

gi|6754568|ref|NP\_034858.1| MRFAWAVLLLGPLQLCPILLRCAP-----GTPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDPSATARRPDGDAASQPRTPILLLRDNRRTASRRARTP-SPSGVAAGRPRPAARHWFQAGFSPSGARDGARRAARRTASF 150  
gi|8393733|ref|NP\_058757.1| MRFAWTVLFLGQLQFCPLLRCAP-----QAPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDSSATAPRADGNAASQPRTPILLLRDNRRTASARARTP-SPSGVAAGRPRPAARHWFQAGFSPSGAGDGAARRAARRTASF 150  
gi|20149540|ref|NP\_002308.2| MRFAWTVLFLGQLCALVHCAPPAAGQOOPPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDPSAAVPGAANASAAQPRTPILLLRDNRRTAARARTP-GSSGVTAGRPRPTARHWFQAGYSTRAREAGASRAENQIAPG 150  
gi|114601320|ref|XP\_001153015.1| MRFAWTVLFLGQLCALVHCAPPAAGQOOPPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDPSAAVPGAANASAAQPRTPILLLRDNRRTAARARTP-GSSGVTAGRPRPTARHWFQAGYSTRAREAGASRAENQIAPG 150  
gi|73970573|ref|XP\_538599.2| MRFAWTVLFLGQLCALVHCAPPAAGQOOPPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDPSAAVPGAANASAAQPRTPILLLRDNRRTAARARTP-GAGG-AGRPRPTARHWFQAGYSTRAREAGASRAENQIAPG 150  
gi|148229898|ref|NP\_776357.2| MRFAWTVLFLGQLCALVHCAPPAAGQOOPPREPPAAPGAWRQTIQWENNGVVFLLSLGAYQPORRRDPSAAVPGAANASAAQPRTPILLLRDNRRTAARARTP-GPSAAAAGRPRPAARHWFQAGYSTRAREAGASRAENQIAPG 150  
gi|46309545|ref|NP\_996972.1| MSMSLIDTFIYFAFVCLLSCIAQTG---GTQRQNTGAAALRQTIQWENNGVVFLLSLGAYQPORRRDPSAAVPGAANASAAQPRTPILLLRDNRRTAARARTP-GPSAAAAGRPRPAARHWFQAGYSTRAREAGASRAENQIAPG 150



gi|6754568|ref|NP\_034858.1| QPPQLSNLRPPSH--IDRMVGGDDPYNPKYSD-DNPNYNYDTYERPRPGSRNRPYGYGTGYFOYGLPDLVDPYIQAATYVQKMSMYNLRCAAEENCLASSAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|8393733|ref|NP\_058757.1| QPPQLSNLRPPSH--VDRMVGGDDPYNPKYSD-DNPNYNYDTYERPRSGSRNRPYGYGTGYFOYGLPDLVDPYIQAATYVQKMSMYNLRCAAEENCLASSAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|20149540|ref|NP\_002308.2| EVPALSNLRPPSR--VDGMVGGDDPYNPKYSD-DNPNYNYDTYERPRPGGRYRPGYGTGYFOYGLPDLVADPYIQAATYVQKMSMYNLRCAAEENCLASTAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|114601320|ref|XP\_001153015.1| DVPALSNLRPPSR--VDGMVGGDDPYNPKYSD-DNPNYNYDTYERPRPGGRYRPGYGTGYFOYGLPDLVADPYIQAATYVQKMSMYNLRCAAEENCLASTAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|73970573|ref|XP\_538599.2| EVPALSNLRPPSR--VDGMVGGDDPYNPKYSD-DNPNYNYDTYERPRPGSRYRPGYGTGYFOYGLPDLVDPYIQAATYVQKMSMYNLRCAAEENCLASSAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|148229898|ref|NP\_776357.2| EVPALSNLRPPSR--VDGMVGGDDPYNPKYSD-DNPNYNYDTYERPRPGSRYRPGYGTGYFOYGLPDLVDPYIQAATYVQKMSMYNLRCAAEENCLASSAYRADVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300  
gi|46309545|ref|NP\_996972.1| RSLNGLDRPAGDD--EVMVGGDDPYNPKYSD-DNPNYNYDTYERPRPQR--PGYGTGYFOYGLPDLVDPYIQAATYVQKMSMYNLRCAAEENCLASSAYRSDVRDYDHRVLLRFPORVKNQGTSDFLPSRPRYSWEWHSCHQHYH 300



gi|6754568|ref|NP\_034858.1| SMDEFSHYDLLDANTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCDIRYTGHHAYASGCTISPY 421  
gi|8393733|ref|NP\_058757.1| SMDEFSHYDLLDASTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCEIRYTGHHAYASGCTISPY 421  
gi|20149540|ref|NP\_002308.2| SMDEFSHYDLLDANTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCDIRYTGHHAYASGCTISPY 421  
gi|114601320|ref|XP\_001153015.1| SMDEFSHYDLLDANTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCDIRYTGHHAYASGCTISPY 421  
gi|73970573|ref|XP\_538599.2| SMDEFSHYDLLDASTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCEIRYTGHHAYASGCTISPY 421  
gi|148229898|ref|NP\_776357.2| SMDEFSHYDLLDASTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCEIRYTGHHAYASGCTISPY 421  
gi|46309545|ref|NP\_996972.1| SMDEFSHYDLLDASTQRRVAEGHKASFLEDTSCDYG YHRRFACTAHTQGLSPGCYDTYAADIDCQWIDIIDVQPGNYILKVS VNP SYLVPESDYTNVVRCDIRYTGHHAYASGCTISPY 421

