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logo

NTDB id 408187 GPY15 RS10750 WP 158161902.1   .....MKKQQGFSLIELMIVIAVIGVLTAVAIPQYQNYVQKSELGAALASVTALKINVEDKIATDGAFPTIANSDMS...   72
NTDB id 1184 GTF74 RS02335 WP 000649326.1     MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA...   77
NTDB id 1167 A1552VC RS11065 WP 000649326.1   MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA...   77
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA...   77
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAV.PTKLSDV.....   69
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.A........AAA   71
NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWS.VKSGTGTEDATK   74
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHT.PSLVSSLP....   74
consensus                                           * **!!*!!!!!!!*!*!!*!*!*!*!*!!!!* *! ** *** ****** *!* *   ** *

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NTDB id 408187 GPY15 RS10750 WP 158161902.1   .....SELGATNTILGTMET.........KQSAGAAGQIIL.TLS..GKTQNSTKKIALER.DANGEWICLTDLTATAIF   134
NTDB id 1184 GTF74 RS02335 WP 000649326.1     .....G.....FTRLGTVEDMGDGKIVIAPTASGALGGTIK.YTF..DAGVVSSSKIQLAR.DANGLWTCSTTVT.SEIA   142
NTDB id 1167 A1552VC RS11065 WP 000649326.1   .....G.....FTRLGTVEDMGDGKIVIAPTASGALGGTIK.YTF..DAGVVSSSKIQLAR.DANGLWTCSTTVT.SEIA   142
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    .....G.....FTRLGTVEDMGDGKIVIAPTASGALGGTIK.YTF..DAGVVSSSKIQLAR.DANGLWTCSTTVT.SEIA   142
NTDB id 1038 HON27 RS01510 WP 000993725.1     .....GFTAATSSVCATYSID.......AFTETDGAGKITCTLK...GNPKIATKIVELTR.STDGAWTCVSDVP.TEFL   132
NTDB id 1292 VP RS12235 WP 005479693.1        TLAQLGVDVAAN.DLGAIDSA............LAGGSPTLTFTFD.ADSSIDGGILTFTRDAAT.GWSCLPTAP.A.IA   134
NTDB id 1192 PAKAF RS23870 WP 024947914.1     KEVPLGVAADAN.KLGTIALK........PDPADGTADITLTFTMGGAGPKNKGKIITLTRTAADGLWKCTSDQD.EQFI   144
NTDB id 1204 PSJM300 03935 AFN76865.1         ..GFIGIGASTN.YC.TVTLI........A...TATGSITC.TTQNGMPDRFNGKTITIAR.TTDGLWDCTSDLD.SRYK   136
consensus                                           *      * *****               * *     **       * *  * *  !  * **!*!** *  * *

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NTDB id 408187 GPY15 RS10750 WP 158161902.1   PKGCNTGTIL.......   144
NTDB id 1184 GTF74 RS02335 WP 000649326.1     PKGCTAGATIN......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   PKGCTAGATIN......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    PKGCTAGATIN......   153
NTDB id 1038 HON27 RS01510 WP 000993725.1     PKGCTAGTPTEGAITSL   149
NTDB id 1292 VP RS12235 WP 005479693.1        VDGCS............   139
NTDB id 1192 PAKAF RS23870 WP 024947914.1     PKGCSK...........   150
NTDB id 1204 PSJM300 03935 AFN76865.1         PGKCL............   141
consensus                                     ***!* *
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X non conserved

X similar

X $\geq$ 50% conserved