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logo

NTDB id 1191 AAC28468.1 484..945( )            ...MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAGNVI    77
NTDB id 1038 HON27 RS01510 WP 000993725.1      .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGA.VPTKLS.......    67
NTDB id 1067431 ACHZDB RS07615 WP 399425124.1  MKTNKQKKQQGFTLIELMIVVAVIGVLAAIAIPQYQKYVAKAEVAAALASMAGVKTNVEAYTVEYGVFPAASQSSALG..    78
NTDB id 1184 GTF74 RS02335 WP 000649326.1      MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAG..    78
NTDB id 1167 A1552VC RS11065 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAG..    78
NTDB id 1160 ABDM36 RS02330 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAG..    78
NTDB id 1292 VP RS12235 WP 005479693.1         MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENG........IAAAAT    72
NTDB id 1192 PAKAF RS23870 WP 024947914.1      .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK    75
consensus                                      **   *** !*!!!!!!!!!!*!*!!*!!!!*!*!*!!*!* ** ** *** * ***! ** **    ****   **   ***

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NTDB id 1191 AAC28468.1 484..945( )            FDS......ANGLDGKYFAPGGVTVTPDTGAINIAFDAGA....NAGKTMTITPTVIAASGQISTWTCAPGAAN.GIETR   146
NTDB id 1038 HON27 RS01510 WP 000993725.1      ..DVGFTAATSSVCATYSIDA.FTETDGAGKITCTLKGN...PKIATKIVEL.....TRS.TDGAWTCVSD.....VPTE   130
NTDB id 1067431 ACHZDB RS07615 WP 399425124.1  VPS......NIPQGTMAFSQ....GTSAAGDIIFTFAGSDVSSLLINKKFAL.....KRD.VNGGWDCTASGASPAVTSD   142
NTDB id 1184 GTF74 RS02335 WP 000649326.1      FTRLGTV.EDMGDGKIVIAP..TASGALGGTIKYTFDAGV....VSSSKIQL.....ARD.ANGLWTCSTT.....VTSE   140
NTDB id 1167 A1552VC RS11065 WP 000649326.1    FTRLGTV.EDMGDGKIVIAP..TASGALGGTIKYTFDAGV....VSSSKIQL.....ARD.ANGLWTCSTT.....VTSE   140
NTDB id 1160 ABDM36 RS02330 WP 000649326.1     FTRLGTV.EDMGDGKIVIAP..TASGALGGTIKYTFDAGV....VSSSKIQL.....ARD.ANGLWTCSTT.....VTSE   140
NTDB id 1292 VP RS12235 WP 005479693.1         LAQLGVDVAANDLGAIDSA...LAGGSPTLT..FTFDAD...SSIDGGILTF.....TRDAAT.GWSCLPT.....AP.A   132
NTDB id 1192 PAKAF RS23870 WP 024947914.1      EVPLGVAADANKLGTIALKPD.PADGTADITLTFTMGGAG..PKNKGKIITL.....TRTAADGLWKCTSD.....QDEQ   142
consensus                                         **         * *  **   * *  **** ******     *    * *      ** * * !*!***     * **

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NTDB id 1191 AAC28468.1 484..945( )            RLPTSCQ*...........   153
NTDB id 1038 HON27 RS01510 WP 000993725.1      FLPKGCTAGTPTEGAITSL   149
NTDB id 1067431 ACHZDB RS07615 WP 399425124.1  LLPKNCQ............   149
NTDB id 1184 GTF74 RS02335 WP 000649326.1      IAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1    IAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1     IAPKGCTAGATIN......   153
NTDB id 1292 VP RS12235 WP 005479693.1         IAVDGCS............   139
NTDB id 1192 PAKAF RS23870 WP 024947914.1      FIPKGCSK...........   150
consensus                                      * ***!*
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X non conserved

X similar

X ≥ 50% conserved