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gi |7657146|ref|NP_055180.1| MADGMPFSCCHYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTASWPDWALPRLSS---AWPGTLRSGMVRGPTATARFG-VPABG---RTPPFPGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |55639039|ref|XP_509417.1| MADGMPFSCCHYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTASWPDWALPRLSS---AWPGTLRSGMVRGPTATARFG-VPABG---RTPPFPGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |13507646|ref|NP_109629.1| MADGQLPFPSCYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTAPWPEWALPRLSS---AWPGTLRSGMVRGPTATARFG-VPABG---RSPFPFGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |16758408|ref|NP_446064.1| MADGQLPFPSCYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTAPWPEWALPRLSS---AWPGTLRSGMVRGPTATARFG-VPABG---RNPFPFGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |62460602|ref|NP_001014955.1| MADGMPFPCHYTSRR-RRDPFRDSSLSSRLDDGFGMDPPDDLTASWPDWALPRLSS---AWPGTLRSGMVRGPTATARFG-VPABG---RSPFPFGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |50978662|ref|NP_001003029.1| MADGMPFSCCHYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTSSWRNWALPRFSI---GWPGTLRSGMVRGPTAAARFG-VPABG---RSPFPFGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |50756697|ref|XP_415280.1| MADSLPFPSCCHYPSRL-RRDPFRDSSLSSRLDDGFGMDPPDDLTADWPDWARPRLTP---TWPGTLRSGMVRGPTAAARFG-VPABG---RSPFPFGEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
gi |155369333|ref|NP_001094427.1| MAEGDY-YTLGNRQRI-RRDPFREQLARRFMDDDFGLPPFPNELSMDWPGWARPRLSHRLDAPWTGSLRSGFPRASMSFQGFSEVYTESPRRATAPPTDSDEFVKVNVHSFKPEELMVKTKDGYVEVSGKHEEKQEGGIVSKNFT 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |7657146|ref|NP_055180.1| KKIQLPAEVDPVTVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDSQEVTCI----- 218
gi |55639039|ref|XP_509417.1| KKIQLPAEVDPVTVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDSQEVTCI----- 218
gi |13507646|ref|NP_109629.1| KKIQLPAEVDPAIVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDNQEVTCI----- 218
gi |16758408|ref|NP_446064.1| KKIQLPAEVDPVTVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDNQEVTCI----- 218
gi |62460602|ref|NP_001014955.1| KKIQLPAEVDPVTVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDSQEVTCI----- 218
gi |50978662|ref|NP_001003029.1| KKIQLPAEVDPVTVFASLSPGGLLIIEAPQVPPYSYTFG---ESSFNNELPQDSQEVTCI----- 218
gi |50756697|ref|XP_415280.1| KKIQLPYEVDPIIVFASLSPGGLLIIEAPQVPPYSYTFG---EGGCSGEIPLSPPEATCA----- 218
gi |155369333|ref|NP_001094427.1| KKIQLPLDVPDPVTVFASLSPGGLLIIEARQTPPYLYLSNEMPAESMEEPPEARQPEPSMSTTFEARE 218
.....160.....170.....180.....190.....200.....210.....

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