

gi|117422433|ref|NP_001070968.1| MRAPHLHLSAASGAR-ALAK--LLP LLMAQLWAAEAALLPQNDTRLDPEAYGSPCARGSPQWQVSLFNGLSFHCAGVLVDQSWVLTAAHCGN-KPLWARVGDHLLLLQGEQLRRTTRSVVHPKYHOGSGPILPRRTDEHDLMLLKLARP 150
gi|114678731|ref|XP_001174085.1| MRAPHLHLSAASGAR-ALAK--LLTLLMAQLWAAEAALLPKNDTRLDPEAYGAPCARGSPQWQVSLFNGLSFHCAGVLVDQSWVLTAAHCGN-KPLWARVGDHLLLLQGEQLRRTTRSVVHPKYHOGSGPILPRRTDEHDLMLLKLARP 150
gi|115497120|ref|NP_001069358.1| MTRHLHLSAASGAQGGGLKLLLLP LLVAQFVSEALLLPANDTSSNFVASGAPCARGSPQWQVSLFNGLSFHCAGVLVDQSWVLTAAHCGNKKPLWARVGDHLLLLQGEQLRRTSRPIVHPKYQOQSGPILPRRTDEHDLMLLKLGRP 150
gi|19526892|ref|NP_598473.1| MRVPLHLSTASGSW-SLVK-LLLPLLMVQLWAAQALLLPGNATRVLDLEASGAQCERDYHPQVSLFHNLFQFCAGVLVDQNWVLTAAHCWRNKPLRARVGDHLLLFQKEQLRSTSSPVFHPKYQACSGPILPHRSDEHDLMLKLSFP 150
gi|51870931|ref|NP_001004100.1| MRVPLHLPTASGSL-SLVK-LLLPLLMQLWAAQALLLPGNATRVLDLEAFGLCPVSPQWQVSLFHNLFQFCAGVLVDQNWVLTAAHCWRNKPLRARVGDHLLLFQSEQLRSTNSPVFHPKYQPCSGPVLPLRSDEHDLMLKLSFP 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



gi|117422433|ref|NP_001070968.1| VVLGPRVRLQLPYRCAQPGDCQVAGWGTAAARRVKYKGLTCSSTIILSPKECEVFYPGVVTNNMICAGLDRGDPQSDSGGGLVCDLTLGILSWGVIYPCGSA-OHPAVYTIQCKYMSWINKVIRSN 281
gi|114678731|ref|XP_001174085.1| VVLGPRARALQLPYRCAQPGDCQVAGWGTAAARRVKYKGLTCSSTIILSPKECEVFYPGVVTNNMICAGLDRGDPQSDSGGGLVCDLTLGILSWGVIYPCGSA-OHPAVYTIQCKYMSWINKVIRSN 281
gi|115497120|ref|NP_001069358.1| VVPGPRVRLRLPFRCQPGDCQIAGWATTSRRRVKYNKGLTCSRVTIILSPKECEVFYPGVITSNMMCAGLDQGDQSDSGGGLVCDLTLGILSWGVIYPCGSA-OHPAVYTIQCKYRSWIEKTIIRSN 281
gi|19526892|ref|NP_598473.1| VMLTSNVHPVQLPFRCSQPGQECQVSGWGTASRRRVKYNRSLSCSRVLLSQKCEFTYPGVITSNMICAEADGNQSDSGGGLVCDLTLHGVLSWGVIYPCGAA-OHPVYSYIICKYTPWIRRVIRFK 281
gi|51870931|ref|NP_001004100.1| VVLTSKVHPVQLPFCQAPRQECQVSGWGTANRRVKYNRSLSCSRVLLSQKCEFTYPGVITSNMICAGMDRQSDSGGGLVCDLTLHGILSWGVIYPCGAATONPAVYAKICNVTNWRVIRSK 281
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280

