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logo

NTDB id 1424 RS RS02795 WP 011000517.1          MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV......   74
NTDB id 1002 ACIAD RS15100 WP 004923779.1       ........MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPST..   70
NTDB id 1184 GTF74 RS02335 WP 000649326.1       ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1167 A1552VC RS11065 WP 000649326.1     ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1160 ABDM36 RS02330 WP 000649326.1      ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1097447 ACM000 RS01580 WP 062850214.1   ........MNAQKGFTLIELMIVVAIIGILAAIAIPQYSKYQAKSKLTASLSEISAGKTVAEERINNGEA..VT......   64
NTDB id 1038 H0N27 RS01510 WP 000993725.1       ........MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAV.PTK......   65
NTDB id 1292 VP RS12235 WP 005479693.1          ...MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.A........   68
NTDB id 1197 PSJM300 03940 AFN76866.1           ......MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKT.PSLTSTD..   71
NTDB id 1192 PAKAF RS23870 WP 024947914.1       ........MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWS.VKSGTGTED   71
consensus                                          **  * * !*!!!!!!!!!!*!**!*!!!*!*!*!**!* ** ** *** ****** *** *  *** * **  *

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NTDB id 1424 RS RS02795 WP 011000517.1          ...GSSV.....FTPTKN.VANLTIAGT.GTVPGQITVTYTT...AAGGGTLALVPTAAGTALPVSSAPSGPIMWTCYAQ  141
NTDB id 1002 ACIAD RS15100 WP 004923779.1       ..AGSSCVGVQEISASNA.TTNVATATCGASSAGQIIVTMDT..TKAKGANITLT..........PTYASGAVTWKCTTT  135
NTDB id 1184 GTF74 RS02335 WP 000649326.1       ...GFTRLGTVEDMGD..GKIVIAP.TASGALGGTIKYTFD..AGVVSSSKIQLAR.............DANGLWTCSTT  136
NTDB id 1167 A1552VC RS11065 WP 000649326.1     ...GFTRLGTVEDMGD..GKIVIAP.TASGALGGTIKYTFD..AGVVSSSKIQLAR.............DANGLWTCSTT  136
NTDB id 1160 ABDM36 RS02330 WP 000649326.1      ...GFTRLGTVEDMGD..GKIVIAP.TASGALGGTIKYTFD..AGVVSSSKIQLAR.............DANGLWTCSTT  136
NTDB id 1097447 ACM000 RS01580 WP 062850214.1   ....LTDLGLQATTSN..CDIAVT.GFAAGSGAGTIACTVKNAPSQVTGAVMTWTR............VGATGIWTCATT  125
NTDB id 1038 H0N27 RS01510 WP 000993725.1       ....LSDVGFTAATSSVCATYSID.AFTETDGAGKITCTLKGNPKIATKIVE..LT............RSTDGAWTCVSD  126
NTDB id 1292 VP RS12235 WP 005479693.1          AAATLAQLGVDVAANDLG.AIDSA..LAGGSPTLTFTFD...ADSSIDGGILTFTR............DA.ATGWSCLPT  129
NTDB id 1197 PSJM300 03940 AFN76866.1           ....EGYIGITDS.TSYC.DVDLD.TAADGHI...ECTAKGGNAGKFDGKTITLNR............TADGEWSCAST  128
NTDB id 1192 PAKAF RS23870 WP 024947914.1       ATKKEVPLGVAADANKLG.TIALKPDPADGTADITLTFTMGGAGPKNKGKIITLTR............TAADGLWKCTSD  138
consensus                                          * * *** *         * **    * ******* **     *  ***  *** *             * **!*!***

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NTDB id 1424 RS RS02795 WP 011000517.1          NKAQAASSVAPSGTMSLAAKYAPAECR............   168
NTDB id 1002 ACIAD RS15100 WP 004923779.1       S................DKKYVPSECRG...........   147
NTDB id 1184 GTF74 RS02335 WP 000649326.1       VT...............SEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1     VT...............SEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1      VT...............SEIAPKGCTAGATIN......   153
NTDB id 1097447 ACM000 RS01580 WP 062850214.1   GITDT...........NNKALAPKGCAAS.........   143
NTDB id 1038 H0N27 RS01510 WP 000993725.1       VP...............TEFLPKGCTAGTPTEGAITSL   149
NTDB id 1292 VP RS12235 WP 005479693.1          AP...............AI.AVDGCS............   139
NTDB id 1197 PSJM300 03940 AFN76866.1           LD...............AKYKPGKCS...........   139
NTDB id 1192 PAKAF RS23870 WP 024947914.1       QD...............EQFIPKGCSK..........   150
consensus                                       *                    *****!**
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X non conserved

X similar

X ≥50% conserved