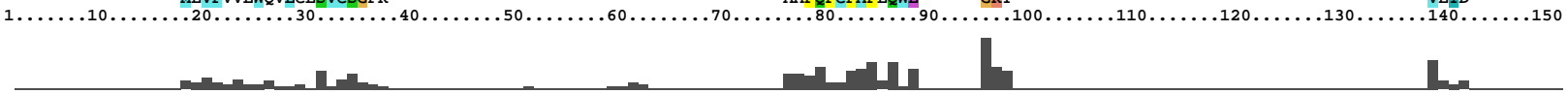


gi	31542612	ref	NP_598826.2	-----MPARAPRRRLVQPPRGTWLLGSL-----WVWVLCGLGMAGS-----LGTFPQPCQAPQWE-----GRQ-----	150
gi	56090361	ref	NP_001007626.1	-----MLTRAPRRRLVQPPRETWLLGGL-----WVWVLCGLGMAGS-----PGTQPQCAQAPQWE-----GRQ-----	150
gi	24475586	ref	NP_060019.1	MPPRMLVAPLPLHFLLAAVSSENRSGRRWQQAVAAGSGPGRNSSRAIHWSLPRALVPAATGTRASGSKRQRPKQGEALWSRPHSRHSAERAGGSHSDPGRLHAPLGLGLPEGRGLHAMPGRAPLRTVPGALGAWLLGGLWAWLLCGL	150
gi	73981923	ref	XP_851744.1	-----MDSPLKSIQVCLYSVVMCQG-----LHWAK-----AAGT-----GQPRLLQRAHSRR-----GGST-----	150
gi	156121221	ref	NP_001095758.1	-----MPPRAPLRVARGSLDANLLGGLWVCALGCLCGVMAAPGAGAGAGGS-----LGAQRPCQAPQWE-----GRQ-----	150
gi	118086182	ref	XP_418830.2	-----MALATGGLLLLGLLVRGG-----AGAEPCRAPQWE-----GRS-----	150
gi	50539892	ref	NP_001002416.1	-----MLVEVVLWQVLCVCSGFR-----AAPQCPAPLQWE-----GRT-----	150



gi	31542612	ref	NP_598826.2	-----GSSGHNNRALVSYDGLNQRVRVLDERKALIPCKRLEFYILLYKDGVMFQIEQATKQCAKIPLVESWDPDLDIPONSTFEDQYSIGGPEQILVQEWSDRRRTARSYETWIGVYTKDCYVPQETFI	300
gi	56090361	ref	NP_001007626.1	-----GSSGHNSRALVSYDGLNQRVRVLDERKALIPCKRLEFYILLYKDGVMFQIEQATKQCAKIPLVESWDPDLDIPONSTFEDQYSIGGPEQIMVQEWSDRRRTARSYETWIGVYTKDCYVPQETFI	300
gi	24475586	ref	NP_060019.1	CSLGAVGAPRRCQAPQQWEGRQVMYQSSGRNSRALLSYDGLNQRVRVLDERKALIPCKRLEFYILLYKDGVMFQIDQATKQCSKMTLTPQWDPDLDIPONSTFEDQYSIGGPEQITVQEWSDRKSARSYETWIGIYTVKDCYVPQETFT	300
gi	73981923	ref	XP_851744.1	-----LFEYILLYEDGVMFQIEQATKQCSKITLTPDWDPLDIPONSTFEDQYSIGGPEQITVQEWSDRKSARSYETWIGIYTVKDCYVPQETFT	300
gi	156121221	ref	NP_001095758.1	-----DSTGRYSRALLSYDGLNQRVRVLDERKALIPCKRLEFYILLYKDGVMFQIEQATKQCSKITLTPDWDPLDIPONSTFEDQYSIGGPEQITVQEWSDRKSARSYETWIGIYTVKDCYVPQETFT	300
gi	118086182	ref	XP_418830.2	-----HGTGRNTRAAVSYDGLNQRRLRILEERKALIPCKKFFFYILLYKDAVMFQIEQVTKLCSKITLTPDWDPPYDIPNTSTFEDQYVIGGPGDQIMVQEWSDRKPARKLESWVGVIYTVKDCYVPQETFT	300
gi	50539892	ref	NP_001002416.1	-----HGTGRNTRAAVSYDAQNQRIRVLECKTGHIPCKKFFFYILLYKFSGLVLFQIEEITKQCAKIALTEAWDPYDIPNTSTFEDQYVIGGPGDMIEVQEWSDRKPARKNEAWVGVIYTVKDCYVPQETFT	300



gi	31542612	ref	NP_598826.2	RNYIVVMSTRFFDVQLSIKIDPSVFTPPSTCQAAQPEKMSDGCSL-	345
gi	56090361	ref	NP_001007626.1	RNYIVVLSTRFFDVQLGIKIDPSVFTPPSTCQAAQPEKMKENCSL-	345
gi	24475586	ref	NP_060019.1	INYSVILSTRFFDIQLGIKIDPSVFTPPSTCQAAQPEKMSDGCSSW-	345
gi	73981923	ref	XP_851744.1	RNYSVLLSTRFFDIQLGIKIDPSVFTPPSTCQAAQPEPMSENCSSW-	345
gi	156121221	ref	NP_001095758.1	KNYSVILSTRFFDIQLGIKIDPSVFTPPSTCQAAQPERMSEECSSW-	345
gi	118086182	ref	XP_418830.2	KNYSVITSTRFFDLKLGIAIDPSVFTPPSTCQAAQPRKMKDEC--	345
gi	50539892	ref	NP_001002416.1	KNYSVITSTRFFDLQLGISDPEVFNPPSTCLALSKRMTSDC--	345

