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logo

NTDB id 1274 LPP RS09530 WP 011214168.1      .....MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT.....   69
NTDB id 1184 GTF74 RS02335 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....   75
NTDB id 1167 A1552VC RS11065 WP 000649326.1  MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....   75
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....   75
NTDB id 393529 FIV02 RS04035 WP 152487868.1 .....MQYARGFTLIELMIVVAIIGILAAIALPAYQNYVAKSQLAAGLSDIAGGRPSYEALVVTENVSSFNLS.......   68
NTDB id 1038 HON27 RS01510 WP 000993725.1    .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNE.GAVPTKLS.......   67
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 1274 LPP RS09530 WP 011214168.1      .GYVSPAATPN.VASIAIGANGVITITYTAAAGGGTIVMTPT.........................LQ.ANGDVTWTCTG   122
NTDB id 1184 GTF74 RS02335 WP 000649326.1    ......AAGFTRLGTV.............EDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLARD.ANGLWTCST   135
NTDB id 1167 A1552VC RS11065 WP 000649326.1  ......AAGFTRLGTV.............EDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLARD.ANGLWTCST   135
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   ......AAGFTRLGTV.............EDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLARD.ANGLWTCST   135
NTDB id 393529 FIV02 RS04035 WP 152487868.1 ..DIGLVSETSRCTVS.........MNYDSADGTGDIT.CTVKG............SPIVNGKTIQLVRS.SSGAWKCQV   123
NTDB id 1038 HON27 RS01510 WP 000993725.1    ..DVGFTAATSSVCAT......YSIDAFTETDGAGKIT.CTLKG............NPKIATKIVELTRS.TDGAWTCVS   125
NTDB id 1192 PAKAF RS23870 WP 024947914.1    EVPLGVAADANKLGTI........ALKPDPADGTADITLTFTMGG...........AGPKNKGKIITLTRTAADGLWKCTS   137
NTDB id 1197 PSJM300 03940 AFN76866.1        .GYIGITDSTS.YCDV........DLD.TAADGH..IE.CTAKGG...........NAGKFDGKTITLNRT.ADGEWSCAS   127
NTDB id 1204 PSJM300 03935 AFN76865.1        .GFIGIGASTN.YCTV........TLI.ATATGS..IT.CTTQNG...........MPDRFNGKTITIART.TDGLWDCTS   129
consensus                                       ** **  ******         *    * ! * ! *  *  *           *  *  * * * ** ****!*!**

logo

NTDB id 1274 LPP RS09530 WP 011214168.1      GTL.LAKYRPASCRP...........   136
NTDB id 1184 GTF74 RS02335 WP 000649326.1    TVT..SEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1  TVT..SEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   TVT..SEIAPKGCTAGATIN......   153
NTDB id 393529 FIV02 RS04035 WP 152487868.1  DASISAIYHPQGCAH...........   138
NTDB id 1038 HON27 RS01510 WP 000993725.1    DVP..IEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1    DQD..EQFIPKGCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1        TLD..AKYKPGKCS............   139
NTDB id 1204 PSJM300 03935 AFN76865.1        DLD..SRYKPGKCL............   141
consensus                                     *    * * !**!*
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X non conserved

X similar

X ≥ 50% conserved