```
logo
NTDB id 1038 HON27 RS01510 WP 000993725.1     ........MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVN.EGAVPTK......   65
NTDB id 1192 PAKAF RS23870 WP 024947914.1     ........MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALS.RGWSVKSGTGTED   71
NTDB id 1191 AAC28468.1 484..945( )           ......MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAG   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 276021 C6N19 RS02755 WP 106405811.1   ........MNAQKGFTLIELMIVVAIIGILAAIAIPAYQDYMVRSKVTEGLNLASSYKTTIAENAGNGAANLAV......   66
NTDB id 1424 RS RS02795 WP 011000517.1        MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV......   74
consensus                                               !!!!!!!!!!!!!*!!*!!!!!!!!*!*!**!***********  *  *  **  *  *      **

logo
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....LSDV.GFTAATSSVCATYSIDAFTETDGAGKITCTL...KGNPKIATKIVELT................RSTDGA   120
NTDB id 1192 PAKAF RS23870 WP 024947914.1     ATK.KEVPL.GVAADANKL.GTIALK.PDPADGTADITLTFTMGGAGPKNKGKIITLT................RTAADGL   132
NTDB id 1191 AAC28468.1 484..945( )           NV........IFDSANGLDGKYFAPGGVTVTPDTGAINIAFDAGAN....AGKTMTITPTVIAA..........SGQIST   132
NTDB id 1002 ACIAD RS15100 WP 004923779.1     SSCVGVQEISASNATTNV..ATATCGA....SSAGQIIVTMDTTKA....KGANITLTPT.............YASGAVT   129
NTDB id 1274 LPP RS09530 WP 011214168.1       V.........SPAATPNV..ASIAIGA......NGVITITY.TAAA....GGGTIVMTPTL............QANGDVT   117
NTDB id 1258 GC085 RS09745 WP 014844212.1     V.........SPAATPNV..QSIAIGA......NGVITITY.TAAA....GGGTIIMTPTL............QANGDVT   117
NTDB id 276021 C6N19 RS02755 WP 106405811.1   ....GVP...NFAATENV..ASITPDG..T...TGKIEILY.TDKA....KGLTLTLTPYDGAEGGIAIAAGTVPTNQIT   127
NTDB id 1424 RS RS02795 WP 011000517.1        ....GSS...VFTPTKNV..ANLTIAG..TGTVPGQITVTY.TTAA....GGGTLALVPTA...AGTALPVSSAPSGPIM   135
consensus                                                ** **  ***  ***       * !**** * **    **  **  ****                * **

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logo
NTDB id 1038 HON27 RS01510 WP 000993725.1     WTCVSDV................PTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1     WKCTSDQ................DEQFIPKGCSK...........   150
NTDB id 1191 AAC28468.1 484..945( )           WTCAPGAANG............IETRRLPTSCQ*...........   153
NTDB id 1002 ACIAD RS15100 WP 004923779.1     WKCTTTS................DKKYVPSECRG...........   147
NTDB id 1274 LPP RS09530 WP 011214168.1       WTCTGGT................LLAKYRPASCRP...........   136
NTDB id 1258 GC085 RS09745 WP 014844212.1     WTCTGGT................LLAKYRPASCRP...........   136
NTDB id 276021 C6N19 RS02755 WP 106405811.1   WVCTVDSAA..............NAKYVPSNCRGS..........   148
NTDB id 1424 RS RS02795 WP 011000517.1        WTCYAQNKAQAASSVAPSGTMSLAAKYAPAECR............   168
consensus                                     !*!*                     ***!   !*
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X non conserved

X similar

X ≥ 50% conserved