.7935	3.511	0.316
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	FKPTLSQQQK	2	812.9804	3.928	0.314
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	GPDVLTATVSGK	2	712.9091	3.601	0.403
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	GSEMVVAGK	2	579.321	3.157	0.35
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	LGVYELLLK	2	664.4211	3.783	0.349
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ANTVQEATFQM[147]ELPK[272]	2	1006.0241	4.838	0.373
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]LGVYELLLK[272]	2	668.4282	2.954	0.252
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	GKSAGLVK	3	393.9239	3.478	0.236
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GYFVHYFAPEGLTTM[147]PK[272]	3	754.7218	4.43	0.402
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]FK[272]PTLSQQQK[272]	2	818.991	3.176	0.309
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GPDVLTATVSGK[272]	2	716.9162	3.112	0.39
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GSEMVVAGK[272]	2	583.3281	3.008	0.227
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GSEM[147]VVAGK[272]	2	591.3256	2.579	0.393
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]NGIDIYSLTVDSR	2	798.9211	3.92	0.395
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	AGFSWIEVTFK	2	782.9298	3.932	0.318
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	YYLQGAK	2	561.819	2.598	0.311
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ANTVQEATFQM[147]ELPK	3	668.3471	3.048	0.284
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]YIFHNFM[147]ER	3	472.8999	2.371	0.238

112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	GSEM[147]VVAGK	2	587.3185	1.915	0.175
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ITFELVYEELLK[272]	2	893.0181	3.429	0.274
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]NVVFVIDK[272]	2	611.3759	2.165	0.166
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]YYLQGAK[272]	2	565.8261	2.446	0.27
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]SPEQQETVLDGNLIIR	2	978.5296	4.528	0.377
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	NVVFVIDK	2	607.3688	2.01	0.191
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]LALDNGGLAR	2	572.3338	2.311	0.156
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	RLGVYELLLK	3	495.3168	3.25	0.11
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	LALDNGGLAR	2	570.3302	2.289	0.043
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]FAHTVVTSR	2	581.3285	2.178	0.236
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]YIFHNFMER	2	700.8487	2.717	0.166
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ETLFSVMPGLK	3	501.2864	2.829	0.124
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ILDDLSPR	2	536.8096	2.837	0.092
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]YIFHNFM[147]ER	2	708.8462	2.401	0.183
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	KPTLSQQQK	3	493.2999	3.31	0.253
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ETLFSVMPGLK[272]	3	503.9578	2.729	0.11
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ETLFSVM[147]PGLK[272]	3	509.2895	2.297	0.107
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	SAGLVK	2	427.7766	2.094	0.081
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GK[272]SAGLVK[272]	3	397.931	3.026	0.097
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]RLGVYELLLK[272]	3	497.9882	2.703	0.069
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	IPKPEASFSPR	3	503.6243	2.035	0.135
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]IPK[272]PEASFSPR	2	758.94	1.694	0.205
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ITFELVYEELLK[272]	3	595.6811	2.701	0.158
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ASPEHVVVTR	3	412.2349	2.659	0.256
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]FAHTVVTSR	3	387.8881	1.829	0.14
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	TGLLLLSDPDK	2	726.4271	1.808	0.124
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]LGVYELLLK[272]	3	445.9545	3.013	0.035
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]EQQETVLDGNLIIR	2	886.4872	4.618	0.339
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GK[272]SAGLVK[272]	2	596.3929	2.669	0.143
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]DVLTATVSGK[272]	2	639.879	3.211	0.364
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	DVLTATVSGK	2	635.8719	3.276	0.313
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ASPEHVVVTR	3	413.5706	2.214	0.09
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GPDVLTATVSGK[272]	3	478.2799	1.851	0.236
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]ASPEHVVVTR	2	619.8523	3.187	0.406
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]AGFSWIEVTFK	2	784.9333	2.937	0.137
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ITFELVYEELLK	3	593.0097	2.541	0.114
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]PEQGVEVTGQYER	2	818.4084	3.286	0.191
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]PTVGETNPR	2	557.8023	2.412	0.24
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]GVYELLLK[272]	2	611.8861	3.001	0.124
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	GKSAGLVK	2	590.3823	1.769	0.117
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]EK[272]NGIDIYSLTVDSR	2	999.541	3.973	0.398
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]NPLVWVH	2	504.791	1.941	0.296
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]LDNGGLAR	2	480.2732	1.943	0.069
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]VPDHAAYHPFR	4	364.1919	2.422	0.099
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]WKETLFSVM[147]PGLK	3	659.3769	4.075	0.334
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]FSSHVGGTLGQF	2	690.8551	2.617	0.303
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	HLQM[147]DIH	3	350.5114	2.378	0.095
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	IPKPEASFSPR	2	754.9329	1.217	0.062

112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	ASPEHVVVTR	2	617.8488	2.549	0.159
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	EKNGIDIYSLTVDSR	3	664.025	2.478	0.158
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	n[145]HLQM[147]DIH	3	351.8471	1.653	0.128
112	IPI00294193	1	36	0.7	0.14	51	68	ITIH4 Isoform 1 of Inter-alpha-trypsin inhibitor heavy chain H4 precu	AEAQAQYSAA	2	575.2804	1.364	0.155
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	GDYTLNNVHACAK	2	866.4189	4.832	0.563
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GDYTLNNVHACAK[272]	2	870.426	5.294	0.551
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	EVSSCHCAPCQGNGVPVLK	3	782.6833	4.543	0.488
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	KPYNVESYTPQTQGK	3	720.7198	4.824	0.404
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]EVSSCHCAPCQGNGVPVLK[272]	3	785.3547	4.723	0.512
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]RLPLEYSYGEYR	3	563.9621	2.326	0.231
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]VEPLYELVTATDFAYSSTVR	3	802.4167	4.892	0.437
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	GDYTLNNVHACAK	3	577.9484	2.895	0.307
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	KNTPIDGK	3	431.5944	3.063	0.267
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GDYTLNNVHACAK[272]	3	580.6198	3.301	0.27
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]YAYLLQPSQFHGEPCNFSDK[272]	3	893.4256	3.493	0.371
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]DTMVEDLVVLVR	2	766.9274	3.748	0.424
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]SGFSFGFK[272]	2	582.8183	2.939	0.332
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	VKVEPLYELVTATDFAYSSTVR	3	923.5005	2.936	0.409
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]CEGFVCAQTGR	2	703.7939	2.973	0.265
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]NTPIDGK[272]	2	516.8001	2.032	0.262
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]EYESYSDFER	2	734.8211	2.427	0.276
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	NTPIDGK	2	512.793	1.814	0.257
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GILNEIK[272]DR	3	449.2729	1.96	0.002
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	KNTPIDGK	2	646.8879	1.919	0.18
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]DTM[147]VEDLVVLVR	2	774.9248	2.032	0.217
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]YEFILK[272]	2	550.8334	2.264	0.073
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GINLFTNSFEGPVLDHR	3	687.3609	3.531	0.311
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GILNEIK[272]	2	537.8417	2.381	0.024
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]QALEEFQK[272]	2	640.8581	2.236	0.053
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	GIFELGISSQSDR	2	774.9045	3.702	0.439
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	YEFILK	2	546.8263	2.149	0.142
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GIFELGISSQSDR	2	776.908	3.399	0.388
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]GDYTLNNVHAC	2	698.8089	2.879	0.278
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]YYAGGCSPH	2	572.7373	2.15	0.321
113	IPI00294395	1	35.9	0.69	0.07	36	30	C8B Complement component C8 beta chain precursor	n[145]LEYSYGEYR	2	662.3205	2.21	0.108
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	IQYYCHEPYYK	2	916.9286	4.326	0.401
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]DYFIATCK[272]	2	647.8244	3.223	0.43
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]FCGQLGSPLGNPPGK[272]K[272]	3	693.3799	4.235	0.425
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]MDVFSQNM[147]FCAGHPSLK[272]	3	754.6855	3.963	0.501
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	FCGQLGSPLGNPPGKK	3	689.3728	2.952	0.375
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]CLPVCGK[272]PVNPVEQR	3	673.6799	2.766	0.155
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]IQYYCHEPYYK[272]	3	614.2929	3.456	0.275
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]TLDEFTIIQNLQPQYQFR	3	800.089	3.389	0.331
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	DYFIATCK	2	643.8173	2.631	0.352
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	IQYYCHEPYYK	3	611.6215	3.299	0.208
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]GGGALLGDR	2	480.2732	3.618	0.204
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]IQYYCHEPYYK[272]	2	920.9357	3.822	0.359
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]VLNYVDWIK[272]	2	719.4209	3.279	0.307

114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	CLPVCGKPVNPVEQR	3	671.0085	2.782	0.1
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]YTTEIIK[272]	2	578.3468	2.366	0.167
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]LGNHPIR	2	475.7863	2.354	0.16
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	NIGEFCGK	2	597.2939	2.304	0.186
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	LGNHPIR	2	473.7827	2.277	0.171
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]NIGEFCGK[272]	2	601.301	2.129	0.161
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	M[147]DVFSQNM[147]FCAGHPSLK	3	757.3457	2.377	0.17
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]EFM[147]SQGNK[272]	2	622.8128	1.901	0.178
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]NLPNGDFR	2	538.7839	1.765	0.022
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	MDVFSQNM[147]FCAGHPSLK	3	752.0141	4.144	0.386
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	VLNYVDWIK[272]	2	717.4173	2.962	0.242
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]QDACQGDSGGVFAVR	2	850.3861	2.362	0.233
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]LFGEVTSPLFPK[272]	2	811.9735	3.62	0.374
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	LFGEVTSPLFPK	2	807.9664	3.501	0.383
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]IAHDLR	2	434.7597	1.881	0.021
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]VSVHPDYR	2	558.7996	1.736	0.009
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]CLPVCGK[272]	2	550.2724	2.234	0.249
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	CLPVCGK	2	546.2653	2.086	0.222
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]PGYELQEDR	2	625.8103	2.491	0.181
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]LGNHPIR	3	317.5266	1.559	0.025
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	GSPLGNPPGKK	3	491.2963	2.415	0.132
114	IPI00296165	1	27.1	0.77	0.09	55	34	C1R;ACYP1;C17orf13 Complement C1r subcomponent precursor	n[145]PVNPVEQR	2	541.8074	1.467	0.18
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	n[145]SALVLQYLR	2	603.87	3.554	0.389
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	n[145]SCEPAVPFPCGR	2	749.8252	2.716	0.235
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	n[145]YGIYTK[272]	2	516.8021	2.207	0.24
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	GKYGIYTK	3	450.6029	2.629	0.155
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	n[145]CAGFHEGGR	3	375.168	2.78	0.259
115	IPI00296176	1	8.2	0.78	0.15	5	3	F9 Coagulation factor IX precursor	CAGFHEGGR	2	560.2448	1.838	0.057
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	GYHLNEEGTR	3	439.2178	2.718	0.267
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	GYHLNEEGTR	2	658.3231	3.381	0.441
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	n[145]TGYYFDGISR	2	661.8285	3.472	0.317
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	SAATLQQEK	2	628.3539	3.12	0.209
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	n[145]GYHLNEEGTR	2	660.3267	3.053	0.323
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	n[145]GYHLNEEGTR	3	440.5535	2.216	0.23
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	TGYYFDGISR	2	659.825	2.693	0.292
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	NVPNCGR	2	473.2233	1.645	0.029
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	SCPSSGYR	2	521.7259	2.538	0.179
116	IPI00296534	1	8	1.01	0.05	2	9	FBLN1 Isoform D of Fibulin-1 precursor	AITPPHPASQAN	2	672.357	1.83	0.085
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	SYTSHTNEIHK	2	798.9101	4.3	0.468
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]ELSHLPSLYDYSAYR	2	979.4925	5.162	0.386
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]GCPTEEGCGER	2	687.2574	3.447	0.49
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]NVVYTCNEGYSLIGNPVAR	2	1130.0546	5.72	0.523
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	SYTSHTNEIHK	3	532.9425	3.737	0.389
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]LIDQYGTHYLQSGSLGGEYR	3	801.0683	4.199	0.425
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]NVVYTCNEGYSLIGNPVAR	3	753.7055	3.914	0.33
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]SCVGETTESTQCEDEELEHLR	3	877.6981	8.093	0.642
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]SYTSHTNEIHK[272]	3	535.6139	4.534	0.385
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]ELSHLPSLYDYSAYR	3	653.3307	3.331	0.166

117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	VLFYVDSEK	2	690.3821	3.477	0.381
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]AASGTQNNVLR	2	637.8503	3.99	0.304
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]VLFYVDSEK[272]	2	694.3892	3.5	0.388
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]SYTSHTNEIHK[272]	2	802.9172	3.51	0.381
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]LTPLYELVK[272]	2	682.4256	2.795	0.296
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]DSCTLPASAEK[272]	2	728.3567	2.876	0.302
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]ILPLTVCK[272]	2	610.8711	2.861	0.3
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]ECNNPPPSGGGR	2	687.8042	3.09	0.296
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	LKQNDFNSVEEK	3	624.3389	4.249	0.244
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]EQTM[147]SECEAGALR	2	815.8532	2.878	0.334
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]LSGNVLSYTFQVK[272]	2	872.4978	3.409	0.304
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	QNDFNSVEEK	2	745.3677	3.397	0.238
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]VFSGDGK[272]	2	499.2815	2.053	0.292
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]LK[272]QNDFNSVEEK[272]	3	628.346	1.983	0.204
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	KVFSGDGK	2	629.3694	2.057	0.274
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	ILPLTVCK	2	606.864	2.412	0.197
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	EVPCASVK	2	579.8041	2.483	0.184
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	LTPLYELVK	2	678.4185	2.289	0.158
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]ELENALK[272]	2	552.8288	2.499	0.106
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]EVPCASVK[272]	2	583.8112	2.171	0.138
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]EQTMSECEAGALR	2	807.8558	3.395	0.318
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]YSAWAESVTNLPQVIK	3	697.385	3.487	0.112
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]SLVPTEFCPSPPALK[272]	2	960.5143	3.727	0.314
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]QNDFNSVEEK[272]	2	749.3748	1.847	0.116
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	DSCTLPASAEK	2	724.3496	1.273	0.096
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]SLVPTEFCPSPPALK[272]	3	640.6786	3.696	0.12
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	n[145]LLEPHCFPL	2	629.8259	2.082	0.281
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	FSSHGCKELENALK	3	677.0168	2.269	0.054
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	SVK[272]	1	617.4104	0.88	0.378
117	IPI00296608	1	31.8	0.66	0.08	54	38	C7 Complement component C7 precursor	EM[147]K	1	703.3808	1.121	0.072
118	IPI00297261	1	4.8	0.54	0	2	4	PTPN1 Tyrosine-protein phosphatase non-receptor type 1	FIMGDSSVQDQWK[272]	3	608.9739	3.285	0.133
118	IPI00297261	1	4.8	0.54	0	2	4	PTPN1 Tyrosine-protein phosphatase non-receptor type 1	FIMGDSSVQDQWK[272]	2	912.9572	2.807	0.01
118	IPI00297261	1	4.8	0.54	0	2	4	PTPN1 Tyrosine-protein phosphatase non-receptor type 1	n[145]GEPSLPEK[272]	2	572.8263	2.009	0.051
118	IPI00297261	1	4.8	0.54	0	2	4	PTPN1 Tyrosine-protein phosphatase non-receptor type 1	GEPSLPEK	2	568.8192	2.047	0.028
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	KDGTHVVENVDATHIGK	3	747.4097	6.428	0.569
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	n[145]DGTHVVENVDATHIGK[272]	3	660.6845	4.619	0.485
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	KDGTHVVENVDATHIGK	4	560.8091	2.928	0.417
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	n[145]STVLTIPEIIIK[272]	2	808.0179	3.36	0.417
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	GTYIPVPIVSELQSGK	3	656.7103	3.434	0.251
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	KETFDVTLEPLSFK	3	692.0576	3.303	0.201
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	n[145]GTYIPVPIVSELQSGK[272]	3	659.3817	2.783	0.237
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	n[145]FQEGQEEER	2	648.3029	2.757	0.169
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	FQEGQEEER	2	646.2993	2.743	0.211
119	IPI00297550	1	10.9	1.04	0.21	11	10	F13A1 Coagulation factor XIII A chain precursor	n[145]VDHHTDK[272]YENNK[272]	3	644.6694	2.437	0.145
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	QDGSVDFGR	1	1120.5382	2.635	0.384
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]QDGSVDFGR	1	1124.5453	2.462	0.446
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEANK	2	908.4605	5.355	0.536
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	DNENVVNEYSSELEK	2	1024.9922	6.266	0.527

120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLRPAPPPISGGGYR	2	1046.0527	3.64	0.47
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GGETSEM[147]YLIQPDSSVKPYR	2	1277.1364	4.691	0.561
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	NYCGLPGEYWLGNDK	2	1027.9768	4.569	0.524
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	QGFGNVATNTDGK	2	794.9076	4.579	0.297
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TM[147]TIHNGM[147]FFSTYDR	2	996.9508	3.904	0.493
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TPCTVSCNIPVVSGK	2	938.9612	3.247	0.376
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEANK[272]	2	912.4676	5.948	0.531
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]DNENVVNEYSSELEK[272]	2	1028.9993	6.476	0.551
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EDGGGWWYNR	2	692.3136	3.715	0.441
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLRPAPPPISGGGYR	2	1048.0563	4.113	0.483
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GGETSEM[147]YLIQPDSSVK[272]PYR	2	1281.1435	4.694	0.544
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]HQLYIDETVNSNIPTNLR	2	1136.0961	4.783	0.51
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]LESDVSAQMEYCR	2	860.8767	4.136	0.428
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]LESDVSAQM[147]EYCR	2	868.8741	3.452	0.502
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]MGPTELLIEMEDWK[272]	2	990.5067	4.709	0.446
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]NYCGLPGEYWLGNDK[272]	2	1031.9839	3.822	0.548
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]QGFGNVATNTDGK[272]	2	798.9147	4.644	0.407
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TMTIHNGM[147]FFSTYDR	2	990.9569	2.932	0.404
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGMFFSTYDR	2	990.9569	4.232	0.435
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TPCTVSCNIPVVSGK[272]	2	942.9683	3.44	0.475
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TVSCNIPVVSGK[272]	2	769.4197	3.835	0.477
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]VYCDM[147]NTENGGWTVIQNR	2	1153.5097	4.447	0.413
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEANK	3	605.9761	4.011	0.21
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEANKYQISVNK	3	930.1559	5.793	0.567
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	DNENVVNEYSSELEK	3	683.6639	3.186	0.313
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLRPAPPPISGGGYR	3	697.7042	3.908	0.532
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GGETSEMYLIQPDSSVKPYR	3	846.4283	3.93	0.264
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KGGETSEMYLIQPDSSVKPYR	3	935.825	4.712	0.294
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KGGETSEM[147]YLIQPDSSVKPYR	3	941.1566	5.851	0.41
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	NSVDELNNNVEAVSQTSSSSFQYMYLLK	3	1149.8961	4.902	0.464
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	NSVDELNNNVEAVSQTSSSSFQYM[147]YLLK	3	1155.2277	6.277	0.573
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TMTIHNGMFFSTYDR	3	654.3063	4.017	0.493
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEANK[272]	3	608.6475	2.103	0.164
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLRPAPPPISGGGYR	3	699.0399	3.204	0.41
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GGETSEMYLIQPDSSVK[272]PYR	3	849.0997	4.811	0.322
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]NSVDELNNNVEAVSQTSSSSFQYM[147]YLLK[27	3	1157.8991	6.071	0.559
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TMTIHNGMFFSTYDR	3	655.642	4.147	0.506
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TMTIHNGM[147]FFSTYDR	3	660.9737	3.235	0.516
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGM[147]FFSTYDR	3	666.3053	2.767	0.168
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KGGETSEMYLIQPDSSVKPYR	4	702.1206	2.699	0.318
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KGGETSEM[147]YLIQPDSSVKPYR	4	706.1193	4.759	0.493
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEANK[272]YQISVNK[272]	4	700.8741	5.246	0.495
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GGETSEM[147]YLIQPDSSVKPYR	3	851.76	2.392	0.21
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TM[147]TIHNGM[147]FFSTYDR	3	664.9696	2.881	0.337
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TPCTVSCNIPVVSGK	3	626.3099	2.45	0.27
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]HQLYIDETVNSNIPTNLR	3	757.7332	3.191	0.287
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]MGPTELLIEMEDWK[272]	3	660.6735	3.872	0.311
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]REEAPSLRPAPPPISGGGYR	3	751.0736	3.536	0.399

120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGMFFSTYDR	3	660.9737	2.777	0.337
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TPCTVSCNIPVVSGK[272]ECEEIIR	3	935.1126	3.785	0.345
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEANKYQISVNK	4	697.8687	4.233	0.382
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]REEAPSLRPAPPPISGGGYR	4	563.5571	3.131	0.334
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGM[147]FFSTYDR	2	998.9543	3.549	0.297
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KAPDAGGCLH	3	432.218	2.985	0.431
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]IQK[272]LESDVSAQM[147]EYCR	3	750.6984	3.85	0.41
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]YQISVNK[272]YR	3	486.9484	3.161	0.284
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]APDAGGCLHADPDLGVLCPTGCQLQEALLQQER	4	1017.486	3.65	0.4
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]QDGSVDFGR	2	562.7763	2.016	0.224
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]NSVDELNNNVEAVSQTSSSSFQYMYLLK[272]	3	1152.5675	3.826	0.26
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	IRPFFPQQ	2	586.8324	2.279	0.288
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	YQISVNK	1	1131.6522	2.524	0.209
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	ECEEIIR	1	1077.5068	2.295	0.165
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]ECEEIIR	2	541.2606	2.866	0.254
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GHRPLDK[272]K[272]	4	346.4703	1.983	0.16
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	YQISVNK	2	566.3297	2.679	0.178
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]CHAANPNGR	2	565.2592	2.269	0.321
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]IRPFFPQQ	2	588.836	2.031	0.171
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	SILENLR	1	984.5837	2.478	0.031
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEANK[272]YQISVNK[272]YR	5	624.7336	2.787	0.136
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	SILENLR	2	492.7955	2.612	0.169
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]ECEEIIR	1	1081.5139	2.224	0.099
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]YQISVNK[272]	2	570.3368	2.442	0.225
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]SILENLR	1	988.5908	2.465	0.025
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]SILENLR	2	494.799	2.897	0.075
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]YQISVNK[272]	1	1139.6664	2.443	0.211
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GHRPLDKK	3	457.6175	2.128	0.194
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GHRPLDK[272]K[272]	3	461.6246	1.833	0.092
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	QDGSVDFGR	2	560.7727	2.235	0.249
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GHRPLDKK	2	685.9226	2.285	0.076
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	REEAPSLR	3	366.5402	2.997	0.151
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	YQISVNKYR	3	484.277	3.472	0.182
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GHRPLDK[272]K[272]	2	691.9333	2.139	0.131
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]QLQEALLQQERPIR	3	622.6941	3.459	0.295
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TPCTVSCNIPVVSGK[272]	3	628.9813	2.557	0.07
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GHRPLDKK	4	343.465	1.415	0.083
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	LESDVSAQM[147]EYCR	3	578.2495	2.518	0.155
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]REEAPSLRPAPPPISGGGYR	2	1126.1068	2.423	0.088
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]YQISVNK[272]YR	2	729.919	2.648	0.253
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TPCTVSCNIPVVSGKECEEIIR	3	932.4412	2.948	0.227
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]ISQLTR	2	431.2674	2.296	0.101
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KGGETSEM[147]YLIQPDSSVK	3	802.4176	4.117	0.253
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	YQISVNKYR	2	725.9119	3.114	0.156
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]PAPPPISGGGYR	2	656.8602	1.971	0.067
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	CTVSCNIPVVSGK	3	560.2764	3.249	0.267
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TMTIHNGM[147]FFSTYDR	2	988.9533	4.224	0.548
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KAPDAGGCLHAD	3	494.2394	2.522	0.301

120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLRPAPPPISG	2	831.4526	2.962	0.257
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]DIWQK[272]	2	489.2866	1.968	0.004
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GFGNVATNTDGK[272]	2	734.8854	3.935	0.325
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]CTVSCNIPVVSGK[272]	2	843.9181	3.066	0.424
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]VNDNEEGFFSAR	2	764.8611	3.719	0.396
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLRPAPPPISGG	2	859.9633	3.223	0.389
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GFGNVATNTDGK	2	730.8783	3.578	0.243
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]MGPTELLIEM[147]EDWK[272]	3	666.0052	2.989	0.109
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]DNEEGFFSAR	2	658.3054	3.405	0.353
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KAPDAGGCLHAD	2	740.8554	2.173	0.345
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLRPAPPPISGG	2	857.9598	3.708	0.32
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEANK[272]YQISVNK[272]YR	3	1040.5511	2.685	0.357
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]REEAPSLR	3	367.8759	2.677	0.109
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GFTVQNEANK[272]	2	698.3772	3.395	0.261
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEA	2	719.3476	2.987	0.324
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLRPAPPPISGGG	2	888.4741	3.195	0.358
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLRPAPPPISG	2	829.4491	2.649	0.395
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	DLWQK	2	485.2795	1.732	0.049
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	HGTDDGVVWM[147]NWK[272]	3	615.6339	3.138	0.146
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GVNDNEEGFFSAR	2	793.3718	3.555	0.311
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]MGPTELLIEM[147]EDWK[272]	2	998.5041	3.143	0.236
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EYSSELEK	2	632.8247	2.709	0.279
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	PAPPPISGGGYR	2	654.8566	2.29	0.359
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	M[147]GPTELLIEM[147]EDWK[272]	2	1004.4981	3.256	0.208
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]NYCGLPGEYWLGNDK[272]	3	688.325	2.39	0.004
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	QVKDNENVVNEYSSELEK	3	848.7696	3.102	0.188
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNE	2	681.8255	2.512	0.263
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	REEAPSLR	2	549.3067	2.701	0.128
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEA	2	717.3441	2.372	0.286
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGFTVQNEANKYQISVNKYR	5	622.3293	2.695	0.053
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EAPSLRPAPPPISGGGYR	3	656.0257	2.729	0.254
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGM[147]FFSTY	2	863.3903	2.46	0.202
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGM[147]F	2	614.2846	1.955	0.296
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]REEAPSLR	2	551.3103	2.998	0.079
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	AHYGGF	2	396.1954	1.875	0.222
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	ECEEIIRK	3	449.2371	1.588	0.046
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	NYCGLPGEYWLGNDK[272]	2	1029.9804	2.315	0.273
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TMTIHNGM[147]F	2	606.2871	2.709	0.112
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	KAPDAGGCLH	2	647.8234	2.118	0.241
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNEAN	2	776.3691	2.353	0.158
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TVQNEANK	2	592.3252	2.364	0.217
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	GSVDFGR	2	439.23	2.195	0.093
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TM[147]TIHNGM[147]FF	2	685.8152	1.844	0.262
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]GSVDFGR	2	441.2335	2.374	0.11
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TM[147]TIHNGM[147]FFSTY	2	861.3868	2.168	0.267
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	CTVSCNIPVVSGK	2	839.911	3.3	0.4
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGMFF	2	679.8213	1.877	0.252
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	TM[147]TIHNGM[147]F	2	612.281	1.818	0.219

120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]APPPISGGGYR	2	608.3338	1.785	0.211
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]HGTDDGVVWM[147]NWK[272]	3	616.9696	2.455	0.13
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLR	1	945.5122	2.196	0.147
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]EEAPSLR	2	473.2597	2.327	0.046
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLR	1	941.5051	2.293	0.057
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]PPPISGGGYR	2	572.8152	2.236	0.082
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	M[147]GPTELLIEM[147]EDWK[272]GDKVK	3	939.1667	2.421	0.179
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGFTVQNE	2	683.8291	2.699	0.141
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]MYLLK[272]	2	478.2981	2.045	0.167
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]TM[147]TIHNGM[147]FF	2	687.8188	2.026	0.078
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	FSTYDR	2	464.7298	1.869	0.113
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	REEAPSLRPAPPPISGG	3	624.3426	2.12	0.172
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]IRPFFPQ	2	524.8067	1.621	0.08
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]SNIPTNLR	2	529.8074	2.172	0.076
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]AHYGGF	2	398.199	1.45	0.107
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]LESDVSAQM[147]EYCR	3	579.5852	2.112	0.002
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	n[145]HYGGFTVQNEANK[272]	3	584.9684	2.514	0.016
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	EEAPSLR	2	471.2562	1.69	0.03
120	IPI00298497	1	75.6	0.86	0.1	405	##	FGB Fibrinogen beta chain precursor	SVK[272]	1	617.4104	0.88	0.378
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	CSYTEDAQCIDGTIEVPK	2	1172.52	3.998	0.537
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TCPKPDDLPFSTVVPLK	2	1162.1313	4.358	0.462
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TFYEPGEEITYSCK	2	996.9577	4.894	0.566
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TFYEPGEEITYSCKPGYVSR	2	1326.6273	5.194	0.572
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]CSYTEDAQCIDGTIEVPK[272]	2	1176.5271	4.191	0.604
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TCPK[272]PDDLPFSTVVPLK[272]	2	1168.142	3.468	0.325
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TFYEPGEEITYSCK[272]	2	1000.9648	4.538	0.507
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TFYEPGEEITYSCK[272]PGYVSR	2	1330.6344	4.588	0.585
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VCPFAGILENGAVR	2	818.427	4.395	0.395
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	ATFGCHDGYSLDGPEEIECTK	3	882.0444	6.589	0.665
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TCPKPDDLPFSTVVPLK	3	775.09	4.772	0.445
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TFYEPGEEITYSCKPGYVSR	3	884.754	5.064	0.537
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]ATFGCHDGYSLDGPEEIECTK[272]	3	884.7158	2.124	0.013
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]GCHDGYSLDGPEEIECTK[272]	3	778.3314	5.223	0.463
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TCPK[272]PDDLPFSTVVPLK[272]	3	779.0971	4.157	0.478
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TFYEPGEEITYSCK[272]PGYVSR	3	887.4254	5.393	0.524
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]NGM[147]LHGDK[272]VSFFCK[272]	4	520.0153	2.528	0.447
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	KATVVYQGER	3	477.6087	2.509	0.244
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]EHSSLAFWK[272]	3	464.922	3.392	0.223
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	ATVVYQGER	2	581.8144	2.433	0.23
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	KATVVYQGER	2	715.9094	3.265	0.311
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	VCPFAGILENGAVR	2	816.4235	3.553	0.41
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	VSFFCK	2	528.7721	2.588	0.327
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]ATVVYQGER	2	583.818	2.676	0.174
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	EHSSLAFWK	3	462.2506	2.983	0.216
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]FK[272]NGM[147]LHGDK[272]	3	532.2961	2.406	0.356
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	EHSSLAFWK	2	692.8723	3.269	0.374
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	TDASDVKPC	2	631.2994	2.035	0.181
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]EHSSLAFWK[272]	2	696.8794	3.281	0.352

121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VSFFCK[272]	2	532.7792	2.374	0.34
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]NGMLHGDK[272]	3	387.2093	2.933	0.211
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]TDASDVK[272]PC	2	635.3065	2.135	0.038
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	KCSYTEDAQCIDGTIEVPK	3	871.4124	2.845	0.254
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	VPVKKATVVYQGER	4	499.3032	2.869	0.241
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VCPFAGILENGAVR	3	545.9538	3.332	0.165
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	CPFPSRPDNGFVNYPAKPTLY	3	903.7821	3.421	0.352
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]CTEEGK[272]	2	500.7377	2	0.254
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]CPFPSRPDNGFVNYPAK[272]PTLY	3	906.4535	3.778	0.307
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	CTEEGK	2	496.7306	2.025	0.182
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	GFVNYPAKPTLYYK	3	694.391	2.945	0.328
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]NGMLHGDK[272]	2	580.3103	1.787	0.225
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]ATVVYQGER	3	389.5477	2.11	0.148
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	ASCKVPVKK	4	392.2401	2.392	0.189
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	VPVKK	2	495.8448	1.377	0.197
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]AGILENGAVR	2	572.3338	3.674	0.279
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	AGILENGAVR	2	570.3302	2.956	0.249
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	VKIQEK	2	582.8768	2.357	0.026
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VK[272]IQEK[272]	3	392.9274	2.283	0.021
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VPVK[272]K[272]	2	501.8555	1.068	0.188
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	EHSSLAFWK[272]	2	694.8758	2.419	0.338
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]GFVNYPAK[272]PTLYYK[272]	3	698.3981	2.929	0.292
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VK[272]IQEK[272]	2	588.8875	1.819	0.055
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	FKNGM[147]LHGDK	3	528.289	1.997	0.146
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	LPFSTVVPLK	2	690.9343	2.678	0.251
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	SCKPGYVSR	3	441.5618	2.16	0.136
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]CSYTEDAQCIDGTIEVPK[272]	3	784.6872	2.055	0.021
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	YPAKPTLYYK	3	555.3239	1.737	0.171
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	KFICPL	2	523.7981	1.565	0.101
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]VYK[272]PSAG	2	505.2997	1.716	0.123
121	IPI00298828	1	55.4	0.83	0.07	120	56	APOH Beta-2-glycoprotein 1 precursor	n[145]ATFGCHDGY	2	580.7347	1.5	0.112
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]CTEGFNVDK[272]K[272]	2	809.9183	3.12	0.41
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]SIAQYWLGCPAPGHL	2	901.9456	1.595	0.091
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]CTEGFNVDK[272]K[272]	3	540.2813	3.202	0.142
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]DWHGVPGQVDAAM[147]AGR	3	609.629	4.981	0.522
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	AVRPGYPK	3	389.9048	2.575	0.27
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	CTEGFNVDKK	3	536.2742	2.8	0.309
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	DWHGVPGQVDAAM[147]AGR	3	608.2933	3.571	0.327
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]AVRPGYPK[272]	3	392.5762	2.304	0.372
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]SIAQYWLGCPAPGHL	3	601.6328	3.192	0.279
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]CTEGFNVDK[272]	2	673.8198	2.955	0.378
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]DWHGVPGQVDAAM[147]AGR	2	913.9399	2.423	0.291
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]GIPDNVDAALALPAHSYSGR	3	723.3787	4.575	0.383
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	SIAQYWLGCPAPGHL	2	899.942	4.337	0.415
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]IYISGM[147]APRPSLAK[272]	3	603.3496	1.966	0.046
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	DWHGVPGQVDAAMAGR	2	903.9389	3.748	0.433
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	AVRPGYPK	2	584.3535	2.974	0.252
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	CTDYTAECKPQVTR	3	662.9683	3.535	0.369

122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]AVRPGYPK[272]	2	588.3606	2.463	0.252
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	RVDTVDPPYPR	3	485.5965	2.764	0.186
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	CTEGFNVDK	2	669.8127	2.469	0.216
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	VYFFK	2	492.2893	1.735	0.167
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	CTEGFNVDKK	2	803.9077	3.969	0.171
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]RVDTVDPPYPR	3	486.9322	2.713	0.095
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]DVWGIEGPIDAAFTR	2	895.9633	3.265	0.512
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]FEDGVLDPDYPR	2	783.8815	2.38	0.226
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]CTDYTAECK[272]PQVTR	3	665.6397	3.061	0.241
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	TYLFK	2	476.2868	2.071	0
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]GQYCYELDEK[272]	2	791.362	4.259	0.322
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]VYFFK[272]GK[272]	2	660.9057	2.113	0.19
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]TYLFK[272]	2	480.2939	1.729	0.177
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	QPQFISR	2	508.2878	1.693	0.036
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	WLGCPAPGHL	2	618.8045	2.634	0.334
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]IYISGMAPR	2	576.3218	2.561	0.315
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]WLGCPAPGHL	2	620.808	2.482	0.359
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	GCPAPGHL	2	469.2228	2.039	0.306
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]VYFFK[272]GK[272]	3	440.9395	2.36	0.04
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]IYISGM[147]APR	2	584.3193	1.948	0.096
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	VRPGYPK	3	366.2257	2.57	0.086
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]GCPAPGHL	2	471.2263	1.735	0.164
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]VRPGYPK[272]	3	368.8971	2.306	0.118
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	RPGYPK	3	333.2029	2.036	0.171
122	IPI00298971	1	34.3	0.65	0.24	72	40	VTN Vitronectin precursor	n[145]VRPGYPK[272]	2	552.8421	1.751	0.138
123	IPI00299435	1	8.1	0.72	0.05	7	4	APOF apolipoprotein F precursor	SGVQQLIQYYQDQK	3	660.018	4.386	0.457
123	IPI00299435	1	8.1	0.72	0.05	7	4	APOF apolipoprotein F precursor	n[145]SGVQQLIQYYQDQK[272]	3	662.6894	4.622	0.323
123	IPI00299435	1	8.1	0.72	0.05	7	4	APOF apolipoprotein F precursor	n[145]DANISQPETTK[272]	2	746.3983	3.739	0.295
123	IPI00299435	1	8.1	0.72	0.05	7	4	APOF apolipoprotein F precursor	DANISQPETTK	2	742.3912	3.294	0.336
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	GIVAAFYSGPSLSDKEK	3	730.4037	4.998	0.454
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	n[145]GIVAAFYSGPSLSDK[272]EK[272]	3	734.4108	3.697	0.379
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	n[145]AQYVLISPEASSR	2	782.9262	3.672	0.365
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	IADVTSGLIGGEDGR	2	800.4205	3.937	0.398
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	n[145]IADVTSGLIGGEDGR	2	802.424	3.186	0.452
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	n[145]LYPTYSTK[272]	2	630.8576	2.516	0.166
124	IPI00299503	1	7.4	0.87	0.1	8	7	GPLD1 Isoform 1 of Phosphatidylinositol-glycan-specific phospholipa	n[145]NQVVIAAGR	2	536.3232	2.762	0.048
125	IPI00299778	1	4.5	0.95	0.04	3	3	PON3 Serum paraoxonase/lactonase 3	ILIGTVFHK	2	654.4136	3.096	0.179
125	IPI00299778	1	4.5	0.95	0.04	3	3	PON3 Serum paraoxonase/lactonase 3	n[145]STVEIFK[272]	2	556.3337	2.331	0.232
125	IPI00299778	1	4.5	0.95	0.04	3	3	PON3 Serum paraoxonase/lactonase 3	n[145]ILIGTVFHK[272]	3	439.2829	3.231	0.154
125	IPI00299778	1	4.5	0.95	0.04	3	3	PON3 Serum paraoxonase/lactonase 3	STVEIFK	2	552.3266	1.676	0.141
125	IPI00299778	1	4.5	0.95	0.04	3	3	PON3 Serum paraoxonase/lactonase 3	n[145]ILIGTVFHK[272]	2	658.4207	2.654	0.121
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]RNDYLDIYAIGVGK[272]	2	943.0248	4.34	0.495
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	ALHQVFEHMLDVSK	3	645.3488	4.757	0.432
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	ALHQVFEHM[147]LDVSK	3	650.6804	4.556	0.419
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	KNQGILEFYGDDIALLK	3	786.4458	4.331	0.399
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]ALHQVFEHMLDVSK[272]	3	648.0202	4.899	0.455
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]RNDYLDIYAIGVGK[272]	3	629.0189	4.135	0.397
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]AVISPGFDVFAK[272]	2	769.9447	3.693	0.295

126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	RNDYLDIYAIGVGK	3	626.3475	3.784	0.335
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]ALHQVFEHM[147]LDVSK[272]	3	653.3518	3.634	0.453
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	AVISPGFDVFAK	2	765.9376	3.816	0.308
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]GESGGAVFLER	2	633.334	3.85	0.379
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]HAIILLTDGK[272]	2	684.9263	3.213	0.395
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	DHENELLNK	2	696.3676	3.624	0.326
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	HAFILQDTK	2	676.8879	2.603	0.358
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	EILNINQK	2	626.3746	1.604	0.035
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	RNDYLDIYAIGVGK	2	939.0177	3.637	0.384
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	HAFILQDTK	3	451.5944	2.877	0.074
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]DFHINLFR	2	603.331	2.727	0.208
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]EILNINQK[272]	2	630.3817	2.486	0.004
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]HAFILQDTK[272]	2	680.895	2.642	0.247
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]VLMSVLNDNSR	2	696.3753	3.072	0.271
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	ELNELGSK	2	585.3299	2.15	0.138
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]DFHINLFR	3	402.5564	2.67	0.035
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]QHLGDVLNFLPL	2	755.4309	1.685	0.172
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]SCPQGLYPSPASR	2	776.8721	3.764	0.399
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]ELNELGSK[272]	2	589.337	1.757	0.036
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]DIAILK[272]	2	480.8203	1.803	1
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]DFHINLF	2	525.2805	1.727	0.093
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	SK[272]K[272]	2	395.7731	0.913	0.136
126	IPI00303963	1	20.9	0.75	0.05	40	26	C2 Complement C2 precursor (Fragment)	n[145]VFAK[272]K[272]	2	512.8476	1.081	0.047
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LGEVNTYAGDLQK	2	844.4544	3.721	0.491
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LNHQLEGLTFQM[147]K	2	927.9984	4.596	0.476
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	SELTQQLNALFQDK	2	958.0179	5.248	0.418
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	SLAPYAQDTQEK	2	815.9254	4.434	0.377
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	TQVNTQAEQLR	2	714.3837	3.594	0.329
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	VNSFFSTFK	2	678.8692	3.514	0.453
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]DK[272]VNSFFSTFK[272]	2	876.4883	3.855	0.43
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LGEVNTYAGDLQK[272]	2	848.4615	4.84	0.492
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LNHQLEGLTFQM[147]K[272]	2	932.0055	4.781	0.455
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]SELTQQLNALFQDK[272]	2	962.025	5.797	0.477
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]SLAELGGHLDQQVEEFR	2	1036.5301	5.457	0.58
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]VEPYGENFNK[272]	2	742.8849	3.807	0.434
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]VNSFFSTFK[272]	2	682.8763	3.489	0.456
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	DKVNSFFSTFK	3	580.6542	1.966	0.183
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	KLVPFATELHER	3	574.0017	3.902	0.396
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LGPHAGDVEGHLSFLEK	3	696.0409	3.248	0.454
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LNHQLEGLTFQMK	3	613.6697	4.61	0.35
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LNHQLEGLTFQM[147]K	3	619.0014	4.682	0.493
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LVPFATELHER	3	484.605	4.923	0.453
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]DK[272]VNSFFSTFK[272]	3	584.6613	4.255	0.464
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LGPHAGDVEGHLSFLEK[272]	3	698.7123	4.094	0.412
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LNHQLEGLTFQMK[272]	3	616.3411	4.275	0.392
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LNHQLEGLTFQM[147]K[272]	3	621.6728	4.357	0.332
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]RVEPYGENFNK[272]	3	547.626	4.476	0.357
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]SLAELGGHLDQQVEEFR	3	691.3558	5.635	0.548

127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]ENADSLQASLRPHADELK[272]	3	761.4042	4.344	0.255
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LGPHAGDVEGHLSFLEK[272]	4	524.286	2.747	0.296
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	EAVEHLQK	2	617.3512	3.055	0.252
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]RVEPYGENFNK[272]	2	820.9354	4.105	0.319
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]TQVNTQAEQLR	2	716.3873	3.897	0.251
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	ENADSLQASLRPHADELK	3	758.7328	4.1	0.247
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	RVEPYGENFNK	2	816.9283	3.964	0.307
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]EAVEHLQK[272]	2	621.3583	3.404	0.267
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	KNAEELK	3	417.9189	2.821	0.192
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LLPHANEVSQK[272]	3	508.631	2.928	0.193
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	ALVQQM[147]EQLR	2	686.3743	2.158	0.207
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	GNTEGLQK	2	563.8144	3.14	0.276
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]GNTEGLQK[272]	2	567.8215	2.806	0.265
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]IDQNVEELK[272]	2	688.3872	3.563	0.266
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	EAVEHLQK	3	411.9032	3.122	0.171
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LK[272]EEIGK[272]ELEELR	3	673.4001	3.349	0.257
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LVPFATELHER	2	726.4039	3.129	0.282
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LLPHANEVSQK[272]	2	762.4429	3.781	0.192
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LLPHANEVSQK	2	758.4358	3.034	0.245
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LAPLAEDVR	2	564.3307	2.768	0.353
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LK[272]EEIGK[272]ELEELR	4	505.3019	2.675	0.245
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	IDQNVEELK	2	684.3801	3.511	0.201
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]ALVQQM[147]EQLR	2	688.3779	3.01	0.238
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LKEEIGKELEELR	3	669.393	3.327	0.382
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]EAVEHLQK[272]	3	414.5746	2.858	0.143
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	DKVNSFFSTFK	2	870.4777	4.097	0.379
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]SLAPYAQDTQEK[272]	2	819.9325	3.277	0.297
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LLPHANEVSQK	3	505.9596	3.223	0.13
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LVPFATELHER	2	728.4075	2.396	0.308
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	VEPYGENFNK	2	738.8778	2.818	0.253
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	KNAEELK	2	626.3747	2.607	0.177
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]SLAELGGHLDQQVEEFRR	4	557.794	2.048	0.137
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	KLVPFATELHER	2	860.4989	3.436	0.396
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]ISASAEELR	2	560.31	3.128	0.138
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LAPLAEDVR	2	562.3271	2.137	0.302
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LTPYADEFK	2	682.3665	2.645	0.146
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	QLTPYAQR	2	558.8117	2.36	0.161
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]QLTPYAQR	2	560.8152	2.425	0.136
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LTPYADEFK[272]	2	686.3736	2.594	0.156
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]IGDNLR	2	416.2439	2.512	0.094
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]ALVQQMEQLR	2	680.3804	2.815	0.147
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	RVEPYGENFNK	3	544.9546	1.951	0.154
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]IDQTVEELRR	2	701.8922	2.305	0.095
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	IDQTVEELRR	2	699.8886	2.168	0.07
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LKEEIGK	2	618.8874	3.068	0.174
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LEPYADQLR	2	622.8353	1.688	0.025
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]LNHQLEGLTF	2	658.36	2.387	0.292
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	AKIDQNVEELK	3	569.6649	1.783	0.155

127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LKEEIGKELEELR	4	502.2966	1.623	0.017
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]IDQTVEELRR	3	468.2639	2.009	0.03
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	LKEEIGK	3	412.9274	2.127	0.074
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	ALFQDK	2	501.2926	2.179	0.087
127	IPI00304273	1	70.2	0.92	0.07	114	78	APOA4 Apolipoprotein A-IV precursor	n[145]ALFQDK[272]	2	505.2997	2.387	0.021
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	ETAVDGELVVLYDVK	2	965.5303	4.978	0.498
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	IYGNQDTSSQLK	2	817.4309	4.238	0.453
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]AEDHFSVIDFNQNIR	2	974.9853	4.223	0.45
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]ETAVDGELVVLYDVK[272]	2	969.5374	4.777	0.497
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]FYNQVSTPLLR	2	741.4153	1.456	0.062
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IQPSGGTNINEALLR	2	863.9821	4.084	0.389
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IYGNQDTSSQLK[272]	2	821.438	4.751	0.415
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	FLHVPDTFEGHFDGVPVISK	3	841.1128	2.902	0.316
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	KFYNQVSTPLLR	3	582.6736	3.667	0.38
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	KLGSYEHR	3	423.9069	3.466	0.417
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	VVNNSPQPQNVVFDVQIPK	3	801.4446	3.565	0.401
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]FLHVPDTFEGHFDGVPVISK[272]	3	843.7842	4.675	0.319
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]VQFELHYQEVK[272]	3	569.9818	4.166	0.265
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]VVNNSPQPQNVVFDVQIPK[272]	3	804.116	3.131	0.318
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]AHVSFK[272]PTVAQQR	4	440.007	3.743	0.453
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]FLHVPDTFEGHFDGVPVISK[272]	4	633.09	2.897	0.411
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	AHVSFKPTVAQQR	3	583.6688	3.298	0.29
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]RLSNENHGIAQR	3	513.6138	3.381	0.315
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	AHVSFKPTVAQQR	4	438.0034	2.453	0.277
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	TEVNVLPGAK	2	654.3878	1.922	0.099
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LSNENHGIAQR	2	691.8665	3.863	0.277
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	LSNENHGIAQR	3	460.2444	2.826	0.268
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LGSYEHR	3	335.846	2.75	0.225
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]NFHNYFGGSEIVVAGK[272]	3	676.3578	3.561	0.428
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]SSALDMENFR	2	657.3175	3.465	0.352
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]TEVNVLPGAK[272]	2	658.3949	3.005	0.268
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]VQSTITSR	2	518.2994	2.802	0.33
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	IYGNQDTSSQLKK	3	634.6863	3.387	0.197
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LSNENHGIAQR	3	461.5801	2.07	0.088
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IYGNQDTSSQLK[272]K[272]	3	638.6934	3.489	0.251
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	IQPSGGTNINEALLR	2	861.9785	3.024	0.35
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	VQFELHYQEVK	2	850.462	3.859	0.292
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]SSALDM[147]ENFR	2	665.3149	1.799	0.102
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	DKHADPDFTR	3	494.5881	2.293	0.294
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]VQFELHYQEVK[272]	2	854.4691	3.7	0.228
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	M[147]ATTMIQSK	2	653.8466	2.733	0.288
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IYLQPGR	2	495.7963	2.136	0.126
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]ALYAQAR	2	468.7728	2.169	0.127
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	M[147]ATTM[147]IQSK	2	661.844	2.446	0.287
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LGSYEHR	2	503.2654	2.095	0.069
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]DK[272]HADPDFTR	3	497.2595	2.557	0.194
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	SLAPTAAAK	2	555.3375	3.043	0.141
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]SLAPTAAAK[272]	2	559.3446	2.918	0.149

128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]HADPDFTR	2	551.7735	2.059	0.254
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]FVHFFAPDNLDPIPK[272]	3	682.3748	4.459	0.255
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IQPSGGTNINEALLR	3	576.3238	3.617	0.095
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	KFYNQVSTPLLR	2	873.5067	2.786	0.236
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	IYLQPGR	1	986.5782	1.789	0.158
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]AHVSFK[272]PTVAQQR	3	586.3402	2.322	0.199
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	MLADAPPQDPSCCSGALYYGSK	3	882.7225	2.674	0.196
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	LGSYEHR	1	1001.5164	1.896	0.122
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]NDLISATK[272]	2	575.3395	3.15	0.212
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	ALYAQAR	1	932.5313	1.941	0.155
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	NDLISATK	2	571.3324	1.603	0.095
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	FVHFFAPDNLDPIPK	3	679.7034	2.99	0.247
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	KLGSYEHR	2	635.3568	2.099	0.128
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]IYLQPGR	1	990.5853	1.759	0.123
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	NFHNYFGGSEIVVAGK	3	673.6864	3.533	0.265
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]ALYAQAR	1	936.5384	1.827	0.177
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]DK[272]HADPDFTR	2	745.3856	1.487	0.137
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	TILDDLR	1	985.5677	1.891	0.092
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	ALYAQAR	2	466.7693	2.068	0.116
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	DKHADPDFTR	2	741.3785	1.966	0.201
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]QTVEAM[147]K[272]	2	555.807	1.931	0.106
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]TILDDLRAEDHFSVIDFNQNIR	4	694.61	2.156	0.151
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]FLHVPDTFEGHFD	3	568.9428	2.684	0.249
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]NVQFNYPH	2	581.7917	2.13	0.355
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LTINQLLAER	2	657.8967	3.229	0.206
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	DPDFTR	2	445.722	1.821	0.22
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]DPDFTR	2	447.7255	1.924	0.186
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]DK[272]HADPDFTR	4	373.1964	1.493	0.118
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LYAQAR	2	433.2543	1.99	0.021
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]PTVGELK[272]	2	516.3206	1.765	0.119
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]LGSYEHR	1	1005.5235	1.741	0.006
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	SVK[272]	1	617.4104	0.88	0.378
128	IPI00305461	1	35.2	0.74	0.12	141	71	ITIH2 Inter-alpha-trypsin inhibitor heavy chain H2 precursor	n[145]DFLK[272]	1	810.4964	1.211	0.02
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	GDATVFFILPNQGK	2	893.9962	3.269	0.389
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	LGFTDLFSK	2	654.3716	3.62	0.319
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]GDATVFFILPNQGK[272]	2	898.0033	2.963	0.381
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	IVDLVSELKK	3	521.9993	3.734	0.407
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]IVDLVSELK[272]K[272]	3	526.0064	1.633	0.098
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]VGSALFLSHNLK[272]	3	525.315	4.151	0.355
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]GFQHLLHTLNLPGHGLETR	4	571.8163	3.335	0.328
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	FYYLIASETPGK	2	834.9535	3.779	0.395
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	VGSALFLSHNLK	2	783.4618	4.447	0.361
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]LGFTDLFSK[272]	2	658.3787	3.094	0.387
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]VGSALFLSHNLK[272]	2	787.4689	3.847	0.409
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	VGSALFLSHNLK	3	522.6436	3.718	0.203
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]FYYLIASETPGK[272]	2	838.9606	3.99	0.241
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]IAPANADFAFR	2	668.8602	3.072	0.339
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]WADLSGITK[272]	2	639.8685	2.91	0.252

129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	IVDLVSELKK	2	782.4953	3.426	0.173
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]GFQHLLHTLNLPGHGLETR	5	457.6545	3.237	0.208
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]DFYVDENTTVR	2	751.8658	2.69	0.248
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]IVDLVSELK[272]K[272]	2	788.506	2.302	0.26
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	IAPANADFAFR	2	666.8566	2.321	0.221
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]YLPCSVLR	2	570.8049	2.18	0.108
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	VGSALFLSH	2	535.8033	2.689	0.372
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	ALWEK[272]PFISSR	3	539.9793	2.316	0.128
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	n[145]VGSALFLSH	2	537.8069	2.389	0.117
129	IPI00328609	1	32.1	0.9	0.15	40	24	SERPINA4 Kallistatin precursor	LNDTMAVYEAK	2	767.9004	2.108	0.091
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]FQSGQVLAALPR	2	715.9155	3.199	0.338
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]SFYANNHCIGTDLNR	3	638.9618	3.811	0.39
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	SKDHEELSLVASEAVR	3	684.0358	3.85	0.383
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]SK[272]DHEELSLVASEAVR	3	686.7072	3.462	0.358
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]EAFAAVSK[272]	2	555.8235	2.467	0.204
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	EAFAAVSK	2	551.8164	2.303	0.225
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]IHIGSSFEK[272]	3	435.9184	2.662	0.162
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]DTGTYGFLLPER	2	756.8944	2.63	0.18
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]IHIGSSFEK[272]	2	653.3739	2.741	0.126
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	HPDM[147]LTK	3	379.8744	2.418	0.128
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	IHIGSSFEK	2	649.3668	2.676	0.075
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]YIK[272]PTCR	2	607.8351	1.941	0.125
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	YIKPTCR	2	603.828	2.002	0.087
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	n[145]YIK[272]PTCR	3	405.5591	2.18	0.006
130	IPI00329775	1	22.5	0.83	0.04	23	14	CPB2 Isoform 1 of Carboxypeptidase B2 precursor	IVFPYSYTR	2	643.3506	1.966	0.027
131	IPI00386502	1	3.9	7.14	0	2	3	BLZF1 Isoform 2 of Golgin-45	AM[147]EVK	2	437.2468	1.996	0.156
131	IPI00386502	1	3.9	7.14	0	2	3	BLZF1 Isoform 2 of Golgin-45	AM[147]EVK[272]	2	439.2503	2.095	0.111
131	IPI00386502	1	3.9	7.14	0	2	3	BLZF1 Isoform 2 of Golgin-45	AM[147]EVK[272]	1	877.4934	1.755	0.105
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	VGAHAGEYGAEALER	2	835.4183	3.652	0.519
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	n[145]TYFPHFDLSHGSAQVK[272]	3	708.0369	4.664	0.444
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	TYFPHFDLSHGSAQVK	4	529.2759	3.298	0.304
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	TYFPHFDLSHGSAQVK	3	705.3655	3.934	0.32
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	M[147]FLSFPTTK	2	684.3733	2.793	0.36
132	IPI00410714	1	33.1	4.26	0.35	6	6	HBA2;HBA1 Hemoglobin subunit alpha	VLSPADK	2	505.3057	1.882	0.178
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	AAHEEICTTNEGVM[147]YR	2	1013.445	4.249	0.543
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	DLQFVEVTDVK	2	786.9353	4.335	0.397
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	EATIPGHLNSYTIK	2	912.5044	4.619	0.452
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	EESPLLIGQQSTVSDVPR	2	1048.0551	6.208	0.553
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GATYNIIVEALK	2	786.4615	4.644	0.398
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	HTSVQTTSSGSGPFTDVR	2	1002.4927	5.367	0.541
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	NLQPASEYTVSLVAIK	2	1007.0726	5.407	0.533
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	SSPVVIDASTAIDAPSNLR	2	1027.0499	4.513	0.553
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	SYTITGLQPGTDYK	2	912.4806	5.245	0.446
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VPGTSTSATLTGLTR	2	801.4465	3.837	0.523
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YSFCTDHTVLVQTR	2	928.4451	4.408	0.526
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]DLQFVEVTDVK[272]	2	790.9424	3.817	0.421
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]ESK[272]PLTAQQTTK[272]	2	882.515	3.731	0.441
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	AAHEEICTTNEGVM[147]YR	3	675.9657	4.757	0.473

133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	AAVYQPQPHPQPPPYGH	3	675.3428	4.483	0.53
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	CDPHEATCYDDGK	3	609.2386	3.275	0.526
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	CHEGGQSYK	3	445.5375	3.466	0.455
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	EATIPGHLNSYTIK	3	608.672	3.73	0.46
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ESKPLTAQQTTK	3	584.672	1.419	0.031
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	FGFCPM[147]AAHEEICTTNEGVM[147]YR	3	924.0469	5.538	0.644
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GDSPASSKPISINYR	3	624.6707	3.871	0.432
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GLKPGVVYEGQLISIQQYGHQEVTR	3	1027.2278	7.563	0.57
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	HTSVQTTSSGSGPFTDVR	3	668.6642	5.394	0.497
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	NTFAEVTGLSPGVTYYFK	3	758.7317	4.889	0.347
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	RPGGEPSPEGTTGQSYNQYSQR	3	846.0652	6.924	0.653
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VDVIPVNLPGEHGQR	3	590.6601	3.558	0.359
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	WCHDNGVNYK	3	521.2395	2.992	0.413
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]EATIPGHLNSYTIK[272]	3	611.3434	3.799	0.275
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]ESK[272]PLTAQQTTK[272]	3	588.6791	2.848	0.287
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]GFNCESK[272]PEAEETCFDK[272]	3	820.0302	4.247	0.517
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]VGDTYERPK[272]	3	451.5853	3.701	0.335
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GLKPGVVYEGQLISIQQYGHQEVTR	4	770.6727	5.478	0.469
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]GLK[272]PGVVYEGQLISIQQYGHQEVTR	4	772.6762	6.638	0.482
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	IYLYTLNDNAR	2	748.3988	3.637	0.35
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	DTLTSRPAQGVVTTLENVSPPR	3	826.7819	4.004	0.299
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GATYNIIVEALKDQQR	3	700.3918	3.99	0.404
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	LGVRPSQGGEAPR	3	488.6074	3.047	0.265
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TEIDKPSQMQVTDVQDNSISVK	3	961.5025	5.157	0.304
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TEIDKPSQM[147]QVTDVQDNSISVK	3	966.8341	3.321	0.315
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VGDTYERPK	3	448.9139	3.126	0.264
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YSFCTDHTVLVQTR	3	619.2992	2.082	0.29
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]NTFAEVTGLSPGVTYYFK[272]	3	761.4031	3.81	0.26
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YQCYCYGR	2	644.2425	2.719	0.44
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]GATYNIIVEALK[272]	2	790.4686	4.168	0.361
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TYHVGEQWQK[272]	3	520.6078	3.672	0.325
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	FTNIGPDTMR	2	646.3268	3.279	0.301
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	LGVRPSQGGEAPR	2	732.4075	2.996	0.404
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	STTPDITGYR	2	625.8224	2.972	0.404
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TFYSCTTEGR	2	675.7945	2.586	0.445
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VDVIPVNLPGEHGQR	2	885.4865	3.693	0.389
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YEVSVYALK	2	676.3847	2.987	0.333
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	CHEGGQSYK	2	667.8027	2.822	0.381
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	FTNIGPDTM[147]R	2	654.3243	2.025	0.012
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VGDTYERPK	2	672.8672	2.957	0.289
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VFAVSHGR	2	506.788	2.684	0.225
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]VGDTYERPK[272]	2	676.8743	2.496	0.256
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GFNCESKPEAEETCFDK	3	816.0231	2.469	0.334
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	DAPIVNK	2	518.8112	2.456	0.241
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]YEK[272]PGSPPR	2	659.8715	2.53	0.262
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	AQITGYR	2	474.7667	2.78	0.109
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ISCTIANR	2	532.273	2.371	0.204
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VTTTPK	2	463.7871	1.887	0.256

133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]ITGYIIK[272]	2	548.3545	2.603	0.139
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	AQITGYR	1	948.5262	1.879	0.158
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YTGNTYR	2	507.7538	2.094	0.198
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TYHVGEQWQK	3	519.2721	3.303	0.089
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]WLPSSSPVTGYR	2	747.3971	3.24	0.289
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ESKPLTAQQTTK	2	876.5044	1.284	0.052
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YEKPGSPPR	2	655.8644	2.521	0.184
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ATITGYR	2	461.2613	2.782	0.315
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YVHGVR	2	435.7509	2.204	0.09
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ITGYIIK	2	544.3474	2.261	0.127
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]DAPIVNK[272]	2	522.8183	2.188	0.103
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VTWAPPPSIDLTNFLVR	3	689.386	2.379	0.182
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	SDTVPSPR	2	499.7669	1.754	0.001
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	IGDQWDK	2	571.3037	1.948	0.097
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TYHVGEQWQK[272]	2	780.408	2.942	0.268
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	YILR	2	352.7263	1.622	0.012
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ALVCTCYGGSR	2	681.2953	2.418	0.367
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	TIKPDVR	2	554.8455	1.495	0.064
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]IGDQWDK[272]	2	575.3108	1.785	0.009
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	IGDTWSK	2	543.8008	1.646	0.01
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	EATIPGHLNSY	2	671.3435	2.706	0.283
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	ITCTSR	2	433.7148	1.323	0.039
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	PQAPITGYR	2	571.8195	2.666	0.201
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	GGNSNGALCHFPF	2	753.8345	2.146	0.363
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	VTPKEK	2	561.3557	1.815	0.078
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	n[145]WLPSSSPVTGYR	3	498.6005	2.09	0
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	SK[272]K[272]	2	395.7731	0.913	0.136
133	IPI00414283	1	28.5	2.07	0.59	145	86	FN1 fibronectin 1 isoform 4 preproprotein	SVK[272]	1	617.4104	0.88	0.378
134	IPI00465436	1	8.3	1.13	0.01	4	4	CAT Catalase	DPILFPSFIHSQK	3	603.6735	4.112	0.311
134	IPI00465436	1	8.3	1.13	0.01	4	4	CAT Catalase	n[145]DPILFPSFIHSQK[272]	3	606.3449	3.277	0.281
134	IPI00465436	1	8.3	1.13	0.01	4	4	CAT Catalase	GAGAFGYFEVTHDITK	3	665.0108	2.991	0.257
134	IPI00465436	1	8.3	1.13	0.01	4	4	CAT Catalase	LGPNYLHIPVNCPYR	3	647.999	2.079	0.018
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]YVTSAPM[147]PEPQAPGR	2	880.9415	3.551	0.496
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]GVALHRPDVYLLPPAR	3	640.043	4.03	0.43
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	DVMQGTDEHVVCK	3	596.281	2.813	0.42
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]LICQATGFSPR	2	691.8557	3.36	0.354
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]VSVFVPPR	2	522.8198	2.287	0.346
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	GVALHRPDVYLLPPAR	3	638.7073	3.419	0.384
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]GQPLSPEK[272]	2	572.3343	2.632	0.271
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]DGFFGNPR	2	527.2654	1.942	0.159
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	GQPLSPEK	2	568.3272	2.526	0.287
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	DGFFGNPR	2	525.2618	2.152	0.04
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]GFPSVLR	2	460.2777	2.301	0.171
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	ESGPTTYK	2	581.8088	1.866	0.142
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]DVMQGTDEHVVCK[272]	3	598.9524	2.486	0.141
135	IPI00477090	1	15.7	0.79	0.09	25	13	IGHM IGHM protein	n[145]ESGPTTYK[272]	2	585.8159	2.021	0.112
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	FDHVITNM[147]NNNYEPR	2	1010.4707	4.395	0.502
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]FDHVITNMNNNYEPR	2	1004.4768	4.294	0.553

136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]FDHVITNM[147]NNNYEPR	2	1012.4743	4.576	0.511
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]GLYYFTYHASSR	2	804.9	3.853	0.544
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]LEQGENVFLQATDK[272]	2	940.5039	6.008	0.416
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]FDHVITNMNNNYEPR	3	669.987	3.999	0.469
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]FDHVITNM[147]NNNYEPR	3	675.3186	4.509	0.311
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]GLYYFTYHASSR	3	536.9358	3.663	0.505
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]VPGLYYFTYHASSR	3	602.3095	4.235	0.492
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	LEQGENVFLQATDK	2	936.4968	4.155	0.409
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]GNLCVNLM[147]R	2	613.3022	2.713	0.194
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]IAFSATR	2	455.2674	2.084	0.148
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]TINVPLR	2	478.8041	2.295	0.098
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]VPGLYYFTYH	2	702.3595	3.448	0.389
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	n[145]LEQGENVFLQATDK	2	938.5003	2.936	0.118
136	IPI00477992	1	28.6	0.8	0.04	19	16	C1QB complement component 1	VPGLYYFTYH	2	700.3559	2.311	0.247
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AIGYLNTGYQR	1	1395.738	3.001	0.369
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AAQVTIQSSGTFSSK	2	896.4836	5.386	0.519
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AFALAGNQDK	2	657.8619	3.909	0.394
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AIGYLNTGYQR	2	698.3726	3.725	0.438
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AVLPTGDVIGDSAK	2	811.9593	4.454	0.485
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DMYSFLEDMGLK	2	864.9204	5.256	0.382
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DM[147]YSFLEDMGLK	2	872.9179	4.565	0.4
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DM[147]YSFLEDM[147]GLK	2	880.9154	3.463	0.551
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DTVIKPLLVEPEGLEK	2	1100.6513	4.275	0.475
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ETTFNSLLCPSGGEVSEELSLK	2	1333.6563	4.998	0.488
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GITYTPVSSTNEK	2	838.9464	4.743	0.469
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLLQQVSLPELPGEYSMK	2	1163.148	4.293	0.521
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLLQQVSLPELPGEYSM[147]K	2	1171.1455	4.607	0.503
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLIYAVLPTGDVIGDSAK	2	1063.117	5.551	0.563
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LTAQPAPTSEDLTSATNIVK	2	1169.1367	6.299	0.542
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LVHVEEPHTETVR	2	843.4521	4.393	0.403
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	M[147]VSGFIPLKPTVK	2	927.0578	2.248	0.234
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	NALFCLESAWK	2	804.4073	3.765	0.425
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	NALFCLESAWK[272]	2	806.4108	3.733	0.329
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QFSFPLSSEPFQGSYK	2	1065.0388	4.921	0.517
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLFFTVLQDVPVR	2	830.9747	4.032	0.488
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSSNEEVMFLTVQVK	2	989.5194	4.364	0.522
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSSNEEVM[147]FLTVQVK	2	997.5168	5.598	0.522
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSTQDTVVALHALSK	2	919.0126	5.939	0.543
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TAQEGDHGSHVYTK	2	905.4476	2.102	0.276
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TEHPFTVEEFVLPK	2	977.0277	4.865	0.505
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TEVSSNHVLIYLDK	2	949.5228	5.754	0.539
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VDLSFSPSQSLPASHAH	2	960.4842	5.456	0.559
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VSVQLEASPAFLAVPVEK	2	1082.6225	6.697	0.526
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VTGEGCVYLQTSLK	2	912.4734	5.139	0.44
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VYDYYETDEFAIAEYNAPCSK	2	1409.1248	5.953	0.618
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YDVENCLANK	2	747.8576	4.475	0.358
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YILNGGTLLGLK	2	771.4744	4.545	0.439
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YSDASDCHGEDSQAFCEK	2	1182.4554	4.985	0.641

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AAQVTIQSSGTFSSK[272]	2	900.4907	5.567	0.42
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AIGYLNTGYQR	2	700.3762	2.433	0.082
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ALLAYAFALAGNQDK[272]	2	927.5218	4.526	0.422
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ATVLNYLPK[272]	2	653.9023	1.422	0.077
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AVLPTGDVIGDSAK[272]	2	815.9664	4.545	0.532
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DMYSFLEDMGLK[272]	2	868.9275	4.838	0.401
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DMYSFLEDM[147]GLK[272]	2	876.925	4.138	0.43
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DM[147]YSFLEDMGLK[272]	2	876.925	4.15	0.469
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DTVIK[272]PLLVEPEGLEK[272]	2	1106.6619	5.104	0.538
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ETTFNSLLCPSGGEVSEELSLK[272]	2	1337.6634	5.364	0.47
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GITYTPVSSTNEK[272]	2	842.9535	4.348	0.485
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]HNVYINGITYTPVSSTNEK[272]	2	1213.1337	5.988	0.476
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLLQQVSLPELPGEYSMK[272]	2	1167.1551	4.097	0.466
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLLQQVSLPELPGEYSM[147]K[272]	2	1175.1526	4.636	0.437
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLIYAVLPTGDVIGDSAK[272]	2	1067.1241	4.811	0.558
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LTAQPAPTSEDLTSATNIVK[272]	2	1173.1438	5.718	0.602
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MCPQLQQYEMHGPEGLR	2	1104.003	5.786	0.58
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MVSGFIPLK[272]PTVK[272]	2	925.0709	3.962	0.447
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]NALFCLESAWK[272]	2	808.4144	3.894	0.406
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]NEDSLVFVQTDK[272]	2	841.9456	4.448	0.3
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QFSFPLSSEPFQGSYK[272]	2	1069.0459	4.952	0.516
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLFTDLEAENDVLH	2	873.937	5.165	0.538
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLFTDLEAENDVLHCVA	2	1033.4882	4.771	0.386
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLFTDLEAENDVLHCVAFAVPK[272]	2	1376.7001	4.566	0.601
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSSNEEVMFLTVQVK[272]	2	993.5265	5.335	0.445
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSSNEEVM[147]FLTVQVK[272]	2	1001.5239	5.759	0.504
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSTQDTVVALHALSK[272]	2	923.0197	5.743	0.53
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TAQEGDHGSHVYTK[272]	2	909.4547	2.842	0.353
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TEHPFTVEEFVLPK[272]	2	981.0348	4.439	0.48
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TEVSSNHVLIYLDK[272]	2	953.5299	6.083	0.491
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASH	2	858.4397	3.958	0.506
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAH	2	962.4877	4.998	0.398
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAHLR	2	1097.0803	6.995	0.408
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VGFYESDVMGR	2	702.3409	2.534	0.168
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VLVPSLLHTETTEK[272]	2	928.0426	4.479	0.424
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VSVQLEASPAFLAVPVEK[272]	2	1086.6296	6.189	0.515
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VTGEGCVYLQTSLK[272]	2	916.4805	5.636	0.528
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VYDYYETDEFAIAEYNAPCSK[272]	2	1413.1319	6.211	0.609
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YDVENCLANK[272]	2	751.8647	4.028	0.462
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YILNGGTLLGLK[272]	2	775.4815	4.182	0.514
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YSDASDCHGEDSQAFCEK[272]	2	1186.4625	5.327	0.62
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AVDQSVLLM[147]KPDAELSASSVYNLLPEK	3	1118.6057	4.875	0.571
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DTVIKPLLVEPEGLEK	3	734.1033	2.481	0.146
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FSGQLNSHGCFYQQVK	3	723.6838	5.014	0.526
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GFSSTQDTVVALHALSK	3	681.0408	5.048	0.411
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	HNVYINGITYTPVSSTNEK	3	806.4202	4.455	0.421
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KPQYM[147]VLVPSLLHTETTEK	3	884.1594	4.808	0.5
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KYSDASDCHGEDSQAFCEK	3	878.036	6.714	0.65

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLLQQVSLPELPGEYSM[147]K	3	781.0994	3.269	0.221
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LVHVEEPHTETVR	3	562.6372	2.792	0.097
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	MVLVPSLLHTETTEK	3	660.0396	4.313	0.432
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	MVSGFIPLKPTVK	3	613.0426	2.28	0.11
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	PQYMVLVPSLLHTETTEK	3	789.4311	4.681	0.366
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	PQYM[147]VLVPSLLHTETTEK	3	794.7628	4.994	0.483
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QFSFPLSSEPFQGSYK	3	710.3616	3.137	0.199
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QLNYKHYDGSYSTFGER	3	782.3818	4.28	0.448
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QQNAQGGFSSTQDTVVALHALSK	3	889.7998	7.236	0.599
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLFTDLEAENDVLHCVAFAVPK	3	915.4644	5.125	0.541
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SIYKPGQTVK	3	514.3119	1.57	0.046
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSTQDTVVALHALSK	3	613.0108	4.485	0.48
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SVSGKPQYM[147]VLVPSLLHTETTEK	3	994.2107	4.756	0.502
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TAQEGDHGSHVYTK	3	603.9675	1.809	0.266
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TEHPFTVEEFVLPK	3	651.6875	4.005	0.341
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TEVSSNHVLIYLDK	3	633.3509	1.817	0.002
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VVSMDENFHPLNELIPLVYIQDPK	3	1030.8826	3.767	0.446
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VVSM[147]DENFHPLNELIPLVYIQDPK	3	1036.2143	5.497	0.465
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YNILPEKEEFPFALGVQTLPQTCDEPK	3	1191.6105	5.641	0.431
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YSDASDCHGEDSQAFCEK	3	788.6394	5.059	0.711
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AGAFCLSEDAGLGISSTASLR	3	739.3651	2.89	0.13
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AQGGFSSTQDTVVALHALSK[272]	3	769.0845	6.566	0.49
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AVDQSVLLM[147]K[272]PDAELSASSVYNLLPEK	3	1122.6128	5.314	0.468
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DTVIK[272]PLLVEPEGLEK[272]	3	738.1104	3.647	0.285
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EQAPHCICANGR	3	512.2216	3.75	0.436
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSGQLNSHGCFYQQVK[272]	3	726.3552	5.518	0.586
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GFSSTQDTVVALHALSK[272]	3	683.7122	5.542	0.468
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LHTEAQIQEEGTVVELTGR	3	752.0647	3.925	0.442
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLEIPLTVTHPVVR	3	577.6932	3.348	0.495
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LVHVEEPHTETVR	3	563.9729	2.842	0.313
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MCPQLQQYEMHGPEGLR	3	736.3377	5.677	0.585
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MCPQLQQYEM[147]HGPEGLR	3	741.6694	4.264	0.493
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MVSGFIPLK[272]PTVK[272]	3	617.0497	2.357	0.273
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]PQYMVLVPSLLHTETTEK[272]	3	792.1025	3.953	0.225
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]PQYM[147]VLVPSLLHTETTEK[272]	3	797.4342	4.834	0.431
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QFSFPLSSEPFQGSYK[272]	3	713.033	3.281	0.132
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QLNYK[272]HYDGSYSTFGER	3	785.0532	4.578	0.48
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QQNAQGGFSSTQDTVVALHALSK[272]	3	892.4712	7.587	0.565
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCGH	3	670.6433	5.013	0.476
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCGHTQTVQAH	3	925.7689	5.528	0.487
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLFTDLEAENDVLHCVAFAVPK[272]	3	918.1358	5.827	0.504
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLNEEAVK[272]K[272]	3	483.9594	3.202	0.377
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SIYK[272]PGQTVK[272]	3	518.319	3.034	0.335
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSTQDTVVALHALSK[272]	3	615.6822	4.539	0.418
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SVSGK[272]PQYMVLVPSLLHTETTEK[272]	3	992.8862	4.696	0.513
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SVSGK[272]PQYM[147]VLVPSLLHTETTEK[272]	3	998.2178	5.435	0.473
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TAQEGDHGSHVYTK[272]	3	606.6389	2.208	0.297
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAHLR	3	731.7226	3.918	0.442

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VSVQLEASPAFLAVPVEK[272]	3	724.7555	3.957	0.249
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVSMDENFHPLNELIPLVYIQDPK[272]	3	1033.554	4.828	0.473
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VYDYYETDEFAIAEYNAPCSK[272]	3	942.4237	2.343	0.043
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YMVLVPSLLHTETTEK[272]	3	717.0654	4.157	0.316
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YNILPEK[272]EEFPFALGVQTLPQTCDEPK[272]	3	1195.6176	6.275	0.546
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YSDASDCHGEDSQAFCEK[272]	3	791.3108	3.857	0.499
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KPQYMVLVPSLLHTETTEK	4	659.3727	3.972	0.435
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KPQYM[147]VLVPSLLHTETTEK	4	663.3714	4.214	0.468
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SVSGKPQYM[147]VLVPSLLHTETTEK	4	745.9099	3.157	0.315
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]SHELPCGH	4	507.233	4.929	0.474
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SGK[272]PQYM[147]VLVPSLLHTETTEK[272]	4	702.3901	4.027	0.459
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SVSGK[272]PQYMVLVPSLLHTETTEK[272]	4	744.9165	4.5	0.534
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SVSGK[272]PQYM[147]VLVPSLLHTETTEK[272]	4	748.9152	3.005	0.176
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TAQEGDHGSHVYTK[272]	4	455.231	2.919	0.507
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCGHTQTVQAH	5	555.8643	3.573	0.334
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ATVLNYLPK	2	649.8952	2.663	0.344
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AAQVTIQSSGTFSSK	3	597.9915	3.143	0.276
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AVDQSVLLMKPDAELSASSVYNLLPEK	3	1113.2741	4.221	0.325
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	M[147]VSGFIPLKPTVK	3	618.3743	1.861	0.035
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SKAIGYLNTGYQR	3	584.3248	3.689	0.339
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLNEEAVKK	3	479.9523	3.168	0.253
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TGTHGLLVKQEDM[147]K	3	665.0348	3.269	0.361
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VSVQLEASPAFLAVPVEK	3	722.0841	3.643	0.31
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AAQVTIQSSGTFSSK[272]	3	600.6629	2.692	0.173
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AVDQSVLLMK[272]PDAELSASSVYNLLPEK[272]	3	1117.2812	4.06	0.327
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]HNVYINGITYTPVSSTNEK[272]	3	809.0916	3.837	0.196
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]HYDGSYSTFGER	3	521.5713	2.617	0.318
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLLQQVSLPELPGEYSMK[272]	3	778.4392	3.875	0.302
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLLQQVSLPELPGEYSM[147]K[272]	3	783.7708	3.395	0.162
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SK[272]AIGYLNTGYQR	3	586.9962	4.013	0.309
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TEHPFTVEEFVLPK[272]	3	654.3589	3.895	0.325
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TEVSSNHVLIYLDK[272]	3	636.0223	2.978	0.199
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVSM[147]DENFHPLNELIPLVYIQDPK[272]	3	1038.8857	3.52	0.306
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QQNAQGGFSSTQDTVVALHALSK	4	667.6017	3.997	0.327
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	NEDSLVFVQTDK	2	837.9385	3.772	0.418
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLNEEAVKK	2	719.4249	3.757	0.317
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SIYKPGQTVK	2	770.9642	1.25	0.016
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSGSLLNNAIK	2	692.4014	3.62	0.366
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EQAPHCICANGR	2	767.8288	3.292	0.47
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLNEEAVK[272]K[272]	2	725.4355	3.846	0.351
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSGSLLNNAIK[272]	2	696.4085	3.957	0.357
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VTAAPQSVCALR	2	703.3742	3.375	0.302
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YGAATFTR	2	515.7756	2.604	0.238
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DMYSFLEDM[147]GLK[272]	3	584.9524	3.078	0.286
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GPTQEFK[272]K[272]	3	456.2734	2.776	0.22
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TGTHGLLVK[272]QEDM[147]K[272]	4	502.0333	2.427	0.288
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAHLR	4	549.0438	2.683	0.231
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LPPNVVEESAR	2	675.8725	2.157	0

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DNGCFR	2	451.1925	2.125	0.346
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SASNMAIVDVK[272]	2	711.8969	2.985	0.155
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SIYK[272]PGQTVK[272]	2	776.9748	1.231	0.01
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KPQYMVLVPSLLHTETTEK	3	878.8278	4.987	0.359
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SVSGKPQYMVLVPSLLHTETTEK	3	988.8791	4.885	0.43
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GHFSISIPVK[272]	3	458.2779	2.568	0.254
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]MVLVPSLLHTETTEK[272]	3	662.711	4.446	0.372
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]STQDTVVALHALSK[272]	3	586.6715	3.75	0.424
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVVQK[272]K[272]	3	378.2641	1.898	0.301
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLNEEAVK	1	1169.6526	2.836	0.185
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FEVQVTVPK	2	663.8927	2.84	0.327
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SASNMAIVDVK	2	707.8898	3.457	0.181
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]HYDGSYSTFGER	2	781.8533	2.146	0.167
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LVHVEEPHTETVR	2	845.4557	2.984	0.34
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VGFYESDVM[147]GR	2	710.3384	2.156	0.263
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DLKPAIVK	3	435.2869	2.515	0.232
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GPTQEFKK	3	452.2663	3.009	0.164
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLNEEAVK[272]	1	1177.6668	2.898	0.199
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LPPNVVEESAR	2	677.876	2.57	0.423
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AQGGFSSTQDTVVALHALSK	3	766.4131	4.827	0.35
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFQPFFVELTM[147]PYSVIR	3	735.7227	3.847	0.203
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YGAATFTR	1	1026.5368	2.424	0.24
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AIGYLNTGYQR	1	1399.7451	2.863	0.318
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LSFYYLIM[147]AK[272]	2	776.938	3.816	0.335
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YNILPEKEEFPFALGVQTLPQTCDEPK	4	893.9597	2.722	0.117
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVVQK[272]	1	860.5808	1.601	0.126
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DNGCFR	2	449.1889	2.036	0.331
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FEVQVTVPK[272]	2	667.8998	2.066	0.151
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FQVDNNNR	2	575.7897	2.631	0.171
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GPTQEFK[272]	2	547.8079	2.84	0.191
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLNEEAVK[272]	2	589.337	2.688	0.141
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KYSDASDCHGEDSQAFCEK	4	658.7789	2.494	0.159
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCGH	4	503.2343	4.251	0.344
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SDIAPVAR	1	972.5595	2.406	0.057
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SASNM[147]AIVDVK	2	715.8873	3.619	0.171
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DNSVHWER	3	394.8599	2.967	0.358
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TITKLSFVK	3	486.3132	2.24	0.314
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YGAATFTR	1	1030.5439	1.848	0.355
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SASNM[147]AIVDVK[272]	2	719.8944	2.986	0.127
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DLKPAIVK	2	652.4267	2.425	0.2
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GEAFTLK	2	523.3057	2.332	0.171
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SLNEEAVK	2	585.3299	2.984	0.159
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GEAFTLK[272]	2	527.3128	2.404	0.191
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SVSGKPQYMVLVPSLLHTETTEK	4	741.9112	3.785	0.221
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GHFSISIPVK	2	682.9061	2.609	0.292
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GPTQEFK	2	543.8008	2.143	0.148
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GEAFTLK	1	1045.6042	2.389	0.044
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GHFSISIPVK[272]	2	686.9132	2.331	0.326

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SDIAPVAR	2	486.7834	2.418	0.026
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GVPIPNK	2	502.8162	2.324	0.225
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DMYSFLEDM[147]GLK	3	582.281	2.511	0.244
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]IAQWQSFQLEGGLK	3	630.3516	3.164	0.265
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QGIPFFGQVR	2	646.8652	2.295	0.022
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VVVQK	1	852.5666	1.813	0.15
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GEAFTLK[272]	1	1053.6184	2.164	0.061
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DMYSFLEDMGLK	3	576.9494	2.477	0.216
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KEYEMK	3	416.5654	2.443	0.128
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LSFVK[272]VDSHFR	3	541.6468	2.831	0.223
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FQVDNNNR	1	1150.5722	2.413	0.164
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFQPFFVELTM[147]PYSVIR	2	1103.0804	4.222	0.318
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DLK[272]PAIVK[272]	2	658.4374	2.086	0.106
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DMYSFLEDM[147]GLK	2	872.9179	2.128	0.298
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TTVM[147]VK[272]	2	491.7959	2.028	0.206
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VDLSFSPSQSLPASHAHL	3	678.3532	4.549	0.255
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YK[272]PGQTVK[272]	3	451.6136	2.483	0.337
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AVDQSVLLMK[272]PDAELSASSVYNLLPEK[272]	4	838.2127	3.477	0.097
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GVPIPNK[272]	2	506.8233	2.066	0.197
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TTVMVK[272]	2	483.7985	2.196	0.162
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VSGFIPLKPTVK	3	569.3625	2.601	0.35
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TK[272]VFQLK[272]	3	432.6186	2.888	0.145
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAHL	3	679.6889	3.5	0.256
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SDIAPVAR	1	968.5524	2.394	0.035
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FQVDNNNR	2	573.7862	2.269	0.042
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AFTNSK	2	474.2691	2.184	0.117
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ALLAYAFALAGNQDK[272]	3	618.6836	1.971	0.133
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VVVQKK	2	560.8819	2.011	0.16
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VDSHFR	1	900.4687	1.824	0.075
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDSHFR	2	452.7415	2.06	0.125
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YM[147]VLVPSLLHTETTEK[272]	3	722.3971	4.156	0.28
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VVVQK	2	426.7869	2.028	0.097
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ITIALLEIPLTVTHPVVR	3	710.4441	3.064	0.339
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YM[147]VLVPSLLHTETTEK	3	719.7257	4.026	0.285
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVVQK[272]	2	430.794	2.001	0.115
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LSFVK[272]VDSHFR	4	406.4869	1.915	0.147
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TTVMVK	2	479.7914	2.049	0.121
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QQNAQGGFSSTQDTVVALHALSK[272]	4	669.6053	3.55	0.153
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EYEMK[272]	2	494.2566	1.826	0.175
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLLQQVSLPELPGEYSMK	3	775.7678	2.814	0.033
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TGTHGLLVK	2	603.3719	2.011	0.149
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KEYEMK	2	624.3445	2.14	0.063
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DMYSFLEDMGLK[272]	3	579.6208	2.252	0.217
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]SHELPCG	3	630.2886	2.983	0.292
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TGTHGLLVK[272]QEDM[147]K[272]	3	669.0419	2.218	0.251
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFTNSK[272]	2	478.2762	1.986	0.087
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TTVM[147]VK	2	487.7888	1.992	0.143
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VDSHFR	2	450.738	1.968	0.107

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLEIPLTVTHPVVR	3	576.3575	3.201	0.273
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VGFYESDVM[147]GR	1	1419.6695	1.382	0.273
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TKVFQLK	3	428.6115	2.301	0.168
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ALLAYAFALAGNQDK[272]R	3	670.7174	2.171	0.196
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDSHFR	1	904.4758	1.669	0.064
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SSEPFQGSYK	2	705.3567	3.74	0.402
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DNSVHWERPQK[272]PK[272]	4	514.0381	2.587	0.076
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]PVPGHVTVSICR	3	485.5958	2.96	0.269
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	M[147]VLVPSLLHTETTEK	3	665.3712	3.404	0.246
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FVELTM[147]PYSVIR	2	807.9377	3.594	0.266
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AVDQSVLLM[147]K[272]PDAELSASSVYNLLPEK	4	842.2114	2.646	0.024
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DNSVHWERPQK[272]PK	2	1023.0617	2.511	0.11
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GPTQEFKK	2	677.8958	2.144	0.088
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VLVPSLLHTETTEK	2	924.0355	4.114	0.397
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SSEPFQGSYK[272]	2	709.3638	3.736	0.372
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VLVPSLLHTETTEK	3	616.3594	2.973	0.257
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LSFYYLIM[147]AK	2	772.9309	3.777	0.323
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KDNSVHWER	3	484.2565	3.05	0.224
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFALAGNQDK[272]	2	661.869	3.826	0.366
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]IAQWQSFQLEGGLK	2	945.0237	3.873	0.152
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LTAQPAPTSEDLTSATNIVK[272]	3	782.4316	2.998	0.324
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSGQLNSHG	2	545.7735	2.749	0.449
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]NQGNTWLTAFVLK	2	888.4999	3.914	0.225
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TK[272]VFQLK[272]	2	648.4242	2.788	0.085
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TGTHGLLVK[272]	3	405.2551	2.524	0.066
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	EYEMK	2	490.2495	1.712	0.086
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSFPLSSEPFQGSYK[272]	2	1005.0166	4.62	0.364
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TKVFQLK	2	642.4136	2.721	0.088
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VTGEGCVYLQTSLK	3	608.6513	2.228	0.122
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TITK[272]LSFVK[272]	3	490.3203	2.559	0.11
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GYLNTGYQR	2	608.3156	3.515	0.388
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ALLAYAFALAGNQDK	3	616.0122	2.028	0.087
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]IGYLNTGYQR	2	664.8576	2.38	0.226
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VDLSFSPSQSLPASHAH	3	641.9942	3.215	0.225
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	QLNYK	2	473.2795	1.725	0.097
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TVSAEALESQELCGTEVPSVPEHGR	3	939.1168	3.305	0.245
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVSMDENFHPLNELIPLVYIQD	2	1365.2025	3.398	0.356
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FVELTMPYSVIR	2	799.9403	3.516	0.329
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VSGFIPLK[272]PTVK[272]	2	859.5507	2.916	0.469
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LLIYAVLPTGDVIGDSAK[272]	3	711.7518	2.556	0.153
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	M[147]VSGFIPLKPTVK	4	464.0325	1.742	0.022
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GPTQEFK[272]K[272]R	2	761.957	2.215	0.227
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FVELTM[147]PYSVIR	2	805.9342	2.746	0.284
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FTVLQDVPVR	2	659.386	2.797	0.307
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLHTETTEK	3	451.2575	2.397	0.261
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VDLSFSPSQSLPASH	2	856.4361	2.9	0.335
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YESDVM[147]GR	2	558.7592	2.22	0.384
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SIYKPGQ	2	536.8112	2.002	0.384

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]RK[272]EYEMK[272]	2	708.4057	2.021	0.116
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]SHELPCGHTQTVQAH	4	698.5772	3.5	0.173
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DLK[272]PAIVK[272]	3	439.294	2.155	0.009
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FVELTMPYSVIR	2	797.9367	3.457	0.208
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VVVQK[272]K[272]	2	566.8926	1.356	0.051
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QLNYK[272]	2	477.2866	1.602	0.005
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SASNMAIVDVK[272]	3	474.9337	2.316	0.017
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TGTHGLLVK	3	402.5837	1.942	0.082
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AGLGISSTASLR	2	638.8707	2.778	0.268
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QLNYK[272]HYDGSYSTFGER	4	589.0418	2.525	0.188
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SASNM[147]AIVDVK	3	477.5939	1.837	0.087
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SNM[147]AIVDVK[272]	2	640.8598	2.796	0.215
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SLFTDLEAENDVLHCVAF	2	1107.0224	3.947	0.387
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LLHTETTEK	2	676.3827	2.554	0.278
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FIPLKPTVK	2	731.9791	1.951	0.3
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ETTFNSLLCPSGGEVSEELSLK[272]	3	892.1113	2.118	0.118
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LK[272]PAIVK[272]	2	600.9239	2.228	0.263
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSGQLNSH	2	517.2628	2.497	0.217
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SIYK[272]PGQ	2	540.8183	1.877	0.309
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EVSEELSLK[272]	2	661.3763	2.602	0.269
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	GPTQEFKKR	2	755.9463	2.267	0.201
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TEVSSNHVLIY	2	703.3758	2.49	0.301
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	HVLIYLDK	2	640.8899	3.014	0.126
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSGQLNSHGC	2	620.272	2.339	0.297
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]APVGHFYEPQAPSAEVEMTSY	2	1227.5738	3.414	0.264
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	RKEYEMK	2	702.3951	2.298	0.116
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FEVQVTVPK	2	665.8962	1.822	0.021
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]SHELPC	3	611.2814	2.286	0.234
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ALLAYAF	2	454.7657	1.983	0.287
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SASNM[147]AIVDVK[272]	3	480.2653	1.961	0.032
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VSGFIPLK[272]PTVK[272]	3	573.3696	2.821	0.167
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DNSVHWERPQK[272]PK	4	512.0345	2.051	0.03
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]NQGNTWLTAFVLK	3	592.669	2.126	0.115
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TITKLSFVK	2	728.9662	2.056	0.277
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LSVSYTGSR	2	557.3047	2.431	0.266
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TVLQDVPVR	2	585.8518	2.304	0.235
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EQAPHCICAN	2	661.2675	2.337	0.238
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	RKEYEMK	4	351.7012	1.948	0.116
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFYYLIM[147]AK[272]	2	720.396	2.494	0.237
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VYDYYETDEFAIAEYNAPCSK	3	939.7523	1.693	0.053
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LSFVKVDSHFR	3	538.9754	2.395	0.044
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GTEVPSVPEHGR	3	470.2482	2.699	0.147
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FSGQLNSH	2	515.2593	2.537	0.147
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	M[147]VSGFIPLK	2	644.3783	2.288	0.258
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GIPFFGQVR	2	582.836	2.391	0.204
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TM[147]PYSVIR	2	563.806	2.066	0.244
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]PVPGHVTVSICR	2	727.8901	2.019	0.177
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GFIPLK[272]PTVK[272]	3	511.3361	1.821	0.239

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SNM[147]AIVDVK	2	636.8527	2.447	0.183
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TITK[272]LSFVK[272]	2	734.9768	1.699	0.132
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TVLQDVPVR	2	583.8482	2.152	0.252
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VNYKDR	2	537.8064	1.831	0.028
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ALAGNQDK	2	548.8091	2.576	0.154
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]EYEM[147]K[272]	2	502.2541	1.265	0.003
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FSGQLNSHGCF	2	693.8062	2.418	0.193
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LNTGYQR	2	498.2732	2.432	0.107
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	IPLKPTVK	2	658.4449	1.879	0.24
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]CPQLQQYEMHGPEGLR	3	692.6576	2.507	0.162
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SVSYTGSR	2	498.7591	1.852	0.243
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FIPLKPTVK	3	488.3218	1.843	0.206
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YYLIM[147]AK	2	599.3387	1.837	0.233
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YLNTGYQR	2	579.8048	2.262	0.156
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LKPAIVK	3	396.9446	2.98	0.033
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YMVLVPSLLHTETTEK	3	714.394	3.231	0.156
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TMPYSVIR	2	555.8085	2.107	0.14
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]PNVVEESAR	2	572.8076	2.455	0.097
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SGGRTEHPFTVEEFVLPK	3	770.7462	3.342	0.143
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FIPLK[272]PTVK[272]	2	737.9898	2.539	0.209
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AVDQSVLLM[147]KPDAELSASSVYNLLPEK	4	839.2061	2.534	0.033
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]IPLK[272]PTVK[272]	2	664.4555	1.94	0.175
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KEYEM[147]K	3	421.8971	1.385	0.042
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]DNSVHWERPQKPK	4	512.0345	2.327	0.08
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	RKEYEM[147]K	2	710.3925	1.696	0.055
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VLVPSLLHTETTEK[272]	3	619.0308	2.716	0.099
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]LVHVEEPHTETVR	4	423.2315	1.959	0.058
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SVSYTGSR	2	500.7626	2.147	0.099
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TQEFK[272]K[272]	2	606.8693	2.049	0.084
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QLEGGLK[272]	2	516.8183	2.208	0.109
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LKPAIVK	2	594.9132	1.839	0.153
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]SHELPCGH	3	675.9749	2.337	0.154
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]APVGHFYEPQAPSAEVEMTSY	3	818.7183	2.558	0.13
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	KLSFYYLIM[147]	2	737.4124	1.736	0.266
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	NALFCLESAWK[272]	3	537.943	2.384	0.006
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]YNILPEK[272]	2	582.847	2.486	0.138
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YNLLPEK	2	578.8399	2.385	0.082
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]HYDGSYSTFGE	2	703.8027	1.999	0.172
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCGHTQTVQAH	4	694.5785	2.873	0.057
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]VTVSICR	2	484.2629	1.981	0.073
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TQEFK[272]K[272]	3	404.9153	2.084	0.079
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	SFVHLEPM[147]	2	558.2814	1.689	0.17
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	HYDGSYSTFGE	2	701.7991	1.601	0.229
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	VLLAYLTAQPAPTSEDLTSATNIVK	3	966.2059	3.037	0.13
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TQEFK[272]	2	470.7708	1.775	0.07
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	RKEYEMK	3	468.5991	1.99	0.052
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]CICANGR	2	486.7038	1.616	0.161
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFQPFFVEL	2	621.338	2.101	0.048

137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]IALLEIPLTVTHPVVR	3	639.0669	3.252	0.245
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]NQGNTWLTAF	2	648.3287	2.02	0.284
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	FVELTMPYSVIR	3	532.2936	2.128	0.093
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ALLAYAF	2	456.7692	1.824	0.061
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	LSVSYTGSR	2	555.3011	2.07	0.071
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SIYK[272]PG	2	476.789	1.742	0.081
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]QFSFPLSSEPFQG	2	807.8997	2.255	0.031
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPC	3	605.9498	1.801	0.157
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFQLEGGLK[272]	2	633.8685	1.98	0.034
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPMSHELPCG	3	624.9569	2.505	0.051
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	ALLAYAF	1	908.5241	1.559	0.282
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]SFVHLEPM[147]	2	560.2849	1.622	0.177
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]GK[272]PQYM[147]VLVPSLLHTETTEK[272]	4	680.6321	2.633	0.17
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	TQEFKK	2	600.8587	1.911	0.008
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]AFQPF	1	753.4052	1.648	0.137
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	HYDGSYSTFG	2	637.2779	1.93	0.158
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	YILNGGTLLGLK	3	514.652	2.005	0.096
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]TPVSSTNEK[272]	2	625.8452	1.763	0.051
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	DTVIKPL	2	533.337	1.613	0.059
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	AFQPF	1	749.3981	1.518	0.155
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]ALLAYAF	1	912.5312	1.571	0.199
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	n[145]FIPLK[272]PTVK[272]	3	492.3289	2.151	0.005
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	EM[147]K	1	703.3808	1.121	0.072
137	IPI00478003	1	74.1	0.77	0.13	851	##	A2M Alpha-2-macroglobulin precursor	RKDTVIK	2	640.4141	1.645	0.035
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]LSNNALSGLPQGVFGK[272]	2	945.538	4.973	0.521
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]LSNNALSGLPQGVFGK[272]	3	630.6944	3.956	0.282
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	GQVVPALNEK	2	667.8932	3.233	0.382
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]GQVVPALNEK[272]	2	671.9003	3.265	0.334
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	LSNNALSGLPQGVFGK	2	941.5309	3.603	0.461
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]LEILSISK[272]	2	595.8836	2.723	0.111
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	LSNNALSGLPQGVFGK	3	628.023	2.955	0.071
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	LEILSISK	2	591.8765	2.187	0.107
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	DLEELVK	2	563.3294	2.153	0.05
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]QLVCPVTR	2	553.3026	1.944	0.025
138	IPI00479116	1	9.2	0.77	0.03	15	11	CPN2 Carboxypeptidase N subunit 2 precursor	n[145]VLPAGLFAH	2	534.8198	2.532	0.351
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	ADLSGITGAR	1	1100.6059	3.215	0.435
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]ADLSGITGAR	1	1104.613	2.977	0.311
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	AVLDVFEEGTEASAATAVK	2	1094.5785	6.621	0.586
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	DEELSCTVVELK	2	845.9232	4.841	0.422
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LYGSEAFATDFQDSAAAK	2	1086.5341	5.596	0.634
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	NLAVSQVVHK	2	687.9145	4.141	0.378
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	VLVNYIFFK	2	711.9291	3.836	0.382
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]AVLDVFEEGTEASAATAVK[272]	2	1098.5856	7.041	0.605
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FRDEELSCTVVELK[272]	2	1001.515	4.74	0.439
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LYGSEAFATDFQDSAAAK[272]	2	1090.5412	7.292	0.601
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]NSPLDEENLTQENQDR	2	1023.4783	5.158	0.482
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]HPNSPLDEENLTQENQDR	3	760.6918	5.583	0.396
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]WEMPFDPQDTHQSR	3	639.9605	3.65	0.49

139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	KLINDYVK	3	471.6257	3.073	0.094
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LYGSEAFATDFQDSAAAK[272]	3	727.3632	2.43	0.117
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]NLAVSQVVHK[272]	3	461.6168	2.04	0.073
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]ITLLSALVETR	2	680.42	3.479	0.271
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]WEM[147]PFDPQDTHQSR	3	645.2922	2.465	0.374
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]ADLSGITGAR	2	552.8101	2.375	0.149
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]NLAVSQVVHK[272]	2	691.9216	4.007	0.332
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LM[147]IIVPTDTQNIFFMSK	3	765.4144	3.45	0.421
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]EIGELYLPK[272]	2	675.3996	3.076	0.252
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]MSLHHLTIPYFR	3	553.6368	2.386	0.227
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	EIGELYLPK	2	671.3925	2.877	0.21
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]DEELSCTVVELK[272]	2	849.9303	4.95	0.24
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	NLAVSQVVHK	3	458.9454	3.075	0.171
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	ADLSGITGAR	2	550.8066	2.86	0.311
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LINDYVK	2	572.8399	3.117	0.171
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	KLINDYVK	2	706.9349	3.276	0.143
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	EQLSLLDR	1	1113.6263	2.352	0.137
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]EQLSLLDR	1	1117.6334	2.357	0.127
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LM[147]IIVPTDTQNIFFM[147]SK[272]	3	773.4175	3.226	0.386
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FTEDAK[272]	1	998.5398	2.124	0.09
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]EQLSLLDR	2	559.3203	3.127	0.117
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]ITLLSALVETR	3	453.9491	3.43	0.117
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LMIIVPTDTQNIFFMSK[272]	3	762.7542	3.615	0.327
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FTEDAK[272]R	3	385.5518	2.51	0.19
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]MEEVEAM[147]LLPETLK[272]	2	969.0143	2.941	0.214
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FTEDAK[272]R	2	577.8241	1.921	0.169
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	FTEDAK	1	990.5256	2.028	0.021
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FRDEELSCTVVELK[272]	3	668.0124	3.662	0.362
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LINDYVK[272]	2	576.847	2.217	0.127
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	M[147]EEVEAM[147]LLPETLK	2	973.0047	2.235	0.221
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	DLDSQTMM[147]VLVNYIFFK	3	787.4058	4.795	0.325
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FTEDAK[272]	2	499.7735	1.974	0.145
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]ITDIIK[272]	2	495.8255	2.225	1
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]LM[147]IIVPTDTQNIFFMSK[272]	3	768.0858	2.799	0.342
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	FTEDAKR	2	573.817	2.365	0.014
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	EQLSLLDR	2	557.3168	2.252	0.024
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]IIVPTDTQNIFFM[147]SK[272]	3	686.711	3.586	0.194
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]HPNSPLDEENLTQENQDR	2	1140.5341	4.59	0.379
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]IIVPTDTQNIFFMSK[272]	3	681.3793	2.999	0.3
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]DSLEFR	2	455.7412	1.898	0.033
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]SLHHLTIPYFR	3	509.9566	2.511	0.291
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LMIIVPTDTQNIFFMSK	3	760.0828	3.038	0.267
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	DSLEFR	1	906.468	1.428	0.085
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]DLDSQTMM[147]VLVNYIFFK[272]	3	790.0772	3.689	0.294
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]VLVNYIFFK[272]	2	715.9362	3.813	0.333
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]FYLSK[272]K[272]	3	406.5921	1.855	0.022
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]TLLSALVETR	2	623.878	3.177	0.186
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]MEEVEAMLLPETLK[272]	2	961.0169	1.417	0.043

139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	FTEDAK	2	495.7664	1.516	0.011
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	IIVPTDTQNIFFM[147]SK	3	684.0396	2.976	0.193
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]MEEVEAM[147]LLPETLK[272]R	3	698.379	1.926	0.079
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]AVLDVFEEGTEASAATAVK[272]	3	732.7261	2.098	0.061
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LYGSEAFATDFQDSAAAK	3	724.6918	2.096	0.09
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	LM[147]IIVPTDTQNIFFM[147]SK	3	770.7461	2.807	0.14
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]DLSGITGAR	2	517.2916	2.405	0.121
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]HLTIPYFR	2	595.8438	2.075	0.168
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]EQLSLLDRFTEDAK[272]R	4	528.044	1.993	0.004
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	n[145]SALVETR	2	460.2701	2.216	0.041
139	IPI00550991	1	52.7	0.66	0.07	129	71	SERPINA3 Alpha-1-antichymotrypsin precursor	SK[272]K[272]	2	395.7731	0.913	0.136
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]EFSHLGK[272]	1	1105.6245	2.704	0.288
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	CCESASEDCMAK	2	847.7958	4.096	0.606
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	CCESASEDCM[147]AK	2	855.7933	3.887	0.695
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	DVCDPGNTK	2	637.797	3.076	0.521
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	GQELCADYSENTFTEYK	2	1162.5141	5.726	0.589
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	SCESNSPFPVHPGTAECCTK	2	1256.5164	6.319	0.642
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	SLGECCDVEDSTTCFNAK	2	1170.4608	4.342	0.627
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	VCSQYAAYGEK	2	772.8655	3.798	0.348
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	VPTADLEDVLPLAEDITNILSK	2	1323.7337	4.834	0.455
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]CCESASEDCMAK[272]	2	851.8029	4.28	0.646
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]GQELCADYSENTFTEYK[272]	2	1166.5212	6.078	0.616
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	2	1260.5235	5.232	0.653
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SLGECCDVEDSTTCFNAK[272]	2	1174.4679	4.977	0.684
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VCSQYAAYGEK[272]	2	776.8726	4.007	0.521
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	2	1327.7408	4.919	0.487
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	3	705.7369	2.6	0.142
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	SCESNSPFPVHPGTAECCTK	3	838.0134	3.191	0.5
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	VPTADLEDVLPLAEDITNILSK	3	882.8249	3.047	0.191
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]ELSSFIDK[272]GQELCADYSENTFTEYK[272]	3	1132.539	7.664	0.627
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	3	840.6848	3.95	0.566
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SPFPVHPGTAECCTK[272]	3	651.9693	4.226	0.492
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	3	885.4963	2.149	0.323
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	AKLPDATPK	3	454.2819	3.171	0.33
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KLCMAALK	3	448.5957	3.176	0.253
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KLCM[147]AALK	3	453.9273	2.708	0.318
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	RTHLPEVFLSK	3	536.3191	3.15	0.347
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]AK[272]LPDATPK[272]	3	458.289	3.556	0.278
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]THLPEVFLSK[272]	3	486.9568	2.858	0.163
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]HQPQEFPTYVEPTNDEICEAFR	4	710.8249	2.721	0.286
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	2	1058.1017	3.827	0.349
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	RTHLPEVFLSK	2	803.9751	3.523	0.431
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]RTHLPEVFLSK[272]	2	807.9822	3.92	0.36
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	LCDNLSTK	2	610.3123	2.919	0.285
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]EDFTSLSLVLYSR	2	837.447	3.736	0.323
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]EFSHLGK[272]	2	553.3159	2.958	0.263
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]HLSLLTTLSNR	2	699.9129	3.504	0.264
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]LCDNLSTK[272]	2	614.3194	2.937	0.277

140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	FPSGTFEQVSQLVK	3	616.3403	2.872	0.234
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KELSSFIDK	3	496.2925	3.188	0.172
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]ELSSFIDK[272]	1	1226.6872	2.552	0.242
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]DVCDPGNTK[272]	2	641.8041	2.877	0.309
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	THLPEVFLSK	3	484.2854	3.114	0.172
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]LCDNLSTK[272]	1	1227.6316	2.532	0.237
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	ELSSFIDK	2	609.8401	3.559	0.188
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]ELSSFIDK[272]	2	613.8472	2.449	0.126
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]FPSGTFEQVSQLVK[272]	3	619.0117	3.158	0.238
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VM[147]DK[272]YTFELSR	3	564.9672	2.108	0.36
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	ELSSFIDK	1	1218.673	2.311	0.242
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	YTFELSR	1	1055.5521	2.382	0.255
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]DVCDPGNTK[272]	1	1282.601	2.107	0.413
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	EFSHLGK	2	549.3088	2.484	0.267
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	ELPEHTVK	2	616.8535	2.508	0.153
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]THLPEVFLSK[272]	2	729.9316	2.826	0.292
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VLEPTLK[272]	1	1087.6966	2.065	0.184
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]RTHLPEVFLSK[272]	3	538.9905	2.521	0.21
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	4	664.374	3.07	0.015
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]ELPEHTVK[272]	2	620.8606	2.22	0.177
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	NSKFEDCCQEK	3	614.9384	2.917	0.249
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]YTFELSR	1	1059.5592	2.028	0.197
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	THLPEVFLSK	2	725.9245	2.577	0.285
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	VLEPTLK	2	540.3448	2.119	0.244
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VLEPTLK[272]	2	544.3519	2.321	0.122
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]FEDCCQEK[272]	1	1381.5499	2.155	0.345
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KLCM[147]AALK	2	680.3873	2.549	0.253
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]TSALSAK[272]	2	483.2972	2.861	0.135
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]YTFELSR	2	530.2832	2.117	0.101
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	FEDCCQEK	2	687.2715	2.27	0.214
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]HLSLLTTLSNR	3	466.9444	2.137	0.188
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KLCMAALK	2	672.3899	2.503	0.13
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	TSALSAK	2	479.2901	2.554	0.043
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]ELPEHTVK[272]	3	414.2428	1.535	0.027
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]AK[272]LPDATPK[272]	2	686.9299	2.806	0.261
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	AKLPDATPK	2	680.9192	2.71	0.3
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	4	529.5545	2.217	0.113
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	YTFELSR	2	528.2797	2.135	0.055
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VM[147]DK[272]YTFELSR	2	846.9471	2.245	0.276
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	ELPEHTVK	3	411.5714	1.93	0.068
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SCESNSPFPVHPG	2	774.3406	3.567	0.441
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]EFSHLGK[272]	3	369.213	1.646	0.126
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]FEDCCQEK[272]	2	691.2786	1.373	0.05
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SLGECCDVEDSTTCFNAK[272]	3	783.3144	1.966	0.144
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	EFSHLGK	3	366.5416	2.061	0.021
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SLLVSYTK[272]	2	599.8679	2.948	0.182
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]NSK[272]FEDCCQEK[272]	3	618.9455	2.241	0.12
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]CCESASEDCM[147]AK[272]	2	859.8004	2.133	0.363

140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]VMDK[272]YTFELSR	3	559.6355	2.342	0.066
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	VMDKYTFELSR	3	556.9641	2.082	0.036
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTA	2	860.383	2.361	0.182
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]GPLLK[272]K[272]	3	363.257	1.257	0.134
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	4	630.7654	2.044	0.025
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	n[145]HQPQEFPTYVEPTNDEICEAFR	3	947.4307	2.939	0.094
140	IPI00555812	1	62.9	0.92	0.02	8	89	GC Vitamin D-binding protein precursor	FM[147]PAAQLPELPDVELPTNK	3	802.7628	2.851	0.009
141	IPI00654755	1	36.2	3.01	0.4	5	4	HBB Hemoglobin subunit beta	VLGAFSDGLAHLDNLK	3	650.6984	4.465	0.518
141	IPI00654755	1	36.2	3.01	0.4	5	4	HBB Hemoglobin subunit beta	GTFATLSELHCDK	3	583.2884	3.451	0.295
141	IPI00654755	1	36.2	3.01	0.4	5	4	HBB Hemoglobin subunit beta	VHLTPEEK	2	616.8535	2.92	0.297
141	IPI00654755	1	36.2	3.01	0.4	5	4	HBB Hemoglobin subunit beta	KVLGAFSDGLAHLDNLK	3	740.0951	2.401	0.091
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	CSVFYGAPSK	2	692.8413	4.156	0.508
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EMSGSPASGIPVK	2	770.4136	4.147	0.436
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GLEEELQFSLGSK	2	858.962	5.125	0.492
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ITQVLHFTK	2	683.9139	3.962	0.391
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LTVAAPPSGGPGFLSIER	2	955.0307	3.152	0.543
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VDVQAGACEGK	2	701.8445	3.485	0.518
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ASAGLLGAHAAAITAY	2	801.442	4.133	0.557
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHY	2	620.8076	3.784	0.508
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHYY	2	702.3393	3.831	0.484
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHYYYM[147]	2	857.3886	3.96	0.462
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]CSVFYGAPSK[272]	2	696.8484	3.605	0.505
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DFALLSLQVPLK[272]	2	816.5024	3.455	0.45
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EELVYELNPLDHR	2	885.9608	4.22	0.472
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EMSGSPASGIPVK[272]	2	774.4207	4.123	0.379
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EM[147]SGSPASGIPVK[272]	2	782.4182	3.423	0.423
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EPFLSCCQFAESLR	2	933.4124	3.714	0.583
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GHLFLQTDQPIYNPGQR	2	1064.5565	5.825	0.54
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GLEEELQFSLGSK[272]	2	862.9691	4.809	0.498
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GSFEFPVGDAVSK[272]	2	814.4322	4.699	0.476
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LSCCQFAESLR	2	746.8305	3.205	0.529
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LTVAAPPSGGPGFLSIER	2	957.0343	4.08	0.218
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]TTNIQGINLLFSSR	2	854.4792	5.063	0.513
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AEFQDALEKLNMGITDLQGLR	3	881.4705	6.043	0.517
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GPEVQLVAHSPWLK	3	614.3526	4.21	0.408
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHYYYM[147]ILSR	3	728.3619	4.226	0.445
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DFALLSLQVPLK[272]DAK[272]	3	697.4244	4.647	0.48
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GHLFLQTDQPIYNPGQR	3	710.0401	5.031	0.376
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GPEVQLVAHSPWLK[272]	3	617.024	3.685	0.406
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLATLCSAEVCQCAEGK[272]	3	722.333	4.18	0.507
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITPGK[272]PYILTVPGHLDEM[147]QLDIQAR	3	1037.2368	5.251	0.365
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]RGHLFLQTDQPIYNPGQR	3	762.0738	5.052	0.404
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VGLSGM[147]AIADVTLLSGFHALR	3	763.4233	4.694	0.422
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]YVSHFETEGPHVLLYFDSVPTSR	3	942.1403	6.174	0.531
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ITPGKPYILTVPGHLDEMQLDIQAR	4	772.1772	3.963	0.414
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ITPGKPYILTVPGHLDEM[147]QLDIQAR	4	776.1759	5.288	0.551
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITPGK[272]PYILTVPGHLDEMQLDIQAR	4	774.1807	6.085	0.461
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITPGK[272]PYILTVPGHLDEM[147]QLDIQAR	4	778.1794	2.699	0.069

142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AEFQDALEK	2	665.8537	3.858	0.355
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FACYYPR	2	555.2551	2.206	0.17
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LGQYASPTAK[272]	2	662.3792	3.844	0.386
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AEFQDALEKLNM[147]GITDLQGLR	3	886.8022	4.25	0.428
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DFALLSLQVPLKDAK	3	693.4173	3.988	0.442
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GPEVQLVAHSPWLK[272]	3	615.6883	3.613	0.322
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	KYVLPNFEVK	3	552.9994	2.879	0.147
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ITPGKPYILTVPGHLDEM[147]QLDIQAR	3	1034.5654	4.917	0.418
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ITQVLHFTK	3	456.2784	2.988	0.284
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DK[272]GQAGLQR	3	420.9136	3.407	0.36
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EPFLSCCQFAESLR	3	622.6107	2.858	0.354
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLLQALVR	3	592.3653	3.268	0.352
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITQVLHFTK[272]	3	458.9498	3.398	0.281
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]TTNIQGINLLFSSR	3	569.9885	2.805	0.291
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VLQIEK[272]EGAIHR	3	561.0048	3.426	0.317
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]YIYGK[272]PVQGVAYVR	3	634.3678	2.501	0.405
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SHKPLNM[147]GK	4	362.7106	2.598	0.371
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]MRPSTDTITVMVENSHGLR	4	572.7957	5.056	0.307
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]STQDTVIALDALSAYWIASHTTEER	4	731.3705	3.342	0.297
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EGAIHREELVYELNPLDHR	5	487.6577	2.881	0.221
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DFALLSLQVPLK	2	812.4953	3.688	0.363
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DHAVDLIQK	2	659.8776	3.61	0.293
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EM[147]SGSPASGIPVK	2	778.4111	3.563	0.457
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LNM[147]GITDLQGLR	2	743.906	3.41	0.381
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VEYGFQVK	2	625.3506	2.739	0.251
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DHAVDLIQK[272]	2	663.8847	3.194	0.253
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GLQDEDGYR	2	598.7869	2.88	0.329
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITQVLHFTK[272]	2	687.921	3.949	0.349
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VEYGFQVK[272]	2	629.3577	2.729	0.346
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VLSLAQEQVGGSPEK[272]	2	915.5142	3.649	0.451
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DKGQAGLQR	3	418.2422	3.421	0.265
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AIADVTLLSGFHALR	3	576.669	4.098	0.39
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ALLHLLLHEGK[272]	3	511.3236	3.916	0.369
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FGLLDEDGK[272]K[272]TFFR	3	702.3967	2.659	0.366
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FGLLDEDGK[272]K[272]TFFR	4	527.0494	2.623	0.329
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNM[147]GK[272]	4	365.7159	1.622	0.034
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LGQYASPTAK	2	658.3721	3.374	0.31
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ADGSYAAWLSR	2	670.8394	3.681	0.33
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ANSFLGEK[272]	2	577.3264	3.063	0.223
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FGLLDEDGK[272]K[272]	2	777.4486	3.327	0.363
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LNM[147]GITDLQGLR	2	745.9095	3.401	0.326
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GLCVATPVQLR	2	673.8739	2.58	0.407
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LNMGITDLQGLR	2	737.9121	3.365	0.22
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SCGLHQLLR	2	608.8242	2.867	0.302
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VGDTLNLNLR	2	629.8654	2.679	0.19
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	FGLLDEDGKKTFFR	3	698.3896	3.087	0.297
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DHAVDLIQK[272]	3	442.9255	3.57	0.152
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GLQDEDGYR	1	1196.5665	2.57	0.3

142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LNMGITDLQGLR	2	735.9085	3.131	0.288
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]TYNVLDM[147]K[272]	2	644.3465	1.719	0.281
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VLQIEKEGAIHR	3	558.3334	3.607	0.24
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ANSFLGEK	2	573.3193	3.156	0.216
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GSFEFPVGDAVSK	2	810.4251	3.529	0.313
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	KYVLPNFEVK	2	828.9955	3.814	0.224
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]TYNVLDMK[272]	2	636.3491	2.97	0.146
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EVQLVAHSPWLK	3	562.9945	3.755	0.333
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SFQDLSPVIHR	3	481.6006	3.483	0.348
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SHKPLNM[147]GK	2	724.4138	2.044	0.214
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	YVLPNFEVK	2	694.9005	2.675	0.216
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AEFQDALEK[272]	2	669.8608	3.67	0.174
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VDVQAGACEGK[272]	2	705.8516	3.878	0.382
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VVEEQESR	2	560.2918	2.084	0.38
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AIADVTLLSGFHALR	3	575.3333	3.071	0.402
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EAPKVVEEQESR	3	560.9687	2.849	0.27
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	FGLLDEDGKK	3	514.6277	2.275	0.294
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AACAQLNDFLQEYGTQGCQV	3	799.0151	2.752	0.27
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LELSVDGAK[272]	2	610.3605	3.274	0.238
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNM[147]GK[272]	2	730.4245	1.698	0.204
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]YVLPNFEVK[272]	2	698.9076	1.903	0.149
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	TYNVLDM[147]K	2	640.3394	2.201	0.094
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LTSLSDR	2	468.2676	2.423	0.146
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LTVAAPPSGGPGFLSIER	3	638.3586	4.106	0.332
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VTASDPLDTLGSEGALSPGGVASLLR	3	876.4718	3.352	0.189
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VEASISK[272]	2	511.3103	2.247	0.173
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VQQPDCR	2	516.2417	2.111	0.272
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	FGLLDEDGKK	2	771.438	3.21	0.188
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	NNVPCSPK	2	592.7994	2.199	0.257
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	TYNVLDMK	2	632.342	2.078	0.119
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DSSTWLTAFVLK[272]	2	828.466	3.967	0.279
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNMGK[272]	2	722.427	2.703	0.232
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VEASISK	2	507.3032	2.326	0.154
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]NVNFQK[272]	2	519.3028	2.35	0.172
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VDFTLSSER	2	599.3152	2.187	0.218
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNMGK[272]	4	361.7171	2.497	0.132
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SHKPLNMGK	3	477.9466	2.093	0.224
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EVQLVAHSPWLK[272]	3	565.6659	3.689	0.225
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]QGSFQGGFR	2	564.2894	2.462	0.184
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VEASISK[272]	1	1021.6133	2.575	0.098
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNM[147]GK[272]	3	487.2854	1.917	0.235
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	FGLLDEDGK	2	637.343	2.87	0.132
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	TEQWSTLPPETK	2	848.9489	2.692	0.186
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LELSVDGAK	2	606.3534	2.333	0.235
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	NVNFQK	2	515.2957	2.077	0.171
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]NNVPCSPK[272]	2	596.8065	2.336	0.204
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EFHLHLR	2	548.3126	2.549	0.118
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SHKPLNM[147]GK	3	483.2783	1.704	0.242

142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]MRPSTDTITVM[147]VENSHGLR	3	768.7234	3.518	0.11
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SHKPLNMGK	2	716.4163	2.134	0.19
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VFALDQK[272]	2	554.8339	1.325	0.127
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EGAIHREELVYELNPLDHR	4	609.3204	2.196	0.158
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DSSTWLTAFVLK	2	824.4589	2.857	0.154
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]VQQPDCR	2	518.2452	2.174	0.135
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]PVAFSVVPTAAAAVSLK[272]	3	639.3868	2.837	0.351
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GQIVFM[147]NR	2	562.8038	2.855	0.244
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DSSTWLTAFVLK	2	826.4625	2.676	0.091
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SHK[272]PLNMGK[272]	3	481.9537	2.203	0.161
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]RGHLFLQTDQPIYNPGQR	4	571.8072	3.249	0.088
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]TEQWSTLPPETK[272]	2	852.956	1.61	0.276
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]CSVFYGAPSK[272]	3	464.9013	1.687	0.199
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GLESQTK	2	521.7982	1.768	0.194
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ILTVPGHLDEMQLDIQAR	3	731.7321	3.846	0.231
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]MRPSTDTITVM[147]VENSHGLR	4	576.7944	2.715	0.134
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DHAVDLIQK	3	440.2541	2.889	0.044
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AVGSGATFSH	2	537.2724	3.193	0.45
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EFHLHLR	3	365.8775	2.519	0.041
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FGLLDEDGK[272]	2	641.3501	2.202	0.081
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GSFQDLSPVIHR	2	750.408	4.043	0.353
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DK[272]GQAGLQR	2	630.8668	2.384	0.214
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LFETK[272]	2	463.2835	1.994	0.166
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]PVQGVAYVR	2	566.8334	3.358	0.382
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]YIYGK[272]PVQGVAY	2	823.4633	3.385	0.399
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHYYY	2	783.8709	3.188	0.505
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSH	2	539.2759	2.899	0.417
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]CCQDGVTR	2	559.2226	1.446	0.03
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	IFETK	2	459.2764	1.916	0.139
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	VFALDQK	2	550.8268	1.851	0.074
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	YIYGKPVQGVAY	2	819.4562	3.234	0.414
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	YIYGKPVQGVAYVR	3	631.6964	2.829	0.034
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	FGLLDEDGKKTFFR	4	524.0441	2.773	0.093
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLLQALVR	2	888.0443	1.687	0.195
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AVGSGATFSHYYYM[147]IL	2	970.4727	2.976	0.414
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AVGSGATFSHYYY	2	781.8674	2.367	0.416
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ACYYPR	2	481.7209	2.41	0.292
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ADLEK	2	428.2504	1.732	0.017
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	DFALLSLQVPLK	3	541.9993	2.954	0.027
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]AIADVTLLSGFHALR	2	864.4999	3.688	0.454
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPF	2	491.2856	1.528	0.243
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AVGSGATFSHYYYM[147]IL	2	968.4691	2.455	0.488
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLLQAL	2	760.4595	3.069	0.309
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ALLHLLLHEGK	2	762.4747	3.934	0.398
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ALLHLLLHEGK[272]	2	766.4818	3.347	0.41
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ASAGLLGAH	2	470.7703	2.721	0.275
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLLHEGK[272]	3	366.5689	2.178	0.215
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]YYHGDHPVANSLR	3	558.2819	2.468	0.251

142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GSFQDLSPVIHR	3	500.6077	2.748	0.189
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	AVGSGATFSHYY	2	700.3357	2.46	0.338
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]EVQLVAHSPWLK[272]	2	847.9953	4.232	0.215
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	GPEVQLVAHSPWLK[272]	2	923.0288	3.57	0.337
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DGSYAAWLSR	2	635.3209	2.097	0.306
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LLLHEGK	3	363.8975	2.328	0.183
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	HLVPGAPFLLQ	2	666.3954	2.52	0.362
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DFALLSLQVPLK[272]	3	544.6707	2.364	0.024
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]SVGVQLQDVPR	2	671.384	3.329	0.196
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LSCPK	2	437.2379	1.377	0.033
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ASAGLLGAH	2	468.7667	2.087	0.309
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	HLVPGAPFLLQA	2	701.9139	2.071	0.363
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]IYLAPTLAASR	2	660.3938	2.653	0.274
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EVQLVAHSPWLK[272]	3	564.3302	3.088	0.226
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EVQLVAHSPWLK	2	843.9882	4.102	0.246
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ASAGLLGAHAAAITA	2	719.9104	2.703	0.233
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GLESQTK[272]	2	525.8053	1.587	0.027
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLLQA	2	703.9175	2.43	0.235
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]CYYPR	2	446.2023	1.893	0.223
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DHAVDLIQK[272]GYM[147]R	3	617.3326	2.33	0.111
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLLHEGK[272]	2	549.3497	2.055	0.216
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LLLHEGK	2	545.3426	2.273	0.165
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLLFSPSVVHL	2	684.9222	3.02	0.269
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLLFSPSVVH	2	628.3802	2.685	0.21
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	IYLAPTLAASR	2	658.3903	2.453	0.254
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLQALVR	2	478.8223	2.929	0.141
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LTVAAPPSGGPGFLSIERPD	3	709.0518	2.138	0.292
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]PNMIPDGDFNSYVR	2	884.9259	2.47	0.185
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLLQ	2	668.3989	2.165	0.178
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LSGM[147]AIADVTLLSGFHALR	3	710.0576	3.096	0.102
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	ALLHLLLHEGK	3	508.6522	3.208	0.027
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]DDFVIPDISEPGTWK[272]	4	502.5133	2.753	0.113
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LLLFSPSVVHLG	2	713.433	2.199	0.322
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LAPTLAASR	2	520.3166	1.803	0.107
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]ITQVLH	2	427.7645	1.675	0.067
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]LAPTLAASR	2	522.3201	1.986	0.042
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]HLVPGAPFLL	2	604.3696	1.739	0.11
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	EVQLVAHSPWLK[272]	2	845.9917	3.805	0.254
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]FNSYVR	2	465.2517	1.541	0.064
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	LTVAAPPSGGPGFLSIER	3	637.0229	3.062	0.261
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	SVK[272]	1	617.4104	0.88	0.378
142	IPI00654875	1	51	1.25	0.2	7	5	C4B Complement C4-B precursor	n[145]GVAHNNLM[147]AM[147]	2	617.2955	1.583	0.15
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	GCLTQLYENAFFR	2	874.4184	4.361	0.506
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	GDSGGPLVCK	2	629.8178	3.324	0.514
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]GCLTQLYENAFFR	2	876.4219	3.76	0.42
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]GDSGGPLVCK[272]	2	633.8249	3.283	0.423
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]GVNVCQETCTK[272]	2	781.3579	4.353	0.511
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]IYSGILNLSDITK[272]	2	863.0055	4.966	0.345

143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]YGTQGSSGYSLR	2	710.3529	3.522	0.545
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	EIIIHQNYK	3	479.9453	2.875	0.336
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	TLPEPCHSK	3	446.5617	3.089	0.245
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]EIIIHQNYK[272]	3	482.6167	3.467	0.291
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]TLPEPCHSK[272]	3	449.2331	3.233	0.078
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]IAYGTQGSSGYSLR	2	802.4135	4.105	0.4
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]VSEGNHDIALIK[272]	2	792.4534	4.29	0.356
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	VSEGNHDIALIK	2	788.4463	4.044	0.334
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]EIIIHQNYK[272]	2	723.4214	3.517	0.306
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]VSSVEECQK[272]	2	671.8329	3.17	0.387
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	DSVTGTLPK	2	599.3456	2.531	0.303
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]LCNTGDNSVCTTK[272]	2	868.3899	3.761	0.407
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	CLLFSFLPASSINDM[147]EK	3	753.0404	2.882	0.227
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	YSPGGTPTAIK	2	686.3852	2.672	0.317
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]LSM[147]DGSPTR	2	562.2803	2.766	0.241
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]DACK[272]GDSGGPLVCK[272]	3	625.305	3.437	0.227
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]VLTPDAFVCR	2	655.8395	1.819	0.171
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]QK[272]PICLPSK[272]	3	497.9632	3.026	0.31
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]DSVTGTLPK[272]	2	603.3527	2.46	0.212
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]EK[272]GEIQNILQK[272]	3	578.0158	3.64	0.133
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	VLTPDAFVCR	2	653.836	2.286	0.152
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	VSSVEECQK	2	667.8258	2.038	0.14
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]TLPEPCHSK[272]	2	673.346	2.632	0.018
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	LSM[147]DGSPTR	2	560.2768	1.966	0.135
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	DTPFSQIK	2	608.3403	2.388	0.056
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]YQDYK[272]	2	502.7682	1.53	0.045
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	TLPEPCHSK	2	669.3389	1.909	0.01
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]FGCFLK[272]	2	524.7818	1.698	0.02
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	IFQHLAFSDVDVAR	3	586.6493	2.723	0.199
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	VSEGNHDIALIK	3	525.9666	2.106	0.015
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	TGAVSGHSLK[272]	3	414.2425	1.66	0.04
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]IFQHLAFSDVDVAR	3	587.985	3.132	0.149
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]YSPGGTPTAIK[272]	2	690.3923	1.648	0.017
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]TICTYHPN	2	569.7607	1.468	0.184
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]AQYCQM[147]R	2	553.2391	1.764	0.189
143	IPI00654888	1	40.3	0.79	0.17	62	39	KLKB1 Plasma kallikrein precursor	n[145]CQFFTY	1	998.4233	1.511	0.238
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]EFSHLGK[272]	1	1105.6245	2.704	0.288
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	CCESASEDCMAK	2	847.7958	4.096	0.606
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	CCESASEDCM[147]AK	2	855.7933	3.887	0.695
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	DVCDPGNTK	2	637.797	3.076	0.521
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	GQELCADYSENTFTEYK	2	1162.5141	5.726	0.589
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	SCESNSPFPVHPGTAECCTK	2	1256.5164	6.319	0.642
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	SLGECCDVEDSTTCFNAK	2	1170.4608	4.342	0.627
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	VCSQYAAYGEK	2	772.8655	3.798	0.348
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	VPTADLEDVLPLAEDITNILSK	2	1323.7337	4.834	0.455
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]CCESASEDCMAK[272]	2	851.8029	4.28	0.646
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]GQELCADYSENTFTEYK[272]	2	1166.5212	6.078	0.616
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	2	1260.5235	5.232	0.653

144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SLGECCDVEDSTTCFNAK[272]	2	1174.4679	4.977	0.684
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VCSQYAAYGEK[272]	2	776.8726	4.007	0.521
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	2	1327.7408	4.919	0.487
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	3	705.7369	2.6	0.142
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	SCESNSPFPVHPGTAECCTK	3	838.0134	3.191	0.5
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	VPTADLEDVLPLAEDITNILSK	3	882.8249	3.047	0.191
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]ELSSFIDK[272]GQELCADYSENTFTEYK[272]	3	1132.539	7.664	0.627
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	3	840.6848	3.95	0.566
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SPFPVHPGTAECCTK[272]	3	651.9693	4.226	0.492
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	3	885.4963	2.149	0.323
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	AKLPDATPTELAK	3	592.3524	3.252	0.276
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KLCMAALK	3	448.5957	3.176	0.253
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KLCM[147]AALK	3	453.9273	2.708	0.318
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	RTHLPEVFLSK	3	536.3191	3.15	0.347
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]THLPEVFLSK[272]	3	486.9568	2.858	0.163
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]HQPQEFPTYVEPTNDEICEAFR	4	710.8249	2.721	0.286
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	2	1058.1017	3.827	0.349
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	RTHLPEVFLSK	2	803.9751	3.523	0.431
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]RTHLPEVFLSK[272]	2	807.9822	3.92	0.36
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	LCDNLSTK	2	610.3123	2.919	0.285
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]EDFTSLSLVLYSR	2	837.447	3.736	0.323
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]EFSHLGK[272]	2	553.3159	2.958	0.263
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]HLSLLTTLSNR	2	699.9129	3.504	0.264
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]LCDNLSTK[272]	2	614.3194	2.937	0.277
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	FPSGTFEQVSQLVK	3	616.3403	2.872	0.234
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KELSSFIDK	3	496.2925	3.188	0.172
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]ELSSFIDK[272]	1	1226.6872	2.552	0.242
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]DVCDPGNTK[272]	2	641.8041	2.877	0.309
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	THLPEVFLSK	3	484.2854	3.114	0.172
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]LCDNLSTK[272]	1	1227.6316	2.532	0.237
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	ELSSFIDK	2	609.8401	3.559	0.188
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]ELSSFIDK[272]	2	613.8472	2.449	0.126
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]FPSGTFEQVSQLVK[272]	3	619.0117	3.158	0.238
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VM[147]DK[272]YTFELSR	3	564.9672	2.108	0.36
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	ELSSFIDK	1	1218.673	2.311	0.242
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	YTFELSR	1	1055.5521	2.382	0.255
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]DVCDPGNTK[272]	1	1282.601	2.107	0.413
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	EFSHLGK	2	549.3088	2.484	0.267
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	ELPEHTVK	2	616.8535	2.508	0.153
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]THLPEVFLSK[272]	2	729.9316	2.826	0.292
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]AK[272]LPDATPTELAK[272]	3	596.3595	2.987	0.269
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VLEPTLK[272]	1	1087.6966	2.065	0.184
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]RTHLPEVFLSK[272]	3	538.9905	2.521	0.21
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VPTADLEDVLPLAEDITNILSK[272]	4	664.374	3.07	0.015
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]ELPEHTVK[272]	2	620.8606	2.22	0.177
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	NSKFEDCCQEK	3	614.9384	2.917	0.249
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]YTFELSR	1	1059.5592	2.028	0.197
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	THLPEVFLSK	2	725.9245	2.577	0.285

144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	VLEPTLK	2	540.3448	2.119	0.244
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VLEPTLK[272]	2	544.3519	2.321	0.122
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]FEDCCQEK[272]	1	1381.5499	2.155	0.345
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KLCM[147]AALK	2	680.3873	2.549	0.253
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]TSALSAK[272]	2	483.2972	2.861	0.135
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]YTFELSR	2	530.2832	2.117	0.101
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	LPDATPTELAK	2	718.4114	2.562	0.232
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	AKLPDATPTELAK	2	888.025	4.185	0.35
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]LPDATPTELAK[272]	2	722.4185	2.549	0.241
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	FEDCCQEK	2	687.2715	2.27	0.214
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]HLSLLTTLSNR	3	466.9444	2.137	0.188
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KLCMAALK	2	672.3899	2.503	0.13
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	TSALSAK	2	479.2901	2.554	0.043
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]ELPEHTVK[272]	3	414.2428	1.535	0.027
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]AK[272]LPDATPTELAK[272]	2	894.0356	3.474	0.189
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	KFPSGTFEQVSQLVK	4	529.5545	2.217	0.113
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	YTFELSR	2	528.2797	2.135	0.055
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VM[147]DK[272]YTFELSR	2	846.9471	2.245	0.276
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	ELPEHTVK	3	411.5714	1.93	0.068
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SCESNSPFPVHPG	2	774.3406	3.567	0.441
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]EFSHLGK[272]	3	369.213	1.646	0.126
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]FEDCCQEK[272]	2	691.2786	1.373	0.05
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SLGECCDVEDSTTCFNAK[272]	3	783.3144	1.966	0.144
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	EFSHLGK	3	366.5416	2.061	0.021
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SLLVSYTK[272]	2	599.8679	2.948	0.182
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]NSK[272]FEDCCQEK[272]	3	618.9455	2.241	0.12
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]CCESASEDCM[147]AK[272]	2	859.8004	2.133	0.363
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]DATPTELAK[272]	2	617.3501	3.013	0.2
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]VMDK[272]YTFELSR	3	559.6355	2.342	0.066
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	DATPTELAK	2	613.343	2.721	0.145
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	VMDKYTFELSR	3	556.9641	2.082	0.036
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTA	2	860.383	2.361	0.182
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]GPLLK[272]K[272]	3	363.257	1.257	0.134
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]SCESNSPFPVHPGTAECCTK[272]	4	630.7654	2.044	0.025
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	n[145]HQPQEFPTYVEPTNDEICEAFR	3	947.4307	2.939	0.094
144	IPI00742696	1	63.7	1	0.06	16	93	GC vitamin D-binding protein precursor	FM[147]PAAQLPELPDVELPTNK	3	802.7628	2.851	0.009
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	n[145]AIFYM[147]NNPSR	2	686.8436	3.072	0.387
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	n[145]IFFESVYGQCK[272]	2	827.9142	3.045	0.373
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	AIFYM[147]NNPSR	2	684.8401	2.707	0.26
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	n[145]LVVLPFPK[272]	2	600.9016	2.487	0.164
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	LVVLPFPK	2	596.8945	2.773	0.139
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	n[145]VNDAQEYR	2	569.7841	1.871	0.113
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	GCNDSDVLAVAGFALR	2	897.4373	2.513	0.126
145	IPI00743766	1	14.7	0.89	0.03	8	7	FETUB Fetuin-B precursor	SK[272]K[272]	2	395.7731	0.913	0.136
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]GQCLITQSPPYYR	2	858.4219	4.031	0.542
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]LK[272]PVDGHCALESK[272]	3	625.6779	3.999	0.414
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]LK[272]PVDGHCALESK[272]	4	469.5103	3.467	0.486
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	KEEFHEQSFR	3	539.6109	3.534	0.301

146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	LKPVDGHCALESK	3	621.6708	3.384	0.339
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]RPGVYTQVTK[272]	3	479.6203	3.296	0.312
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	RPGVYTQVTK	3	476.9489	3.542	0.197
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	LKPVDGHCALESK	4	466.505	3.397	0.185
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	NPDADEKPWCFIK	3	677.006	2.807	0.245
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]RPGVYTQVTK[272]	2	718.9269	2.751	0.189
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]LIANTLCNSR	2	647.84	2.672	0.269
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	VVLGDQDLK	2	633.8745	2.765	0.243
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]QLLDAK[272]	2	488.3075	1.76	0.087
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]VVLGDQDLK[272]	2	637.8816	2.115	0.126
146	IPI00746623	1	16.1	0.69	0.06	15	14	HABP2 Hyaluronan-binding protein 2 precursor	n[145]DIAILK[272]	2	480.8203	1.803	1
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AELQCPQPAA	1	1213.5704	2.41	0.511
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GYTQQLAFR	1	1223.6532	2.674	0.365
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VVLVAVDK	1	1122.7246	2.728	0.362
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AELQCPQPAA	1	1217.5775	2.368	0.49
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VVLVAVDK[272]	1	1130.7388	2.805	0.36
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ACEPGVDYVYK	2	785.3757	3.48	0.417
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ADIGCTPGSGK	2	666.326	3.782	0.458
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AYYENSPQQVFSTEFEVK	2	1223.5999	5.239	0.474
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	CAEENCFIQK	2	778.8401	4.004	0.496
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	DICEEQVNSLPGSITK	2	1030.0136	4.598	0.506
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	DSCVGSLVVK	2	666.8544	3.015	0.402
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ENEGFTVTAEGK	2	781.3965	5.037	0.468
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EVVADSVWVDVK	2	813.4485	5.332	0.423
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EYVLPSFEVIVEPTEK	2	1080.0854	4.231	0.456
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KQELSEAEQATR	2	835.4471	2.255	0.197
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	M[147]VFQALAQYQK	2	811.9398	4.192	0.408
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SEETKENEGFTVTAEGK	2	1138.5739	6.352	0.496
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGIPIVTSPYQIHFTK	2	1034.5832	4.849	0.534
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SITTDFIPSFR	2	712.3827	3.362	0.442
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SSLSVPYVIVPLK	2	841.5162	2.652	0.28
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TVMVNIENPEGIPVK	2	960.5348	3.991	0.463
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VELLHNPAFCSLATTK	2	1035.5474	4.87	0.573
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VHQYFNVELIQPGAVK	2	1061.5941	6.151	0.544
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VPVAVQGEDTVQSLTQGDGVAK	2	1239.6636	6.357	0.568
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VQLSNDFDEYIMAIEQTIK	2	1269.1515	7.288	0.584
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VQLSNDFDEYIM[147]AIEQTIK	2	1277.1489	6.258	0.58
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VSHSEDDCLAFK	2	838.8922	2.222	0.054
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISDGVR	2	808.4267	4.137	0.487
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ACEPGVDYVYK[272]	2	789.3828	3.475	0.467
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AGDFLEANYMNLQR	2	893.4392	4.428	0.397
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AVYHHFISDGVR	2	772.9082	4.119	0.458
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AYYENSPQQVFSTEFEVK[272]	2	1227.607	6.043	0.594
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AYYTLIGASGQR	2	722.3893	4.625	0.437
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]CAEENCFIQK[272]	2	782.8472	3.311	0.414
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DICEEQVNSLPGSITK[272]	2	1034.0207	3.822	0.479
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DSCVGSLVVK[272]	2	670.8615	3.391	0.186
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DYAGVFSDAGLTFTSSSGQQTAQR	2	1319.6287	4.738	0.524

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ENEGFTVTAEGK[272]	2	785.4036	4.941	0.435
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EPGQDLVVLPLSITTDFIPSFR	2	1294.7083	4.537	0.466
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EVVADSVWVDVK[272]	2	817.4556	4.995	0.466
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EYVLPSFEVIVEPTEK[272]	2	1084.0925	5.461	0.455
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FVTVQATFGTQVVEK[272]	2	971.5481	5.407	0.499
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQDLVVLPLSITTDFIPSFR	2	1181.6606	5.343	0.499
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQGTLSVVTM[147]YH	2	726.8673	4.106	0.459
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQGTLSVVTM[147]YHAK[272]	2	898.4844	4.314	0.458
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]HLIVTPSGCGE	2	651.8188	2.835	0.427
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IAGYALAQM[147]GR	2	655.854	3.719	0.448
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ILLQGTPVAQMTEDAVDAER	2	1151.0975	5.661	0.6
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ILLQGTPVAQM[147]TEDAVDAER	2	1159.0949	5.103	0.579
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LVAYYTLIGASGQR	2	828.4655	4.811	0.555
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SEETK[272]ENEGFTVTAEGK[272]	2	1144.5846	5.79	0.544
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGIPIVTSPYQIHFTK[272]	2	1038.5903	4.279	0.527
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGSDEVQVGQQR	2	717.3587	3.21	0.243
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SNLDEDIIAEENIVSR	2	981.0008	6.356	0.5
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SSLSVPYVIVPLK[272]	2	845.5233	3.383	0.419
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]STVGNSNNYLHLSVLR	2	959.5168	5.668	0.519
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SYTVAIAGYALAQM[147]GR	2	916.4783	5.168	0.524
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TVMVNIENPEGIPVK[272]	2	964.5419	4.124	0.403
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VAYYTLIGASGQR	2	771.9235	4.727	0.485
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAFCSLATTK[272]	2	1039.5545	5.263	0.522
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VHQYFNVELIQPGAVK[272]	2	1065.6012	5.634	0.547
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VPVAVQGEDTVQSLTQGDGVAK[272]	2	1243.6707	6.157	0.621
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VQLSNDFDEYIMAIEQTIK[272]	2	1273.1586	6.908	0.485
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VQLSNDFDEYIM[147]AIEQTIK[272]	2	1281.156	5.951	0.551
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VSHSEDDCLAFK[272]	2	842.8993	4.652	0.584
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VYAYYNLEESCTR	2	900.9063	3.743	0.522
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYGGGYGSTQATF	2	758.3473	3.458	0.487
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYTYLIMNK[272]	2	748.9067	4.03	0.356
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHFISDGVRK	3	627.3479	4.471	0.475
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AYYENSPQQVFSTEFEVK	3	816.069	3.965	0.416
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EGVQKEDIPPADLSDQVPDTESETR	3	1012.4983	5.866	0.515
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GQGTLSVVTM[147]YHAK	3	596.654	4.61	0.476
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GVFVLNKK	3	442.287	2.198	0.315
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KGYTQQLAFR	3	497.9526	5.033	0.445
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KQELSEAEQATR	3	557.3005	2.949	0.265
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KVLLDGVQNPR	3	506.9754	3.503	0.351
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LESEETMVLEAHDAQGDVPVTVTVHDFPGK	3	1177.5873	8.351	0.625
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	3	1182.9189	7.751	0.658
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	IHWESASLLR	3	451.2542	3.596	0.371
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QKPDGVFQEDAPVIHQEMIGGLR	3	948.8324	6.463	0.465
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QKPDGVFQEDAPVIHQEM[147]IGGLR	3	954.164	4.165	0.476
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SEETKENEGFTVTAEGK	3	759.385	2.32	0.054
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGIPIVTSPYQIHFTK	3	690.0579	2.891	0.147
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGQSEDRQPVPGQQMTLK	3	756.0585	4.256	0.373
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGQSEDRQPVPGQQM[147]TLK	3	761.3901	3.876	0.437

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SSLSVPYVIVPLKTGLQEVEVK	3	935.8878	3.211	0.513
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VELLHNPAFCSLATTK	3	690.7007	2.892	0.232
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VPVAVQGEDTVQSLTQGDGVAK	3	826.7781	5.18	0.516
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VQLSNDFDEYIMAIEQTIK	3	846.4367	4.054	0.423
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VQLSNDFDEYIM[147]AIEQTIK	3	851.7684	4.873	0.464
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VSHSEDDCLAFK	3	559.5972	2.177	0.215
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YFKPGM[147]PFDLM[147]VFVTNPDGSPAYR	3	1021.1699	5.48	0.596
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISDGVR	3	539.2869	3.115	0.265
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISDGVRK[272]	3	630.0193	5.173	0.421
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AQGDVPVTVTVHDFPGK[272]	3	685.7086	5.518	0.531
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AYYENSPQQVFSTEFEVK[272]	3	818.7404	3.847	0.462
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DAPDHQELNLDVSLQLPSR	3	764.3965	5.583	0.478
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EGVQK[272]EDIPPADLSDQVPDTESETR	3	1015.1697	6.435	0.463
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FISLGEACK[272]K[272]	3	525.2984	3.721	0.41
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQGTLSVVTM[147]YHAK[272]	3	599.3254	4.515	0.371
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETMVLEAHDAQGDVPVTVTVHDFPGK[272	3	1180.2587	7.564	0.512
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	3	1185.5903	7.226	0.557
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IHWESASLLR	3	452.5899	2.955	0.259
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ILLQGTPVAQM[147]TEDAVDAER	3	773.0657	2.77	0.252
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LSINTHPSQK[272]PLSITVR	3	727.0982	4.644	0.445
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QK[272]PDGVFQEDAPVIHQEMIGGLR	3	951.5038	4.253	0.527
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QK[272]PDGVFQEDAPVIHQEM[147]IGGLR	3	956.8354	5.633	0.473
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RIPIEDGSGEVVLSR	3	590.9992	2.041	0.121
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SEETK[272]ENEGFTVTAEGK[272]	3	763.3921	5.936	0.567
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGIPIVTSPYQIHFTK[272]	3	692.7293	2.917	0.306
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGQSEDRQPVPGQQMTLK[272]	3	758.7299	4.298	0.372
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGQSEDRQPVPGQQM[147]TLK[272]	3	764.0615	2.456	0.098
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SSLSVPYVIVPLK[272]TGLQEVEVK[272]	3	939.8949	4.716	0.485
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]STVGNSNNYLHLSVLR	3	640.0136	4.213	0.442
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SYTVAIAGYALAQM[147]GR	3	611.3213	4.546	0.492
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TELRPGETLNVNFLLR	3	672.7168	3.972	0.382
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAFCSLATTK[272]	3	693.3721	4.542	0.387
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VHQYFNVELIQPGAVK[272]	3	710.7365	3.883	0.204
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VPVAVQGEDTVQSLTQGDGVAK[272]	3	829.4495	4.34	0.381
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VQLSNDFDEYIMAIEQTIK[272]	3	849.1081	4.034	0.391
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VQLSNDFDEYIM[147]AIEQTIK[272]	3	854.4398	2.569	0.339
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VSHSEDDCLAFK[272]	3	562.2686	4.225	0.468
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VTIK[272]PAPETEK[272]	3	549.0014	2.089	0.184
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VVLVAVDK[272]GVFVLNK[272]	3	678.1014	3.558	0.562
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YFK[272]PGM[147]PFDLM[147]VFVTNPDGSPAY	3	1023.8413	5.582	0.569
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QKPDGVFQEDAPVIHQEM[147]IGGLR	4	715.8748	4.565	0.456
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETMVLEAHDAQGDVPVTVTVHDFPGK[272	4	885.4458	5.766	0.601
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	4	889.4446	5.131	0.579
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETMVLEAHDAQGDVPVTVTVHDFPGK[272	5	762.9975	4.019	0.414
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYYIYNEK	2	710.369	2.997	0.376
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YYTYLIM[147]NK	2	752.8971	4.08	0.332
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHFISDGVR	3	537.9512	3.208	0.351
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AKDQLTCNK	3	496.2672	2.515	0.197

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FISLGEACKK	3	521.2913	3.174	0.336
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LSINTHPSQKPLSITVR	3	724.4268	3.942	0.376
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VHQYFNVELIQPGAVK	3	708.0651	4.032	0.125
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VTIKPAPETEK	3	544.9943	2.023	0.179
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AGDFLEANYM[147]NLQR	3	601.2935	4.267	0.342
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AK[272]DQLTCNK[272]	3	500.2743	2.552	0.065
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DSITTWEILAVSM[147]SDK[272]K[272]	3	791.4296	3.659	0.359
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DTWVEHWPEEDECQDEENQK[272]	3	960.7371	4.143	0.331
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DYAGVFSDAGLTFTSSSGQQTAQR	3	880.0882	2.982	0.316
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]EDGK[272]LNK[272]	3	727.7432	3.346	0.157
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GVFVLNK[272]K[272]	3	446.2941	2.933	0.348
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]HQQTVTIPPK[272]	3	479.6203	3.102	0.362
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LK[272]GPLLNK[272]	3	438.966	3.308	0.267
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LVAYYTLIGASGQR	3	552.6461	3.167	0.362
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SLYVSATVILHSGSDM[147]VQAER	3	808.4167	4.337	0.431
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SNLDEDIIAEENIVSR	3	654.3363	2.958	0.308
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LESEETMVLEAHDAQGDVPVTVTVHDFPGKK	4	950.4898	3.991	0.315
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YFKPGM[147]PFDLM[147]VFVTNPDGSPAYR	4	766.1292	3.341	0.259
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISDGVRK[272]	4	472.7663	3.08	0.337
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RQGALELIK[272]K[272]	4	397.7622	2.559	0.248
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VHQYFNVELIQPGAVK[272]	4	533.3042	3.158	0.32
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LESEETMVLEAHDAQGDVPVTVTVHDFPGKK	5	760.5933	2.948	0.319
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHFISDGVR	2	806.4232	3.496	0.401
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KGYTQQLAFR	2	746.4252	1.377	0.107
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VTIKPAPETEK	2	816.9879	3.117	0.31
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YYTYLIMNK	2	744.8996	3.672	0.303
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ADIGCTPGSGK[272]	2	670.3331	3.808	0.377
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FISLGEACK[272]	2	651.3455	2.642	0.303
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYYIYNEK[272]	2	714.3761	2.818	0.297
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GVFVLNK[272]	2	532.839	2.995	0.338
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GVFVLNK[272]K[272]	2	668.9375	2.816	0.413
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GYTQQLAFR	2	614.3338	2.789	0.366
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NTLIIYLDK[272]	2	690.9207	3.447	0.248
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NTMILEICTR	2	692.3494	2.963	0.352
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NTM[147]ILEICTR	2	700.3468	2.329	0.215
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QPSSAFAAFVK[272]	2	720.9081	3.656	0.429
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RIPIEDGSGEVVLSR	2	885.9952	4.715	0.354
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TELRPGETLNVNFLLR	2	1008.5716	4.098	0.333
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TVM[147]VNIENPEGIPVK[272]	2	972.5394	3.959	0.418
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VTIK[272]PAPETEK[272]	2	822.9985	2.521	0.285
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYTYLIM[147]NK[272]	2	756.9042	3.97	0.352
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VVLVAVDKGVFVLNK	3	674.0943	2.88	0.381
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AK[272]DQLTCNK[272]FDLK[272]	3	716.0664	2.89	0.32
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SEFPESWLWNVEDLK[272]EPPK[272]	3	921.4849	3.257	0.412
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TK[272]K[272]QELSEAEQATR	3	684.3868	3.678	0.309
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VYHHFISDGVR	3	491.9288	2.334	0.25
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FISLGEACK	2	647.3384	2.776	0.315
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GLEVTITAR	2	550.3271	2.776	0.391

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GVFVLNKK	2	662.9269	3.117	0.357
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KVLLDGVQNPR	2	759.9594	3.2	0.352
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	NTLIIYLDK	2	686.9136	3.033	0.244
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TGLQEVEVK	2	641.8719	3.451	0.265
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VVLVAVDK	2	561.8659	2.724	0.179
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ASHLGLAR	2	484.7915	2.53	0.264
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GLEVTITAR	2	552.3307	1.398	0.198
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]HQQTVTIPPK[272]	2	718.9269	2.913	0.374
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TGLQEVEVK[272]	2	645.879	2.543	0.253
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VLLDGVQNPR	2	627.868	3.047	0.154
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VVLVAVDK[272]	2	565.873	3.18	0.27
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KVEGTAFVIFGIQDGEQR	3	758.7462	2.854	0.322
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ISLPESLK[272]R	3	444.2814	2.821	0.213
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TIYTPGSTVLYR	3	505.616	2.364	0.301
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LSINTHPSQK[272]PLSITVR	4	545.5755	1.824	0.032
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TK[272]K[272]QELSEAEQATR	4	513.5419	3.989	0.247
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DFDFVPPVVR	1	1334.7226	2.234	0.409
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ASHLGLAR	2	482.788	2.922	0.306
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FISLGEACKK	2	781.4333	3.398	0.224
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGSDEVQVGQQR	2	715.3552	2.962	0.207
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AGDFLEANYM[147]NLQR	2	901.4366	2.923	0.284
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AVLYNYR	2	521.7938	2.78	0.255
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FISLGEACK[272]K[272]	2	787.444	3.788	0.287
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TIYTPGSTVLYR	2	757.9204	2.458	0.048
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LKGPLLNK	3	434.9589	3.126	0.243
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VPVTVTVHDFPGK	3	559.3225	3.131	0.406
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	DFDFVPPVVR	1	1330.7155	1.95	0.397
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LVLSSEK	1	1055.646	2.581	0.215
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GYTQQLAFR	1	1227.6603	2.579	0.294
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	DQLTCNK	2	574.2836	2.809	0.303
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GVFVLNK	2	528.8319	2.981	0.247
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]CCEDGMR	2	525.1768	2.188	0.399
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYYIYNEK[272]	3	476.5865	2.43	0.224
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISDGVR	4	404.717	2.218	0.184
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AEDLVGK	1	1011.5834	2.548	0.166
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AVLYNYR	1	1038.5732	2.408	0.212
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GLEVTITAR	1	1103.6541	2.44	0.314
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LVLSSEK[272]	1	1063.6602	2.45	0.21
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IPIEDGSGEVVLSR	2	807.9446	2.622	0.286
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	IPIEDGSGEVVLSRK	3	627.0264	3.063	0.214
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TM[147]QALPYSTVGNSNNYLHLSVLR	4	685.3568	2.436	0.253
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GLEVTITAR	1	1099.647	2.249	0.271
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AVLYNYR	1	1042.5803	2.166	0.254
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SVQLTEK[272]	1	1092.6504	2.282	0.185
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	HQQTVTIPPK	2	714.9198	1.681	0.343
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DFDFVPPVVR	2	667.8649	2.619	0.238
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DQLTCNK[272]	2	578.2907	2.601	0.245
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IHWESASLLR	2	678.3813	2.683	0.272

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QELSEAEQATR	2	703.3556	3.029	0.313
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GEK[272]PNLSYIIGK[272]	3	584.3525	3.921	0.336
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AQGDVPVTVTVHDFPGK[272]K[272]	4	582.5826	3.919	0.345
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TIYTPGSTVLYR	1	1510.8265	1.55	0.358
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AEDLVGK[272]	1	1019.5976	2.551	0.159
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SVQLTEK	2	542.8217	2.79	0.188
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AEDLVGK[272]	2	510.3024	2.555	0.184
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AEDLVGK	2	506.2953	2.251	0.08
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SVQLTEK[272]	2	546.8288	2.395	0.144
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHF	3	330.1756	2.925	0.324
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AVYHHFISDGVR	3	515.6079	3.264	0.389
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IWDVVEK[272]	2	588.847	2.621	0.14
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VVPEGIR	2	457.283	2.376	0.193
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SVQLTEK	1	1084.6362	2.258	0.098
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	IWDVVEK	2	584.8399	2.529	0.184
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VLLDGVQNPR	2	625.8644	2.488	0.157
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VPVTVTVHDFPGK[272]	3	561.9939	3.16	0.35
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SDDKVTLEER	3	491.2632	1.537	0.064
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]EDGK[272]	3	561.3019	2.295	0.208
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IFTVNHK[272]	3	382.9006	2.443	0.207
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NK[272]LTQSK[272]	3	417.598	2.35	0.265
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	IHWESASLLR	2	676.3777	2.923	0.193
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TKKQELSEAEQATR	3	680.3797	2.875	0.242
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SSLSVPYVIVPLK[272]	3	564.018	3.094	0.144
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	5	711.7571	2.569	0.12
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EKPNLSYIIGK	3	561.3382	3.108	0.334
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEKEDGK	3	557.2948	2.364	0.196
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SDDK[272]VTLEER	1	1479.7894	2.383	0.226
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VFLDCCNYITELR	2	912.9173	2.821	0.266
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SDDKVTLEER	2	736.3912	3.225	0.297
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VPVTVTVHDFPGK[272]K[272]	4	489.7966	1.969	0.08
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VVPEGIR	1	913.5588	1.771	0.215
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AQGDVPVTVTVHDFPGK	3	683.0372	3.014	0.356
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AKDQLTCNK	2	743.8971	3.237	0.321
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AK[272]DQLTCNK[272]	2	749.9078	3.337	0.238
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	5	766.1965	3.224	0.154
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LM[147]NIFLK[272]	2	591.8616	2.495	0.131
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RAPSTWLTAYVVK	3	592.6812	3.507	0.282
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGKK	5	763.7923	2.446	0.068
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QLANGVDR	2	508.7839	2.116	0.205
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EYVLPSFEVIVEPTEK[272]	3	723.0641	2.84	0.189
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SNNYLHLSVLR	3	487.2761	3.104	0.315
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	DSITTWEILAVSM[147]SDK	2	1046.5352	3.073	0.285
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEKEDGK	2	835.4385	3.26	0.281
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]EDGK[272]LNK[272]	4	546.0593	2.242	0.125
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LVLSSEK	2	528.3266	2.076	0.087
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VVPEGIR	2	455.2795	2.015	0.237
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VVPEGIR	1	909.5517	1.756	0.185

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VQLSNDFDEYIM[147]AIEQTIK	4	639.0781	2.941	0.114
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LVLSSEK[272]	2	532.3337	1.882	0.125
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QPSSAFAAFVK	2	716.901	2.204	0.246
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LKGPLLNK	2	651.9347	2.784	0.226
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	RQGALELIKK	4	394.7569	2.131	0.086
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LM[147]NIFLK	2	587.8545	2.594	0.105
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QGALELIK	2	576.361	1.929	0.088
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LK[272]GPLLNK[272]	2	657.9453	2.651	0.226
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TIYTPGSTVLYR	1	1514.8336	2.441	0.436
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LCRDELCR	3	415.1894	1.44	0.108
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEK	2	550.798	1.708	0.023
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AKDQLTCNKFDLK	3	710.7236	2.694	0.201
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	STVGNSNNYLHLSVLR	3	638.6779	3.467	0.296
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	RQGALELIKK	3	526.0068	3.009	0.065
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RPQDAK[272]NTM[147]ILEICTR	4	560.5455	3.168	0.351
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KLVLSSEK	3	441.9502	2.618	0.163
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FLTTAK[272]	2	484.8046	2.068	0.107
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ELLHNPAFCSLATTK[272]	3	660.3493	3.941	0.25
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TVM[147]VNIENPEGIPVK	2	968.5323	3.762	0.407
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]EDGK[272]	2	841.4492	3.512	0.192
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QK[272]PDGVFQEDAPVIHQEMIGGLR	4	713.8797	2.917	0.059
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LMNIFLK[272]	2	583.8641	2.13	0.115
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RQGALELIK[272]K[272]	3	530.0139	3.137	0.051
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TFISPIK	2	543.3395	2.318	0.092
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AK[272]DQLTCNK[272]FDLK[272]	4	537.3017	2.351	0.143
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FLYGK[272]K[272]	2	594.3793	2	0.116
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QPVPGQQM[147]TLK	2	761.9242	2.456	0.117
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TFISPIK[272]	2	547.3466	2.153	0.126
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEKEDGKLNK	4	542.0522	2.018	0.191
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ISLPESLK	2	583.8608	2.635	0.19
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ISLPESLK[272]	2	587.8679	2.624	0.099
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SDDK[272]VTLEER	2	740.3983	3.095	0.214
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]	2	554.8051	1.941	0.042
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QGALELIK[272]	2	580.3681	2.563	0.019
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LSINTHPSQKPLSITVR	4	543.572	1.703	0.066
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]APSTWLTAYVVK	2	810.4675	1.94	0.202
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EYVLPSFEVIVEPTEK	3	720.3927	2.539	0.12
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]EDGK[272]	4	421.2282	2.038	0.007
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	IPIEDGSGEVVLSR	2	805.9411	2.476	0.122
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]WEDPGK[272]	2	510.2737	2.053	0.132
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]WEDPGK[272]	1	1019.5401	1.65	0.073
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	NKLTQSK	3	413.5909	2.096	0.203
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GPLLNK	2	461.2977	2.18	0.04
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FISLGEACK[272]	3	434.5661	2.15	0.113
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RQGALELIK[272]	2	658.4187	2.666	0.032
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]APVIHQEM[147]IGGLR	3	527.6282	2.925	0.257
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFISD	2	652.3312	3.35	0.399
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGIPIVTSPYQIHF	2	849.9643	3.635	0.449

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	WEDPGK	2	506.2666	1.587	0.192
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEKEDGKLNK	3	722.4004	2.733	0.176
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YTLIGASGQR	2	605.3391	3.501	0.361
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TVLTPATNHM[147]G	2	651.3357	2.759	0.158
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SLK[272]VVPEGIR	3	462.6288	2.54	0.135
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KLVLSSEK	2	662.4216	3.209	0.294
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DVPVTVTVHDFPGK[272]	3	600.3362	2.532	0.285
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	WEDPGK	1	1011.5259	1.792	0.045
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EVVADSVWVDVK[272]	2	815.4521	3.24	0.186
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VYHHFISDGVR	2	737.3896	3.818	0.388
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EALK[272]LEEK[272]	2	696.4272	2.747	0.149
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QALAQYQK	2	615.3537	3.289	0.31
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SNNYLHLSVLR	2	730.4105	3.798	0.334
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YFK[272]PGM[147]PFDLM[147]VFVTNPDGSPAY	4	768.1328	3.329	0.073
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NEQVEIR	2	516.2837	2.256	0.021
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QVPDTESETR	2	653.3238	2.53	0.477
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GYALAQM[147]GR	2	563.7934	2.852	0.414
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LMNIFLK	2	579.857	2.015	0.067
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FYHPEKEDGK	4	418.2229	2.248	0.079
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EALKLEEK	2	690.4165	2.973	0.08
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KDQLTCNK	3	472.5881	2.836	0.269
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DSITTWEILAVSM[147]SDKK	3	788.7582	3.095	0.162
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SITTDFIPSFR	2	714.3862	3.564	0.373
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IPYSVVR	2	489.2987	1.877	0.088
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YTQQLAFR	2	585.823	2.933	0.303
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VELLHNPAFCSL	2	764.8862	2.821	0.378
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TIK[272]PAPETEK[272]	2	773.4643	2.865	0.34
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGSDMVQAER	2	612.294	3.037	0.317
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VYHHFISDGVR	2	735.3861	2.995	0.365
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FYHPEK[272]	3	370.2058	1.968	0.048
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SPMYSIITPNILR	2	824.9643	3.176	0.355
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FSCQR	2	415.6921	1.625	0.018
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	NKLTQSK	2	619.8827	2.192	0.172
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IEDGSGEVVLSR	2	702.8762	3.367	0.343
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QALAQYQK[272]	2	619.3608	3.112	0.274
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGIPIVTSPYQIH	2	778.4337	3.517	0.383
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ASHLGLAR	3	323.5301	1.951	0.108
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FLTTAK	2	480.7975	1.642	0.035
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGIPIVTSPYQIHF	2	851.9679	3.163	0.312
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YYGGGYGSTQATF	2	756.3437	2.483	0.393
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TVLTPATNH	2	549.3072	2.935	0.306
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHFISD	2	650.3277	2.739	0.386
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VLISLQEAK[272]	2	644.9076	2.303	0.159
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ASHLGLAR	1	964.5687	1.819	0.008
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TVLTPATNHM[147]	2	622.8249	3.275	0.264
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SLK[272]VVPEGIR	2	693.4396	1.835	0.238
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VGKYPK	2	556.353	1.803	0.16
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYTYLIM[147]NK[272]	3	504.9385	2.321	0.002

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LCRDELCR	2	622.2804	2.554	0.026
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHF	3	328.8399	2.305	0.206
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FGLEK[272]	2	441.2704	1.569	0.107
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	YYTYLIM[147]NK	3	502.2671	3.248	0.002
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AKDQLTCNKFDLK	4	533.2946	2.098	0.044
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]QPVPGQQM[147]TLK[272]	2	765.9313	2.308	0.04
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQGTLSVVTMYH	2	718.8699	3.091	0.339
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TLIIYLDK[272]	2	633.8992	2.661	0.238
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SDEVQVGQQR	2	645.332	3.362	0.223
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GSGEVVLSR	2	524.2994	3.102	0.201
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]STEFEVK[272]	2	564.313	2.806	0.212
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YHHFISDGVR	2	687.8554	2.967	0.206
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]CCEDGM[147]R	2	533.1742	1.132	0.171
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	STEFEVK	2	560.3059	2.934	0.188
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SDDK[272]VTLEER	3	493.9346	2.009	0.044
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ELSEAEQATR	2	639.3263	2.902	0.196
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]DSITTWEILAVSM[147]SDK	2	1048.5388	3.763	0.268
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FVTVQATFGTQVVEK[272]	3	648.0345	2.311	0.061
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	ALLALLQLK	2	631.9316	2.845	0.222
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ALLALLQLK[272]	2	635.9387	2.894	0.173
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAFCSL	2	766.8897	2.705	0.382
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SPM[147]YSIITPNILR	2	832.9618	2.883	0.36
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LSINTHPSQKPL	3	538.9824	2.921	0.135
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SSAFAAFVK	2	604.3454	2.607	0.233
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	EALKLEEK	3	460.6134	2.139	0.072
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAFCSLA	2	802.4083	1.838	0.429
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAFC	2	666.8317	2.173	0.309
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EK[272]PNLSYIIGK[272]	3	565.3453	2.61	0.154
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]NK[272]LTQSK[272]	2	625.8933	1.948	0.157
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KNKLTQSK	4	377.4925	1.418	0.048
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LESEETM[147]VLEAHDAQGDVPVTVTVHDFPGK	4	957.4938	2.633	0.009
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SPM[147]YSIITPNILR	3	555.6436	2.852	0.163
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YHHFISDGVR	3	458.906	2.332	0.17
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SSAFAAFVK[272]	2	608.3525	2.559	0.208
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SGIPIVTSPYQIH	2	776.4301	2.527	0.241
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VLYNYR	2	486.2752	2.175	0.166
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VTNPDGSPAYR	2	660.8369	2.797	0.246
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHF	2	494.7597	1.98	0.196
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	LSYIIGK	2	537.3395	2.591	0.093
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TM[147]QALPYSTVGNSNNYLH	3	724.0167	3.387	0.125
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]EALK[272]LEEK[272]	3	464.6205	2.046	0.024
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KLM[147]NIFLK	3	481.6354	2.356	0.216
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IPIEDGSGEVVLSR	3	538.9655	2.305	0.077
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHF	2	492.7562	1.757	0.252
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VFLDCCNYITELR	3	608.9473	1.699	0.069
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHHFIS	2	594.8178	1.783	0.308
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GALELIK[272]	2	516.3388	2.193	0.147
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYTLIGASGQR	2	686.8707	2.59	0.192

147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KQELSEAEQA	2	706.8727	2.317	0.199
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	SEFPESWLWNVEDLKEPPK	3	917.4778	2.767	0.09
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KGPLLNK	3	397.2642	2.681	0.133
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FDFVPPVVR	2	610.3514	2.085	0.208
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	QKPDGVFQED	2	721.8674	2.159	0.147
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YFK[272]PGMPFDL	2	751.9015	2.096	0.175
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VELLHNPAF	2	592.3332	1.634	0.267
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VGKYPK	3	371.2378	1.282	0.115
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GALELIK	2	512.3317	2.205	0.094
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAFVK	2	408.2606	1.708	0.136
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YYIYNEK[272]	2	640.8419	2.4	0.022
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]ILEICTR	2	519.2838	2.103	0.069
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	TIKPAPETEK	3	511.9715	2.56	0.065
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YSVVR	2	384.2303	1.803	0.049
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TELRPGETLNVN	2	743.9027	2.537	0.06
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]GQGTLSVVTM[147]Y	2	658.3379	2.356	0.131
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]AAVYHH	2	421.2255	1.503	0.193
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VTIK[272]PAPE	2	571.8548	1.867	0.129
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YSVVR	1	767.4533	1.967	0.106
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YM[147]NLQR	2	492.7563	1.672	0.14
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	GEKPNLSYIIGK	3	580.3454	3.147	0.058
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]TIYTPGSTVLY	2	679.8699	2.056	0.163
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LSINTHPSQK[272]	3	471.6082	2.633	0.091
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VYAYYNL	1	1049.5425	1.726	0.248
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGEVVLSR	2	495.7887	2.053	0.028
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]HLIVTPSGCG	2	587.2975	1.728	0.131
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	VTIKPAPE	2	567.8477	1.903	0.195
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	KVEGTAF	2	516.2979	1.723	0.155
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	FISDGVR	2	467.2613	1.879	0.07
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]FISDGVR	2	469.2648	2.187	0.054
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IPIVTSPYQIHFTK	2	964.56	2.995	0.43
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]VPGQQM[147]TLK[272]	2	653.3756	1.773	0.086
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]IWDVVEK	2	586.8435	2.029	0.058
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]RPQDAK[272]NTM[147]ILEICTR	3	747.0582	2.737	0.13
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	AAVYHHF	1	984.5051	2.314	0.401
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]YFK[272]PGM[147]PF	2	645.8434	1.63	0.117
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]LSYIIGK[272]	2	541.3466	1.895	0.016
147	IPI00783987	1	76.9	0.72	0.09	992	##	C3 Complement C3 precursor (Fragment)	n[145]SGIPIVTSPYQIHF	3	568.3143	1.905	0.079
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	n[145]VEHSDLSFSK[272]	2	718.8849	3.779	0.445
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	n[145]VEHSDLSFSK[272]	3	479.5923	3.37	0.283
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	VEHSDLSFSK	2	714.8778	3.47	0.434
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	VSGFHPSDIEVDLLK	3	646.0228	3.84	0.332
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	VNHVTLSQPK	3	468.277	2.417	0.194
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	n[145]IQVYSR	2	455.2674	1.816	0.037
148	IPI00796379	1	34.5	1.51	0.35	8	7	B2M B2M protein	VEHSDLSFSK	3	476.9209	2.522	0.059
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	ATEDEGSEQKIPEATNR	2	1078.035	5.122	0.474
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	FATTFYQHLADSK	2	904.9702	5.586	0.442
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	NDNDNIFLSPLSISTAFAM[147]TK	2	1298.1598	5.61	0.604

149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	TSDQIHFFFAK	2	810.9303	4.615	0.494
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]AFLEVNEEGSEAAASTAVVIAGR	2	1218.1304	6.161	0.519
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]ATEDEGSEQK[272]IPEATNR	2	1082.0421	5.633	0.526
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]FATTFYQHLADSK[272]	2	908.9773	4.136	0.498
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]NDNDNIFLSPLSISTAFAM[147]TK[272]	2	1302.1669	6.123	0.525
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]TSDQIHFFFAK[272]	2	814.9374	4.303	0.452
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	ATEDEGSEQKIPEATNR	3	719.0257	4.316	0.432
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	FATTFYQHLADSK	3	603.6492	3.653	0.463
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	TSDQIHFFFAK	3	540.956	3.174	0.359
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]FATTFYQHLADSK[272]	3	606.3206	4.512	0.378
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	KATEDEGSEQK	3	547.953	3.234	0.319
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	KATEDEGSEQKIPEATNR	3	808.4224	4.807	0.39
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]ATEDEGSEQK[272]IPEATNR	3	721.6971	4.138	0.313
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]NDNDNIFLSPLSISTAFAM[147]TK[272]	3	868.447	2.584	0.132
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]TSDQIHFFFAK[272]	3	543.6274	2.621	0.225
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]EVPLNTIIFMGR	2	767.4326	2.112	0.039
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	KATEDEGSEQKIPEATNR	4	606.5686	2.92	0.36
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	EVPLNTIIFMGR	2	765.4291	3.288	0.34
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]EVPLNTIIFM[147]GR	2	775.4301	2.835	0.306
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	ATEDEGSEQK	2	687.3308	2.896	0.377
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]DIPMNPMCIYR	2	771.8565	3.245	0.246
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]DIPM[147]NPM[147]CIYR	2	787.8514	1.997	0.196
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]ATEDEGSEQK[272]	2	691.3379	2.563	0.266
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]DIPMNPM[147]CIYR	2	779.854	3.33	0.249
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	EVPLNTIIFM[147]GR	2	773.4265	2.727	0.081
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	FDTISEK	2	560.3059	2.545	0.082
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]EVPLNTIIFM[147]GR	3	517.2891	3.049	0.102
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]VANPCVK[272]	2	532.7954	1.904	0.15
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]DIPM[147]NPMCIYR	2	779.854	2.327	0.297
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	IPEATNR	2	470.7642	1.64	0.169
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	DIPM[147]NPMCIYR	2	777.8504	2.326	0.363
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]FATTFYQH	2	579.7887	3.023	0.328
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]VWELSK[272]	2	525.3153	1.647	0.042
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]IPEATNR	2	472.7677	1.449	0.001
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]TSDQIHFFF	2	643.3203	2.113	0.38
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]TSDQIHFFFA	2	678.8389	2.73	0.412
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	KATEDEGSEQK	2	821.4258	1.385	0.211
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]YQHLADSK[272]	2	625.3426	2.422	0.254
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]EVPLNTIIFMGR	3	511.9575	1.933	0.035
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]YQHLADSK[272]	3	417.2308	2.906	0.067
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	ATTFYQHLADSK	3	554.6264	2.574	0.145
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]ANRPFLVF	2	554.3252	2.016	0.199
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]LYVSDAFHK[272]	3	456.5902	2.185	0.148
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	YQHLADSK	2	621.3355	1.942	0.138
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	n[145]ATTFYQHLADSK[272]	3	557.2978	2.985	0.043
149	IPI00844156	1	56.4	0.77	0.04	24	48	SERPINC1 SERPINC1 protein	NDNDNIFLSPLSISTAFAM[147]TK	3	865.7756	1.6	0.062
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]APEPISTQSHSVLILFHSDNSGENR	4	720.6156	2.845	0.208
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]AYAPLK[272]K[272]	3	408.2677	2.161	0.223

150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]VETEDQVLATFCGR	2	879.4196	3.852	0.335
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]AYAPLK[272]K[272]	2	611.8979	2.093	0.181
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]SDFSNEER	2	564.2579	2.197	0.145
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	n[145]SDENEQHLGVK[272]	3	515.27	2.488	0.212
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	AYAPLKK	2	605.8872	1.642	0.182
150	IPI00871139	1	8	0.68	0.06	7	7	MASP1 92 kDa protein	AYAPLKK	3	404.2606	1.388	0
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DNCCILDER	1	1316.5224	2.197	0.387
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	AIQLTYNPDESSKPNMIDAATLK	2	1470.7787	5.094	0.532
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	DTVQIHDITGK	2	753.9174	2.746	0.228
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	EGFGHLSPTGTTEFWLGNEK	2	1244.1188	6.265	0.484
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	EGFGHLSPTGTTEFWLGNEK[272]	2	1246.1224	4.795	0.417
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	GIADFLSTYQTK	2	812.4407	3.051	0.398
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLGGAK	2	913.5052	6.014	0.506
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYFIKPLK	2	857.5244	1.858	0.052
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	TSTADYAMFK	2	707.8554	4.457	0.417
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	TSTADYAM[147]FK	2	715.8529	3.863	0.469
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YLQEIYNSNNQK	2	897.4627	4.713	0.457
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PN	2	983.012	4.094	0.355
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PNMIDAATLK[272]	2	1476.7893	4.506	0.499
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DTVQIHDITGK[272]	2	757.9245	2.184	0.264
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]EGFGHLSPTGTTEF	2	812.3922	4.052	0.387
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]EGFGHLSPTGTTEFWLGNEK[272]	2	1248.1259	5.966	0.53
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]GIADFLSTYQTK[272]	2	816.4478	4.5	0.481
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGGAK[272]	2	917.5123	5.644	0.542
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYFIK[272]PLK[272]	2	863.5351	4.383	0.491
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TCGIADFLSTYQTK[272]	2	941.4701	4.266	0.52
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TSTADYAMFK[272]	2	711.8625	4.331	0.491
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TSTADYAM[147]FK[272]	2	719.86	3.398	0.494
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TVQIHDITGK[272]	2	700.411	4.305	0.445
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VAQLEAQCQEPCK[272]	2	913.9292	3.131	0.474
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YEASILTHDSSIR	2	818.4266	2.532	0.345
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YLQEIYNSNNQK[272]	2	901.4698	4.827	0.399
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	AIQLTYNPDESSKPNMIDAATLK	3	980.8549	5.053	0.466
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	AIQLTYNPDESSKPNM[147]IDAATLK	3	986.1865	4.989	0.469
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	CHAGHLNGVYYQGGTYSK	3	761.0262	6.364	0.582
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	DTVQIHDITGK	3	502.9474	3.135	0.345
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	EGFGHLSPTGTTEFWLGNEK	3	829.7483	4.738	0.529
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	FAGGDAGDAFDGFDFGDDPSDK	3	835.0277	4.955	0.548
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KMLEEIMKYEASILTHDSSIR	3	972.1892	4.613	0.483
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLGGAK	3	609.3392	2.59	0.105
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VAQLEAQCQEPCKDTVQIHDITGK	3	1056.1887	5.282	0.512
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YLQEIYNSNNQK	3	598.6442	4.373	0.402
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PNMIDAATLK[272]	3	984.862	3.571	0.279
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	3	990.1936	4.764	0.525
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]CHAGHLNGVYYQGGTYSK[272]	3	763.6976	6.804	0.526
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DTVQIHDITGK[272]	3	505.6188	2.543	0.318
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FAGGDAGDAFDGFDFGDDPSDK[272]	3	837.6991	5.273	0.599
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IHLISTQSAIPYALR	3	609.6918	3.428	0.234

151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLG	3	478.5921	4.393	0.484
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGGAK[272]	3	612.0106	3.661	0.316
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYFIK[272]PLK[272]	3	576.0258	3.663	0.352
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VAQLEAQCQEPCK[272]DTVQIHDITGK[272]	3	1060.1958	6.647	0.576
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YEASILTHDSSIR	3	545.9535	1.619	0.018
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KMLEEIMKYEASILTHDSSIR	4	729.3937	5.105	0.517
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLGGAK	4	457.2563	3.972	0.446
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYFIKPLK	4	429.2659	1.97	0.306
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VAQLEAQCQEPCKDTVQIHDITGK	4	792.3934	4.279	0.383
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGGAK[272]	4	459.2598	2.972	0.326
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYFIK[272]PLK[272]	4	432.2712	2.567	0.337
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VAQLEAQCQEPCK[272]DTVQIHDITGK[272]	4	795.3987	5.295	0.4
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KMLEEIMKYEASILTHDSSIR	5	583.7164	4.728	0.502
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYFIKPLK	3	572.0187	2.77	0.129
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	RLDGSVDFK	3	439.5822	4.1	0.236
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VGPEADKYR	3	438.9104	3.208	0.4
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YLQEIYNSNNQK[272]	3	601.3156	3.845	0.254
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	4	742.897	3.57	0.303
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]RLDGSVDFK[272]K[272]	4	399.9913	1.638	0.129
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LDGSVDFKK	2	714.9142	2.941	0.408
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YEASILTHDSSIR	2	816.423	3.879	0.4
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IHLISTQSAIPYALR	2	914.0341	1.996	0.145
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIMK	3	486.6122	1.855	0.026
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]RLDGSVDFK[272]	3	442.2536	2.947	0.247
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VAQLEAQCQEPCK	2	909.9221	2.805	0.445
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DNCCILDER	2	658.7648	2.981	0.348
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]RLDGSVDFK[272]K[272]	2	798.9754	3.28	0.353
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	EGFGHLSPTGTTEFWLGNEK[272]	3	831.084	4.4	0.389
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	AIQLTYNPDESSKPNMIDAATLK	4	735.893	3.439	0.235
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIM[147]KYEASILTHDSSIR	4	737.3912	3.365	0.256
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	IHLISTQSAIPYALR	3	608.3561	2.856	0.191
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LDGSVDFK[272]	1	1168.6453	2.342	0.262
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	RLDGSVDFKK	2	792.9647	3.362	0.24
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VELEDWNGR	2	629.3148	2.308	0.088
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YEASILTHDSSIR	3	544.6178	2.921	0.077
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]EGFGHLSPTGTTEFWLGNEK[272]	3	832.4197	3.443	0.259
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VGPEADK[272]YR	3	441.5818	2.251	0.103
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VELEDWNGR	1	1257.6223	2.928	0.147
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VELEDWNGR	1	1261.6294	2.673	0.18
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VELEDWNGR	2	631.3183	3.031	0.313
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LDGSVDFK	1	1160.6311	2.327	0.233
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLG	3	477.2564	3.871	0.321
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGGA	3	521.2783	3.566	0.346
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LDGSVDFK[272]	2	584.8263	2.144	0.151
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VGPEADK[272]	2	502.2868	2.179	0.28
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TVQIHDITGK[272]	3	467.2764	2.782	0.376
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LDGSVDFK	2	580.8192	1.873	0.032
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHL	3	459.5849	3.542	0.315

151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]MLEEIMK[272]	2	591.3293	2.962	0.134
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	RLDGSVDFK	2	658.8697	2.103	0.095
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VGPEADKYR	2	657.8619	2.808	0.375
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VGPEADK[272]YR	1	1322.7308	2.449	0.2
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KMLEEIMK	2	721.4172	2.78	0.129
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]RLDGSVDFK[272]	2	662.8768	2.291	0.16
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]VGPEADK[272]YR	2	661.869	2.769	0.276
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]RLDGSVDFK[272]K[272]	3	532.986	1.735	0.026
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIM[147]K	2	737.4121	2.285	0.204
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	VGPEADK	2	498.2797	2.189	0.169
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]SSK[272]PNM[147]IDAATLK[272]	3	608.6802	2.767	0.347
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]AIQLTYNPDESSK[272]PNM[147]IDAATLK[272]	2	1484.7868	3.126	0.486
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	RLDGSVDFKK	4	396.986	2.457	0.278
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	MLEEIMK	2	587.3222	2.417	0.101
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	M[147]LEEIM[147]K	2	603.3171	2.223	0.107
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHH	3	421.8903	2.967	0.282
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TIGEGQQHHLGGAK[272]	3	574.316	3.612	0.27
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LDGSVDFKK	3	476.9452	1.837	0.188
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]MLEEIM[147]K[272]	1	1197.6462	1.765	0.072
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	IVNLKEK	2	632.4111	2.819	0.118
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	MLEEIM[147]K	1	1189.632	1.934	0.082
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]MLEEIM[147]K[272]YEASILTHDSSIR	4	668.3485	2.549	0.25
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LDGSVDFK[272]K[272]	2	720.9248	2.054	0.087
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LDGSVDFK[272]K[272]	3	480.9523	1.955	0.135
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	MLEEIM[147]K	2	595.3196	2.315	0.031
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]TSTADYAM[147]FK[272]	3	480.2424	1.865	0.191
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLGG	3	496.2636	3.146	0.254
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LISTQSAIPYALR	2	788.9626	4.069	0.373
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	TVQIHDITGK	2	696.4039	3.838	0.335
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	TSTADYAM[147]FK	3	477.571	2.856	0.018
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]GIADFLSTYQTK[272]	3	544.6343	2.582	0.283
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DCQDIANK[272]	2	620.7989	2.82	0.28
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LYFIKPLK	2	721.4684	3.112	0.352
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGG	2	745.8953	3.614	0.358
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQHHLG	2	715.381	3.461	0.335
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	DCQDIANK	2	616.7918	2.496	0.194
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IVNLK[272]EK[272]	2	638.4217	2.263	0.11
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLG	2	717.3845	3.463	0.384
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	GLYFIKPLK	2	749.9791	2.89	0.437
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYFIK[272]	2	622.3681	2.709	0.342
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]SGLYFIK[272]PLK[272]	3	533.3396	3.119	0.258
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	AIQLTYNPDESSKPN	2	979.0049	4.288	0.405
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	FFTSHNGM[147]QF	2	686.3112	2.747	0.326
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	SGLYFIKPLK	2	793.4951	3.781	0.265
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYFIK	2	618.361	2.567	0.286
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]GLYFIK[272]PLK[272]	2	755.9898	3.012	0.296
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LYFIK[272]PLK[272]	2	727.479	3.091	0.256
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DTVQIHDITG	2	621.826	2.288	0.391

151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LYFIKPLK	3	481.3147	1.885	0.271
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]SGLYFIK[272]PLK[272]	2	799.5058	3.404	0.239
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	DTVQIHDITG	2	619.8224	2.759	0.286
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHLGG	3	497.5993	2.822	0.184
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FFTSHNGM[147]QF	2	688.3147	2.877	0.265
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	SGLYFIKPLK	3	529.3325	2.977	0.136
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]ISTQSAIPYALR	2	732.4206	3.391	0.321
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQHHL	2	688.8738	2.869	0.216
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YEASILTHDSSIR	4	409.7169	1.687	0.015
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YEASILTHDS	2	638.3144	1.862	0.352
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIMKYEASILTHDSSIR	4	733.3924	3.879	0.427
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IHLISTQ	2	478.2883	2.054	0.266
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KMLEEIM[147]KYEASILTHDSSIR	4	733.3924	3.519	0.472
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FFTSHN	2	448.7228	2.161	0.188
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYFIK[272]PL	2	727.4365	2.447	0.395
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]EGFGHLSPTGTT	2	674.3367	2.533	0.223
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YEASILTHDS	2	640.318	2.22	0.234
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FFTSH	2	391.7013	1.92	0.227
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIM[147]KYEASILTHDSSIR	3	982.8525	2.812	0.121
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]DCQDIANK[272]	1	1240.5905	2.453	0.308
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IHLISTQSAIPY	2	743.923	2.414	0.205
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	DCQDIANK	1	1232.5763	2.406	0.352
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	KM[147]LEEIMKYEASILTHDSSIR	3	977.5208	3.306	0.46
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYFIKPL	2	723.4294	1.875	0.23
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YFIK[272]PLK[272]	2	670.937	2.033	0.13
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YVATR	2	375.2189	1.994	0.107
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	GLYFIKPLK	3	500.3218	2.139	0.144
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]HLISTQSAIPYALR	2	857.4921	2.743	0.248
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	M[147]IDAATLK	2	579.8312	2.801	0.147
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	YFIKPLK	2	664.9263	2.041	0.169
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]SILTHDSSIR	3	424.9058	2.689	0.065
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	FFTSHN	2	446.7192	1.767	0.192
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]IHLISTQSAIPYAL	2	835.9835	1.722	0.273
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	IHLISTQSAIPY	2	741.9194	2.17	0.169
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	MIDAATLK	2	571.8338	2.63	0.131
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	FFTSH	2	389.6978	1.74	0.137
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]GLYFIK[272]PLK[272]	3	504.3289	2.159	0.114
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	IHLISTQ	2	476.2848	1.79	0.173
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FFTSH	1	782.3954	1.633	0.216
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]MIDAATLK[272]	2	575.8409	2.197	0.172
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	TCGIADFLSTYQTK	2	937.463	4.226	0.314
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	SAIPYALR	2	515.8059	2.267	0.008
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LYFIK[272]PLK[272]	3	485.3218	2.426	0.045
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]YEASILTHD	2	596.802	1.94	0.09
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTIGEGQQH	2	561.7988	1.874	0.079
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	QSGLYF	1	854.4407	1.393	0.317
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]FIK[272]PLK[272]	3	393.2726	2.208	0.024
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	IHLISTQSAIPYAL	2	833.98	1.798	0.058

151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	FFTSH	1	778.3883	1.635	0.134
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]QSGLYF	1	858.4478	1.427	0.281
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	n[145]LTIGEGQQH	2	563.8023	1.31	0.145
151	IPI00877703	1	62.2	N_A	N_A	N_A	99	FGG 52 kDa protein	LTYAYF	1	917.4768	1.334	0.289
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ETLFSVMPGLK	2	751.426	4.357	0.421
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ETLFSVM[147]PGLK	2	759.4235	3.992	0.419
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ITFELVYEELLK	2	889.011	3.679	0.459
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ETLFSVMPGLK[272]	2	755.4331	3.501	0.447
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ETLFSVM[147]PGLK[272]	2	763.4306	3.517	0.46
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]QLGLPGPPDVPDHAAYHPFR	2	1164.6039	4.527	0.561
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	DQFNLIVFSTEATQWRPSLVPASAENVNK	3	1181.2861	5.92	0.573
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ITFELVYEELLKR	3	645.0434	3.128	0.455
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ITFELVYEELLK[272]R	3	647.7148	3.584	0.375
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]NPLVWVHASPEHVVVTR	3	695.3889	3.303	0.383
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]QLGLPGPPDVPDHAAYHPFR	3	776.7384	3.683	0.368
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]VPDHAAYHPFR	3	485.2534	4.443	0.388
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]YIFHNFMER	3	467.5682	3.928	0.357
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	NPLVWVHASPEHVVVTR	4	520.7917	4.709	0.506
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]QLGLPGPPDVPDHAAYHPFR	4	582.8056	2.51	0.108
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]NPLVWVHASPEHVVVTR	4	521.7935	3.511	0.316
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	GPDVLTATVSGK	2	712.9091	3.601	0.403
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	GSEMVVAGK	2	579.321	3.157	0.35
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	LGVYELLLK	2	664.4211	3.783	0.349
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]LGVYELLLK[272]	2	668.4282	2.954	0.252
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GYFVHYFAPEGLTTM[147]PK[272]	3	754.7218	4.43	0.402
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GPDVLTATVSGK[272]	2	716.9162	3.112	0.39
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GSEMVVAGK[272]	2	583.3281	3.008	0.227
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GSEM[147]VVAGK[272]	2	591.3256	2.579	0.393
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	AGFSWIEVTFK	2	782.9298	3.932	0.318
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]YIFHNFM[147]ER	3	472.8999	2.371	0.238
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	GSEM[147]VVAGK	2	587.3185	1.915	0.175
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ITFELVYEELLK[272]	2	893.0181	3.429	0.274
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]NVVFVIDK[272]	2	611.3759	2.165	0.166
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]SPEQQETVLDGNLIIR	2	978.5296	4.528	0.377
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	NVVFVIDK	2	607.3688	2.01	0.191
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]LALDNGGLAR	2	572.3338	2.311	0.156
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	RLGVYELLLK	3	495.3168	3.25	0.11
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	LALDNGGLAR	2	570.3302	2.289	0.043
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]YIFHNFMER	2	700.8487	2.717	0.166
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ETLFSVMPGLK	3	501.2864	2.829	0.124
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ILDDLSPR	2	536.8096	2.837	0.092
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]YIFHNFM[147]ER	2	708.8462	2.401	0.183
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ETLFSVMPGLK[272]	3	503.9578	2.729	0.11
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ETLFSVM[147]PGLK[272]	3	509.2895	2.297	0.107
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]RLGVYELLLK[272]	3	497.9882	2.703	0.069
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ITFELVYEELLK[272]	3	595.6811	2.701	0.158
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ASPEHVVVTR	3	412.2349	2.659	0.256
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	TGLLLLSDPDK	2	726.4271	1.808	0.124

152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]LGVYELLLK[272]	3	445.9545	3.013	0.035
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]EQQETVLDGNLIIR	2	886.4872	4.618	0.339
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]DVLTATVSGK[272]	2	639.879	3.211	0.364
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	DVLTATVSGK	2	635.8719	3.276	0.313
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ASPEHVVVTR	3	413.5706	2.214	0.09
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GPDVLTATVSGK[272]	3	478.2799	1.851	0.236
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]ASPEHVVVTR	2	619.8523	3.187	0.406
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]AGFSWIEVTFK	2	784.9333	2.937	0.137
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ITFELVYEELLK	3	593.0097	2.541	0.114
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]PEQGVEVTGQYER	2	818.4084	3.286	0.191
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]PTVGETNPR	2	557.8023	2.412	0.24
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]GVYELLLK[272]	2	611.8861	3.001	0.124
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]NPLVWVH	2	504.791	1.941	0.296
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]LDNGGLAR	2	480.2732	1.943	0.069
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]VPDHAAYHPFR	4	364.1919	2.422	0.099
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]WKETLFSVM[147]PGLK	3	659.3769	4.075	0.334
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]FSSHVGGTLGQF	2	690.8551	2.617	0.303
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	HLQM[147]DIH	3	350.5114	2.378	0.095
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	ASPEHVVVTR	2	617.8488	2.549	0.159
152	IPI00878609	1	33.5	0.69	0.02	8	3	ITIH4 Protein	n[145]HLQM[147]DIH	3	351.8471	1.653	0.128
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	CSVFYGAPSK	2	692.8413	4.156	0.508
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EMSGSPASGIPVK	2	770.4136	4.147	0.436
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GLEEELQFSLGSK	2	858.962	5.125	0.492
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ITQVLHFTK	2	683.9139	3.962	0.391
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LTVAAPPSGGPGFLSIER	2	955.0307	3.152	0.543
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VDVQAGACEGK	2	701.8445	3.485	0.518
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ASAGLLGAHAAAITAY	2	801.442	4.133	0.557
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHY	2	620.8076	3.784	0.508
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHYY	2	702.3393	3.831	0.484
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHYYYM[147]	2	857.3886	3.96	0.462
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]CSVFYGAPSK[272]	2	696.8484	3.605	0.505
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DFALLSLQVPLK[272]	2	816.5024	3.455	0.45
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EELVYELNPLDHR	2	885.9608	4.22	0.472
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EMSGSPASGIPVK[272]	2	774.4207	4.123	0.379
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EM[147]SGSPASGIPVK[272]	2	782.4182	3.423	0.423
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EPFLSCCQFAESLR	2	933.4124	3.714	0.583
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GHLFLQTDQPIYNPGQR	2	1064.5565	5.825	0.54
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GLEEELQFSLGSK[272]	2	862.9691	4.809	0.498
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GSFEFPVGDAVSK[272]	2	814.4322	4.699	0.476
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LSCCQFAESLR	2	746.8305	3.205	0.529
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LTVAAPPSGGPGFLSIER	2	957.0343	4.08	0.218
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]TTNIQGINLLFSSR	2	854.4792	5.063	0.513
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AEFQDALEKLNMGITDLQGLR	3	881.4705	6.043	0.517
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GPEVQLVAHSPWLK	3	614.3526	4.21	0.408
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHYYYM[147]ILSR	3	728.3619	4.226	0.445
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DFALLSLQVPLK[272]DAK[272]	3	697.4244	4.647	0.48
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GHLFLQTDQPIYNPGQR	3	710.0401	5.031	0.376
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GPEVQLVAHSPWLK[272]	3	617.024	3.685	0.406

153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLATLCSAEVCQCAEGK[272]	3	722.333	4.18	0.507
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITPGK[272]PYILTVPGHLDEM[147]QLDIQAR	3	1037.2368	5.251	0.365
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]RGHLFLQTDQPIYNPGQR	3	762.0738	5.052	0.404
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VGLSGM[147]AIADVTLLSGFHALR	3	763.4233	4.694	0.422
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]YVSHFETEGPHVLLYFDSVPTSR	3	942.1403	6.174	0.531
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ITPGKPYILTVPGHLDEMQLDIQAR	4	772.1772	3.963	0.414
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ITPGKPYILTVPGHLDEM[147]QLDIQAR	4	776.1759	5.288	0.551
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITPGK[272]PYILTVPGHLDEMQLDIQAR	4	774.1807	6.085	0.461
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITPGK[272]PYILTVPGHLDEM[147]QLDIQAR	4	778.1794	2.699	0.069
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AEFQDALEK	2	665.8537	3.858	0.355
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FACYYPR	2	555.2551	2.206	0.17
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LGQYASPTAK[272]	2	662.3792	3.844	0.386
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AEFQDALEKLNM[147]GITDLQGLR	3	886.8022	4.25	0.428
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DFALLSLQVPLKDAK	3	693.4173	3.988	0.442
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GPEVQLVAHSPWLK[272]	3	615.6883	3.613	0.322
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	KYVLPNFEVK	3	552.9994	2.879	0.147
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ITPGKPYILTVPGHLDEM[147]QLDIQAR	3	1034.5654	4.917	0.418
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ITQVLHFTK	3	456.2784	2.988	0.284
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DK[272]GQAGLQR	3	420.9136	3.407	0.36
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EPFLSCCQFAESLR	3	622.6107	2.858	0.354
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLLQALVR	3	592.3653	3.268	0.352
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITQVLHFTK[272]	3	458.9498	3.398	0.281
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]TTNIQGINLLFSSR	3	569.9885	2.805	0.291
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VLQIEK[272]EGAIHR	3	561.0048	3.426	0.317
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]YIYGK[272]PVQGVAYVR	3	634.3678	2.501	0.405
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SHKPLNM[147]GK	4	362.7106	2.598	0.371
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]MRPSTDTITVMVENSHGLR	4	572.7957	5.056	0.307
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]STQDTVIALDALSAYWIASHTTEER	4	731.3705	3.342	0.297
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EGAIHREELVYELNPLDHR	5	487.6577	2.881	0.221
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DFALLSLQVPLK	2	812.4953	3.688	0.363
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DHAVDLIQK	2	659.8776	3.61	0.293
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EM[147]SGSPASGIPVK	2	778.4111	3.563	0.457
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LNM[147]GITDLQGLR	2	743.906	3.41	0.381
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VEYGFQVK	2	625.3506	2.739	0.251
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DHAVDLIQK[272]	2	663.8847	3.194	0.253
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GLQDEDGYR	2	598.7869	2.88	0.329
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITQVLHFTK[272]	2	687.921	3.949	0.349
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VEYGFQVK[272]	2	629.3577	2.729	0.346
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VLSLAQEQVGGSPEK[272]	2	915.5142	3.649	0.451
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DKGQAGLQR	3	418.2422	3.421	0.265
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AIADVTLLSGFHALR	3	576.669	4.098	0.39
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ALLHLLLHEGK[272]	3	511.3236	3.916	0.369
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FGLLDEDGK[272]K[272]TFFR	3	702.3967	2.659	0.366
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FGLLDEDGK[272]K[272]TFFR	4	527.0494	2.623	0.329
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNM[147]GK[272]	4	365.7159	1.622	0.034
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LGQYASPTAK	2	658.3721	3.374	0.31
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ADGSYAAWLSR	2	670.8394	3.681	0.33
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AEMADQAAAWLTR	2	789.3968	3.115	0.376

153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FGLLDEDGK[272]K[272]	2	777.4486	3.327	0.363
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LNM[147]GITDLQGLR	2	745.9095	3.401	0.326
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AEM[147]ADQAAAWLTR	2	797.3942	3.361	0.321
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GLCVATPVQLR	2	673.8739	2.58	0.407
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LNMGITDLQGLR	2	737.9121	3.365	0.22
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SCGLHQLLR	2	608.8242	2.867	0.302
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VGDTLNLNLR	2	629.8654	2.679	0.19
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	FGLLDEDGKKTFFR	3	698.3896	3.087	0.297
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DHAVDLIQK[272]	3	442.9255	3.57	0.152
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GLQDEDGYR	1	1196.5665	2.57	0.3
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LNMGITDLQGLR	2	735.9085	3.131	0.288
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ASSFLGEK[272]	2	563.821	2.836	0.232
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]TYNVLDM[147]K[272]	2	644.3465	1.719	0.281
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VLQIEKEGAIHR	3	558.3334	3.607	0.24
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ASSFLGEK	2	559.8139	1.671	0.158
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GSFEFPVGDAVSK	2	810.4251	3.529	0.313
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	KYVLPNFEVK	2	828.9955	3.814	0.224
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]TYNVLDMK[272]	2	636.3491	2.97	0.146
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EVQLVAHSPWLK	3	562.9945	3.755	0.333
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SFQDLSPVIHR	3	481.6006	3.483	0.348
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SHKPLNM[147]GK	2	724.4138	2.044	0.214
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	YVLPNFEVK	2	694.9005	2.675	0.216
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AEFQDALEK[272]	2	669.8608	3.67	0.174
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VDVQAGACEGK[272]	2	705.8516	3.878	0.382
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VVEEQESR	2	560.2918	2.084	0.38
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AIADVTLLSGFHALR	3	575.3333	3.071	0.402
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EAPKVVEEQESR	3	560.9687	2.849	0.27
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	FGLLDEDGKK	3	514.6277	2.275	0.294
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AACAQLNDFLQEYGTQGCQV	3	799.0151	2.752	0.27
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LELSVDGAK[272]	2	610.3605	3.274	0.238
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNM[147]GK[272]	2	730.4245	1.698	0.204
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]YVLPNFEVK[272]	2	698.9076	1.903	0.149
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	TYNVLDM[147]K	2	640.3394	2.201	0.094
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LTSLSDR	2	468.2676	2.423	0.146
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LTVAAPPSGGPGFLSIER	3	638.3586	4.106	0.332
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VTASDPLDTLGSEGALSPGGVASLLR	3	876.4718	3.352	0.189
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VEASISK[272]	2	511.3103	2.247	0.173
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VQQPDCR	2	516.2417	2.111	0.272
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	FGLLDEDGKK	2	771.438	3.21	0.188
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	NNVPCSPK	2	592.7994	2.199	0.257
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	TYNVLDMK	2	632.342	2.078	0.119
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNMGK[272]	2	722.427	2.703	0.232
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VEASISK	2	507.3032	2.326	0.154
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]NVNFQK[272]	2	519.3028	2.35	0.172
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VDFTLSSER	2	599.3152	2.187	0.218
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNMGK[272]	4	361.7171	2.497	0.132
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SHKPLNMGK	3	477.9466	2.093	0.224
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EVQLVAHSPWLK[272]	3	565.6659	3.689	0.225

153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]QGSFQGGFR	2	564.2894	2.462	0.184
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VEASISK[272]	1	1021.6133	2.575	0.098
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNM[147]GK[272]	3	487.2854	1.917	0.235
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	FGLLDEDGK	2	637.343	2.87	0.132
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	TEQWSTLPPETK	2	848.9489	2.692	0.186
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LELSVDGAK	2	606.3534	2.333	0.235
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	NVNFQK	2	515.2957	2.077	0.171
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]NNVPCSPK[272]	2	596.8065	2.336	0.204
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EFHLHLR	2	548.3126	2.549	0.118
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SHKPLNM[147]GK	3	483.2783	1.704	0.242
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]MRPSTDTITVM[147]VENSHGLR	3	768.7234	3.518	0.11
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SHKPLNMGK	2	716.4163	2.134	0.19
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VFALDQK[272]	2	554.8339	1.325	0.127
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EGAIHREELVYELNPLDHR	4	609.3204	2.196	0.158
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]VQQPDCR	2	518.2452	2.174	0.135
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]PVAFSVVPTAAAAVSLK[272]	3	639.3868	2.837	0.351
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GQIVFM[147]NR	2	562.8038	2.855	0.244
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SHK[272]PLNMGK[272]	3	481.9537	2.203	0.161
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]RGHLFLQTDQPIYNPGQR	4	571.8072	3.249	0.088
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]TEQWSTLPPETK[272]	2	852.956	1.61	0.276
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]CSVFYGAPSK[272]	3	464.9013	1.687	0.199
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GLESQTK	2	521.7982	1.768	0.194
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ILTVPGHLDEMQLDIQAR	3	731.7321	3.846	0.231
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]MRPSTDTITVM[147]VENSHGLR	4	576.7944	2.715	0.134
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DHAVDLIQK	3	440.2541	2.889	0.044
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AVGSGATFSH	2	537.2724	3.193	0.45
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EFHLHLR	3	365.8775	2.519	0.041
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FGLLDEDGK[272]	2	641.3501	2.202	0.081
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GSFQDLSPVIHR	2	750.408	4.043	0.353
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DK[272]GQAGLQR	2	630.8668	2.384	0.214
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LFETK[272]	2	463.2835	1.994	0.166
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]PVQGVAYVR	2	566.8334	3.358	0.382
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]YIYGK[272]PVQGVAY	2	823.4633	3.385	0.399
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHYYY	2	783.8709	3.188	0.505
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSH	2	539.2759	2.899	0.417
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]CCQDGVTR	2	559.2226	1.446	0.03
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	IFETK	2	459.2764	1.916	0.139
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	VFALDQK	2	550.8268	1.851	0.074
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	YIYGKPVQGVAY	2	819.4562	3.234	0.414
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	YIYGKPVQGVAYVR	3	631.6964	2.829	0.034
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	FGLLDEDGKKTFFR	4	524.0441	2.773	0.093
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AEM[147]ADQAAAWLTR	2	795.3907	2.504	0.016
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLLQALVR	2	888.0443	1.687	0.195
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AVGSGATFSHYYYM[147]IL	2	970.4727	2.976	0.414
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AVGSGATFSHYYY	2	781.8674	2.367	0.416
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ACYYPR	2	481.7209	2.41	0.292
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ADLEK	2	428.2504	1.732	0.017
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	DFALLSLQVPLK	3	541.9993	2.954	0.027

153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]AIADVTLLSGFHALR	2	864.4999	3.688	0.454
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPF	2	491.2856	1.528	0.243
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AVGSGATFSHYYYM[147]IL	2	968.4691	2.455	0.488
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLLQAL	2	760.4595	3.069	0.309
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ALLHLLLHEGK	2	762.4747	3.934	0.398
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ALLHLLLHEGK[272]	2	766.4818	3.347	0.41
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ASAGLLGAH	2	470.7703	2.721	0.275
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLLHEGK[272]	3	366.5689	2.178	0.215
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]YYHGDHPVANSLR	3	558.2819	2.468	0.251
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GSFQDLSPVIHR	3	500.6077	2.748	0.189
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	AVGSGATFSHYY	2	700.3357	2.46	0.338
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]EVQLVAHSPWLK[272]	2	847.9953	4.232	0.215
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	GPEVQLVAHSPWLK[272]	2	923.0288	3.57	0.337
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DGSYAAWLSR	2	635.3209	2.097	0.306
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LLLHEGK	3	363.8975	2.328	0.183
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	HLVPGAPFLLQ	2	666.3954	2.52	0.362
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DFALLSLQVPLK[272]	3	544.6707	2.364	0.024
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]SVGVQLQDVPR	2	671.384	3.329	0.196
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LSCPK	2	437.2379	1.377	0.033
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ASAGLLGAH	2	468.7667	2.087	0.309
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	HLVPGAPFLLQA	2	701.9139	2.071	0.363
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]IYLAPTLAASR	2	660.3938	2.653	0.274
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EVQLVAHSPWLK[272]	3	564.3302	3.088	0.226
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EVQLVAHSPWLK	2	843.9882	4.102	0.246
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ASAGLLGAHAAAITA	2	719.9104	2.703	0.233
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GLESQTK[272]	2	525.8053	1.587	0.027
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLLQA	2	703.9175	2.43	0.235
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]CYYPR	2	446.2023	1.893	0.223
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DHAVDLIQK[272]GYM[147]R	3	617.3326	2.33	0.111
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ALTLTK[272]	2	467.8124	2.063	0.195
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLLHEGK[272]	2	549.3497	2.055	0.216
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LLLHEGK	2	545.3426	2.273	0.165
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLLFSPSVVHL	2	684.9222	3.02	0.269
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLLFSPSVVH	2	628.3802	2.685	0.21
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	IYLAPTLAASR	2	658.3903	2.453	0.254
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLQALVR	2	478.8223	2.929	0.141
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LTVAAPPSGGPGFLSIERPD	3	709.0518	2.138	0.292
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]PNMIPDGDFNSYVR	2	884.9259	2.47	0.185
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLLQ	2	668.3989	2.165	0.178
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LSGM[147]AIADVTLLSGFHALR	3	710.0576	3.096	0.102
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ALTLTK	2	463.8053	2.316	0.037
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	ALLHLLLHEGK	3	508.6522	3.208	0.027
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]DDFVIPDISEPGTWK[272]	4	502.5133	2.753	0.113
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LLLFSPSVVHLG	2	713.433	2.199	0.322
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LAPTLAASR	2	520.3166	1.803	0.107
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]GVAHNNLM[147]AM[147]	2	617.2955	1.583	0.15
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]ITQVLH	2	427.7645	1.675	0.067
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]LAPTLAASR	2	522.3201	1.986	0.042

153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]HLVPGAPFLL	2	604.3696	1.739	0.11
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	EVQLVAHSPWLK[272]	2	845.9917	3.805	0.254
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	n[145]FNSYVR	2	465.2517	1.541	0.064
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	LTVAAPPSGGPGFLSIER	3	637.0229	3.062	0.261
153	IPI00892604	1	51.4	0.53	0.08	12	7	C4B Complement component C4B	SVK[272]	1	617.4104	0.88	0.378
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	GNVATEISTER	2	658.8439	3.649	0.443
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KGNVATEISTER	2	792.9389	4.231	0.48
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	NSEEFAAAMSR	2	676.8168	3.568	0.463
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	TKNSEEFAAAM[147]SR	2	869.4331	3.669	0.494
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	VLVDHFGYTK	2	729.9089	4.134	0.517
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EQHLFLPFSYK[272]	2	848.9687	4.402	0.386
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]GFEPTLEALFGK[272]	2	798.9475	4.344	0.474
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]NSEEFAAAM[147]SR	2	686.8178	3.694	0.466
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]TK[272]NSEEFAAAM[147]SR	2	873.4402	3.187	0.546
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]VLVDHFGYTK[272]	2	733.916	3.578	0.49
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]YTYNYEAESSSGVPGTADSR	2	1149.5176	6.378	0.589
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	DKDQEVLLQTFLDDASPGDKR	3	937.4943	3.571	0.507
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KGNVATEISTER	3	528.9617	4.645	0.413
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KLTISEQNIQR	3	537.3177	2.091	0.229
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KM[147]GLAFESTK	3	516.6253	3.103	0.375
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KYTYNYEAESSSGVPGTADSR	3	854.7418	4.207	0.361
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	ILGEELGFASLHDLQLLGK	3	778.4458	6.146	0.581
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	NKYGM[147]VAQVTQTLK	3	673.047	4.326	0.415
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	QVFLYPEKDEPTYILNIK	3	877.4913	5.05	0.47
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	QVFLYPEKDEPTYILNIKR	3	929.525	4.681	0.468
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	TSQCTLKEVYGFNPEGK	3	789.7327	5.047	0.45
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	VLVDHFGYTK	3	486.9417	2.617	0.145
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]NK[272]YGM[147]VAQVTQTLK[272]	3	677.0541	4.763	0.466
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QVFLYPEK[272]DEPTYILNIK[272]	3	881.4984	4.386	0.416
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]TK[272]NSEEFAAAM[147]SR	3	582.6292	4.825	0.439
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	DKDQEVLLQTFLDDASPGDKR	4	703.3726	4.134	0.508
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	QVFLYPEKDEPTYILNIKR	4	697.3956	3.337	0.305
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QVFLYPEK[272]DEPTYILNIK[272]R	4	700.4009	4.749	0.499
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	GFEPTLEALFGK	2	794.9404	3.631	0.434
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QGFFPDSVNK[272]	2	713.8821	3.574	0.335
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	EQHLFLPFSYK	3	563.6435	3.969	0.286
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	STSPPKQAEAVLK	3	592.6841	3.349	0.342
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	TKNSEEFAAAM[147]SR	3	579.9578	3.608	0.358
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]FFGEGTK[272]K[272]	3	449.2662	2.616	0.357
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]ILGEELGFASLHDLQLLGK[272]	3	781.1172	3.059	0.318
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QVFLYPEK[272]DEPTYILNIK[272]R	3	933.5321	4.364	0.342
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]STSPPK[272]QAEAVLK[272]	3	596.6912	2.72	0.044
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]VLVDHFGYTK[272]	3	489.6131	1.98	0.076
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	KLTISEQNIQR	2	805.4729	3.461	0.406
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	MGLAFESTK	2	632.342	3	0.397
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	SVSLPSLDPASAK	2	776.4407	3.38	0.189
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]LAAYLM[147]LM[147]R	2	629.3444	2.395	0.372
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]SVSLPSLDPASAK[272]	2	780.4478	3.103	0.09

154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]CVQSTK[272]PSLM[147]IQK[272]	3	653.0264	2.76	0.273
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	HVAEAICK	2	598.8176	3.014	0.372
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	QGFFPDSVNK	2	709.875	3.353	0.37
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	TGISPLALIK	2	646.9187	3.34	0.295
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]NSEEFAAAMSR	2	678.8204	3.733	0.309
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]SPSQADINK[272]	2	624.3454	3.905	0.214
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]TGISPLALIK[272]	2	650.9258	3.718	0.363
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	EQHLFLPFSYK	2	844.9616	2.892	0.404
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	NSEEFAAAM[147]SR	2	684.8143	3.185	0.395
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]HVAEAICK[272]	2	602.8247	2.907	0.329
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]IEGNLIFDPNNYLPK[272]	2	1018.0588	2.363	0.27
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]ANLFNK[272]LVTELR	3	569.345	2.717	0.309
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]LTISEQNIQR	2	673.3815	3.436	0.212
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]MGLAFESTK[272]	2	636.3491	1.653	0.178
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]DDK[272]HEQDMVNGIM[147]LSVEK[272]	3	846.0965	3.136	0.28
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	M[147]GLAFESTK	2	640.3394	2.501	0.282
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	QAEAVLK	2	519.819	2.581	0.242
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]GNVATEISTER	2	660.8474	3.534	0.321
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QAEAVLK[272]	2	523.8261	2.561	0.223
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]FFGEGTK[272]	2	537.2972	2.293	0.228
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	LTISEQNIQR	2	671.3779	3.166	0.137
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EIFNMAR	2	512.772	2.305	0.191
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	EVYGFNPEGK	2	710.367	2.774	0.194
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]NYQLYK[272]	2	558.8183	2.13	0.218
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]LAIPEGK[272]	2	508.3232	2.195	0.235
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]ESQLPTVM[147]DFR	2	741.8726	2.63	0.182
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EVYGFNPEGK[272]	2	714.3741	2.975	0.14
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	FFGEGTK	2	533.2901	2.14	0.191
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	LAIPEGK	2	504.3161	2.626	0.1
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]DLGQCDR	2	498.7216	2.179	0.167
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]TLQELK[272]K[272]	2	646.4192	2.349	0.01
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]SK[272]EVPEAR	3	401.9027	2.821	0.112
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	DLGQCDR	2	496.7181	1.967	0.133
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	NYQLYK	2	554.8112	1.753	0.125
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	IEGNLIFDPNNYLPK	2	1014.0517	2.112	0.102
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]LEDTPK[272]	2	495.7892	2.092	0.064
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]SK[272]EVPEAR	2	602.3504	2.183	0.282
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	LVTELR	2	435.774	2.059	0.15
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	SPSQADINK	2	620.3383	3.167	0.238
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EEEMLENVSLVCPK[272]	2	977.4823	4.208	0.351
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	CVQSTKPSLM[147]IQK	3	649.0193	2.691	0.076
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]TSQCTLK	2	555.792	1.863	0.032
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	LAAYLM[147]LM[147]R	2	627.3409	1.722	0.048
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	DDKHEQDM[147]VNGIM[147]LSVEK	3	847.421	2.89	0.106
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]ALYWVNGQVPDGVSK[272]	3	641.0189	3.123	0.043
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	EEEMLENVSLVCPK	2	973.4752	3.33	0.434
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]ATLYALSH	2	510.2858	2.74	0.27
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EQHLFLPFSYK[272]	3	566.3149	2.044	0.048

154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	ATLYALSH	2	508.2822	3.065	0.218
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	FFGEGTKK	3	445.2591	1.762	0.109
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	SKEVPEAR	2	598.3433	1.844	0.155
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	SKEVPEAR	3	399.2313	2.434	0.047
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]EEEM[147]LENVSLVCPK[272]	2	985.4798	3.986	0.414
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]AYLM[147]LM[147]R	2	537.2838	2.319	0.279
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	EEEM[147]LENVSLVCPK	2	981.4727	2.402	0.241
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	SVSLPSLDPASAK	3	517.9629	2.004	0.003
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]QVFLYPEK[272]DEPTYILNIK[272]	4	661.3756	2.866	0.043
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]SQADINK[272]	2	532.303	1.904	0.19
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	TKNSEEFAAAMSR	3	574.6262	2.263	0.018
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	AYLM[147]LM[147]R	2	535.2803	2.007	0.139
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]NFVASHIAN	2	558.7996	2.175	0.234
154	IPI00894122	1	52.1	0.89	0.01	4	4	APOB APOB protein	n[145]VLVDHFG	2	465.7619	1.4	0.109
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		LGEHNIDVLEGNEQFIN	2	1041.0186	3.842	0.484
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		LGEHNIDVLEGNEQFINAARI	2	1246.6483	5.866	0.412
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		LGEHNIDVLEGNEQFINAARI	3	831.4346	5.699	0.468
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		LSSPATLNSR	2	593.333	2.68	0.163
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPNFNGNTLDNDIMLIRL	3	855.46	4.571	0.415
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]LGEHNIDVLEGNEQFINAAR	3	795.0756	4.17	0.324
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]LSSPATLNSR	2	595.3365	2.256	0.182
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPNFNGNTLDNDIM[147]LIRL	3	860.7916	4.917	0.404
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPNFNGN	2	633.8331	2.738	0.385
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]IITHPNFNGN	2	635.8367	3.182	0.296
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]LGEHNIDVLEGNEQFIN	2	1043.0221	4.559	0.292
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPNFNG	2	576.8117	2.015	0.207
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]LGEHNIDVLEGNEQ	2	855.9244	3.136	0.289
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]IITHPN	2	419.7488	1.648	0.267
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPNFN	2	548.3009	2.141	0.14
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]IITHPNFN	2	550.3045	2.019	0.122
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]IITHPNFNGNTLD	2	800.416	2.367	0.171
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		IITHPN	2	417.7453	1.522	0.113
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		LSSPATLN	1	942.5255	1.542	0.3
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]SSPATLNSR	2	538.7945	2.045	0.016
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]IITHPNFNG	2	578.8152	1.145	0.193
155	Trypa5|PromTA	1	100	7.25	3.16	13	5		n[145]LSSPATLN	1	946.5326	1.639	0.276
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		CCTESLVNR	1	1256.5255	2.625	0.448
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]CCTESLVNR	1	1260.5326	3.271	0.418
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		CCTESLVNR	2	628.7664	2.692	0.374
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KVPQVSTPTLVEVSR	2	960.5675	3.012	0.399
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		VPQVSTPTLVEVSR	2	826.4725	2.141	0.242
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KVPQVSTPTLVEVSR	3	640.7141	2.558	0.274
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KYLYEIAR	3	445.931	2.961	0.374
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]CCTESLVNR	2	630.7699	3.019	0.386
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]LVNELTEFAK[272]	2	726.4211	2.988	0.274
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		LVNELTEFAK	2	722.414	3.313	0.308
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KQTALVELLK	3	521.6713	1.714	0.005
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		YLYEIAR	2	534.2979	2.505	0.117

156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		YLYEIAR	1	1067.5885	2.485	0.127
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]YLYEIAR	1	1071.5956	2.423	0.119
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]DLGEEQFK[272]	2	627.3345	1.947	0.047
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]LVTDLTK[272]	2	539.3416	1.491	0.104
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		LVTDLTK	1	1069.6617	2.119	0.085
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		LVTDLTK	2	535.3345	2.415	0.102
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]LVTDLTK[272]	1	1077.6759	1.923	0.042
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]YLYEIAR	2	536.3014	1.823	0.031
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]FK[272]DLGEEQFK[272]	3	558.3139	3.045	0.179
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		DLGEEQFK	2	623.3274	2.108	0.077
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		AWSVAR	2	415.2376	2.019	0.07
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]CASIQK[272]	2	492.2665	1.47	0.143
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		ADIAK	2	399.2477	1.927	0.203
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		CASIQK	2	488.2594	1.741	0.13
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]CTESLVNR	2	556.2715	1.944	0.168
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		FKDLGEEQFK	3	554.3068	2.251	0.074
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		GACLLPK	2	514.2932	2.195	0.008
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		QVSTPTLVEVSR	2	728.4119	2.45	0.147
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KYLYEIAR	2	668.3928	2.621	0.313
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		DVCK	2	395.7012	1.198	0.182
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]QVSTPTLVEVSR	2	730.4155	2.58	0.171
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		RHPYFYAPELL	2	773.4143	2.366	0.06
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]CVLHEK[272]	2	531.7876	2.428	0.209
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		HPYFYAPELL	2	695.3637	2.509	0.257
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		HPYFYAPEL	2	638.8217	1.934	0.114
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KQTALVELLK	2	782.0033	2.027	0.066
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		CVLHEK	2	527.7805	2.238	0.165
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		FQNALIVR	2	550.8324	3.137	0.218
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KVPQVSTPTLV	2	724.9454	2.074	0.255
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KVPQVSTPTLVEV	2	839.0009	2.257	0.265
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		TESLVNR	2	479.7695	2.317	0.127
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		ETYVPK	2	508.7924	2.07	0.15
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]LLECADDR	2	562.7635	2.137	0.129
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		VSTPTLVEVSR	2	664.3827	2.392	0.141
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]FQNALIVR	2	552.836	2.972	0.193
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]RHPYFYAPEL	2	718.8758	2.125	0.089
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		PLLEK	2	440.2868	1.519	0.055
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		RHPYFYAPEL	2	716.8723	2.058	0.158
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KPLLEK	3	383.2569	2.599	0.063
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]VTDLTK[272]	2	482.7995	2.013	0.049
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]IYEIAR	2	454.7697	2.144	0.122
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		KVPQVSTPTLVEVS	2	882.517	2.651	0.424
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]PLLEK[272]	2	444.2939	1.942	0
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		RPCFSAL	2	490.2462	1.689	0.117
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		n[145]FQNALIVR	1	1104.6647	2.248	0.187
156	gi|113574|sp|P(	1	23.7	1.35	0.32	39	11		ALIVR	2	356.2474	1.693	0.002
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LGEHNIDVLEGNEQFIN	2	1041.0186	3.842	0.484
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LGEHNIDVLEGNEQFINAAK	2	1246.1507	6.144	0.496

157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNGNTLDNDIM[147]LIK	2	1290.1862	4.83	0.566
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LGEHNIDVLEGNEQFINAAK[272]	2	1250.1578	6.109	0.487
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGNTLDNDIMLIK[272]	2	1286.1958	4.412	0.558
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGNTLDNDIM[147]LIK[272]	2	1294.1933	4.503	0.562
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LGEHNIDVLEGNEQFINAAK	3	831.1029	6.369	0.494
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNGNTLDNDIMLIK	3	855.1282	2.71	0.297
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNGNTLDNDIM[147]LIK	3	860.4599	4.422	0.287
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LGEHNIDVLEGNEQFINAAK[272]	3	833.7743	6.943	0.319
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGNTLDNDIMLIK[272]	3	857.7996	6.747	0.56
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGNTLDNDIM[147]LIK[272]	3	863.1313	2.288	0.017
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LGEHNIDVLEGNEQFINAAK	4	623.579	3.098	0.277
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		VATVSLPR	2	491.8059	1.302	0.217
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]VATVSLPR	2	493.8094	2.148	0.174
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		NKPGVYTK	3	442.9346	2.869	0.197
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]NK[272]PGVYTK[272]	3	446.9417	2.284	0.241
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LSSPATLNSR	2	593.333	2.68	0.163
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]NK[272]PGVYTK[272]	2	669.909	2.625	0.204
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		SSGSSYPSLLQCLK	2	898.4577	2.31	0.268
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LSSPATLNSR	2	595.3365	2.256	0.182
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		NKPGVYTK	2	663.8983	2.102	0.174
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNGN	2	633.8331	2.738	0.385
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGN	2	635.8367	3.182	0.296
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LGEHNIDVLEGNEQFIN	2	1043.0221	4.559	0.292
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNGNTLDNDIMLIK	4	641.598	2.179	0.01
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFNG	2	576.8117	2.015	0.207
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LGEHNIDVLEGNEQ	2	855.9244	3.136	0.289
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPN	2	419.7488	1.648	0.267
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		TLDNDIMLIK	2	728.4157	3.628	0.324
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPNFN	2	548.3009	2.141	0.14
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFN	2	550.3045	2.019	0.122
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]TLDNDIM[147]LIK[272]	2	740.4202	2.934	0.251
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNGNTLD	2	800.416	2.367	0.171
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]TVSLPR	2	408.7566	2.007	0.09
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		IITHPN	2	417.7453	1.522	0.113
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		LSSPATLN	1	942.5255	1.542	0.3
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]SSPATLNSR	2	538.7945	2.045	0.016
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]IITHPNFNG	2	578.8152	1.145	0.193
157	gi|136429|sp|P(	1	34.6	0.92	0.08	61	30		n[145]LSSPATLN	1	946.5326	1.639	0.276
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		SEIDNVKK	3	451.6014	3.001	0.334
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		n[145]DVDGAYM[147]TK[272]	2	652.3258	2.767	0.294
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		DVDGAYM[147]TK	2	648.3187	2.857	0.22
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		n[145]YEDEINK[272]R	3	452.245	2.209	0.161
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		AQYEDIAQK	2	673.3592	2.645	0.041
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		n[145]SEIDNVK[272]K[272]	3	455.6085	2.233	0.027
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		IDNVKK	2	568.8612	2.026	0.076
158	gi|71536|pir||KF	1	6.9	1.81	0.67	6	6		IDNVK	2	434.7662	1.944	0.066
159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	LGEHNIEVLEGNEQFINAAK	3	835.7747	4.736	0.39
159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	NKPGVYTK	3	442.9346	2.869	0.197

159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	n[145]NK[272]PGVYTK[272]	3	446.9417	2.284	0.241
159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	n[145]NK[272]PGVYTK[272]	2	669.909	2.625	0.204
159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	n[145]LGEHNIEVLEGNEQFINAAK[272]	3	838.4461	4.24	0.264
159	IPI00011694	0.9999	11.3	0.78	0.12	2	2	PRSS1 Trypsin-1 precursor	NKPGVYTK	2	663.8983	2.102	0.174
160	IPI00168728	0.9999	3.2	0.57	0.02	2	2	IGHM FLJ00385 protein (Fragment)	n[145]TPLGDTTHTCPR	3	496.9067	2.644	0.357
160	IPI00168728	0.9999	3.2	0.57	0.02	2	2	IGHM FLJ00385 protein (Fragment)	n[145]TPLGDTTHTCPR	2	744.8564	2.091	0.184
161	IPI00063523	0.9998	1.2	0.65	0	2	2	FLJ40176 Conserved hypothetical protein	ALHLQEEK	2	624.359	2.594	0.14
161	IPI00063523	0.9998	1.2	0.65	0	2	2	FLJ40176 Conserved hypothetical protein	n[145]ALHLQEEK[272]	2	628.3661	2.578	0.143
161	IPI00063523	0.9998	1.2	0.65	0	2	2	FLJ40176 Conserved hypothetical protein	SK[272]K[272]	2	395.7731	0.913	0.136
161	IPI00063523	0.9998	1.2	0.65	0	2	2	FLJ40176 Conserved hypothetical protein	SVK[272]	1	617.4104	0.88	0.378
162	IPI00171678	0.9998	4.8	5.21	6.97	4	2	DBH Dopamine beta-hydroxylase	n[145]LEVHYHNPLVIEGR	3	607.336	3.52	0.257
162	IPI00171678	0.9998	4.8	5.21	6.97	4	2	DBH Dopamine beta-hydroxylase	n[145]TPEGLTLLFK[272]	2	703.9285	2.298	0.122
162	IPI00171678	0.9998	4.8	5.21	6.97	4	2	DBH Dopamine beta-hydroxylase	LEEPT	2	364.6949	1.495	0.082
163	IPI00439446	0.9998	5.5	2.61	1.87	3	2	MAN1A1 MAN1A1 protein	n[145]HEFEEAK[272]	2	589.3082	2.111	0.212
163	IPI00439446	0.9998	5.5	2.61	1.87	3	2	MAN1A1 MAN1A1 protein	n[145]GGLLEHK[272]	2	521.3184	2.435	0.063
163	IPI00439446	0.9998	5.5	2.61	1.87	3	2	MAN1A1 MAN1A1 protein	FDGGVEAIATRQNEK	3	639.0059	1.933	0.056
163	IPI00439446	0.9998	5.5	2.61	1.87	3	2	MAN1A1 MAN1A1 protein	EM[147]K	1	703.3808	1.121	0.072
163	IPI00439446	0.9998	5.5	2.61	1.87	3	2	MAN1A1 MAN1A1 protein	YILR	2	352.7263	1.622	0.012
164	IPI00412492	0.9996	0.6	0.73	0.03	7	2	PLXND1 Isoform 1 of Plexin-D1 precursor	n[145]EIPEYRK[272]	3	408.2393	2.148	0.035
164	IPI00412492	0.9996	0.6	0.73	0.03	7	2	PLXND1 Isoform 1 of Plexin-D1 precursor	EIPEYRK	3	405.5679	2.036	0.061
164	IPI00412492	0.9996	0.6	0.73	0.03	7	2	PLXND1 Isoform 1 of Plexin-D1 precursor	YILR	2	352.7263	1.622	0.012
165	IPI00328765	0.9995	7.4	0.81	0.02	3	3	PIBF1 Isoform 1 of Progesterone-induced-blocking factor 1	VLQLEK	2	505.3239	2.639	0.029
165	IPI00328765	0.9995	7.4	0.81	0.02	3	3	PIBF1 Isoform 1 of Progesterone-induced-blocking factor 1	n[145]VLQLEK[272]	2	509.331	2.26	0.039
165	IPI00328765	0.9995	7.4	0.81	0.02	3	3	PIBF1 Isoform 1 of Progesterone-induced-blocking factor 1	n[145]QTQQPYR	2	532.7839	2.248	0.137
165	IPI00328765	0.9995	7.4	0.81	0.02	3	3	PIBF1 Isoform 1 of Progesterone-induced-blocking factor 1	SK[272]K[272]	2	395.7731	0.913	0.136
166	IPI00431645	0.9995	3.2	0.65	0	3	2	HP HP protein	n[145]ILGGHLDAK[272]	3	404.5832	2.622	0.139
166	IPI00431645	0.9995	3.2	0.65	0	3	2	HP HP protein	ILGGHLDAK	3	401.9118	2.614	0.074
167	gi|547749|sp|P1	0.9993	5.2	1.19	0.06	2	2	KRT10 Keratin	n[145]ALEESNYELEGK[272]	2	835.4298	5.194	0.465
167	gi|547749|sp|P1	0.9993	5.2	1.19	0.06	2	2	KRT10 Keratin	n[145]SLLEGEGSSGGGGR	2	703.8533	2.937	0.026
167	gi|547749|sp|P1	0.9993	5.2	1.19	0.06	2	2	KRT10 Keratin	EM[147]K	1	703.3808	1.121	0.072
168	IPI00301143	0.9993	5.2	0.8	0	2	2	PI16 Isoform 1 of Peptidase inhibitor 16 precursor	n[145]AQVSPTASDM[147]LHM[147]R	3	573.9473	3.873	0.383
168	IPI00301143	0.9993	5.2	0.8	0	2	2	PI16 Isoform 1 of Peptidase inhibitor 16 precursor	AQVSPTASDM[147]LHM[147]R	3	572.6116	2.36	0.108
169	IPI00004373	0.9989	10.1	1.03	0.04	2	2	MBL2 Mannose-binding protein C precursor	n[145]TEGQFVDLTGNR	2	740.8793	2.702	0.274
169	IPI00004373	0.9989	10.1	1.03	0.04	2	2	MBL2 Mannose-binding protein C precursor	n[145]SPDGDSSLAASER	2	718.3427	3.606	0.29
170	IPI00030205	0.9986	8.3	0.78	0	2	2	IGKV3-20 Ig kappa chain V-III region HAH precursor	n[145]LLIYGASSR	2	562.3332	2.762	0.131
170	IPI00030205	0.9986	8.3	0.78	0	2	2	IGKV3-20 Ig kappa chain V-III region HAH precursor	LLIYGASSR	2	560.3297	2.16	0.09
171	IPI00064667	0.9985	5.5	1.14	0.09	3	3	CNDP1 Beta-Ala-His dipeptidase precursor	KPAITYGTR	2	643.8826	1.796	0.206
171	IPI00064667	0.9985	5.5	1.14	0.09	3	3	CNDP1 Beta-Ala-His dipeptidase precursor	n[145]HLEDVFSK[272]	2	631.8528	2.197	0.141
171	IPI00064667	0.9985	5.5	1.14	0.09	3	3	CNDP1 Beta-Ala-His dipeptidase precursor	n[145]AITYGTR	2	463.2648	2.27	0.083
172	IPI00719373	0.998	11.7	0.8	0	1	2	IGL@ IGL@ protein	SYSCQVTHEGSTVEK	3	660.9767	4.441	0.439
172	IPI00719373	0.998	11.7	0.8	0	1	2	IGL@ IGL@ protein	n[145]AGVETTK[272]PSK[272]	3	483.9594	2.503	0.283
173	IPI00167638	0.9976	3.3	0.77	0.08	5	2	GTPBP10 Isoform 1 of GTP-binding protein 10	n[145]ELELYK[272]	2	541.8205	2.222	0.081
173	IPI00167638	0.9976	3.3	0.77	0.08	5	2	GTPBP10 Isoform 1 of GTP-binding protein 10	ELELYK	2	537.8134	1.954	0.098
174	IPI00413626	0.9969	3.4	0.93	0.01	4	2	SUNC1 Isoform 1 of Sad1/unc-84 domain-containing protein 1	n[145]EQLELLK[272]	2	580.8601	1.949	0.036
174	IPI00413626	0.9969	3.4	0.93	0.01	4	2	SUNC1 Isoform 1 of Sad1/unc-84 domain-containing protein 1	EQLELLK	2	576.853	2.393	0.051
175	IPI00022389	0.9967	16.5	0.44	0.11	2	2	CRP Isoform 1 of C-reactive protein precursor	KAFVFPK	3	419.6008	2.912	0.172
175	IPI00022389	0.9967	16.5	0.44	0.11	2	2	CRP Isoform 1 of C-reactive protein precursor	n[145]APLTK[272]PLK[272]	3	433.9623	1.693	0.029
176	IPI00293011	0.9959	2.1	0.79	0	2	2	TCP11L1 cDNA FLJ11386 fis	ELEEVAIK	2	605.8558	2.906	0.029

176	IPI00293011	0.9959	2.1	0.79	0	2	2	TCP11L1 cDNA FLJ11386 fis	n[145]ELEEVAIK[272]	2	609.8629	2.489	0.067
177	IPI00218795	0.9958	4.9	0.74	0.04	2	2	SELL L-selectin precursor	AEIEYLEK	2	637.8532	2.578	0.23
177	IPI00218795	0.9958	4.9	0.74	0.04	2	2	SELL L-selectin precursor	NKEDCVEIYIK	3	607.3197	2.921	0.127
178	IPI00215977	0.9957	2.7	0.74	0	2	2	IGF2 Isoform 2 of Insulin-like growth factor II precursor	GFYFR	2	415.2214	1.957	0.155
178	IPI00215977	0.9957	2.7	0.74	0	2	2	IGF2 Isoform 2 of Insulin-like growth factor II precursor	n[145]GFYFR	2	417.225	1.91	0.02
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	KVSPHSGVVALTKPVPEPR	4	605.3641	2.189	0.036
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	GYHLNEEGTR	3	439.2178	2.718	0.267
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	GYHLNEEGTR	2	658.3231	3.381	0.441
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	n[145]TGYYFDGISR	2	661.8285	3.472	0.317
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	n[145]GYHLNEEGTR	2	660.3267	3.053	0.323
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	n[145]GYHLNEEGTR	3	440.5535	2.216	0.23
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	TGYYFDGISR	2	659.825	2.693	0.292
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	NVPNCGR	2	473.2233	1.645	0.029
179	IPI00296537	0.9899	7.9	1.03	0	2	1	FBLN1 Isoform C of Fibulin-1 precursor	SCPSSGYR	2	521.7259	2.538	0.179
180	IPI00444845	0.9854	1.6	0.44	0.09	2	2	ZBTB47 Zinc finger and BTB domain-containing protein 47	n[145]VK[272]LEEK[272]	3	393.2554	2.257	0.17
180	IPI00444845	0.9854	1.6	0.44	0.09	2	2	ZBTB47 Zinc finger and BTB domain-containing protein 47	n[145]VK[272]LEEK[272]	2	589.3795	2.115	0.146
181	IPI00025829	0.9824	2.7	0.63	0	2	2	CYP2B6 Cytochrome P450 2B6	n[145]FHYQDQEFLK[272]	2	821.9271	2.738	0.044
181	IPI00025829	0.9824	2.7	0.63	0	2	2	CYP2B6 Cytochrome P450 2B6	FHYQDQEFLK	2	817.92	2.788	0.012
182	IPI00296099	0.9819	1.6	0.2	0	1	1	THBS1 Thrombospondin-1 precursor	n[145]FVFGTTPEDILR	2	769.9204	3.407	0.267
182	IPI00296099	0.9819	1.6	0.2	0	1	1	THBS1 Thrombospondin-1 precursor	n[145]YECR	2	380.6656	1.459	0.182
182	IPI00296099	0.9819	1.6	0.2	0	1	1	THBS1 Thrombospondin-1 precursor	SVK[272]	1	617.4104	0.88	0.378
183	IPI00000856	0.9804	0.7	0.14	0.08	3	2	FERMT2 Isoform 1 of Fermitin family homolog 2	n[145]YDAIR	1	781.4325	1.827	0.181
183	IPI00000856	0.9804	0.7	0.14	0.08	3	2	FERMT2 Isoform 1 of Fermitin family homolog 2	n[145]DVTIR	2	374.2277	1.629	0.02
184	IPI00304189	0.9779	2.5	1	0.1	3	2	OPTN Isoform 1 of Optineurin	LKEEIGK	2	618.8874	3.068	0.174
184	IPI00304189	0.9779	2.5	1	0.1	3	2	OPTN Isoform 1 of Optineurin	n[145]LK[272]EELGK[272]	2	624.8981	2.905	0.186
184	IPI00304189	0.9779	2.5	1	0.1	3	2	OPTN Isoform 1 of Optineurin	n[145]LK[272]EELGK[272]	3	416.9345	2.515	0.138
184	IPI00304189	0.9779	2.5	1	0.1	3	2	OPTN Isoform 1 of Optineurin	SVK[272]	1	617.4104	0.88	0.378
184	IPI00304189	0.9779	2.5	1	0.1	3	2	OPTN Isoform 1 of Optineurin	EM[147]K	1	703.3808	1.121	0.072
185	IPI00010390	0.9776	1.4	29.1	5.74	5	2	SAMD4A SAMD4A protein	VEYM[147]KLLPK	2	778.9653	2.387	0.035
185	IPI00010390	0.9776	1.4	29.1	5.74	5	2	SAMD4A SAMD4A protein	VEYM[147]K[272]LLPK[272]	2	782.9724	2.441	0.065
185	IPI00010390	0.9776	1.4	29.1	5.74	5	2	SAMD4A SAMD4A protein	VEYM[147]KL	1	1078.5966	1.723	0.131
186	IPI00014826	0.976	2.7	0.76	0	3	2	BNC2 Isoform 2 of Zinc finger protein basonuclin-2	EGQAVAVPSSK	2	676.8803	2.398	0.012
186	IPI00014826	0.976	2.7	0.76	0	3	2	BNC2 Isoform 2 of Zinc finger protein basonuclin-2	n[145]EGQAVAVPSSK[272]	2	680.8874	2.383	0.02
187	IPI00216704	0.9748	1	0.75	0.01	3	2	SPTB Isoform 2 of Spectrin beta chain	n[145]DSPDVTHR	2	535.771	1.665	0.009
187	IPI00216704	0.9748	1	0.75	0.01	3	2	SPTB Isoform 2 of Spectrin beta chain	GAEILGLIDEK[272]	3	481.2843	2.752	0.22
187	IPI00216704	0.9748	1	0.75	0.01	3	2	SPTB Isoform 2 of Spectrin beta chain	EM[147]K	1	703.3808	1.121	0.072
188	IPI00154742	0.9738	10.7	0.7	0	1	1	IGL@ IGL@ protein	SYSCQVTHEGSTVEK	3	660.9767	4.441	0.439
188	IPI00154742	0.9738	10.7	0.7	0	1	1	IGL@ IGL@ protein	n[145]AGVETTTPSK[272]	2	639.8608	3.005	0.303
189	gi|1082558|pir||	0.9719	2.6	0.67	0	1	1	KRT9 Keratin	n[145]FSSSSGYGGGSSR	2	690.319	3.905	0.485
189	gi|1082558|pir||	0.9719	2.6	0.67	0	1	1	KRT9 Keratin	SK[272]K[272]	2	395.7731	0.913	0.136
190	IPI00032293	0.9719	11.6	0.64	0	3	1	CST3 Cystatin-C precursor	n[145]TQPNLDNCPFHDQPHLK[272]	4	585.2882	3.915	0.414
191	IPI00418437	0.9719	2.8	12.8	0	1	1	KLHL17 Kelch-like protein 17	SCSPYFHAM[147]FTNEM[147]SESR	5	469.1927	2.676	0.33
192	IPI00005292	0.9692	1.8	N_A	N_A	N_A	1	SPOCK1 Testican-1 precursor	n[145]CNEEGYYK	3	445.8658	2.811	0.33
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	AILK	2	362.7577	1.441	1
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	SLAVPIVAK[272]	3	394.5961	1.278	0.236
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	SVK[272]	1	617.4104	0.88	0.378
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	SK[272]K[272]	2	395.7731	0.913	0.136
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	VVEVK	2	427.279	2.118	0.091

193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]VSWYSK	2	527.2961	1.237	0.091
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]LEILSISK[272]	2	595.8836	2.723	0.111
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]FAIVR	2	375.2432	2.189	0.062
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]YVFR	2	364.7142	1.594	0.143
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]DSMRGDSGR	2	562.7654	1.804	0.097
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]EALLK[272]	2	431.2861	1.702	0.145
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]ELLVK[272]	2	445.3017	2.078	0.148
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	LEILSISK	2	591.8765	2.187	0.107
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]IVEYR	2	412.2434	1.932	0.032
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	EAIIK	2	427.279	1.917	0.224
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	EM[147]K	1	703.3808	1.121	0.072
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	ELLVK	2	441.2946	1.881	0.208
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	IVEYR	2	410.2398	1.544	0.072
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	YILR	2	352.7263	1.622	0.012
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]VPVPVPK[272]	2	512.3439	1.021	0.025
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	INGSEPLQ	2	499.2693	1.559	0.112
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]TYDGGSK	2	506.265	1.936	0.094
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]EVVLK[272]	2	438.2939	1.642	0.018
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	LELK	2	391.7604	1.578	1
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]YLVK[272]	2	405.77	1.349	0.275
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	n[145]YLVK	2	403.7665	1.33	0.146
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	GDLR	1	600.3464	1.375	0.017
193	IPI00023283	0.967	0.8	0.68	0.4	11	5	TTN Isoform 2 of Titin	SVAK	2	342.7238	0.841	0.121
194	IPI00099131	0.9662	2.1	0.88	0.04	2	2	ZC3H7B Isoform 1 of Zinc finger CCCH domain-containing protein 7	n[145]ALNELGR	2	458.7703	1.884	0.163
194	IPI00099131	0.9662	2.1	0.88	0.04	2	2	ZC3H7B Isoform 1 of Zinc finger CCCH domain-containing protein 7	n[145]ADIEK[272]	2	432.2575	2.035	0.149
194	IPI00099131	0.9662	2.1	0.88	0.04	2	2	ZC3H7B Isoform 1 of Zinc finger CCCH domain-containing protein 7	SK[272]K[272]	2	395.7731	0.913	0.136
195	IPI00167547	0.9638	1.9	0	0	3	1	BRWD3 Isoform 1 of Bromodomain and WD repeat-containing prote	n[145]AAAPTQIEAELYYLIAR	4	510.0339	2.868	0.054
195	IPI00167547	0.9638	1.9	0	0	3	1	BRWD3 Isoform 1 of Bromodomain and WD repeat-containing prote	n[145]TAVNNFLLK[272]	2	654.3999	1.98	0.033
196	IPI00646328	0.9589	1.1	1.04	0.06	3	2	FRMPD1 FERM and PDZ domain-containing protein 1	DIILTVK	2	541.3527	1.97	0.171
196	IPI00646328	0.9589	1.1	1.04	0.06	3	2	FRMPD1 FERM and PDZ domain-containing protein 1	n[145]DIILTVK[272]	2	545.3598	1.479	0.064
196	IPI00646328	0.9589	1.1	1.04	0.06	3	2	FRMPD1 FERM and PDZ domain-containing protein 1	n[145]YSISR	2	385.2199	1.799	0.061
196	IPI00646328	0.9589	1.1	1.04	0.06	3	2	FRMPD1 FERM and PDZ domain-containing protein 1	SVK[272]	1	617.4104	0.88	0.378
196	IPI00646328	0.9589	1.1	1.04	0.06	3	2	FRMPD1 FERM and PDZ domain-containing protein 1	EM[147]K	1	703.3808	1.121	0.072
197	IPI00384280	0.9558	1.8	0.59	0.02	2	1	PCYOX1 Prenylcysteine oxidase 1 precursor	n[145]CPSIILHDR	3	415.2184	3.436	0.196
198	IPI00022331	0.9531	3.2	0.88	0	1	1	LCAT Phosphatidylcholine-sterol acyltransferase precursor	n[145]SSGLVSNAPGVQIR	2	764.9318	4.02	0.414
199	IPI00168611	0.9453	4.3	0.77	0	1	1	SLC17A8 Isoform 1 of Vesicular glutamate transporter 3	n[145]IMNCGGFGM[147]EATLLLVVGFSHTK[272]	4	694.6041	2.885	0.288
200	IPI00000111	0.945	1.1	0.64	0	2	2	CHST9 GalNAc-4-sulfotransferase 2	n[145]AIIK[272]K[272]	2	502.8633	1.788	0.11
200	IPI00000111	0.945	1.1	0.64	0	2	2	CHST9 GalNAc-4-sulfotransferase 2	AIIKK	2	496.8526	1.656	0.26
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	n[145]DLTQEM[147]EVHAEK	3	577.2898	2.202	0.098
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	n[145]LPDK[272]K[272]	2	516.8426	1.673	0.019
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	SK[272]K[272]	2	395.7731	0.913	0.136
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	AILK	2	362.7577	1.441	1
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	SVK[272]	1	617.4104	0.88	0.378
201	IPI00009329	0.9443	1.1	0.7	0	1	2	UTRN Utrophin	EM[147]K	1	703.3808	1.121	0.072
202	IPI00328317	0.9427	10	0.25	0	1	1	CPLX4 Complexin-4 precursor	n[145]EEGGASDPAAAQGM[147]TR	3	569.929	2.686	0.281
203	IPI00294926	0.9376	9.1	N_A	N_A	N_A	1	CBX6 Chromobox protein homolog 6	EFCNPEDFEKVAAGVAGAAGGGGSIGASK[272]	5	634.1142	2.702	0.297
203	IPI00294926	0.9376	9.1	N_A	N_A	N_A	1	CBX6 Chromobox protein homolog 6	SK[272]K[272]	2	395.7731	0.913	0.136
203	IPI00294926	0.9376	9.1	N_A	N_A	N_A	1	CBX6 Chromobox protein homolog 6	n[145]YLVK[272]	2	405.77	1.349	0.275

203	IPI00294926	0.9376	9.1	N_A	N_A	N_A	1	CBX6 Chromobox protein homolog 6	n[145]YLVK	2	403.7665	1.33	0.146
204	IPI00000276	0.9305	4.4	0.75	0	2	2	CYorf15B Uncharacterized protein CYorf15B	n[145]VFQIK[272]	2	461.8019	1.87	0.006
204	IPI00000276	0.9305	4.4	0.75	0	2	2	CYorf15B Uncharacterized protein CYorf15B	VFQIK	2	457.7948	1.499	0.056
204	IPI00000276	0.9305	4.4	0.75	0	2	2	CYorf15B Uncharacterized protein CYorf15B	SVK[272]	1	617.4104	0.88	0.378
205	IPI00292807	0.9289	3.3	0.85	0	1	1	PHF8 Putative uncharacterized protein	M[147]DTYSHQALK	2	745.3771	3.476	0.325
206	IPI00031410	0.9275	0.6	N_A	N_A	N_A	1	FRAP1 FKBP12-rapamycin complex-associated protein	STLVESRCCRDLM[147]EEK[272]	3	764.3529	4.485	0.278
207	IPI00013179	0.925	10	0.81	0	1	1	PTGDS Prostaglandin-H2 D-isomerase precursor	n[145]TM[147]LLQPAGSLGSYSYR	2	952.4889	3.484	0.325
208	IPI00006675	0.9243	1.6	1.5	0.41	3	3	ABCC4 Multidrug resistance-associated protein 4	n[145]AEAAALTETAK	2	680.3837	1.931	0.029
208	IPI00006675	0.9243	1.6	1.5	0.41	3	3	ABCC4 Multidrug resistance-associated protein 4	AEAAALTETAK	2	678.3801	1.943	0.006
208	IPI00006675	0.9243	1.6	1.5	0.41	3	3	ABCC4 Multidrug resistance-associated protein 4	LSAVDAEVSR	2	593.825	2.041	0.101
209	IPI00514002	0.9213	1.1	0.9	0	2	2	STARD9 Isoform 1 of StAR-related lipid transfer protein 9	NNGQFVK	2	543.8064	2.185	0.061
209	IPI00514002	0.9213	1.1	0.9	0	2	2	STARD9 Isoform 1 of StAR-related lipid transfer protein 9	n[145]NNGQFVK[272]	2	547.8135	1.803	0.076
209	IPI00514002	0.9213	1.1	0.9	0	2	2	STARD9 Isoform 1 of StAR-related lipid transfer protein 9	SK[272]K[272]	2	395.7731	0.913	0.136
210	IPI00026216	0.916	1.8	1.62	0.31	3	2	NPEPPS Puromycin-sensitive aminopeptidase	AGIISTVEVLK	2	705.44	2.989	0.053
210	IPI00026216	0.916	1.8	1.62	0.31	3	2	NPEPPS Puromycin-sensitive aminopeptidase	n[145]AGIISTVEVLK[272]	2	709.4471	2.311	0.022
210	IPI00026216	0.916	1.8	1.62	0.31	3	2	NPEPPS Puromycin-sensitive aminopeptidase	n[145]GHLDAL	2	385.2199	1.797	0.033
211	IPI00218414	0.9152	3.8	0.84	0	1	1	CA2 Carbonic anhydrase 2	n[145]VGSAK[272]PGLQK[272]	3	472.9682	2.483	0.245
212	IPI00384419	0.9098	1.5	0.71	0.02	4	2	TDRD9 tudor domain containing 9	n[145]SSEVEYINK	2	676.8626	1.681	0.095
212	IPI00384419	0.9098	1.5	0.71	0.02	4	2	TDRD9 tudor domain containing 9	n[145]NVDFQK[272]	2	519.7948	2.392	0
212	IPI00384419	0.9098	1.5	0.71	0.02	4	2	TDRD9 tudor domain containing 9	NVDFQK	2	515.7877	2.103	0.012
212	IPI00384419	0.9098	1.5	0.71	0.02	4	2	TDRD9 tudor domain containing 9	SEILK	2	435.2764	1.848	0.099
213	IPI00008603	0.9079	2.7	0.74	0	2	1	ACTA2 Actin	AGFAGDDAPR	2	558.7753	2.499	0.405
214	IPI00178743	0.9074	1.4	0.57	0	1	2	ALMS1 ALMS1	n[145]QIQVNISDFEGHSNPEGTPVFADRLPEK[272]MK	5	763.1994	3.083	0.285
214	IPI00178743	0.9074	1.4	0.57	0	1	2	ALMS1 ALMS1	SPLQEAESK	2	634.8459	2.084	0.006
214	IPI00178743	0.9074	1.4	0.57	0	1	2	ALMS1 ALMS1	SVK[272]	1	617.4104	0.88	0.378
214	IPI00178743	0.9074	1.4	0.57	0	1	2	ALMS1 ALMS1	SK[272]K[272]	2	395.7731	0.913	0.136
214	IPI00178743	0.9074	1.4	0.57	0	1	2	ALMS1 ALMS1	EM[147]K	1	703.3808	1.121	0.072
215	IPI00019447	0.9039	1.3	0.93	0.05	2	2	FANCI Isoform 1 of Fanconi anemia group I protein	n[145]SLNYTGEK[272]K[272]	2	736.4277	2.47	0.029
215	IPI00019447	0.9039	1.3	0.93	0.05	2	2	FANCI Isoform 1 of Fanconi anemia group I protein	n[145]SLNYTGEK[272]	2	600.3292	1.852	0.047
215	IPI00019447	0.9039	1.3	0.93	0.05	2	2	FANCI Isoform 1 of Fanconi anemia group I protein	LLLLK	2	440.3232	1.8	1
215	IPI00019447	0.9039	1.3	0.93	0.05	2	2	FANCI Isoform 1 of Fanconi anemia group I protein	SK[272]K[272]	2	395.7731	0.913	0.136

[a]accession number, protein accession number under human International Protein Index database; [b]SD, standard deviation; [c]XNP, the number of a quantified peptide by EXPRESS; [d]USP, the number of a unique stripped peptide;