

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 PIIA_HUMAN	VSFELFADK	528.274032	2
sp P62937 PIIA_HUMAN	VSFELFADK	532.281132	2
sp P62937 PIIA_HUMAN	FEDENFILK	577.790046	2
sp P62937 PIIA_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