```
logo
NTDB id 153 SP RS04730 WP 000387344.1       MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVAAVSKDSSTEKEVKKEEKEEPLEQDLITVDVKGAVKSP   85
NTDB id 1310 DR RS09495 WP 010888490.1      ......MLPHERLWQVALAAG.......................................................................   15
NTDB id 140111 TO73 RS11490 WP 003049113.1  .............MVLGYLLA.......................................................................    8
NTDB id 1034 TT RS08100 WP 024119555.1      .............MVLGYLLA.......................................................................    8
consensus                                           *   *********

logo
NTDB id 153 SP RS04730 WP 000387344.1       GIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGSG.TASSTSKEKKVNLNKASLEELKQVK  169
NTDB id 1310 DR RS09495 WP 010888490.1      ..............ALLVGGLTLGPALLPRPQLPTVTRTAL..PPVQADANAEPRTYPTTASVTPLISGRVNLNTASQEQLEALP   84
NTDB id 140111 TO73 RS11490 WP 003049113.1  ..............VALLALLSLWPKLAPRPLP............VRVEAL.......SEASFTPPAPEPVSLNGASLEELEALP   60
NTDB id 1034 TT RS08100 WP 024119555.1      ..............VALLGLLALWPKLAPAPLP............VRVEAL.......GKVAPLPQAQTPVSLNEASLEELMALP   60
consensus                                                 *****  !************  **  **   *******    *  * *  ** *  *! !! !!*!*!****

logo
NTDB id 153 SP RS04730 WP 000387344.1       GLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 1310 DR RS09495 WP 010888490.1      KVGPAMAKKIIAGR....PYHSLADLDRVKGVGEATLRTLTPLVKW.   126
NTDB id 140111 TO73 RS11490 WP 003049113.1  GIGPTLAQRIVEGR....PYGTVEDLLKVKGIGPRTLERIRPYVRP.   102
NTDB id 1034 TT RS08100 WP 024119555.1      GIGPVLARRIVEGR....PYARVEDLLKVKGIGPATLERLRPYLRP.   102
consensus                                   **!* *!**!***!     *******!**!*!*!**!**********
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X non conserved

X similar

X ≥ 50% conserved