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logo

NTDB id 1191 AAC28468.1 484..945( )   ...MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAGN..   75
NTDB id 1184 GTF74 RS02335 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA..AG   78
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA..AG   78
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA..AG   78
NTDB id 383921 BSR23 RS03665 WP 025536236.1   MKHSKQKKQQGFTLIELMIVVAVIGVLAAIAMPQYQKYVAKSEVASVLATLTGAKTNVEAYTVENGLFPDGSASDA..T.   77
NTDB id 1292 VP RS12235 WP 005479693.1   MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAA.....A..T.   72
NTDB id 1038 HON27 RS01510 WP 000993725.1   .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLSDV.....   69
NTDB id 1192 PAKAF RS23870 WP 024947914.1   .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK   75
NTDB id 1197 PSJM300 03940 AFN76866.1   ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTDE.....   72
NTDB id 1204 PSJM300 03935 AFN76865.1   .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....   74
consensus   **  *** **!!*!!!!!!!*!*!!*!!!*!*!*!!*!* ** ****** ****** ** ** * ****  **  *

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NTDB id 1191 AAC28468.1 484..945( )   ...VIF.DSANGLDGKYFAPGGVTVTPDTGAINIAFDAGA.......NAGKTMTITPTVIAASGQISTWTCAPGAANGIE   144
NTDB id 1184 GTF74 RS02335 WP 000649326.1   FTRLGTV.EDMGDGKIVIAPTA.SGALG.GTIKYTFDA....G...VVSSSKIQLAR.....D.ANGLWTCSTT....VT   138
NTDB id 1167 A1552VC RS11065 WP 000649326.1   FTRLGTV.EDMGDGKIVIAPTA.SGALG.GTIKYTFDA....G...VVSSSKIQLAR.....D.ANGLWTCSTT....VT   138
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   FTRLGTV.EDMGDGKIVIAPTA.SGALG.GTIKYTFDA....G...VVSSSKIQLAR.....D.ANGLWTCSTT....VT   138
NTDB id 383921 BSR23 RS03665 WP 025536236.1   PTALGVP..SMHLGTV..AFTD.QSADG.GKISFTFATTASAGVSALVSGKKLTLTR.....TATSGGWDCSSSD...LG   143
NTDB id 1292 VP RS12235 WP 005479693.1   LAQLGVDVAANDLGA.....ID.SALAG.GSPTLTFTFDAD....SSIDGGILTFTR.....D.AATGWSCLPTA...PA   132
NTDB id 1038 HON27 RS01510 WP 000993725.1   ....GFTAATSSVCATYSIDAF.TETDGAGKIT.CTLK...G..NPKIATKIVELTR.....S.TDGAWTCVSD....VP   128
NTDB id 1192 PAKAF RS23870 WP 024947914.1   EVPLGVAADANKLGTIALKP...DPADGTADITLTFTM...GGAGPKNKGKITTLTR.....TAADGLWKCTSD....QD   140
NTDB id 1197 PSJM300 03940 AFN76866.1   .GYIGITDSTS.YCDVDLD....TAADGH..IE.CTAK...GGNAGKFDGKTITLNR.....T.ADGEWSCAST....LD   130
NTDB id 1204 PSJM300 03935 AFN76865.1   .GFIGIGASTN.YCTVTLI....ATATGS..IT.CTTQ...NGMPDRFNGKTITIAR.....T.TDGLWDCTSD....LD   132
consensus      ***       ** * ***    *** * * ** **       *    * ** *****      * * **!*!***    *

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NTDB id 1191 AAC28468.1 484..945( )   TRRLPTSCQ*.........   153
NTDB id 1184 GTF74 RS02335 WP 000649326.1   SEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   SEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   SEIAPKGCTAGATIN......   153
NTDB id 383921 BSR23 RS03665 WP 025536236.1   SELLPKTCK.........   152
NTDB id 1292 VP RS12235 WP 005479693.1   I..AVDGCS.........   139
NTDB id 1038 HON27 RS01510 WP 000993725.1   TEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1   EQFIPKGCSK........   150
NTDB id 1197 PSJM300 03940 AFN76866.1   AKYKPGKCS.........   139
NTDB id 1204 PSJM300 03935 AFN76865.1   SRYKPGKCL.........   141
consensus   **  ***!*
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X non conserved

X similar

X ≥ 50% conserved