```
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 27323 GALLO RS09400 WP 009854820.1  .MSKKILIIEDEKNLARFVSLELQHEGYSVVVETNGRLGLQTALDDDFDLILLDLMLPDMDGFEITRRLRLEKETSIIMMTARDS   84
NTDB id 471 HSISS4 RS01430 WP 002883757.1   .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMMTARDS   84
consensus                                    *!*!!!**!!!********** *!!!***** *!!**!* * ****!*!*!!!!!!**!!*!***********!*!**!*!!

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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......AENTSFRDLVIDKTNRTVHRGKKVIDLTRREYDLL   164
NTDB id 27323 GALLO RS09400 WP 009854820.1  IMDIVAGLDRGADDYIVKPFAIEELLARVRAVFRRQDVESKREQD....SQGKEGLLGLRLNPQNRSAVRGDDEISLTKREYDLL   165
NTDB id 471 HSISS4 RS01430 WP 002883757.1   IMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKNAP....AK.ASTYRDLKLDVQNRTVVRGDEAIPLTKREFDLL   164
consensus                                    *!*!*!!**!!!!!**!!!***!!*!!**! *!! *                * *! * * **** ** ****!!*!!**!!

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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 27323 GALLO RS09400 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