

gi|300116232|ref|NP\_001177828.1|MAWQGLVLAACLMLFPE-TADCLSRCSLCAVKTDGPKPINPLICSLCQAALLPSEEWERCQSFLFFFPSTLGLNDRKEDLGSKSVGEGPYSELAKLSGSFLKELEKSKFLPSISTKENTLSKSLKLRGLSDGFRGAESELMRDA 149  
gi|114680523|ref|XP\_001154951.1|MAWQGLVLAACLMLFPE-ATADCLSRCSLCAVKTDGPKPINPLICSLCQAALLPSEEWERCQSFLFFFPSTLGLNDRKEDLGSKSVGEGPYSELAKLSGSFLKELEKSKFLPSISTKENTLSKSLKLRGLSDGFRGAESELMRDA 149  
gi|73992010|ref|XP\_542932.2|MAWQGLVLAACLMLFPE-ATADCLSRCSLCAVKTDGPKPINPLICSLCQAALLPSEEWERCQSFLFFFPSTLGLNDRKEDLGSKSVGEGPYSELAKLSGSFLKELEKSKFLPSISTKENTLSKSLKLRGLSDGFRGAESELMRDA 149  
gi|9256622|ref|NP\_061351.1|MAWSRLMLAACLLVMPNVMADCLSLCSLCAVRIQDGPRPINPLICSLCQDLVPPSEEWETCRGFSFLTLTVSGLRGKDDLEDEVALEEGISAHAKLLEPVLEKEKSRLLTSVP-----EKFRLGSSSFNGKESSELGAD 140  
gi|9506967|ref|NP\_062247.1|MAWSRLMLAACLLVMPSEVAADCLSLCSLCAVRIQDGPHPINPLICSLCQDLVPPSEEWETCRGFSFLTLTVSGLRGKDDLENEVALEEGYATLTKLLEPLLKELEKGLLTSVS-----EKLRLGSSRFGNGRESSELLGTD 140



gi|300116232|ref|NP\_001177828.1|QLNDGAMETGILYLAEEDPKEQVKRYGGFLRKYPKRSSEVAGEG---DGDSMGHEDLYKRYGGFLRRIRPKLKWDNQKRYGGFLRRQFKVVTRSQEDPNAYSSELFD 254  
gi|114680523|ref|XP\_001154951.1|QLNDGAMETGILYLAEEDPKEQVKRYGGFLRKYPKRSSEVAGEG---DGDSMGHEDLYKRYGGFLRRIRPKLKWDNQKRYGGFLRRQFKVVTRSQEDPNAYSSELFD 254  
gi|73992010|ref|XP\_542932.2|QLNDDAMEAGALDENEEDPKEQVKRYGGFLRKYPKRSSEVAGEGNG--DGDAVGHEDLYKRYGGFLRRIRPKLKWDNQKRYGGFLRRQFKVVTRSQEDPNAYSSELLDG 256  
gi|9256622|ref|NP\_061351.1|RMNDEAAQGRIVHNEEDLRKQAKRYGGFLRKYPKRSSEMARDEDGGDGDQVGHEDLYKRYGGFLRRIRPKLKWDNQKRYGGFLRRQFKVVTRSQENPNAYS-EDLDV 248  
gi|9506967|ref|NP\_062247.1|LMNDEAAQAGILHFNEEDLRKQAKRYGGFLRKYPKRSSEMTGDEDRGQDGDQVGHEDLYKRYGGFLRRIRPKLKWDNQKRYGGFLRRQFKVVTRSQENPNAYS-EDLDV 248

