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logo

NTDB id 1119 NGFG RS11465 WP 017147169.1      ....MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGKWPENNTSAGVAS.    75
NTDB id 128850 IX91 RS02210 WP 004742603.1    MQSLQSRKTKGFTLIELMIVVAVIGVLAAIAVPQYQTYVAKSEASSALASITGHRANVETYVLTEGSFPANSSALAIPS.    79
NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGF.    79
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGF.    79
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAAGF.    79
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPAYQNYIAKSQASEAFTLADGLKTTINTNLQAGTCFAGGATAVTAADK    75
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLSDV......    69
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDA.TK    74
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....    74
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTDE.....    72
consensus                                           * **!!*!!!!!!!*!**!*!!!*!*!*!!*!****  * *** ****** *****  ****** *** ***

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NTDB id 1119 NGFG RS11465 WP 017147169.1      ..PPTDI...KGKYVK.......EVEVKNGVVTATMLSSGVNNEIKGKKLSLWGRRENGSVKWFCGQPVTRADDDTVADA   143
NTDB id 128850 IX91 RS02210 WP 004742603.1    ..SPLGIISY............DNVSSAAGDILFTFSSSGVSPDVVSKKVSLKRD.GNG..AWTCQ...............   128
NTDB id 1184 GTF74 RS02335 WP 000649326.1     ..TRLGTVEDMG.DGKIVIAP.TASGALGGTIKYTFDAGV....VSSSKIQLARD.ANG..LWTCS...............   134
NTDB id 1167 A1552VC RS11065 WP 000649326.1   ..TRLGTVEDMG.DGKIVIAP.TASGALGGTIKYTFDAGV....VSSSKIQLARD.ANG..LWTCS...............   134
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    ..TRLGTVEDMG.DGKIVIAP.TASGALGGTIKYTFDAGV....VSSSKIQLARD.ANG..LWTCS...............   134
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     VSG........KYGDAEI....GGTAPNCTITYTFKSSGVSNKLTSTKIVMNVS.ETG..ILTKN..............   125
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....GFTAATSSVCATYSIDAFTETDGAGKIT.CTLKG..NPKIATKIVELTRS.TDG..AWTCV..............   124
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KEVPLGVAADANKLGTIALKP..DPADGTADITLTFTMGGAGPKNKGKIITLTRTAADG..LWKCT..............   136
NTDB id 1204 PSJM300 03935 AFN76865.1         ..GFIGIGASTN.YCTVTLI...ATATGS..IT.CTTQNGMPDRFNGKTITIART.TDG..LWDCT..............   128
NTDB id 1197 PSJM300 03940 AFN76866.1         ..GYIGITDSTS.YCDVDLD...TAADGH..IE.CTAKGGNAGKFDGKTITLNRT.ADG..EWSCA..............   126
consensus                                        ***        * * *    * * *** *****  **   ** ****** ** **!  *****

logo

NTDB id 1119 NGFG RS11465 WP 017147169.1      KDGKEIDTKHLPSTCRDKASDAK.....   166
NTDB id 128850 IX91 RS02210 WP 004742603.1    ...SDVDSDLKPKGCS............   141
NTDB id 1184 GTF74 RS02335 WP 000649326.1     ...TTVTSEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   ...TTVTSEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    ...TTVTSEIAPKGCTAGATIN......   153
NTDB id 1079 FDQ49 RS11140 WP 022575996.1     .SGTDTPVELLPQSFVAS..........   142
NTDB id 1038 HON27 RS01510 WP 000993725.1     ...SDVPTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1     ...SDQDEQFIPKGCSK...........   150
NTDB id 1204 PSJM300 03935 AFN76865.1         ...SDLDSRYKPGKCL............   141
NTDB id 1197 PSJM300 03940 AFN76866.1         ...STLDAKYKPGKCS............   139
consensus                                        ******* !*****
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X non conserved

X similar

X ≥ 50% conserved