

Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1	53	170	1790	170	4563	515.3	7.05	36856
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	65	79	1429	79	1663	187	6.4	30950
P0C0L5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	62	76	1668	3	1744	192.6	7.27	39091
P0C0L4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	61	75	1642	2	1744	192.7	7.08	36936
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	47	65	837	65	2386	262.5	5.71	18992
Q9HC84	Mucin-5B OS=Homo sapiens GN=MUC5B PE=1 SV=3	19	63	803	63	5762	596	6.64	18452
Q8WX17	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	13	57	699	57	22152	2351.2	6	11622
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1	47	47	590	41	1474	163.2	6.46	11459
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	52	38	637	38	1065	122.1	5.72	15016
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1	69	30	969	30	491	55.9	8.27	26298
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1	62	31	5272	12	452	49.3	6.77	133954
P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	43	47	457	47	1676	188.2	6.52	7831
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	59	32	587	32	609	69.3	6.28	9968
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	43	29	435	29	946	106.4	6.86	8084
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	36	23	440	23	911	101.3	6.79	10912
P04220	Ig mu heavy chain disease protein OS=Homo sapiens PE=1	57	20	3860	1	391	43	5.24	100484
P15144	Aminopeptidase N OS=Homo sapiens GN=ANPEP PE=1	41	30	278	30	967	109.5	5.48	5395
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	34	18	698	18	500	55.1	6.55	14656
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	57	24	461	24	597	67	7.3	8533
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1	59	22	664	22	453	51.5	5.62	13179
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1	35	31	248	31	1231	139	6.61	3682
P12111	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3	18	36	184	36	3177	343.5	6.68	3868
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1	14	28	154	28	2813	309.1	5.48	3771
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	18	19	385	19	1404	151	9.5	7080
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1	50	12	238	7	375	41.7	5.48	4635
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1	53	18	471	18	267	30.8	5.76	11559
P08519	Apolipoprotein(a) OS=Homo sapiens GN=LPA PE=1 SV=1	33	23	216	23	4548	501	5.88	4161
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	43	23	224	15	586	69.4	6.27	3564
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1	71	14	388	7	353	37.6	6.51	8742
P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1	26	25	193	19	1482	163.8	6.38	3290

P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2	11	26	137	26	4391	468.5	6.51	3193
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	54	16	370	16	347	38.1	5.47	7394
P00736	Complement C1r subcomponent OS=Homo sapiens	39	16	152	16	705	80.1	6.21	3190
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1	94	10	1157	10	106	11.6	5.87	33064
Q4LDE5	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 OS=Homo sapiens GN=SVEP1	12	26	133	26	3571	389.9	5.5	2209
P02671	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1	26	18	369	18	866	94.9	6.01	5129
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP	37	13	163	13	493	54.7	7.01	3869
P07355	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	57	17	167	17	339	38.6	7.75	4629
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	31	19	170	19	930	103.3	6.98	3265
P02748	Complement component C9 OS=Homo sapiens GN=C9	43	19	205	19	559	63.1	5.59	4095
P02649	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	41	12	242	12	317	36.1	5.73	5744
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	35	11	218	11	478	54.3	5.8	4524
P27169	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	48	10	158	10	355	39.7	5.22	3022
P00734	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	44	19	128	19	622	70	5.9	2359
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1	46	10	260	6	330	36.1	8.19	5475
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	34	11	161	11	481	53.4	6.7	3987
P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2	59	11	278	4	340	36.5	6.1	6126
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1	52	18	144	18	418	46.7	5.59	2967
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1	6	15	170	15	3396	372.6	4.51	3191
Q7Z7G0	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1	18	13	144	13	1075	118.6	9.44	3399
P10909	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	34	14	201	14	449	52.5	6.27	4421
P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	49	14	166	7	406	45.2	6.58	2873
Q99102	Mucin-4 OS=Homo sapiens GN=MUC4 PE=1 SV=4	13	15	99	15	2169	231.4	6.27	2122
Q06033	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITIH3 PE=1 SV=2	23	13	96	13	890	99.8	5.74	2093
P09871	Complement C1s subcomponent OS=Homo sapiens	31	13	103	13	688	76.6	4.96	1987
P05546	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1	37	13	115	13	499	57	6.9	1924
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3	36	18	144	10	577	67.8	6.4	1905
P01597	Ig kappa chain V-I region DEE OS=Homo sapiens PE=1	27	5	196	4	108	11.7	9.36	5531
P04083	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	50	13	90	13	346	38.7	7.02	2823

P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	29	13	95	13	676	75.1	5.67	1847
O43490	Prominin-1 OS=Homo sapiens GN=PROM1 PE=1 SV=1	24	13	77	13	865	97.1	7.27	1942
Q9Y6R7	IgGfc-binding protein OS=Homo sapiens GN=FCGBP PE=1	9	17	67	17	5405	571.6	5.34	824
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2	85	7	583	2	106	11.3	7.24	14263
P14618	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1	38	14	84	14	531	57.9	7.84	1900
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	41	7	529	2	214	23	8.84	12337
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR	53	12	143	5	348	39	7.09	2272
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2	41	10	163	4	326	35.9	7.59	2548
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	49	11	91	11	320	35.9	5.05	2433
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	32	9	67	9	434	47.1	7.39	1734
P06396	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	26	12	70	12	782	85.6	6.28	1323
P12109	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1	18	10	57	10	1028	108.5	5.43	1324
Q08380	Galectin-3-binding protein OS=Homo sapiens	30	11	68	11	585	65.3	5.27	1677
P21589	5'-nucleotidase OS=Homo sapiens GN=NT5E PE=1 SV=1	34	12	67	12	574	63.3	7.03	1378
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1	11	12	61	12	1786	197.9	4.94	1265
P46940	Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1	14	14	52	14	1657	189.1	6.48	776
P04899	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3	39	9	78	7	355	40.4	5.54	2032
P00747	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	29	14	56	14	810	90.5	7.24	568
P27105	Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=1 SV=3	47	8	60	8	288	31.7	7.88	1471
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1	11	12	59	12	1609	177.5	5.12	1514
P10643	Complement component C7 OS=Homo sapiens GN=C7	22	11	57	11	843	93.5	6.48	1158
P01042	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	18	10	78	10	644	71.9	6.81	1491
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	25	11	51	11	703	77.2	5.22	874
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	37	7	53	7	335	36	8.46	1570
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens	31	10	67	10	423	47.6	5.52	1522
P05187	Alkaline phosphatase, placental type OS=Homo sapiens GN=ALPP PE=1 SV=2	38	14	58	12	535	57.9	6.29	1076
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1	37	3	100	1	114	12.6	7.93	3886
P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1	67	8	196	8	159	18.1	5.24	5147

P35858	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	21	9	65	9	605	66	6.79	1679
P06727	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1	30	10	57	10	396	45.4	5.38	1287
P00558	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1	25	8	59	8	417	44.6	8.1	1281
P08571	Monocyte differentiation antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	29	8	37	8	375	40.1	6.23	790
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1	36	6	150	5	115	12.4	9.25	2910
P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1	31	3	146	2	108	11.8	8.91	6178
P35579	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4	9	12	33	12	1960	226.4	5.6	523
P27487	Dipeptidyl peptidase 4 OS=Homo sapiens GN=DPP4 PE=1	15	9	36	9	766	88.2	6.04	870
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1	31	5	75	4	108	11.6	9.25	2148
P04206	Ig kappa chain V-III region GOL OS=Homo sapiens PE=1	49	3	107	1	109	11.8	9.25	5492
P13671	Complement component C6 OS=Homo sapiens GN=C6	11	8	50	8	934	104.7	6.76	1006
Q9BXX0	EMILIN-2 OS=Homo sapiens GN=EMILIN2 PE=1 SV=3	15	9	41	9	1053	115.6	6.46	928
P68032	Actin, alpha cardiac muscle 1 OS=Homo sapiens GN=ACTC1	19	6	109	1	377	42	5.39	1193
Q96PD5	N-acetylmuramoyl-L-alanine amidase OS=Homo sapiens GN=PGLYRP2 PE=1 SV=1	30	8	43	8	576	62.2	7.55	1080
P01606	Ig kappa chain V-I region OU OS=Homo sapiens PE=1 SV=1	17	2	124	1	108	11.8	9.91	5489
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1	60	4	173	1	109	11.8	8.48	6067
P0DJ18	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1	41	4	67	4	122	13.5	6.79	1853
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3	27	8	147	3	377	41.3	7.9	2860
P01609	Ig kappa chain V-I region Scw OS=Homo sapiens PE=1	28	2	124	1	108	11.8	6	5528
P62879	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Homo sapiens GN=GNB2 PE=1 SV=3	34	8	55	4	340	37.3	6	888
P01596	Ig kappa chain V-I region CAR OS=Homo sapiens PE=1	34	2	80	1	107	11.7	9.41	2651
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	87	1	108	11.8	8.44	2835
Q8NFI5	Retinoic acid-induced protein 3 OS=Homo sapiens GN=GPRC5A PE=1 SV=2	21	4	51	4	357	40.2	8.15	1198
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	25	10	32	8	646	70.9	5.52	692
P01594	Ig kappa chain V-I region AU OS=Homo sapiens PE=1 SV=1	39	2	122	1	108	11.9	5.33	5478
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1	24	2	128	1	108	11.8	5.36	5587
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	8	10	44	10	2067	217.1	6.39	485
P01019	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	24	8	43	8	485	53.1	6.32	1196

P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	19	8	40	8	591	67	8.13	617
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG	19	7	47	7	525	59.5	7.5	1181
P18136	Ig kappa chain V-III region HIC OS=Homo sapiens PE=2	57	4	162	1	129	14.1	6.6	4598
P62873	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3	34	8	53	4	340	37.4	6	685
O14791	Apolipoprotein L1 OS=Homo sapiens GN=APOL1 PE=1	30	6	44	6	398	43.9	5.81	701
P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	50	8	42	8	147	15.9	5.76	1015
O00299	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4	44	6	52	6	241	26.9	5.17	1170
Q9Y696	Chloride intracellular channel protein 4 OS=Homo sapiens GN=CLIC4 PE=1 SV=4	43	6	30	6	253	28.8	5.59	490
Q8WUM4	Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1	10	6	27	6	868	96	6.52	813
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	7	8	44	8	2201	240.7	4.89	649
P07357	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	19	7	21	7	584	65.1	6.47	346
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	30	5	41	5	253	26.7	8.63	726
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3	41	5	25	2	181	20.6	7.43	628
P68104	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1	15	5	31	5	462	50.1	9.01	758
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	24	6	52	6	352	39	6.25	857
P09525	Annexin A4 OS=Homo sapiens GN=ANXA4 PE=1 SV=4	20	5	23	5	319	35.9	6.13	1236
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ	26	5	45	3	245	27.7	4.79	567
P01613	Ig kappa chain V-I region Ni OS=Homo sapiens PE=1 SV=1	30	2	45	2	112	12.2	5.36	1616
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1	33	2	80	1	113	12.3	6	2925
P01781	Ig heavy chain V-III region GAL OS=Homo sapiens PE=1	33	3	42	3	116	12.7	8.48	978
P07437	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	26	7	36	7	444	49.6	4.89	409
Q5JWF2	Guanine nucleotide-binding protein G(s) subunit alpha isoforms XLas OS=Homo sapiens GN=GNAS PE=1 SV=2	10	7	42	6	1037	111	5.03	678
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5	13	6	33	4	641	70	5.66	680
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4	7	8	40	8	1823	202.4	6.28	278
P50148	Guanine nucleotide-binding protein G(q) subunit alpha OS=Homo sapiens GN=GNAQ PE=1 SV=4	19	5	27	2	359	42.1	5.68	611

P23083	Ig heavy chain V-I region V35 OS=Homo sapiens PE=1	34	3	51	3	117	13	9.55	932
P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1	25	5	45	3	255	29.2	4.74	335
P68363	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B	25	7	32	7	451	50.1	5.06	440
P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1	34	3	123	2	120	13.2	6.57	3806
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	21	4	34	4	245	25.8	8.41	1012
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	53	6	42	6	165	18	7.81	609
P51884	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	20	5	40	5	338	38.4	6.61	469
Q08431	Lactadherin OS=Homo sapiens GN=MFGE8 PE=1 SV=2	24	6	35	6	387	43.1	8.15	522
P08697	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2	20	5	24	5	491	54.5	6.29	356
P18085	ADP-ribosylation factor 4 OS=Homo sapiens GN=ARF4	32	4	16	1	180	20.5	7.14	340
P13797	Plastin-3 OS=Homo sapiens GN=PLS3 PE=1 SV=4	11	5	27	5	630	70.8	5.6	577
P01611	Ig kappa chain V-I region Wes OS=Homo sapiens PE=1	31	2	38	2	108	11.6	7.28	1069
Q9UQB8	Brain-specific angiogenesis inhibitor 1-associated protein 2 OS=Homo sapiens GN=BAIAP2 PE=1 SV=1	15	7	20	7	552	60.8	8.9	356
O00159	Unconventional myosin-Ic OS=Homo sapiens GN=MYO1C	10	7	33	7	1063	121.6	9.41	477
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	9	5	32	5	724	83.2	5.03	877
P61224	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B	41	6	35	6	184	20.8	5.78	775
P36955	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	23	6	21	6	418	46.3	6.38	258
P01768	Ig heavy chain V-III region CAM OS=Homo sapiens PE=1	31	4	99	2	122	13.7	9.63	2176
Q9H4M9	EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2	22	7	29	7	534	60.6	6.83	280
P00338	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2	20	6	28	6	332	36.7	8.27	515
P01743	Ig heavy chain V-I region HG3 OS=Homo sapiens PE=4	56	4	47	4	117	12.9	8.92	1458
P22792	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2	19	6	25	6	545	60.5	5.99	335
Q8TE68	Epidermal growth factor receptor kinase substrate 8-like protein 1 OS=Homo sapiens GN=EPS8L1 PE=1 SV=3	13	6	21	6	723	80.2	6.04	310
P02774	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1	16	5	21	5	474	52.9	5.54	471
P20851	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	30	6	26	6	252	28.3	5.14	407
P08133	Annexin A6 OS=Homo sapiens GN=ANXA6 PE=1 SV=3	12	5	37	5	673	75.8	5.6	612

P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	34	4	28	4	202	22.3	8.31	759
P61225	Ras-related protein Rap-2b OS=Homo sapiens GN=RAP2B	40	5	23	5	183	20.5	4.81	581
P01764	Ig heavy chain V-III region VH26 OS=Homo sapiens PE=1	54	4	74	3	117	12.6	8.28	1251
P35542	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4	27	3	56	3	130	14.7	9.07	783
O43707	Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	9	6	16	6	911	104.8	5.44	275
P01763	Ig heavy chain V-III region WEA OS=Homo sapiens PE=1	36	3	82	2	114	12.2	8.5	2046
P08754	Guanine nucleotide-binding protein G(k) subunit alpha OS=Homo sapiens GN=GNAI3 PE=1 SV=3	16	4	38	2	354	40.5	5.69	954
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	47	3	50	2	100	10.7	6.52	884
Q95604	HLA class I histocompatibility antigen, Cw-17 alpha chain OS=Homo sapiens GN=HLA-C PE=1 SV=1	17	4	17	1	372	41.2	6.8	496
P01891	HLA class I histocompatibility antigen, A-68 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=4	18	4	25	2	365	40.9	6.7	561
P04075	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	19	5	26	5	364	39.4	8.09	586
O00560	Syntenin-1 OS=Homo sapiens GN=SDCBP PE=1 SV=1	24	4	20	4	298	32.4	7.53	276
Q562R1	Beta-actin-like protein 2 OS=Homo sapiens GN=ACTBL2	17	4	39	2	376	42	5.59	522
P15941	Mucin-1 OS=Homo sapiens GN=MUC1 PE=1 SV=3	3	4	32	4	1255	122	7.47	615
Q9HCB6	Spondin-1 OS=Homo sapiens GN=SPON1 PE=1 SV=2	8	5	27	5	807	90.9	6.11	419
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1	27	2	55	2	116	12.4	9.73	1049
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2	21	3	28	3	191	21.2	6.55	627
P02656	Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=1	27	2	11	2	99	10.8	5.41	431
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	20	4	24	4	240	25.8	6.61	618
P29992	Guanine nucleotide-binding protein subunit alpha-11 OS=Homo sapiens GN=GNA11 PE=1 SV=2	19	5	17	2	359	42.1	5.69	394
P03952	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	7	4	20	4	638	71.3	8.22	605
Q9NQ84	G-protein coupled receptor family C group 5 member C OS=Homo sapiens GN=GPRC5C PE=1 SV=2	11	4	19	4	441	48.2	8.43	657
P01008	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1	11	4	18	4	464	52.6	6.71	614
P10314	HLA class I histocompatibility antigen, A-32 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=2	19	4	19	2	365	41	6.47	437

P01861	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4	26	5	51	2	327	35.9	7.36	1026
P06313	Ig kappa chain V-IV region JI OS=Homo sapiens PE=4 SV=1	27	3	56	1	133	14.6	6.58	1122
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	21	4	39	4	224	25	5.63	536
P23528	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3	23	3	12	3	166	18.5	8.09	232
Q9NZA1	Chloride intracellular channel protein 5 OS=Homo sapiens GN=CLIC5 PE=1 SV=3	16	4	20	4	410	46.5	4.82	320
O00391	Sulfhydryl oxidase 1 OS=Homo sapiens GN=QSOX1 PE=1	7	3	23	3	747	82.5	8.92	462
P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1	43	3	33	3	100	11.2	6.62	290
P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4	17	2	43	2	146	16.2	8.28	1223
P08174	Complement decay-accelerating factor OS=Homo sapiens GN=CD55 PE=1 SV=4	16	3	19	3	381	41.4	7.59	284
P31946	14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB	15	3	34	1	246	28.1	4.83	263
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	22	3	39	3	156	18	9.64	675
P09543	2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Homo sapiens GN=CNP PE=1 SV=2	13	4	13	4	421	47.5	9.07	202
P01769	Ig heavy chain V-III region GA OS=Homo sapiens PE=1	31	4	26	4	122	13.2	9.77	261
O95436	Sodium-dependent phosphate transport protein 2B OS=Homo sapiens GN=SLC34A2 PE=1 SV=3	12	4	17	4	690	75.7	8.15	404
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	46	5	24	5	140	15	8.27	197
Q9H223	EH domain-containing protein 4 OS=Homo sapiens GN=EHD4 PE=1 SV=1	9	4	7	4	541	61.1	6.76	151
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	52	3	31	3	115	12.6	4.96	620
O14745	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Homo sapiens GN=SLC9A3R1 PE=1 SV=4	15	3	18	3	358	38.8	5.77	376
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS	21	4	30	4	223	25.4	6.54	530
P22105	Tenascin-X OS=Homo sapiens GN=TNXB PE=1 SV=3	3	3	9	3	4289	464	5.34	153
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1	22	2	34	2	111	11.9	5.08	1574
P60174	Triosephosphate isomerase OS=Homo sapiens GN=TPI1	22	3	10	3	286	30.8	5.92	131
P02790	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	15	4	13	4	462	51.6	7.02	184
P02749	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1	24	4	23	4	345	38.3	7.97	161
P05362	Intercellular adhesion molecule 1 OS=Homo sapiens GN=ICAM1 PE=1 SV=2	11	4	26	4	532	57.8	7.99	492

P13164	Interferon-induced transmembrane protein 1 OS=Homo sapiens GN=IFITM1 PE=1 SV=3	30	2	15	2	125	14	7.93	304
Q15485	Ficolin-2 OS=Homo sapiens GN=FCN2 PE=1 SV=2	15	3	14	3	313	34	6.77	336
P11233	Ras-related protein Ral-A OS=Homo sapiens GN=RALA	17	3	21	3	206	23.6	7.11	375
P04434	Ig kappa chain V-III region VH (Fragment) OS=Homo sapiens PE=4 SV=1	42	3	34	2	116	12.7	5.94	629
P61586	Transforming protein RhoA OS=Homo sapiens GN=RHOA	23	4	14	1	193	21.8	6.1	170
P00488	Coagulation factor XIII A chain OS=Homo sapiens GN=F13A1 PE=1 SV=4	8	4	21	4	732	83.2	6.09	172
Q8WUT4	Leucine-rich repeat neuronal protein 4 OS=Homo sapiens GN=LRRN4 PE=1 SV=3	9	3	13	3	740	78.8	7.2	137
Q9NQ79	Cartilage acidic protein 1 OS=Homo sapiens GN=CRTAC1	6	3	16	3	661	71.4	5.12	408
P01903	HLA class II histocompatibility antigen, DR alpha chain OS=Homo sapiens GN=HLA-DRA PE=1 SV=1	18	3	14	3	254	28.6	5	268
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	8	4	11	4	698	77	7.12	153
P60033	CD81 antigen OS=Homo sapiens GN=CD81 PE=1 SV=1	25	3	18	3	236	25.8	5.29	263
Q8IWA5	Choline transporter-like protein 2 OS=Homo sapiens GN=SLC44A2 PE=1 SV=3	8	4	16	4	706	80.1	8.57	250
Q15833	Syntaxin-binding protein 2 OS=Homo sapiens GN=STXBP2	12	4	15	4	593	66.4	6.55	224
P01892	HLA class I histocompatibility antigen, A-2 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=1	15	3	17	1	365	40.9	6.99	370
P12110	Collagen alpha-2(VI) chain OS=Homo sapiens GN=COL6A2	4	3	11	3	1019	108.5	6.21	305
P15924	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	2	4	10	4	2871	331.6	6.81	102
P01602	Ig kappa chain V-I region HK102 (Fragment) OS=Homo sapiens GN=IGKV1-5 PE=4 SV=1	15	1	24	1	117	12.8	6.51	404
Q8N512	Arrestin domain-containing protein 1 OS=Homo sapiens GN=ARRDC1 PE=2 SV=1	20	5	13	5	433	46	7.02	143
Q9UEW3	Macrophage receptor MARCO OS=Homo sapiens GN=MARCO PE=1 SV=1	5	2	16	2	520	52.6	8.78	639
O60635	Tetraspanin-1 OS=Homo sapiens GN=TSPAN1 PE=1 SV=2	5	1	12	1	241	26.3	5.25	583
P08134	Rho-related GTP-binding protein RhoC OS=Homo sapiens GN=RHOA PE=1 SV=1	23	4	13	1	193	22	6.58	126
Q96TA1	Niban-like protein 1 OS=Homo sapiens GN=FAM129B PE=1	8	4	13	4	746	84.1	6.19	179
P01717	Ig lambda chain V-IV region H1I OS=Homo sapiens PE=1	18	1	16	1	107	11.5	6.51	397

P33908	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IA OS=Homo sapiens GN=MAN1A1 PE=1 SV=3	9	4	13	4	653	72.9	6.47	116
P61981	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG	13	3	27	1	247	28.3	4.89	155
P21926	CD9 antigen OS=Homo sapiens GN=CD9 PE=1 SV=4	15	3	14	3	228	25.4	7.15	223
P14923	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3	5	3	7	3	745	81.7	6.14	141
P08670	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4	9	4	13	4	466	53.6	5.12	160
Q9P2B2	Prostaglandin F2 receptor negative regulator OS=Homo sapiens GN=PTGFRN PE=1 SV=2	6	4	14	4	879	98.5	6.61	104
P21980	Protein-glutamine gamma-glutamyltransferase 2 OS=Homo sapiens GN=TGM2 PE=1 SV=2	7	3	15	3	687	77.3	5.22	264
P10301	Ras-related protein R-Ras OS=Homo sapiens GN=RRAS	16	3	17	3	218	23.5	6.93	438
P06702	Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1	38	3	11	3	114	13.2	6.13	144
P04430	Ig kappa chain V-I region BAN OS=Homo sapiens PE=1	17	1	11	1	108	11.8	8.44	364
P01780	Ig heavy chain V-III region JON OS=Homo sapiens PE=1	17	1	16	1	115	12.6	9.29	396
P09758	Tumor-associated calcium signal transducer 2 OS=Homo sapiens GN=TACSTD2 PE=1 SV=3	10	2	11	2	323	35.7	8.87	493
P01624	Ig kappa chain V-III region POM OS=Homo sapiens PE=1	25	2	32	1	109	11.9	8.94	925
P01833	Polymeric immunoglobulin receptor OS=Homo sapiens GN=PIGR PE=1 SV=4	5	3	13	3	764	83.2	5.74	253
P01605	Ig kappa chain V-I region Lay OS=Homo sapiens PE=1	25	2	42	1	108	11.8	7.96	635
P27701	CD82 antigen OS=Homo sapiens GN=CD82 PE=1 SV=1	11	3	16	3	267	29.6	5.24	128
Q9BXR6	Complement factor H-related protein 5 OS=Homo sapiens GN=CFHR5 PE=1 SV=1	5	3	10	3	569	64.4	7.06	145
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1	17	1	9	1	108	11.4	6.52	380
Q99828	Calcium and integrin-binding protein 1 OS=Homo sapiens GN=CIB1 PE=1 SV=4	15	2	17	2	191	21.7	4.78	390
Q53GD3	Choline transporter-like protein 4 OS=Homo sapiens GN=SLC44A4 PE=2 SV=2	2	1	8	1	710	79.2	8.59	171
Q9UK55	Protein Z-dependent protease inhibitor OS=Homo sapiens GN=SERPINA10 PE=1 SV=1	6	2	8	2	444	50.7	8.27	385
O95445	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1	32	3	14	3	188	21.2	6.01	139
P61006	Ras-related protein Rab-8A OS=Homo sapiens GN=RAB8A	17	3	8	1	207	23.7	9.07	162
O95832	Claudin-1 OS=Homo sapiens GN=CLDN1 PE=1 SV=1	15	2	14	2	211	22.7	8.09	230
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3	7	2	15	2	381	41.9	5.01	419
P30041	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3	21	3	11	3	224	25	6.38	200

Q12805	EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 PE=1 SV=2	7	2	10	2	493	54.6	5.07	107
P27348	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1	15	3	26	1	245	27.7	4.78	101
P01616	Ig kappa chain V-II region MIL OS=Homo sapiens PE=1	33	2	44	1	112	12	9.29	2159
P62820	Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A	19	3	10	1	205	22.7	6.21	155
P06309	Ig kappa chain V-II region GM607 (Fragment) OS=Homo sapiens PE=4 SV=1	33	2	46	1	117	12.7	7.12	2159
P43353	Aldehyde dehydrogenase family 3 member B1 OS=Homo sapiens GN=ALDH3B1 PE=1 SV=1	8	3	11	3	468	51.8	7.62	248
P05023	Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1 PE=1 SV=1	2	2	12	2	1023	112.8	5.49	255
P09211	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1	15	2	17	2	210	23.3	5.64	348
P00751	Complement factor B OS=Homo sapiens GN=CFB PE=1	4	2	13	2	764	85.5	7.06	210
P00740	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	5	2	13	2	461	51.7	5.47	325
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1	19	2	34	1	129	14.3	8.51	543
P04208	Ig lambda chain V-I region WAH OS=Homo sapiens PE=1	28	2	24	1	109	11.7	6.54	618
P06889	Ig lambda chain V-IV region MOL OS=Homo sapiens PE=1	24	1	16	1	106	11.3	4.42	199
P09923	Intestinal-type alkaline phosphatase OS=Homo sapiens GN=ALPI PE=1 SV=2	8	3	11	1	528	56.8	5.86	183
P05164	Myeloperoxidase OS=Homo sapiens GN=MPO PE=1 SV=1	3	2	17	2	745	83.8	8.97	449
Q14112	Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3	3	2	16	2	1375	151.2	5.29	134
Q06830	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	15	3	11	3	199	22.1	8.13	146
Q96IY4	Carboxypeptidase B2 OS=Homo sapiens GN=CPB2 PE=1	4	2	17	2	423	48.4	7.71	473
P01703	Ig lambda chain V-I region NEWM OS=Homo sapiens PE=1	27	2	11	2	103	10.9	9.29	219
O75351	Vacuolar protein sorting-associated protein 4B OS=Homo sapiens GN=VPS4B PE=1 SV=2	7	2	12	2	444	49.3	7.23	194
P02655	Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1	32	2	5	2	101	11.3	4.72	111
P06319	Ig lambda chain V-VI region EB4 OS=Homo sapiens PE=2	15	1	9	1	131	14.1	5.05	391
P09972	Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	6	1	10	1	364	39.4	6.87	130
P22352	Glutathione peroxidase 3 OS=Homo sapiens GN=GPX3	14	2	11	2	226	25.5	8.13	119
P01601	Ig kappa chain V-I region HK101 (Fragment) OS=Homo sapiens PE=4 SV=1	14	1	9	1	117	12.8	8.16	529
P04216	Thy-1 membrane glycoprotein OS=Homo sapiens	16	2	10	2	161	17.9	8.73	277
Q969X1	Protein lifeguard 3 OS=Homo sapiens GN=TMBIM1 PE=1	5	1	9	1	311	34.6	7.72	524

O15551	Claudin-3 OS=Homo sapiens GN=CLDN3 PE=1 SV=1	13	2	17	2	220	23.3	8.05	170
P01772	Ig heavy chain V-III region KOL OS=Homo sapiens PE=1	13	1	9	1	126	13.7	5.87	600
P01615	Ig kappa chain V-II region FR OS=Homo sapiens PE=1 SV=1	35	2	44	2	113	12.7	7.94	801
P15169	Carboxypeptidase N catalytic chain OS=Homo sapiens GN=CPN1 PE=1 SV=1	3	1	6	1	458	52.3	7.34	207
Q9NZH0	G-protein coupled receptor family C group 5 member B OS=Homo sapiens GN=GPRC5B PE=2 SV=2	3	1	7	1	403	44.8	8.24	339
K7EJ46	Small integral membrane protein 22 OS=Homo sapiens GN=SMIM22 PE=2 SV=1	13	1	6	1	135	14.6	6.52	257
Q9UBI6	Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-12 OS=Homo sapiens GN=GNG12 PE=1 SV=3	22	1	8	1	72	8	8.97	134
Q86YZ3	Hornerin OS=Homo sapiens GN=HRNR PE=1 SV=2	2	1	5	1	2850	282.2	10.04	169
P01612	Ig kappa chain V-I region Mev OS=Homo sapiens PE=1	17	1	4	1	109	11.9	6.57	174
P60903	Protein S100-A10 OS=Homo sapiens GN=S100A10 PE=1	18	2	9	2	97	11.2	7.37	107
P04431	Ig kappa chain V-I region Walker OS=Homo sapiens PE=1	14	1	8	1	129	14.1	8.97	231
P06311	Ig kappa chain V-III region IARC/BL41 OS=Homo sapiens	12	1	9	1	128	14.1	6.61	127
A6NJS3	Putative V-set and immunoglobulin domain-containing-like protein IGHV1OR21-1 OS=Homo sapiens GN=IGHV1OR21-	13	1	6	1	120	13.4	7.81	130
P00742	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	2	1	5	1	488	54.7	5.94	163
P17931	Galectin-3 OS=Homo sapiens GN=LGALS3 PE=1 SV=5	7	2	10	2	250	26.1	8.56	119
Q15907	Ras-related protein Rab-11B OS=Homo sapiens	6	1	6	1	218	24.5	5.94	170
Q5D862	Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1	0	1	6	1	2391	247.9	8.31	205
P01771	Ig heavy chain V-III region HIL OS=Homo sapiens PE=1	9	1	12	1	121	13.6	9.36	332
P11597	Cholesteryl ester transfer protein OS=Homo sapiens GN=CETP PE=1 SV=2	3	1	6	1	493	54.7	6.09	162
P10599	Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3	12	1	9	1	105	11.7	4.92	235
Q96CX2	BTB/POZ domain-containing protein KCTD12 OS=Homo sapiens GN=KCTD12 PE=1 SV=1	3	1	7	1	325	35.7	5.64	174
P01775	Ig heavy chain V-III region LAY OS=Homo sapiens PE=1	16	1	9	1	119	12.9	8.48	150
P05556	Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2	1	1	8	1	798	88.4	5.39	116
P01762	Ig heavy chain V-III region TRO OS=Homo sapiens PE=1	8	1	5	1	122	13.5	9.72	134
P06326	Ig heavy chain V-I region Mot OS=Homo sapiens PE=1	10	1	9	1	125	13.6	6.29	100
A6NMB1	Sialic acid-binding Ig-like lectin 16 OS=Homo sapiens GN=SIGLEC16 PE=2 SV=3	1	1	42	1	481	53	9.03	454
P06316	Ig lambda chain V-I region BL2 OS=Homo sapiens PE=2	6	1	23	1	130	13.6	7.77	313

P32119	Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	9	1	8	1	198	21.9	5.97	188
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Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	61	207	3333	207	4563	515.3	7.05	74280
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	71	94	3263	94	1663	187	6.4	71963
P0C0L5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	61	80	2992	3	1744	192.6	7.27	73844
P0C0L4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	60	80	2964	3	1744	192.7	7.08	71295
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	51	74	1331	74	2386	262.5	5.71	30880
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	62	48	1659	48	1065	122.1	5.72	41707
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	53	56	974	49	1474	163.2	6.46	20263
P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	55	63	854	63	1676	188.2	6.52	16147
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	72	36	1576	36	491	55.9	8.27	43592
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	61	34	862	34	609	69.3	6.28	18104
Q9HC84	Mucin-5B OS=Homo sapiens GN=MUC5B PE=1 SV=3	16	50	415	50	5762	596	6.64	10550
P08519	Apolipoprotein(a) OS=Homo sapiens GN=LPA PE=1 SV=1	46	36	508	36	4548	501	5.88	9797
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	43	27	597	27	911	101.3	6.79	14322
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	63	29	897	29	597	67	7.3	18896
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	59	27	3257	10	452	49.3	6.77	86563
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	45	30	672	30	946	106.4	6.86	13117
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	74	27	1090	27	453	51.5	5.62	21862
P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	40	38	372	31	1482	163.8	6.38	6955
P00734	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	50	25	349	25	622	70	5.9	8887
Q8WX17	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	9	43	353	43	22152	2351.2	6	5802
P04220	Ig mu heavy chain disease protein OS=Homo sapiens PE=1 SV=1	55	18	2888	1	391	43	5.24	81879
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	22	40	230	40	2813	309.1	5.48	5554
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	33	18	772	18	500	55.1	6.55	16570
Q4LDE5	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 OS=Homo sapiens GN=SVEP1 PE=1 SV=3	20	41	212	41	3571	389.9	5.5	4564
P00736	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	44	20	315	19	705	80.1	6.21	8640
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	63	24	1160	24	267	30.8	5.76	27587
Q9Y6R7	IgGf-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3	21	36	234	36	5405	571.6	5.34	4544
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	38	31	281	30	1231	139	6.61	5420

P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	61	22	541	12	406	45.2	6.58	9270
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	41	16	269	16	493	54.7	7.01	5982
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	71	14	594	7	353	37.6	6.51	12811
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	37	23	320	23	930	103.3	6.98	6777
Q06033	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITIH3 PE=1 SV=2	38	22	192	22	890	99.8	5.74	3840
P09871	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	46	21	308	21	688	76.6	4.96	6003
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	54	20	259	20	418	46.7	5.59	5029
P02649	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	58	16	364	16	317	36.1	5.73	10225
P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2 PE=1 SV=3	59	11	403	4	340	36.5	6.1	9246
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	49	21	304	21	559	63.1	5.59	6916
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	37	13	414	13	478	54.3	5.8	8936
P10909	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	35	17	414	17	449	52.5	6.27	10453
P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	39	20	204	20	676	75.1	5.67	4027
P27169	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	60	13	302	12	355	39.7	5.22	6777
P02671	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2	25	19	576	19	866	94.9	6.01	8784
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	54	16	306	16	347	38.1	5.47	5862
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	94	10	795	10	106	11.6	5.87	22655
P15144	Aminopeptidase N OS=Homo sapiens GN=ANPEP PE=1 SV=4	36	25	164	25	967	109.5	5.48	3618
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	48	12	321	7	330	36.1	8.19	6569
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	16	16	267	16	1404	151	9.5	5311
P00747	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	42	21	167	21	810	90.5	7.24	2347
P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	69	10	154	10	147	15.9	5.76	5442
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	21	25	123	25	2201	240.7	4.89	2300
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	59	16	286	6	348	39	7.09	4134
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	46	13	146	13	423	47.6	5.52	3655
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	45	10	180	6	375	41.7	5.48	3654

Q08380	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	30	13	163	13	585	65.3	5.27	4404
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1 SV=3	8	16	126	16	3396	372.6	4.51	2921
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	31	15	108	15	703	77.2	5.22	2271
P05546	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	46	15	170	15	499	57	6.9	3220
P22792	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	41	13	120	13	545	60.5	5.99	2679
P06727	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	49	18	137	18	396	45.4	5.38	2318
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	51	10	236	4	326	35.9	7.59	3142
P35858	Insulin-like growth factor-binding protein complex acid labile subunit OS=Homo sapiens GN=IGFALS PE=1 SV=1	32	14	106	14	605	66	6.79	2452
P01597	Ig kappa chain V-I region DEE OS=Homo sapiens PE=1 SV=1	17	4	115	3	108	11.7	9.36	2861
P10643	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	29	14	90	14	843	93.5	6.48	2514
P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=4	7	20	106	20	4391	468.5	6.51	1782
Q7Z7G0	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1 SV=1	21	15	123	15	1075	118.6	9.44	2397
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	14	15	92	15	1786	197.9	4.94	2256
P13671	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	28	15	111	15	934	104.7	6.76	1872
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	33	9	137	9	352	39	6.25	2393
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	27	8	137	8	481	53.4	6.7	3550
P02774	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	46	14	80	14	474	52.9	5.54	1619
O14791	Apolipoprotein L1 OS=Homo sapiens GN=APOL1 PE=1 SV=5	35	8	88	8	398	43.9	5.81	2060
P22105	Tenascin-X OS=Homo sapiens GN=TNXB PE=1 SV=3	9	18	93	18	4289	464	5.34	1642
P01042	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	18	12	129	12	644	71.9	6.81	2980
P01019	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	26	9	79	9	485	53.1	6.32	2282
PODJ18	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	64	8	243	8	122	13.5	6.79	8464
O00391	Sulfhydryl oxidase 1 OS=Homo sapiens GN=QSOX1 PE=1 SV=3	22	10	70	10	747	82.5	8.92	1577
POCG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	75	5	453	2	106	11.3	7.24	10865
P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1 SV=4	60	7	188	7	159	18.1	5.24	5533
P06396	Gelsolin OS=Homo sapiens GN=GSN PE=1 SV=1	25	11	76	11	782	85.6	6.28	1443
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	27	13	94	9	586	69.4	6.27	1748
P14618	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	32	12	73	12	531	57.9	7.84	1498

B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	36	5	423	2	214	23	8.84	9223
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3	11	12	73	12	1609	177.5	5.12	1718
P51884	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	34	9	85	9	338	38.4	6.61	1362
P12111	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=5	11	18	62	18	3177	343.5	6.68	868
P07357	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	32	10	60	10	584	65.1	6.47	993
P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	26	10	56	10	591	67	8.13	1171
P07355	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	45	11	67	11	339	38.6	7.75	1770
P20851	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	44	8	104	8	252	28.3	5.14	1505
P00740	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	32	10	61	10	461	51.7	5.47	1361
P03952	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	27	11	58	11	638	71.3	8.22	1190
Q96PD5	N-acetylmuramoyl-L-alanine amidase OS=Homo sapiens GN=PGLYRP2 PE=1 SV=1	28	8	67	8	576	62.2	7.55	1707
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2	37	3	69	1	114	12.6	7.93	2801
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	33	6	53	6	335	36	8.46	1524
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	21	7	60	7	525	59.5	7.5	1605
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	43	11	149	5	377	41.3	7.9	2315
P04206	Ig kappa chain V-III region GOL OS=Homo sapiens PE=1 SV=1	49	3	84	1	109	11.8	9.25	4151
Q9UK55	Protein Z-dependent protease inhibitor OS=Homo sapiens GN=SERPINA10 PE=1 SV=1	30	9	38	9	444	50.7	8.27	1052
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1	60	4	117	1	109	11.8	8.48	4239
P08697	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	26	7	53	7	491	54.5	6.29	1133
P15169	Carboxypeptidase N catalytic chain OS=Homo sapiens GN=CPN1 PE=1 SV=1	30	7	38	7	458	52.3	7.34	725
P01606	Ig kappa chain V-I region OU OS=Homo sapiens PE=1 SV=1	17	2	73	1	108	11.8	9.91	2751
P01609	Ig kappa chain V-I region Scw OS=Homo sapiens PE=1 SV=1	28	2	76	1	108	11.8	6	2786
P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1 SV=1	31	2	75	1	108	11.8	8.91	2802

Q14520	Hyaluronan-binding protein 2 OS=Homo sapiens GN=HABP2 PE=1 SV=1	26	9	48	9	560	62.6	6.54	682
P01594	Ig kappa chain V-I region AU OS=Homo sapiens PE=1 SV=1	39	2	75	1	108	11.9	5.33	2734
Q9BXX0	EMILIN-2 OS=Homo sapiens GN=EMILIN2 PE=1 SV=3	20	11	48	11	1053	115.6	6.46	895
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1 SV=1	24	2	73	1	108	11.8	5.36	2726
P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	50	7	45	7	202	22.3	8.31	1130
P08571	Monocyte differentiation antigen CD14 OS=Homo sapiens GN=CD14 PE=1 SV=2	29	7	47	7	375	40.1	6.23	919
P01008	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	19	7	42	7	464	52.6	6.71	1060
O95445	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1 SV=2	38	5	50	5	188	21.2	6.01	954
P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1	67	6	138	6	100	11.2	6.62	2392
P04083	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	26	6	21	6	346	38.7	7.02	1009
P18136	Ig kappa chain V-III region HIC OS=Homo sapiens PE=2 SV=2	57	4	102	1	129	14.1	6.6	3028
P68032	Actin, alpha cardiac muscle 1 OS=Homo sapiens GN=ACTC1 PE=1 SV=1	23	6	89	2	377	42	5.39	1316
P35442	Thrombospondin-2 OS=Homo sapiens GN=THBS2 PE=1 SV=2	15	11	47	11	1172	129.9	4.83	488
P43251	Biotinidase OS=Homo sapiens GN=BTD PE=1 SV=2	16	5	36	5	543	61.1	6.25	382
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	7	8	37	8	2067	217.1	6.39	532
Q96IY4	Carboxypeptidase B2 OS=Homo sapiens GN=CPB2 PE=1 SV=2	34	9	46	9	423	48.4	7.71	751
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1 SV=1	36	5	88	4	115	12.4	9.25	1700
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4	9	9	39	9	1823	202.4	6.28	394
P33908	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IA OS=Homo sapiens GN=MAN1A1 PE=1 SV=3	21	7	47	7	653	72.9	6.47	576
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	22	6	32	6	434	47.1	7.39	665
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1 SV=1	31	5	43	4	108	11.6	9.25	1497
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	19	9	31	9	698	77	7.12	334
P19652	Alpha-1-acid glycoprotein 2 OS=Homo sapiens GN=ORM2 PE=1 SV=2	31	5	34	5	201	23.6	5.11	685
P00558	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3	27	8	45	8	417	44.6	8.1	855
P01613	Ig kappa chain V-I region Ni OS=Homo sapiens PE=1 SV=1	30	2	43	2	112	12.2	5.36	2026
P02790	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	21	6	29	6	462	51.6	7.02	845
P12259	Coagulation factor V OS=Homo sapiens GN=F5 PE=1 SV=4	4	7	31	7	2224	251.5	6.05	653
Q9NQ79	Cartilage acidic protein 1 OS=Homo sapiens GN=CRTAC1 PE=1 SV=2	14	6	40	6	661	71.4	5.12	926
P35542	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2	27	3	132	3	130	14.7	9.07	2208

P01861	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1	26	5	66	2	327	35.9	7.36	1137
P07437	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	30	8	41	8	444	49.6	4.89	534
P01764	Ig heavy chain V-III region VH26 OS=Homo sapiens PE=1 SV=1	54	4	45	3	117	12.6	8.28	774
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1	33	2	74	1	113	12.3	6	2611
O43490	Prominin-1 OS=Homo sapiens GN=PROM1 PE=1 SV=1	15	8	19	8	865	97.1	7.27	254
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	46	1	108	11.8	8.44	1697
P68363	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	27	7	30	7	451	50.1	5.06	493
P01596	Ig kappa chain V-I region CAR OS=Homo sapiens PE=1 SV=1	34	2	36	1	107	11.7	9.41	1371
Q9UHG3	Prenylcysteine oxidase 1 OS=Homo sapiens GN=PCYOX1 PE=1 SV=3	14	5	19	5	505	56.6	6.18	618
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	21	5	78	5	224	25	5.63	1580
P35579	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4	6	7	15	7	1960	226.4	5.6	287
P35443	Thrombospondin-4 OS=Homo sapiens GN=THBS4 PE=1 SV=2	13	6	28	5	961	105.8	4.68	668
P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1	25	2	89	1	120	13.2	6.57	2656
P00742	Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	15	6	18	6	488	54.7	5.94	445
Q99102	Mucin-4 OS=Homo sapiens GN=MUC4 PE=1 SV=4	7	7	40	7	2169	231.4	6.27	479
P21589	5'-nucleotidase OS=Homo sapiens GN=NT5E PE=1 SV=1	20	6	18	6	574	63.3	7.03	285
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	30	5	27	5	253	26.7	8.63	287
P05090	Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	26	4	52	4	189	21.3	5.15	734
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	9	5	24	2	724	83.2	5.03	808
P43652	Afamin OS=Homo sapiens GN=AFM PE=1 SV=1	16	6	36	6	599	69	5.9	653
P02656	Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=1 SV=1	27	2	28	2	99	10.8	5.41	994
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1 SV=1	27	2	49	2	116	12.4	9.73	961
P01781	Ig heavy chain V-III region GAL OS=Homo sapiens PE=1 SV=1	33	3	27	3	116	12.7	8.48	640
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	29	6	63	6	223	25.4	6.54	1190
Q8NFJ5	Retinoic acid-induced protein 3 OS=Homo sapiens GN=GPRC5A PE=1 SV=2	18	3	29	3	357	40.2	8.15	860
P01763	Ig heavy chain V-III region WEA OS=Homo sapiens PE=1 SV=1	36	3	71	2	114	12.2	8.5	1543
P02749	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	28	5	35	5	345	38.3	7.97	318
P01768	Ig heavy chain V-III region CAM OS=Homo sapiens PE=1 SV=1	18	3	58	1	122	13.7	9.63	1500
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	11	2	25	2	245	25.8	8.41	975

P02765	Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1	24	4	22	4	367	39.3	5.72	507
P04899	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3	21	5	15	5	355	40.4	5.54	231
P22352	Glutathione peroxidase 3 OS=Homo sapiens GN=GPX3 PE=1 SV=2	29	5	29	5	226	25.5	8.13	385
P00751	Complement factor B OS=Homo sapiens GN=CFB PE=1 SV=2	9	5	16	5	764	85.5	7.06	276
Q9HDC9	Adipocyte plasma membrane-associated protein OS=Homo sapiens GN=APMAP PE=1 SV=2	20	6	26	6	416	46.5	6.16	314
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	47	3	41	2	100	10.7	6.52	619
P49747	Cartilage oligomeric matrix protein OS=Homo sapiens GN=COMP PE=1 SV=2	7	3	19	2	757	82.8	4.6	578
P23083	Ig heavy chain V-I region V35 OS=Homo sapiens PE=1 SV=1	34	3	30	3	117	13	9.55	629
P80108	Phosphatidylinositol-glycan-specific phospholipase D OS=Homo sapiens GN=GPLD1 PE=1 SV=3	11	5	11	5	840	92.3	6.37	308
P07900	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	10	5	22	2	732	84.6	5.02	713
P06313	Ig kappa chain V-IV region JI OS=Homo sapiens PE=4 SV=1	27	3	30	1	133	14.6	6.58	590
Q03591	Complement factor H-related protein 1 OS=Homo sapiens GN=CFHR1 PE=1 SV=2	15	3	16	1	330	37.6	7.39	407
P07858	Cathepsin B OS=Homo sapiens GN=CTSB PE=1 SV=3	16	3	13	3	339	37.8	6.3	274
Q9HCB6	Spondin-1 OS=Homo sapiens GN=SPON1 PE=1 SV=2	10	5	25	5	807	90.9	6.11	346
P01880	Ig delta chain C region OS=Homo sapiens GN=IGHD PE=1 SV=2	16	4	14	4	384	42.2	7.93	203
P36955	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4	13	4	8	4	418	46.3	6.38	132
Q9NZP8	Complement C1r subcomponent-like protein OS=Homo sapiens GN=C1RL PE=1 SV=2	6	2	23	1	487	53.5	7.2	535
P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4 SV=1	17	2	37	2	146	16.2	8.28	851
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3	10	3	21	3	381	41.9	5.01	654
P48740	Mannan-binding lectin serine protease 1 OS=Homo sapiens GN=MASP1 PE=1 SV=3	12	5	13	5	699	79.2	5.49	120
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	20	4	26	4	240	25.8	6.61	538
P01743	Ig heavy chain V-I region HG3 OS=Homo sapiens PE=4 SV=1	22	2	34	2	117	12.9	8.92	725

P04180	Phosphatidylcholine-sterol acyltransferase OS=Homo sapiens GN=LCAT PE=1 SV=1	12	3	16	3	440	49.5	6.11	420
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	52	3	28	3	115	12.6	4.96	370
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3	10	5	29	1	577	67.8	6.4	377
P12109	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3	7	4	10	4	1028	108.5	5.43	209
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2	23	3	9	3	181	20.6	7.43	173
A1L4H1	Soluble scavenger receptor cysteine-rich domain-containing protein SSC5D OS=Homo sapiens GN=SSC5D PE=2 SV=3	4	3	11	3	1573	165.6	6.13	201
Q8TDL5	BPI fold-containing family B member 1 OS=Homo sapiens GN=BPIFB1 PE=1 SV=1	12	3	11	3	484	52.4	7.23	167
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1 SV=1	17	1	5	1	108	11.4	6.52	218
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	9	4	9	2	646	70.9	5.52	179
P02655	Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1 SV=1	32	2	22	2	101	11.3	4.72	519
P62879	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Homo sapiens GN=GNB2 PE=1 SV=3	18	4	11	3	340	37.3	6	167
P07585	Decorin OS=Homo sapiens GN=DCN PE=1 SV=1	18	4	16	4	359	39.7	8.54	157
P01717	Ig lambda chain V-IV region Hil OS=Homo sapiens PE=1 SV=1	18	1	18	1	107	11.5	6.51	527
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	22	2	30	2	111	11.9	5.08	1978
Q16610	Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2	14	5	17	5	540	60.6	6.71	146
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	32	4	18	4	165	18	7.81	362
Q08431	Lactadherin OS=Homo sapiens GN=MFGE8 PE=1 SV=2	14	4	19	4	387	43.1	8.15	194
Q9UEW3	Macrophage receptor MARCO OS=Homo sapiens GN=MARCO PE=1 SV=1	5	2	17	2	520	52.6	8.78	649
Q15485	Ficolin-2 OS=Homo sapiens GN=FCN2 PE=1 SV=2	12	2	15	2	313	34	6.77	387
P02654	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	27	4	36	4	83	9.3	8.47	221
P49908	Selenoprotein P OS=Homo sapiens GN=SEPP1 PE=1 SV=3	12	3	16	3	381	43.2	7.87	235
P06681	Complement C2 OS=Homo sapiens GN=C2 PE=1 SV=2	6	4	20	4	752	83.2	7.42	236
P02753	Retinol-binding protein 4 OS=Homo sapiens GN=RBP4 PE=1 SV=3	16	2	11	2	201	23	6.07	149
P22891	Vitamin K-dependent protein Z OS=Homo sapiens GN=PROZ PE=1 SV=2	20	5	20	5	400	44.7	5.97	149

P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	20	4	28	2	245	27.7	4.79	273
Q12805	EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 PE=1 SV=2	12	3	14	3	493	54.6	5.07	151
P04430	Ig kappa chain V-I region BAN OS=Homo sapiens PE=1 SV=1	17	1	8	1	108	11.8	8.44	330
P55056	Apolipoprotein C-IV OS=Homo sapiens GN=APOC4 PE=1 SV=1	32	4	12	4	127	14.5	8.92	113
P01769	Ig heavy chain V-III region GA OS=Homo sapiens PE=1 SV=1	31	3	19	3	122	13.2	9.77	206
P00488	Coagulation factor XIII A chain OS=Homo sapiens GN=F13A1 PE=1 SV=4	6	3	23	3	732	83.2	6.09	240
P34931	Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2	6	3	14	1	641	70.3	6.02	106
P01616	Ig kappa chain V-II region MIL OS=Homo sapiens PE=1 SV=1	33	2	47	1	112	12	9.29	1987
P04434	Ig kappa chain V-III region VH (Fragment) OS=Homo sapiens PE=4 SV=1	23	2	32	1	116	12.7	5.94	411
Q13790	Apolipoprotein F OS=Homo sapiens GN=APOF PE=1 SV=2	15	2	17	2	326	35.4	5.64	317
P01602	Ig kappa chain V-I region HK102 (Fragment) OS=Homo sapiens GN=IGKV1-5 PE=4 SV=1	15	1	13	1	117	12.8	6.51	331
P06319	Ig lambda chain V-VI region EB4 OS=Homo sapiens PE=2 SV=1	15	1	12	1	131	14.1	5.05	427
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	16	2	11	2	156	18	9.64	285
P01611	Ig kappa chain V-I region Wes OS=Homo sapiens PE=1 SV=1	17	1	15	1	108	11.6	7.28	283
Q14112	Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3	4	3	11	3	1375	151.2	5.29	131
P06310	Ig kappa chain V-II region RPMI 6410 OS=Homo sapiens PE=4 SV=1	29	2	48	1	133	14.7	9.25	1960
Q9NQ84	G-protein coupled receptor family C group 5 member C OS=Homo sapiens GN=GPRC5C PE=1 SV=2	3	1	4	1	441	48.2	8.43	338
P09211	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	15	2	12	2	210	23.3	5.64	207
O75636	Ficolin-3 OS=Homo sapiens GN=FCN3 PE=1 SV=2	11	2	14	2	299	32.9	6.67	179
P06889	Ig lambda chain V-IV region MOL OS=Homo sapiens PE=1 SV=1	24	1	15	1	106	11.3	4.42	176
P01601	Ig kappa chain V-I region HK101 (Fragment) OS=Homo sapiens PE=4 SV=1	14	1	7	1	117	12.8	8.16	402
P01605	Ig kappa chain V-I region Lay OS=Homo sapiens PE=1 SV=1	25	2	31	1	108	11.8	7.96	347
O00187	Mannan-binding lectin serine protease 2 OS=Homo sapiens GN=MASP2 PE=1 SV=4	5	3	11	3	686	75.7	5.63	127
P04208	Ig lambda chain V-I region WAH OS=Homo sapiens PE=1 SV=1	28	2	24	1	109	11.7	6.54	535

P27105	Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=1 SV=3	17	2	5	2	288	31.7	7.88	100
P41222	Prostaglandin-H2 D-isomerase OS=Homo sapiens GN=PTGDS PE=1 SV=1	17	2	7	2	190	21	7.8	210
Q07954	Prolow-density lipoprotein receptor-related protein 1 OS=Homo sapiens GN=LRP1 PE=1 SV=2	1	2	7	2	4544	504.3	5.39	106
P19971	Thymidine phosphorylase OS=Homo sapiens GN=TYMP PE=1 SV=2	3	1	8	1	482	49.9	5.53	287
P01772	Ig heavy chain V-III region KOL OS=Homo sapiens PE=1 SV=1	13	1	3	1	126	13.7	5.87	155
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1 SV=2	19	2	33	1	129	14.3	8.51	298
Q15166	Serum paraoxonase/lactonase 3 OS=Homo sapiens GN=PON3 PE=1 SV=3	6	2	12	1	354	39.6	5.41	143
P09758	Tumor-associated calcium signal transducer 2 OS=Homo sapiens GN=TACSTD2 PE=1 SV=3	5	1	8	1	323	35.7	8.87	264
P01780	Ig heavy chain V-III region JON OS=Homo sapiens PE=1 SV=1	17	1	8	1	115	12.6	9.29	169
P04075	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	5	1	7	1	364	39.4	8.09	249
A6NIZ1	Ras-related protein Rap-1b-like protein OS=Homo sapiens PE=2 SV=1	7	1	8	1	184	20.9	5.48	285
P01833	Polymeric immunoglobulin receptor OS=Homo sapiens GN=PIGR PE=1 SV=4	2	1	6	1	764	83.2	5.74	219
Q6UX71	Plexin domain-containing protein 2 OS=Homo sapiens GN=PLXDC2 PE=1 SV=1	2	1	3	1	529	59.5	6.46	106
Q9Y5Y7	Lymphatic vessel endothelial hyaluronic acid receptor 1 OS=Homo sapiens GN=LYVE1 PE=1 SV=2	6	2	8	2	322	35.2	8.28	107
O00560	Syntenin-1 OS=Homo sapiens GN=SDCBP PE=1 SV=1	8	1	7	1	298	32.4	7.53	165
P14625	Endoplasmin OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	2	1	6	1	803	92.4	4.84	212
P01762	Ig heavy chain V-III region TRO OS=Homo sapiens PE=1 SV=1	8	2	10	2	122	13.5	9.72	147
P01771	Ig heavy chain V-III region HIL OS=Homo sapiens PE=1 SV=1	9	1	11	1	121	13.6	9.36	171
P08174	Complement decay-accelerating factor OS=Homo sapiens GN=CD55 PE=1 SV=4	3	1	9	1	381	41.4	7.59	124
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	10	1	7	1	140	15	8.27	111
P07711	Cathepsin L1 OS=Homo sapiens GN=CTSL PE=1 SV=2	4	1	5	1	333	37.5	5.45	114
P49913	Cathelicidin antimicrobial peptide OS=Homo sapiens GN=CAMP PE=1 SV=1	5	1	8	1	170	19.3	9.41	193
P06326	Ig heavy chain V-I region Mot OS=Homo sapiens PE=1 SV=1	10	1	9	1	125	13.6	6.29	117

P18827	Syndecan-1 OS=Homo sapiens GN=SDC1 PE=1 SV=3	5	1	9	1	310	32.4	4.63	151
P01615	Ig kappa chain V-II region FR OS=Homo sapiens PE=1 SV=1	12	1	32	1	113	12.7	7.94	664

Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	34	115	929	115	4563	515.3	7.05	26194
POCOL5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	49	58	662	4	1744	192.6	7.27	17479
POCOL4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	47	55	630	1	1744	192.7	7.08	16061
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	49	51	526	44	1474	163.2	6.46	14145
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	61	29	2167	12	452	49.3	6.77	53780
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	18	27	266	27	2386	262.5	5.71	6671
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	28	33	216	33	1663	187	6.4	5758
P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	28	30	236	23	1482	163.8	6.38	6245
P04220	Ig mu heavy chain disease protein OS=Homo sapiens PE=1 SV=1	58	19	1392	2	391	43	5.24	28485
Q8WXI7	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	9	32	258	32	22152	2351.2	6	5363
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	33	16	327	16	500	55.1	6.55	7631
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	89	8	653	8	106	11.6	5.87	24001
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	30	15	206	15	597	67	7.3	3731
P15924	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	9	19	43	19	2871	331.6	6.81	1243
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	46	13	158	13	347	38.1	5.47	5117
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	10	14	116	14	2201	240.7	4.89	2626
P07355	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	48	15	105	15	339	38.6	7.75	3158
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	29	15	104	15	609	69.3	6.28	2586
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1 SV=3	5	12	158	12	3396	372.6	4.51	4489
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	31	15	107	8	586	69.4	6.27	2216
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	50	12	144	9	330	36.1	8.19	2951
P01597	Ig kappa chain V-I region DEE OS=Homo sapiens PE=1 SV=1	17	3	120	2	108	11.7	9.36	3892
P12111	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=5	6	13	96	13	3177	343.5	6.68	2897
P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1 SV=4	67	8	147	8	159	18.1	5.24	5321
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3	8	10	87	10	1609	177.5	5.12	2905
Q08380	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	23	10	79	10	585	65.3	5.27	1758
P11226	Mannose-binding protein C OS=Homo sapiens GN=MBL2 PE=1 SV=2	45	11	99	11	248	26.1	5.49	2220
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	41	6	279	3	214	23	8.84	7586
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1 SV=1	31	5	87	4	108	11.6	9.25	2819

P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	29	10	85	5	406	45.2	6.58	1862
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	75	5	289	1	106	11.3	7.24	8801
P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=4	3	9	65	9	4391	468.5	6.51	1575
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	30	6	60	6	240	25.8	6.61	1910
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	33	9	69	5	375	41.7	5.48	1592
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	44	10	76	10	267	30.8	5.76	2169
Q9Y6R7	IgGfc-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3	4	11	79	11	5405	571.6	5.34	2395
P26038	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3	20	10	80	4	577	67.8	6.4	1802
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	37	9	83	4	353	37.6	6.51	2012
P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1 SV=1	31	2	91	1	108	11.8	8.91	4459
P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	10	11	58	11	1676	188.2	6.52	1161
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	30	8	62	3	348	39	7.09	1244
P01594	Ig kappa chain V-I region AU OS=Homo sapiens PE=1 SV=1	31	2	75	1	108	11.9	5.33	3906
P01606	Ig kappa chain V-I region OU OS=Homo sapiens PE=1 SV=1	17	2	83	1	108	11.8	9.91	4422
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2	37	3	61	3	114	12.6	7.93	2749
P12110	Collagen alpha-2(VI) chain OS=Homo sapiens GN=COL6A2 PE=1 SV=4	11	7	40	7	1019	108.5	6.21	1267
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	12	8	59	8	930	103.3	6.98	1497
P01609	Ig kappa chain V-I region Scw OS=Homo sapiens PE=1 SV=1	28	2	79	1	108	11.8	6	4006
P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	17	9	51	9	676	75.1	5.67	1279
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1 SV=1	36	6	139	5	115	12.4	9.25	3139
O95436	Sodium-dependent phosphate transport protein 2B OS=Homo sapiens GN=SLC34A2 PE=1 SV=3	13	5	45	5	690	75.7	8.15	1191
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	37	8	40	8	320	35.9	5.05	1356
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	70	2	108	11.8	8.44	3458
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1 SV=1	24	2	79	1	108	11.8	5.36	3862
P12109	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3	9	6	46	6	1028	108.5	5.43	1461
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	6	7	47	7	1786	197.9	4.94	1519
P48740	Mannan-binding lectin serine protease 1 OS=Homo sapiens GN=MASP1 PE=1 SV=3	15	7	54	7	699	79.2	5.49	1525

P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2 PE=1 SV=3	23	6	51	1	340	36.5	6.1	1552
P35241	Radixin OS=Homo sapiens GN=RDX PE=1 SV=1	15	8	56	1	583	68.5	6.37	1143
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1	39	3	78	1	109	11.8	8.48	3188
P04206	Ig kappa chain V-III region GOL OS=Homo sapiens PE=1 SV=1	49	3	59	1	109	11.8	9.25	2889
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	14	7	46	7	559	63.1	5.59	1387
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	5	5	37	5	1404	151	9.5	869
P01611	Ig kappa chain V-I region Wes OS=Homo sapiens PE=1 SV=1	31	2	43	2	108	11.6	7.28	2335
O00560	Syntenin-1 OS=Homo sapiens GN=SDCBP PE=1 SV=1	29	4	22	4	298	32.4	7.53	411
Q02413	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	7	5	14	5	1049	113.7	5.03	451
A0M8Q6	Ig lambda-7 chain C region OS=Homo sapiens GN=IGLC7 PE=1 SV=2	51	3	102	1	106	11.3	8.28	2152
P01613	Ig kappa chain V-I region Ni OS=Homo sapiens PE=1 SV=1	30	2	51	2	112	12.2	5.36	2434
P01768	Ig heavy chain V-III region CAM OS=Homo sapiens PE=1 SV=1	39	6	121	4	122	13.7	9.63	2337
Q9UGT4	Sushi domain-containing protein 2 OS=Homo sapiens GN=SUSD2 PE=1 SV=1	9	4	18	4	822	90.2	6.28	801
Q9HC84	Mucin-5B OS=Homo sapiens GN=MUC5B PE=1 SV=3	2	6	30	6	5762	596	6.64	744
P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1	34	3	107	2	120	13.2	6.57	3181
P14618	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	11	5	34	5	531	57.9	7.84	776
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	25	5	33	3	245	27.7	4.79	705
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1	33	2	33	1	113	12.3	6	1383
P62873	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3	22	5	26	2	340	37.4	6	633
P04083	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	16	4	27	4	346	38.7	7.02	1355
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	14	5	25	4	646	70.9	5.52	647
P62879	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Homo sapiens GN=GNB2 PE=1 SV=3	22	5	23	2	340	37.3	6	723
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	22	6	73	3	326	35.9	7.59	1313
P01769	Ig heavy chain V-III region GA OS=Homo sapiens PE=1 SV=1	25	4	30	4	122	13.2	9.77	480
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	20	4	28	4	223	25.4	6.54	989
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1 SV=1	27	2	55	2	116	12.4	9.73	1199
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	22	3	46	3	156	18	9.64	836
P62736	Actin, aortic smooth muscle OS=Homo sapiens GN=ACTA2 PE=1 SV=1	16	5	30	1	377	42	5.39	539

P01833	Polymeric immunoglobulin receptor OS=Homo sapiens GN=PIGR PE=1 SV=4	7	5	23	5	764	83.2	5.74	422
P05362	Intercellular adhesion molecule 1 OS=Homo sapiens GN=ICAM1 PE=1 SV=2	9	4	25	4	532	57.8	7.99	731
P50995	Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1	14	6	18	6	505	54.4	7.65	377
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	13	4	29	4	423	47.6	5.52	655
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	5	4	36	4	911	101.3	6.79	840
P12429	Annexin A3 OS=Homo sapiens GN=ANXA3 PE=1 SV=3	19	4	15	4	323	36.4	5.92	460
P01763	Ig heavy chain V-III region WEA OS=Homo sapiens PE=1 SV=1	36	3	99	2	114	12.2	8.5	1908
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	2	4	15	4	2067	217.1	6.39	708
O00159	Unconventional myosin-Ic OS=Homo sapiens GN=MYO1C PE=1 SV=4	5	4	21	4	1063	121.6	9.41	398
P01743	Ig heavy chain V-I region HG3 OS=Homo sapiens PE=4 SV=1	44	3	39	3	117	12.9	8.92	859
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2	26	3	25	3	181	20.6	7.43	634
P09525	Annexin A4 OS=Homo sapiens GN=ANXA4 PE=1 SV=4	12	3	22	3	319	35.9	6.13	836
P60174	Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=3	17	3	8	3	286	30.8	5.92	233
P01616	Ig kappa chain V-II region MIL OS=Homo sapiens PE=1 SV=1	33	2	36	1	112	12	9.29	1397
P01781	Ig heavy chain V-III region GAL OS=Homo sapiens PE=1 SV=1	31	3	21	3	116	12.7	8.48	478
Q08431	Lactadherin OS=Homo sapiens GN=MFGE8 PE=1 SV=2	15	4	26	4	387	43.1	8.15	470
P15941	Mucin-1 OS=Homo sapiens GN=MUC1 PE=1 SV=3	4	4	31	4	1255	122	7.47	649
P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4 SV=1	17	2	40	2	146	16.2	8.28	1229
Q15485	Ficolin-2 OS=Homo sapiens GN=FCN2 PE=1 SV=2	15	3	12	3	313	34	6.77	436
Q01628	Interferon-induced transmembrane protein 3 OS=Homo sapiens GN=IFITM3 PE=1 SV=2	12	1	27	1	133	14.6	7.01	1495
P05186	Alkaline phosphatase, tissue-nonspecific isozyme OS=Homo sapiens GN=ALPL PE=1 SV=4	13	5	26	5	524	57.3	6.67	487
P14923	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3	7	5	16	5	745	81.7	6.14	327
P04899	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3	11	3	16	2	355	40.4	5.54	528
Q8IWA5	Choline transporter-like protein 2 OS=Homo sapiens GN=SLC44A2 PE=1 SV=3	8	4	24	4	706	80.1	8.57	410
O95832	Claudin-1 OS=Homo sapiens GN=CLDN1 PE=1 SV=1	15	2	23	2	211	22.7	8.09	625

Q4LDE5	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 OS=Homo sapiens GN=SVEP1 PE=1 SV=3	1	3	16	3	3571	389.9	5.5	665
P08519	Apolipoprotein(a) OS=Homo sapiens GN=LPA PE=1 SV=1	26	4	22	4	4548	501	5.88	398
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	15	3	17	3	335	36	8.46	414
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	11	4	22	4	453	51.5	5.62	564
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	47	3	18	3	100	10.7	6.52	469
P83593	Ig kappa chain V-IV region STH (Fragment) OS=Homo sapiens PE=1 SV=1	28	2	24	2	109	12.1	7.94	678
P56747	Claudin-6 OS=Homo sapiens GN=CLDN6 PE=1 SV=2	13	2	23	1	220	23.3	8	592
Q9NQ84	G-protein coupled receptor family C group 5 member C OS=Homo sapiens GN=GPRC5C PE=1 SV=2	9	3	12	3	441	48.2	8.43	863
P21926	CD9 antigen OS=Homo sapiens GN=CD9 PE=1 SV=4	10	4	52	4	228	25.4	7.15	658
P01624	Ig kappa chain V-III region POM OS=Homo sapiens PE=1 SV=1	25	2	24	1	109	11.9	8.94	698
P30501	HLA class I histocompatibility antigen, Cw-2 alpha chain OS=Homo sapiens GN=HLA-C PE=1 SV=1	13	3	16	3	366	41.1	6.11	333
P21589	5'-nucleotidase OS=Homo sapiens GN=NT5E PE=1 SV=1	12	4	8	4	574	63.3	7.03	208
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITI2 PE=1 SV=2	5	3	16	3	946	106.4	6.86	376
P06744	Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4	7	3	13	3	558	63.1	8.32	288
O15551	Claudin-3 OS=Homo sapiens GN=CLDN3 PE=1 SV=1	13	2	19	1	220	23.3	8.05	450
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1 SV=1	17	1	15	1	108	11.4	6.52	700
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	22	2	20	2	111	11.9	5.08	1203
P07737	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	31	3	27	3	140	15	8.27	519
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	12	3	15	3	352	39	6.25	352
Q8WUM4	Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1	6	4	10	4	868	96	6.52	113
P27105	Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=1 SV=3	10	2	18	2	288	31.7	7.88	501
Q8NFJ5	Retinoic acid-induced protein 3 OS=Homo sapiens GN=GPRC5A PE=1 SV=2	9	2	19	2	357	40.2	8.15	671
P04434	Ig kappa chain V-III region VH (Fragment) OS=Homo sapiens PE=4 SV=1	23	2	20	1	116	12.7	5.94	394

P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1	12	3	16	1	255	29.2	4.74	146
P61224	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1	18	3	18	3	184	20.8	5.78	571
P05090	Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	20	3	24	3	189	21.3	5.15	300
P01764	Ig heavy chain V-III region VH26 OS=Homo sapiens PE=1 SV=1	38	3	55	2	117	12.6	8.28	1167
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	7	2	10	2	434	47.1	7.39	374
P63000	Ras-related C3 botulinum toxin substrate 1 OS=Homo sapiens GN=RAC1 PE=1 SV=1	17	3	19	3	192	21.4	8.5	279
P15144	Aminopeptidase N OS=Homo sapiens GN=ANPEP PE=1 SV=4	4	3	13	3	967	109.5	5.48	285
Q08722	Leukocyte surface antigen CD47 OS=Homo sapiens GN=CD47 PE=1 SV=1	9	3	31	3	323	35.2	7.21	672
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	9	3	22	3	418	46.7	5.59	424
P01717	Ig lambda chain V-IV region H11 OS=Homo sapiens PE=1 SV=1	18	1	12	1	107	11.5	6.51	267
Q9H4G4	Golgi-associated plant pathogenesis-related protein 1 OS=Homo sapiens GN=GLIPR2 PE=1 SV=3	17	2	14	2	154	17.2	9.41	622
Q08188	Protein-glutamine gamma-glutamyltransferase E OS=Homo sapiens GN=TGM3 PE=1 SV=4	5	3	7	3	693	76.6	5.86	140
P01602	Ig kappa chain V-I region HK102 (Fragment) OS=Homo sapiens GN=IGKV1-5 PE=4 SV=1	14	1	13	1	117	12.8	6.51	718
Q5JWF2	Guanine nucleotide-binding protein G(s) subunit alpha isoforms XLas OS=Homo sapiens GN=GNAS PE=1 SV=2	2	2	13	1	1037	111	5.03	342
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1 SV=2	19	2	20	1	129	14.3	8.51	480
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	7	3	18	3	481	53.4	6.7	322
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	7	4	12	4	703	77.2	5.22	225
P22735	Protein-glutamine gamma-glutamyltransferase K OS=Homo sapiens GN=TGM1 PE=1 SV=4	4	2	15	2	817	89.7	6.04	544
P06319	Ig lambda chain V-VI region EB4 OS=Homo sapiens PE=2 SV=1	15	1	9	1	131	14.1	5.05	716
O15162	Phospholipid scramblase 1 OS=Homo sapiens GN=PLSCR1 PE=1 SV=1	6	1	16	1	318	35	4.94	322
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	11	2	7	2	253	26.7	8.63	319
P62330	ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2	18	2	13	2	175	20.1	8.95	372
P06311	Ig kappa chain V-III region IARC/BL41 OS=Homo sapiens PE=1 SV=1	12	1	15	1	128	14.1	6.61	351
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	12	2	17	2	245	25.8	8.41	427

P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	4	2	12	2	591	67	8.13	264
P04208	Ig lambda chain V-I region WAH OS=Homo sapiens PE=1 SV=1	28	2	16	2	109	11.7	6.54	492
P04430	Ig kappa chain V-I region BAN OS=Homo sapiens PE=1 SV=1	17	1	14	1	108	11.8	8.44	531
O43490	Prominin-1 OS=Homo sapiens GN=PROM1 PE=1 SV=1	4	2	10	2	865	97.1	7.27	175
P17066	Heat shock 70 kDa protein 6 OS=Homo sapiens GN=HSPA6 PE=1 SV=2	5	2	14	1	643	71	6.14	346
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	1	2	8	2	2813	309.1	5.48	209
P59190	Ras-related protein Rab-15 OS=Homo sapiens GN=RAB15 PE=1 SV=1	10	2	9	1	212	24.4	5.71	164
P01605	Ig kappa chain V-I region Lay OS=Homo sapiens PE=1 SV=1	25	2	21	1	108	11.8	7.96	329
P09758	Tumor-associated calcium signal transducer 2 OS=Homo sapiens GN=TACSTD2 PE=1 SV=3	5	1	9	1	323	35.7	8.87	503
P06889	Ig lambda chain V-IV region MOL OS=Homo sapiens PE=1 SV=1	24	1	15	1	106	11.3	4.42	189
P01703	Ig lambda chain V-I region NEWM OS=Homo sapiens PE=1 SV=1	27	2	12	2	103	10.9	9.29	275
P51153	Ras-related protein Rab-13 OS=Homo sapiens GN=RAB13 PE=1 SV=1	11	2	4	1	203	22.8	9.19	129
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	9	2	7	2	224	25	5.63	178
P01701	Ig lambda chain V-I region NEW OS=Homo sapiens PE=1 SV=1	27	2	18	2	111	11.4	8	359
P06576	ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3	3	1	8	1	529	56.5	5.4	405
P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	7	1	9	1	202	22.3	8.31	512
Q53GD3	Choline transporter-like protein 4 OS=Homo sapiens GN=SLC44A4 PE=2 SV=2	3	2	15	2	710	79.2	8.59	196
O60635	Tetraspanin-1 OS=Homo sapiens GN=TSPAN1 PE=1 SV=2	5	1	6	1	241	26.3	5.25	402
P01903	HLA class II histocompatibility antigen, DR alpha chain OS=Homo sapiens GN=HLA-DRA PE=1 SV=1	7	1	9	1	254	28.6	5	441
O00187	Mannan-binding lectin serine protease 2 OS=Homo sapiens GN=MASP2 PE=1 SV=4	4	2	8	2	686	75.7	5.63	202
P01762	Ig heavy chain V-III region TRO OS=Homo sapiens PE=1 SV=1	8	2	13	2	122	13.5	9.72	191
Q92485	Acid sphingomyelinase-like phosphodiesterase 3b OS=Homo sapiens GN=SMPDL3B PE=2 SV=2	8	2	7	2	455	50.8	5.64	199
P04216	Thy-1 membrane glycoprotein OS=Homo sapiens GN=THY1 PE=1 SV=2	16	2	10	2	161	17.9	8.73	356
P01040	Cystatin-A OS=Homo sapiens GN=CSTA PE=1 SV=1	18	1	4	1	98	11	5.5	182

Q96S97	Myeloid-associated differentiation marker OS=Homo sapiens GN=MYADM PE=1 SV=2	10	2	8	2	322	35.3	8.15	158
P01612	Ig kappa chain V-I region Mev OS=Homo sapiens PE=1 SV=1	17	1	8	1	109	11.9	6.57	259
Q9NZH0	G-protein coupled receptor family C group 5 member B OS=Homo sapiens GN=GPRC5B PE=2 SV=2	3	1	9	1	403	44.8	8.24	622
P01772	Ig heavy chain V-III region KOL OS=Homo sapiens PE=1 SV=1	13	1	9	1	126	13.7	5.87	658
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	26	2	13	2	115	12.6	4.96	374
P01601	Ig kappa chain V-I region HK101 (Fragment) OS=Homo sapiens PE=4 SV=1	14	1	2	1	117	12.8	8.16	108
Q08554	Desmocollin-1 OS=Homo sapiens GN=DSC1 PE=1 SV=2	2	1	2	1	894	99.9	5.43	118
P16422	Epithelial cell adhesion molecule OS=Homo sapiens GN=EPCAM PE=1 SV=2	5	1	4	1	314	34.9	7.46	169
P10599	Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3	12	1	9	1	105	11.7	4.92	421
P24043	Laminin subunit alpha-2 OS=Homo sapiens GN=LAMA2 PE=1 SV=4	0	1	6	1	3122	343.7	6.4	259
Q06830	Peroxisome oxidoreductin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	6	1	7	1	199	22.1	8.13	256
P35030	Trypsin-3 OS=Homo sapiens GN=PRSS3 PE=1 SV=2	4	1	3	1	304	32.5	7.49	105
Q86X29	Lipolysis-stimulated lipoprotein receptor OS=Homo sapiens GN=LSR PE=1 SV=4	2	1	5	1	649	71.4	7.97	154
P01780	Ig heavy chain V-III region JON OS=Homo sapiens PE=1 SV=1	17	1	23	1	115	12.6	9.29	419
P01771	Ig heavy chain V-III region HIL OS=Homo sapiens PE=1 SV=1	9	1	4	1	121	13.6	9.36	159
Q9BVK6	Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=2	5	1	6	1	235	27.3	8.02	129
P23528	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3	7	1	6	1	166	18.5	8.09	139
P81605	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	10	1	7	1	110	11.3	6.54	187
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	5	1	6	1	165	18	7.81	198
P23083	Ig heavy chain V-I region V35 OS=Homo sapiens PE=1 SV=1	11	1	9	1	117	13	9.55	166
Q05639	Elongation factor 1-alpha 2 OS=Homo sapiens GN=EEF1A2 PE=1 SV=1	2	1	7	1	463	50.4	9.03	245
P01911	HLA class II histocompatibility antigen, DRB1-15 beta chain OS=Homo sapiens GN=HLA-DRB1 PE=1 SV=2	3	1	6	1	266	29.9	7.74	130
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	2	1	9	1	493	54.7	7.01	120
P01603	Ig kappa chain V-I region Ka OS=Homo sapiens PE=1 SV=1	15	1	9	1	108	11.9	8.85	218
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	3	1	4	1	478	54.3	5.8	110

Q9BXR6	Complement factor H-related protein 5 OS=Homo sapiens GN=CFHR5 PE=1 SV=1	2	1	4	1	569	64.4	7.06	158
P29992	Guanine nucleotide-binding protein subunit alpha-11 OS=Homo sapiens GN=GNA11 PE=1 SV=2	3	1	3	1	359	42.1	5.69	100
P08174	Complement decay-accelerating factor OS=Homo sapiens GN=CD55 PE=1 SV=4	3	1	9	1	381	41.4	7.59	129
P01615	Ig kappa chain V-II region FR OS=Homo sapiens PE=1 SV=1	12	1	8	1	113	12.7	7.94	154
P29373	Cellular retinoic acid-binding protein 2 OS=Homo sapiens GN=CRABP2 PE=1 SV=2	7	1	9	1	138	15.7	5.4	143
P04211	Ig lambda chain V region 4A OS=Homo sapiens PE=4 SV=1	8	1	10	1	117	12.4	7.03	212
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3	2	1	5	1	381	41.9	5.01	113
P06316	Ig lambda chain V-I region BL2 OS=Homo sapiens PE=2 SV=1	6	1	24	1	130	13.6	7.77	243

Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	53	188	2481	188	4563	515.3	7.05	62411
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	71	98	2809	98	1663	187	6.4	69432
P0C0L5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	58	81	2942	4	1744	192.6	7.27	70717
P0C0L4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	57	80	2835	3	1744	192.7	7.08	67303
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	46	66	1235	66	2386	262.5	5.71	34719
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	52	51	746	44	1474	163.2	6.46	18662
P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	48	53	648	53	1676	188.2	6.52	14758
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	53	27	1416	27	491	55.9	8.27	38657
P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=4	21	55	316	55	4391	468.5	6.51	8875
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	68	37	655	37	609	69.3	6.28	14094
P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	39	41	438	34	1482	163.8	6.38	9853
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	39	30	566	30	1065	122.1	5.72	15139
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	62	30	2258	11	452	49.3	6.77	53104
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	26	38	305	38	2201	240.7	4.89	8154
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	55	26	796	26	597	67	7.3	16593
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITI1 PE=1 SV=3	37	23	389	23	911	101.3	6.79	10220
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	36	21	1370	21	500	55.1	6.55	29297
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	55	24	840	24	453	51.5	5.62	17286
Q9Y6R7	IgGf-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3	15	28	218	28	5405	571.6	5.34	6107
P04220	Ig mu heavy chain disease protein OS=Homo sapiens PE=1 SV=1	59	21	1483	2	391	43	5.24	35187
P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	37	19	238	19	676	75.1	5.67	7630
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	18	36	228	36	2813	309.1	5.48	5156
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	33	27	202	27	1231	139	6.61	4244
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITI4 PE=1 SV=4	24	16	206	16	930	103.3	6.98	5951
P00736	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	38	18	153	18	705	80.1	6.21	3810
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	52	12	335	5	353	37.6	6.51	7316
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	63	21	454	21	267	30.8	5.76	10273

P09871	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	36	17	142	17	688	76.6	4.96	4743
P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	48	18	271	9	406	45.2	6.58	5715
Q08380	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	30	13	161	13	585	65.3	5.27	4588
Q06033	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITIH3 PE=1 SV=2	27	18	179	18	890	99.8	5.74	5050
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	55	13	273	7	330	36.1	8.19	5604
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	47	17	217	17	418	46.7	5.59	4401
P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2 PE=1 SV=3	59	11	191	4	340	36.5	6.1	4542
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	80	7	663	7	106	11.6	5.87	22023
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	17	19	99	19	2067	217.1	6.39	2660
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	39	16	165	16	559	63.1	5.59	3675
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	47	14	198	5	348	39	7.09	3516
P08519	Apolipoprotein(a) OS=Homo sapiens GN=LPA PE=1 SV=1	31	17	146	17	4548	501	5.88	3189
P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	69	9	112	9	147	15.9	5.76	4050
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	49	14	256	14	347	38.1	5.47	6159
P01008	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	37	13	125	13	464	52.6	6.71	3283
Q8WX17	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	7	22	164	22	22152	2351.2	6	1744
P00747	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	30	19	127	19	810	90.5	7.24	2109
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	39	13	122	13	423	47.6	5.52	2829
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	25	16	213	16	946	106.4	6.86	3014
P02774	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	53	16	74	16	474	52.9	5.54	2083
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	37	7	123	7	240	25.8	6.61	4212
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1 SV=3	6	15	122	15	3396	372.6	4.51	3291
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	24	7	111	7	352	39	6.25	2962
P01597	Ig kappa chain V-I region DEE OS=Homo sapiens PE=1 SV=1	17	4	98	3	108	11.7	9.36	3667
P22105	Tenascin-X OS=Homo sapiens GN=TNXB PE=1 SV=3	8	19	82	19	4289	464	5.34	1753
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	12	14	79	14	1786	197.9	4.94	2231
P01042	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	22	11	102	11	644	71.9	6.81	2876

P12111	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=5	7	16	71	16	3177	343.5	6.68	1371
P11226	Mannose-binding protein C OS=Homo sapiens GN=MBL2 PE=1 SV=2	45	11	117	11	248	26.1	5.49	3316
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3	12	13	85	13	1609	177.5	5.12	2273
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4	13	16	54	16	1823	202.4	6.28	934
P13671	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	13	9	64	9	934	104.7	6.76	1655
Q4LDE5	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 OS=Homo sapiens GN=SVEP1 PE=1 SV=3	7	13	62	13	3571	389.9	5.5	1539
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	75	5	413	1	106	11.3	7.24	10400
P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1 SV=4	51	6	150	6	159	18.1	5.24	6193
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	27	11	88	11	703	77.2	5.22	1860
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	36	5	358	2	214	23	8.84	7383
Q99715	Collagen alpha-1(XII) chain OS=Homo sapiens GN=COL12A1 PE=1 SV=2	6	14	63	14	3063	332.9	5.53	941
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	8	8	60	8	1404	151	9.5	1413
Q15063	Periostin OS=Homo sapiens GN=POSTN PE=1 SV=2	19	9	57	9	836	93.3	7.53	1379
Q15582	Transforming growth factor-beta-induced protein ig-h3 OS=Homo sapiens GN=TGFB1 PE=1 SV=1	26	11	51	11	683	74.6	7.71	1024
P50895	Basal cell adhesion molecule OS=Homo sapiens GN=BCAM PE=1 SV=2	22	10	48	10	628	67.4	5.81	1463
P48740	Mannan-binding lectin serine protease 1 OS=Homo sapiens GN=MASP1 PE=1 SV=3	21	11	66	11	699	79.2	5.49	1488
O95445	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1 SV=2	50	6	40	6	188	21.2	6.01	710
P10909	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	19	6	58	6	449	52.5	6.27	1648
P10643	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	24	12	36	12	843	93.5	6.48	504
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	32	7	41	7	375	41.7	5.48	887
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2	37	3	62	3	114	12.6	7.93	3114
P20851	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	40	7	56	7	252	28.3	5.14	725
P01019	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	19	7	58	7	485	53.1	6.32	1620

P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1 SV=1	31	3	63	2	108	11.8	8.91	4022
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	28	9	126	1	377	41.3	7.9	2099
P04206	Ig kappa chain V-III region GOL OS=Homo sapiens PE=1 SV=1	55	4	60	2	109	11.8	9.25	3085
P01609	Ig kappa chain V-I region Scw OS=Homo sapiens PE=1 SV=1	28	2	56	1	108	11.8	6	3745
Q14520	Hyaluronan-binding protein 2 OS=Homo sapiens GN=HABP2 PE=1 SV=1	13	6	50	6	560	62.6	6.54	1379
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	28	8	93	3	326	35.9	7.59	1414
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1 SV=1	24	2	65	1	108	11.8	5.36	3799
P05546	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	20	8	76	8	499	57	6.9	1197
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1	33	2	46	1	113	12.3	6	1810
P02671	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2	9	8	143	8	866	94.9	6.01	2026
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1	60	4	74	1	109	11.8	8.48	3157
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	48	2	108	11.8	8.44	2610
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1 SV=1	31	4	64	3	108	11.6	9.25	1777
Q14112	Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3	7	6	35	6	1375	151.2	5.29	1034
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	30	7	81	7	223	25.4	6.54	1840
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1 SV=1	36	5	121	4	115	12.4	9.25	3498
P08697	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	21	5	26	5	491	54.5	6.29	514
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	17	6	44	6	478	54.3	5.8	1312
P01611	Ig kappa chain V-I region Wes OS=Homo sapiens PE=1 SV=1	31	2	33	2	108	11.6	7.28	1416
P18136	Ig kappa chain V-III region HIC OS=Homo sapiens PE=2 SV=2	57	4	52	1	129	14.1	6.6	2108
P12109	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3	9	6	36	6	1028	108.5	5.43	879
P24043	Laminin subunit alpha-2 OS=Homo sapiens GN=LAMA2 PE=1 SV=4	4	9	16	9	3122	343.7	6.4	350
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	12	5	52	5	525	59.5	7.5	1619
P01833	Polymeric immunoglobulin receptor OS=Homo sapiens GN=PIGR PE=1 SV=4	14	7	26	7	764	83.2	5.74	665
P22792	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	20	7	31	7	545	60.5	5.99	683
A0M8Q6	Ig lambda-7 chain C region OS=Homo sapiens GN=IGLC7 PE=1 SV=2	51	3	167	1	106	11.3	8.28	3339

P07357	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	13	6	33	6	584	65.1	6.47	521
P02749	Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3	32	6	34	6	345	38.3	7.97	323
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	21	4	33	4	245	25.8	8.41	1055
P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1	34	3	128	2	120	13.2	6.57	4180
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	15	5	42	5	493	54.7	7.01	956
P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	34	4	23	4	202	22.3	8.31	842
P01613	Ig kappa chain V-I region Ni OS=Homo sapiens PE=1 SV=1	30	2	34	2	112	12.2	5.36	1510
P01768	Ig heavy chain V-III region CAM OS=Homo sapiens PE=1 SV=1	34	5	117	4	122	13.7	9.63	3052
P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	12	6	29	6	591	67	8.13	868
O00187	Mannan-binding lectin serine protease 2 OS=Homo sapiens GN=MASP2 PE=1 SV=4	8	4	29	4	686	75.7	5.63	1080
P01763	Ig heavy chain V-III region WEA OS=Homo sapiens PE=1 SV=1	36	3	107	2	114	12.2	8.5	2858
P22891	Vitamin K-dependent protein Z OS=Homo sapiens GN=PROZ PE=1 SV=2	29	8	23	8	400	44.7	5.97	182
P35542	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2	27	3	75	3	130	14.7	9.07	830
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	13	5	29	5	481	53.4	6.7	638
P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1	66	5	30	5	100	11.2	6.62	326
P05090	Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	26	4	44	4	189	21.3	5.15	829
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1 SV=1	27	2	41	1	116	12.4	9.73	785
P01616	Ig kappa chain V-II region MIL OS=Homo sapiens PE=1 SV=1	33	2	30	1	112	12	9.29	1006
Q7Z7G0	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1 SV=1	5	4	29	4	1075	118.6	9.44	808
Q12805	EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 PE=1 SV=2	15	4	10	4	493	54.6	5.07	163
P01765	Ig heavy chain V-III region TIL OS=Homo sapiens PE=1 SV=1	26	2	84	1	115	12.3	9.13	1457
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	47	3	21	2	100	10.7	6.52	449
Q92496	Complement factor H-related protein 4 OS=Homo sapiens GN=CFHR4 PE=1 SV=3	9	4	23	4	578	65.3	4.94	345

P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4 SV=1	17	2	32	2	146	16.2	8.28	1066
P01743	Ig heavy chain V-I region HG3 OS=Homo sapiens PE=4 SV=1	22	2	42	2	117	12.9	8.92	1212
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	8	4	18	4	698	77	7.12	310
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	21	4	37	4	224	25	5.63	743
P01776	Ig heavy chain V-III region WAS OS=Homo sapiens PE=1 SV=1	24	2	36	1	117	13.1	7.99	543
P01602	Ig kappa chain V-I region HK102 (Fragment) OS=Homo sapiens GN=IGKV1-5 PE=4 SV=1	29	2	14	2	117	12.8	6.51	665
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	52	3	19	3	115	12.6	4.96	437
P04430	Ig kappa chain V-I region BAN OS=Homo sapiens PE=1 SV=1	17	1	15	1	108	11.8	8.44	857
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1 SV=1	17	1	15	1	108	11.4	6.52	716
Q9BXR6	Complement factor H-related protein 5 OS=Homo sapiens GN=CFHR5 PE=1 SV=1	8	3	12	3	569	64.4	7.06	330
P35442	Thrombospondin-2 OS=Homo sapiens GN=THBS2 PE=1 SV=2	5	3	11	3	1172	129.9	4.83	320
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	22	2	16	2	111	11.9	5.08	731
Q15485	Ficolin-2 OS=Homo sapiens GN=FCN2 PE=1 SV=2	12	2	13	2	313	34	6.77	373
P0DJ18	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	28	2	16	2	122	13.5	6.79	785
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3	9	2	8	2	381	41.9	5.01	153
P01717	Ig lambda chain V-IV region Hil OS=Homo sapiens PE=1 SV=1	18	1	14	1	107	11.5	6.51	449
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	10	1	15	1	156	18	9.64	432
P27169	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	8	2	12	2	355	39.7	5.22	389
P01781	Ig heavy chain V-III region GAL OS=Homo sapiens PE=1 SV=1	19	2	22	2	116	12.7	8.48	396
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	19	3	20	3	253	26.7	8.63	638
P00734	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	6	2	8	2	622	70	5.9	277
P19652	Alpha-1-acid glycoprotein 2 OS=Homo sapiens GN=ORM2 PE=1 SV=2	13	2	10	1	201	23.6	5.11	296
P07333	Macrophage colony-stimulating factor 1 receptor OS=Homo sapiens GN=CSF1R PE=1 SV=2	2	1	9	1	972	107.9	6.37	175
P15169	Carboxypeptidase N catalytic chain OS=Homo sapiens GN=CPN1 PE=1 SV=1	6	2	11	2	458	52.3	7.34	185

P06727	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	7	2	10	2	396	45.4	5.38	429
Q01628	Interferon-induced transmembrane protein 3 OS=Homo sapiens GN=IFITM3 PE=1 SV=2	12	1	11	1	133	14.6	7.01	533
P03952	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	4	2	8	2	638	71.3	8.22	523
P06889	Ig lambda chain V-IV region MOL OS=Homo sapiens PE=1 SV=1	24	1	11	1	106	11.3	4.42	334
P02654	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	27	4	30	4	83	9.3	8.47	188
Q9Y5Y7	Lymphatic vessel endothelial hyaluronic acid receptor 1 OS=Homo sapiens GN=LYVE1 PE=1 SV=2	9	3	20	3	322	35.2	8.28	426
P01780	Ig heavy chain V-III region JON OS=Homo sapiens PE=1 SV=1	17	1	9	1	115	12.6	9.29	144
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1 SV=2	19	2	18	1	129	14.3	8.51	283
P06311	Ig kappa chain V-III region IARC/BL41 OS=Homo sapiens PE=1 SV=1	12	1	10	1	128	14.1	6.61	533
P01612	Ig kappa chain V-I region Mev OS=Homo sapiens PE=1 SV=1	17	1	7	1	109	11.9	6.57	314
Q15848	Adiponectin OS=Homo sapiens GN=ADIPOQ PE=1 SV=1	20	3	10	3	244	26.4	5.74	185
Q04756	Hepatocyte growth factor activator OS=Homo sapiens GN=HGFA PE=1 SV=1	4	2	6	2	655	70.6	7.24	174
P83593	Ig kappa chain V-IV region STH (Fragment) OS=Homo sapiens PE=1 SV=1	28	2	9	2	109	12.1	7.94	174
P14543	Nidogen-1 OS=Homo sapiens GN=NID1 PE=1 SV=3	2	2	7	2	1247	136.3	5.29	229
Q9HC84	Mucin-5B OS=Homo sapiens GN=MUC5B PE=1 SV=3	1	3	8	3	5762	596	6.64	101
P02649	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	5	1	7	1	317	36.1	5.73	340
P00740	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	4	2	9	2	461	51.7	5.47	191
P35052	Glypican-1 OS=Homo sapiens GN=GPC1 PE=1 SV=2	3	1	6	1	558	61.6	7.3	269
Q9UGM3	Deleted in malignant brain tumors 1 protein OS=Homo sapiens GN=DMBT1 PE=1 SV=2	7	1	5	1	2413	260.6	5.44	144
O60635	Tetraspanin-1 OS=Homo sapiens GN=TSPAN1 PE=1 SV=2	5	1	3	1	241	26.3	5.25	123
P01769	Ig heavy chain V-III region GA OS=Homo sapiens PE=1 SV=1	16	2	10	2	122	13.2	9.77	194
P01771	Ig heavy chain V-III region HIL OS=Homo sapiens PE=1 SV=1	9	1	9	1	121	13.6	9.36	189
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2	6	1	7	1	181	20.6	7.43	101
P01700	Ig lambda chain V-I region HA OS=Homo sapiens PE=1 SV=1	12	1	11	1	112	11.9	8.91	592
P81605	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	10	1	8	1	110	11.3	6.54	129
P61626	Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1	8	1	3	1	148	16.5	9.16	102
P06316	Ig lambda chain V-I region BL2 OS=Homo sapiens PE=2 SV=1	6	1	12	1	130	13.6	7.77	118

Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	39	127	777	127	4563	515.3	7.05	13336
P0COL5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	48	63	731	4	1744	192.6	7.27	14596
P0COL4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	47	61	711	2	1744	192.7	7.08	13892
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	33	48	423	48	2386	262.5	5.71	9129
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	40	47	289	47	1663	187	6.4	5169
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	38	25	613	25	911	101.3	6.79	13653
P12111	Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=1 SV=5	17	39	202	39	3177	343.5	6.68	4151
P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=4	13	36	202	36	4391	468.5	6.51	4148
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	56	21	1038	7	452	49.3	6.77	22425
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	35	36	196	31	1474	163.2	6.46	3308
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3	16	20	133	20	1609	177.5	5.12	2783
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	36	17	240	17	597	67	7.3	3655
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	41	21	174	21	609	69.3	6.28	2866
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	56	17	380	17	267	30.8	5.76	8059
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	30	14	347	14	500	55.1	6.55	5481
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	16	23	124	23	2067	217.1	6.39	2459
P04220	Ig mu heavy chain disease protein OS=Homo sapiens PE=1 SV=1	57	15	599	1	391	43	5.24	10975
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	28	18	203	18	946	106.4	6.86	2954
O15230	Laminin subunit alpha-5 OS=Homo sapiens GN=LAMA5 PE=1 SV=8	9	22	114	22	3695	399.5	7.02	2339
Q8WXI7	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	7	25	135	25	22152	2351.2	6	1970

P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	18	24	133	24	1676	188.2	6.52	2002
P55268	Laminin subunit beta-2 OS=Homo sapiens GN=LAMB2 PE=1 SV=2	16	21	101	21	1798	195.9	6.52	1735
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	50	18	112	18	491	55.9	8.27	1614
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	80	6	266	6	106	11.6	5.87	6772
P07355	Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	40	13	89	13	339	38.6	7.75	2460
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	15	18	79	18	1786	197.9	4.94	1129
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	33	14	94	14	559	63.1	5.59	1808
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	41	11	82	11	453	51.5	5.62	1532
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	43	11	181	5	353	37.6	6.51	4116
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1 SV=3	4	12	106	12	3396	372.6	4.51	2771
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3	32	10	66	10	381	41.9	5.01	1369
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	10	13	58	13	2201	240.7	4.89	913
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	29	8	48	7	434	47.1	7.39	1164
P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	19	10	59	10	676	75.1	5.67	1368
P15311	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	25	13	87	6	586	69.4	6.27	1134
P50895	Basal cell adhesion molecule OS=Homo sapiens GN=BCAM PE=1 SV=2	27	12	63	12	628	67.4	5.81	1225
P26038	Moiesin OS=Homo sapiens GN=MSN PE=1 SV=3	23	12	90	5	577	67.8	6.4	1253
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	34	10	62	10	418	46.7	5.59	1039
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	34	9	111	4	330	36.1	8.19	1651
Q06033	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITI1H3 PE=1 SV=2	13	8	65	8	890	99.8	5.74	1377

P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	40	10	72	7	375	41.7	5.48	1110
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4	9	12	42	12	1823	202.4	6.28	582
P22735	Protein-glutamine gamma-glutamyltransferase K OS=Homo sapiens GN=TGM1 PE=1 SV=4	13	8	39	8	817	89.7	6.04	903
Q7Z7G0	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1 SV=1	14	10	46	10	1075	118.6	9.44	733
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	15	10	39	10	1065	122.1	5.72	851
P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2 PE=1 SV=3	23	7	98	1	340	36.5	6.1	2263
P01597	Ig kappa chain V-I region DEE OS=Homo sapiens PE=1 SV=1	17	2	58	1	108	11.7	9.36	887
P14543	Nidogen-1 OS=Homo sapiens GN=NID1 PE=1 SV=3	10	10	34	10	1247	136.3	5.29	842
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	7	7	45	7	1404	151	9.5	902
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	35	9	60	9	347	38.1	5.47	873
POCG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	75	5	136	2	106	11.3	7.24	2303
P01593	Ig kappa chain V-I region AG OS=Homo sapiens PE=1 SV=1	31	2	46	1	108	12	5.99	1100
P15924	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	4	9	31	9	2871	331.6	6.81	642
P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	8	10	44	5	1482	163.8	6.38	558
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2	37	3	45	3	114	12.6	7.93	1070
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	14	5	69	5	478	54.3	5.8	1323
P02649	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	30	7	36	7	317	36.1	5.73	613
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	36	5	105	2	214	23	8.84	1448
P12110	Collagen alpha-2(VI) chain OS=Homo sapiens GN=COL6A2 PE=1 SV=4	13	9	44	9	1019	108.5	6.21	512
P14618	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	20	8	40	8	531	57.9	7.84	875
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	19	7	40	7	481	53.4	6.7	875

P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1 SV=1	31	3	53	2	108	11.8	8.91	1169
P05164	Myeloperoxidase OS=Homo sapiens GN=MPO PE=1 SV=1	17	10	35	10	745	83.8	8.97	523
P13671	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	11	7	22	7	934	104.7	6.76	529
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	25	8	92	2	377	41.3	7.9	1367
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	22	6	30	3	348	39	7.09	770
P12109	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3	13	9	36	9	1028	108.5	5.43	473
P10909	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	18	5	42	5	449	52.5	6.27	1015
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	5	10	32	10	2813	309.1	5.48	334
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1	39	3	46	2	109	11.8	8.48	1674
Q8WUM4	Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1	8	6	30	6	868	96	6.52	639
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1 SV=1	24	2	47	1	108	11.8	5.36	983
P00734	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	14	7	18	7	622	70	5.9	375
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	27	7	71	2	326	35.9	7.59	1117
P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	14	6	26	3	406	45.2	6.58	709
P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	11	6	20	6	591	67	8.13	511
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	34	2	108	11.8	8.44	763
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	23	5	25	5	335	36	8.46	601
P05362	Intercellular adhesion molecule 1 OS=Homo sapiens GN=ICAM1 PE=1 SV=2	16	6	27	6	532	57.8	7.99	546
P02671	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2	5	4	30	4	866	94.9	6.01	542

P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1 SV=4	37	5	47	5	159	18.1	5.24	974
P10643	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	12	6	18	6	843	93.5	6.48	166
P05546	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	9	5	36	5	499	57	6.9	604
P27701	CD82 antigen OS=Homo sapiens GN=CD82 PE=1 SV=1	24	5	30	5	267	29.6	5.24	511
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	15	5	19	5	493	54.7	7.01	290
P15144	Aminopeptidase N OS=Homo sapiens GN=ANPEP PE=1 SV=4	9	6	29	6	967	109.5	5.48	523
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	18	6	32	6	423	47.6	5.52	389
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1 SV=1	17	2	22	1	108	11.6	9.25	632
P04083	Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	13	4	19	4	346	38.7	7.02	656
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	21	4	30	4	245	25.8	8.41	523
P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1	25	2	39	1	120	13.2	6.57	738
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1 SV=1	36	3	31	2	115	12.4	9.25	585
P14923	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3	9	5	15	5	745	81.7	6.14	271
P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	48	4	16	4	147	15.9	5.76	358
P27105	Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=1 SV=3	16	3	21	3	288	31.7	7.88	474
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITI4 PE=1 SV=4	5	4	36	4	930	103.3	6.98	624
Q29963	HLA class I histocompatibility antigen, Cw-6 alpha chain OS=Homo sapiens GN=HLA-C PE=2 SV=2	20	5	14	2	366	40.9	6	178
P01861	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1	18	5	39	1	327	35.9	7.36	585
P11142	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	9	4	16	3	646	70.9	5.52	362

P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1	59	4	40	4	100	11.2	6.62	513
P01765	Ig heavy chain V-III region TIL OS=Homo sapiens PE=1 SV=1	26	2	28	0	115	12.3	9.13	657
P01891	HLA class I histocompatibility antigen, A-68 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=4	17	4	17	1	365	40.9	6.7	154
P11226	Mannose-binding protein C OS=Homo sapiens GN=MBL2 PE=1 SV=2	13	4	21	4	248	26.1	5.49	563
P0DJ18	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	25	3	11	2	122	13.5	6.79	390
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2	31	4	16	4	181	20.6	7.43	362
P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	31	4	17	4	202	22.3	8.31	230
P05023	Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1 PE=1 SV=1	5	4	19	4	1023	112.8	5.49	308
P62979	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	16	2	30	2	156	18	9.64	514
P16190	HLA class I histocompatibility antigen, A-33 alpha chain OS=Homo sapiens GN=HLA-A PE=1 SV=3	16	4	14	0	365	40.9	6.7	112
P04899	Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2 PE=1 SV=3	15	4	21	4	355	40.4	5.54	376
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1 SV=1	27	2	18	1	116	12.4	9.73	482
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	4	4	17	3	1231	139	6.61	251
P39059	Collagen alpha-1(XV) chain OS=Homo sapiens GN=COL15A1 PE=1 SV=2	4	5	21	5	1388	141.6	5	297
P09758	Tumor-associated calcium signal transducer 2 OS=Homo sapiens GN=TACSTD2 PE=1 SV=3	10	2	12	2	323	35.7	8.87	425
P60033	CD81 antigen OS=Homo sapiens GN=CD81 PE=1 SV=1	20	3	11	3	236	25.8	5.29	171
P01042	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	9	4	10	4	644	71.9	6.81	152
O95832	Claudin-1 OS=Homo sapiens GN=CLDN1 PE=1 SV=1	15	2	21	2	211	22.7	8.09	181
Q8NFJ5	Retinoic acid-induced protein 3 OS=Homo sapiens GN=GPRC5A PE=1 SV=2	9	2	12	2	357	40.2	8.15	436

Q14112	Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3	3	3	16	3	1375	151.2	5.29	348
P07900	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	5	3	13	3	732	84.6	5.02	208
P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4 SV=1	17	2	10	2	146	16.2	8.28	257
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	16	3	12	3	223	25.4	6.54	394
Q9NQ84	G-protein coupled receptor family C group 5 member C OS=Homo sapiens GN=GPRC5C PE=1 SV=2	7	3	15	3	441	48.2	8.43	406
P01764	Ig heavy chain V-III region VH26 OS=Homo sapiens PE=1 SV=1	19	2	22	1	117	12.6	8.28	302
P61224	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1	18	3	13	3	184	20.8	5.78	267
P62736	Actin, aortic smooth muscle OS=Homo sapiens GN=ACTA2 PE=1 SV=1	13	4	26	1	377	42	5.39	210
P68363	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	12	4	13	4	451	50.1	5.06	209
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	7	3	7	3	703	77.2	5.22	121
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1 SV=2	19	2	24	2	129	14.3	8.51	585
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	11	2	11	2	253	26.7	8.63	250
P68104	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1	7	3	18	3	462	50.1	9.01	479
Q9H4M9	EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2	7	3	11	3	534	60.6	6.83	117
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1	33	2	15	2	113	12.3	6	409
P21926	CD9 antigen OS=Homo sapiens GN=CD9 PE=1 SV=4	7	3	33	3	228	25.4	7.15	483
Q06830	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	15	3	19	3	199	22.1	8.13	331
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	9	3	9	3	320	35.9	5.05	157
O15551	Claudin-3 OS=Homo sapiens GN=CLDN3 PE=1 SV=1	13	2	21	2	220	23.3	8.05	388
P09104	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3	7	2	8	1	434	47.2	5.03	283
P07357	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	5	2	6	2	584	65.1	6.47	106

Q9P2B2	Prostaglandin F2 receptor negative regulator OS=Homo sapiens GN=PTGFRN PE=1 SV=2	4	3	12	3	879	98.5	6.61	174
P09871	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	5	3	13	3	688	76.6	4.96	187
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5	6	3	9	2	641	70	5.66	138
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	7	3	10	3	525	59.5	7.5	216
O60635	Tetraspanin-1 OS=Homo sapiens GN=TSPAN1 PE=1 SV=2	5	1	8	1	241	26.3	5.25	330
Q99102	Mucin-4 OS=Homo sapiens GN=MUC4 PE=1 SV=4	1	2	13	2	2169	231.4	6.27	304
Q96P63	Serpin B12 OS=Homo sapiens GN=SERPINB12 PE=1 SV=1	8	3	6	3	405	46.2	5.53	107
P20160	Azurocidin OS=Homo sapiens GN=AZU1 PE=1 SV=3	16	2	9	2	251	26.9	9.5	159
P02654	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	24	3	25	3	83	9.3	8.47	303
P63104	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	13	3	16	3	245	27.7	4.79	164
P62873	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3	10	2	7	2	340	37.4	6	196
Q02413	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	3	2	5	2	1049	113.7	5.03	129
P81605	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	13	2	13	2	110	11.3	6.54	329
P00338	L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2	8	3	19	3	332	36.7	8.27	211
P35542	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2	20	2	23	2	130	14.7	9.07	236
O95436	Sodium-dependent phosphate transport protein 2B OS=Homo sapiens GN=SLC34A2 PE=1 SV=3	6	2	7	2	690	75.7	8.15	154
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	12	2	11	2	240	25.8	6.61	257
P12273	Prolactin-inducible protein OS=Homo sapiens GN=PIP PE=1 SV=1	14	2	7	2	146	16.6	8.05	245
Q96IY4	Carboxypeptidase B2 OS=Homo sapiens GN=CPB2 PE=1 SV=2	5	2	8	2	423	48.4	7.71	232
P01714	Ig lambda chain V-III region SH OS=Homo sapiens PE=1 SV=1	17	1	14	1	108	11.4	6.52	223

P08134	Rho-related GTP-binding protein RhoC OS=Homo sapiens GN=RHOC PE=1 SV=1	10	2	5	2	193	22	6.58	122
Q9BXX0	EMILIN-2 OS=Homo sapiens GN=EMILIN2 PE=1 SV=3	2	2	11	2	1053	115.6	6.46	142
Q8N271	Prominin-2 OS=Homo sapiens GN=PROM2 PE=1 SV=1	3	2	10	2	834	91.8	6.15	157
O00299	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4	12	2	13	2	241	26.9	5.17	188
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	5	2	9	2	352	39	6.25	228
Q03591	Complement factor H-related protein 1 OS=Homo sapiens GN=CFHR1 PE=1 SV=2	7	2	5	1	330	37.6	7.39	156
Q15485	Ficolin-2 OS=Homo sapiens GN=FCN2 PE=1 SV=2	7	2	8	2	313	34	6.77	139
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2	13	2	7	2	191	21.2	6.55	105
P05090	Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	13	2	12	2	189	21.3	5.15	147
P62805	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2	17	2	10	2	103	11.4	11.36	152
P11166	Solute carrier family 2, facilitated glucose transporter member 1 OS=Homo sapiens GN=SLC2A1 PE=1 SV=2	4	2	14	2	492	54	8.72	170
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	26	2	11	2	115	12.6	4.96	226
P15153	Ras-related C3 botulinum toxin substrate 2 OS=Homo sapiens GN=RAC2 PE=1 SV=1	9	2	12	2	192	21.4	7.61	128
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	31	2	20	2	100	10.7	6.52	245
Q9NZH0	G-protein coupled receptor family C group 5 member B OS=Homo sapiens GN=GPRC5B PE=2 SV=2	3	1	5	1	403	44.8	8.24	235
P20851	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	13	2	8	2	252	28.3	5.14	148
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	9	2	13	2	224	25	5.63	176
P59190	Ras-related protein Rab-15 OS=Homo sapiens GN=RAB15 PE=1 SV=1	10	2	6	2	212	24.4	5.71	103
P01772	Ig heavy chain V-III region KOL OS=Homo sapiens PE=1 SV=1	13	1	4	1	126	13.7	5.87	236
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	5	1	8	1	165	18	7.81	315

Q15758	Neutral amino acid transporter B(0) OS=Homo sapiens GN=SLC1A5 PE=1 SV=2	2	1	6	1	541	56.6	5.48	177
P16422	Epithelial cell adhesion molecule OS=Homo sapiens GN=EPCAM PE=1 SV=2	4	1	4	1	314	34.9	7.46	117
P00740	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	2	1	8	1	461	51.7	5.47	259
P09211	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	5	1	9	1	210	23.3	5.64	264
P02655	Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1 SV=1	11	1	7	1	101	11.3	4.72	127
Q08722	Leukocyte surface antigen CD47 OS=Homo sapiens GN=CD47 PE=1 SV=1	2	1	9	1	323	35.2	7.21	130
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	7	1	5	1	111	11.9	5.08	112

Accession	Description	Coverage [%]	# Peptides	# PSMs	# Unique Peptides	# AAs	MW [kDa]	calc. pI	Score Mascot: Mascot
P04114	Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2	39	124	879	124	4563	515.3	7.05	17588
P0C0L5	Complement C4-B OS=Homo sapiens GN=C4B PE=1 SV=2	53	74	1705	4	1744	192.6	7.27	35591
P0C0L4	Complement C4-A OS=Homo sapiens GN=C4A PE=1 SV=2	52	72	1678	2	1744	192.7	7.08	34833
P02751	Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	38	58	939	58	2386	262.5	5.71	19111
P01024	Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	49	58	554	58	1663	187	6.4	12253
P01031	Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4	35	40	343	40	1676	188.2	6.52	6778
P01023	Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3	41	41	345	35	1474	163.2	6.46	6289
P98160	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2 PE=1 SV=4	12	34	189	34	4391	468.5	6.51	3891
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	68	26	368	26	491	55.9	8.27	6900
P19827	Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3	35	24	329	24	911	101.3	6.79	7828
P02768	Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	48	25	258	25	609	69.3	6.28	4604
P05155	Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=1 SV=2	34	18	1130	18	500	55.1	6.55	22405
P04003	C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2	55	22	416	22	597	67	7.3	7330
P01871	Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3	53	19	614	19	452	49.3	6.77	11252
O00468	Agrin OS=Homo sapiens GN=AGRN PE=1 SV=5	19	25	176	25	2067	217.1	6.39	3410
P00450	Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1	29	20	177	20	1065	122.1	5.72	3751
P24821	Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	18	23	128	23	2201	240.7	4.89	2640
P02679	Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3	53	18	239	18	453	51.5	5.62	5199
P02647	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	56	18	388	18	267	30.8	5.76	9819
P19823	Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2	24	15	194	15	946	106.4	6.86	2862
P11047	Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3	16	19	117	19	1609	177.5	5.12	2462
P00734	Prothrombin OS=Homo sapiens GN=F2 PE=1 SV=2	35	18	106	18	622	70	5.9	2331

P20742	Pregnancy zone protein OS=Homo sapiens GN=PZP PE=1 SV=4	16	18	123	12	1482	163.8	6.38	1880
P02748	Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	34	15	108	15	559	63.1	5.59	2128
P01876	Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	52	12	292	5	353	37.6	6.51	6620
P07942	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	13	16	85	16	1786	197.9	4.94	1400
P07225	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	20	12	97	12	676	75.1	5.67	1933
P01009	Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	31	12	101	12	418	46.7	5.59	1980
P08603	Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4	20	16	80	15	1231	139	6.61	1173
P01834	Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1	82	7	304	7	106	11.6	5.87	6626
P00738	Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1	38	11	82	7	406	45.2	6.58	1690
P10909	Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	28	10	117	10	449	52.5	6.27	2388
P01857	Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	47	10	130	5	330	36.1	8.19	1909
P02671	Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2	10	9	96	9	866	94.9	6.01	1620
P02766	Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1	69	10	60	10	147	15.9	5.76	1770
P01877	Ig alpha-2 chain C region OS=Homo sapiens GN=IGHA2 PE=1 SV=3	33	8	174	1	340	36.5	6.1	3905
P13611	Versican core protein OS=Homo sapiens GN=VCAN PE=1 SV=3	3	9	71	9	3396	372.6	4.51	1883
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Homo sapiens GN=ITIH4 PE=1 SV=4	12	8	63	8	930	103.3	6.98	1409
P00739	Haptoglobin-related protein OS=Homo sapiens GN=HPR PE=1 SV=2	36	8	60	4	348	39	7.09	1332
P09871	Complement C1s subcomponent OS=Homo sapiens GN=C1S PE=1 SV=1	16	7	53	7	688	76.6	4.96	1424
P15924	Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	5	10	38	10	2871	331.6	6.81	734

P50895	Basal cell adhesion molecule OS=Homo sapiens GN=BCAM PE=1 SV=2	25	11	38	11	628	67.4	5.81	744
O43866	CD5 antigen-like OS=Homo sapiens GN=CD5L PE=1 SV=1	40	9	60	9	347	38.1	5.47	781
P01042	Kininogen-1 OS=Homo sapiens GN=KNG1 PE=1 SV=2	17	9	48	9	644	71.9	6.81	885
P00736	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2	19	9	48	9	705	80.1	6.21	727
O15230	Laminin subunit alpha-5 OS=Homo sapiens GN=LAMA5 PE=1 SV=8	4	9	37	9	3695	399.5	7.02	664
P04004	Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	14	5	132	5	478	54.3	5.8	2964
P0CG05	Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1	75	5	122	2	106	11.3	7.24	1652
P55058	Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=1 SV=1	16	6	46	6	493	54.7	7.01	841
P01859	Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2	32	7	65	2	326	35.9	7.59	1229
Q16363	Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4	7	9	25	9	1823	202.4	6.28	357
P27169	Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=1 SV=3	35	7	44	7	355	39.7	5.22	684
P55268	Laminin subunit beta-2 OS=Homo sapiens GN=LAMB2 PE=1 SV=2	8	9	30	9	1798	195.9	6.52	411
P01011	Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	20	7	37	7	423	47.6	5.52	641
Q06033	Inter-alpha-trypsin inhibitor heavy chain H3 OS=Homo sapiens GN=ITIH3 PE=1 SV=2	11	7	42	7	890	99.8	5.74	951
P14618	Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	20	8	30	8	531	57.9	7.84	541
Q92954	Proteoglycan 4 OS=Homo sapiens GN=PRG4 PE=1 SV=2	5	5	32	5	1404	151	9.5	859
P01860	Ig gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	23	7	90	2	377	41.3	7.9	1600
B9A064	Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	36	5	103	2	214	23	8.84	1111
P26022	Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3	20	6	39	6	381	41.9	5.01	1140

P20851	C4b-binding protein beta chain OS=Homo sapiens GN=C4BPB PE=1 SV=1	35	6	36	6	252	28.3	5.14	494
P01593	Ig kappa chain V-I region AG OS=Homo sapiens PE=1 SV=1	31	2	32	1	108	12	5.99	852
P01610	Ig kappa chain V-I region WEA OS=Homo sapiens PE=1 SV=1	31	3	34	2	108	11.8	8.91	825
P01625	Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2	37	3	32	3	114	12.6	7.93	783
P01008	Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1	18	6	32	6	464	52.6	6.71	547
P01591	Immunoglobulin J chain OS=Homo sapiens GN=IGJ PE=1 SV=4	51	5	42	5	159	18.1	5.24	895
P11226	Mannose-binding protein C OS=Homo sapiens GN=MBL2 PE=1 SV=2	26	6	30	6	248	26.1	5.49	705
P01019	Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	14	4	18	4	485	53.1	6.32	621
P04196	Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1	12	5	29	5	525	59.5	7.5	558
P01608	Ig kappa chain V-I region Roy OS=Homo sapiens PE=1 SV=1	24	2	27	1	108	11.8	5.36	606
Q02413	Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	8	5	17	5	1049	113.7	5.03	306
P02649	Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1	24	5	29	5	317	36.1	5.73	786
P07358	Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=1 SV=3	10	5	16	5	591	67	8.13	375
P51884	Lumican OS=Homo sapiens GN=LUM PE=1 SV=2	18	5	21	5	338	38.4	6.61	239
P04275	von Willebrand factor OS=Homo sapiens GN=VWF PE=1 SV=4	3	6	25	6	2813	309.1	5.48	320
P01861	Ig gamma-4 chain C region OS=Homo sapiens GN=IGHG4 PE=1 SV=1	16	4	39	1	327	35.9	7.36	767
P00747	Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2	10	5	22	5	810	90.5	7.24	257
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	11	5	23	1	724	83.2	5.03	521
P01619	Ig kappa chain V-III region B6 OS=Homo sapiens PE=1 SV=1	31	3	21	2	108	11.6	9.25	557
Q4LDE5	Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 OS=Homo sapiens GN=SVEP1 PE=1 SV=3	2	6	14	6	3571	389.9	5.5	163
P02652	Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1	59	4	40	4	100	11.2	6.62	453

P07900	Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	10	5	23	1	732	84.6	5.02	521
P03952	Plasma kallikrein OS=Homo sapiens GN=KLKB1 PE=1 SV=1	12	5	18	5	638	71.3	8.22	421
P60709	Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	19	4	12	4	375	41.7	5.48	182
P01620	Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1	39	3	32	2	109	11.8	8.48	1338
P23142	Fibulin-1 OS=Homo sapiens GN=FBLN1 PE=1 SV=4	9	4	23	4	703	77.2	5.22	251
P02743	Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2	17	4	26	4	223	25.4	6.54	761
P0DJ18	Serum amyloid A-1 protein OS=Homo sapiens GN=SAA1 PE=1 SV=1	41	4	20	3	122	13.5	6.79	469
P01598	Ig kappa chain V-I region EU OS=Homo sapiens PE=1 SV=1	27	2	18	2	108	11.8	8.44	535
Q9Y6R7	IgGfc-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3	2	4	23	4	5405	571.6	5.34	555
P14543	Nidogen-1 OS=Homo sapiens GN=NID1 PE=1 SV=3	4	4	12	4	1247	136.3	5.29	362
P02741	C-reactive protein OS=Homo sapiens GN=CRP PE=1 SV=1	14	3	45	3	224	25	5.63	1060
P07360	Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3	34	4	20	4	202	22.3	8.31	288
P13671	Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3	6	4	19	4	934	104.7	6.76	470
P01767	Ig heavy chain V-III region BUT OS=Homo sapiens PE=1 SV=1	36	3	40	2	115	12.4	9.25	938
P10643	Complement component C7 OS=Homo sapiens GN=C7 PE=1 SV=2	8	4	14	4	843	93.5	6.48	256
P05546	Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	7	4	31	4	499	57	6.9	591
Q7Z7G0	Target of Nesh-SH3 OS=Homo sapiens GN=ABI3BP PE=1 SV=1	4	4	13	4	1075	118.6	9.44	112
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1	14	4	27	4	352	39	6.25	419
Q8WX17	Mucin-16 OS=Homo sapiens GN=MUC16 PE=1 SV=2	3	4	11	4	22152	2351.2	6	144
P35542	Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2	27	3	44	3	130	14.7	9.07	528
P49788	Retinoic acid receptor responder protein 1 OS=Homo sapiens GN=RARRES1 PE=1 SV=2	14	3	18	3	294	33.3	8.51	464

P01766	Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1	25	2	38	1	120	13.2	6.57	913
P14923	Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3	6	3	8	3	745	81.7	6.14	242
P08294	Extracellular superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD3 PE=1 SV=2	15	3	19	3	240	25.8	6.61	392
P01764	Ig heavy chain V-III region VH26 OS=Homo sapiens PE=1 SV=1	19	2	13	1	117	12.6	8.28	207
P01765	Ig heavy chain V-III region TIL OS=Homo sapiens PE=1 SV=1	26	2	16	0	115	12.3	9.13	508
P05090	Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	16	3	19	3	189	21.3	5.15	346
P02774	Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1	9	2	12	2	474	52.9	5.54	159
O95445	Apolipoprotein M OS=Homo sapiens GN=APOM PE=1 SV=2	18	3	12	3	188	21.2	6.01	187
P01779	Ig heavy chain V-III region TUR OS=Homo sapiens PE=1 SV=1	27	2	17	1	116	12.4	9.73	474
P22792	Carboxypeptidase N subunit 2 OS=Homo sapiens GN=CPN2 PE=1 SV=3	6	3	10	3	545	60.5	5.99	182
P02747	Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3	16	3	19	3	245	25.8	8.41	348
Q08380	Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	5	2	14	2	585	65.3	5.27	286
P02790	Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2	12	3	12	3	462	51.6	7.02	145
P01621	Ig kappa chain V-III region NG9 (Fragment) OS=Homo sapiens PE=1 SV=1	47	3	16	3	100	10.7	6.52	155
Q9HCB6	Spondin-1 OS=Homo sapiens GN=SPON1 PE=1 SV=2	5	3	14	3	807	90.9	6.11	112
P39059	Collagen alpha-1(XV) chain OS=Homo sapiens GN=COL15A1 PE=1 SV=2	3	3	17	3	1388	141.6	5	165
P05164	Myeloperoxidase OS=Homo sapiens GN=MPO PE=1 SV=1	6	3	8	3	745	83.8	8.97	109
P02746	Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=1 SV=3	11	2	11	2	253	26.7	8.63	363
P80748	Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	22	2	15	2	111	11.9	5.08	765
P01617	Ig kappa chain V-II region TEW OS=Homo sapiens PE=1 SV=1	33	2	14	2	113	12.3	6	462

Q96IY4	Carboxypeptidase B2 OS=Homo sapiens GN=CPB2 PE=1 SV=2	5	2	10	2	423	48.4	7.71	288
P00740	Coagulation factor IX OS=Homo sapiens GN=F9 PE=1 SV=2	4	2	14	2	461	51.7	5.47	327
P07357	Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2	5	2	8	2	584	65.1	6.47	135
P06331	Ig heavy chain V-II region ARH-77 OS=Homo sapiens PE=4 SV=1	11	1	15	1	146	16.2	8.28	476
P81605	Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	13	2	14	2	110	11.3	6.54	430
P01781	Ig heavy chain V-III region GAL OS=Homo sapiens PE=1 SV=1	17	2	17	2	116	12.7	8.48	231
P06727	Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3	7	2	7	2	396	45.4	5.38	122
P02787	Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	4	2	7	2	698	77	7.12	118
Q96P63	Serpin B12 OS=Homo sapiens GN=SERPINB12 PE=1 SV=1	6	2	6	2	405	46.2	5.53	113
P04207	Ig kappa chain V-III region CLL OS=Homo sapiens PE=1 SV=2	19	2	13	2	129	14.3	8.51	168
P02654	Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1	13	2	18	2	83	9.3	8.47	262
P18428	Lipopolysaccharide-binding protein OS=Homo sapiens GN=LBP PE=1 SV=3	6	2	9	2	481	53.4	6.7	230
PODJI9	Serum amyloid A-2 protein OS=Homo sapiens GN=SAA2 PE=1 SV=1	25	2	6	1	122	13.5	9.14	109
Q14112	Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3	2	2	7	2	1375	151.2	5.29	134
P04406	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	8	2	7	2	335	36	8.46	268
Q03591	Complement factor H-related protein 1 OS=Homo sapiens GN=CFHR1 PE=1 SV=2	7	2	10	1	330	37.6	7.39	317
P01880	Ig delta chain C region OS=Homo sapiens GN=IGHD PE=1 SV=2	7	2	7	2	384	42.2	7.93	216
P04433	Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1	26	2	13	2	115	12.6	4.96	240
P08697	Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3	6	2	9	2	491	54.5	6.29	104
P01717	Ig lambda chain V-IV region Hil OS=Homo sapiens PE=1 SV=1	18	1	7	1	107	11.5	6.51	118

P39060	Collagen alpha-1(XVIII) chain OS=Homo sapiens GN=COL18A1 PE=1 SV=5	2	1	6	1	1754	178.1	6.01	252
P35030	Trypsin-3 OS=Homo sapiens GN=PRSS3 PE=1 SV=2	4	1	6	1	304	32.5	7.49	213
P35052	Glypican-1 OS=Homo sapiens GN=GPC1 PE=1 SV=2	3	1	5	1	558	61.6	7.3	233
P12273	Prolactin-inducible protein OS=Homo sapiens GN=PIP PE=1 SV=1	8	1	8	1	146	16.6	8.05	302
Q9NPY3	Complement component C1q receptor OS=Homo sapiens GN=CD93 PE=1 SV=3	2	1	9	1	652	68.5	5.44	229
P01700	Ig lambda chain V-l region HA OS=Homo sapiens PE=1 SV=1	12	1	9	1	112	11.9	8.91	372
Q9UGM3	Deleted in malignant brain tumors 1 protein OS=Homo sapiens GN=DMBT1 PE=1 SV=2	7	1	5	1	2413	260.6	5.44	113
P09211	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	5	1	6	1	210	23.3	5.64	129