```
logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTA....   76
NTDB id 343798 EPB58 RS19015 WP 050936251.1   MKTNKQKKQQGFTLIELMIVVAIIGVLSAIAVPAYQNYVAKGEAATALGSLRALVTPAELKLQQDGELSGVVS.DL....   75
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAATLAQL....   76
NTDB id 1002 ACIAD RS15100 WP 004923779.1     .....MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTAGS..   73
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK   75
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....   74
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTDE.....   72
consensus                                     **  *** **!!*!!!!!!!*!*!!*!*!*!*!*!**!* ** ** *!* ****** *** *   **  * *

logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1     .......AGFTRLGTVEDMGDGKIVIAPTA.SGALGGTIKYTFDA...G.VVSSSKIQLARDA...NGLWTCSTTVTSEI   141
NTDB id 1167 A1552VC RS11065 WP 000649326.1   .......AGFTRLGTVEDMGDGKIVIAPTA.SGALGGTIKYTFDA...G.VVSSSKIQLARDA...NGLWTCSTTVTSEI   141
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    .......AGFTRLGTVEDMGDGKIVIAPTA.SGALGGTIKYTFDA...G.VVSSSKIQLARDA...NGLWTCSTTVTSEI   141
NTDB id 343798 EPB58 RS19015 WP 050936251.1   ....GG.SASHALGN..........ITTTG.TGVSAAVLTFTFTK...G.SLSGDAITL.TKS...TSGWVCNDGTT...   128
NTDB id 1292 VP RS12235 WP 005479693.1        ....GVDVAANDLGA..........IDSAL.AG.GSPTLTFTFDA...DSSIDGGILTFTRDA...ATGWSCLPTAPA.I   133
NTDB id 1002 ACIAD RS15100 WP 004923779.1     .SCVGVQEISASNAT........TNVATATCGASSAGQII..VTM..DTTKAKGANITLTPTYASGAVTWKCTTTSDKKY   140
NTDB id 1192 PAKAF RS23870 WP 024947914.1     EVPLGVAADANKLGT........IALKPD..PADGTADITLTFTMGGAGPKNKGKIITLTRTAA..DGLWKCTSDQDEQF   143
NTDB id 1204 PSJM300 03935 AFN76865.1         .GFIGIGASTN.YCT........VTLI.A..TATGS..IT.CTTQNGMPDRFNGKTITIART.T..DGLWDCTSDLDSRY   135
NTDB id 1197 PSJM300 03940 AFN76866.1         .GYIGITDSTS.YCD........VDLD.T..AADGH..IE.CTAKGGNAGKFDGKTITLNRT.A..DGEWSCASTLDAKY   133
consensus                                         ** **   ***         * *  *  **  ** * ***     *  * *  *** * *    **!*!****

logo
NTDB id 1184 GTF74 RS02335 WP 000649326.1     APKGCTAGATIN   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   APKGCTAGATIN   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    APKGCTAGATIN   153
NTDB id 343798 EPB58 RS19015 WP 050936251.1   ILDNCN......   134
NTDB id 1292 VP RS12235 WP 005479693.1        AVDGCS......   139
NTDB id 1002 ACIAD RS15100 WP 004923779.1     VPSECRG.....   147
NTDB id 1192 PAKAF RS23870 WP 024947914.1     IPKGCSK.....   150
NTDB id 1204 PSJM300 03935 AFN76865.1         KPGKCL......   141
NTDB id 1197 PSJM300 03940 AFN76866.1         KPGKCS......   139
consensus                                      * *!*
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X non conserved

X similar

X ≥ 50% conserved