

Protein Name	Peptide Sequence	Precursor Mz	Precursor Charge
sp P17931 LEG3_HUMAN	IALDFQR	431.742702	2
sp P17931 LEG3_HUMAN	IALDFQR	436.746837	2
sp P17931 LEG3_HUMAN	GNDVAFHFNPR	425.207266	3
sp P17931 LEG3_HUMAN	GNDVAFHFNPR	428.543355	3
sp P10451 OSTP_HUMAN	AIPVAQDLNAPSDWDSR	927.952671	2
sp P10451 OSTP_HUMAN	AIPVAQDLNAPSDWDSR	932.956806	2
sp P10451 OSTP_HUMAN	ANDESNEHSDVIDSQELSK	706.313693	3
sp P10451 OSTP_HUMAN	ANDESNEHSDVIDSQELSK	708.985092	3
sp P35222 CTNB1_HUMAN	LLNDEDQVVVNK	693.367181	2
sp P35222 CTNB1_HUMAN	LLNDEDQVVVNK	697.37428	2
sp P35222 CTNB1_HUMAN	LVQLLVR	420.786913	2
sp P35222 CTNB1_HUMAN	LVQLLVR	425.791047	2
sp P30041 PRDX6_HUMAN	DFTPVCCTELGR	698.332284	2
sp P30041 PRDX6_HUMAN	DFTPVCCTELGR	703.336418	2
sp P30041 PRDX6_HUMAN	LSILYPATTGR	596.340238	2
sp P30041 PRDX6_HUMAN	LSILYPATTGR	601.344372	2
sp O00299 CLIC1_HUMAN	LAALNPESNTAGLDIFAK	922.991265	2
sp O00299 CLIC1_HUMAN	LAALNPESNTAGLDIFAK	926.998365	2
sp O00299 CLIC1_HUMAN	NSNPALNDNLEK	664.82568	2
sp O00299 CLIC1_HUMAN	NSNPALNDNLEK	668.832779	2
sp P14618 KPYM_HUMAN	NTGIICTIGPASR	680.356093	2
sp P14618 KPYM_HUMAN	NTGIICTIGPASR	685.360228	2
sp P14618 KPYM_HUMAN	APIIAVTR	420.76872	2
sp P14618 KPYM_HUMAN	APIIAVTR	425.772854	2
sp P07355 ANXA2_HUMAN	GVDEVTIVNILTNR	771.927937	2
sp P07355 ANXA2_HUMAN	GVDEVTIVNILTNR	776.932071	2
sp P07355 ANXA2_HUMAN	QDIAFAYQR	556.28018	2
sp P07355 ANXA2_HUMAN	QDIAFAYQR	561.284315	2
sp P15941 MUC1_HUMAN	QGGFLGLSNIK	567.319305	2
sp P15941 MUC1_HUMAN	QGGFLGLSNIK	571.326405	2
sp P15941 MUC1_HUMAN	EGTINVHDTVETQFNQYK	674.656954	3
sp P15941 MUC1_HUMAN	EGTINVHDTVETQFNQYK	677.328354	3
sp P55060 XPO2_HUMAN	ANIVHMLMLSSPEQIQK	603.329344	3
sp P55060 XPO2_HUMAN	ANIVHMLMLSSPEQIQK	606.000744	3
sp P55060 XPO2_HUMAN	LLQTDDEEEAGLLELLK	965.006778	2
sp P55060 XPO2_HUMAN	LLQTDDEEEAGLLELLK	969.013878	2
sp P50454 SERPH_HUMAN	GVVEVTHDLQK	408.890943	3
sp P50454 SERPH_HUMAN	GVVEVTHDLQK	411.562343	3
sp P50454 SERPH_HUMAN	DTQSGSLLFIGR	647.343509	2
sp P50454 SERPH_HUMAN	DTQSGSLLFIGR	652.347643	2
sp P16035 TIMP2_HUMAN	EVDSGNDIYGNIPIK	760.865002	2
sp P16035 TIMP2_HUMAN	EVDSGNDIYGNIPIK	764.872102	2
sp P09382 LEG1_HUMAN	SFVLNLGK	439.260728	2
sp P09382 LEG1_HUMAN	SFVLNLGK	443.267827	2
sp P09382 LEG1_HUMAN	LPDGYEFK	484.739825	2
sp P09382 LEG1_HUMAN	LPDGYEFK	488.746925	2
sp Q13938 CAYP1_HUMAN	EAVIAAAFAK	495.784567	2
sp Q13938 CAYP1_HUMAN	EAVIAAAFAK	499.791666	2
sp Q13938 CAYP1_HUMAN	SGDGVTVDDLR	616.809499	2
sp Q13938 CAYP1_HUMAN	SGDGVTVDDLR	621.813633	2
sp P42574 CASP3_HUMAN	SGTDVDAANLR	559.775459	2
sp P42574 CASP3_HUMAN	SGTDVDAANLR	564.779593	2
sp P16949 STMN1_HUMAN	SHEAEVLK	456.742899	2
sp P16949 STMN1_HUMAN	SHEAEVLK	460.749999	2
sp P16949 STMN1_HUMAN	AIEENNNFSK	583.277834	2
sp P16949 STMN1_HUMAN	AIEENNNFSK	587.284934	2
sp P14780 MMP9_HUMAN	SLGPALLLLQK	576.871174	2
sp P14780 MMP9_HUMAN	SLGPALLLLQK	580.878273	2
sp P14780 MMP9_HUMAN	AFALWSAVTPLTFTFR	840.95904	2
sp P14780 MMP9_HUMAN	AFALWSAVTPLTFTFR	845.963174	2
sp P08670 VIME_HUMAN	ILLAELEQLK	585.360639	2
sp P08670 VIME_HUMAN	ILLAELEQLK	589.367738	2
sp P08670 VIME_HUMAN	ISLPLPNFSSLNLR	785.951215	2
sp P08670 VIME_HUMAN	ISLPLPNFSSLNLR	790.955349	2
sp P16070 CD44_HUMAN	TEAADLCK	454.213117	2

sp P16070 CD44_HUMAN	TEAADLCK	458.220217	2
sp P15328 FOLR1_HUMAN	TELLNVCMNAK	646.820498	2
sp P15328 FOLR1_HUMAN	TELLNVCMNAK	650.827597	2
sp P15328 FOLR1_HUMAN	FNWNHCGEMAPACK	574.570893	3
sp P15328 FOLR1_HUMAN	FNWNHCGEMAPACK	577.242293	3
sp O43852 CALU_HUMAN	TFDQLTPEESK	647.811707	2
sp O43852 CALU_HUMAN	TFDQLTPEESK	651.818806	2
sp P09603 CSF1_HUMAN	AFLLVQDIMEDTMR	841.418033	2
sp P09603 CSF1_HUMAN	AFLLVQDIMEDTMR	846.422168	2
sp P09603 CSF1_HUMAN	TFYETPLQLLEK	741.39795	2
sp P09603 CSF1_HUMAN	TFYETPLQLLEK	745.405049	2
sp Q06830 PRDX1_HUMAN	ADEGISFR	447.719424	2
sp Q06830 PRDX1_HUMAN	ADEGISFR	452.723558	2
sp Q06830 PRDX1_HUMAN	LVQAFQFTDK	598.819138	2
sp Q06830 PRDX1_HUMAN	LVQAFQFTDK	602.826238	2
sp O75556 SG2A1_HUMAN	TINSDISIPEYK	690.356282	2
sp O75556 SG2A1_HUMAN	TINSDISIPEYK	694.363381	2
sp O75556 SG2A1_HUMAN	ELLQEFIDSDAAAEAMGK	646.643546	3
sp O75556 SG2A1_HUMAN	ELLQEFIDSDAAAEAMGK	649.314945	3
sp P22626 ROA2_HUMAN	TLETVPLER	529.298038	2
sp P22626 ROA2_HUMAN	TLETVPLER	534.302173	2
sp P22626 ROA2_HUMAN	GGNFGFGDSR	507.225405	2
sp P22626 ROA2_HUMAN	GGNFGFGDSR	512.229539	2
sp P09211 GSTP1_HUMAN	ASCLYGQLPK	568.792066	2
sp P09211 GSTP1_HUMAN	ASCLYGQLPK	572.799166	2
sp P09211 GSTP1_HUMAN	TLGLYGK	376.221071	2
sp P09211 GSTP1_HUMAN	TLGLYGK	380.228171	2
sp Q01995 TAGL_HUMAN	TLMALGSLAVTK	602.852124	2
sp Q01995 TAGL_HUMAN	TLMALGSLAVTK	606.859223	2
sp Q01995 TAGL_HUMAN	EFTESQLQEGK	648.309331	2
sp Q01995 TAGL_HUMAN	EFTESQLQEGK	652.316431	2
sp P00441 SODC_HUMAN	AVCVLK	345.204367	2
sp P00441 SODC_HUMAN	AVCVLK	349.211466	2
sp P00441 SODC_HUMAN	TLVVHEK	413.245078	2
sp P00441 SODC_HUMAN	TLVVHEK	417.252177	2
sp P04083 ANXA1_HUMAN	GGPGSAVSPYPTFNPSSDVAALHK	786.057118	3
sp P04083 ANXA1_HUMAN	GGPGSAVSPYPTFNPSSDVAALHK	788.728517	3
sp P04083 ANXA1_HUMAN	DITSDTSGDFR	607.270206	2
sp P04083 ANXA1_HUMAN	DITSDTSGDFR	612.274341	2
sp Q01469 FABP5_HUMAN	ELGVGIALR	464.284734	2
sp Q01469 FABP5_HUMAN	ELGVGIALR	469.288869	2
sp Q01469 FABP5_HUMAN	LVVECMNNTCTR	847.904767	2
sp Q01469 FABP5_HUMAN	LVVECMNNTCTR	852.908902	2
sp P08253 MMP2_HUMAN	IDAVYEAPQEEK	696.338089	2
sp P08253 MMP2_HUMAN	IDAVYEAPQEEK	700.345188	2
sp P08253 MMP2_HUMAN	VDAAFNWSK	519.256174	2
sp P08253 MMP2_HUMAN	VDAAFNWSK	523.263273	2
sp P07237 PDIA1_HUMAN	ALAPEYAK	431.737085	2
sp P07237 PDIA1_HUMAN	ALAPEYAK	435.744185	2
sp P07237 PDIA1_HUMAN	ILEFFGLK	483.786578	2
sp P07237 PDIA1_HUMAN	ILEFFGLK	487.793678	2
sp P12830 CADH1_HUMAN	NLVQIK	357.729063	2
sp P12830 CADH1_HUMAN	NLVQIK	361.736163	2
sp P12830 CADH1_HUMAN	VFYSITGQGADTPPVGVFIIR	789.417875	3
sp P12830 CADH1_HUMAN	VFYSITGQGADTPPVGVFIIR	792.753965	3
sp P61604 CH10_HUMAN	FLPLFDR	454.255445	2
sp P61604 CH10_HUMAN	FLPLFDR	459.25958	2
sp P61604 CH10_HUMAN	VLLPEYGGTK	538.802957	2
sp P61604 CH10_HUMAN	VLLPEYGGTK	542.810056	2
sp P30086 PEBP1_HUMAN	VLTPQVK	443.273835	2
sp P30086 PEBP1_HUMAN	VLTPQVK	447.280935	2
sp P30086 PEBP1_HUMAN	LYEQLSGK	469.2531	2
sp P30086 PEBP1_HUMAN	LYEQLSGK	473.260199	2
sp P09466 PAEP_HUMAN	VLVEDDEIMQGFIR	832.42163	2
sp P09466 PAEP_HUMAN	VLVEDDEIMQGFIR	837.425764	2
sp P80188 NGAL_HUMAN	VPLQQNFQDNQFQGGK	597.632192	3

sp P80188 NGAL_HUMAN	VPLQQNFQDNQFQ GK	600.303591	3
sp P80188 NGAL_HUMAN	ELTSELK	410.22655	2
sp P80188 NGAL_HUMAN	ELTSELK	414.23365	2
sp Q14508 WFDC2_HUMAN	EGSCPQVNINFPQLGLCR	697.003303	3
sp Q14508 WFDC2_HUMAN	EGSCPQVNINFPQLGLCR	700.339392	3
sp Q14508 WFDC2_HUMAN	VSCVTPNF	462.218203	2
sp Q14508 WFDC2_HUMAN	VSCVTPNF	467.231817	2
sp P62937 PPIA_HUMAN	VSFELFADK	528.274032	2
sp P62937 PPIA_HUMAN	VSFELFADK	532.281132	2
sp P62937 PPIA_HUMAN	FEDENFILK	577.790046	2
sp P62937 PPIA_HUMAN	FEDENFILK	581.797146	2
sp P04792 HSPB1_HUMAN	LFDQAFGLPR	582.314023	2
sp P04792 HSPB1_HUMAN	LFDQAFGLPR	587.318157	2
sp P04792 HSPB1_HUMAN	LATQSN EITIPVTFESR	953.499455	2
sp P04792 HSPB1_HUMAN	LATQSN EITIPVTFESR	958.503589	2
sp P07339 CATD_HUMAN	VSTLPAILTK	521.828974	2
sp P07339 CATD_HUMAN	VSTLPAILTK	525.836074	2
sp P07339 CATD_HUMAN	VGFAEAAR	410.719227	2
sp P07339 CATD_HUMAN	VGFAEAAR	415.723361	2
sp P11166 GTR1_HUMAN	TFDEIASGFR	571.77747	2
sp P11166 GTR1_HUMAN	TFDEIASGFR	576.781605	2
sp P00338 LDHA_HUMAN	LVITAGAR	457.295102	2
sp P00338 LDHA_HUMAN	LVITAGAR	462.299236	2
sp P00338 LDHA_HUMAN	VTLTSEEEAR	567.785492	2
sp P00338 LDHA_HUMAN	VTLTSEEEAR	572.789627	2
sp P60174 TPIS_HUMAN	VVLAYEPVWAIGTGK	801.948141	2
sp P60174 TPIS_HUMAN	VVLAYEPVWAIGTGK	805.95524	2
sp P60174 TPIS_HUMAN	SNVSDAVAQSTR	617.804748	2
sp P60174 TPIS_HUMAN	SNVSDAVAQSTR	622.808882	2
sp P05164 PERM_HUMAN	IANVFTNAFR	576.811647	2
sp P05164 PERM_HUMAN	IANVFTNAFR	581.815782	2
sp P05164 PERM_HUMAN	VVLEGGIDPILR	640.88227	2
sp P05164 PERM_HUMAN	VVLEGGIDPILR	645.886404	2
sp P14174 MIF_HUMAN	LLCGLLAER	522.797151	2
sp P14174 MIF_HUMAN	LLCGLLAER	527.801286	2
sp P14174 MIF_HUMAN	VYINYDDMNAANVGWNNSTFA	809.695191	3
sp P14174 MIF_HUMAN	VYINYDDMNAANVGWNNSTFA	813.037601	3
sp P01833 PIGR_HUMAN	GGCITLISSEGYVSSK	829.40872	2
sp P01833 PIGR_HUMAN	GGCITLISSEGYVSSK	833.415819	2
sp P01833 PIGR_HUMAN	VYTVDLGR	461.753267	2
sp P01833 PIGR_HUMAN	VYTVDLGR	466.757401	2
sp P05787 K2C8_HUMAN	WSLLQQQK	515.787641	2
sp P05787 K2C8_HUMAN	WSLLQQQK	519.79474	2
sp P05787 K2C8_HUMAN	LSELEAALQR	565.31422	2
sp P05787 K2C8_HUMAN	LSELEAALQR	570.318354	2
sp O43278 SPIT1_HUMAN	WYYDPTEQICK	751.834659	2
sp O43278 SPIT1_HUMAN	WYYDPTEQICK	755.841759	2
sp O43278 SPIT1_HUMAN	SFVYGGCLGNK	601.287148	2
sp O43278 SPIT1_HUMAN	SFVYGGCLGNK	605.294247	2
sp P06733 ENOA_HUMAN	TIAPALVSK	450.28166	2
sp P06733 ENOA_HUMAN	TIAPALVSK	454.28876	2
sp P06733 ENOA_HUMAN	YISPDQLADLYK	713.366649	2
sp P06733 ENOA_HUMAN	YISPDQLADLYK	717.373749	2
sp Q16658 FSCN1_HUMAN	FLIVAHDDGR	381.536612	3
sp Q16658 FSCN1_HUMAN	FLIVAHDDGR	384.872701	3
sp Q16658 FSCN1_HUMAN	YLAPSGPSGTLK	595.82442	2
sp Q16658 FSCN1_HUMAN	YLAPSGPSGTLK	599.83152	2
sp P43490 NAMPT_HUMAN	YDGHLP I EIK	395.548517	3
sp P43490 NAMPT_HUMAN	YDGHLP I EIK	398.219917	3
sp P43490 NAMPT_HUMAN	YLLETSGNLDGLE YK	857.930342	2
sp P43490 NAMPT_HUMAN	YLLETSGNLDGLE YK	861.937442	2
sp P40121 CAPG_HUMAN	YQEGGVESAFHK	451.214086	3
sp P40121 CAPG_HUMAN	YQEGGVESAFHK	453.885486	3
sp P40121 CAPG_HUMAN	EGNPEEDLTADK	659.293878	2
sp P40121 CAPG_HUMAN	EGNPEEDLTADK	663.300978	2