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logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK   75
NTDB id 1197 PSJM300 03940 AFN76866.1         ...MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFEQAINEGKTPSLTSTDE.....   72
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....   74
NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATT........   72
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATT........   72
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATT........   72
NTDB id 1054861 BCT18 RS01325 WP 016790035.1  .MNNKRTNQKGFTLIELMIVVAIIGVLSAIAVPAYKDYVSKSEASSALATLKSLQTPAELTFQEEGALSS.AI.......   71
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAATL.......   73
consensus                                         * * **!!*!!!!!!!!!*!!*!*!*!*! !**!* *!  * *!! ** ***  !  *   **    *

logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     EVPLGVAADANKLGTIALK.......PDPADGTADITLTFTMGGAGPKNKGKIITLTRTAADGLWKCTSDQDEQFIPKGC   148
NTDB id 1197 PSJM300 03940 AFN76866.1         .GYIGITDSTS.YCDVDLD........TAADGH..IE.CTAKGGNAGKFDGKTITLNRT.ADGEWSCASTLDAKYKPGKC   138
NTDB id 1204 PSJM300 03935 AFN76865.1         .GFIGIGASTN.YCTVTLI........ATATGS..IT.CTTQNGMPDRFNGKTITIART.TDGLWDCTSDLDSRYKPGKC   140
NTDB id 1184 GTF74 RS02335 WP 000649326.1     ...AGTAAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.G.VVSSSKIQLARD.ANGLWTCSTTVTSEIAPKGC   146
NTDB id 1167 A1552VC RS11065 WP 000649326.1   ...AGTAAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.G.VVSSSKIQLARD.ANGLWTCSTTVTSEIAPKGC   146
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    ...AGTAAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.G.VVSSSKIQLARD.ANGLWTCSTTVTSEIAPKGC   146
NTDB id 1054861 BCT18 RS01325 WP 016790035.1  .SALGVSSGSNTLGTLS........ISNVGSSTVSPEILFTFNG.G.SLNSATLTLSRD.ATTGWACTKST.AVPDLDGC   138
NTDB id 1292 VP RS12235 WP 005479693.1        .AQLGVDVAANDLGAID........SALAG...GSPTLTFTFDA.DSSIDGGILTFTRD.AATGWSCLPTA.PAIAVDGC   138
consensus                                        *!****   *****            ***    ** *** * *  *    *** !* * **!*!****     * *!

logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     SK.....   150
NTDB id 1197 PSJM300 03940 AFN76866.1         S......   139
NTDB id 1204 PSJM300 03935 AFN76865.1         L......   141
NTDB id 1184 GTF74 RS02335 WP 000649326.1     TAGATIN   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   TAGATIN   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    TAGATIN   153
NTDB id 1054861 BCT18 RS01325 WP 016790035.1  R......   139
NTDB id 1292 VP RS12235 WP 005479693.1        S......   139
consensus                                     *
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X non conserved

X similar

X $\geq$ 50% conserved