```
logo
NTDB id 1400 DSB67 RS12660 WP 010643261.1    MKTNKQKKQQGFTLIELMIVVAIIGILAALAIPAYQDYTKKATMAEFPKIASSTKLAVELCAHEHAADAGAFKTKCHQ..    78
NTDB id 1184 GTF74 RS02335 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATE....GSFPATTAG..    74
NTDB id 1167 A1552VC RS11065 WP 000649326.1  MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATE....GSFPATTAG..    74
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATE....GSFPATTAG..    74
NTDB id 359465 EA719 RS01555 WP 000993717.1 .....MNAQKGFTLIELMIVVAIIGILAAIAIPAYQNYIAKSQVSTGLADITAGKTNAETKLA.E....GLTAALTDV..    68
NTDB id 1038 HON27 RS01510 WP 000993725.1    .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVN.E....GAVPT..KL..    66
NTDB id 1002 ACIAD RS15100 WP 004923779.1    .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILN.....AGALVAGTPST    70
NTDB id 1292 VP RS12235 WP 005479693.1       MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQE.....NGIAA......    69
NTDB id 1192 PAKAF RS23870 WP 024947914.1    .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSR.....GWSVKSGTGTE    70
NTDB id 1197 PSJM300 03940 AFN76866.1        ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINE.....GKTPSLTSTDE    72
NTDB id 1204 PSJM300 03935 AFN76865.1        .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISE.....GHTPSLVSSLP    74
consensus                                        * * **!!*!!!!!!!*!*!!*!!!*!*!*!**!* ** ** *** ** *** **  *        *  *
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logo
NTDB id 1400 DSB67 RS12660 WP 010643261.1    ......TANGIPAAF.TLNNLTVTPAAGA..ASGAISITVNAAND.........KGPIKQNEKYILTATYA....SNGVT   136
NTDB id 1184 GTF74 RS02335 WP 000649326.1    ......TAAGFTRLG...........TVEDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLAR.......DANGL   130
NTDB id 1167 A1552VC RS11065 WP 000649326.1  ......TAAGFTRLG...........TVEDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLAR.......DANGL   130
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   ......TAAGFTRLG...........TVEDMGDGKIVIAPTASGALGGTIKYTFDAGVVSSSKIQLAR.......DANGL   130
NTDB id 359465 EA719 RS01555 WP 000993717.1 ......TTLGLQQST....NACAITANIGTNGASNITCTLK......GT.......SQINGKKIEWIRDADNATNGTTGA   125
NTDB id 1038 HON27 RS01510 WP 000993725.1    ......SDVGFTAATSSVCATYSIDAFTETDGAGKITCTLK......GN.......PKIATKIVELTR.......STDGA   120
NTDB id 1002 ACIAD RS15100 WP 004923779.1    AGS...SCVGVQEISASNATTNVATATCG..ASSAGQIIVTM............DTTKAKGANITLTPTYA....SGAVT   129
NTDB id 1292 VP RS12235 WP 005479693.1       ..AATLAQLGVDVAANDLGAIDS..ALAG..GSPTLTFTFD.............ADSSIDGGILTFTRD.......AATG   123
NTDB id 1192 PAKAF RS23870 WP 024947914.1    DATKKEVPLGVAADANKLGTIALKPDPAD..GTADITLTFTM..........GGAGPKNKGKIITLTRTA......ADGL   132
NTDB id 1197 PSJM300 03940 AFN76866.1        ......GYIGITDSTS.YCDVDLD.TAAD..GH..IE.CTAK..........GGNAGKFDGKTITLNRT.......ADGE   122
NTDB id 1204 PSJM300 03935 AFN76865.1        ......GFIGIGASTN.YCTVTLI.ATAT..GS..IT.CTTQ..........NGMPDRFNGKTITIART.......TDGL   124
consensus                                          * *!*               *  *  * * ***  *             *  * *  * * *         * *
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logo
NTDB id 1400 DSB67 RS12660 WP 010643261.1    WSSECKDATNAVQTSYCP................   154
NTDB id 1184 GTF74 RS02335 WP 000649326.1    WTCST.....TVTSEIAPKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1  WTCST.....TVTSEIAPKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1   WTCST.....TVTSEIAPKGCTAGATIN......   153
NTDB id 359465 EA719 RS01555 WP 000993717.1 WRCKT.....DVAENLRPKSCGAS..........   144
NTDB id 1038 HON27 RS01510 WP 000993725.1    WTCVS.....DVPTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1002 ACIAD RS15100 WP 004923779.1    WKCTT.....TSDKKYVPSECRG...........   147
NTDB id 1292 VP RS12235 WP 005479693.1       WSCLP.....TAPAI.AVDGCS............   139
NTDB id 1192 PAKAF RS23870 WP 024947914.1    WKCTS.....DQDEQFIPKGCSK...........   150
NTDB id 1197 PSJM300 03940 AFN76866.1        WSCAS.....TLDAKYKPGKCS............   139
NTDB id 1204 PSJM300 03935 AFN76865.1        WDCTS.....DLDSRYKPGKCL............   141
consensus                                    !****      ** * * ******
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X non conserved

X similar

X ≥ 50% conserved