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logo

NTDB id 1191 AAC28468.1 484..945( )          ...MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAGNVI   77
NTDB id 1077917 ACJZU6 RS03635 WP 408900657.1 .....MKKQQGFTLIELMIVVAVIGVLSAIAIPKYQQYVEKGAIAAALSSATALKTNYEDYLAVSGAAPTTSA.......   68
NTDB id 1184 GTF74 RS02335 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTA.......   73
NTDB id 1167 A1552VC RS11065 WP 000649326.1  MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTA.......   73
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTA.......   73
NTDB id 1292 VP RS12235 WP 005479693.1       MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAA.........   71
NTDB id 1038 HON27 RS01510 WP 000993725.1    .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLSD.......   68
NTDB id 1192 PAKAF RS23870 WP 024947914.1    .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATK.   74
NTDB id 1197 PSJM300 03940 AFN76866.1        ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTD......   71
consensus                                        * * !*!!!!!!!!!!*!*!!*!*!*!*!!*!* ** ** *** ****** ******  ** * ***

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NTDB id 1191 AAC28468.1 484..945( )          FDSANGLDGKYFAP.............GGVTV..T.....PDTGAINIAFDAGANAGKTMTITPTVIAASGQISTWTCAP   137
NTDB id 1077917 ACJZU6 RS03635 WP 408900657.1 .....AIGATKFSLGTVT.....VT.SGAIAV..AVTEGGGKGAG...............VTL.T......RADNAWTCAV   114
NTDB id 1184 GTF74 RS02335 WP 000649326.1    .....GTAAGFTRLGTVE.....DMGDGKIVI..APTASGALGGTIKYTFDAGVVSSSKIQLAR......DANGLWTCST   135
NTDB id 1167 A1552VC RS11065 WP 000649326.1  .....GTAAGFTRLGTVE.....DMGDGKIVI..APTASGALGGTIKYTFDAGVVSSSKIQLAR......DANGLWTCST   135
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   .....GTAAGFTRLGTVE.....DMGDGKIVI..APTASGALGGTIKYTFDAGVVSSSKIQLAR......DANGLWTCST   135
NTDB id 1292 VP RS12235 WP 005479693.1       TLAQLGVDVAANDLGAIDSAL....AGGSPTL..TFT.FD..........ADSSIDGGILTFTR......DAATGWSCLP   128
NTDB id 1038 HON27 RS01510 WP 000993725.1    ....VGFTAATSSVCATYSIDAFTETDGAGKI.TCTLK............GNPKIATKIVELTR......STDGAWTCVS   125
NTDB id 1192 PAKAF RS23870 WP 024947914.1    KEVPLGVAADANKLGTIA.LKP.DPADGTADITLTFTMGG..........AGPKNKGKIITLTR.....TAADGLWKCTS   137
NTDB id 1197 PSJM300 03940 AFN76866.1        .EGYIGITDS.TSYCDVD.LD..TAADGHI...ECTAKGG..........NAGKFDGKTITLNR......TADGEWSCAS   127
consensus                                        ***** * *****       **! * *    *  *   *       ** *    *** *        * * !*! *

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NTDB id 1191 AAC28468.1 484..945( )          GAANGIETRRLPTSCQ*...........   153
NTDB id 1077917 ACJZU6 RS03635 WP 408900657.1 SASSA...TIKLNGCTTNETSS......   133
NTDB id 1184 GTF74 RS02335 WP 000649326.1    TVTS....EIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1  TVTS....EIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   TVTS....EIAPKGCTAGATIN......   153
NTDB id 1292 VP RS12235 WP 005479693.1       TAPA....IA.VDGCS............   139
NTDB id 1038 HON27 RS01510 WP 000993725.1    DVPT....EFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1    DQDE....QFIPKGCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1        TLDA....KYKPGKCS............   139
consensus                                    ** *        ***!*
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X non conserved

X similar

X ≥ 50% conserved