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logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     .....MKAQKGFTLIELMIVVAIIGILAAIAIPQYQNYVARSEGASALASVNPLKTTVEEALSRGWSVKSGTGTEDATKK   75
NTDB id 1204 PSJM300 03935 AFN76865.1         .MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFEQAISEGHTPSLVSSLP.....   74
NTDB id 1258 GCO85 RS09745 WP 014844212.1     ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQ........   66
NTDB id 1274 LPP RS09530 WP 011214168.1       ......MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQ........   66
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 1168121 AL542 RS07470 WP 040528180.1 .....MKKQQGFSLVELMIVVAIIGALTAFAIPAYQDYVQRSEAMAGLATLKSLQTPAQLEEQETGVFPT.S........   66
NTDB id 1292 VP RS12235 WP 005479693.1        MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGIAAAAT........   72
consensus                                           * **!!*!*!!!!!!!*!!*!*!*!*!*!!*!* ** ** *!* *  *** *** * * ** ****
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logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     EVPLGVAAD.ANKLGTIA.......LKPDPADGTADITLTFTMGGAGPKNKGKIITLTRTAADGLWKCTSDQD..EQFIP   145
NTDB id 1204 PSJM300 03935 AFN76865.1         .GFIGIGAS.TNYC.TVT.......LIA...TATGSITC.TTQNGMPDRFNGKTITIAR.TTDGLWDCTSDLD..SRYKP   137
NTDB id 1258 GCO85 RS09745 WP 014844212.1     AATGYVSPAATPNVQSI........AI....GANGVITITYTAAA.GGGTIIMTPTLQA.NGDVTWTCTGGT.LLAKYRP   131
NTDB id 1274 LPP RS09530 WP 011214168.1       AATGYVSPAATPNVASI........AI....GANGVITITYTAAA.GGGTIVMTPTLQA.NGDVTWTCTGGT.LLAKYRP   131
NTDB id 1184 GTF74 RS02335 WP 000649326.1     AG...TAAG.FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.GVVSS.SKIQLAR.DANGLWTCSTTVT..SEIAP   143
NTDB id 1167 A1552VC RS11065 WP 000649326.1   AG...TAAG.FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.GVVSS.SKIQLAR.DANGLWTCSTTVT..SEIAP   143
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    AG...TAAG.FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFDA.GVVSS.SKIQLAR.DANGLWTCSTTVT..SEIAP   143
NTDB id 1168121 AL542 RS07470 WP 040528180.1 LADIGARSN.MNALGTV......SLVAATAGDDATSAGLQFAFDA.TNSSLGGLILTTR.LENTGWDCTPSGTIPTGLDL   137
NTDB id 1292 VP RS12235 WP 005479693.1        LAQLGVDVA.ANDLGAI......DSA.....LAGGSPTLTFTFDA.DSSIDGGILTFTR.DAATGWSCLPTAP...AIAV   135
consensus                                     **   ****    ****        ***  * ***  ** ***** *  *    *** *  * **!*!***    * * *
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logo

NTDB id 1192 PAKAF RS23870 WP 024947914.1     KGCSK.....   150
NTDB id 1204 PSJM300 03935 AFN76865.1         GKCL......   141
NTDB id 1258 GCO85 RS09745 WP 014844212.1     ASCRP.....   136
NTDB id 1274 LPP RS09530 WP 011214168.1       ASCRP.....   136
NTDB id 1184 GTF74 RS02335 WP 000649326.1     KGCTAGATIN   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   KGCTAGATIN   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    KGCTAGATIN   153
NTDB id 1168121 AL542 RS07470 WP 040528180.1  EGCRQ.....   142
NTDB id 1292 VP RS12235 WP 005479693.1        DGCS......   139
consensus                                      *!*
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X non conserved

X similar

X ≥ 50% conserved