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logo

NTDB id 1191 AAC28468.1 484..945( )      ...MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAGNVI    77
NTDB id 1184 GTF74 RS02335 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....  75
NTDB id 1167 A1552VC RS11065 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....  75
NTDB id 1160 ABDM36 RS02330 WP 000649326.1 MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGT.....  75
NTDB id 103949 QUF32 RS13290 WP 143683926.1 MNANQRKQQQGFTLIELMIVVAIIGVLSAIAVPAYKNYVSKSELASGLATLKSLITPAELYIQETGSLSGASASA.....  75
NTDB id 1292 VP RS12235 WP 005479693.1   MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAATLAQ.....  75
NTDB id 1038 HON27 RS01510 WP 000993725.1 .....MNAQKGFTLIELMIVVAIIGILAAIAIPQYQNYIAKSQVNSGFSEISSIKTGYENAVNEGAVPTKLSDVGFT...  72
NTDB id 1192 PAKAF RS23870 WP 024947914.1 .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATK.  74
NTDB id 1204 PSJM300 03935 AFN76865.1    .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....  74
consensus                                *     * **!!*!!!!!!!*!*!!*!*!*!*!*!**!* ** ****** ****** ** ** * **  * **

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NTDB id 1191 AAC28468.1 484..945( )      FDSANGL..DGKYFAPGGVTVT.....PDTGAINIAFDAG....ANAGKTMTITPTVIAASGQISTWTCAPGAANGIETR  146
NTDB id 1184 GTF74 RS02335 WP 000649326.1 .AAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA....GVVSSSKIQLAR......DANGLWTCSTTVTSEIA..  142
NTDB id 1167 A1552VC RS11065 WP 000649326.1 .AAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA....GVVSSSKIQLAR......DANGLWTCSTTVTSEIA..  142
NTDB id 1160 ABDM36 RS02330 WP 000649326.1 .AAGFTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA....GVVSSSKIQLAR......DANGLWTCSTTVTSEIA..  142
NTDB id 103949 QUF32 RS13290 WP 143683926.1 ....FGVVSGGVALGEIT...VP.....TASKLTFTFN....DSSINGSTIDYDR.......SSGGWVCKVTLASGITL.  131
NTDB id 1292 VP RS12235 WP 005479693.1   ....LGVDVAANDLGAID...SALA..GGSPTLTFTFDA...DSSIDGGILTFTR......DAATGWSCLPTAPA.IAV.  135
NTDB id 1038 HON27 RS01510 WP 000993725.1 .......AATSSVCATYSI..DAFTETDGAGKITCTLKG...NPKIATKIVELTR......STDGAWTCVSDVPTEFL..  132
NTDB id 1192 PAKAF RS23870 WP 024947914.1 KEVPLGVAADANKLGTIAL..KP.DPADGTADITLTFTMGGAGPKNKGKITLTR.....TAADGLWKCTSDQDEQFI..  144
NTDB id 1204 PSJM300 03935 AFN76865.1    ..GFIGIGASTNYC.TVTL..IA....TATGSITC.TTQNGMPDRFNGKTITIAR......TTDGLWDCTSDLDSRYK..  136
consensus                                     ** **   **   *   *      **********      * *  * * *       * **!*!**** * *

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NTDB id 1191 AAC28468.1 484..945( )      RLPTSCQ*...........  153
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 103949 QUF32 RS13290 WP 143683926.1 DIPKSCQ...........  138
NTDB id 1292 VP RS12235 WP 005479693.1   ...DGCS............  139
NTDB id 1038 HON27 RS01510 WP 000993725.1 ..PKGCTAGTPTEGAITSL  149
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 ≥ 50% conserved