```
logo
NTDB id 669470 MTP24 RS13885 WP 279388985.1  ...MKTMTQKQQGFTLIELMIVVAIIGILAAVALPAYQNYTKKAYFSEVILATAPYKAGVQTCILTN...AIADCDTGSN   74
NTDB id 1383 A4U84 RS08530 WP 005711781.1    MKKIRLIRPLAKAFTLIELMIVIAIIAILATIAIPSYTSYTQKAALSELLSASASYKSDVEICIYNVGDKKISECSSGKN   80
NTDB id 1365 NTHI RS01955 WP 011271973.1     .MKLTTQQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNE..TTNCTGGKN   77
NTDB id 1354 HI 0299 AAC21963.1              .MKLTTLQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNE..TTSCTGGKN   77
NTDB id 1292 VP RS12235 WP 005479693.1       ..MKHSKQKKQQGFTLIELMIVVAIIGVLAAAAIPAYQNYVTRSEVTSGLATVKALITPAELHYQENGI.AAAATL.AQL   76
NTDB id 1191 AAC28468.1 484..945( )          .....MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAAL.VAG   74
NTDB id 1274 LPP RS09530 WP 011214168.1      ........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQA.....   67
NTDB id 1258 GC085 RS09745 WP 014844212.1    ........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQA.....   67
consensus                                       ***!!!!!!!!!!*!!!*!!**!*!*!* !* ** *** * ***  *  *     *      *    *
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logo
NTDB id 669470 MTP24 RS13885 WP 279388985.1  GIPATTA....TKVVAS.........VSVTDGVILVTPKAVNGVLATDTYTLTPVGATNAGDQITSWTETCANGI.....   136
NTDB id 1383 A4U84 RS08530 WP 005711781.1    GVRADKSDVPDAKYLKS.........ISVASGVITVAG...KGNIDGYGYTMTP.KFANNN...ITWATSCQGADISLFP   144
NTDB id 1365 NTHI RS01955 WP 011271973.1     GIAADITT..AKGYVKS.........VTTSNGAITVKG...DGTLANMEYILQATGNAATG...VTWTTTCKGTDASLFP   140
NTDB id 1354 HI 0299 AAC21963.1              GIAADIKT..AKGYVAS.........VITQSGGITVKG...NGTLANMEYILQAKGNAAAG...VTWTTTCKGTDASLFP   140
NTDB id 1292 VP RS12235 WP 005479693.1       GV...........DVAANDLGAIDSALAGGSPTLTFTFDAD..SSIDGGIL......TFTRDAATGWS..CLPTAPAI..   133
NTDB id 1191 AAC28468.1 484..945( )          NVIFDSANGLDGKYFAPGGVT.....VTPDTGAINIAFDAG..ANAGKTMTITPTV.IAASGQISTWT..CAPGAANGIE   144
NTDB id 1274 LPP RS09530 WP 011214168.1      ..........ATGYVSPAATPNVASIAIGANGVITITYTAA..AG.GGTIVMTPTL.QA..NGDVTWT..CTGGTLL...   126
NTDB id 1258 GC085 RS09745 WP 014844212.1    ..........ATGYVSPAATPNVQSIAIGANGVITITYTAA..AG.GGTIIMTPTL.QA..NGDVTWT..CTGGTLL...   126
consensus                                    **          **           *    * ***   *          ****    *     **!*  ! **
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logo
NTDB id 669470 MTP24 RS13885 WP 279388985.1  ..YC......   138
NTDB id 1383 A4U84 RS08530 WP 005711781.1    ANFCGQKATP   154
NTDB id 1365 NTHI RS01955 WP 011271973.1     ANFCGSVTQ.   149
NTDB id 1354 HI 0299 AAC21963.1              ANFCGSVTK.   149
NTDB id 1292 VP RS12235 WP 005479693.1       ...AVDGCS.   139
NTDB id 1191 AAC28468.1 484..945( )          TRRLPTSCQ*   153
NTDB id 1274 LPP RS09530 WP 011214168.1      AKYRPASCRP   136
NTDB id 1258 GC085 RS09745 WP 014844212.1    AKYRPASCRP   136
consensus                                    * *
```

X non conserved
X similar
X ≥50% conserved