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                                 *     .     :  * :*     *   ::.**::********:*****:************.**:*  *******  **:**:** *.** *** ** ** **.*:**:* *.. *** ***** ** *** ***********::**
gi|13128972|ref|NP_076932.1|     -MRGNLALVGVLISLAFLSLLPSGHPQPAGDDACSVQILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDMGDKGQKGSVGRHGKIGPIGSKGEKGDSGDIGPPGPNGEPGLPCECSQLRKAIGEMDNQVSQLTSELKFIKNAVAGVRET   150
gi|114575925|ref|XP_001150689.1| -MRGNLALAGVLISLAFLSLLPSGHPQPAGDDACSVQILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDMGDKGQKGSVGRHGKIGPIGSKGEKGDSGDIGPPGPNGEPGLPCECSQLRKAIGEMDNQVSQLTSELKFIKNAVAGVRET   150
gi|103472134|ref|NP_082142.1|    MMMRDLALAGMLISLAFLSLLPSGCPQQTTEDACSVQILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDMGDKGQKGTVGRHGKIGPIGAKGEKGDSGDIGPPGPSGEPGIPCECSQLRKAIGEMDNQVTQLTTELKFIKNAVAGVRET   150
gi|73979698|ref|XP_848349.1|     -MKRDLALIG----LAFLWLLPLGCPQQTAEDACSLHILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDMGDKGQKGSVGRHGKIGPIGSKGEKGDSGDIGPPGPNGEPGIP-ESCQLRKGIGEMDNQVTQLTTELKFIKNAVAGVRET   150
gi|115495093|ref|NP_001069771.1| -MKRALALMG----LAFLCVLRAGAAQQTVDDACSVQILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDVGDKGQKGGVGRHGKIGPIGSKGEKGDSGDIGPPGPNGEPGIPCECSQLRKAIGEMDNQVTQLTAELKFIKNAVAGVRET   150
gi|113206116|ref|NP_001038130.1| -MKRDLVFLVTLISLAFLSLLRSGYPQHIAEESCSVQILVPGLKGEAGEKGEKGAPGRPGRVGPPGEKGEMGDKGQKGSMGRHGKIGPIGSKGEKGDNGDMGPPGPNGDPGIPCECSQLRKAIGEMDIQVAQLTTELKFIKNAVAGVRET   150
gi|55925245|ref|NP_001007332.1|  -MVGEKLVAYMLVSVLGLALLRSVFGQHMPEEPCSVQILVPGLKGEAGEKGEKGAPGRPGRVGPTGEQGPPGDKGQKGSPGRYGKMGPTGPKGLKGDMGDPGPKGPNGEPGVPCECAPLRKMIGEMDIQVVQLTNELKFIKNAVAGIKET   150
                                 1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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                                 :.*:********** :*:: *:.*** *.****  **  :*.*: :***:**:******:.** *** ::*** **.:**.********:*****:*:** * ****: **:*:*****: :
gi|13128972|ref|NP_076932.1|     ESKIYLLVKEEKRYADAQLSCQGRGGTLSMPKDEAANGLMAAYLAQAGLARVFIGINDLEKEGAFVYSDHSPMRTFNKWRSGEPNNAYDEEDCVEMVASGGWNDVACHTTMYFMCEFDKENM   272
gi|114575925|ref|XP_001150689.1| ESKIYLLVKEEKRYTDAQLSCQGRGGTLSMPKDEAANGLMAAYLAQAGLARVFIGINDLEKEGAFVYSDRSPMRTFNKWRSGEPNNAYDEEDCVEMVASGGWNDVACHTTMYFMCEFDKENM   272
gi|103472134|ref|NP_082142.1|    ESKIYLLVKEEKRYADAQLSCQARGGTLSMPKDEAANGLMASYLAQAGLARVFIGINDLEKEGAFVYSDRSPMQTFNKWRSGEPNNAYDEEDCVEMVASGGWNDVACHITMYFMCEFDKENL   272
gi|73979698|ref|XP_848349.1|     DSKIYLLVKEEKRYVDAQLSCQGRGGTLGMPKDEAANGLMAAYIAQAGLARVFIGINDLDREGTFVYSDRSPMQTFNKWRSGEPNNAYDEEDCVEMVASGGWNDVACHVTMHFLCEFDKEHV   272
gi|115495093|ref|NP_001069771.1| EQKMYLLVKEEKRYLDAQLACQGRGGTLSMPKDEAANALLAAYITQAGLARVFIGINDLEREGAFVYADRSPMQTFSKWRSGEPNNAYDEEDCVELVASGGWNDVACHLTMHFLCEFDKEHV   272
gi|113206116|ref|NP_001038130.1| DNKIYLLVKEEKRYKEAQLYCHGRGGTLSMPKDENANNLIASYINQAGLTRVFIGINDLEKEGNFVYSDRSPMQTFNKWRSGEPNNAYDEEDCVEMVASGGWNDVACHITMYFVCEFDKENV   272
gi|55925245|ref|NP_001007332.1|  DSKVYLLVKEEKRYREAEVFCQGRGGHLAMPKDAAANRAIAGYVTDAGLSRVYIGINDLEREGHFVYVERSPMTTFSRWREGEPNNAYDDEDCVEMVSSGEWIDVACQLTMYFVCEFDKDTV   272
                                 .......160.......170.......180.......190.......200.......210.......220.......230.......240.......250.......260.......270..
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