```
logo
NTDB id 374 SMU RS06885 WP 002262930.1      ..MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRKNSHVPIIMLSAKDS   83
NTDB id 413 AAK55817.1 141..845( )          ..MKKILIVDDEKPISDIIKFNMTKEGYEVVTAFNGREALEQFEAEQPDIIILDLMLPEIDGLEVAKTIRKTSSVPILMLSAKDS   83
NTDB id 606 V4T04 RS01910 WP 003130756.1    MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMMTARDS   85
NTDB id 73507 FNL60 RS01520 WP 002262121.1  .MAKDILIIEDEKKLAKFVSLELQHEGYEVEVTYDGRSGLERALEKDFDLILLDLMLPEMDGYEVIRRLQLEKTTYVIIITARDS   84
NTDB id 471 HSISS4 RS01430 WP 002883757.1   .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMMTARDS   84
consensus                                    !*!!!**!!! ******** *!!!*******!!**!* * ****!*!*!!!!!!!*!!*!!******** *******!*!!

logo
NTDB id 374 SMU RS06885 WP 002262930.1      EFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELTHREFELL   168
NTDB id 413 AAK55817.1 141..845( )          EFDKVIGLELGADDYVTKPFSNRELQARVKALLRRSQPMPVDGQ.EADSKPQPIQIGDLEIVPDAYVAKKYGEELDLTHREFELL   167
NTDB id 606 V4T04 RS01910 WP 003130756.1    TMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVE.RAEN.....TSFRDLVIDKTNRTVHRGKKVIDLTREYDLL   164
NTDB id 73507 FNL60 RS01520 WP 002262121.1  IMDIVTGLDRGADGYIVKPFAIEELLARVRAIFRRQDIEKAKHSSKND.....TTYRDLKLDVHNRAAIRNGETIPLTKREFDLL   164
NTDB id 471 HSISS4 RS01430 WP 002883757.1   IMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKNAPAKA.....STYRDLKLDVQNRTVVRGDEAIPLTKREFDLL   164
consensus                                    *!*! !!**!!!*!**!!!***!!*!!**! *!! *   *           ****!! **  ** * * ** **!!*!!**!!

   <0
logo
NTDB id 374 SMU RS06885 WP 002262930.1      HHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD.   235
NTDB id 413 AAK55817.1 141..845( )          YHLASHTGQVITREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMRNNA*   234
NTDB id 606 V4T04 RS01910 WP 003130756.1    LTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKI.DVEGQDSYIQTVRGLGYVMRERK.   230
NTDB id 73507 FNL60 RS01520 WP 002262121.1  NALVENINQVMTREELLARVWKYDQEAETNVVDVYIRYLRGKI.DVPGKESYIQTVRGMGYIIRENN.   230
NTDB id 471 HSISS4 RS01430 WP 002883757.1   NTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKI.DVPGKESYIQTVRGMGYVIREK..   229
consensus                                    !******!*!!!*!*  !!*!*  *****!!!**!*!! !! !*******!!*!*!!*!!****
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X non conserved

X similar

X ≥ 50% conserved