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                                  : :: .  :  . **  ..:*  :.* *:  *:  ** :..  **.*  *: :* **: **  :. .: *:***  **.**: ** :** .*. *** ::***. ** *** :** ****:*.****::
gi|5032019|ref|NP_005846.1|       MARALCRLPRRGLWLLLAHHLFMTTACQEANYGALLRELCLTQFQVDMEAVGETLWCDWGRTIRSYRELADCTWHMAEKLGCFWPNAEVDRFFLAVHGRYFRSCPISGRAVRDPPGSILYPFIVVPITVTLLVTALVVWQS---------   150
gi|114584083|ref|XP_516183.2|     MARALCRLPRRGLWLLLAHHLFMATACQEANYGALLQELCLTQFQVDMEAVGETLWCDWGRTIRSYRELADCTWHMAEKLGCFWPNAEVDRFFLAVHGRYFRSCPISGRAVRDPPGSILYPFIMVPITVTLLVTALVVWQS---------   150
gi|139948257|ref|NP_001077246.1|  MAPGLRGLRCLGLWLLLVTHLCLATACQHADHGTLLQEFCLPQFQAHMEAVGRTLWCDWGKTIRSYRELSDCTRHVVQKLDCFWPNAEVDKFFLVVHQRYFRNCPVSGRALQDPPSSILCPFIVVPILVTLLVTVLVVWRASAQRASCRP   150
gi|8394145|ref|NP_058590.1|       MAPGLRGLPRCGLWLLLAHHLFMVTACRDPDYGTLIQELCLSRFKENMETIGKTLWCDWGKTIQSYGELTYCTKHVAHTIGCFWPNPEVDRFFIAVHHRYFSKCPISGRALRDPPNSILCPFIALPITVTLLMTALVVWRS---------   150
gi|13928890|ref|NP_113833.1|      MAPGLRGLPRRGLWLLLAHHLFMVTACRDPDYGTLIQELCLSRFKEDMETIGKTLWCDWGKTIGSYGELTHCTKLVANKIGCFWPNPEVDKFFIAVHHRYFSKCPVSGRALRDPPNSILCPFIVLPITVTLLMTALVVWRS---------   150
gi|118093300|ref|XP_421873.2|     ----MALLPRRFLCFVLAHHFIAATACHEAEYGRQIRERCLRPFKLSMEGIGQRLWCDWDETMGTYGELTNCTVAVAENLTCYWPNRLVDEFFVAVHSHYFRNCSPSGRALRDPPNSILCPFILVPILVTLLMTALVVWRS---------   150
gi|50540458|ref|NP_001002693.1|   MAS--VSWRHALFVLVAVNLLVLARACS-SHYGSAIEEFCMAKFKLDMEVLDQRQWCSWEDTVESYGELTNCTFLVALKMNCFWPNRMVDEFFIRVHRHYFHDCSLSGRLLHDPPNRILGPFIVVPILVTLLMTALVVWRS---------   150
gi|162139008|ref|NP_001104721.1|  MAS--VSWRHALFVLVAVNLLVLARACS-SHYGSAIEEFCLAKFKLDMEVLDQRQWCSWEDTVESYGELTNCTFLVALKMNCFWPNRMVDEFFIRVHRHYFHDCSLSGRLLHDPPNRILGPFIVVPILVTLLMTALVVWRS---------   150
                                  1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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                                            *:: :
gi|5032019|ref|NP_005846.1|       ----------KRTEGIV-------------------------------------------------------------------------------------------   267
gi|114584083|ref|XP_516183.2|     ----------KRTEGIV-------------------------------------------------------------------------------------------   267
gi|139948257|ref|NP_001077246.1|  RSVCGGRGRAARSQKLQSRWEVGTEARPDQGFWSPNGRADPRAPPPPRKSPLLLRRAHPGMPARVVKHTRWPGKAAQGLGGSGHSCWSCWGWSELSPNNPHHSRVLPVARLWRSICD   267
gi|8394145|ref|NP_058590.1|       ----------KRTEGIV-------------------------------------------------------------------------------------------   267
gi|13928890|ref|NP_113833.1|      ----------KRTEGIV-------------------------------------------------------------------------------------------   267
gi|118093300|ref|XP_421873.2|     ----------KRSEGIV-------------------------------------------------------------------------------------------   267
gi|50540458|ref|NP_001002693.1|   ----------KRSEGIV-------------------------------------------------------------------------------------------   267
gi|162139008|ref|NP_001104721.1|  ----------KRSEGIV-------------------------------------------------------------------------------------------   267
                                  .......160.......170.......180.......190.......200.......210.......220.......230.......240.......250.......260.......
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