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logo
NTDB id 1188 NELON RS01755 WP 003772458.1     .......MKAMQKGFTLIELMIVIAIIGILAAIALPAYQDYTVRSKVSEALIAASSPKSALSEAFQTDGITGMTAAAKAF    73
NTDB id 269187 C3F34 RS08220 WP 104441389.1  ........MNAQKGFTLIELMIVVAIIGILAAIAIPAYQDYIVRSKVTEGLNLASSYKTIIAENAGNGASNLTL......    66
NTDB id 1424 RS RS02795 WP 011000517.1        MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV......    74
NTDB id 1002 ACIAD RS15100 WP 004923779.1    ........MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTAG    72
NTDB id 1274 LPP RS09530 WP 011214168.1      .........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY    71
NTDB id 1258 GC085 RS09745 WP 014844212.1    .........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY    71
NTDB id 1038 HON27 RS01510 WP 000993725.1    ........MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLSD....    68
NTDB id 1192 PAKAF RS23870 WP 024947914.1    ........MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDA    72
NTDB id 1197 PSJM300 03940 AFN76866.1        ......MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTD...    71
consensus                                              !!!!!!!!!!!!*!!*!!!!!!!!*!*!**!***********  ***  !*****     *
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logo
NTDB id 1188 NELON RS01755 WP 003772458.1     NKTAAAGGGAGGAAAAGTQHASKYVKSV..KVTEASPWDITVAISATEANGIPTGLDGNTIRLFPLVNKA....V.PAAT   146
NTDB id 269187 C3F34 RS08220 WP 104441389.1  ...........GIPNF...AATDNVASITP...NATTGAIVIVYT.DK.......AKSVELTLTPFDGKEGGTAIKNKTV   121
NTDB id 1424 RS RS02795 WP 011000517.1        ...........GSSVF...TPTKNVANLTIAGTGTVPGQITVTYT.TA.......AGGGTLALVPT...AAGTALPVSSA   129
NTDB id 1002 ACIAD RS15100 WP 004923779.1    .......SSCVGVQEISASNATTNVATA..TCGASSAGQIIVTMDTTK.......AKGANITLTPT.............Y   123
NTDB id 1274 LPP RS09530 WP 011214168.1      .......V.........SPAATPNVASI..AIGA..NGVITITY.TAA.......AGGGTIVMTPT...L.........Q   111
NTDB id 1258 GC085 RS09745 WP 014844212.1    .......V.........SPAATPNVQSI..AIGA..NGVITITY.TAA.......AGGGTIIMTPT...L.........Q   111
NTDB id 1038 HON27 RS01510 WP 000993725.1    ..........VGFTAATSSV..CATYSI..DAFTETDGAGKI.TCTLK...GNPKIATKIVELT...............R   115
NTDB id 1192 PAKAF RS23870 WP 024947914.1    ....TKKEVPLGVAADANKL..GTIA.L..KP.DPADGTADITLTFTM.GGAGPKNKGKIITLT...............R   126
NTDB id 1197 PSJM300 03940 AFN76866.1        .......EGYIGITDS.TSY..CDVD.L..D..TAADGHI...ECTAK.GGNAGKFDGKTITLN...............R   117
consensus                                               *     *  ** *****          * *  **   *          * *  ** ***
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logo
NTDB id 1188 NELON RS01755 WP 003772458.1     STGAIDWACASAAKETADKRFGTGSGVAVGTLPAKYAPSECK............   188
NTDB id 269187 C3F34 RS08220 WP 104441389.1  PTNQITWVCKVKQPA................DNAKYVPANCRG...........   148
NTDB id 1424 RS RS02795 WP 011000517.1        PSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAECR............   168
NTDB id 1002 ACIAD RS15100 WP 004923779.1    ASGAVTWKCTTT...................S.DKKYVPSECRG...........   147
NTDB id 1274 LPP RS09530 WP 011214168.1      ANGDVTWTCTGG...................TLLAKYRPASCRP...........   136
NTDB id 1258 GC085 RS09745 WP 014844212.1    ANGDVTWTCTGG...................TLLAKYRPASCRP...........   136
NTDB id 1038 HON27 RS01510 WP 000993725.1    .STDGAWTCVSD...................VPTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1    TAADGLWKCTSD...................QDEQFIPKGCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1        .TADGEWSCAST...................LDAKYKPGKCS............   139
consensus                                     **** !*! *                    ** *** !* !*
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X non conserved

X similar

X ≥ 50% conserved