


gi |14150201|ref|NP_115752.1|
gi |114666894|ref|XP_511540.2|
gi |73965584|ref|XP_848774.1|
gi |21313232|ref|NP_083925.1|
gi |27689721|ref|XP_220940.1|
gi |114051189|ref|NP_001039667.1|
gi |292619271|ref|XP_700929.4|
gi |118121219|ref|XP_001236361.1|

MASKIGSRRWMLQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAAIIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MASKIGSRRWMLQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAAIIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MASKIGSRRWMLQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAAIIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MASKIGSRRWMPQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAYIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MASKIGSRRWMLQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAAIIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MASKIGSRRWMLQLIMQLGSEVLLTRCPFWGCFSQLMLYAERAEARRKPDIPVPPYLYFDMGAAVLCASFMSTGKRRRWFALGAALQLAI STYAAIIGGVVHYGDWLKVR-MYSRTVAIIGGFVLVLSGAGELYRRKPRSRSLQSTGQVFLG 150
MAAAM--RKKALRFFSIFGAFILTRFGFWNCFAMLMFAERADVRRKPDIQVPPYLYIDLGAAVLCASFMSTGKRRRWFALAAAHALALSTYYSYVGGQVHYGDWLKVR-MYSRAMAIIIGGFLLLSGAGELYRRKPRSRSLQSTGQVFLG 150
-----CPSS-----GKPNHADILLVNLQVSEVELIN-----DRTETFPPLAHLNV----- 150

1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



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IYLIICVAYSLSQHSKEDRLAYLNHLPGGELMIQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLIICVAYSLSQHSKEDRLAYLNHLPGGELMIQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLIICVAYSLSQHSKEDRLAYLNHLPGGELMIQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLIICVAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLIICVAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
VVLVAVAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLNQVAVAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
IYLNQVAVAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258
-----KAYSLSQHSKEDRLAYLNHLPGGELMVQLFFVLYGVLALAFLSGYVVTLAAQILAVLLPPVMLLIDGNVAYWHTRRVEFWNQMKLLGESVGFIGAIVILATDG 258

.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....

