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                                 *       .    *     .        .  *  :    .  * :**  :  . : ::: .*     :*. .: .:::.  **: :.*:::**   ::* * : .       :..:*: :  *   : :*:*::.*:: .  ::.:*  :
gi|10835141|ref|NP_000563.1|     MHSSALLCC-LVLLTGVRASPGQGTQSENSCTHFPGNLPNMLRDLRDAFSRVKTFFQMKD-QLDNLLLKESLLEDFKGYLGCQALSEMIQFYLEEVMPQAENQD--PDIKAHVNSLGENLKTLRLRLRRCHRFLPCENKSKAVEQVKNAF   150
gi|55589346|ref|XP_525040.1|     MHSSALLCC-LVLLTGVRASPGQGTQSENSCTHFPGNLPNMLRDLRDAFSRVKTFFQMKD-QLDNLLLKESLLEDFKGYLGCQALSEMIQFYLEEVMPQAENQD--PDIKVHVNSLGENLKTLRLRLRRCHRFLPCENKSKAVEQVKNAF   150
gi|73960302|ref|XP_855560.1|     MPSPALLCCCLVLLAGVGASRHQSTLPEDDCTHFPASLPHMLRELRAAFGRVKTFFQMKD-KLDNILLTGSLLEDFKSYLGCQALSEMIQFYLEEVMPRAENHD--PDIKNHVNSLGEKLKTLRLRLRRCHRFLPCENKSKAVEQVKSAF   150
gi|41386772|ref|NP_776513.1|     MHSSALLCC-LVFLAGVAASRDASTLSDSSCIHLPTSLPHMLRELRAAFGEAKTFFQMKD-QLHSLLLTQSLLDDFKGYLGCQALSEMIQFYLEEVMPQAENHG--PDIKEHVNSLGEKLKTLRLRLRRCHRFLPCENKSKAVEKVKRVF   150
gi|6754318|ref|NP_034678.1|      MPGSALLCC-LLLLTGMRISRGQYSREDNNCTHFPVGQSHMLLELRTAFSQVKTFFQTKD-QLDNILLTDSLMQDFKGYLGCQALSEMIQFYLVEVMPQAEKHG--PEIKEHLNSLGEKLKTLRMRLRRCHRFLPCENKSKAVEQVKSDF   150
gi|7549767|ref|NP_036986.1|      MLGSALLCC-LLLLAGVKTSKGHSIRGDNNCTHFPVSQTHMLRELRAAFSQVKTFFQKKD-QLDNIVLTDSLLQDFKGYLGCQALSEMIKFYLVEVMPQAENHG--PEIKEHLNSLGEKLKTLWIQLRRCHRFLPCENKSKAVEQVKNDF   150
gi|52138717|ref|NP_001004414.1|  MQT----CC-QALLLLLAACTLPAHCLEPTCLHFSELLPARLRELRVKFEEIKDYFQSRDDELNIQLLSSELLDEFKGTFGCQSVSEMLRFYTDEVLPRAMQTS--TSHQQSMGDLGNMLLGLKATIRRCHRFFTCEKRSKAIKQIKETF   150
gi|67459916|ref|NP_001018621.2|  MIFSGVILSALLTLLLCDCAQSRRVECKTDCCSFVEGFPLRLRELRSAYKEIQKFYESND--DLEPLLNEDIKHNINSPYGCHVMNEILHFYLETILPTALQKNPLKHSTTPIDSIGNIFQELKRDMVKCKRYFSCQN-PFEVNSLKNSY   150
                                 1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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gi|10835141|ref|NP_000563.1|     NKLQEKGIYKAMSEFDIFINYIEAYMTMKIRN-   183
gi|55589346|ref|XP_525040.1|     NKLQEKGIYKAMSEFDIFINYIEAYMTMKIRN-   183
gi|73960302|ref|XP_855560.1|     SKLQEKGVYKAMSEFDIFINYIETYMTMRMKI-   183
gi|41386772|ref|NP_776513.1|     SELQERGVYKAMSEFDIFINYIETYMTTKMQK-   183
gi|6754318|ref|NP_034678.1|      NKLQDQGVYKAMNEFDIFINCIEAYMMIKMKS-   183
gi|7549767|ref|NP_036986.1|      NKLQDKGVYKAMNEFDIFINCIEAYVTLKMKN-   183
gi|52138717|ref|NP_001004414.1|  EKMDENGIYKAMGEFDIFINYIEEYLLMRRRK-   183
gi|67459916|ref|NP_001018621.2|  EKMKEKGVYKAMGELDLLFRYIEQYLASKRVKH   183
                                 .......160.......170.......180...
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