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logo

NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT   80
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT   80
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGFT   80
NTDB id 1002 ACIAD RS15100 WP 004923779.1     .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTA....   71
NTDB id 167327 AL533 RS19025 WP 104919228.1  .....MNAQKGFTLIELMIVVAIIGILAAIAIPAYQNYIAKSQVSTGLTDITAGKANTETKLAE.GLTANLTNATD....   70
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNE.GAVPT..KLSD....   68
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSR.GWSVKSGTGTEDATK   74
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISE.GHTPSLVSSLP....   74
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINE.GKTPSLTSTDE....   72
consensus                                          **!!*!!!!!!!*!*!!*!!!!!*!*!**!* ** ** *** ** *!* *** *   !  *  * **
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logo

NTDB id 1184 GTF74 RS02335 WP 000649326.1     R...LGTVED...MGDGKIVIAPTASGALGGTIKYTF....DAGVVSSSKIQLARD.......ANGLWTCSTTVTSEIAP   143
NTDB id 1167 A1552VC RS11065 WP 000649326.1   R...LGTVED...MGDGKIVIAPTASGALGGTIKYTF....DAGVVSSSKIQLARD.......ANGLWTCSTTVTSEIAP   143
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    R...LGTVED...MGDGKIVIAPTASGALGGTIKYTF....DAGVVSSSKIQLARD.......ANGLWTCSTTVTSEIAP   143
NTDB id 1002 ACIAD RS15100 WP 004923779.1     GSSCVGVQEISASNATTNVATATCGASSA.GQII..VTM..DTTKAKGANITLTPT....YASGAVTWKCTTTSDKKYVP   142
NTDB id 167327 AL533 RS19025 WP 104919228.1  ....VGLQAS.SNACSS..ISVSVATTGL.TSIE.CTLQG..TSQISGKKISWERTADSATAGTTGIWRCKTDVADNLRP   139
NTDB id 1038 HON27 RS01510 WP 000993725.1     ....VGFTAATSSVCATYSIDAFTETDGA.GKIT.CTLKG..NPKIATKIVELT.......RSTDGAWTCVSDVPTEFLP   133
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KEVPLGVAADANKLGTIALK..PDPADGT.ADITLTFTMGGAGPKNKGKIITLTRT......AADGLWKCTSDQDEQFIP   145
NTDB id 1204 PSJM300 03935 AFN76865.1         ..GFIGIGASTN.YCTVTLI...ATATGS...IT.CTTQNGMPDRFNGKTITIART.......TDGLWDCTSDLDSRYKP   137
NTDB id 1197 PSJM300 03940 AFN76866.1         ..GYIGITDSTS.YCDVDLD...TAADGH...IE.CTAKGGNAGKFDGKTITLNRT.......ADGEWSCASTLDAKYKP   135
consensus                                         *!* *    **   ** * *** *  * !         * ** ** *** **        * **!*!**** * * !
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logo

NTDB id 1184 GTF74 RS02335 WP 000649326.1     KGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   KGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    KGCTAGATIN......   153
NTDB id 1002 ACIAD RS15100 WP 004923779.1     SECRG...........   147
NTDB id 167327 AL533 RS19025 WP 104919228.1  KSCGAS..........   145
NTDB id 1038 HON27 RS01510 WP 000993725.1     KGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KGCSK...........   150
NTDB id 1204 PSJM300 03935 AFN76865.1         GKCL............   141
NTDB id 1197 PSJM300 03940 AFN76866.1         GKCS............   139
consensus                                     **!**
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X non conserved

X similar

X ≥ 50% conserved