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logo

NTDB id 1002 ACIAD RS15100 WP 004923779.1     .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTAGSS.    74
NTDB id 1274 LPP RS09530 WP 011214168.1       ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGYV..    72
NTDB id 1258 GC085 RS09745 WP 014844212.1     ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGYV..    72
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPAYQNYIAKSQASEAFTLADGLKTTINTNLQAGTCFAGGATAVTAADK    75
NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT    80
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT    80
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT    80
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENG.IAAAA........    71
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGA.VPTKLSDV.....    69
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGW.SVKSGTGTEDATK    74
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGK.TPSLTSTDE....    72
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consensus                                              **!!*!!!!!!!*!*!!*!!!*!*!*!**!* ** *  *** ** *** *** *       **   **

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NTDB id 1002 ACIAD RS15100 WP 004923779.1     ...CVGVQEISASNATTNVATA.TCGASSAGQIIVTMDT....TKAKGANITLTPT.YASGAVTWKCTT.TS.DKKYVPS   143
NTDB id 1274 LPP RS09530 WP 011214168.1       ...........SPAATPNVASI.AIGA..NGVITITY.T....AAAGGGTIVMTPTLQANGDVTWTCTG.GTLLAKYRPA   132
NTDB id 1258 GC085 RS09745 WP 014844212.1     ...........SPAATPNVQSI.AIGA..NGVITITY.T....AAAGGGTIIMTPTLQANGDVTWTCTG.GTLLAKYRPA   132
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     V........SGKYGDAEIG.....GTAPNCTITYTFKSSGVSNKLTSTKIVMNVS....ETGILTKNSGTDTPVELLPQ   137
NTDB id 1184 GTF74 RS02335 WP 000649326.1     R...LGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDA....GVVSSSKIQLARD....ANGLWTCS..TTVTSEIAPK   144
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    R...LGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDA....GVVSSSKIQLARD....ANGLWTCS..TTVTSEIAPK   144
NTDB id 1167 A1552VC RS11065 WP 000649326.1   R...LGTVEDMG.DGKIVIAPTASGALGG..TIKYTFDA....GVVSSSKIQLARD....ANGLWTCS..TTVTSEIAPK   144
NTDB id 1292 VP RS12235 WP 005479693.1        TLAQLGVDVAANDLGAIDSA.....LAGGSPTLTFTF...DADSSIDGGILTFTRD....AATGWSCL..PTAPAIA.VD   136
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....GFTAATSSVCATYSIDAF.TETDGAGKIT.CTLKG..NPKIATKIVELTRS....TDGAWTCV..SDVPTEFLPK   134
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KEVPLGVAADANKLGTIALKP...DPADGTADITLTFTMGGAGPKNKGKIITLTRT...AADGLWKCT..SDQDEQFIPK   146
NTDB id 1197 PSJM300 03940 AFN76866.1         ..GYIGITDSTS.YCDVDLD....TAADGH..IE.CTAKGGNAGKFDGKTITLNRT....ADGEWSCA..STLDAKYKPG   136
NTDB id 1204 PSJM300 03935 AFN76865.1         ..GFIGIGASTN.YCTVTLI....ATATGS..IT.CTTQNGMPDRFNGKTITIART....TDGLWDCT..SDLDSRYKPG   138
consensus                                        **         * * **      ** *   ** **          *  * * **    * * ****  ***   * *

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NTDB id 1002 ACIAD RS15100 WP 004923779.1     ECRG...........   147
NTDB id 1274 LPP RS09530 WP 011214168.1       SCRP...........   136
NTDB id 1258 GC085 RS09745 WP 014844212.1     SCRP...........   136
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     SFVAS..........   142
NTDB id 1184 GTF74 RS02335 WP 000649326.1     GCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    GCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   GCTAGATIN......   153
NTDB id 1292 VP RS12235 WP 005479693.1        GCS............   139
NTDB id 1038 HON27 RS01510 WP 000993725.1     GCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1     GCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1         KCS............   139
NTDB id 1204 PSJM300 03935 AFN76865.1         KCL............   141
consensus                                      **
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X non conserved

X similar

X ≥ 50% conserved