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gi |7661704|ref|NP_054776.1| MKTLOSTLLLLLVPLIKPAPPFOQDSRIIYDYGTDNFEE-SIFSQDYEDKYLDGKNIKEKET-VIIPNEKSLQLOKDEAITPLPPKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADIPNLRRL 150
gi |114625540|ref|XP_001145422.1| MKTLOSTLLLLLVPLIKPAPPFOQDSRIIYDYGTDNFEE-SIFSQDYEDKYLDGKNIKEKET-VIIPNEKSLQLOKDEAITPLPPKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADIPNLRRL 150
gi |27806829|ref|NP_776371.1| MKTLOSTLLLLLVPLIKPAPPFOQDSRIIYDYGTDNFEE-SIFSQDYEDKYLDGKNIKEKET-VIIPNEKSLQLOKDEAITPLPPKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADIPNLRRL 150
gi |6679166|ref|NP_032786.1| METVHSTFLLLLLVPLTQQAPQSDLSHVNSKYGTDNSEE-TKFSQDYEDKYLDGKSIKEKET-MIIPDEKSLQLOKDEVIPSLPTKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADMPNLRRL 150
gi |27683465|ref|XP_214441.1| MKTVHPTLLLLLVPLTQQTQQLDSHVNSKYGTDNSEE-TKFIQDYEDKYLDGKNIKEKET-MIIPDEKSLQLOKDEVIPSPVPTKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADMPNLRRL 150
gi |73947214|ref|XP_853340.1| MKTLOSTLFLLLFIPLTEPAPPFLQDSRMTYDYGTDNFEE-TLFSQDFEDKYLDGKNIKEKET--ILPEEKSLQLOKDERVMLPPLKKNDEMPTCLLCVCLSGSVYCEEVDIDAVPPLPKESAYLYARFNKIKKLTAKDFADIPNLRRL 150
gi |45383706|ref|NP_989540.1| MKTLQAFFLVAFVPLVKPAPPIQDSKPKFYEVVDADFATGSLIQDYE--MLPKDTIKDGTN---VSLDTALRLQADDSLSARPTKDTN-LPTCLLCVCLSGSVYCEEVDIDAVPPLPKETAYLYARFNKIKRIAVSDFADITTLRRL 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |7661704|ref|NP_054776.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLKL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNALES VPLNLPESLRVIHLQFN NIASITDDTFCKANDTSWIRDRIEIRLEGNP IVLGKHPNSFICLKR LPIGSYF 300
gi |114625540|ref|XP_001145422.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLKL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNALES VPLNLPESLRVIHLQFN NIASITDDTFCKANDTSWIRDRIEIRLEGNP IVLGKHPNSFICLKR LPIGSYF 300
gi |27806829|ref|NP_776371.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLKL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNALES VPLNLPESLRVIHLQFN NIASITDDTFCKANDTSWIRDRIEIRLEGNP IVLGKHPNSFICLKR LPIGSYF 300
gi |6679166|ref|NP_032786.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLRL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNDLES VPPNLPESLRVIHLQFN SSSITDDTFCKANDTRVIRERIEIRLEGNP IALGKHPNSFICLKR LPIGSYF 300
gi |27683465|ref|XP_214441.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLRL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNDLES VPPNLPESLRVIHLQFN SSSITDDTFCKANDTRVIRERIEIRLEGNP IALGKHPNSFICLKR LPIGSYF 300
gi |73947214|ref|XP_853340.1| DFTGNLIEDIEDGTFSKLSLLEELS LAENQLLKL PVLPPKLT LFN AKYNIKSRG IKA NAFK KLN NLF LYL D HNALES VPLNLPESLRVIHLQFN NITSDITDDTFCKANDTRVIRDRIEIRLEGNP IVLGKHPNSFICLKR LPIGSYF 300
gi |45383706|ref|NP_989540.1| DFTGNMIEIEDGAFSKLLLEELS LAENRLV KLPVLPPKLT LFN ANQNR IKS RGIK N NAFK KLT N LAYLYLGHNALES VPLNLPESLRILHLQHN NITINDTDFCKSNTRVIRTRMDEIRMEGNP ILLAKHVNAFSCLRTL PVGTY 300
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....290.....300

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