```
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 48928 SGGB RS09260 WP 009854820.1   .MSKKILIIEDEKNLARFVSLELQHEGYSVVVETNGRLGLQTALDDDFDLILLDLMLPDMDGFEITRRLRLEKETSIIMMTARDS   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    TMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAVER......AENTSFRDLVIDKTNRTVHRGKKVIDLTREYDLL   164
NTDB id 48928 SGGB RS09260 WP 009854820.1   IMDIVAGLDRGADDYIVKPFAIEELLARVRAVFRRQDVESKREQD....SQGKEGLLGLRLNPQNRSAVRGDDEISLTKREYDLL   165
NTDB id 471 HSISS4 RS01430 WP 002883757.1   IMDIVAGLDRGADDYIVKPFAIEELLARTRATFRRQDIEAAKNAP....AK.ASTYRDLKLDVQNRTVVRGDEAIPLTKREFDLL   164
consensus                                   *!*!*!!**!!!!!**!!!***!!*!!**! *!! *                 * *! * * **** ** ****!!*!!**!!

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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 48928 SGGB RS09260 WP 009854820.1   SVLLSNVNRVMTREELLASVWRYDTDIETNVVDVYIRYLRGKV.DIPGKESYIQTVRGMGYIIREK..   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