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logo
NTDB id 1109 OK783 RS10670 WP 157147416.1     ....MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTSAG....   72
NTDB id 1002 ACIAD RS15100 WP 004923779.1     .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTA.GS.   73
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 422226 G0030 RS01380 WP 180182068.1   .....MNAQKGFTLIELMIVVAIIGILAAIAIPAYQNYTAKSQVAAGLAEISPAKTNAEAKLAEGVS.ATITSA......   68
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAV.PTKLSD......   68
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.AA........AA   71
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWS.VKSGTGTEDATK   74
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKT.PSLTSTDE....   72
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHT.PSLVSSLP....   74
consensus                                              **!!*!!!!!!!*!**!*!!!*!*!*!**!* ** ** *** ** *** *** *  *** *  **
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logo
NTDB id 1109 OK783 RS10670 WP 157147416.1     ......VASASTIKGKYVQK.....VEVAKGVVTAQMASTGVNKEIQDKKLSLWAK...RQDGSVKWFCGQPVTRTGDND   138
NTDB id 1002 ACIAD RS15100 WP 004923779.1     ..SCVGVQEISASNATTNVA.TATCGASSAGQIIVTMD....TTKAKGANITLTPT...YASGAVTWKCTTT........   135
NTDB id 1184 GTF74 RS02335 WP 000649326.1     R...LGTVEDMG.DGKIVIAPTASGALG..GTIKYTFD....AGVVSSSKIQLAR......DANGLWTCSTT........   136
NTDB id 1167 A1552VC RS11065 WP 000649326.1   R...LGTVEDMG.DGKIVIAPTASGALG..GTIKYTFD....AGVVSSSKIQLAR......DANGLWTCSTT........   136
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    R...LGTVEDMG.DGKIVIAPTASGALG..GTIKYTFD....AGVVSSSKIQLAR......DANGLWTCSTT........   136
NTDB id 422226 G0030 RS01380 WP 180182068.1   ....SALGISTSKRCTALSVS...VAKTGESSITCTLA...GTTSINGRKIQWVRGADNTDGTIGTWTCKTN........   130
NTDB id 1038 HON27 RS01510 WP 000993725.1     ....VGFTAATSSVCATYSIDA.FTETDGAGKITCTLK...GNPKIATKIVELT......RSTDGAWTCVSD........   126
NTDB id 1292 VP RS12235 WP 005479693.1        TLAQLGVDVAANDLGAIDS.....ALAGGSPTLTFTFD...ADSSIDGGILTFTR.....D.AATGWSCLPT........   129
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KEVPLGVAADANKLGTIALKP...DPADGTADITLTFTMGGAGPKNKGKIITLTR.....TAADGLWKCTSD........   138
NTDB id 1197 PSJM300 03940 AFN76866.1         ..GYIGITDSTS.YCDVDLD....TAADGH..IE.CTAKGGNAGKFDGKTITLNR.....T.ADGEWSCAST........   128
NTDB id 1204 PSJM300 03935 AFN76865.1         ..GFIGIGASTN.YCTVTLI....ATATGS..IT.CTTQNGMPDRFNGKTITIAR.....T.TDGLWDCTSD........   130
consensus                                        ***   *   * * *      *  * * ** *         * ** *** *        * * !*!***
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logo
NTDB id 1109 OK783 RS10670 WP 157147416.1     DTVADANNAIDTKHLPSTCRDESSAT......   164
NTDB id 1002 ACIAD RS15100 WP 004923779.1     .........SDKKYVPSECRG...........   147
NTDB id 1184 GTF74 RS02335 WP 000649326.1     .........VTSEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   .........VTSEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    .........VTSEIAPKGCTAGATIN......   153
NTDB id 422226 G0030 RS01380 WP 180182068.1   .........VADDFIPKTCESASIS.......   146
NTDB id 1038 HON27 RS01510 WP 000993725.1     .........VPTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1292 VP RS12235 WP 005479693.1        .........APAI.AVDGCS............   139
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .........QDEQFIPKGCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1         .........LDAKYKPGKCS............   139
NTDB id 1204 PSJM300 03935 AFN76865.1         .........LDSRYKPGKCL............   141
consensus                                              * * * ***!*
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X non conserved

X similar

X ≥ 50% conserved