Fraction	Count	Protein	Reference	Peptide	Count	MaxOfXC	MaxOfDe	First(	Accessior
90									
	12								
		Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alp							
			CO3_HUMAN	K.GYTQQLAFR.Q	1	2.6592	0.2376		P01024
			CO3_HUMAN	R.SNLDEDIIAEENIVSR.S	2	4.5692	0.5706		P01024
			CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	3.9318	0.6333		P01024
			CO3_HUMAN	K.VLLDGVQNPR.A	2	3.4394	0.5621		P01024
			CO3_HUMAN	K.SGSDEVQVGQQR.T	1	3.8062	0.3043		P01024
			CO3_HUMAN	K.SDDKVTLEER.L	1	2.994	0.2781		P01024
			CO3_HUMAN	K.KQELSEAEQATR.T	2	3.7276	0.4363		P01024
			CO3_HUMAN	K.AGDFLEANYMNLQR.S	1	3.9738	0.3282		P01024
			CO3_HUMAN	K.QELSEAEQATR.T	1	2.8849	0.4095		P01024
	8								
		Complement component C9 precursor [Contains: Complement component C9a; Compl							
			CO9_HUMAN	R.VVEESELAR.T	2	2.8556	0.4705		P02748
			CO9_HUMAN	R.DRVVEESELAR.T	1	3.7524	0.4644		P02748
			CO9_HUMAN	R.DGNTLTYYR.R	1	2.5406	0.283		P02748
			CO9_HUMAN	R.AIEDYINEFSVR.K	2	4.1252	0.5592		P02748
			CO9_HUMAN	K.TSNFNAAISLK.F	2	3.3066	0.4952		P02748
	7								
		Serum albumin precursor - Homo sapiens (Human)							
			ALBU_HUMAN	K.FQNALLVR.Y	1	2.602	0.2615	2	P02768
			ALBU_HUMAN	K.AVMDDFAAFVEK.C	1	3.3546	0.5289	2	P02768
			ALBU_HUMAN	K.LVNEVTEFAK.T	1	3.2914	0.5163	2	P02768
			ALBU_HUMAN	K.KLVAASQAALGL.-	1	2.7916	0.4564	1	P02768
			ALBU_HUMAN	K.VPQVSTPTLVEVSR.N	1	3.4955	0.522	2	P02768
			ALBU_HUMAN	K.LVAASQAALGL.-	1	2.8618	0.6146	1	P02768
			ALBU_HUMAN	K.KQTALVELVK.H	1	2.4147	0.3919	2	P02768
	6								
		Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H							
			APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	2	4.2535	0.5512		P02647
			APOA1_HUMAN	K.DLATVYVDVLK.D	1	2.9673	0.5165		P02647
			APOA1_HUMAN	K.WQEEM*ELYR.Q	1	3.2165	0.4247		P02647
			APOA1_HUMAN	R.DYVSQFEGSALGK.Q	1	4.0101	0.5763		P02647
			APOA1_HUMAN	K.ATEHLSTLSEK.A	1	2.7592	0.4283		P02647
	3								
		Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S							
			SAMP_HUMAN	R.VGEYSLYIGR.H	1	2.9871	0.5439		P02743
			SAMP_HUMAN	R.QGYFVEAQPK.I	1	2.6943	0.4268		P02743
			SAMP_HUMAN	R.AYSLFSYNTQGR.D	1	2.8854	0.4212		P02743
	2								
		Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)							
			A2MG_HUMAN	K.AIGYLNTGYQR.Q	1	2.9575	0.3723		P01023
			A2MG_HUMAN	R.VGFYESDVMGR.G	1	3.2578	0.4841		P01023
		Ig alpha-2 chain C region - Homo sapiens (Human)							
			IGHA2_HUMAN	R.WLQGSQELPR.E	1	3.1245	0.3647	23	P01877
			IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	1	3.5351	0.508	21	P01877
		IGHA1 protein - Homo sapiens (Human)							

Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	1	4.5705	0.4937	12	Q8N5K4
Q8N5K4_HUMAN	K.TPLTATLSK.S	1	2.5677	0.4692	11	Q8N5K4

Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-

ITIH4_HUMAN	R.GPDVLTATVSGK.L	1	3.2619	0.4863	1	Q14624
ITIH4_HUMAN	K.GSEMVVAGK.L	1	2.4321	0.3519	1	Q14624

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Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr

FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.7801	0.5268		P02765

Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -

APOA2_HUMAN	K.SPELQAEAK.S	1	2.7774	0.3417		P02652

Ciliary dynein heavy chain 5 (Axonemal beta dynein heavy chain 5) (HL1) - Homo sapie

DYH5_HUMAN	K.RTYSGVSQDLLDVSSGSQWK	1	3.0295	0.1935		Q8TE73

Haptoglobin-related protein precursor - Homo sapiens (Human)

HPTR_HUMAN	R.ILGGHLDAK.G	1	2.4098	0.1878	3	P00739

Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	1	4.3425	0.5963	20	A0A5E5

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.1102	0.2857		P01591

Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	R.PPGFSPFR.S	1	2.4683	0.3416	1	P01042

Myosin-reactive immunoglobulin heavy chain variable region (Fragment) - Homo sapier

Q9UL94_HUMAN	-.EVQLVESGAEVK.K	1	2.9892	0.3708	2	Q9UL94

Postreplication repair protein RAD18 (hRAD18) (hHR18) (RING finger protein 73) - Hor

RAD18_HUMAN	K.EASPAAK.T	1	2.2602	0.1046		Q9NS91

Transthyretin precursor (Prealbumin) (TBPA) (TTR) (ATTR) - Homo sapiens (Human)

TTHY_HUMAN	K.AADDTWEPFASGK.T	1	4.8619	0.6225		P02766

Tribbles homolog 3 (TRB-3) (Neuronal cell death-inducible putative kinase) (p65-interac

TRIB3_HUMAN	K.VYPVQEALAVLEPYAR.L	1	2.6096	0.0819		Q96RU7

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Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alp

CO3_HUMAN	K.RIPIEDGSGEVVLSRK.V	1	3.8845	0.343		P01024
CO3_HUMAN	K.TGLQEVEVK.A	1	3.5366	0.4136		P01024
CO3_HUMAN	K.SLYVSATVILHSGSDMVQAER	1	5.493	0.6292		P01024
CO3_HUMAN	K.SGSDEVQVGQQR.T	2	4.0746	0.3779		P01024
CO3_HUMAN	K.SGQSEDRQPVPGQQMTLK.I	1	4.7844	0.5098		P01024
CO3_HUMAN	K.SDDKVTLEERLDK.A	2	3.5921	0.2749		P01024
CO3_HUMAN	K.GYTQQLAFRQPSSAFAAFVK.	1	4.2829	0.6315		P01024
CO3_HUMAN	K.YFKPGMPFDLMVFVTNPDGS	1	3.7322	0.3458		P01024
CO3_HUMAN	K.SDDKVTLEER.L	1	2.8225	0.2462		P01024
CO3_HUMAN	K.VLLDGVQNPR.A	1	3.6992	0.5598		P01024
CO3_HUMAN	K.RIPIEDGSGEVVLSR.K	2	4.1091	0.4762		P01024
CO3_HUMAN	K.QELSEAEQATR.T	1	3.3061	0.4311		P01024
CO3_HUMAN	K.LMNIFLK.D	1	2.4936	0.1912		P01024
CO3_HUMAN	K.KVEGTAFVIFGIQDGEQR.I	1	4.2464	0.5103		P01024
CO3_HUMAN	K.KGYTQQLAFR.Q	1	2.9878	0.4496		P01024
CO3_HUMAN	R.AYYENSPQQVFSTEFEVK.E	1	2.8735	0.4886		P01024
CO3_HUMAN	K.GYTQQLAFR.Q	1	2.6313	0.1501		P01024
CO3_HUMAN	K.GQGTLSVVTMYHAK.A	1	3.7409	0.5828		P01024
CO3_HUMAN	K.EDIPPADLSDQVPDTESETR.I	1	4.3681	0.6092		P01024

CO3_HUMAN	K.AGDFLEANYM*NLQR.S	1	3.8626	0.3722	P01024
CO3_HUMAN	R.VVLVAVDKGVFVLNKK.N	1	2.4002	0.2343	P01024
CO3_HUMAN	K.KQELSEAEQATR.T	2	4.1268	0.3936	P01024
CO3_HUMAN	R.SEETKENEGFTVTAEGK.G	1	4.8514	0.5502	P01024
CO3_HUMAN	R.VPVAVQGEDTVQSLTQGDGV	1	4.6844	0.6958	P01024
CO3_HUMAN	R.TKKQELSEAEQATR.T	4	4.6184	0.5866	P01024
CO3_HUMAN	R.SYTVAIAGYALAQMGR.L	1	3.3546	0.5733	P01024
CO3_HUMAN	R.TVMVNIENPEGIPVK.Q	1	2.6338	0.361	P01024
CO3_HUMAN	R.SGIPIVTSPYQIHFTK.T	1	3.5273	0.5743	P01024
CO3_HUMAN	R.EGVQKEDIPPADLSDQVPDTE	6	5.5359	0.6864	P01024
CO3_HUMAN	R.RHQQTVTIPPK.S	1	3.5754	0.3989	P01024
CO3_HUMAN	R.QVREPGQDLVVLPLSITTDFIF	1	4.3441	0.5452	P01024
CO3_HUMAN	R.MDKVGKYPK.E	1	2.424	0.2547	P01024
CO3_HUMAN	R.HQQTVTIPPK.S	2	2.8683	0.4194	P01024
CO3_HUMAN	R.KVLLDGVQNPR.A	1	3.8239	0.3776	P01024
CO3_HUMAN	R.IPIEDGSGEVVLSRK.V	1	3.7483	0.3919	P01024
CO3_HUMAN	R.IPIEDGSGEVVLSR.K	6	4.3342	0.6088	P01024
CO3_HUMAN	R.ILLQGTPVAQMTEDAVDAERL	2	4.4566	0.6632	P01024
CO3_HUMAN	R.ILLQGTPVAQMTEDAVDAER.I	2	5.8418	0.6211	P01024
CO3_HUMAN	R.ILLQGTPVAQM*TEDAVDAER	1	2.8474	0.5221	P01024
CO3_HUMAN	R.ILLQGTPVAQM*TEDAVDAER	1	5.154	0.661	P01024
CO3_HUMAN	R.LVAYYTLIGASGQR.E	1	2.5677	0.4693	P01024
CO3_HUMAN	R.SNLDEDIIAEENIVSR.S	3	5.5038	0.5868	P01024

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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H

APOA1_HUMAN	K.VSFLSALEEYTKK.L	1	3.1183	0.4384	P02647
APOA1_HUMAN	R.AELQEGAR.Q	1	2.4038	0.1764	P02647
APOA1_HUMAN	R.AHVDALRTHLAPYSDELRQR.	1	4.5136	0.5032	P02647
APOA1_HUMAN	R.DYVSQFEGSALGK.Q	2	4.4857	0.6616	P02647
APOA1_HUMAN	R.EQLGPVTQEFWDNLEK.E	2	3.9443	0.5417	P02647
APOA1_HUMAN	R.LAARLEALKENGGAR.L	1	3.8259	0.3314	P02647
APOA1_HUMAN	R.VKDLATVYVDVLK.D	3	4.0019	0.4235	P02647
APOA1_HUMAN	K.VSFLSALEEYTK.K	2	4.3547	0.6234	P02647
APOA1_HUMAN	R.THLAPYSDELRQR.L	1	3.6811	0.4258	P02647
APOA1_HUMAN	K.DLATVYVDVLK.D	2	3.2638	0.4581	P02647
APOA1_HUMAN	K.ATEHLSTLSEK.A	2	3.7751	0.4809	P02647
APOA1_HUMAN	K.DLATVYVDVLKDSGRDYVSQI	1	4.6121	0.5144	P02647
APOA1_HUMAN	K.DSGRDYVSQFEGSALGK.Q	3	5.2861	0.5919	P02647
APOA1_HUMAN	K.VQPYLDDFQKKWQEEM*ELY	2	3.3599	0.6735	P02647
APOA1_HUMAN	K.KWQEEM*ELYR.Q	2	3.2763	0.4351	P02647
APOA1_HUMAN	K.LREQLGPVTQEFWDNLEK.E	2	5.3065	0.6059	P02647
APOA1_HUMAN	K.QLNLKLLDNWDSVTSTFSK.L	1	3.9994	0.6633	P02647
APOA1_HUMAN	K.VEPLRAELQEGAR.Q	2	2.7029	0.2483	P02647
APOA1_HUMAN	K.VQPYLDDFQKK.W	1	2.4652	0.1965	P02647
APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	6	4.3298	0.6109	P02647

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Apolipoprotein B-100 precursor (Apo B-100) [Contains: Apolipoprotein B-48 (Apo B-48)

APOB_HUMAN	R.TGISPLALIK.G	1	2.457	0.3216	P04114
APOB_HUMAN	K.LLSGGNTLHLVSTTK.T	1	2.5825	0.4951	P04114
APOB_HUMAN	K.LTISEQNIQR.A	1	3.1386	0.3844	P04114
APOB_HUMAN	K.M*TSNFPVDLSDYPK.S	1	4.0575	0.4488	P04114

APOB_HUMAN	K.SVSDGIAALDLNAVANK.I	2	4.7745	0.4534		P04114
APOB_HUMAN	R.ALVDTLKFVTQAEGAK.Q	1	4.2753	0.5563		P04114
APOB_HUMAN	R.VNDESTEGKTSYR.L	1	3.3399	0.5718		P04114
APOB_HUMAN	R.NLQNNAEWVYQGAIR.Q	1	3.707	0.505		P04114
APOB_HUMAN	K.LIVAM*SSWLQK.A	1	3.0196	0.364		P04114
APOB_HUMAN	R.ITENDIQIALDDAK.I	1	3.2275	0.3468		P04114
APOB_HUMAN	R.IGQDGISTSATTNLK.C	1	3.4779	0.5022		P04114
APOB_HUMAN	K.FSVPAGIVIPSFQALTAR.F	1	3.9445	0.4292		P04114
APOB_HUMAN	K.AVSM*PSFSILGSDVR.V	1	4.398	0.5742		P04114
APOB_HUMAN	K.ALVEQGFTVPEIK.T	1	3.5082	0.5635		P04114
APOB_HUMAN	R.VPSYTLILPSLELPVLHVPR.N	1	5.7891	0.6071		P04114
APOB_HUMAN	R.SEYQADYESLR.F	1	3.17	0.6075		P04114
APOB_HUMAN	K.IEGNLIFDPNNYLPK.E	1	3.4298	0.5119		P04114

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Serum albumin precursor - Homo sapiens (Human)

ALBU_HUMAN	K.VFDEFKPLVEEPQNLIK.Q	1	4.9386	0.4405	2	P02768
ALBU_HUMAN	K.KVPQVSTPTLVEVSR.N	2	3.6357	0.5665	2	P02768
ALBU_HUMAN	K.EQLKAVMDDFAAFVEK.C	1	3.9159	0.4812	2	P02768
ALBU_HUMAN	K.ATKEQLKAVMDDFAAFVEK.C	1	4.752	0.5466	2	P02768
ALBU_HUMAN	K.AEFAEVSKLVTDLTK.V	2	3.8622	0.474	1	P02768
ALBU_HUMAN	K.YLYEIAR.R	1	2.4492	0.2762	1	P02768
ALBU_HUMAN	K.YICENQDSISSK.L	1	4.5453	0.5127	1	P02768
ALBU_HUMAN	K.VPQVSTPTLVEVSR.N	1	3.4104	0.3995	2	P02768
ALBU_HUMAN	R.YTKKVPQVSTPTLVEVSR.N	1	4.0806	0.5159	2	P02768
ALBU_HUMAN	R.RHPYFYAPELLFFAKR.Y	1	3.9906	0.4702	1	P02768
ALBU_HUMAN	R.HPYFYAPELLFFAK.R	1	4.1486	0.4288	1	P02768
ALBU_HUMAN	K.LDELRDEGKASSAK.Q	1	3.2119	0.1527	1	P02768
ALBU_HUMAN	R.RHPDYSVVLLLR.L	3	4.1724	0.4272	1	P02768

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Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)

MUCB_HUMAN	R.GQPLSPEKYVTSAPM*PEPQ/	1	3.5311	0.3627	5	P04220
MUCB_HUMAN	R.FTCTVTHTDLPSPLK.Q	1	3.3488	0.309	5	P04220
MUCB_HUMAN	K.QVGSGVTTDEVEAEAK.E	1	4.0349	0.5004		P04220
MUCB_HUMAN	K.VSVFVPPRDGFFGNPR.K	1	3.3851	0.3571	5	P04220
MUCB_HUMAN	K.YVTSAPMPEPQAPGR.Y	1	3.7569	0.602	5	P04220
MUCB_HUMAN	K.YVTSAPM*PEPQAPGR.Y	1	3.6615	0.5322	5	P04220
MUCB_HUMAN	K.SKLICQATGFSPR.Q	1	4.2717	0.5342	5	P04220
MUCB_HUMAN	R.GQPLSPEKYVTSAPMPEPQA	2	4.3777	0.7014	5	P04220
MUCB_HUMAN	K.VSVFVPPR.D	1	2.5872	0.4801	5	P04220
MUCB_HUMAN	R.VFAIPPSFASIFLTK.S	2	3.1096	0.5035	5	P04220
MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	2	3.9674	0.5169	5	P04220
MUCB_HUMAN	R.EGKQVGSGVTTDEVEAEAK.I	1	4.1688	0.4546		P04220

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Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis

CLUS_HUMAN	R.ELDESLQVAER.L	1	2.8979	0.448	1	P10909
CLUS_HUMAN	K.FMETVAEK.A	1	2.1179	0.3197	1	P10909
CLUS_HUMAN	K.LFDSDPITVTVPVEVSR.K	2	5.0899	0.5805	1	P10909
CLUS_HUMAN	K.TLIEKTNEER.K	2	2.843	0.297	1	P10909
CLUS_HUMAN	K.TLLSNLEEAK.K	1	2.5787	0.4135	1	P10909
CLUS_HUMAN	R.ASSIIDELFQDR.F	1	3.5899	0.5459	1	P10909
CLUS_HUMAN	R.KTLLSNLEEAK.K	1	3.3317	0.2658	1	P10909

	CLUS_HUMAN	R.VTTVASHTSDSDVPSGVTEV\	1	4.6901	0.5539		P10909
	CLUS_HUMAN	K.FMETVAEKALQEYR.K	2	4.1294	0.6003	1	P10909
	CLUS_HUMAN	K.TLLSNLEEAKK.K	1	2.5176	0.2833	1	P10909

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Ig alpha-2 chain C region - Homo sapiens (Human)

IGHA2_HUMAN	R.GFSPKDVLVR.W	1	2.7016	0.3494	23	P01877
IGHA2_HUMAN	R.WLQGSQELPREK.Y	3	3.3133	0.3932	23	P01877
IGHA2_HUMAN	R.WLQGSQELPR.E	1	2.6642	0.3089	23	P01877
IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	2	4.4006	0.6047	21	P01877
IGHA2_HUMAN	R.EKYLTWASRQEPSQGTTTFA	1	4.5972	0.5535	21	P01877
IGHA2_HUMAN	R.DASGATFTWTPSSGK.S	1	2.8191	0.4866	10	P01877
IGHA2_HUMAN	K.YLTWASRQEPSQGTTTFAVT	1	4.3049	0.5664	21	P01877
IGHA2_HUMAN	K.SAVQGPPER.D	1	2.5202	0.4097	18	P01877
IGHA2_HUMAN	K.KGDTFSCMVGHEALPLAFTQI	1	3.428	0.2605	19	P01877

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Complement component C9 precursor [Contains: Complement component C9a; Compl

CO9_HUMAN	R.AIEDYINEFSVR.K	1	3.9962	0.5562		P02748
CO9_HUMAN	R.DRDGNTLTYYR.R	1	3.0928	0.3691		P02748
CO9_HUMAN	R.RPWNVASLIYETKGEK.N	2	4.7545	0.4625		P02748
CO9_HUMAN	R.TEHYEEQIEAFK.S	1	3.2369	0.5499		P02748
CO9_HUMAN	R.TEHYEEQIEAFKSIIQEK.T	1	4.6678	0.3905		P02748
CO9_HUMAN	K.TSNFNAAISLK.F	1	3.1394	0.5546		P02748
CO9_HUMAN	K.LSPIYNLVPVK.M	2	3.4207	0.4793		P02748
CO9_HUMAN	K.LLRGTVIDVTDFVNWASSIND/	1	5.821	0.5178		P02748
CO9_HUMAN	R.RPWNVASLIYETK.G	1	3.9729	0.4318		P02748

Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-

ITIH4_HUMAN	R.ANTVQEATFQMELPK.K	1	4.6503	0.569	1	Q14624
ITIH4_HUMAN	R.FKPTLSQQQK.S	1	2.5183	0.3032	1	Q14624
ITIH4_HUMAN	K.EKAEAQAQYSAAVAK.G	1	2.8786	0.5068	1	Q14624
ITIH4_HUMAN	R.VVNRANTVQEATFQMELPK.K	1	4.8987	0.5652	1	Q14624
ITIH4_HUMAN	K.SPEQQETVLDGNLIIR.Y	2	4.5809	0.5234	1	Q14624
ITIH4_HUMAN	K.NVVFVIDKSGSM*SGR.K	1	3.2488	0.4923	1	Q14624
ITIH4_HUMAN	K.GSEMVVAGK.L	1	2.8148	0.4082	1	Q14624
ITIH4_HUMAN	R.LGVYELLLK.V	1	3.3176	0.4585	1	Q14624
ITIH4_HUMAN	K.NVVFVIDKSGSMSGR.K	1	3.8487	0.5405	1	Q14624
ITIH4_HUMAN	K.ILDDLSPR.D	1	2.6441	0.1592	1	Q14624

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IGLV2-14 protein - Homo sapiens (Human)

A0A5C9_HUMAN	K.QSNNKYAASSYLSLTPEQWK	1	2.7104	0.494	29	A0A5C9
A0A5C9_HUMAN	K.ADSSPVKAGVETTTPSKQSNI	1	4.7316	0.4236	17	A0A5C9
A0A5C9_HUMAN	K.AGVETTTPSKQSNNK.Y	1	3.5849	0.4585	17	A0A5C9
A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	4.5	0.6039	29	A0A5C9
A0A5C9_HUMAN	K.ADSSPVKAGVETTTPSK.Q	2	3.3869	0.5092	17	A0A5C9
A0A5C9_HUMAN	R.SYSCQVTHEGSTVEK.T	1	3.7037	0.5175	28	A0A5C9
A0A5C9_HUMAN	K.AGVETTTPSK.Q	1	3.3397	0.3695	17	A0A5C9
A0A5C9_HUMAN	K.AAPSVTLFPPSSEELQANK.A	2	3.9877	0.5214	18	A0A5C9

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Haptoglobin-related protein precursor - Homo sapiens (Human)

HPTR_HUMAN	K.QLVEIEKVVLHPNYHQVDIGLI	1	3.8899	0.4594		P00739
HPTR_HUMAN	R.VGYVSGWGQSDNFKLTDHLK	1	3.4728	0.4822		P00739
HPTR_HUMAN	R.VGYVSGWGQSDNFK.L	1	4.3648	0.6056		P00739

HPTR_HUMAN	R.TEGDGVYTLNDK.K	1	2.7191	0.4119	2	P00739
HPTR_HUMAN	R.ILGGHLDAK.G	1	2.731	0.2804	3	P00739
HPTR_HUMAN	K.YVMLPVADQYDCITHYEGSTC	1	3.2017	0.3918		P00739
HPTR_HUMAN	K.VTSIQDWVQKTIAEN.-	1	2.8735	0.3053	3	P00739
HPTR_HUMAN	K.SPVGVQPILNEHTFCVGMSK.	1	3.8356	0.4283		P00739
HPTR_HUMAN	K.VTSIQDWVQK.T	1	3.0847	0.4541	3	P00739

Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	R.DIPTNSPELEETLTHTITK.L	1	4.3517	0.6488	1	P01042
KNG1_HUMAN	R.ITEATKTVGSDTFYSFK.Y	1	3.3091	0.3804		P01042
KNG1_HUMAN	K.KYFIDFVAR.E	1	3.1867	0.4748	1	P01042
KNG1_HUMAN	K.AATGECTATVGKR.S	1	2.733	0.5113		P01042
KNG1_HUMAN	K.KYNSQNQSNNQFVLYR.I	2	5.0919	0.4841		P01042
KNG1_HUMAN	K.TVGSDTFYSFK.Y	1	2.9558	0.4534		P01042
KNG1_HUMAN	K.YNSQNQSNNQFVLYR.I	1	3.0483	0.3148		P01042
KNG1_HUMAN	K.AATGECTATVGK.R	1	2.6874	0.4648		P01042

Transthyretin precursor (Prealbumin) (TBPA) (TTR) (ATTR) - Homo sapiens (Human)

TTHY_HUMAN	R.YTIAALLSPYSYSTTAVVTNPK	1	3.5895	0.5996		P02766
TTHY_HUMAN	R.RYTIAALLSPYSYSTTAVVTNP	1	4.3222	0.5579		P02766
TTHY_HUMAN	R.KAADDTWEPFASGK.T	2	4.7011	0.6127		P02766
TTHY_HUMAN	R.GSPAINVAVHVFR.K	1	3.923	0.4536		P02766
TTHY_HUMAN	K.VLDAVRGSPAINVAVHVFR.K	1	3.1378	0.3863		P02766
TTHY_HUMAN	K.TSESGELHGLTTEEEFVEGIY	1	4.001	0.4704		P02766
TTHY_HUMAN	K.AADDTWEPFASGK.T	1	4.1696	0.6489		P02766
TTHY_HUMAN	K.TSESGELHGLTTEEEFVEGIY	1	5.397	0.5343		P02766

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Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -

APOA2_HUMAN	K.SKEQLTPLIK.K	1	2.8448	0.2979		P02652
APOA2_HUMAN	K.AGTELVNFLSYFVELGTQPAT	1	3.5458	0.492		P02652
APOA2_HUMAN	K.KAGTELVNFLSYFVELGTQPA	2	4.275	0.554		P02652
APOA2_HUMAN	K.SKEQLTPLIKK.A	1	3.3715	0.4314		P02652
APOA2_HUMAN	K.SPELQAEAK.S	2	2.7384	0.3556		P02652
APOA2_HUMAN	K.VKSPELQAEAKSYFEK.S	1	3.6005	0.5889		P02652

Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	R.DSSTWLTAFVLK.V	1	2.7378	0.4208	2	P0C0L4
CO4A_HUMAN	R.GLQDEDGYR.M	1	2.5422	0.3907	3	P0C0L4
CO4A_HUMAN	R.STQDTVIALDALSAYWIASHT	1	4.4498	0.4475	3	P0C0L4
CO4A_HUMAN	R.TLEIPGNSDPNMIPDGDFNSY	1	2.5262	0.5223	3	P0C0L4
CO4A_HUMAN	R.VTASDPLDTLGSEGALSPGG\	1	3.3517	0.3474	3	P0C0L4
CO4A_HUMAN	K.VLSLAQEQVGGSPEK.L	1	4.1215	0.5488	3	P0C0L4
CO4A_HUMAN	K.DHAVDLIQKGYMR.I	1	3.3116	0.4813	3	P0C0L4
CO4A_HUMAN	K.DHAVDLIQK.G	1	2.683	0.4144	3	P0C0L4

Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) - Homo sapiens (Human)

FINC_HUMAN	K.WLPSSSPVTGYR.V	1	2.4251	0.3914	1	P02751
FINC_HUMAN	R.WSRPQAPITGYR.I	1	3.4503	0.2879	1	P02751
FINC_HUMAN	R.VTWAPPPSIDLTNFLVR.Y	1	2.6481	0.3702	1	P02751
FINC_HUMAN	R.SYTITGLQPGTDYK.I	1	3.6172	0.5387	1	P02751
FINC_HUMAN	R.NTFAEVTGLSPGVTYYFK.V	1	5.3946	0.6546	1	P02751
FINC_HUMAN	K.LGVRPSQGGEAPR.E	1	3.5762	0.4593	1	P02751
FINC_HUMAN	K.IYLYTLNDNAR.S	1	2.9985	0.4347	1	P02751
FINC_HUMAN	R.RPGGEPSPEGTTGQSYNQYS	1	5.6805	0.5459		P02751

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Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	K.VQWKVDNALQSGNSQESVTI	1	5.3758	0.5969	20	A0A5E5
A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	1	4.0113	0.6022	21	A0A5E5
A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	3	5.5862	0.6307	20	A0A5E5
A0A5E5_HUMAN	K.HKVYACEVTHQGLSSPVTK.S	1	3.52	0.5303	21	A0A5E5
A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	1	5.2253	0.6447	20	A0A5E5

IGHM protein - Homo sapiens (Human)

Q6GMY2_HUMAN	K.YAATSQVLLPSK.D	1	3.1227	0.3667	3	Q6GMY2
Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAKESGF	1	3.9437	0.1085	4	Q6GMY2
Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	2	4.5632	0.5367	4	Q6GMY2
Q6GMY2_HUMAN	K.NVPLPVIAELPPK.V	1	2.4138	0.5893	4	Q6GMY2
Q6GMY2_HUMAN	R.EGKQVGSGVTTDQVQAEAK.I	2	4.8742	0.4518	4	Q6GMY2

Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-

ITIH1_HUMAN	R.VQSWKGSLVQASEANLQAAC	1	4.5809	0.5685		P19827
ITIH1_HUMAN	K.LDAQASFLPK.E	1	3.0334	0.333		P19827
ITIH1_HUMAN	K.AAISGENAGLVR.A	1	3.7583	0.4753		P19827
ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	2	3.663	0.3879	1	P19827
ITIH1_HUMAN	R.GFSLDEATNLNGGLLR.G	1	3.0512	0.1493		P19827
ITIH1_HUMAN	R.KAAISGENAGLVR.A	1	3.6166	0.4187		P19827

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Alpha-1-antitrypsin precursor (Alpha-1 protease inhibitor) (Alpha-1-antiproteinase) - Ho

A1AT_HUMAN	K.VFSNGADLSGVTEEAPLKLSK	1	3.4378	0.2754	1	P01009
A1AT_HUMAN	K.SVLGQLGITK.V	2	3.2497	0.4665	1	P01009
A1AT_HUMAN	K.LSITGTYDLK.S	1	2.9039	0.3452	1	P01009
A1AT_HUMAN	K.KLSSWVLLMK.Y	1	3.0234	0.3783		P01009
A1AT_HUMAN	K.ITPNLAEFAFSLYR.Q	1	2.6544	0.4234		P01009

Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr

FETUA_HUMAN	R.HTFM*GVVSLGSPSGEVSHPI	1	4.0314	0.5471		P02765
FETUA_HUMAN	R.HTFMGVVSLGSPSGEVSHPR	1	4.3221	0.5783		P02765
FETUA_HUMAN	K.LDGKFSVVYAK.C	1	3.3296	0.4844		P02765
FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.5552	0.5094		P02765
FETUA_HUMAN	R.TVVQPSVGAAAGPVVPPCPG	2	4.2867	0.7059		P02765

IGHA1 protein - Homo sapiens (Human)

Q8N5K4_HUMAN	K.GTTVTVSSASPTSPK.V	1	3.5773	0.5901	1	Q8N5K4
Q8N5K4_HUMAN	K.TPLTATLSK.S	1	2.6	0.4406	11	Q8N5K4
Q8N5K4_HUMAN	K.TFTCTAAYPESK.T	2	3.0622	0.377	12	Q8N5K4
Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	2	4.1387	0.6122	12	Q8N5K4

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Complement C1r subcomponent precursor (EC 3.4.21.41) (Complement component 1,

C1R_HUMAN	K.TLDEFTIIQNLQPQYQFR.D	1	3.6375	0.6273		P00736
C1R_HUMAN	R.HSCQAECSSELYTEASGYISS	1	3.6651	0.485		P00736
C1R_HUMAN	K.VLNYVDWIKKEMEEED.-	1	4.5723	0.4846		P00736
C1R_HUMAN	K.QRPPDLDTSSNAVDLLFFTDE	1	3.2118	0.2014		P00736
C1R_HUMAN	R.YTTTMGVNTYK.A	1	2.9805	0.4952		P00736

KNG1 protein - Homo sapiens (Human)

Q05CF8_HUMAN	R.IGEIKEETTSHLR.S	4	4.0135	0.4737		Q05CF8
Q05CF8_HUMAN	K.AGAEPASEREVS.-	1	2.5076	0.3872		Q05CF8

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Apolipoprotein A-IV precursor (Apo-AIV) (ApoA-IV) - Homo sapiens (Human)

APOA4_HUMAN	K.KLVPFATELHER.L	1	3.2616	0.2567		P06727

APOA4_HUMAN	K.IDQNVEELKGR.L	1	3.8271	0.3118	1	P06727
APOA4_HUMAN	K.LGEVNTYAGDLQK.K	1	3.3428	0.5157		P06727
APOA4_HUMAN	K.SELTQQLNALFQDK.L	1	3.9987	0.4331		P06727

Apolipoprotein E precursor (Apo-E) - Homo sapiens (Human)

APOE_HUMAN	R.LGPLVEQGR.V	1	2.5321	0.3513		P02649
APOE_HUMAN	R.LSKELQAAQAR.L	1	3.5135	0.4371		P02649
APOE_HUMAN	R.AATVGSLAGQPLQER.A	2	4.2196	0.5271		P02649

Apolipoprotein-L1 precursor (Apolipoprotein L-I) (Apolipoprotein L) (ApoL-I) (Apo-L) (Ap

APOL1_HUMAN	R.VTEPISAESGEQVER.V	1	4.4356	0.5909		O14791
APOL1_HUMAN	K.ILQADQEL.-	1	2.4127	0.346		O14791
APOL1_HUMAN	K.VAQELEEKLNILNNNYK.I	2	4.2498	0.416		O14791

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.TRFVYHLSDLCK.K	1	3.4027	0.1156		P01591
IGJ_HUMAN	R.IIRSSEDPNEDIVER.N	2	4.8629	0.3044		P01591
IGJ_HUMAN	R.SSEDPNEDIVER.N	1	2.9104	0.2989		P01591

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Apolipoprotein(a) precursor (EC 3.4.21.-) (Apo(a)) (Lp(a)) - Homo sapiens (Human)

APOA_HUMAN	R.TPENYPNAGLTR.N	1	3.1129	0.3994	1	P08519
APOA_HUMAN	R.GTYSTTVTGR.T	1	3.2904	0.5102	39	P08519
APOA_HUMAN	R.GSFSTTVTGR.T	1	2.8014	0.3728	4	P08519

C4b-binding protein alpha chain precursor (C4bp) (Proline-rich protein) (PRP) - Homo s

C4BP_HUMAN	R.FSAICQGDGTWSPR.T	1	2.7546	0.4094		P04003
C4BP_HUMAN	K.LNNGEITQHR.K	1	2.4541	0.1835		P04003
C4BP_HUMAN	K.LSLEIEQLELQR.D	1	3.9841	0.2668		P04003

Carboxypeptidase N catalytic chain precursor (EC 3.4.17.3) (CPN) (Carboxypeptidase

CBPN_HUMAN	R.IVQLIQDTR.I	1	3.2179	0.3247		P15169
CBPN_HUMAN	K.YVGNMHGNEALGR.E	1	3.3119	0.3647		P15169
CBPN_HUMAN	R.NFPDLNTYIYYNEK.Y	1	4.138	0.4766		P15169

Carboxypeptidase N subunit 2 precursor (Carboxypeptidase N polypeptide 2) (Carboxy

CPN2_HUMAN	K.LSNNALSGLPQGVFGK.L	2	3.4141	0.5334		P22792
CPN2_HUMAN	K.AGGSWDLAVQER.A	1	2.6148	0.1893		P22792

Fibulin-1 precursor - Homo sapiens (Human)

FBLN1_HUMAN	R.SAATLQQEKTDTVR.C	2	4.2136	0.5613		P23142
FBLN1_HUMAN	R.GYHLNEEGTR.C	1	2.5434	0.3321		P23142

Heparin cofactor 2 precursor (Heparin cofactor II) (HC-II) (Protease inhibitor leuserpin 2

HEP2_HUMAN	R.LNEREVVKVSMMQTK.G	1	3.7793	0.4084		P05546
HEP2_HUMAN	R.LNEREVVK.V	1	3.0082	0.1334		P05546
HEP2_HUMAN	K.TLEAQLTPR.V	1	2.8729	0.3895		P05546

Ig lambda chain V-III region LOI - Homo sapiens (Human)

LV302_HUMAN	-.YVLTQPPSVSVAPGETAR.L	2	3.8701	0.5159		P80748
LV302_HUMAN	R.FSGSNSGNTATLTISR.V	1	4.5371	0.4265	4	P80748

IgGFc-binding protein precursor (FcgammaBP) (Fcgamma-binding protein antigen) - H

FCGBP_HUMAN	R.VNGVLTALPVSVADGR.I	1	2.4859	0.5101	2	Q9Y6R7
FCGBP_HUMAN	R.GSQAVSYTR.S	1	2.9468	0.3523	2	Q9Y6R7
FCGBP_HUMAN	R.GATTSPGVYELSSR.C	1	3.5498	0.6082		Q9Y6R7

Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-

ITIH2_HUMAN	K.FYNQVSTPLLR.N	1	3.1558	0.4017	1	P19823
ITIH2_HUMAN	K.IQPSGGTNINEALLR.A	1	3.0001	0.5474	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLK.K	1	3.211	0.3976	1	P19823

Plasma kallikrein precursor (EC 3.4.21.34) (Plasma prekallikrein) (Kininogenin) (Fletche

KLKB1_HUMAN	K.TQSSDGKAQMQSPA.-	1	3.4332	0.5063	1	P03952
KLKB1_HUMAN	K.VAEYMDWILEK.T	1	3.7517	0.5278	1	P03952
KLKB1_HUMAN	R.IAYGTQGSSGYSLR.L	1	3.2432	0.5338	1	P03952

Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk

PON1_HUMAN	K.IFFYDSENPPASEVLR.I	2	5.2769	0.5703		P27169
PON1_HUMAN	R.IQNILTEEPK.V	1	3.2441	0.1978		P27169

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Alpha-1-antichymotrypsin precursor (ACT) [Contains: Alpha-1-antichymotrypsin His-Prc

AACT_HUMAN	R.NLAVSQVVHK.A	1	2.8192	0.2593		P01011
AACT_HUMAN	R.LYGSEAFATDFQDSAAAK.K	1	4.9881	0.5788		P01011

Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)

A2MG_HUMAN	K.HYDGSYSTFGER.Y	1	2.8728	0.4901		P01023
A2MG_HUMAN	K.QFSFPLSSEPFQGSYK.V	1	3.1793	0.4241		P01023

Apolipoprotein C-III precursor (Apo-CIII) (ApoC-III) - Homo sapiens (Human)

APOC3_HUMAN	K.DALSSVQESQVAQQAR.G	2	3.9931	0.4816		P02656

CD44 antigen precursor (Phagocytic glycoprotein I) (PGP-1) (HUTCH-I) (Extracellular r

CD44_HUMAN	R.YGFIEGHVVIPR.I	2	3.3144	0.435	1	P16070

Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	K.GAYPLSIEPIGVR.F	1	3.4241	0.4348		P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPMK	1	3.2593	0.3704		P00450

Histidine-rich glycoprotein precursor (Histidine-proline-rich glycoprotein) (HPRG) - Hom

HRG_HUMAN	R.RDGYLFQLLR.I	1	2.6295	0.3525		P04196
HRG_HUMAN	R.DGYLFQLLR.I	1	3.3483	0.2502		P04196

Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	2.8538	0.3073	20	Q6N094
Q6N094_HUMAN	K.VSNKALPAPIEK.T	1	2.662	0.1861	24	Q6N094

Ig kappa chain V-III region VG precursor (Fragment) - Homo sapiens (Human)

KV309_HUMAN	R.ASQSVSSYLAWYQQKPGQAI	1	4.4015	0.5622		P04433
KV309_HUMAN	R.LLIYDASNR.A	1	2.4233	0.2887	1	P04433

Ig kappa chain V-IV region Len - Homo sapiens (Human)

KV402_HUMAN	-.DIVMTQSPDSLAVSLGER.A	2	4.9703	0.4704		P01625

Kinesin heavy chain isoform 5A (Neuronal kinesin heavy chain) (NKHC) (Kinesin heavy

KIF5A_HUMAN	K.LEAELSRWR.N	2	2.5325	0.1639		Q12840

Latrophilin-2 (Fragment) - Homo sapiens (Human)

Q9UJ49_HUMAN	-.MTGNYLLTNPLLR.P	1	2.9367	0.2794		Q9UJ49
Q9UJ49_HUMAN	-.M*TGNYLLTNPLLR.P	1	2.5212	0.1882		Q9UJ49

Rheumatoid factor D5 light chain (Fragment) - Homo sapiens (Human)

A0N5G5_HUMAN	-.EIVLTQSPATLSLSPGER.A	2	5.2878	0.5822		A0N5G5

Vitronectin precursor (Serum-spreading factor) (S-protein) (V75) [Contains: Vitronectin

VTNC_HUMAN	K.LIRDVWGIEGPIDAAFTR.I	2	4.9046	0.6092		P04004

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28S ribosomal protein S15, mitochondrial precursor (S15mt) (MRP-S15) - Homo sapier

RT15_HUMAN	K.IRSYEEHLEKHR.K	1	2.454	0.1838		P82914

Acidic repeat-containing protein - Homo sapiens (Human)

Q96QF7_HUMAN	K.MVKTAGLCSTGEM*WYPKWI	1	2.8106	0.1087		Q96QF7

Alpha-1-acid glycoprotein 2 precursor (AGP 2) (Orosomucoid-2) (OMD 2) - Homo sapie

A1AG2_HUMAN	R.SDVMYTDWKK.D	1	2.9949	0.3864		P19652

Apolipoprotein D precursor (Apo-D) (ApoD) - Homo sapiens (Human)

APOD_HUMAN	K.VLNQELR.A	1	2.5001	0.1715		P05090

Biotinidase precursor (EC 3.5.1.12) - Homo sapiens (Human)

BTD_HUMAN	R.LSSGLVTAALYGR.L	1	2.7985	0.3954		P43251

Cadherin-13 precursor (Truncated-cadherin) (T-cadherin) (T-cad) (Heart-cadherin) (H-c

CAD13_HUMAN	R.INENTGSVSVTR.T	1	3.5656	0.5068		P55290

Calcium/calmodulin-dependent protein kinase type IV (EC 2.7.11.17) (CAM kinase-GR)

KCC4_HUMAN	K.GAQAELMKVQALEK.V	1	2.4358	0.1535		Q16566

CDNA FLJ45949 fis, clone PLACE7007973 - Homo sapiens (Human)

Q6ZRZ8_HUMAN	R.DQDERDRAER.K	1	2.4075	0.2236		Q6ZRZ8

C-jun-amino-terminal kinase-interacting protein 1 (JNK-interacting protein 1) (JIP-1) (JN

JIP1_HUMAN	R.RSTATQMAPPGGPPAAPPG(	1	3.3722	0.2048		Q9UQF2

Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co

C1S_HUMAN	K.TMQENSTPRED.-	1	3.3084	0.4528		P09871

Complement C4-B precursor (Basic complement C4) [Contains: Complement C4 beta c

CO4B_HUMAN	K.LQETSNWLLSQQQADGSFQ[	1	6.0448	0.6761	1	P0C0L5

DENN domain-containing protein 2A - Homo sapiens (Human)

DEN2A_HUMAN	R.GFIQERELRR.Q	1	2.7601	0.1102		Q9ULE3

Eukaryotic translation initiation factor 2 subunit 1 (Eukaryotic translation initiation factor

IF2A_HUMAN	R.EVLINNINRR.L	1	2.7885	0.0817		P05198

Eukaryotic translation initiation factor 2C 1 (eIF2C 1) (eIF-2C 1) (Argonaute-1) (Putative

I2C1_HUMAN	K.RVGDTLLGM*ATQCVQVK.N	1	2.5899	0.0981	4	Q9UL18

Haptoglobin precursor [Contains: Haptoglobin alpha chain; Haptoglobin beta chain] - H

HPT_HUMAN	K.LRTEGDGVYTLNNEKQWINK.	1	5.2291	0.4517		P00738

HECT domain and RCC1-like domain-containing protein 2 - Homo sapiens (Human)

HERC2_HUMAN	R.KDDSVEPSGTKK.E	1	2.4712	0.166		O95714

Hemopexin precursor (Beta-1B-glycoprotein) - Homo sapiens (Human)

HEMO_HUMAN	R.WKNFPSPVDAAFR.Q	1	3.7199	0.5157		P02790

HGSC289 (OTTHUMP00000016309) - Homo sapiens (Human)

Q6UXB8_HUMAN	R.LMVELHNLYR.A	1	2.8209	0.3497		Q6UXB8

Hypothetical protein DKFZp686C15213 - Homo sapiens (Human)

Q6MZU6_HUMAN	K.TTPPMLDSDGSFFLYSK.L	1	2.4399	0.289	1	Q6MZU6

Ig heavy chain V-I region V35 precursor - Homo sapiens (Human)

HV103_HUMAN	R.QAPGQGLEWMGR.I	1	2.4154	0.5231	1	P23083

Ig heavy chain V-III region TIL - Homo sapiens (Human)

HV304_HUMAN	-.EVQLLESGGGLVQPGGSLR.L	1	5.0057	0.3081	5	P01765

Ig kappa chain V-II region Cum - Homo sapiens (Human)

KV201_HUMAN	R.FSGSGSGTDFTLK.I	1	3.7227	0.5767	9	P01614

Ig kappa chain V-III region SIE - Homo sapiens (Human)

KV302_HUMAN	R.LLIYGASSR.A	1	3.0887	0.3968	11	P01620

Ig kappa chain V-IV region precursor (Fragment) - Homo sapiens (Human)

KV401_HUMAN	K.LLIYWASTR.E	1	3.159	0.3234	4	P06312

Ig lambda chain V region 4A precursor - Homo sapiens (Human)

LV001_HUMAN	R.FSGSLLGGK.A	1	2.4491	0.2758	2	P04211

Ig lambda chain V-I region HA - Homo sapiens (Human)

LV102_HUMAN	K.SGTSASLAISGLR.S	1	2.7397	0.4354	4	P01700

Ig lambda chain V-IV region Hil - Homo sapiens (Human)

LV403_HUMAN	-.SYELTQPPSVSVSPGQTAR.I	1	4.4722	0.576	1	P01717

IGHG4 protein - Homo sapiens (Human)

Q8TC63_HUMAN	K.YGPPCPSCPAPEFLGGPSVF[	1	3.9828	0.2976	1	Q8TC63

IQ motif and Sec7 domain-containing protein 1 (ADP-ribosylation factors guanine nucle

IQEC1_HUMAN	R.EIFLFNDLLVVTKIFQKK.K	1	3.6184	0.2207		Q6DN90

MORC family CW-type zinc finger protein 4 (Zinc finger CW-type coiled-coil domain pro

MORC4_HUMAN	K.VEELEQER.N	1	2.7648	0.1018	1	Q8TE76

mTERF domain-containing protein 3, mitochondrial precursor (Mitochondrial transcriptio

MTER3_HUMAN	R.PFLACFTYTTDKQSSK.E	1	2.6832	0.2263		Q49AM1

Mucin-16 (Ovarian carcinoma antigen CA125) (Ovarian cancer related tumor marker CA

MUC16_HUMAN	R.LDPKSPGLNR.E	1	2.4008	0.1221	2	Q8WXI7

Myosin-reactive immunoglobulin heavy chain variable region (Fragment) - Homo sapien

Q9UL88_HUMAN	K.SKTDGGTTDYAAPVKGR.L	1	4.0633	0.5484	1	Q9UL88
Q9UL94_HUMAN	-.EVQLVESGAEVK.K	1	3.6363	0.1634	2	Q9UL94

Myosin-reactive immunoglobulin light chain variable region (Fragment) - Homo sapiens

Q9UL83_HUMAN	-.EIVMTQSPATLSVSPGER.A	1	4.7812	0.5546	1	Q9UL83

PDZ domain-containing RING finger protein 4 (Ligand of Numb-protein X 4) (SEMACAI

PZRN4_HUMAN	K.TTEQGCSAESKEKVLEGSK.L	1	2.6071	0.253		Q6ZMN7

Plasma protease C1 inhibitor precursor (C1 Inh) (C1Inh) (C1 esterase inhibitor) (C1-inh

IC1_HUMAN	R.LEDMEQALSPSVFK.A	1	3.2741	0.4692	1	P05155

Protein enabled homolog - Homo sapiens (Human)

ENAH_HUMAN	K.IQDHQVVINCAIPKGLK.Y	1	2.5702	0.3103		Q8N8S7

Protein FAM32A - Homo sapiens (Human)

FA32A_HUMAN	K.GVAELGVTKR.K	1	2.8578	0.0911		Q9Y421

Protein LAP4 (Protein scribble homolog) (hScrib) - Homo sapiens (Human)

LAP4_HUMAN	R.ELCIQKAPGERLGISIR.G	1	2.4801	0.2534		Q14160

Protein Z-dependent protease inhibitor precursor (PZ-dependent protease inhibitor) (PZ

ZPI_HUMAN	R.IFSPFADLSELSATGR.N	1	4.2283	0.565		Q9UK55

Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II) [Contains: Activation peptid

THRB_HUMAN	R.TATSEYQTFFNPR.T	1	3.4867	0.547		P00734

Scaffold attachment factor B (Scaffold attachment factor B1) (SAF-B) (HSP27 estroger

SAFB1_HUMAN	K.ILDILGETCKSEPVK.E	1	2.4975	0.3008	2	Q15424

Serine/threonine-protein kinase ULK2 (EC 2.7.11.1) (Unc-51-like kinase 2) - Homo sap

ULK2_HUMAN	R.M*FYEKNRSLMPSIPR.E	1	2.9302	0.1474		Q8IYT8

Serine/threonine-protein kinase ULK4 (EC 2.7.11.1) (Unc-51-like kinase 4) - Homo sap

ULK4_HUMAN	K.FDAKILHLPTYSVDK.L	1	2.9066	0.2866		Q96C45

Serotransferrin precursor (Transferrin) (Siderophilin) (Beta-1-metal-binding globulin) - H

TRFE_HUMAN	K.MYLGYEYVTAIR.N	1	3.7743	0.5799	1	P02787

Serum amyloid A-4 protein precursor (Constitutively expressed serum amyloid A protein

SAA4_HUMAN	R.SFFKEALQGVGDMGR.A	1	4.0702	0.4166		P35542

Sulfhydryl oxidase 1 precursor (EC 1.8.3.2) (Quiescin Q6) (hQSOX) - Homo sapiens (H

QSCN6_HUMAN	K.AHFSPSNIILDFPAAGSAARR.	1	2.4101	0.2458		O00391

Tenascin-X precursor (TN-X) (Hexabrachion-like protein) - Homo sapiens (Human)

TENX_HUMAN	K.ILISGLEPSTPYR.F	1	2.8274	0.3557	3	P22105

von Willebrand factor precursor (vWF) [Contains: von Willebrand antigen 2 (von Willebr

VWF_HUMAN	R.EGGPSQIGDALGFAVR.Y	1	2.8034	0.3595		P04275

Zinc finger CCHC domain-containing protein 11 - Homo sapiens (Human)

ZCH11_HUMAN	R.M*DDFQLKGIVEEKFVK.W	1	2.6253	0.2739	1	Q5TAX3

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Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	R.KAEEEHLGILGPQLHADVGDK	3	7.6451	0.6431		P00450
CERU_HUMAN	R.IYHSHIDAPK.D	2	3.3499	0.3644		P00450

CERU_HUMAN	R.HYYIAAEEIIWNYAPSGIDIFTK	2	5.7136	0.5543		P00450
CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEMF	1	4.5513	0.6441		P00450
CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEM*	1	5.1265	0.6163		P00450
CERU_HUMAN	R.GPEEEHLGILGPVIWAEVGDT	3	6.09	0.6182		P00450
CERU_HUMAN	R.FNKNNEGTYYSPNYNPQSR.S	2	5.6801	0.5983		P00450
CERU_HUMAN	K.VNKDDEEFIESNK.M	4	4.6014	0.3931		P00450
CERU_HUMAN	K.VDKDNEDFQESNR.M	1	4.1359	0.4584		P00450
CERU_HUMAN	K.WYLFGMGNEVDVHAAFFHG(	2	5.1022	0.6856		P00450
CERU_HUMAN	R.KAEEEHLGILGPQLHADVGDK	2	6.0298	0.6325		P00450
CERU_HUMAN	R.EYTDASFTNR.K	1	2.6978	0.4603		P00450
CERU_HUMAN	R.KERGPEEEHLGILGPVIWAEV	1	5.9261	0.5534		P00450
CERU_HUMAN	R.M*FTTAPDQVDKEDEDFQESI	3	4.1471	0.5684		P00450
CERU_HUMAN	R.MFTTAPDQVDKEDEDFQESN	4	4.1487	0.6296		P00450
CERU_HUMAN	R.PYSIHAHGVQTESSTVTPTLP	2	4.6263	0.6516		P00450
CERU_HUMAN	R.QSEDSTFYLGER.T	2	3.2444	0.4133		P00450
CERU_HUMAN	R.RPYLKVFNPR.R	1	3.221	0.229		P00450
CERU_HUMAN	R.RQSEDSTFYLGER.T	1	3.6975	0.3014		P00450
CERU_HUMAN	R.SGAGTEDSACIPWAYYSTVD(	1	4.0327	0.4785		P00450
CERU_HUMAN	R.SVPPSASHVAPTETFTYEWT\	3	3.3365	0.4812		P00450
CERU_HUMAN	R.TYYIAAVEVEWDYSPQR.E	3	5.3405	0.5371		P00450
CERU_HUMAN	K.ALYLQYTDETFR.T	7	4.3867	0.5526		P00450
CERU_HUMAN	K.TYSDHPEKVNKDDEEFIESNK	2	5.4274	0.5208		P00450
CERU_HUMAN	R.TTIEKPVWLGFLGPIIK.A	3	4.935	0.4856		P00450
CERU_HUMAN	K.DNEDFQESNR.M	1	3.4326	0.3808		P00450
CERU_HUMAN	K.DIASGLIGPLIICK.K	1	3.3087	0.5941		P00450
CERU_HUMAN	K.TYSDHPEK.V	3	2.5164	0.3672		P00450
CERU_HUMAN	K.DDEEFIESNK.M	1	2.8312	0.4821		P00450
CERU_HUMAN	K.DIFTGLIGPMK.I	3	3.2612	0.3867		P00450
CERU_HUMAN	K.DVDKEFYLFPTVFDENESLLLI	1	5.5442	0.5752		P00450
CERU_HUMAN	K.ERGPEEEHLGILGPVIWAEVG	5	7.2271	0.7235		P00450
CERU_HUMAN	K.EVGPTNADPVCLAK.M	3	3.503	0.6483		P00450
CERU_HUMAN	K.GAYPLSIEPIGVR.F	3	3.5501	0.4686		P00450
CERU_HUMAN	K.HYYIGIIETTWDYASDHGEK.K	16	6.7611	0.6821		P00450
CERU_HUMAN	K.HYYIGIIETTWDYASDHGEKK.	1	5.2626	0.5921		P00450
CERU_HUMAN	K.KLISVDTEHSNIYLQNGPDR.I	2	4.8661	0.581		P00450
CERU_HUMAN	K.NMATRPYSIHAHGVQTESST\	2	5.883	0.6773		P00450
CERU_HUMAN	K.TYCSEPEKVDKDNEDFQESN	1	4.7784	0.4739		P00450
CERU_HUMAN	K.DIFTGLIGPM*K.I	1	3.2531	0.3962		P00450
CERU_HUMAN	K.LISVDTEHSNIYLQNGPDR.I	3	4.5699	0.6463		P00450
CERU_HUMAN	K.NNEGTYYSPNYNPQSR.S	2	4.6069	0.5651		P00450
CERU_HUMAN	K.PVWLGFLGPIIK.A	4	4.2241	0.5872		P00450
CERU_HUMAN	K.NM*ATRPYSIHAHGVQTESST	1	5.2609	0.6473		P00450
CERU_HUMAN	K.NLASRPYTFHSHGITYYK.E	3	4.6355	0.6072		P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPMK	2	5.6308	0.6386		P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPM*K	2	4.5463	0.6487		P00450
CERU_HUMAN	K.MYYSAVDPTK.D	1	2.548	0.4306		P00450
CERU_HUMAN	K.M*YYSAVDPTKDIFTGLIGPM*	3	4.3103	0.6216		P00450
CERU_HUMAN	K.M*YYSAVDPTK.D	1	3.0841	0.3815		P00450

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Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-

ITIH1_HUMAN	R.GFSLDEATNLNGGLLR.G	5	4.3301	0.4879		P19827
ITIH1_HUMAN	R.IYEDHDATQQLQGFYSQVAK.	1	5.5506	0.6376	1	P19827

ITIH1_HUMAN	R.GHMLENHVER.L	1	3.3255	0.5669	1	P19827
ITIH1_HUMAN	R.GM*ADQDGLKPTIDKPSEDSF	1	3.9006	0.5573	1	P19827
ITIH1_HUMAN	R.GMADQDGLKPTIDKPSEDSPI	1	5.7865	0.5211	1	P19827
ITIH1_HUMAN	R.GMADQDGLKPTIDKPSEDSPI	1	3.627	0.5442	1	P19827
ITIH1_HUMAN	R.IADNKQSSFK.A	1	2.8838	0.2609	1	P19827
ITIH1_HUMAN	R.GHM*LENHVER.L	1	2.91	0.51	1	P19827
ITIH1_HUMAN	R.KAAISGENAGLVR.A	2	3.885	0.45		P19827
ITIH1_HUMAN	R.LWAYLTIQELLAK.R	6	5.2387	0.57	1	P19827
ITIH1_HUMAN	R.NHM*QYEIVIK.V	1	2.7384	0.3458		P19827
ITIH1_HUMAN	R.PTVSQQQSCPTCSTSLLNGH	1	3.3497	0.3206		P19827
ITIH1_HUMAN	R.QAVDTAVDGVFIR.S	2	4.0839	0.3928		P19827
ITIH1_HUMAN	R.RTFVLSALQPSPTHSSSNTQF	1	4.3761	0.5457	1	P19827
ITIH1_HUMAN	R.FAHYVVTSQVVNTANEAR.E	2	6.9636	0.5764		P19827
ITIH1_HUMAN	K.ILGDM*QPGDYFDLVLFGTR.V	1	4.43	0.3887		P19827
ITIH1_HUMAN	R.NHMQYEIVIK.V	2	3.3445	0.4197		P19827
ITIH1_HUMAN	R.TMEQFTIHLTVNPQSK.V	1	3.9388	0.4388		P19827
ITIH1_HUMAN	K.TAFISDFAVTADGNAFIGDIK.D	4	5.4606	0.6804		P19827
ITIH1_HUMAN	K.AAISGENAGLVR.A	2	3.8781	0.5339		P19827
ITIH1_HUMAN	K.ADVQAHGEGQEFSITCLVDEI	1	5.246	0.4996	1	P19827
ITIH1_HUMAN	K.GSLVQASEANLQAAQDFVR.C	5	6.3263	0.6909		P19827
ITIH1_HUMAN	K.ILGDMQPGDYFDLVLFGTR.V	1	5.0308	0.5904		P19827
ITIH1_HUMAN	K.NVVFVIDISGSMR.G	1	3.0241	0.5528		P19827
ITIH1_HUMAN	K.LDAQASFLPK.E	2	3.1253	0.3793		P19827
ITIH1_HUMAN	K.QLVHHFEIDVDIFEPQGISK.L	2	3.6058	0.6141		P19827
ITIH1_HUMAN	R.EVAFDLEIPK.T	2	2.9051	0.4001		P19827
ITIH1_HUMAN	K.RNHMQYEIVIK.V	1	3.3047	0.3758		P19827
ITIH1_HUMAN	K.TAFISDFAVTADGNAFIGDIKD	3	4.0042	0.5208		P19827
ITIH1_HUMAN	K.TAFISDFAVTADGNAFIGDIKD	1	4.3978	0.3599		P19827
ITIH1_HUMAN	K.VTFQLTYEEVLK.R	1	2.4282	0.3985		P19827
ITIH1_HUMAN	K.VTFQLTYEEVLKR.N	1	3.7568	0.5164		P19827
ITIH1_HUMAN	R.ERGHMLENHVER.L	1	4.1689	0.2965	1	P19827
ITIH1_HUMAN	K.PLLVDVDLQYPQDAVLALTQN	1	4.6244	0.5786	1	P19827
ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	4	3.9088	0.4557	1	P19827

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Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-

ITIH2_HUMAN	K.KFYNQVSTPLLR.N	1	3.5614	0.4479	1	P19823
ITIH2_HUMAN	K.M*KQTVEAMK.T	1	2.7198	0.3226	1	P19823
ITIH2_HUMAN	K.RLSNENHGIAQR.I	2	3.5736	0.3534	1	P19823
ITIH2_HUMAN	K.TILDDLRAEDHFSVIDFNQNIR	1	3.1296	0.5058	1	P19823
ITIH2_HUMAN	K.VQFELHYQEVK.W	1	3.8888	0.3393	2	P19823
ITIH2_HUMAN	K.VQSTITSR.M	1	2.432	0.413	2	P19823
ITIH2_HUMAN	K.VVNNSPQPQNVVFDVQIPK.G	1	5.2792	0.5828	2	P19823
ITIH2_HUMAN	R.AEDHFSVIDFNQNIR.T	1	4.3171	0.4332	1	P19823
ITIH2_HUMAN	R.FLHVPDTFEGHFDGVPVISK.C	2	4.8977	0.6632	2	P19823
ITIH2_HUMAN	K.HLEVDVWVIEPQGLR.F	2	5.3972	0.5897	2	P19823
ITIH2_HUMAN	R.KLWAYLTINQLLAER.S	3	5.6257	0.5946	1	P19823
ITIH2_HUMAN	R.ETAVDGELVVLYDVK.R	14	5.0392	0.6434	1	P19823
ITIH2_HUMAN	R.LSNENHGIAQR.I	2	3.7332	0.4238	1	P19823
ITIH2_HUMAN	R.M*ATTMIQSK.V	1	2.5179	0.464	2	P19823
ITIH2_HUMAN	R.M*LADAPPQDPSCCSGALYY(	2	4.2002	0.5912	1	P19823
ITIH2_HUMAN	R.MATTMIQSK.V	1	2.6594	0.4986	2	P19823
ITIH2_HUMAN	R.MLADAPPQDPSCCSGALYYG	1	3.5959	0.4354	1	P19823

ITIH2_HUMAN	R.NVQFNYPHTSVTDVTQNNFH	1	5.7687	0.5076	1	P19823
ITIH2_HUMAN	R.SILQM*SLDHHIVTPLTSLVIEN	1	6.4134	0.6228	1	P19823
ITIH2_HUMAN	R.SILQMSLDHHIVTPLTSLVIENI	1	4.8709	0.5002	1	P19823
ITIH2_HUMAN	R.SSALDMENFR.T	1	3.3624	0.3791	2	P19823
ITIH2_HUMAN	R.TEVNVLPGAK.V	1	3.008	0.4963	2	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLKK.F	1	4.0493	0.4489	1	P19823
ITIH2_HUMAN	K.FYNQVSTPLLR.N	3	2.8129	0.3361	1	P19823
ITIH2_HUMAN	K.DKHADPDFTR.K	1	2.6516	0.4015	1	P19823
ITIH2_HUMAN	K.AHVSFKPTVAQQR.I	2	3.4951	0.4652	2	P19823
ITIH2_HUMAN	K.AGELEVFNGYFVHFFAPDNLI	3	5.5355	0.5734	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLK.K	2	3.7803	0.4743	1	P19823
ITIH2_HUMAN	K.HADPDFTR.K	1	2.4434	0.3433	1	P19823

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Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alp

CO3_HUMAN	K.QLYNVEATSYALLALLQLK.D	1	7.0806	0.534	P01024
CO3_HUMAN	K.VLLDGVQNPR.A	1	3.5023	0.56	P01024
CO3_HUMAN	K.VQLSNDFDEYIM*AIEQTIK.S	1	5.0867	0.6095	P01024
CO3_HUMAN	K.VQLSNDFDEYIMAIEQTIK.S	1	5.9006	0.6729	P01024
CO3_HUMAN	R.AYYENSPQQVFSTEFEVK.E	1	3.6456	0.5693	P01024
CO3_HUMAN	R.EGVQKEDIPPADLSDQVPDTI	2	5.4599	0.548	P01024
CO3_HUMAN	R.EPGQDLVVLPLSITTDFIPSFR	1	4.5304	0.63	P01024
CO3_HUMAN	R.EVVADSVWVDVK.D	1	3.415	0.5514	P01024
CO3_HUMAN	R.ILLQGTPVAQMTEDAVDAER.I	3	5.3076	0.65	P01024
CO3_HUMAN	K.VHQYFNVELIQPGAVK.V	1	4.1212	0.4023	P01024
CO3_HUMAN	R.KVLLDGVQNPR.A	1	2.8749	0.3083	P01024
CO3_HUMAN	K.KVEGTAFVIFGIQDGEQR.I	1	4.5362	0.5451	P01024
CO3_HUMAN	R.LVAYYTLIGASGQR.E	1	3.7569	0.5578	P01024
CO3_HUMAN	R.SGIPIVTSPYQIHFTK.T	1	4.1842	0.5855	P01024
CO3_HUMAN	R.SNLDEDIIAEENIVSR.S	1	3.416	0.4836	P01024
CO3_HUMAN	R.SYTVAIAGYALAQMGR.L	1	2.709	0.327	P01024
CO3_HUMAN	R.TELRPGETLNVNFLLR.M	1	5.172	0.5062	P01024
CO3_HUMAN	R.TVMVNIENPEGIPVK.Q	1	3.9038	0.4554	P01024
CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	3.2239	0.4655	P01024
CO3_HUMAN	K.DICEEQVNSLPGSITK.A	1	2.7751	0.4603	P01024
CO3_HUMAN	K.AGDFLEANYM*NLQR.S	2	4.0918	0.4953	P01024
CO3_HUMAN	K.AGDFLEANYMNLQR.S	1	4.4307	0.4475	P01024
CO3_HUMAN	K.TGLQEVEVK.A	1	3.3167	0.3936	P01024
CO3_HUMAN	K.SSLSVPYVIVPLK.T	1	3.0082	0.3409	P01024
CO3_HUMAN	R.SEETKENEGFTVTAEGK.G	1	4.6189	0.594	P01024
CO3_HUMAN	K.SGQSEDRQPVPGQQMTLK.I	1	4.5269	0.5515	P01024
CO3_HUMAN	R.VPVAVQGEDTVQSLTQGDGV	2	5.8495	0.7364	P01024
CO3_HUMAN	K.QELSEAEQATR.T	1	3.3018	0.2988	P01024
CO3_HUMAN	K.RIPIEDGSGEVVLSR.K	2	5.1026	0.5708	P01024
CO3_HUMAN	K.TIYTPGSTVLYR.I	1	2.6206	0.5093	P01024
CO3_HUMAN	K.EDIPPADLSDQVPDTESETR.I	1	4.5135	0.6871	P01024
CO3_HUMAN	K.EYVLPSFEVIVEPTEK.F	2	5.0068	0.5148	P01024
CO3_HUMAN	K.FYYIYNEK.G	1	2.6681	0.3195	P01024
CO3_HUMAN	K.GQGTLSVVTMYHAK.A	3	3.446	0.5516	P01024
CO3_HUMAN	K.KGYTQQLAFR.Q	1	3.3841	0.4461	P01024
CO3_HUMAN	K.KQELSEAEQATR.T	2	4.161	0.4019	P01024
CO3_HUMAN	K.LSINTHPSQKPLSITVR.T	1	4.9425	0.5476	P01024
CO3_HUMAN	K.AAVYHHFISDGVRK.S	1	3.3434	0.4289	P01024

CO3_HUMAN	K.SDDKVTLEER.L	1	2.9032	0.1551		P01024
CO3_HUMAN	R.YYGGGYGSTQATFMVFQALA	2	5.5775	0.6776		P01024
CO3_HUMAN	R.YYTYLIMNK.G	1	2.804	0.4252		P01024

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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H

APOA1_HUMAN	R.QEMSKDLEEVK.A	2	3.0563	0.2414		P02647
APOA1_HUMAN	R.QEM*SKDLEEVK.A	1	2.4545	0.193		P02647
APOA1_HUMAN	R.LEALKENGGAR.L	1	3.2411	0.3159		P02647
APOA1_HUMAN	R.EQLGPVTQEFWDNLEK.E	1	4.0239	0.5692		P02647
APOA1_HUMAN	R.DYVSQFEGSALGK.Q	2	3.6179	0.5883		P02647
APOA1_HUMAN	K.WQEEM*ELYR.Q	1	3.0536	0.4112		P02647
APOA1_HUMAN	R.THLAPYSDELR.Q	1	2.9938	0.4554		P02647
APOA1_HUMAN	K.VEPLRAELQEGAR.Q	1	2.9413	0.3149		P02647
APOA1_HUMAN	K.LREQLGPVTQEFWDNLEK.E	2	5.776	0.6025		P02647
APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	5	4.7755	0.6181		P02647
APOA1_HUMAN	K.ETEGLRQEM*SKDLEEVK.A	1	3.6306	0.478		P02647
APOA1_HUMAN	K.ETEGLRQEM*SK.D	1	2.9893	0.434		P02647
APOA1_HUMAN	K.DSGRDYVSQFEGSALGK.Q	2	5.1248	0.5602		P02647
APOA1_HUMAN	K.DLATVYVDVLK.D	1	3.449	0.5452		P02647
APOA1_HUMAN	K.VSFLSALEEYTK.K	2	4.5202	0.5629		P02647
APOA1_HUMAN	K.KWQEEM*ELYR.Q	1	2.482	0.1893		P02647
APOA1_HUMAN	R.VKDLATVYVDVLK.D	2	3.6441	0.4916		P02647

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Serum albumin precursor - Homo sapiens (Human)

ALBU_HUMAN	K.VFDEFKPLVEEPQNLIK.Q	2	5.2812	0.4844	2	P02768
ALBU_HUMAN	K.AAFTECCQAADK.A	1	2.7008	0.2506	1	P02768
ALBU_HUMAN	K.ADDKETCFAEEGKK.L	1	4.0691	0.3857	2	P02768
ALBU_HUMAN	K.AVM*DDFAAFVEK.C	1	3.4885	0.5582	2	P02768
ALBU_HUMAN	K.DVFLGM*FLYEYAR.R	1	3.0526	0.4635	1	P02768
ALBU_HUMAN	K.FQNALLVR.Y	2	2.9109	0.2511	2	P02768
ALBU_HUMAN	K.KQTALVELVK.H	2	3.6835	0.3782	2	P02768
ALBU_HUMAN	K.KVPQVSTPTLVEVSR.N	2	5.4103	0.5659	2	P02768
ALBU_HUMAN	K.LVAASQAALGL.-	1	2.8613	0.6112	1	P02768
ALBU_HUMAN	K.SLHTLFGDKLCTVATLR.E	1	4.0645	0.4032	2	P02768
ALBU_HUMAN	R.RHPDYSVVLLLR.L	1	4.6162	0.4266	1	P02768
ALBU_HUMAN	R.RHPYFYAPELLFFAK.R	1	5.0501	0.4556	1	P02768
ALBU_HUMAN	K.LVNEVTEFAK.T	1	2.9605	0.4236	2	P02768

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Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	K.TWQDCEYKDAAK.A	2	3.5975	0.4724		P01042
KNG1_HUMAN	R.DIPTNSPELEETLTHTITK.L	1	4.0214	0.6718	1	P01042
KNG1_HUMAN	K.YFIDFVAR.E	2	3.2398	0.4482	1	P01042
KNG1_HUMAN	K.TVGSDTFYSFK.Y	1	3.7619	0.6191		P01042
KNG1_HUMAN	K.SVSEINPTTQMK.E	1	3.265	0.4728		P01042
KNG1_HUMAN	K.LDDDLEHQGGHVLDHGHK.H	1	4.0196	0.5616		P01042
KNG1_HUMAN	K.KYNSQNQSNNQFVLYR.I	1	5.3583	0.5685		P01042
KNG1_HUMAN	K.KYFIDFVAR.E	1	3.3106	0.422	1	P01042
KNG1_HUMAN	K.ICVGCPRDIPTNSPELEETLTH	1	3.8333	0.5063	1	P01042
KNG1_HUMAN	K.AVDAALKK.Y	1	2.5985	0.3204		P01042
KNG1_HUMAN	K.AATGECTATVGK.R	1	3.0333	0.4774		P01042
KNG1_HUMAN	K.YNSQNQSNNQFVLYR.I	2	4.7637	0.4967		P01042

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Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)

A2MG_HUMAN	R.VGFYESDVMGR.G	1	3.4327	0.453		P01023
A2MG_HUMAN	K.AIGYLNTGYQR.Q	2	3.2163	0.4049		P01023
A2MG_HUMAN	K.DTVIKPLLVEPEGLEK.E	1	4.5551	0.3496		P01023
A2MG_HUMAN	K.HYDGSYSTFGER.Y	1	3.2909	0.4886		P01023
A2MG_HUMAN	K.LPPNVVEESAR.A	1	3.2042	0.5496		P01023
A2MG_HUMAN	K.NEDSLVFVQTDK.S	1	3.7339	0.5924		P01023
A2MG_HUMAN	K.QQNAQGGFSSTQDTVVALHA	1	3.6958	0.4238		P01023
A2MG_HUMAN	R.LLLQQVSLPELPGEYSMK.V	1	4.8947	0.5549		P01023
A2MG_HUMAN	R.TEHPFTVEEFVLPK.F	1	3.9112	0.3843		P01023
A2MG_HUMAN	R.VTAAPQSVCALR.A	1	2.6093	0.3587		P01023
A2MG_HUMAN	R.VVSM*DENFHPLNELIPLVYIQ	1	3.5563	0.3329		P01023
A2MG_HUMAN	R.SASNMAIVDVK.M	1	2.6079	0.3071		P01023

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Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	K.LHLETDSLALVALGALDTALY/	1	4.7529	0.5953	3	P0C0L4
CO4A_HUMAN	R.VGDTLNLNLR.A	1	3.4732	0.4319	3	P0C0L4
CO4A_HUMAN	R.VFALDQK.M	1	2.5503	0.1977	3	P0C0L4
CO4A_HUMAN	R.STQDTVIALDALSAYWIASHT	1	4.2578	0.6131	3	P0C0L4
CO4A_HUMAN	R.HLVPGAPFLLQALVR.E	1	2.7498	0.4302	3	P0C0L4
CO4A_HUMAN	R.GPEVQLVAHSPWLK.D	1	2.8358	0.4485	3	P0C0L4
CO4A_HUMAN	R.GLQDEDGYR.M	1	2.5968	0.5206	3	P0C0L4
CO4A_HUMAN	R.DSSTWLTAFVLK.V	1	2.6235	0.3397	2	P0C0L4
CO4A_HUMAN	K.VLSLAQEQVGGSPEK.L	2	5.1059	0.6161	3	P0C0L4
CO4A_HUMAN	K.LNMGITDLQGLR.L	1	2.5586	0.2846	3	P0C0L4
CO4A_HUMAN	R.YIYGKPVQGVAYVR.F	1	3.2959	0.5991	3	P0C0L4

Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) - Homo sapiens (Human)

FINC_HUMAN	K.WLPSSSPVTGYR.V	1	3.1163	0.4837	1	P02751
FINC_HUMAN	R.RPGGEPSPEGTTGQSYNQY	1	5.4137	0.5425		P02751
FINC_HUMAN	R.TEIDKPSQMQVTDVQDNSISV	1	5.5515	0.3436	1	P02751
FINC_HUMAN	R.VPGTSTSATLTGLTR.G	1	3.1377	0.4841	1	P02751
FINC_HUMAN	K.IYLYTLNDNAR.S	1	3.8327	0.416	1	P02751
FINC_HUMAN	K.VTIMWTPPESAVTGYR.V	1	3.4243	0.5166	1	P02751
FINC_HUMAN	R.ITYGETGGNSPVQEFTVPGSK	1	4.488	0.5134	1	P02751
FINC_HUMAN	R.NLQPASEYTVSLVAIK.G	1	3.227	0.4345	1	P02751
FINC_HUMAN	R.NTFAEVTGLSPGVTYYFK.V	1	3.7594	0.5942	1	P02751
FINC_HUMAN	R.WKCDPVDQCQDSETGTFYQI	1	4.5905	0.4025	2	P02751
FINC_HUMAN	R.WSRPQAPITGYR.I	1	3.6001	0.346	1	P02751
FINC_HUMAN	K.LGVRPSQGGEAPR.E	1	2.6399	0.265	1	P02751

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Apolipoprotein B-100 precursor (Apo B-100) [Contains: Apolipoprotein B-48 (Apo B-48)

APOB_HUMAN	K.YNALDLTNNGK.L	1	2.7654	0.1942		P04114
APOB_HUMAN	K.AVSM*PSFSILGSDVR.V	1	4.0603	0.5483		P04114
APOB_HUMAN	K.ENFAGEATLQR.I	1	2.5091	0.4312		P04114
APOB_HUMAN	K.FPEVDVLTK.Y	1	2.9504	0.3317		P04114
APOB_HUMAN	K.ALVEQGFTVPEIK.T	1	2.5383	0.2644		P04114
APOB_HUMAN	R.M*YQM*DIQQELQR.Y	1	3.4837	0.3835		P04114
APOB_HUMAN	K.NLTDFAEQYSIQDWAK.R	1	2.4487	0.2359		P04114
APOB_HUMAN	K.IAELSATAQEIIK.S	1	2.8708	0.3884		P04114
APOB_HUMAN	K.NFVASHIANILNSEELDIQDLK	1	6.4638	0.4767		P04114

	APOB_HUMAN	R.SPSQADINK.I	1	3	0.3132		P04114
	APOB_HUMAN	R.SEYQADYESLR.F	1	3.1846	0.5405		P04114

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Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-

	ITIH4_HUMAN	K.LALDNGGLAR.R	1	2.6888	0.3256	1	Q14624
	ITIH4_HUMAN	K.EKAEAQAQYSAAVAK.G	1	4.3265	0.6481	1	Q14624
	ITIH4_HUMAN	K.AEAQAQYSAAVAK.G	1	4.6531	0.5173	1	Q14624
	ITIH4_HUMAN	K.ILDDLSPR.D	1	2.8798	0.1937	1	Q14624
	ITIH4_HUMAN	R.LGVYELLLK.V	1	2.725	0.5061	1	Q14624
	ITIH4_HUMAN	K.LQDRGPDVLTATVSGK.L	1	3.0593	0.496	1	Q14624
	ITIH4_HUMAN	K.SPEQQETVLDGNLIIR.Y	1	4.7218	0.4493	1	Q14624
	ITIH4_HUMAN	R.DQFNLIVFSTEATQWRPSLVF	1	5.3314	0.4925	1	Q14624
	ITIH4_HUMAN	R.GPDVLTATVSGK.L	1	3.1508	0.5252	1	Q14624

Lumican precursor (Keratan sulfate proteoglycan lumican) (KSPG lumican) - Homo sap

	LUM_HUMAN	R.LKEDAVSAAFK.G	1	3.3321	0.3572		P51884
	LUM_HUMAN	R.LPSGLPVSLLTLYLDNNK.I	2	5.5251	0.6226		P51884
	LUM_HUMAN	R.FNALQYLR.L	1	3.0957	0.2872		P51884
	LUM_HUMAN	K.SLEYLDLSFNQIAR.L	1	4.4958	0.4373		P51884
	LUM_HUMAN	K.SLEDLQLTHNK.I	2	2.8885	0.473		P51884
	LUM_HUMAN	K.RFNALQYLR.L	1	3.3418	0.2566		P51884
	LUM_HUMAN	R.NNQIDHIDEK.A	1	3.2258	0.3067		P51884

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Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)

	MUCB_HUMAN	K.YVTSAPM*PEPQAPGR.Y	1	3.2687	0.5713	5	P04220
	MUCB_HUMAN	K.YVTSAPMPEPQAPGR.Y	2	3.7309	0.6551	5	P04220
	MUCB_HUMAN	K.VSVFVPPR.D	1	2.5387	0.4638	5	P04220
	MUCB_HUMAN	K.QVGSGVTTDEVEAEAK.E	1	3.0672	0.433		P04220
	MUCB_HUMAN	K.LICQATGFSPR.Q	1	3.6477	0.534	5	P04220
	MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	1	4.452	0.5188	5	P04220
	MUCB_HUMAN	R.VFAIPPSFASIFLTK.S	1	2.88	0.5408	5	P04220

IGLV2-14 protein - Homo sapiens (Human)

	A0A5C9_HUMAN	R.SYSCQVTHEGSTVEK.T	2	4.8361	0.611	28	A0A5C9
	A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	5.2299	0.6322	29	A0A5C9
	A0A5C9_HUMAN	K.AGVETTTPSK.Q	1	3.3702	0.3627	17	A0A5C9
	A0A5C9_HUMAN	K.AAPSVTLFPPSSEELQANK.A	4	4.1104	0.5395	18	A0A5C9

Transthyretin precursor (Prealbumin) (TBPA) (TTR) (ATTR) - Homo sapiens (Human)

	TTHY_HUMAN	R.KAADDTWEPFASGK.T	1	4.6936	0.6245		P02766
	TTHY_HUMAN	K.AADDTWEPFASGK.T	1	3.7585	0.6033		P02766
	TTHY_HUMAN	K.TSESGELHGLTTEEEFVEGIY	3	5.3731	0.6916		P02766
	TTHY_HUMAN	R.YTIAALLSPYSYSTTAVVTNPK	1	3.5282	0.4192		P02766
	TTHY_HUMAN	K.TSESGELHGLTTEEEFVEGIY	2	4.5014	0.5666		P02766

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Ig alpha-2 chain C region - Homo sapiens (Human)

	IGHA2_HUMAN	K.YLTWASR.Q	1	2.5058	0.3501	21	P01877
	IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	2	4.0791	0.6277	21	P01877
	IGHA2_HUMAN	K.KGDTFSCMVGHEALPLAFTQ	1	4.466	0.3794	19	P01877
	IGHA2_HUMAN	K.SAVQGPPER.D	1	2.5867	0.3631	18	P01877
	IGHA2_HUMAN	R.WLQGSQELPR.E	2	3.3314	0.4244	23	P01877

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Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk

PON1_HUMAN	K.ILLMDLNEEDPTVLELGITGSK	1	3.5078	0.5758		P27169
PON1_HUMAN	R.IQNILTEEPK.V	1	3.8464	0.213		P27169
PON1_HUMAN	K.IFFYDSENPPASEVLR.I	3	5.8656	0.6284		P27169
PON1_HUMAN	R.VVAEGFDFANGINISPDGK.Y	1	3.026	0.2793		P27169

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Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co

C1S_HUMAN	K.TMQENSTPRED.-	1	3.4685	0.4953		P09871
C1S_HUMAN	R.TNFDNDIALVR.L	2	3.2065	0.3518		P09871
C1S_HUMAN	R.SSNNPHSPIVEEFQVPYNK.L	1	4.1189	0.4624	1	P09871
C1S_HUMAN	K.SDFSNEER.F	1	2.4732	0.3737	4	P09871

Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	3	4.0315	0.481	21	A0A5E5
A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	1	5.2453	0.6423	20	A0A5E5
A0A5E5_HUMAN	K.DSTYSLSSTLTLSK.A	1	2.7193	0.2932	21	A0A5E5

Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S

SAMP_HUMAN	K.IVLGQEQDSYGGKFDR.S	1	4.1137	0.5074		P02743
SAMP_HUMAN	R.VGEYSLYIGR.H	1	3.1572	0.497		P02743
SAMP_HUMAN	R.QGYFVEAQPK.I	1	2.594	0.4894		P02743
SAMP_HUMAN	R.AYSLFSYNTQGR.D	1	2.6535	0.5099		P02743
SAMP_HUMAN	R.GYVIIKPLVWV.-	1	2.4318	0.4151		P02743

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Alpha-1-antitrypsin precursor (Alpha-1 protease inhibitor) (Alpha-1-antiproteinase) - Ho

A1AT_HUMAN	K.TDTSHHDQDHPTFNK.I	1	3.4864	0.3271		P01009
A1AT_HUMAN	K.ITPNLAEFAFSLYR.Q	1	3.3167	0.6353		P01009
A1AT_HUMAN	K.VFSNGADLSGVTEEAPLK.L	1	3.1252	0.3499	1	P01009
A1AT_HUMAN	K.ELDRDTVFALVNYIFFK.G	1	3.4486	0.3319		P01009

AMBP protein precursor [Contains: Alpha-1-microglobulin (Protein HC) (Complex-formi

AMBP_HUMAN	K.KEDSCQLGYSAGPCMGMTSF	1	2.8148	0.371		P02760
AMBP_HUMAN	R.AFIQLWAFDAVK.G	1	2.6677	0.5306		P02760
AMBP_HUMAN	R.TVAACNLPIVR.G	2	3.0014	0.5124		P02760

Haptoglobin-related protein precursor - Homo sapiens (Human)

HPTR_HUMAN	R.LRTEGDGVYTLNDK.K	1	2.7808	0.2445	2	P00739
HPTR_HUMAN	R.ILGGHLDAK.G	1	2.6734	0.2857	3	P00739
HPTR_HUMAN	K.GSFPWQAK.M	2	2.7904	0.385	3	P00739

Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	R.VVSVLTVLHQDWLNGK.E	1	3.4603	0.5173	25	Q6N094
Q6N094_HUMAN	R.EPQVYTLPPSRDELTK.N	1	3.7999	0.4731	15	Q6N094
Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	2.8519	0.46	20	Q6N094
Q6N094_HUMAN	K.FNWYVDGVEVHNAK.T	1	5.0571	0.5892	20	Q6N094

IGHM protein - Homo sapiens (Human)

Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	2	4.3392	0.4953	4	Q6GMY2
Q6GMY2_HUMAN	K.YAATSQVLLPSKDVMQGTDE	1	6.3012	0.6623	3	Q6GMY2
Q6GMY2_HUMAN	R.EGKQVGSGVTTDQVQAEAK.	1	3.729	0.0937	4	Q6GMY2

Inter-alpha-trypsin inhibitor heavy chain H3 precursor (ITI heavy chain H3) (Inter-alpha-

ITIH3_HUMAN	R.LVDEDMNSFK.A	2	3.5178	0.5765		Q06033
ITIH3_HUMAN	R.DYIFGNYIER.L	1	3.2084	0.5288		Q06033
ITIH3_HUMAN	R.LVDEDM*NSFK.A	1	3.0464	0.4253		Q06033

Vitronectin precursor (Serum-spreading factor) (S-protein) (V75) [Contains: Vitronectin

VTNC_HUMAN	R.FEDGVLDPDYPR.N	1	3.2864	0.4262		P04004
VTNC_HUMAN	R.IYISGMAPR.P	1	3.1957	0.3745		P04004

VTNC_HUMAN	R.DVWGIEGPIDAAFTR.I	1	4.2401	0.4257		P04004
VTNC_HUMAN	R.DWHGVPGQVDAAM*AGR.I	1	3.3588	0.1622		P04004

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Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -

APOA2_HUMAN	K.VKSPELQAEAK.S	2	3.5731	0.3545		P02652
APOA2_HUMAN	K.SKEQLTPLIK.K	1	3.1002	0.4085		P02652

Apolipoprotein F precursor (Apo-F) (Lipid transfer inhibitor protein) (LTIP) - Homo sapie

APOF_HUMAN	R.SGVQQLIQYYQDQK.D	2	4.5022	0.5114		Q13790
APOF_HUMAN	K.SYDLDPGAGSLEI.-	1	2.6485	0.3479		Q13790

Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis

CLUS_HUMAN	R.ASSIIDELFQDR.F	1	3.7493	0.4862	1	P10909
CLUS_HUMAN	K.FM*ETVAEK.A	1	2.5768	0.241	1	P10909
CLUS_HUMAN	R.ELDESLQVAER.L	1	3.3064	0.354	1	P10909

Complement C1r subcomponent precursor (EC 3.4.21.41) (Complement component 1,

C1R_HUMAN	K.QRPPDLDTSSNAVDLLFFTDE	1	4.3028	0.4447		P00736
C1R_HUMAN	R.YTTTMGVNTYK.A	2	3.058	0.4901		P00736

Complement component C9 precursor [Contains: Complement component C9a; Compl

CO9_HUMAN	R.VVEESELAR.T	1	3.1114	0.4767		P02748
CO9_HUMAN	R.TEHYEEQIEAFK.S	1	3.373	0.5047		P02748
CO9_HUMAN	R.AIEDYINEFSVR.K	1	3.9139	0.5553		P02748

Hypothetical protein - Homo sapiens (Human)

Q6P5S3_HUMAN	K.ANPTVTLFPPSSEELQANK.A	2	4.1542	0.5369	10	Q6P5S3
Q6P5S3_HUMAN	K.AGVETTKPSK.Q	1	2.5983	0.2523	10	Q6P5S3

IGHA1 protein - Homo sapiens (Human)

Q8N5K4_HUMAN	K.TPLTATLSK.S	1	2.4973	0.4579	11	Q8N5K4
Q8N5K4_HUMAN	K.GTTVTVSSASPTSPK.V	1	2.9674	0.5291	1	Q8N5K4
Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	1	4.1428	0.5953	12	Q8N5K4

2

Apolipoprotein E precursor (Apo-E) - Homo sapiens (Human)

APOE_HUMAN	R.AATVGSLAGQPLQER.A	1	3.7601	0.4751		P02649
APOE_HUMAN	R.LGPLVEQGR.V	1	2.5078	0.3433		P02649

Apolipoprotein(a) precursor (EC 3.4.21.-) (Apo(a)) (Lp(a)) - Homo sapiens (Human)

APOA_HUMAN	R.GTYSTTVTGR.T	1	2.6927	0.4594	39	P08519
APOA_HUMAN	R.TPENYPNAGLTR.N	1	2.4386	0.1752	1	P08519

Fibulin-1 precursor - Homo sapiens (Human)

FBLN1_HUMAN	K.TGYYFDGISR.M	1	3.0621	0.4106		P23142
FBLN1_HUMAN	R.SAATLQQEK.T	1	2.6468	0.3794		P23142

Heparin cofactor 2 precursor (Heparin cofactor II) (HC-II) (Protease inhibitor leuserpin 2

HEP2_HUMAN	K.NYNLVESLK.L	1	2.4719	0.3412		P05546
HEP2_HUMAN	R.SVNDLYIQK.Q	1	3.0269	0.4432		P05546

Histidine-rich glycoprotein precursor (Histidine-proline-rich glycoprotein) (HPRG) - Hom

HRG_HUMAN	K.YKEENDDFASFR.V	1	3.7937	0.6083		P04196
HRG_HUMAN	K.DSPVLIDFFEDTER.Y	1	4.5195	0.575		P04196

Ig kappa chain V-IV region precursor (Fragment) - Homo sapiens (Human)

KV401_HUMAN	K.LLIYWASTR.E	1	3.3492	0.3688	4	P06312
KV401_HUMAN	K.SSQSVLYSSNNK.N	1	3.0441	0.3912	2	P06312

Ig lambda chain V-III region LOI - Homo sapiens (Human)

LV302_HUMAN	-.YVLTQPPSVSVAPGETAR.L	2	4.0006	0.4775		P80748

IGHG4 protein - Homo sapiens (Human)

Q8TC63_HUMAN	K.YGPPCPSCPAPEFLGGPSVF	1	4.0683	0.3936	1	Q8TC63

	Q8TC63_HUMAN	R.EPQVYTLPPSQEEMTK.N	1	2.4362	0.2292	1	Q8TC63

Myosin-reactive immunoglobulin light chain variable region (Fragment) - Homo sapiens

	Q9UL83_HUMAN	-.EIVM*TQSPATLSVSPGER.A	1	3.955	0.4591	1	Q9UL83
	Q9UL83_HUMAN	-.EIVMTQSPATLSVSPGER.A	1	4.8316	0.5908	1	Q9UL83

Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II) [Contains: Activation peptid

	THRB_HUMAN	R.TATSEYQTFFNPR.T	2	3.5803	0.5196		P00734

1

26S proteasome non-ATPase regulatory subunit 14 (26S proteasome regulatory subun

	PSDE_HUMAN	K.EMLELAKNYNKAVEEEDK.M	1	3.4661	0.2207		O00487

Actin-related protein 2/3 complex subunit 1A (SOP2-like protein) - Homo sapiens (Hum

	ARC1A_HUMAN	R.DRTQIALSPNNHEVHIYK.K	1	2.4562	0.0883		Q92747

Adenylate cyclase type 6 (EC 4.6.1.1) (Adenylate cyclase type VI) (ATP pyrophosphate

	ADCY6_HUMAN	R.FLLTFQREDLEKK.Y	1	2.5032	0.1113		O43306

Adipose differentiation-related protein - Homo sapiens (Human)

	Q5SYF3_HUMAN	R.IEERLPILNQPSTQVSLGQKC.	1	2.4897	0.0977		Q5SYF3

Alpha-1-acid glycoprotein 2 precursor (AGP 2) (Orosomucoid-2) (OMD 2) - Homo sapie

	A1AG2_HUMAN	R.SDVMYTDWK.K	1	3.174	0.4515		P19652

Aminopeptidase O (EC 3.4.11.-) (AP-O) - Homo sapiens (Human)

	AMPO_HUMAN	K.HYVLDLDVDFESQVIEGTIVLF	1	3.6182	0.1689		Q8N6M6

Angiotensinogen precursor [Contains: Angiotensin-1 (Angiotensin I) (Ang I); Angiotensi

	ANGT_HUMAN	R.ADSQAQLLLSTVVGVFTAPGL	1	5.2888	0.5924		P01019

Apolipoprotein C-III precursor (Apo-CIII) (ApoC-III) - Homo sapiens (Human)

	APOC3_HUMAN	K.DALSSVQESQVAQQAR.G	1	4.9755	0.3963		P02656

Apolipoprotein D precursor (Apo-D) (ApoD) - Homo sapiens (Human)

	APOD_HUMAN	R.NPNLPPETVDSLK.N	1	2.4472	0.3612		P05090

Apolipoprotein-L1 precursor (Apolipoprotein L-I) (Apolipoprotein L) (ApoL-I) (Apo-L) (Ap

	APOL1_HUMAN	R.VTEPISAESGEQVER.V	1	4.3195	0.5684		O14791

Basic helix-loop-helix domain containing, class B, 9 (P60-like protein) - Homo sapiens (

	Q6PI77_HUMAN	K.LLLNMSENPTAARDM*INM*K.	1	2.6218	0.0955		Q6PI77

Brefeldin A-inhibited guanine nucleotide-exchange protein 1 (Brefeldin A-inhibited GEP

	BIG1_HUMAN	K.QRRLLYNLEMEQMAK.T	1	2.465	0.1352	1	Q9Y6D6

C4B1 (Complement component 4B) (Childo blood group) - Homo sapiens (Human)

	Q6U2E9_HUMAN	R.GSSTWLTAFVLK.V	1	3.6371	0.4513		Q6U2E9

Casein kinase I isoform gamma-1 (EC 2.7.11.1) (CKI-gamma 1) - Homo sapiens (Huma

	KC1G1_HUMAN	K.RNTPIEALCENFPEEMATYLR	1	3.4684	0.3322	1	Q9HCP0

Catenin alpha-3 (Alpha T-catenin) (Cadherin-associated protein) - Homo sapiens (Hum

	CTNA3_HUMAN	K.M*ISESGSRMDVLAR.Q	1	2.4713	0.1267		Q9UI47

CDNA FLJ13244 fis, clone OVARC1000679, highly similar to Homo sapiens myosin-IX

	Q9H8T5_HUMAN	K.VFLDEYM*NEFKTSDCTATKV	1	3.2005	0.2713		Q9H8T5

CLIP-associating protein 2 (Cytoplasmic linker-associated protein 2) (hOrbit2) - Homo s

	CLAP2_HUMAN	K.VKVAILKYIETLAK.Q	1	2.9489	0.3212	1	O75122

Collagen type IX alpha 2 (Fragment) - Homo sapiens (Human)

	Q5QPF8_HUMAN	-.SPFRGPPGAAGAR.R	1	2.7687	0.2401		Q5QPF8

Contactin-1 precursor (Neural cell surface protein F3) (Glycoprotein gp135) - Homo sap

	CNTN1_HUMAN	R.WRKVLEPM*PSTAEISTSGAV	1	2.4518	0.2187		Q12860

Crumbs homolog 1 precursor - Homo sapiens (Human)

	CRUM1_HUMAN	K.IDWNHITLENISSGSSLNVK.A	1	2.9722	0.1379		P82279

DNA ligase 4 (EC 6.5.1.1) (DNA ligase IV) (Polydeoxyribonucleotide synthase [ATP] 4)

	DNL4_HUMAN	K.NEGTRLAIKALELR.F	1	3.0536	0.1861		P49917

Eomesodermin homolog - Homo sapiens (Human)

EOMES_HUMAN	R.RLSPSNSSNENSPSIK.C	1	2.428	0.1136		Q95936

ETS-related transcription factor Elf-1 (E74-like factor 1) - Homo sapiens (Human)

ELF1_HUMAN	-.M*AAVVQQNDLVFEFASNVM*I	1	2.4511	0.1836	2	P32519

Fetal beta-MHC binding factor - Homo sapiens (Human)

Q14297_HUMAN	K.KRIIYDSDSESEETVQVK.N	1	2.5178	0.2887		Q14297

Glutamate dehydrogenase 2, mitochondrial precursor (EC 1.4.1.3) (GDH) - Homo sapie

DHE4_HUMAN	R.SARQIM*HTAMKYNLGLDLR.T	1	3.3058	0.1232		P49448

Haptoglobin precursor [Contains: Haptoglobin alpha chain; Haptoglobin beta chain] - H

HPT_HUMAN	R.VGYVSGWGR.N	1	3.1823	0.413	2	P00738

Hypothetical protein DKFZp686E23209 - Homo sapiens (Human)

Q68CN4_HUMAN	K.NQVSLTCLVK.G	1	2.6481	0.4324	30	Q68CN4

Hypothetical protein DKFZp686I15212 - Homo sapiens (Human)

Q6N030_HUMAN	K.WYVDGVEVHNAK.T	1	2.6759	0.4194	4	Q6N030

Hypothetical protein FL25415 - Homo sapiens (Human)

Q7Z5J8_HUMAN	R.AM*KCKSIPFGMK.S	1	2.649	0.295		Q7Z5J8

Hypothetical protein pp12301 - Homo sapiens (Human)

Q8WYX2_HUMAN	R.AEVGGWLEPR.S	1	2.4279	0.0925		Q8WYX2

Ig heavy chain V-III region JON - Homo sapiens (Human)

HV319_HUMAN	-.DVQLVESGGGLVKPGGSLR.L	1	2.636	0.149		P01780

Ig heavy chain V-III region TIL - Homo sapiens (Human)

HV304_HUMAN	-.EVQLLESGGGLVQPGGSLR.L	1	4.8803	0.3725	5	P01765

Ig kappa chain V-III region SIE - Homo sapiens (Human)

KV302_HUMAN	R.FSGSGSGTDFTLTISR.L	1	4.0281	0.6388	12	P01620

Ig kappa chain V-IV region Len - Homo sapiens (Human)

KV402_HUMAN	-.DIVMTQSPDSLAVSLGER.A	1	2.9568	0.3584		P01625

Ig lambda chain V-III region SH - Homo sapiens (Human)

LV301_HUMAN	-.SELTQDPAVSVALGQTVR.I	1	2.9339	0.275		P01714

IGHG1 protein - Homo sapiens (Human)

Q6PJ95_HUMAN	R.GTLVTVSSASTK.G	1	2.9184	0.2997		Q6PJ95

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.0961	0.3203		P01591

Integrin alpha-E precursor (Mucosal lymphocyte 1 antigen) (HML-1 antigen) (Integrin al

ITAE_HUMAN	R.FAIGVGEEFKSAR.T	1	2.5436	0.2094		P38570

Kinesin heavy chain isoform 5A (Neuronal kinesin heavy chain) (NKHC) (Kinesin heavy

KIF5A_HUMAN	K.LEAELSRWR.N	1	2.4136	0.146		Q12840

KNG1 protein - Homo sapiens (Human)

Q05CF8_HUMAN	K.AGAEPASEREVS.-	1	2.9131	0.4265		Q05CF8

Leucine-rich repeat and fibronectin type III domain-containing protein 1 (Fragment) - H

Q9P244_HUMAN	R.PQQRYSFDGDYGALFQSHSY	1	3.2362	0.114		Q9P244

Low-density lipoprotein receptor-related protein 2 precursor (Megalin) (Glycoprotein 33

LRP2_HUMAN	R.CDRHNDCGDYSDER.G	1	2.5889	0.2861		P98164

Lysosomal-trafficking regulator (Beige homolog) - Homo sapiens (Human)

LYST_HUMAN	K.DKAASESIR.V	1	2.1447	0.3083		Q99698

MRNA clone with similarity to L-glycerol-3-phosphate:NAD oxidoreductase and albumin

Q13138_HUMAN	-.M*SVLM*GANIASEVADEK.F	1	2.6931	0.0971		Q13138

Myosin-reactive immunoglobulin heavy chain variable region (Fragment) - Homo sapier

Q9UL94_HUMAN	-.EVQLVESGAEVK.K	1	4.1798	0.4812	2	Q9UL94

Myotubularin-related protein 4 (FYVE domain-containing dual specificity protein phosph

Q8IV27_HUMAN	R.SM*DDLLSACDTSSPLTR.T	1	2.5071	0.1381		Q8IV27

N-acetylglutamate synthase, mitochondrial precursor (EC 2.3.1.1) (Amino-acid acetyltra

NAGS_HUMAN	R.SLVQRDIQAFLNQCGASPGE/	1	2.4605	0.1971		Q8N159

NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial precursor (EC 1

NDUS2_HUMAN	R.AEM*KTSM*ESLIHHFK.L	1	3.315	0.1556		O75306

Nesprin-2 (Nuclear envelope spectrin repeat protein 2) (Syne-2) (Synaptic nuclear enve

SYNE2_HUMAN	R.KIQHLEQLLESITESENK.I	1	2.8835	0.1425		Q8WXH0

Non-structural maintenance of chromosomes element 1 homolog (Non-SMC element 1

NSE1_HUMAN	K.FVQNKWLIEK.E	1	2.4635	0.1302		Q8WV22

Novel protein similar to hemicentin (LOC392395) (Fragment) - Homo sapiens (Human)

A2A3K3_HUMAN	R.ASVWLLVR.E	1	2.4768	0.1172		A2A3K3

P2X purinoceptor 6 (ATP receptor) (P2X6) (Purinergic receptor) (P2XM) (Purinergic rec

P2RX6_HUMAN	R.LLQFGIVVYVVGWALLAK.K	1	3.2257	0.2436		O15547

PHACTR4 protein - Homo sapiens (Human)

Q6NUN6_HUMAN	K.LNHRPSEPELNLNSWPCKSK	1	3.4059	0.1456	2	Q6NUN6

Protein Z-dependent protease inhibitor precursor (PZ-dependent protease inhibitor) (PZ

ZPI_HUMAN	R.IFSPFADLSELSATGR.N	1	3.2171	0.596		Q9UK55

Pyruvate kinase isozymes M1/M2 (EC 2.7.1.40) (Pyruvate kinase muscle isozyme) (Pyr

KPYM_HUMAN	K.ITLDNAYMEKCDENILWLDYK	1	3.6846	0.2175	1	P14618

RcADH5 (Fragment) - Homo sapiens (Human)

Q2VIM7_HUMAN	K.FARAKEFGATECINPQGFSK.I	1	3.2711	0.1739		Q2VIM7

Receptor-type tyrosine-protein phosphatase kappa precursor (EC 3.1.3.48) (Protein-tyr

PTPRK_HUMAN	R.KGYNIYFQAM*SSVEKETK.T	1	2.4473	0.1347	4	Q15262

Rheumatoid factor D5 light chain (Fragment) - Homo sapiens (Human)

A0N5G5_HUMAN	-.EIVLTQSPATLSLSPGER.A	1	3.9939	0.5844		A0N5G5

Rho GTPase-activating protein 29 (PTPL1-associated Rho GTPase-activating protein 1

O15463_HUMAN	K.MNSKNLGVIFGPSLIR.P	1	3.3165	0.1942		O15463

Ribosomal RNA upstream binding transcription factor (Fragment) - Homo sapiens (Hun

O00164_HUMAN	K.TTTRM*TMR.M	1	2.638	0.2708		O00164

RNA polymerase-associated protein RTF1 homolog - Homo sapiens (Human)

RTF1_HUMAN	K.DKDLNSKSASDLSEDLFK.V	1	2.7092	0.1737		Q92541

Selenoprotein P precursor (SeP) - Homo sapiens (Human)

SEPP1_HUMAN	K.LPTDSELAPR.S	1	2.565	0.4091		P49908

Short transient receptor potential channel 3 (TrpC3) (Htrp-3) (Htrp3) - Homo sapiens (H

TRPC3_HUMAN	K.QDISSLRYELLEDK.S	1	2.4231	0.0937	1	Q13507

Solute carrier family 22 member 11 (Organic anion transporter 4) - Homo sapiens (Hum

OAT4_HUMAN	R.INGHKEAKNLTIEVLMSSVK.E	1	2.6246	0.1469	1	Q9NSA0

Succinyl-CoA ligase [GDP-forming] beta-chain, mitochondrial precursor (EC 6.2.1.4) (S

SUCB2_HUMAN	K.VPLVVRLEGTNVQEAQK.I	1	2.5292	0.1748	1	Q96I99

Synaptotagmin-like protein 5 - Homo sapiens (Human)

SYTL5_HUMAN	-.M*SKNSEFINLSFLLDHEK.E	1	2.5595	0.233		Q8TDW5

Transcription elongation factor B polypeptide 1 (RNA polymerase II transcription factor

ELOC_HUMAN	K.AMLSGPGQFAENETNEVNFR	1	2.4776	0.1316		Q15369

Transcription factor-like 5 protein (Cha transcription factor) (HPV-16 E2-binding protein

TCFL5_HUMAN	R.MMLLSEAGAAEKTSGGGDG/	1	3.2678	0.1332	1	Q9UL49

Tumor protein p73 (p53-like transcription factor) (p53-related protein) - Homo sapiens (

P73_HUMAN	R.QRVM*EAVHFRVR.H	1	2.4785	0.138		O15350

Uncharacterized protein C12orf52 (Fragment) - Homo sapiens (Human)

Q53GM3_HUMAN	R.AIHPAGPSKTEPGPAADSQK.	1	2.7129	0.0949		Q53GM3

Uncharacterized protein C9orf144 - Homo sapiens (Human)

Q6ZU69_HUMAN	K.DPFLFNELSFPQLLPK.T	1	3.3084	0.1028		Q6ZU69

Uracil-DNA glycosylase (EC 3.2.2.-) (UDG) - Homo sapiens (Human)

UNG_HUMAN	-.M*IGQKTLYSFFSPSPAR.K	1	2.5035	0.1067		P13051

V5-6 protein (Fragment) - Homo sapiens (Human)

Q5NV92_HUMAN	R.FSGSSSGAER.Y	1	2.5372	0.3321	1	Q5NV92

VH1 protein precursor (Fragment) - Homo sapiens (Human)

O95978_HUMAN	K.GPSVFPLAPCSR.S	1	2.7329	0.3025	11	O95978

von Willebrand factor precursor (vWF) [Contains: von Willebrand antigen 2 (von Willeb

VWF_HUMAN	K.GLYLETEAGYYK.L	1	2.5119	0.4371	1	P04275

Zinc finger and BTB domain-containing protein 22 (Zinc finger protein 297) (Protein BIN

ZBT22_HUMAN	R.HRLGGVGAVPGPGTPTGPSL	1	2.4507	0.0988	1	O15209

Zinc finger CCHC domain-containing protein 11 - Homo sapiens (Human)

ZCH11_HUMAN	R.M*DDFQLKGIVEEKFVK.W	1	2.4006	0.3023	1	Q5TAX3

Zinc finger protein 638 (Nuclear protein 220) (Zinc-finger matrin-like protein) (Cutaneou

ZN638_HUMAN	R.DITKQSQETEARPSIM*K.R	1	2.6796	0.1963	1	Q14966

Zinc finger, MYM-type 4 (Fragment) - Homo sapiens (Human)

Q5T5E1_HUMAN	K.PIVTINTNSISTK.C	1	2.6026	0.1306	1	Q5T5E1

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Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-

ITIH2_HUMAN	R.NDLISATKTQVADAKR.Y	1	3.5041	0.4864	1	P19823
ITIH2_HUMAN	R.MLADAPPQDPSCCSGALYYG	2	3.8781	0.5745	1	P19823
ITIH2_HUMAN	R.FLHVPDTFEGHFDGVPVISK.(	1	3.2089	0.4492	2	P19823
ITIH2_HUMAN	R.SSALDM*ENFR.T	1	2.6144	0.2459	2	P19823
ITIH2_HUMAN	R.MATTMIQSK.V	2	2.7744	0.5394	2	P19823
ITIH2_HUMAN	R.SSALDM*ENFRTEVNVLPGAK	1	4.0958	0.3378	2	P19823
ITIH2_HUMAN	R.SSIKEKTVGR.A	1	2.5765	0.2159	2	P19823
ITIH2_HUMAN	R.TEVNVLPGAK.V	2	2.9321	0.5081	2	P19823
ITIH2_HUMAN	R.TWRNDLISATKTQVADAK.R	1	4.6325	0.5939	1	P19823
ITIH2_HUMAN	R.YIEKIQPSGGTNINEALLR.A	1	5.4076	0.4787	1	P19823
ITIH2_HUMAN	K.VVNNSPQPQNVVFDVQIPK.G	3	5.5189	0.5838	2	P19823
ITIH2_HUMAN	K.IQPSGGTNINEALLR.A	1	2.7373	0.474	1	P19823
ITIH2_HUMAN	K.AGELEVFNGYFVHFFAPDNLI	2	6.0186	0.6118	1	P19823
ITIH2_HUMAN	K.AHVSFKPTVAQQR.I	2	3.6466	0.653	2	P19823
ITIH2_HUMAN	K.FYNQVSTPLLR.N	3	3.3421	0.4424	1	P19823
ITIH2_HUMAN	K.HADPDFTR.K	1	2.5453	0.2499	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLKK.F	1	3.7726	0.4463	1	P19823
ITIH2_HUMAN	K.VQFELHYQEVK.W	1	3.7944	0.3312	2	P19823
ITIH2_HUMAN	R.MATTM*IQSK.V	1	2.9539	0.4924	2	P19823
ITIH2_HUMAN	R.AEDHFSVIDFNQNIR.T	3	4.734	0.5106	1	P19823
ITIH2_HUMAN	R.ETAVDGELVVLYDVK.R	1	3.7825	0.5206	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLK.K	1	3.5248	0.3853	1	P19823
ITIH2_HUMAN	R.IYLQPGRLAK.H	1	2.4259	0.1464	2	P19823
ITIH2_HUMAN	R.LSNENHGIAQR.I	1	3.8566	0.4369	1	P19823
ITIH2_HUMAN	R.M*ATTM*IQSK.V	2	2.7928	0.46	2	P19823
ITIH2_HUMAN	K.KFYNQVSTPLLR.N	1	3.3843	0.5348	1	P19823
ITIH2_HUMAN	K.TVGRALYAQAR.A	1	2.7259	0.2828	2	P19823

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Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEM*	1	3.22	0.5322	P00450
CERU_HUMAN	K.MYYSAVDPTK.D	1	3.2211	0.542	P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPM*K	1	4.0388	0.5273	P00450
CERU_HUMAN	K.NNEGTYYSPNYNPQSR.S	1	4.0667	0.5753	P00450
CERU_HUMAN	K.VDKDNEDFQESNR.M	1	3.3991	0.2614	P00450
CERU_HUMAN	K.VNKDDEEFIESNK.M	1	4.416	0.3636	P00450
CERU_HUMAN	K.WYLFGMGNEVDVHAAFFHG(	1	5.5071	0.5384	P00450
CERU_HUMAN	K.M*YYSAVDPTKDIFTGLIGPMK	1	4.776	0.5259	P00450
CERU_HUMAN	R.FNKNNEGTYYSPNYNPQSR.S	1	4.6774	0.546	P00450
CERU_HUMAN	R.EYTDASFTNR.K	1	2.9746	0.5085	P00450
CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEMF	2	5.0619	0.6325	P00450
CERU_HUMAN	R.HYYIAAEEIIWNYAPSGIDIFTK	2	6.1043	0.538	P00450
CERU_HUMAN	R.IYHSHIDAPK.D	1	2.4952	0.3266	P00450
CERU_HUMAN	R.M*FTTAPDQVDKEDEDFQESI	1	3.6744	0.5139	P00450
CERU_HUMAN	R.MFTTAPDQVDKEDEDFQESN	1	3.6808	0.5136	P00450
CERU_HUMAN	R.QSEDSTFYLGER.T	1	3.1407	0.3037	P00450
CERU_HUMAN	R.RQSEDSTFYLGER.T	1	3.9411	0.4426	P00450
CERU_HUMAN	R.TTIEKPVWLGFLGPIIK.A	1	3.5758	0.3707	P00450
CERU_HUMAN	K.ALYLQYTDETFR.T	1	3.8859	0.4773	P00450
CERU_HUMAN	R.GPEEEHLGILGPVIWAEVGDT	1	6.2697	0.3884	P00450
CERU_HUMAN	K.LISVDTEHSNIYLQNGPDR.I	1	3.9109	0.4612	P00450
CERU_HUMAN	K.HYYIGIIETTWDYASDHGEK.K	1	6.3779	0.6495	P00450
CERU_HUMAN	K.GAYPLSIEPIGVR.F	1	2.7902	0.48	P00450
CERU_HUMAN	K.DVDKEFYLFPTVFDENESLLLI	1	4.5011	0.426	P00450
CERU_HUMAN	K.DNEDFQESNR.M	1	3.2689	0.3686	P00450
CERU_HUMAN	K.DIFTGLIGPMK.I	1	3.0047	0.3554	P00450
CERU_HUMAN	K.DIFTGLIGPM*K.I	1	2.4105	0.1466	P00450
CERU_HUMAN	K.DDEEFIESNK.M	1	3.2952	0.439	P00450
CERU_HUMAN	K.M*YYSAVDPTK.D	1	2.5848	0.3368	P00450
CERU_HUMAN	K.AETGDKVYVHLK.N	1	3.2214	0.3758	P00450
CERU_HUMAN	K.AEEEHLGILGPQLHADVGDK.\	1	4.1972	0.5292	P00450
CERU_HUMAN	K.M*YYSAVDPTKDIFTGLIGPM*	1	3.9448	0.4931	P00450
CERU_HUMAN	K.DIASGLIGPLIICK.K	2	4.3401	0.6494	P00450

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Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alpl

CO3_HUMAN	R.ILLQGTPVAQM*TEDAVDAER	1	2.9235	0.4754	P01024
CO3_HUMAN	K.RIPIEDGSGEVVLSRK.V	1	4.5227	0.5182	P01024
CO3_HUMAN	K.QELSEAEQATR.T	1	2.7298	0.3657	P01024
CO3_HUMAN	K.QLYNVEATSYALLALLQLK.D	1	5.5373	0.5384	P01024
CO3_HUMAN	R.SNLDEDIIAEENIVSR.S	2	5.1859	0.5862	P01024
CO3_HUMAN	R.SEETKENEGFTVTAEGK.G	1	4.7076	0.4226	P01024
CO3_HUMAN	R.TKKQELSEAEQATR.T	1	4.7115	0.5599	P01024
CO3_HUMAN	R.ILLQGTPVAQMTEDAVDAER.I	2	5.7638	0.6789	P01024
CO3_HUMAN	K.SGSDEVQVGQQR.T	1	4.19	0.3442	P01024
CO3_HUMAN	R.HQQTVTIPPK.S	1	2.5841	0.3512	P01024
CO3_HUMAN	K.AGDFLEANYMNLQR.S	3	4.3313	0.4882	P01024
CO3_HUMAN	K.EDIPPADLSDQVPDTESETR.I	2	3.7817	0.557	P01024
CO3_HUMAN	K.EYVLPSFEVIVEPTEK.F	1	2.5923	0.362	P01024
CO3_HUMAN	K.GYTQQLAFRQPSSAFAAFVK.	1	4.7648	0.497	P01024
CO3_HUMAN	K.KQELSEAEQATR.T	1	3.4753	0.3278	P01024
CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	2.7258	0.1234	P01024
CO3_HUMAN	R.VPVAVQGEDTVQSLTQGDGV	1	5.1963	0.7144	P01024

CO3_HUMAN	K.VLLDGVQNPR.A	1	3.4662	0.573		P01024
CO3_HUMAN	K.RIPIEDGSGEVVLSR.K	1	3.7351	0.4347		P01024
CO3_HUMAN	R.YYGGGYGSTQATFMVFQALA	1	4.0807	0.3507		P01024
CO3_HUMAN	R.EGVQKEDIPPADLSDQVPDTE	3	5.2856	0.59		P01024

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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H

APOA1_HUMAN	K.ETEGLRQEM*SK.D	1	2.7655	0.3825		P02647
APOA1_HUMAN	K.VSFLSALEEYTK.K	1	4.6188	0.5201		P02647
APOA1_HUMAN	K.VQPYLDDFQKK.W	1	2.7915	0.2288		P02647
APOA1_HUMAN	K.VEPLRAELQEGAR.Q	3	2.9112	0.2793		P02647
APOA1_HUMAN	K.LREQLGPVTQEFWDNLEK.E	1	5.5864	0.5742		P02647
APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	5	4.179	0.6194		P02647
APOA1_HUMAN	K.ATEHLSTLSEK.A	2	3.2232	0.4615		P02647
APOA1_HUMAN	K.KWQEEM*ELYR.Q	1	2.7015	0.4586		P02647
APOA1_HUMAN	R.QEM*SKDLEEVK.A	1	2.4179	0.1487		P02647
APOA1_HUMAN	R.THLAPYSDELR.Q	2	2.8422	0.5336		P02647
APOA1_HUMAN	R.QGLLPVLESFK.V	1	2.5099	0.1701		P02647
APOA1_HUMAN	K.WQEEM*ELYR.Q	1	3.0301	0.412		P02647
APOA1_HUMAN	R.DYVSQFEGSALGK.Q	1	4.3233	0.6111		P02647
APOA1_HUMAN	K.DSGRDYVSQFEGSALGK.Q	1	4.8341	0.512		P02647
APOA1_HUMAN	R.EQLGPVTQEFWDNLEK.E	2	4.3223	0.6572		P02647

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Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-

ITIH1_HUMAN	R.NHMQYEIVIK.V	1	3.1952	0.376		P19827
ITIH1_HUMAN	R.QAVDTAVDGVFIR.S	2	3.8576	0.3897		P19827
ITIH1_HUMAN	R.KAAISGENAGLVR.A	1	3.928	0.4798		P19827
ITIH1_HUMAN	R.IYEDHDATQQLQGFYSQVAK.	1	4.3672	0.333	1	P19827
ITIH1_HUMAN	R.IADNKQSSFK.A	1	3.055	0.4421	1	P19827
ITIH1_HUMAN	R.GHMLENHVER.L	1	3.0745	0.5011	1	P19827
ITIH1_HUMAN	R.RTFVLSALQPSPTHSSSNTQF	1	3.7329	0.2596	1	P19827
ITIH1_HUMAN	K.AAISGENAGLVR.A	1	3.8435	0.4867		P19827
ITIH1_HUMAN	R.ERGHMLENHVER.L	1	3.383	0.3872	1	P19827
ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	5	4.2898	0.5559	1	P19827
ITIH1_HUMAN	K.GSLVQASEANLQAAQDFVR.C	2	6.6478	0.6669		P19827
ITIH1_HUMAN	K.ILGDMQPGDYFDLVLFGTR.V	1	5.1778	0.6129		P19827
ITIH1_HUMAN	R.EVAFDLEIPK.T	1	2.6577	0.4386		P19827
ITIH1_HUMAN	K.LDAQASFLPK.E	2	3.6937	0.3315		P19827

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Inter-alpha-trypsin inhibitor heavy chain H3 precursor (ITI heavy chain H3) (Inter-alpha-

ITIH3_HUMAN	R.FAHNVVTMR.A	1	2.7646	0.3475		Q06033
ITIH3_HUMAN	R.DYIFGNYIER.L	1	2.5602	0.3856		Q06033
ITIH3_HUMAN	K.SMEDKGMTNINDGLLR.G	2	4.2129	0.5338		Q06033
ITIH3_HUMAN	K.SM*EDKGMTNINDGLLR.G	1	3.9832	0.1095		Q06033
ITIH3_HUMAN	K.GMTNINDGLLR.G	1	3.2815	0.2333		Q06033
ITIH3_HUMAN	K.ALQERDYIFGNYIER.L	1	3.6437	0.4662		Q06033
ITIH3_HUMAN	R.LVDEDMNSFK.A	1	3.3033	0.508		Q06033
ITIH3_HUMAN	R.LVDEDMNSFKADVK.G	2	4.1082	0.4389		Q06033
ITIH3_HUMAN	R.SLPEGVANGIEVYSTK.I	2	4.1435	0.5732		Q06033
ITIH3_HUMAN	R.LVDEDM*NSFK.A	1	3.4448	0.4596		Q06033
ITIH3_HUMAN	K.AVSQGKTAGLVK.A	1	2.8527	0.378		Q06033

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Serum albumin precursor - Homo sapiens (Human)

ALBU_HUMAN	R.FKDLGEENFK.A	1	2.5611	0.217	2	P02768
ALBU_HUMAN	R.YTKKVPQVSTPTLVEVSR.N	1	4.284	0.5055	2	P02768
ALBU_HUMAN	R.RHPYFYAPELLFFAK.R	1	4.976	0.4894	1	P02768
ALBU_HUMAN	R.RHPDYSVVLLLR.L	1	4.1363	0.3765	1	P02768
ALBU_HUMAN	K.VPQVSTPTLVEVSR.N	1	3.2789	0.5043	2	P02768
ALBU_HUMAN	K.TPVSDRVTK.C	1	2.4732	0.2131	2	P02768
ALBU_HUMAN	K.KQTALVELVK.H	1	2.5916	0.4004	2	P02768
ALBU_HUMAN	K.AVMDDFAAFVEK.C	2	3.7959	0.5896	2	P02768
ALBU_HUMAN	K.AVM*DDFAAFVEK.C	1	3.6565	0.5647	2	P02768
ALBU_HUMAN	K.KVPQVSTPTLVEVSR.N	2	4.1777	0.6051	2	P02768

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Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	K.SHALQLNNR.Q	1	2.718	0.4437	3	P0C0L4
CO4A_HUMAN	K.ADGSYAAWLSR.D	1	2.7223	0.4981	3	P0C0L4
CO4A_HUMAN	K.AEFQDALEK.L	1	2.4961	0.3418	3	P0C0L4
CO4A_HUMAN	K.LNMGITDLQGLR.L	1	3.3964	0.5059	3	P0C0L4
CO4A_HUMAN	K.TEQWSTLPPETK.D	1	3.3432	0.448	3	P0C0L4
CO4A_HUMAN	K.VLSLAQEQVGGSPEK.L	1	4.5882	0.5913	3	P0C0L4
CO4A_HUMAN	R.EMSGSPASGIPVK.V	1	2.7882	0.2252	3	P0C0L4
CO4A_HUMAN	R.GHLFLQTDQPIYNPGQR.V	1	4.7246	0.3634	3	P0C0L4
CO4A_HUMAN	R.TTNIQGINLLFSSR.R	1	4.5155	0.5392	3	P0C0L4
CO4A_HUMAN	R.VFALDQK.M	1	2.5915	0.1934	3	P0C0L4
CO4A_HUMAN	K.LNM*GITDLQGLR.L	1	2.5929	0.4385	3	P0C0L4

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Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	K.DSTYSLSSTLTLSK.A	1	2.6737	0.4473	21	A0A5E5
A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	6	4.6175	0.6026	21	A0A5E5
A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	2	5.7341	0.6742	20	A0A5E5

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Ig alpha-2 chain C region - Homo sapiens (Human)

IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	2	3.8729	0.5346	21	P01877
IGHA2_HUMAN	R.WLQGSQELPR.E	1	2.9744	0.3431	23	P01877
IGHA2_HUMAN	R.GFSPKDVLVR.W	1	2.8553	0.2066	23	P01877
IGHA2_HUMAN	R.DASGATFTWTPSSGK.S	1	3.5943	0.5914	10	P01877
IGHA2_HUMAN	R.WLQGSQELPREK.Y	1	3.132	0.346	23	P01877

Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)

MUCB_HUMAN	R.VFAIPPSFASIFLTK.S	2	3.1533	0.5221	5	P04220
MUCB_HUMAN	R.DGFFGNPR.K	1	2.5819	0.3389	5	P04220
MUCB_HUMAN	K.YVTSAPMPEPQAPGR.Y	1	3.6408	0.5394	5	P04220
MUCB_HUMAN	K.LICQATGFSPR.Q	1	3.195	0.5084	5	P04220
MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	1	4.098	0.4257	5	P04220

Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	K.KYNSQNQSNNQFVLYR.I	1	3.342	0.3446		P01042
KNG1_HUMAN	K.YNSQNQSNNQFVLYR.I	1	4.1565	0.491		P01042
KNG1_HUMAN	K.TVGSDTFYSFK.Y	1	3.1324	0.4782		P01042
KNG1_HUMAN	K.KYFIDFVAR.E	1	3.207	0.3921	1	P01042
KNG1_HUMAN	K.AATGECTATVGKR.S	1	2.5896	0.5163		P01042
KNG1_HUMAN	K.YFIDFVAR.E	1	2.4007	0.3726	1	P01042

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Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	K.FNWYVDGVEVHNAK.T	1	5.01	0.5468	20	Q6N094
Q6N094_HUMAN	K.VSNKALPAPIEKTISK.A	1	3.9401	0.4434	24	Q6N094
Q6N094_HUMAN	K.VSNKALPAPIEK.T	1	3.2937	0.3461	24	Q6N094
Q6N094_HUMAN	K.TTPPVLDSDGSFFLYSK.L	1	2.8457	0.4517	19	Q6N094
Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	2.809	0.3973	20	Q6N094

Lumican precursor (Keratan sulfate proteoglycan lumican) (KSPG lumican) - Homo sap

LUM_HUMAN	K.ISNIPDEYFK.R	1	2.9713	0.3354		P51884
LUM_HUMAN	R.NNQIDHIDEK.A	1	3.3039	0.3087		P51884
LUM_HUMAN	R.FNALQYLR.L	1	2.8459	0.2834		P51884
LUM_HUMAN	K.RFNALQYLR.L	1	3.6585	0.3249		P51884
LUM_HUMAN	K.SLEDLQLTHNK.I	1	2.9502	0.3713		P51884

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Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)

A2MG_HUMAN	R.VGFYESDVMGR.G	1	3.5296	0.5215		P01023
A2MG_HUMAN	R.SSGSLLNNAIK.G	1	2.8061	0.2666	2	P01023
A2MG_HUMAN	K.DTVIKPLLVEPEGLEK.E	1	3.928	0.2964		P01023
A2MG_HUMAN	K.AIGYLNTGYQR.Q	1	2.7937	0.3476		P01023

Apolipoprotein B-100 precursor (Apo B-100) [Contains: Apolipoprotein B-48 (Apo B-48)

APOB_HUMAN	K.GNVATEISTER.D	1	2.7243	0.5333		P04114
APOB_HUMAN	K.NSEEFAAAMSR.Y	1	2.8829	0.1011		P04114
APOB_HUMAN	R.SPSQADINK.I	1	3.1368	0.3502		P04114
APOB_HUMAN	K.ALVEQGFTVPEIK.T	1	2.9936	0.4148		P04114

Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis

CLUS_HUMAN	R.ASSIIDELFQDR.F	1	3.813	0.5125	1	P10909
CLUS_HUMAN	K.FM*ETVAEKALQEYR.K	1	3.9544	0.4796	1	P10909
CLUS_HUMAN	R.VTTVASHTSDSDVPSGVTEV\	1	3.3325	0.3064		P10909
CLUS_HUMAN	R.RELDESLQVAER.L	1	3.1621	0.4634	1	P10909

Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S

SAMP_HUMAN	R.VGEYSLYIGR.H	1	3.6209	0.4889		P02743
SAMP_HUMAN	R.AYSLFSYNTQGR.D	1	4.1331	0.4475		P02743
SAMP_HUMAN	K.IVLGQEQDSYGGKFDR.S	1	4.4388	0.5523		P02743
SAMP_HUMAN	K.IVLGQEQDSYGGK.F	1	3.2995	0.482		P02743

Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk

PON1_HUMAN	R.IQNILTEEPK.V	2	3.6421	0.2234		P27169
PON1_HUMAN	K.IFFYDSENPPASEVLR.I	2	5.7075	0.5951		P27169

Vitronectin precursor (Serum-spreading factor) (S-protein) (V75) [Contains: Vitronectin

VTNC_HUMAN	R.DWHGVPGQVDAAMAGR.I	1	3.5161	0.3501		P04004
VTNC_HUMAN	R.DVWGIEGPIDAAFTR.I	1	3.9927	0.4187		P04004
VTNC_HUMAN	R.FEDGVLDPDYPR.N	2	4.0638	0.5264		P04004

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AMBP protein precursor [Contains: Alpha-1-microglobulin (Protein HC) (Complex-formi

AMBP_HUMAN	R.AFIQLWAFDAVK.G	1	4.0773	0.4822		P02760
AMBP_HUMAN	R.AFIQLWAFDAVKGK.C	2	3.5754	0.4304		P02760

Haptoglobin-related protein precursor - Homo sapiens (Human)

HPTR_HUMAN	R.TEGDGVYTLNDK.K	1	3.7554	0.4976	2	P00739
HPTR_HUMAN	K.GSFPWQAK.M	1	2.594	0.3497	3	P00739
HPTR_HUMAN	R.VGYVSGWGQSDNFK.L	1	3.0026	0.4906		P00739

IGLV2-14 protein - Homo sapiens (Human)

A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	4.7331	0.6171	29	A0A5C9

	A0A5C9_HUMAN	K.AGVETTTPSK.Q	2	3.186	0.3683	17	A0A5C9
2							
Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr							
	FETUA_HUMAN	R.HTFMGVVSLGSPSGEVSHPR	1	4.4754	0.5378		P02765
	FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.2658	0.444		P02765
Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -							
	APOA2_HUMAN	K.AGTELVNFLSYFVELGTQPAT	1	4.2802	0.508		P02652
	APOA2_HUMAN	K.VKSPELQAEAK.S	1	2.8377	0.314		P02652
Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co							
	C1S_HUMAN	K.VEDPESTLFGSVIR.Y	1	4.1359	0.5773		P09871
	C1S_HUMAN	K.TMQENSTPRED.-	1	3.5263	0.5241		P09871
Complement-activating component of Ra-reactive factor precursor (EC 3.4.21.-) (Ra-re							
	MASP1_HUMAN	R.APGELEHGLITFSTR.N	1	3.2582	0.4819	1	P48740
	MASP1_HUMAN	R.TGVITSPDFPNPYPK.S	1	4.3089	0.4875	2	P48740
Hypothetical protein DKFZp686C15213 - Homo sapiens (Human)							
	Q6MZU6_HUMAN	K.TTPPMLDSDGSFFLYSK.L	2	3.0566	0.5256	1	Q6MZU6
IGHA1 protein - Homo sapiens (Human)							
	Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	2	4.3007	0.5209	12	Q8N5K4
IGHG4 protein - Homo sapiens (Human)							
	Q8TC63_HUMAN	K.YGPPCPSCPAPEFLGGPSVF	2	4.3162	0.2631	1	Q8TC63
IGHM protein - Homo sapiens (Human)							
	Q6GMY2_HUMAN	K.YAATSQVLLPSK.D	1	3.2688	0.381	3	Q6GMY2
	Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	1	3.5728	0.578	4	Q6GMY2
Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-							
	ITIH4_HUMAN	R.GPDVLTATVSGK.L	1	3.5018	0.5418	1	Q14624
	ITIH4_HUMAN	K.AEAQAQYSAAVAK.G	1	4.7849	0.511	1	Q14624
Myosin-reactive immunoglobulin heavy chain variable region (Fragment) - Homo sapier							
	Q9UL94_HUMAN	-.EVQLVESGAEVK.K	2	4.0683	0.4836	2	Q9UL94
Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II) [Contains: Activation peptid							
	THRB_HUMAN	R.ELLESYIDGR.I	1	2.4766	0.5165		P00734
	THRB_HUMAN	R.TATSEYQTFFNPR.T	1	3.632	0.5593		P00734
1							
Adiponectin precursor (Adipocyte, C1q and collagen domain-containing protein) (30 kD							
	ADIPO_HUMAN	K.GDIGETGVPGAEGPR.G	1	3.0273	0.4112		Q15848
Apolipoprotein A-IV precursor (Apo-AIV) (ApoA-IV) - Homo sapiens (Human)							
	APOA4_HUMAN	R.TQVNTQAEQLR.R	1	3.0352	0.4069	1	P06727
Apolipoprotein D precursor (Apo-D) (ApoD) - Homo sapiens (Human)							
	APOD_HUMAN	K.VLNQELR.A	1	2.5101	0.1762		P05090
Apolipoprotein E precursor (Apo-E) - Homo sapiens (Human)							
	APOE_HUMAN	R.AATVGSLAGQPLQER.A	1	4.1129	0.5627		P02649
Apolipoprotein F precursor (Apo-F) (Lipid transfer inhibitor protein) (LTIP) - Homo sapie							
	APOF_HUMAN	R.SGVQQLIQYYQDQK.D	1	4.9573	0.4634		Q13790
Apolipoprotein(a) precursor (EC 3.4.21.-) (Apo(a)) (Lp(a)) - Homo sapiens (Human)							
	APOA_HUMAN	R.GTYSTTVTGR.T	1	2.9724	0.4457	39	P08519
Basic helix-loop-helix domain containing, class B, 9 (P60-like protein) - Homo sapiens (							
	Q6PI77_HUMAN	K.LLLNMSENPTAARDMINM*K.A	1	2.5985	0.2209		Q6PI77
CDNA FLJ32471 fis, clone SKNMC2000322, weakly similar to MAJOR CENTROMERE							
	Q96ME4_HUMAN	R.KAITLEMK.L	1	2.5761	0.096		Q96ME4
Cell division control protein 42 homolog precursor (G25K GTP-binding protein) - Homo							

CDC42_HUMAN	K.NKQKPITPETAEK.L	1	2.7211	0.2472		P60953

Centromeric protein E (CENP-E) - Homo sapiens (Human)

CENPE_HUMAN	K.FQESQEEIKSLTK.E	1	2.5151	0.1822	1	Q02224

Chromodomain-helicase-DNA-binding protein 8 (EC 3.6.1.-) (ATP-dependent helicase (

CHD8_HUMAN	R.IPVINKVDGTLLVGEDAPR.R	1	2.89	0.1966		Q9HCK8

Chromosome 9 open reading frame 142 - Homo sapiens (Human)

Q9BUH6_HUMAN	R.LRALTLGLAKR.V	1	2.6364	0.0967		Q9BUH6

Ciliary dynein heavy chain 7 - Homo sapiens (Human)

Q8WXX0_HUMAN	R.LDFDNDGM*VEADDLR.S	1	2.8601	0.1359		Q8WXX0

Collagen type IX alpha 2 (Fragment) - Homo sapiens (Human)

Q5QPF8_HUMAN	-.SPFRGPPGAAGAR.R	1	3.2364	0.1537		Q5QPF8

Complement component 4A (Rodgers blood group) - Homo sapiens (Human)

Q5JNX2_HUMAN	K.AEMADQAAAWLTR.Q	1	2.9762	0.353	1	Q5JNX2

Complement component C9 precursor [Contains: Complement component C9a; Compl

CO9_HUMAN	R.AIEDYINEFSVR.K	1	2.9034	0.3973		P02748

Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) - Homo sapiens (Human)

FINC_HUMAN	R.WSRPQAPITGYR.I	1	3.2159	0.4075	1	P02751

Ficolin-3 precursor (Collagen/fibrinogen domain-containing protein 3) (Collagen/fibrinog

FCN3_HUMAN	R.RQDGSVDFFR.S	1	2.6846	0.2293	1	O75636

Haptoglobin precursor [Contains: Haptoglobin alpha chain; Haptoglobin beta chain] - H

HPT_HUMAN	R.HYEGSTVPEK.K	1	2.523	0.3065	2	P00738

HECT, UBA and WWE domain-containing protein 1 (EC 6.3.2.-) (E3 ubiquitin protein lig

HUWE1_HUMAN	R.ELQLPSMSMLTSK.T	1	2.6009	0.1288	2	Q7Z6Z7

Heparin cofactor 2 precursor (Heparin cofactor II) (HC-II) (Protease inhibitor leuserpin 2

HEP2_HUMAN	K.TLEAQLTPR.V	1	3.0036	0.4042		P05546

Histidine-rich glycoprotein precursor (Histidine-proline-rich glycoprotein) (HPRG) - Hom

HRG_HUMAN	R.DGYLFQLLR.I	1	3.1929	0.1649		P04196

Hypothetical protein - Homo sapiens (Human)

Q6P5S3_HUMAN	K.ANPTVTLFPPSSEELQANK.A	1	2.9752	0.4715	10	Q6P5S3

Hypothetical protein DKFZp686E23209 - Homo sapiens (Human)

Q68CN4_HUMAN	R.EPQVYTLPPSR.E	1	2.4094	0.1179	28	Q68CN4

Hypothetical protein FL25415 - Homo sapiens (Human)

Q7Z5J8_HUMAN	R.AM*KCKSIPFGMK.S	1	2.6364	0.2184		Q7Z5J8

Ig heavy chain V-III region HIL - Homo sapiens (Human)

HV310_HUMAN	-.QVKLVQAGGGVVQPGR.S	1	3.0651	0.3446		P01771

Ig heavy chain V-III region TIL - Homo sapiens (Human)

HV304_HUMAN	-.EVQLLESGGGLVQPGGSLR.L	1	4.8836	0.3024	5	P01765

Ig kappa chain V-II region Cum - Homo sapiens (Human)

KV201_HUMAN	R.FSGSGSGTDFTLK.I	1	3.3547	0.5799	9	P01614

Ig kappa chain V-III region SIE - Homo sapiens (Human)

KV302_HUMAN	-.EIVLTQSPGTLSLSPGER.A	1	3.8722	0.4388	4	P01620

Ig kappa chain V-III region VG precursor (Fragment) - Homo sapiens (Human)

KV309_HUMAN	R.LLIYDASNR.A	1	2.6447	0.329	1	P04433

Ig lambda chain V-III region LOI - Homo sapiens (Human)

LV302_HUMAN	-.YVLTQPPSVSVAPGETAR.L	1	4.2178	0.5052		P80748

Ig lambda chain V-IV region Hil - Homo sapiens (Human)

LV403_HUMAN	-.SYELTQPPSVSVSPGQTAR.I	1	5.064	0.6278	1	P01717

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.2484	0.3165		P01591

KIAA1529 - Homo sapiens (Human)

Q5VV25_HUMAN	K.M*DESKEGSIQGLEEMQVER.	1	3.346	0.1416	1	Q5VV25

Large neutral amino acids transporter small subunit 2 (L-type amino acid transporter 2)

LAT2_HUMAN	K.GEYFWLEPK.N	1	2.9531	0.3508	3	Q9UHI5

Latrophilin-2 (Fragment) - Homo sapiens (Human)

Q9UJ49_HUMAN	-.MTGNYLLTNPLLR.P	1	2.9573	0.2862		Q9UJ49

Low-density lipoprotein receptor-related protein 6 precursor - Homo sapiens (Human)

LRP6_HUMAN	K.DGATELLLLAR.R	1	2.7099	0.0908		O75581

LW-1 (Heat shock transcription factor family, X linked 1) - Homo sapiens (Human)

Q9UBD0_HUMAN	R.SLSMARCEER.N	1	2.6242	0.223		Q9UBD0

MRNA clone with similarity to L-glycerol-3-phosphate:NAD oxidoreductase and albumin

Q13138_HUMAN	-.M*SVLM*GANIASEVADEK.F	1	2.4654	0.1366		Q13138

Myosin-reactive immunoglobulin light chain variable region (Fragment) - Homo sapiens

Q9UL83_HUMAN	-.EIVMTQSPATLSVSPGER.A	1	4.2188	0.5949	1	Q9UL83

Myotubularin-related protein 4 (FYVE domain-containing dual specificity protein phosph

Q8IV27_HUMAN	R.SM*DDLLSACDTSSPLTR.T	1	2.6669	0.1252		Q8IV27

PDZ domain-containing RING finger protein 4 (Ligand of Numb-protein X 4) (SEMACAI

PZRN4_HUMAN	K.TTEQGCSAESKEKVLEGSK.L	1	2.7356	0.2832		Q6ZMN7

Prostaglandin F2 receptor negative regulator precursor (Prostaglandin F2-alpha recept

FPRP_HUMAN	K.VSSKNIKSPR.Y	1	3.1948	0.2248		Q9P2B2

Protein S100-A7 (S100 calcium-binding protein A7) (Psoriasin) - Homo sapiens (Humar

S10A7_HUMAN	K.GTNYLADVFEK.K	1	2.6418	0.236		P31151

Ras-related protein Rab-23 - Homo sapiens (Human)

RAB23_HUMAN	K.MVVVGNGAVGK.S	1	2.4116	0.0955		Q9ULC3

Rheumatoid factor D5 light chain (Fragment) - Homo sapiens (Human)

A0N5G5_HUMAN	-.EIVLTQSPATLSLSPGER.A	1	4.739	0.5655		A0N5G5

Rhomboid 5 homolog 2 (Rhomboid family member 2) (Rhomboid veinlet-like protein 5)

Q6PJF5_HUMAN	R.SVLDATGQRCR.V	1	2.409	0.1536	2	Q6PJF5

Serum amyloid A-4 protein precursor (Constitutively expressed serum amyloid A protei

SAA4_HUMAN	K.EALQGVGDMGR.A	1	2.9208	0.3914		P35542

Transthyretin precursor (Prealbumin) (TBPA) (TTR) (ATTR) - Homo sapiens (Human)

TTHY_HUMAN	K.AADDTWEPFASGK.T	1	4.035	0.6265		P02766

VH1 protein precursor (Fragment) - Homo sapiens (Human)

O95978_HUMAN	K.GPSVFPLAPCSR.S	1	2.5089	0.3022	11	O95978

VPS29 protein - Homo sapiens (Human)

Q8N0S8_HUMAN	R.QGPGAGLGK.A	1	2.4154	0.1155		Q8N0S8

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Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	K.M*RPSTDTITVM*VENSHGLR.	1	5.3455	0.4948	3	P0C0L4
CO4A_HUMAN	K.VDFTLSSER.D	1	2.6247	0.4715	3	P0C0L4
CO4A_HUMAN	R.EAPKVVEEQESR.V	1	2.5773	0.5395	3	P0C0L4
CO4A_HUMAN	R.DFALLSLQVPLKDAK.S	1	3.8599	0.4941	3	P0C0L4
CO4A_HUMAN	R.AVGSGATFSHYYYMILSR.G	1	3.6985	0.6084	3	P0C0L4
CO4A_HUMAN	R.ALERGLQDEDGYR.M	1	2.5935	0.2722	3	P0C0L4
CO4A_HUMAN	K.YVLPNFEVK.I	1	2.4632	0.3441	3	P0C0L4
CO4A_HUMAN	K.VVEEQESR.V	4	2.5262	0.3543	3	P0C0L4
CO4A_HUMAN	K.VLSLAQEQVGGSPEK.L	1	4.731	0.5709	3	P0C0L4
CO4A_HUMAN	K.VGGNSKGTLKVLR.T	1	2.8159	0.2124	3	P0C0L4

CO4A_HUMAN	R.DKGQAGLQR.A	11	2.7415	0.2568	3	P0C0L4
CO4A_HUMAN	K.TFFRGLESQTK.L	1	2.4056	0.2849	3	P0C0L4
CO4A_HUMAN	K.TEQWSTLPPETK.D	1	2.8973	0.4007	3	P0C0L4
CO4A_HUMAN	K.PVQGVAYVR.F	1	2.3957	0.4719	3	P0C0L4
CO4A_HUMAN	K.MRPSTDTITVMVENSHGLR.V	1	5.427	0.6066	3	P0C0L4
CO4A_HUMAN	R.EELVYELNPLDHR.G	1	3.8013	0.4943	3	P0C0L4
CO4A_HUMAN	K.M*RPSTDTITVMVENSHGLR.\	2	4.748	0.3489	3	P0C0L4
CO4A_HUMAN	R.GLQDEDGYR.M	2	2.8566	0.5383	3	P0C0L4
CO4A_HUMAN	K.LVNGQSHISLSK.A	1	3.7374	0.4945	3	P0C0L4
CO4A_HUMAN	K.LNMGITDLQGLR.L	2	3.3918	0.4391	3	P0C0L4
CO4A_HUMAN	K.LGQYASPTAKR.C	1	3.1966	0.3869	3	P0C0L4
CO4A_HUMAN	K.ITQVLHFTK.D	1	2.4765	0.4482	4	P0C0L4
CO4A_HUMAN	K.AINEKLGQYASPTAK.R	2	4.3174	0.543	3	P0C0L4
CO4A_HUMAN	K.MRPSTDTITVM*VENSHGLR.\	1	3.6823	0.374	3	P0C0L4
CO4A_HUMAN	R.NVNFQKAINEK.L	1	3.6832	0.2978	3	P0C0L4
CO4A_HUMAN	R.YVSHFETEGPHVLLYFDSVPT	1	4.5682	0.5779	3	P0C0L4
CO4A_HUMAN	R.YIYGKPVQGVAYVR.F	1	3.9092	0.5945	3	P0C0L4
CO4A_HUMAN	R.VTASDPLDTLGSEGALSPGG\	1	4.5595	0.6608	3	P0C0L4
CO4A_HUMAN	R.VTASDPLDTLGSEGALSPGG\	2	4.2656	0.6063	3	P0C0L4
CO4A_HUMAN	R.TYNVLDMK.N	1	3.3693	0.4127	3	P0C0L4
CO4A_HUMAN	R.TTNIQGINLLFSSR.R	4	4.5898	0.5961	3	P0C0L4
CO4A_HUMAN	R.TLEIPGNSDPNMIPDGDFNSY	1	3.5815	0.63	3	P0C0L4
CO4A_HUMAN	R.SFFPENWLWR.V	1	2.6393	0.3122	3	P0C0L4
CO4A_HUMAN	R.RGHLFLQTDQPIYNPGQR.V	2	4.6473	0.5456	3	P0C0L4
CO4A_HUMAN	R.RCSVFYGAPSKSR.L	1	3.3448	0.3473	3	P0C0L4
CO4A_HUMAN	R.GLEEELQFSLGSK.I	1	4.2515	0.4403	3	P0C0L4
CO4A_HUMAN	R.NVNFQKAINEKLGQYASPTAK	1	4.5351	0.5425	3	P0C0L4
CO4A_HUMAN	R.EMSGSPASGIPVK.V	1	2.6797	0.3507	3	P0C0L4
CO4A_HUMAN	R.LTVAAPPSGGPGFLSIER.P	1	3.3765	0.4992	3	P0C0L4
CO4A_HUMAN	R.HLVPGAPFLLQALVR.E	2	3.8238	0.4893	3	P0C0L4
CO4A_HUMAN	R.GSFEFPVGDAVSK.V	1	2.8299	0.4985	3	P0C0L4
CO4A_HUMAN	R.GPEVQLVAHSPWLKDSLSR.T	1	3.4913	0.4631	3	P0C0L4
CO4A_HUMAN	R.GPEVQLVAHSPWLK.D	1	4.5666	0.5095	3	P0C0L4
CO4A_HUMAN	R.GLQDEDGYRM*K.F	1	2.6251	0.3491	3	P0C0L4
CO4A_HUMAN	K.AEFQDALEK.L	1	2.8694	0.3723	3	P0C0L4
CO4A_HUMAN	R.GLESQTKLVNGQSHISLSK.A	1	3.4579	0.1749	3	P0C0L4
CO4A_HUMAN	R.GHLFLQTDQPIYNPGQR.V	2	6.3752	0.5835	3	P0C0L4
CO4A_HUMAN	R.FGLLDEDGKKTFFR.G	1	3.3017	0.3036	3	P0C0L4
CO4A_HUMAN	R.RCSVFYGAPSK.S	1	2.8722	0.2106	3	P0C0L4
CO4A_HUMAN	K.AINEKLGQYASPTAKR.C	1	4.4912	0.5899	3	P0C0L4

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Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-

ITIH2_HUMAN	K.RLSNENHGIAQR.I	2	3.7835	0.4039	1	P19823
ITIH2_HUMAN	R.ETAVDGELVVLYDVK.R	1	3.8402	0.6318	1	P19823
ITIH2_HUMAN	R.FLHVPDTFEGHFDGVPVISK.(	1	3.3018	0.5154	2	P19823
ITIH2_HUMAN	K.VVNNSPQPQNVVFDVQIPK.G	1	4.9614	0.4964	2	P19823
ITIH2_HUMAN	K.KFYNQVSTPLLR.N	1	3.8649	0.5346	1	P19823
ITIH2_HUMAN	K.HADPDFTR.K	1	2.512	0.3838	1	P19823
ITIH2_HUMAN	K.GQQKAHVSFKPTVAQQR.I	1	4.0853	0.4042	2	P19823
ITIH2_HUMAN	K.FYNQVSTPLLR.N	1	3.4113	0.4264	1	P19823
ITIH2_HUMAN	R.MATTM*IQSK.V	1	2.9762	0.4598	2	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLK.K	1	3.5901	0.4237	1	P19823

ITIH2_HUMAN	R.IYGNQDTSSQLKK.F	2	3.8157	0.4006	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLKKFYNQVSTF	1	4.3299	0.4929	1	P19823
ITIH2_HUMAN	K.AHVSFKPTVAQQR.I	1	3.5895	0.6213	2	P19823
ITIH2_HUMAN	R.M*LADAPPQDPSCCSGALYY(	1	3.4703	0.5436	1	P19823
ITIH2_HUMAN	R.AEDHFSVIDFNQNIR.T	1	4.6563	0.288	1	P19823
ITIH2_HUMAN	R.LSNENHGIAQR.I	2	4.1516	0.3894	1	P19823
ITIH2_HUMAN	R.MATTMIQSKVVNNSPQPQNV	1	4.56	0.4861	2	P19823
ITIH2_HUMAN	R.MLADAPPQDPSCCSGALYYG	2	3.6164	0.4878	1	P19823
ITIH2_HUMAN	R.NDLISATKTQVADAKR.Y	1	3.7345	0.5161	1	P19823
ITIH2_HUMAN	R.SSALDMENFRTEVNVLPGAK.	1	4.0338	0.4172	2	P19823
ITIH2_HUMAN	R.SSIKEKTVGR.A	1	2.4603	0.1843	2	P19823
ITIH2_HUMAN	R.TWRNDLISATK.T	1	2.521	0.1842	1	P19823
ITIH2_HUMAN	R.TWRNDLISATKTQVADAK.R	1	4.2409	0.5001	1	P19823
ITIH2_HUMAN	R.YIEKIQPSGGTNINEALLR.A	1	5.4825	0.5517	1	P19823
ITIH2_HUMAN	R.MATTMIQSK.V	1	2.8126	0.5382	2	P19823

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Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alp

CO3_HUMAN	R.SNLDEDIIAEENIVSR.S	1	5.4557	0.5715	P01024
CO3_HUMAN	K.SDDKVTLEER.L	1	3.2588	0.2563	P01024
CO3_HUMAN	K.SGSDEVQVGQQR.T	1	3.3117	0.3516	P01024
CO3_HUMAN	K.VLLDGVQNPR.A	1	3.3183	0.5598	P01024
CO3_HUMAN	R.EGVQKEDIPPADLSDQVPDTE	3	5.5142	0.5336	P01024
CO3_HUMAN	R.HQQTVTIPPK.S	1	2.505	0.3936	P01024
CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	3.1751	0.4506	P01024
CO3_HUMAN	R.VPVAVQGEDTVQSLTQGDGV	2	4.4772	0.6429	P01024
CO3_HUMAN	K.RIPIEDGSGEVVLSRK.V	1	4.4831	0.5354	P01024
CO3_HUMAN	R.SEETKENEGFTVTAEGK.G	1	4.4223	0.3828	P01024
CO3_HUMAN	R.ILLQGTPVAQM*TEDAVDAER	1	3.3808	0.4497	P01024
CO3_HUMAN	K.KQELSEAEQATR.T	2	3.718	0.3724	P01024
CO3_HUMAN	K.GYTQQLAFRQPSSAFAAFVK.	1	4.5889	0.5592	P01024
CO3_HUMAN	K.AGDFLEANYMNLQR.S	1	4.1106	0.4254	P01024
CO3_HUMAN	K.AGDFLEANYM*NLQR.S	1	2.6288	0.2629	P01024
CO3_HUMAN	R.TKKQELSEAEQATR.T	2	4.237	0.5487	P01024
CO3_HUMAN	K.QELSEAEQATR.T	1	2.7646	0.3225	P01024

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Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	K.NNEGTYYSPNYNPQSR.S	1	4.7426	0.606	P00450
CERU_HUMAN	K.GAYPLSIEPIGVR.F	1	3.5718	0.4988	P00450
CERU_HUMAN	K.HYYIGIIETTWDYASDHGEK.K	1	6.6025	0.6253	P00450
CERU_HUMAN	K.LVYREYTDASFTNRK.E	1	3.5225	0.4532	P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPM*H	1	3.6968	0.5466	P00450
CERU_HUMAN	K.VNKDDEEFIESNK.M	1	4.1833	0.3575	P00450
CERU_HUMAN	R.EYTDASFTNR.K	1	2.8128	0.4277	P00450
CERU_HUMAN	R.M*FTTAPDQVDKEDEDFQESI	1	4.0628	0.5182	P00450
CERU_HUMAN	K.M*YYSAVDPTKDIFTGLIGPMH	1	4.2403	0.3812	P00450
CERU_HUMAN	R.QSEDSTFYLGER.T	2	3.2006	0.4715	P00450
CERU_HUMAN	R.EYTDASFTNRK.E	1	2.5209	0.3365	P00450
CERU_HUMAN	R.IYHSHIDAPK.D	1	2.4698	0.3693	P00450
CERU_HUMAN	K.ALYLQYTDETFR.T	1	4.0149	0.5188	P00450
CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEMF	1	4.5534	0.564	P00450
CERU_HUMAN	R.FNKNNEGTYYSPNYNPQSR.S	1	4.9563	0.5036	P00450

CERU_HUMAN	R.TTIEKPVWLGFLGPIIK.A	2	3.5467	0.4755		P00450

Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-

ITIH1_HUMAN	K.GSLVQASEANLQAAQDFVR.C	2	6.1873	0.6434		P19827
ITIH1_HUMAN	R.GMADQDGLKPTIDKPSEDSPI	1	4.2484	0.5302	1	P19827
ITIH1_HUMAN	R.EVAFDLEIPK.T	1	2.7124	0.4066		P19827
ITIH1_HUMAN	R.ERGHMLENHVER.L	1	3.6744	0.3971	1	P19827
ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	1	4.0929	0.4873	1	P19827
ITIH1_HUMAN	K.LDAQASFLPKELAAQTIKK.S	1	3.3189	0.4358		P19827
ITIH1_HUMAN	K.ILGDM*QPGDYFDLVLFGTR.V	1	4.1008	0.4885		P19827
ITIH1_HUMAN	R.GMADQDGLKPTIDKPSEDSPI	1	4.9796	0.4231	1	P19827
ITIH1_HUMAN	K.AAISGENAGLVR.A	1	3.7029	0.4929		P19827
ITIH1_HUMAN	R.IADNKQSSFK.A	1	3.0309	0.4321	1	P19827
ITIH1_HUMAN	R.KAAISGENAGLVR.A	2	4.1054	0.47		P19827
ITIH1_HUMAN	R.QAVDTAVDGVFIR.S	3	3.3237	0.4209		P19827
ITIH1_HUMAN	K.LDAQASFLPK.E	2	3.5069	0.3116		P19827

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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H

APOA1_HUMAN	K.ATEHLSTLSEK.A	1	3.1331	0.4471		P02647
APOA1_HUMAN	K.DSGRDYVSQFEGSALGK.Q	1	2.6999	0.1416		P02647
APOA1_HUMAN	K.ETEGLRQEM*SK.D	1	2.438	0.3291		P02647
APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	3	4.1332	0.6337		P02647
APOA1_HUMAN	K.LREQLGPVTQEFWDNLEK.E	1	5.1349	0.5009		P02647
APOA1_HUMAN	K.QLNLKLLDNWDSVTSTFSK.L	1	3.6643	0.5707		P02647
APOA1_HUMAN	K.VEPLRAELQEGAR.Q	2	2.8214	0.2719		P02647
APOA1_HUMAN	K.VQPYLDDFQK.K	1	2.4122	0.2945		P02647
APOA1_HUMAN	K.VQPYLDDFQKK.W	1	2.6481	0.1298		P02647
APOA1_HUMAN	K.VSFLSALEEYTK.K	1	4.3079	0.4842		P02647
APOA1_HUMAN	R.LAEYHAKATEHLSTLSEK.A	1	4.5138	0.5435		P02647
APOA1_HUMAN	R.EQLGPVTQEFWDNLEK.E	2	4.3563	0.5978		P02647

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Serum albumin precursor - Homo sapiens (Human)

ALBU_HUMAN	K.AEFAEVSK.L	1	2.6498	0.1925	1	P02768
ALBU_HUMAN	R.FKDLGEENFK.A	1	2.7715	0.2678	2	P02768
ALBU_HUMAN	K.KQTALVELVK.H	1	2.7693	0.3609	2	P02768
ALBU_HUMAN	K.AVM*DDFAAFVEK.C	1	3.4279	0.5199	2	P02768
ALBU_HUMAN	K.VFDEFKPLVEEPQNLIK.Q	1	4.4292	0.4409	2	P02768
ALBU_HUMAN	K.TPVSDRVTK.C	1	2.436	0.3475	2	P02768
ALBU_HUMAN	R.AFKAWAVAR.L	1	2.5932	0.3575	1	P02768
ALBU_HUMAN	R.LAKTYETTLEK.C	1	2.7136	0.4055	1	P02768
ALBU_HUMAN	R.RHPDYSVVLLLR.L	1	4.6035	0.4053	1	P02768
ALBU_HUMAN	R.RHPYFYAPELLFFAK.R	1	3.2116	0.2868	1	P02768
ALBU_HUMAN	R.YTKKVPQVSTPTLVEVSR.N	1	3.8278	0.5115	2	P02768
ALBU_HUMAN	K.KVPQVSTPTLVEVSR.N	3	4.6438	0.5704	2	P02768

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Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co

C1S_HUMAN	K.M*LTPEHVFIHPGWK.L	1	3.6098	0.4597		P09871
C1S_HUMAN	K.TM*QENSTPRED.-	1	2.5095	0.4583		P09871
C1S_HUMAN	K.GDSGGAFAVQDPNDK.T	1	4.6296	0.5134		P09871
C1S_HUMAN	K.TMQENSTPRED.-	2	3.3133	0.5043		P09871
C1S_HUMAN	R.VKNYVDWIMK.T	1	3.4796	0.311		P09871
C1S_HUMAN	R.SSNNPHSPIVEEFQVPYNK.L	1	4.0838	0.3753	1	P09871

C1S_HUMAN	K.VEDPESTLFGSVIR.Y	1	3.6472	0.5109		P09871
C1S_HUMAN	K.SDFSNEER.F	1	2.5573	0.3803	4	P09871
C1S_HUMAN	K.EDTPNSVWEPAK.A	1	3.2727	0.3885	1	P09871

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Inter-alpha-trypsin inhibitor heavy chain H3 precursor (ITI heavy chain H3) (Inter-alpha-

ITIH3_HUMAN	K.SMEDKGMTNINDGLLR.G	1	4.4179	0.5161		Q06033
ITIH3_HUMAN	R.SLPEGVANGIEVYSTK.I	1	3.95	0.5713		Q06033
ITIH3_HUMAN	R.LVDEDMNSFKADVK.G	1	4.2542	0.4803		Q06033
ITIH3_HUMAN	R.DYIFGNYIER.L	1	2.9156	0.4235		Q06033
ITIH3_HUMAN	K.KGHVSFKPSLDQQR.S	1	3.2329	0.4348		Q06033
ITIH3_HUMAN	K.ALQERDYIFGNYIER.L	2	3.9918	0.5458		Q06033
ITIH3_HUMAN	R.LVDEDMNSFK.A	1	3.496	0.5126		Q06033

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Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)

A2MG_HUMAN	K.DTVIKPLLVEPEGLEK.E	1	3.8172	0.4013		P01023
A2MG_HUMAN	R.VGFYESDVMGR.G	2	3.685	0.534		P01023
A2MG_HUMAN	R.QTVSWAVTPK.S	1	2.401	0.3288		P01023
A2MG_HUMAN	K.LPPNVVEESAR.A	1	2.9241	0.3693		P01023
A2MG_HUMAN	K.AIGYLNTGYQR.Q	1	2.7452	0.2976		P01023
A2MG_HUMAN	K.NEDSLVFVQTDK.S	1	3.9914	0.5453		P01023

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Ig alpha-2 chain C region - Homo sapiens (Human)

IGHA2_HUMAN	K.SAVQGPPER.D	1	2.5989	0.4698	18	P01877
IGHA2_HUMAN	R.WLQGSQELPR.E	1	2.7773	0.4151	23	P01877
IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	1	4.2826	0.5297	21	P01877
IGHA2_HUMAN	R.GFSPKDVLVR.W	1	2.4818	0.2204	23	P01877
IGHA2_HUMAN	K.YLTWASRQEPSQGTTTFAVT	1	6.6449	0.6461	21	P01877
IGHA2_HUMAN	R.WLQGSQELPREK.Y	1	3.0378	0.2827	23	P01877

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AMBP protein precursor [Contains: Alpha-1-microglobulin (Protein HC) (Complex-formi

AMBP_HUMAN	R.TVAACNLPIVR.G	1	3.4799	0.4835		P02760
AMBP_HUMAN	R.AFIQLWAFDAVKGK.C	2	4.1325	0.4926		P02760
AMBP_HUMAN	R.AFIQLWAFDAVK.G	2	3.8073	0.5939		P02760

Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	4	3.6829	0.5561	21	A0A5E5
A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	1	5.8573	0.63	20	A0A5E5

Lumican precursor (Keratan sulfate proteoglycan lumican) (KSPG lumican) - Homo sap

LUM_HUMAN	K.RFNALQYLR.L	1	3.0978	0.2047		P51884
LUM_HUMAN	R.FNALQYLR.L	1	2.5981	0.3308		P51884
LUM_HUMAN	K.SLEDLQLTHNK.I	1	3.2307	0.4638		P51884
LUM_HUMAN	K.ISNIPDEYFK.R	1	2.9579	0.2464		P51884
LUM_HUMAN	K.NIPTVNENLENYYLEVNQLEK	1	4.1032	0.3938		P51884

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Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -

APOA2_HUMAN	K.VKSPELQAEAKSYFEK.S	1	3.6573	0.4774		P02652
APOA2_HUMAN	K.SPELQAEAK.S	1	3.4355	0.3032		P02652
APOA2_HUMAN	K.VKSPELQAEAK.S	2	3.7409	0.1945		P02652

Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	3.0124	0.4499	20	Q6N094
Q6N094_HUMAN	K.TTPPVLDSDGSFFLYSK.L	1	2.9721	0.4571	19	Q6N094

Q6N094_HUMAN	K.FNWYVDGVEVHNAK.T	1	4.7663	0.5151	20	Q6N094
Q6N094_HUMAN	K.VSNKALPAPIEK.T	1	3.4142	0.3664	24	Q6N094

Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S

SAMP_HUMAN	K.IVLGQEQDSYGGK.F	2	3.2867	0.4962		P02743
SAMP_HUMAN	K.IVLGQEQDSYGGKFDR.S	1	4.6182	0.578		P02743
SAMP_HUMAN	R.AYSLFSYNTQGR.D	1	2.9559	0.5446		P02743

Vitronectin precursor (Serum-spreading factor) (S-protein) (V75) [Contains: Vitronectin

VTNC_HUMAN	R.DVWGIEGPIDAAFTR.I	1	4.667	0.2991		P04004
VTNC_HUMAN	R.FEDGVLDPDYPR.N	1	3.6219	0.4228		P04004
VTNC_HUMAN	R.IYISGMAPRPSLAK.K	1	3.4761	0.363		P04004
VTNC_HUMAN	K.TYLFKGSQYWR.F	1	2.4056	0.4457		P04004

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Apolipoprotein B-100 precursor (Apo B-100) [Contains: Apolipoprotein B-48 (Apo B-48)

APOB_HUMAN	R.ALVDTLKFVTQAEGAK.Q	1	5.2569	0.6109		P04114
APOB_HUMAN	R.DLKVEDIPLAR.I	1	2.5225	0.1314		P04114
APOB_HUMAN	R.SPSQADINK.I	1	2.6767	0.3156		P04114

Haptoglobin-related protein precursor - Homo sapiens (Human)

HPTR_HUMAN	R.TEGDGVYTLNDK.K	1	3.1341	0.4401	2	P00739
HPTR_HUMAN	R.VGYVSGWGQSDNFK.L	1	4.2868	0.6115		P00739
HPTR_HUMAN	R.ILGGHLDAKGSFPWQAK.M	1	3.4527	0.3699	3	P00739

IGHA1 protein - Homo sapiens (Human)

Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	2	4.4406	0.6319	12	Q8N5K4
Q8N5K4_HUMAN	K.SAVQGPPDR.D	1	2.5366	0.3242	4	Q8N5K4

IGHM protein - Homo sapiens (Human)

Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	1	3.5819	0.5938	4	Q6GMY2
Q6GMY2_HUMAN	K.YAATSQVLLPSK.D	1	2.8675	0.3813	3	Q6GMY2
Q6GMY2_HUMAN	R.EGKQVGSGVTTDQVQAEAK.	1	4.1758	0.101	4	Q6GMY2

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.IIRSSEDPNEDIVERNIR.I	1	3.7276	0.393		P01591
IGJ_HUMAN	R.IIRSSEDPNEDIVER.N	1	4.04	0.3934		P01591
IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.1595	0.2789		P01591

Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	K.YNSQNQSNNQFVLYR.I	1	4.2439	0.5298		P01042
KNG1_HUMAN	K.YFIDFVAR.E	1	2.8595	0.4346	1	P01042
KNG1_HUMAN	K.KYFIDFVAR.E	1	2.982	0.4953	1	P01042

2

Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr

FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.2736	0.4267		P02765
FETUA_HUMAN	R.HTFMGVVSLGSPSGEVSHPR	1	5.0783	0.6454		P02765

Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis

CLUS_HUMAN	R.ASSIIDELFQDR.F	1	4.0437	0.481	1	P10909
CLUS_HUMAN	R.ELDESLQVAER.L	1	2.8502	0.425	1	P10909

Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) - Homo sapiens (Human)

FINC_HUMAN	R.WSRPQAPITGYR.I	1	3.2693	0.29	1	P02751
FINC_HUMAN	R.FTNIGPDTMR.V	1	2.4988	0.2377	1	P02751

Ig lambda chain V-IV region Hil - Homo sapiens (Human)

LV403_HUMAN	-.SYELTQPPSVSVSPGQTAR.I	2	5.3552	0.6273	1	P01717

Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)

MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	1	3.9092	0.4529	5	P04220
MUCB_HUMAN	K.YVTSAPMPEPQAPGR.Y	1	3.7289	0.5464	5	P04220

IGLV2-14 protein - Homo sapiens (Human)

A0A5C9_HUMAN	K.AGVETTTPSK.Q	1	3.3895	0.3475	17	A0A5C9
A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	5.3057	0.6077	29	A0A5C9

Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-

ITIH4_HUMAN	R.GPDVLTATVSGK.L	1	3.0093	0.4413	1	Q14624
ITIH4_HUMAN	K.AEAQAQYSAAVAK.G	1	4.4999	0.5246	1	Q14624

Plasma protease C1 inhibitor precursor (C1 Inh) (C1Inh) (C1 esterase inhibitor) (C1-inh

IC1_HUMAN	R.LEDMEQALSPSVFK.A	1	4.578	0.5767	1	P05155
IC1_HUMAN	K.TNLESILSYPK.D	1	3.5952	0.4402	1	P05155

Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II) [Contains: Activation peptid

THRB_HUMAN	R.TATSEYQTFFNPR.T	2	3.9466	0.4962		P00734

Thiopurine S-methyltransferase (EC 2.1.1.67) (Thiopurine methyltransferase) - Homo s

TPMT_HUMAN	R.VFFPLCGKAVEM*KWFADR.C	2	2.9926	0.0932		P51580

Zinc finger CCHC domain-containing protein 11 - Homo sapiens (Human)

ZCH11_HUMAN	R.M*DDFQLKGIVEEKFVK.W	2	3.2114	0.3501	1	Q5TAX3

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60S ribosomal protein L3-like - Homo sapiens (Human)

RL3L_HUMAN	K.VACIGAWHPARVGCSIAR.A	1	2.6599	0.2128		Q92901

Anti-folate binding protein (Fragment) - Homo sapiens (Human)

A2NYQ9_HUMAN	-.QVQLVESGGGLVQPGR.S	1	2.457	0.4229		A2NYQ9

Antithrombin III variant - Homo sapiens (Human)

Q7KZ97_HUMAN	R.DDLYVSDAFHK.A	1	2.4967	0.4915		Q7KZ97

Apolipoprotein E precursor (Apo-E) - Homo sapiens (Human)

APOE_HUMAN	R.LSKELQAAQAR.L	1	3.3117	0.4569		P02649

C4B1 (Complement component 4B) (Childo blood group) - Homo sapiens (Human)

Q6U2E9_HUMAN	R.GSSTWLTAFVLK.V	1	2.6087	0.3873		Q6U2E9

CD180 antigen precursor (Lymphocyte antigen 64) (Radioprotective 105 kDa protein) -

CD180_HUMAN	K.ECPQLELLDLAFTR.L	1	2.999	0.4318		Q99467

CDNA FLJ40176 fis, clone TESTI2017102. (Fragment) - Homo sapiens (Human)

Q8N800_HUMAN	K.QLKAHLLQKEEK.D	1	2.8733	0.0945		Q8N800

Ciliary dynein heavy chain 7 - Homo sapiens (Human)

Q8WXX0_HUMAN	K.TQTM*RGSPFIKPYEK.Q	1	2.473	0.1831		Q8WXX0

Complement C4-B precursor (Basic complement C4) [Contains: Complement C4 beta c

CO4B_HUMAN	K.LQETSNWLLSQQQADGSFQL	1	3.6095	0.3175	1	P0C0L5

Complement component C9 precursor [Contains: Complement component C9a; Compl

CO9_HUMAN	R.AIEDYINEFSVR.K	1	3.339	0.4863		P02748

Complement-activating component of Ra-reactive factor precursor (EC 3.4.21.-) (Ra-re

MASP1_HUMAN	R.TGVITSPDFPNPYPK.S	1	3.7292	0.4799	2	P48740

D site-binding protein (Albumin D box-binding protein) (Albumin D-element-binding prot

DBP_HUMAN	R.TPAPLLLGGPAGTPPGGGALI	1	3.2803	0.1027	1	Q10586

ELKS/RAB6-interacting/CAST family member 1 (RAB6-interacting protein 2) (ERC prot

RB6I2_HUMAN	R.EM*VLAQEESAR.T	1	2.5384	0.1349		Q8IUD2

Endothelial zinc finger protein induced by tumor necrosis factor alpha (Zinc finger prote

ZNF71_HUMAN	K.PFECSECGKAFSKNSSLTQH	1	3.3151	0.2297		Q9NQZ8

GRIP and coiled-coil domain-containing protein 2 (Golgi coiled coil protein GCC185) (C

GCC2_HUMAN	K.AIHQEEVKELMCQIEASAK.E	1	2.7128	0.1074	1	Q8IWJ2

Histidine-rich glycoprotein precursor (Histidine-proline-rich glycoprotein) (HPRG) - Hom

HRG_HUMAN	R.DGYLFQLLR.I	1	2.5636	0.1762		P04196

Hypothetical protein DKFZp686N1815 - Homo sapiens (Human)

Q5HYI3_HUMAN	-.MSDALIKVSM*ENTENSVDSK.	1	3.2663	0.1666		Q5HYI3

Ig heavy chain V-III region TIL - Homo sapiens (Human)

HV304_HUMAN	-.EVQLLESGGGLVQPGGSLR.L	1	5.0079	0.2959	5	P01765

Ig lambda chain V-III region LOI - Homo sapiens (Human)

LV302_HUMAN	-.YVLTQPPSVSVAPGETAR.L	1	3.9234	0.4707		P80748

IGHG4 protein - Homo sapiens (Human)

Q8TC63_HUMAN	K.YGPPCPSCPAPEFLGGPSVFI	1	3.2921	0.2131	1	Q8TC63

KIAA1729 protein (Fragment) - Homo sapiens (Human)

Q9C0D4_HUMAN	K.QGRLLSRSLSISR.N	1	2.7735	0.0979		Q9C0D4

KNG1 protein - Homo sapiens (Human)

Q05CF8_HUMAN	R.IGEIKEETTSHLR.S	1	3.7015	0.458		Q05CF8

Makorin-2 (RING finger protein 62) - Homo sapiens (Human)

MKRN2_HUMAN	K.ACKYFEQGK.G	1	2.9893	0.2316		Q9H000

Myosin-10 (Myosin heavy chain 10) (Myosin heavy chain, nonmuscle IIb) (Nonmuscle n

MYH10_HUMAN	R.ALEEALEAK.E	1	2.7396	0.1632	4	P35580

Myosin-reactive immunoglobulin light chain variable region (Fragment) - Homo sapiens

Q9UL83_HUMAN	-.EIVMTQSPATLSVSPGER.A	1	4.3905	0.5531	1	Q9UL83

Neuron navigator 1 - Homo sapiens (Human)

Q5VUY9_HUMAN	-.M*DARIVHALLAGRM*LGSSVK	1	2.4134	0.1138		Q5VUY9

OTTHUMP00000016950 (Fragment) - Homo sapiens (Human)

Q5T699_HUMAN	R.ELESCLRREYTPENLPLLLLQ`	1	3.4942	0.2002		Q5T699

PDZ domain-containing RING finger protein 4 (Ligand of Numb-protein X 4) (SEMACAI

PZRN4_HUMAN	K.TTEQGCSAESKEKVLEGSK.L	1	2.8186	0.3041		Q6ZMN7

Protein FAM71A - Homo sapiens (Human)

FA71A_HUMAN	K.YAPIFESDFIQITK.R	1	2.459	0.2482		Q8IYT1

Round spermatid basic protein 1 - Homo sapiens (Human)

Q5VWQ0_HUMAN	K.QENGERTGGVPLIKAPK.R	1	2.4472	0.1388		Q5VWQ0

Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk

PON1_HUMAN	K.IFFYDSENPPASEVLR.I	1	5.35	0.6121		P27169

Synapsin-1 (Synapsin I) (Brain protein 4.1) - Homo sapiens (Human)

SYN1_HUMAN	K.MAQALPRQR.Q	1	2.4339	0.2295	1	P17600

U3 small nucleolar RNA-associated protein 15 homolog - Homo sapiens (Human)

UTP15_HUMAN	K.NYMKQRDDILINR.P	1	2.5718	0.0921		Q8TED0

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Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	R.GLQDEDGYRMK.F	1	2.4071	0.3129	3	P0C0L4
CO4A_HUMAN	R.GQIVFM*NR.E	1	2.492	0.3471	3	P0C0L4
CO4A_HUMAN	R.GQIVFMNR.E	1	2.4333	0.3426	3	P0C0L4
CO4A_HUMAN	R.TTNIQGINLLFSSR.R	1	4.7631	0.5765	3	P0C0L4
CO4A_HUMAN	R.TYNVLDM*K.N	1	2.694	0.5051	3	P0C0L4
CO4A_HUMAN	R.VFALDQK.M	1	2.4999	0.2052	3	P0C0L4
CO4A_HUMAN	R.GLQDEDGYRM*K.F	1	2.7311	0.4259	3	P0C0L4
CO4A_HUMAN	K.LNM*GITDLQGLR.L	1	3.4167	0.4309	3	P0C0L4
CO4A_HUMAN	R.TYNVLDMK.N	1	3.4537	0.455	3	P0C0L4
CO4A_HUMAN	R.FGLLDEDGKK.T	1	3.1154	0.3077	3	P0C0L4
CO4A_HUMAN	R.EMSGSPASGIPVK.V	1	2.6469	0.1944	3	P0C0L4
CO4A_HUMAN	R.EAPKVVEEQESR.V	1	2.5727	0.5202	3	P0C0L4
CO4A_HUMAN	K.VVEEQESR.V	6	3.0124	0.4802	3	P0C0L4

CO4A_HUMAN	K.VLSLAQEQVGGSPEK.L	1	4.6859	0.5801	3	P0C0L4
CO4A_HUMAN	K.LNMGITDLQGLR.L	1	2.9992	0.4069	3	P0C0L4
CO4A_HUMAN	R.VGDTLNLNLR.A	1	3.0392	0.4279	3	P0C0L4
CO4A_HUMAN	K.AEFQDALEK.L	2	3.0637	0.3898	3	P0C0L4
CO4A_HUMAN	R.VTASDPLDTLGSEGALSPGG\	1	4.7875	0.4026	3	P0C0L4
CO4A_HUMAN	K.LVNGQSHISLSK.A	1	3.5957	0.3682	3	P0C0L4

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Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	K.NNEGTYYSPNYNPQSR.S	1	4.5834	0.6147		P00450
CERU_HUMAN	R.RQSEDSTFYLGER.T	1	3.9292	0.4297		P00450
CERU_HUMAN	R.QSEDSTFYLGER.T	1	3.1714	0.3961		P00450
CERU_HUMAN	R.MFTTAPDQVDKEDEDFQESN	1	3.5121	0.4899		P00450
CERU_HUMAN	R.M*FTTAPDQVDKEDEDFQESI	1	3.9187	0.5423		P00450
CERU_HUMAN	R.HYYIAAEEIIWNYAPSGIDIFTk	1	4.1791	0.4869		P00450
CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEMF	2	4.4138	0.6055		P00450
CERU_HUMAN	R.GPEEEHLGILGPVIWAEVGDT	1	5.9732	0.3634		P00450
CERU_HUMAN	R.EYTDASFTNRK.E	1	2.6647	0.1886		P00450
CERU_HUMAN	R.TTIEKPVWLGFLGPIIK.A	1	3.4838	0.4146		P00450
CERU_HUMAN	K.VNKDDEEFIESNK.M	1	4.3337	0.3829		P00450
CERU_HUMAN	K.MYYSAVDPTK.D	1	2.9549	0.5316		P00450
CERU_HUMAN	K.NLASRPYTFHSHGITYYK.E	1	4.3157	0.5078		P00450
CERU_HUMAN	K.MYYSAVDPTKDIFTGLIGPMK	1	3.9247	0.4754		P00450
CERU_HUMAN	K.M*YYSAVDPTKDIFTGLIGPMk	1	4.6273	0.3653		P00450
CERU_HUMAN	K.KALYLQYTDETFR.T	1	4.2346	0.5446		P00450
CERU_HUMAN	K.HYYIGIIETTWDYASDHGEK.K	1	5.6986	0.5691		P00450
CERU_HUMAN	K.GAYPLSIEPIGVR.F	1	3.282	0.5201		P00450
CERU_HUMAN	K.DVDKEFYLFPTVFDENESLLLI	1	4.5865	0.5302		P00450
CERU_HUMAN	K.ALYLQYTDETFR.T	1	3.832	0.5919		P00450
CERU_HUMAN	R.EYTDASFTNR.K	1	2.6524	0.4441		P00450
CERU_HUMAN	K.LISVDTEHSNIYLQNGPDR.I	1	3.7583	0.3084		P00450

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Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-

ITIH1_HUMAN	R.KAAISGENAGLVR.A	1	3.8053	0.4364		P19827
ITIH1_HUMAN	K.AAISGENAGLVR.A	2	3.7241	0.5011		P19827
ITIH1_HUMAN	K.GSLVQASEANLQAAQDFVR.C	2	5.9813	0.6263		P19827
ITIH1_HUMAN	K.LDAQASFLPK.E	1	3.4022	0.3892		P19827
ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	2	3.9365	0.4775	1	P19827
ITIH1_HUMAN	R.IYEDHDATQQLQGFYSQVAK.	1	3.6277	0.379	1	P19827
ITIH1_HUMAN	R.LWAYLTIQELLAK.R	1	4.9746	0.6001	1	P19827
ITIH1_HUMAN	R.QAVDTAVDGVFIR.S	1	4.1716	0.3856		P19827
ITIH1_HUMAN	R.EVAFDLEIPK.T	1	2.9459	0.3283		P19827

Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-

ITIH2_HUMAN	R.SSALDMENFR.T	1	3.0546	0.3267	2	P19823
ITIH2_HUMAN	R.LSNENHGIAQR.I	1	4.061	0.4168	1	P19823
ITIH2_HUMAN	R.TWRNDLISATK.T	1	2.5672	0.1646	1	P19823
ITIH2_HUMAN	R.TEVNVLPGAK.V	1	2.598	0.4778	2	P19823
ITIH2_HUMAN	R.SSALDM*ENFR.T	1	2.6962	0.3241	2	P19823
ITIH2_HUMAN	R.MATTM*IQSK.V	1	2.8153	0.4635	2	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLK.K	1	3.4642	0.4347	1	P19823
ITIH2_HUMAN	R.IYGNQDTSSQLKK.F	1	4.2318	0.4001	1	P19823
ITIH2_HUMAN	K.FYNQVSTPLLR.N	1	2.8481	0.4364	1	P19823

ITIH2_HUMAN	R.AEDHFSVIDFNQNIR.T	1	4.4015	0.2253	1	P19823
ITIH2_HUMAN	K.VVNNSPQPQNVVFDVQIPK.G	1	5.2367	0.5132	2	P19823
ITIH2_HUMAN	K.IQPSGGTNINEALLR.A	1	2.6386	0.3213	1	P19823

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Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H

APOA1_HUMAN	K.VQPYLDDFQKK.W	1	2.6372	0.1761		P02647
APOA1_HUMAN	R.EQLGPVTQEFWDNLEK.E	1	3.5264	0.5807		P02647
APOA1_HUMAN	R.DYVSQFEGSALGK.Q	2	4.1499	0.6571		P02647
APOA1_HUMAN	K.WQEEM*ELYR.Q	1	2.5422	0.4107		P02647
APOA1_HUMAN	K.ATEHLSTLSEK.A	1	2.9371	0.4577		P02647
APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	2	4.348	0.5849		P02647
APOA1_HUMAN	R.THLAPYSDELR.Q	1	2.4994	0.5258		P02647

Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alpl

CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	2.7685	0.4295		P01024
CO3_HUMAN	K.SGSDEVQVGQQR.T	1	3.7891	0.3803		P01024
CO3_HUMAN	R.SEETKENEGFTVTAEGK.G	1	4.1686	0.4056		P01024
CO3_HUMAN	K.TGLQEVEVK.A	1	2.6887	0.3767		P01024
CO3_HUMAN	R.TKKQELSEAEQATR.T	1	4.0671	0.5069		P01024
CO3_HUMAN	K.VLLDGVQNPR.A	1	3.426	0.436		P01024
CO3_HUMAN	R.LVAYYTLIGASGQR.E	1	3.4572	0.4591		P01024
CO3_HUMAN	K.QELSEAEQATR.T	1	2.4884	0.4596		P01024
CO3_HUMAN	R.ILLQGTPVAQM*TEDAVDAER	1	4.5116	0.613		P01024

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Serum albumin precursor - Homo sapiens (Human)

ALBU_HUMAN	K.KVPQVSTPTLVEVSR.N	1	3.2926	0.3997	2	P02768
ALBU_HUMAN	R.RHPYFYAPELLFFAK.R	1	3.7014	0.5109	1	P02768
ALBU_HUMAN	K.VPQVSTPTLVEVSR.N	1	3.2986	0.5401	2	P02768
ALBU_HUMAN	K.KQTALVELVK.H	1	2.6422	0.3424	2	P02768
ALBU_HUMAN	K.AVMDDFAAFVEK.C	2	3.7478	0.5739	2	P02768
ALBU_HUMAN	K.AVM*DDFAAFVEK.C	1	3.4831	0.635	2	P02768
ALBU_HUMAN	R.RHPDYSVVLLLR.L	1	3.8562	0.3682	1	P02768

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Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S

SAMP_HUMAN	K.IVLGQEQDSYGGK.F	1	3.9718	0.4492		P02743
SAMP_HUMAN	R.AYSLFSYNTQGR.D	1	4.04	0.4496		P02743
SAMP_HUMAN	R.DNELLVYK.E	1	2.4053	0.3164		P02743
SAMP_HUMAN	R.QGYFVEAQPK.I	1	2.5672	0.4379		P02743
SAMP_HUMAN	R.VGEYSLYIGR.H	1	3.6042	0.4342		P02743

3

Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co

C1S_HUMAN	R.TNFDNDIALVR.L	1	3.5425	0.3835		P09871
C1S_HUMAN	K.GDSGGAFAVQDPNDK.T	1	4.3845	0.546		P09871
C1S_HUMAN	K.TMQENSTPRED.-	1	3.3437	0.5281		P09871

Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	K.VDNALQSGNSQESVTEQDSK	1	5.5712	0.6261	20	A0A5E5
A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	2	3.917	0.5906	21	A0A5E5

Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	2.8869	0.5022	20	Q6N094
Q6N094_HUMAN	K.TTPPVLDSDGSFFLYSK.L	1	3.4781	0.3901	19	Q6N094
Q6N094_HUMAN	K.FNWYVDGVEVHNAK.T	1	3.2459	0.4806	20	Q6N094

Ig alpha-2 chain C region - Homo sapiens (Human)							
	IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	1	3.9947	0.5029	21	P01877
	IGHA2_HUMAN	R.WLQGSQELPR.E	1	3.0348	0.3545	23	P01877
	IGHA2_HUMAN	R.DASGATFTWTPSSGK.S	1	3.496	0.5053	10	P01877
IGHM protein - Homo sapiens (Human)							
	Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	1	4.418	0.5227	4	Q6GMY2
	Q6GMY2_HUMAN	K.YAATSQVLLPSK.D	2	3.7846	0.4407	3	Q6GMY2
IGLV2-14 protein - Homo sapiens (Human)							
	A0A5C9_HUMAN	K.AGVETTTPSK.Q	2	2.5807	0.377	17	A0A5C9
	A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	4.8145	0.6578	29	A0A5C9
Inter-alpha-trypsin inhibitor heavy chain H3 precursor (ITI heavy chain H3) (Inter-alpha-							
	ITIH3_HUMAN	K.GM*TNINDGLLR.G	1	3.0132	0.5054		Q06033
	ITIH3_HUMAN	R.LVDEDM*NSFK.A	1	3.5135	0.3997		Q06033
	ITIH3_HUMAN	R.LVDEDMNSFK.A	1	3.1914	0.4437		Q06033
Lumican precursor (Keratan sulfate proteoglycan lumican) (KSPG lumican) - Homo sap							
	LUM_HUMAN	K.ISNIPDEYFK.R	1	3.2376	0.4606		P51884
	LUM_HUMAN	K.SLEDLQLTHNK.I	1	3.3638	0.4227		P51884
	LUM_HUMAN	R.FNALQYLR.L	1	2.6056	0.3009		P51884
2							
Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr							
	FETUA_HUMAN	R.HTFMGVVSLGSPSGEVSHPR	1	4.6129	0.6253		P02765
	FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.5472	0.3899		P02765
Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -							
	APOA2_HUMAN	K.SPELQAEAK.S	1	2.8151	0.3461		P02652
	APOA2_HUMAN	K.VKSPELQAEAK.S	1	3.4516	0.2175		P02652
Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis							
	CLUS_HUMAN	R.VTTVASHTSDSDVPSGVTEV\	1	3.3538	0.2748		P10909
	CLUS_HUMAN	R.ELDESLQVAER.L	1	2.9854	0.4245	1	P10909
Haptoglobin-related protein precursor - Homo sapiens (Human)							
	HPTR_HUMAN	R.VGYVSGWGQSDNFK.L	1	4.5866	0.5893		P00739
	HPTR_HUMAN	R.TEGDGVYTLNDK.K	1	2.5299	0.1702	2	P00739
Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)							
	MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	1	3.9999	0.3893	5	P04220
	MUCB_HUMAN	K.LICQATGFSPR.Q	1	3.368	0.4031	5	P04220
Immunoglobulin J chain - Homo sapiens (Human)							
	IGJ_HUMAN	R.IIRSSEDPNEDIVER.N	1	4.1415	0.3555		P01591
	IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.0844	0.3986		P01591
Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy							
	KNG1_HUMAN	K.YNSQNQSNNQFVLYR.I	1	4.3897	0.4959		P01042
	KNG1_HUMAN	K.TVGSDTFYSFK.Y	1	3.4538	0.5855		P01042
Myosin-reactive immunoglobulin heavy chain variable region (Fragment) - Homo sapier							
	Q9UL94_HUMAN	-.EVQLVESGAEVK.K	2	4.1458	0.4717	2	Q9UL94
Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk							
	PON1_HUMAN	K.IFFYDSENPPASEVLR.I	2	5.6229	0.621		P27169
Vitronectin precursor (Serum-spreading factor) (S-protein) (V75) [Contains: Vitronectin							
	VTNC_HUMAN	R.IYISGMAPR.P	1	2.475	0.3139		P04004
	VTNC_HUMAN	R.FEDGVLDPDYPR.N	1	3.1655	0.4206		P04004
1							
AMBP protein precursor [Contains: Alpha-1-microglobulin (Protein HC) (Complex-formi							
	AMBP_HUMAN	R.AFIQLWAFDAVK.G	1	4.3525	0.5211		P02760

Apolipoprotein(a) precursor (EC 3.4.21.-) (Apo(a)) (Lp(a)) - Homo sapiens (Human)

APOA_HUMAN	R.GTYSTTVTGR.T	1	2.4154	0.5344	39	P08519

Cadherin-20 precursor - Homo sapiens (Human)

CAD20_HUMAN	K.IQDINDNEPK.F	1	2.9307	0.1572	2	Q9HBT6

Carboxypeptidase N subunit 2 precursor (Carboxypeptidase N polypeptide 2) (Carboxy

CPN2_HUMAN	K.AGGSWDLAVQER.A	1	2.8187	0.4298		P22792

CDNA FLJ39060 fis, clone NT2RP7013729 (Novel protein) - Homo sapiens (Human)

Q8N8P6_HUMAN	R.HSASVPM*ATGANGLETKDE	1	2.6044	0.0904		Q8N8P6

CDNA FLJ40176 fis, clone TESTI2017102. (Fragment) - Homo sapiens (Human)

Q8N800_HUMAN	K.QLKAHLLQKEEK.D	1	2.5621	0.196		Q8N800

Ciliary dynein heavy chain 7 - Homo sapiens (Human)

Q8WXX0_HUMAN	K.TQTM*RGSPFIKPYEK.Q	1	2.8472	0.2339		Q8WXX0

Complement component C9 precursor [Contains: Complement component C9a; Compl

CO9_HUMAN	R.VVEESELAR.T	1	2.6919	0.5073		P02748

Fibulin-1 precursor - Homo sapiens (Human)

FBLN1_HUMAN	R.SAATLQQEKTDTVR.C	1	3.201	0.3533		P23142

Ig kappa chain V-III region VG precursor (Fragment) - Homo sapiens (Human)

KV309_HUMAN	R.LLIYDASNR.A	1	2.8136	0.4602	1	P04433

IGHA1 protein - Homo sapiens (Human)

Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	1	4.8828	0.5256	12	Q8N5K4

Inter-alpha-trypsin inhibitor heavy chain H4 precursor (ITI heavy chain H4) (Inter-alpha-

ITIH4_HUMAN	K.GSEMVVAGK.L	1	2.5364	0.2952	1	Q14624

Latrophilin-2 (Fragment) - Homo sapiens (Human)

Q9UJ49_HUMAN	-.MTGNYLLTNPLLR.P	1	2.8956	0.3433		Q9UJ49

Nesprin-1 (Nuclear envelope spectrin repeat protein 1) (Synaptic nuclear envelope prot

SYNE1_HUMAN	R.LQQILNFQNDLK.V	1	2.5953	0.1957		Q8NF91

Olfactory receptor 17-1 - Homo sapiens (Human)

Q9UL14_HUMAN	R.GQGRPEKGAQM*EGCTPR.G	1	3.2623	0.2951		Q9UL14

Prothrombin precursor (EC 3.4.21.5) (Coagulation factor II) [Contains: Activation peptid

THRB_HUMAN	R.TATSEYQTFFNPR.T	1	3.0985	0.558		P00734

Signal peptide peptidase-like 2B (EC 3.4.23.-) (Protein SPP-like 2B) (Protein SPPL2b)

PSL1_HUMAN	R.TVRAALYAPK.E	1	2.5145	0.2658		Q8TCT7

Sodium bicarbonate cotransporter 3 (Sodium bicarbonate cotransporter 2) (Sodium bic

S4A7_HUMAN	R.TGLSASNLSLR.G	1	2.9495	0.0945		Q9Y6M7

Transcription elongation factor B polypeptide 1 (RNA polymerase II transcription factor

ELOC_HUMAN	K.AMLSGPGQFAENETNEVNFR	1	2.861	0.09		Q15369

Tribbles homolog 3 (TRB-3) (Neuronal cell death-inducible putative kinase) (p65-intera

TRIB3_HUMAN	K.VYPVQEALAVLEPYAR.L	1	2.8921	0.0852		Q96RU7

Zinc finger CCHC domain-containing protein 11 - Homo sapiens (Human)

ZCH11_HUMAN	R.M*DDFQLKGIVEEKFVK.W	1	2.6587	0.185	1	Q5TAX3

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Ceruloplasmin precursor (EC 1.16.3.1) (Ferroxidase) - Homo sapiens (Human)

CERU_HUMAN	R.GVYSSDVFDIFPGTYQTLEMF	1	4.5017	0.5373		P00450
CERU_HUMAN	R.QSEDSTFYLGER.T	1	3.1777	0.4158		P00450
CERU_HUMAN	R.EYTDASFTNR.K	1	2.8435	0.4899		P00450
CERU_HUMAN	K.NNEGTYYSPNYNPQSR.S	1	4.3791	0.5262		P00450
CERU_HUMAN	K.NLASRPYTFHSHGITYYK.E	1	3.3318	0.3666		P00450
CERU_HUMAN	K.MYYSAVDPTK.D	1	2.8316	0.5446		P00450

	CERU_HUMAN	R.M*FTTAPDQVDKEDEDFQESI	1	3.6194	0.5255		P00450
	CERU_HUMAN	K.KALYLQYTDETFR.T	1	3.9241	0.4069		P00450
	CERU_HUMAN	R.HYYIAAEEIIWNYAPSGIDIFTK	1	3.7563	0.4767		P00450
	CERU_HUMAN	K.GAYPLSIEPIGVR.F	1	2.8385	0.5027		P00450
	CERU_HUMAN	K.DIFTGLIGPMK.I	1	2.5124	0.3608		P00450
	CERU_HUMAN	K.ALYLQYTDETFR.T	1	3.7459	0.5887		P00450
	CERU_HUMAN	R.MFTTAPDQVDKEDEDFQESN	1	3.2052	0.4359		P00450
	CERU_HUMAN	R.GPEEEHLGILGPVIWAEVGDT	1	5.0116	0.2546		P00450
10							
Inter-alpha-trypsin inhibitor heavy chain H1 precursor (ITI heavy chain H1) (Inter-alpha-							
	ITIH1_HUMAN	R.LWAYLTIQELLAK.R	1	4.4657	0.579	1	P19827
	ITIH1_HUMAN	K.AAISGENAGLVR.A	1	3.4562	0.5003		P19827
	ITIH1_HUMAN	K.GSLVQASEANLQAAQDFVR.C	2	5.7131	0.6455		P19827
	ITIH1_HUMAN	K.LDAQASFLPK.E	1	3.1268	0.3438		P19827
	ITIH1_HUMAN	K.QYYEGSEIVVAGR.I	1	3.6842	0.4616	1	P19827
	ITIH1_HUMAN	R.FAHYVVTSQVVNTANEAR.E	1	3.6668	0.3095		P19827
	ITIH1_HUMAN	R.NHMQYEIVIK.V	1	2.9186	0.3214		P19827
	ITIH1_HUMAN	R.QAVDTAVDGVFIR.S	1	2.8834	0.3834		P19827
	ITIH1_HUMAN	R.EVAFDLEIPK.T	1	2.781	0.419		P19827
7							
Inter-alpha-trypsin inhibitor heavy chain H2 precursor (ITI heavy chain H2) (Inter-alpha-							
	ITIH2_HUMAN	R.SSALDMENFR.T	1	3.0438	0.3373	2	P19823
	ITIH2_HUMAN	R.SSALDM*ENFR.T	1	2.6937	0.3523	2	P19823
	ITIH2_HUMAN	R.IYGNQDTSSQLK.K	1	3.5273	0.4715	1	P19823
	ITIH2_HUMAN	K.VVNNSPQPQNVVFDVQIPK.G	2	5.1604	0.5597	2	P19823
	ITIH2_HUMAN	K.IQPSGGTNINEALLR.A	2	3.5258	0.5773	1	P19823
6							
Apolipoprotein A-I precursor (Apo-AI) (ApoA-I) [Contains: Apolipoprotein A-I(1-242)] - H							
	APOA1_HUMAN	K.DSGRDYVSQFEGSALGK.Q	1	4.6894	0.356		P02647
	APOA1_HUMAN	K.WQEEM*ELYR.Q	1	3.1586	0.2961		P02647
	APOA1_HUMAN	K.VSFLSALEEYTK.K	1	4.0551	0.5126		P02647
	APOA1_HUMAN	K.LLDNWDSVTSTFSK.L	1	3.9495	0.532		P02647
	APOA1_HUMAN	K.ATEHLSTLSEK.A	1	2.7128	0.4119		P02647
	APOA1_HUMAN	R.DYVSQFEGSALGK.Q	1	4.2544	0.6043		P02647
5							
Complement C3 precursor [Contains: Complement C3 beta chain; Complement C3 alp							
	CO3_HUMAN	K.VQLSNDFDEYIMAIEQTIK.S	1	3.7883	0.3539		P01024
	CO3_HUMAN	R.ILLQGTPVAQMTEDAVDAER.I	1	4.3936	0.532		P01024
	CO3_HUMAN	K.SGSDEVQVGQQR.T	1	3.8284	0.3151		P01024
	CO3_HUMAN	R.TKKQELSEAEQATR.T	1	4.2335	0.5286		P01024
	CO3_HUMAN	R.IPIEDGSGEVVLSR.K	1	3.7892	0.5929		P01024
Ig mu heavy chain disease protein (BOT) - Homo sapiens (Human)							
	MUCB_HUMAN	K.QVGSGVTTDEVEAEAK.E	1	2.9155	0.4035		P04220
	MUCB_HUMAN	K.GVALHRPDVYLLPPAR.E	1	3.9153	0.4453	5	P04220
	MUCB_HUMAN	R.GQPLSPEKYVTSAPMPEPQA	1	3.2128	0.3236	5	P04220
	MUCB_HUMAN	K.YVTSAPMPEPQAPGR.Y	1	3.4655	0.5179	5	P04220
	MUCB_HUMAN	R.EGKQVGSGVTTDEVEAEAK.E	1	4.4872	0.2898		P04220
4							
Serum albumin precursor - Homo sapiens (Human)							
	ALBU_HUMAN	R.RHPDYSVVLLLR.L	1	3.6792	0.334	1	P02768

ALBU_HUMAN	K.AVMDDFAAFVEK.C	1	3.371	0.5933	2	P02768
ALBU_HUMAN	K.FQNALLVR.Y	1	2.496	0.253	2	P02768
ALBU_HUMAN	K.LVNEVTEFAK.T	1	2.9488	0.492	2	P02768

3

Complement C4-A precursor (Acidic complement C4) [Contains: Complement C4 beta

CO4A_HUMAN	R.GLQDEDGYR.M	1	2.8407	0.4582	3	P0C0L4
CO4A_HUMAN	R.VTASDPLDTLGSEGALSPGG\	1	3.7844	0.5837	3	P0C0L4
CO4A_HUMAN	R.VGDTLNLNLR.A	1	3.3181	0.3626	3	P0C0L4

Ig alpha-2 chain C region - Homo sapiens (Human)

IGHA2_HUMAN	R.QEPSQGTTTFAVTSILR.V	1	3.4094	0.5602	21	P01877
IGHA2_HUMAN	R.GFSPKDVLVR.W	1	2.7185	0.3404	23	P01877
IGHA2_HUMAN	R.WLQGSQELPR.E	1	2.7793	0.3408	23	P01877

Kininogen-1 precursor (Alpha-2-thiol proteinase inhibitor) [Contains: Kininogen-1 heavy

KNG1_HUMAN	K.TVGSDTFYSFK.Y	1	3.2349	0.5678		P01042
KNG1_HUMAN	K.KYFIDFVAR.E	1	2.4739	0.3494	1	P01042
KNG1_HUMAN	K.YFIDFVAR.E	1	2.7463	0.4936	1	P01042

2

Apolipoprotein A-II precursor (Apo-AII) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)] -

APOA2_HUMAN	K.VKSPELQAEAK.S	1	3.2396	0.3399		P02652
APOA2_HUMAN	K.SPELQAEAK.S	1	3.151	0.3134		P02652

Hypothetical protein - Homo sapiens (Human)

A0A5E5_HUMAN	R.TVAAPSVFIFPPSDEQLK.S	2	2.8274	0.4395	21	A0A5E5

Hypothetical protein DKFZp686O01196 - Homo sapiens (Human)

Q6N094_HUMAN	K.TTPPVLDSDGSFFLYSK.L	1	2.9694	0.4475	19	Q6N094
Q6N094_HUMAN	K.GPSVFPLAPSSK.S	1	2.5384	0.4339	20	Q6N094

IGHM protein - Homo sapiens (Human)

Q6GMY2_HUMAN	K.QVGSGVTTDQVQAEAK.E	1	4.1407	0.6249	4	Q6GMY2
Q6GMY2_HUMAN	K.YAATSQVLLPSK.D	1	3.3104	0.4058	3	Q6GMY2

Serum amyloid P-component precursor (SAP) (9.5S alpha-1-glycoprotein) [Contains: S

SAMP_HUMAN	K.IVLGQEQDSYGGK.F	1	3.4325	0.381		P02743
SAMP_HUMAN	R.QGYFVEAQPK.I	1	2.982	0.4275		P02743

1

Alpha-2-HS-glycoprotein precursor (Fetuin-A) (Alpha-2-Z-globulin) (Ba-alpha-2-glycopr

FETUA_HUMAN	K.HTLNQIDEVK.V	1	3.3442	0.4234		P02765

Alpha-2-macroglobulin precursor (Alpha-2-M) - Homo sapiens (Human)

A2MG_HUMAN	K.AIGYLNTGYQR.Q	1	2.6834	0.4142		P01023

AMBP protein precursor [Contains: Alpha-1-microglobulin (Protein HC) (Complex-formi

AMBP_HUMAN	R.AFIQLWAFDAVK.G	1	3.8391	0.4943		P02760

Clusterin precursor (Complement-associated protein SP-40,40) (Complement cytolysis

CLUS_HUMAN	R.ELDESLQVAER.L	1	3.2778	0.4208	1	P10909

Complement C1s subcomponent precursor (EC 3.4.21.42) (C1 esterase) [Contains: Co

C1S_HUMAN	R.TNFDNDIALVR.L	1	2.9965	0.4325		P09871

IGHA1 protein - Homo sapiens (Human)

Q8N5K4_HUMAN	R.DASGVTFTWTPSSGK.S	1	4.8686	0.5423	12	Q8N5K4

IGLV2-14 protein - Homo sapiens (Human)

A0A5C9_HUMAN	K.YAASSYLSLTPEQWK.S	1	4.9738	0.6582	29	A0A5C9

Immunoglobulin J chain - Homo sapiens (Human)

IGJ_HUMAN	R.SSEDPNEDIVER.N	1	3.0033	0.3247		P01591

Latrophilin-2 (Fragment) - Homo sapiens (Human)

Q9UJ49_HUMAN	-.MTGNYLLTNPLLR.P	1	2.6732	0.3671		Q9UJ49
Serum paraoxonase/arylesterase 1 (EC 3.1.1.2) (EC 3.1.8.1) (PON 1) (Serum aryldialk						
PON1_HUMAN	K.IFFYDSENPPASEVLR.I	1	5.4049	0.5888		P27169
Transcription factor-like 5 protein (Cha transcription factor) (HPV-16 E2-binding protein						
TCFL5_HUMAN	R.MMLLSEAGAAEKTSGGGDG/	1	3.4634	0.1908	1	Q9UL49

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

lement component C9b] - Homo sapiens (Human)

lomo sapiens (Human)

erum amyloid P-component(1-203)] - Homo sapiens (Human)

·inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

otein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

Homo sapiens (Human)

ens (Human)

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

ns (Human)

mo sapiens (Human)

cting inhibitor of NF-kappaB) (SINK) - Homo sapiens (Human)

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

Ɨomo sapiens (Human)

] - Homo sapiens (Human)

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

lement component C9b] - Homo sapiens (Human)

·inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

' chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

Homo sapiens (Human)

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma c

·inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

mo sapiens (Human)

otein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

r subcomponent) [Contains: Complement C1r subcomponent heavy chain; Complement C1r subcor

ɔoL) - Homo sapiens (Human)

ʂapiens (Human)

N polypeptide 1) (Carboxypeptidase N small subunit) (Lysine carboxypeptidase) (Arginine carboxype

peptidase N 83 kDa chain) (Carboxypeptidase N regulatory subunit) (Carboxypeptidase N large sub

2) (HLS2) - Homo sapiens (Human)

omo sapiens (Human)

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

er factor) [Contains: Plasma kallikrein heavy chain; Plasma kallikrein light chain] - Homo sapiens (Hu

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

ɔ-less] - Homo sapiens (Human)

natrix receptor-III) (ECMR-III) (GP90 lymphocyte homing/adhesion receptor) (Hermes antigen) (Hya

ıo sapiens (Human)

' chain neuron-specific 1) - Homo sapiens (Human)

V65 subunit; Vitronectin V10 subunit; Somatomedin B] - Homo sapiens (Human)

ıs (Human)

ɛns (Human)

cadherin) (P105) - Homo sapiens (Human)

) (CaMK IV) - Homo sapiens (Human)

NK MAP kinase scaffold protein 1) (Islet-brain 1) (IB-1) (Mitogen-activated protein kinase 8-interactin

omplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

chain; Complement C4-B alpha chain; C4a anaphylatoxin; C4b-B; C4d-B; Complement C4 gamma c

2 subunit alpha) (eIF-2-alpha) (EIF-2alpha) (EIF-2A) - Homo sapiens (Human)

e RNA-binding protein Q99) - Homo sapiens (Human)

omo sapiens (Human)

eotide-exchange protein 100) (ADP-ribosylation factors guanine nucleotide-exchange protein 2) (Bref

ɔtein 2) - Homo sapiens (Human)

ɔn termination factor-like protein) (mTERFL) - Homo sapiens (Human)

A125) (CA-125) - Homo sapiens (Human)

ıs (Human)

(Human)

P3-like protein) - Homo sapiens (Human)

ıibiting factor) - Homo sapiens (Human)

ZI) (Serpin A10) - Homo sapiens (Human)

le fragment 1; Activation peptide fragment 2; Thrombin light chain; Thrombin heavy chain] - Homo sa

ı response element-TATA box-binding protein) (HSP27 ERE-TATA-binding protein) - Homo sapiens

iens (Human)

iens (Human)

ɬomo sapiens (Human)

ɲ) (C-SAA) - Homo sapiens (Human)

ɬuman)

rand antigen II)] - Homo sapiens (Human)

·inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

Homo sapiens (Human)

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma c

] - Homo sapiens (Human)

·inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

›iens (Human)

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

›mplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

erum amyloid P-component(1-203)] - Homo sapiens (Human)

mo sapiens (Human)

ng glycoprotein heterogeneous in charge) (Alpha-1 microglycoprotein); Inter-alpha-trypsin inhibitor li

·inhibitor heavy chain 3) (Serum-derived hyaluronan-associated protein) (SHAP) - Homo sapiens (Hu

V65 subunit; Vitronectin V10 subunit; Somatomedin B] - Homo sapiens (Human)

Homo sapiens (Human)

ens (Human)

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

r subcomponent) [Contains: Complement C1r subcomponent heavy chain; Complement C1r subcor

lement component C9b] - Homo sapiens (Human)

2) (HLS2) - Homo sapiens (Human)

o sapiens (Human)

(Human)

le fragment 1; Activation peptide fragment 2; Thrombin light chain; Thrombin heavy chain] - Homo sa

it rpn11) (26S proteasome-associated PAD1 homolog 1) - Homo sapiens (Human)

an)

-lyase 6) (Adenylyl cyclase 6) (Ca(2+)-inhibitable adenylyl cyclase) - Homo sapiens (Human)

ens (Human)

n-2 (Angiotensin II) (Ang II); Angiotensin-3 (Angiotensin III) (Ang III) (Des-Asp[1]-angiotensin II)] - H

oL) - Homo sapiens (Human)

Human)

1) (p200 ARF-GEP1) (p200 ARF guanine nucleotide exchange factor) - Homo sapiens (Human)

an)

an)

a mRNA. (Fragment) - Homo sapiens (Human)

sapiens (Human)

piens (Human)

- Homo sapiens (Human)

ens (Human)

omo sapiens (Human)

lpha-IEL) (CD103 antigen) [Contains: Integrin alpha-E light chain; Integrin alpha-E heavy chain] - Ho

chain neuron-specific 1) - Homo sapiens (Human)

omo sapiens (Human)

0) (gp330) - Homo sapiens (Human)

gene sequences - Homo sapiens (Human)

ns (Human)

natase FYVE-DSP2) - Homo sapiens (Human)

ansferase) [Contains: N-acetylglutamate synthase long form; N-acetylglutamate synthase short form

.6.5.3) (EC 1.6.99.3) (NADH-ubiquinone oxidoreductase 49 kDa subunit) (Complex I-49kD) (CI-49kD

elope protein 2) (Nucleus and actin connecting element protein) (Protein NUANCE) - Homo sapiens

homolog) - Homo sapiens (Human)

ceptor P2X-like 1) - Homo sapiens (Human)

ZI) (Serpin A10) - Homo sapiens (Human)

ruvate kinase 2/3) (Cytosolic thyroid hormone-binding protein) (CTHBP) (THBP1) - Homo sapiens (H

osine phosphatase kappa) (R-PTP-kappa) - Homo sapiens (Human)

L) - Homo sapiens (Human)

nan)

łuman)

nan)

uccinyl-CoA synthetase, betaG chain) (SCS-betaG) (GTP-specific succinyl-CoA synthetase subunit

SIII subunit C) (SIII p15) (Elongin-C) (EloC) (Elongin 15 kDa subunit) - Homo sapiens (Human)

1) (E2BP-1) - Homo sapiens (Human)

Human)

rand antigen II)] - Homo sapiens (Human)

NG1) (Zinc finger and BTB domain-containing protein 22A) - Homo sapiens (Human)

ıs T-cell lymphoma-associated antigen se33-1) (CTCL tumor antigen se33-1) - Homo sapiens (Huma

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

Homo sapiens (Human)

-inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

-inhibitor heavy chain 3) (Serum-derived hyaluronan-associated protein) (SHAP) - Homo sapiens (Hu

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma c

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

)iens (Human)

] - Homo sapiens (Human)

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

erum amyloid P-component(1-203)] - Homo sapiens (Human)

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

V65 subunit; Vitronectin V10 subunit; Somatomedin B] - Homo sapiens (Human)

ng glycoprotein heterogeneous in charge) (Alpha-1 microglycoprotein); Inter-alpha-trypsin inhibitor li

otein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

Homo sapiens (Human)

mplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

active factor serine protease p100) (RaRF) (Mannan-binding lectin serine protease 1) (Mannose-bin

inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

ns (Human)

le fragment 1; Activation peptide fragment 2; Thrombin light chain; Thrombin heavy chain] - Homo sa

a adipocyte complement-related protein) (Adipocyte complement-related 30 kDa protein) (ACRP30)

ens (Human)

Human)

AUTOANTIGEN B - Homo sapiens (Human)

sapiens (Human)

CHD8) (CHD-8) (Helicase with SNF2 domain 1) - Homo sapiens (Human)

lement component C9b] - Homo sapiens (Human)

gen domain-containing lectin 3 p35) (Hakata antigen) - Homo sapiens (Human)

omo sapiens (Human)

gase URE-B1) (Mcl-1 ubiquitin ligase E3) (Mule) (ARF-binding protein 1) (ARF-BP1) - Homo sapiens

2) (HLS2) - Homo sapiens (Human)

o sapiens (Human)

(hLAT2) - Homo sapiens (Human)

ı gene sequences - Homo sapiens (Human)

(Human)

ıatase FYVE-DSP2) - Homo sapiens (Human)

P3-like protein) - Homo sapiens (Human)

or regulatory protein) (Prostaglandin F2-alpha receptor-associated protein) (CD9 partner 1) (CD9P-1

n)

(Rhomboid veinlet-like protein 6) - Homo sapiens (Human)

n) (C-SAA) - Homo sapiens (Human)

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma c

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

·inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

Homo sapiens (Human)

)mplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

·inhibitor heavy chain 3) (Serum-derived hyaluronan-associated protein) (SHAP) - Homo sapiens (Hu

ng glycoprotein heterogeneous in charge) (Alpha-1 microglycoprotein); Inter-alpha-trypsin inhibitor li

)iens (Human)

Homo sapiens (Human)

erum amyloid P-component(1-203)] - Homo sapiens (Human)

V65 subunit; Vitronectin V10 subunit; Somatomedin B] - Homo sapiens (Human)

] - Homo sapiens (Human)

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

otein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

·inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

ıibiting factor) - Homo sapiens (Human)

le fragment 1; Activation peptide fragment 2; Thrombin light chain; Thrombin heavy chain] - Homo sa

apiens (Human)

Homo sapiens (Human)

chain; Complement C4-B alpha chain; C4a anaphylatoxin; C4b-B; C4d-B; Complement C4 gamma c

lement component C9b] - Homo sapiens (Human)

active factor serine protease p100) (RaRF) (Mannan-binding lectin serine protease 1) (Mannose-bin

ein) (TAXREB302) - Homo sapiens (Human)

ein 1) - Homo sapiens (Human)

in 71) (ZNF47) - Homo sapiens (Human)

TCL tumor antigen se1-1) (CLL-associated antigen KW-11) (Renal carcinoma antigen NY-REN-53)

o sapiens (Human)

nyosin heavy chain IIb) (NMMHC II-b) (NMMHC-IIB) (Cellular myosin heavy chain, type B) (Nonmus

(Human)

P3-like protein) - Homo sapiens (Human)

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma

·inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

Homo sapiens (Human)

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

erum amyloid P-component(1-203)] - Homo sapiens (Human)

mplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

·inhibitor heavy chain 3) (Serum-derived hyaluronan-associated protein) (SHAP) - Homo sapiens (Hu

›iens (Human)

›tein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

Homo sapiens (Human)

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

ıs (Human)

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

V65 subunit; Vitronectin V10 subunit; Somatomedin B] - Homo sapiens (Human)

ng glycoprotein heterogeneous in charge) (Alpha-1 microglycoprotein); Inter-alpha-trypsin inhibitor li

peptidase N 83 kDa chain) (Carboxypeptidase N regulatory subunit) (Carboxypeptidase N large sub

lement component C9b] - Homo sapiens (Human)

inhibitor heavy chain 4) (Inter-alpha-trypsin inhibitor family heavy chain-related protein) (IHRP) (Plas

ein 1) (Syne-1) (Myocyte nuclear envelope protein 1) (Myne-1) (Enaptin) - Homo sapiens (Human)

le fragment 1; Activation peptide fragment 2; Thrombin light chain; Thrombin heavy chain] - Homo sa

(Intramembrane protease 4) (IMP4) (Presenilin-like protein 1) - Homo sapiens (Human)

arbonate cotransporter 2b) (Bicarbonate transporter) (Solute carrier family 4 member 7) - Homo sap

SIII subunit C) (SIII p15) (Elongin-C) (EloC) (Elongin 15 kDa subunit) - Homo sapiens (Human)

cting inhibitor of NF-kappaB) (SINK) - Homo sapiens (Human)

·inhibitor heavy chain 1) (Inter-alpha-trypsin inhibitor complex component III) (Serum-derived hyaluro

·inhibitor heavy chain 2) (Inter-alpha-trypsin inhibitor complex component II) (Serum-derived hyaluro

Homo sapiens (Human)

ha chain; C3a anaphylatoxin; Complement C3b alpha' chain; Complement C3c alpha' chain fragmer

chain; Complement C4-A alpha chain; C4a anaphylatoxin; C4b-A; C4d-A; Complement C4 gamma c

chain; Bradykinin (Kallidin I); Lysyl-bradykinin (Kallidin II); Kininogen-1 light chain; Low molecular w

Homo sapiens (Human)

erum amyloid P-component(1-203)] - Homo sapiens (Human)

otein) [Contains: Alpha-2-HS-glycoprotein chain A; Alpha-2-HS-glycoprotein chain B] - Homo sapien

ng glycoprotein heterogeneous in charge) (Alpha-1 microglycoprotein); Inter-alpha-trypsin inhibitor li

inhibitor) (CLI) (NA1/NA2) (Apolipoprotein J) (Apo-J) (Testosterone-repressed prostate message 2)

mplement C1s subcomponent heavy chain; Complement C1s subcomponent light chain] - Homo sa

ylphosphatase 1) (A-esterase 1) (Aromatic esterase 1) (K-45) - Homo sapiens (Human)

1) (E2BP-1) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

ɜma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

s (Human)

eight growth-promoting factor] - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

ɜma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

eight growth-promoting factor] - Homo sapiens (Human)

chain] - Homo sapiens (Human)

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

s (Human)

ɱponent light chain] - Homo sapiens (Human)

eptidase) (Kininase-1) (Serum carboxypeptidase N) (SCPN) (Anaphylatoxin inac

unit) - Homo sapiens (Human)

nan-associated protein) (SHAP) - Homo sapiens (Human)

uman)

luronate receptor) (Heparan sulfate proteoglycan) (Epican) (CDw44) - Homo s

g protein 1) - Homo sapiens (Human)

piens (Human)

hain] - Homo sapiens (Human)

feldin-resistant Arf-GEF 2 protein) - Homo sapiens (Human)

apiens (Human)

(Human)

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

nan-associated protein) (SHAP) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

eight growth-promoting factor] - Homo sapiens (Human)

chain] - Homo sapiens (Human)

ɜma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

ıpiens (Human)

ght chain (ITI-LC) (Bikunin) (HI-30)] - Homo sapiens (Human)

ıman)

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

nponent light chain] - Homo sapiens (Human)

apiens (Human)

omo sapiens (Human)

ɔmo sapiens (Human)

; N-acetylglutamate synthase conserved domain form] - Homo sapiens (Human

D) - Homo sapiens (Human)

(Human)

luman)

beta) - Homo sapiens (Human)

an)

nan-associated protein) (SHAP) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

ɹman)

chain] - Homo sapiens (Human)

eight growth-promoting factor] - Homo sapiens (Human)

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

ght chain (ITI-LC) (Bikunin) (HI-30)] - Homo sapiens (Human)

s (Human)

piens (Human)

ding protein-associated serine protease) (MASP-1) (Serine protease 5) [Con

sma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

apiens (Human)

(Adipose most abundant gene transcript 1) (apM-1) (Gelatin-binding protein)

s (Human)

l) (CD315 antigen) - Homo sapiens (Human)

chain] - Homo sapiens (Human)

nan-associated protein) (SHAP) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

ɹpiens (Human)

ɹman)

ght chain (ITI-LC) (Bikunin) (HI-30)] - Homo sapiens (Human)

eight growth-promoting factor] - Homo sapiens (Human)

s (Human)

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

sma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

apiens (Human)

chain] - Homo sapiens (Human)

ding protein-associated serine protease) (MASP-1) (Serine protease 5) [Con

- Homo sapiens (Human)

cle myosin heavy chain-B) (NMMHC-B) - Homo sapiens (Human)

chain] - Homo sapiens (Human)

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

nan-associated protein) (SHAP) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

ıpiens (Human)

ɹman)

s (Human)

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

eight growth-promoting factor] - Homo sapiens (Human)

ght chain (ITI-LC) (Bikunin) (HI-30)] - Homo sapiens (Human)

unit) - Homo sapiens (Human)

ɜma kallikrein sensitive glycoprotein 120) (PK-120) (GP120) [Co

apiens (Human)

iens (Human)

ɔnan-associated protein) (SHAP) - Homo sapiens (Human)

nan-associated protein) (SHAP) - Homo sapiens (Human)

ıt 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Com

chain] - Homo sapiens (Human)

eight growth-promoting factor] - Homo sapiens (Human)

s (Human)

ght chain (ITI-LC) (Bikunin) (HI-30)] - Homo sapiens (Human)

(TRPM-2) [Contains: Clusterin beta chain (ApoJalpha) (Complement cytolysis

piens (Human)