```
logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1        MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 1167 A1552VC RS11065 WP 000649326.1      MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 1160 ABDM36 RS02330 WP 000649326.1       MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 1292 VP RS12235 WP 005479693.1           MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.AAAAIL......   73
NTDB id 1038 HON27 RS01510 WP 000993725.1        .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAV.PTKLSD......   68
NTDB id 1092564 ACLSSQ RS14695 WP 014236553.1   ....MNKVQQGFTLIELMVVVAIIGILAAVALPAYSDYQAKGKIAAGLAEISAVRTAYEMIANDGGT.PTE.........   66
NTDB id 1192 PAKAF RS23870 WP 024947914.1        .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWS.VKSGTGTEDATK   74
NTDB id 1197 PSJM300 03940 AFN76866.1            ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKT.PSLTSTDE....   72
NTDB id 1204 PSJM300 03935 AFN76865.1            .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHT.PSLVSSLP....   74
consensus                                            *** **!!*!!!!!*!!!*!!*!!!*!*!*!**!* ** ** *** ******  !  * **** *   *
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logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1        ........AGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD..AGVVSSSKIQLARD.ANGLWTCST.TVTSEIAPK   144
NTDB id 1167 A1552VC RS11065 WP 000649326.1      ........AGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD..AGVVSSSKIQLARD.ANGLWTCST.TVTSEIAPK   144
NTDB id 1160 ABDM36 RS02330 WP 000649326.1       ........AGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD..AGVVSSSKIQLARD.ANGLWTCST.TVTSEIAPK   144
NTDB id 1292 VP RS12235 WP 005479693.1           ..AQLGVDVAANDLGAIDS.........ALAGGS..PTLTFTFD.ADSSIDGGILTFTRD.AATGWSCLPTA..PAIAVD   136
NTDB id 1038 HON27 RS01510 WP 000993725.1        ....VGFTAATSSVCATYSID.....AFTETDGAG..KITCTLKGN.PKIATKIVELTRS.TDGAWTCVS.DVPTEFLPK   134
NTDB id 1092564 ACLSSQ RS14695 WP 014236553.1   ..ATIGLDTSKSGHCTYT.VD.....N.T.......GISCALKNAPTKVNGLAIKLARA.AGGAWSCDASALPTDMKPK   128
NTDB id 1192 PAKAF RS23870 WP 024947914.1        KEVPLGVAADANKLGTIA.LK.....P.DPADGTADITLTFTMGGAGPKNKGKIITLTRTAADGLWKCTS.DQDEQFIPK   146
NTDB id 1197 PSJM300 03940 AFN76866.1            ..GYIGITDSTS.YCDVD.LD.......TAADGH..IE.CTAKGGNAGKFDGKTITLNRT.ADGEWSCAS.TLDAKYKPG   136
NTDB id 1204 PSJM300 03935 AFN76865.1            ..GFIGIGASTN.YCTVT.LI.......ATATGS..IT.CTTQNGMPDRFNGKTITIART.TDGLWDCTS.DLDSRYKPG   138
consensus                                            *** ** * ***** *          * *    ** **    * ** *  * **!  * **!*!**  * *** **
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logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1        GCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1      GCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 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 1092564 ACLSSQ RS14695 WP 014236553.1   GCS............   131
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                                        *!*
```

X non conserved

X similar

X ≥ 50% conserved