```
logo
NTDB id 92857 ADU72 RS04850 WP 046872131.1  ..MQKILVVDDEKPISDIVKFNLTKEGYEVITAYDGEEALQKVEDESPDLILLDLMLPKIDGLEVARRVRAKHSIPIIMVTAKDS   83
NTDB id 413 AAK55817.1 141..845( )          ..MKKILIVDDEKPISDIIKFNMTKEGYEVVTAFNGREALEQFEAEQPDIIILDLMLPEIDGLEVAKTIRKTSSVPILMLSAKDS   83
NTDB id 374 SMU RS06885 WP 002262930.1      ..MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRKNSHVPIIMLSAKDS   83
NTDB id 606 V4T04 RS01910 WP 003130756.1    MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMMTARDS   85
NTDB id 471 HSISS4 RS01430 WP 002883757.1   .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMMTARDS   84
consensus                                     ***!!***!!!********** *!!!*******!***!*  *** *!*!*!!!!!**!!*!!******   **!*!**!*!!

logo
NTDB id 92857 ADU72 RS04850 WP 046872131.1  ELDKVVGLELGADDYVTKPFSNRELVARVKANLRRQDQLVSAEQEN.....NDIKVGDLDINPDAYAVSRHGEQLDLTHREFELL   163
NTDB id 413 AAK55817.1 141..845( )          EFDKVIGLELGADDYVTKPFSNRELQARVKALLRRSQPMPVDGQ.EADSKPQPIQIGDLEIVPDAYVAKKYGEELDLTHREFELL   167
NTDB id 374 SMU RS06885 WP 002262930.1      EFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELTHREFELL   168
NTDB id 606 V4T04 RS01910 WP 003130756.1    TMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV.ERAEN.....TSFRDLVIDKTNRTVHRGKKVIDLTRREYDLL   164
NTDB id 471 HSISS4 RS01430 WP 002883757.1   IMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKNAPAKA.....STYRDLKLDVQNRTVVRGDEAIPLTKREFDLL   164
consensus                                   **!*!*!!**!!!!!**!!!***!!*!!**! *!! * *             * **!! * **** * * ** **!!*!!**!!

   <0
logo
NTDB id 92857 ADU72 RS04850 WP 046872131.1  YYLAQHMGQVMTREHLLQTVWGYDYFGDVRTVDVTVRRLREKIEENPSNPVYLVTRRGVGYYLRNPENEK   233
NTDB id 413 AAK55817.1 141..845( )          YHLASHTGQVITREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMRNNA*..   234
NTDB id 374 SMU RS06885 WP 002262930.1      HHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD...   235
NTDB id 606 V4T04 RS01910 WP 003130756.1    LTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKID.VEGQDSYIQTVRGLGYVMRERK...   230
NTDB id 471 HSISS4 RS01430 WP 002883757.1   NTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKID.VPGKESYIQTVRGMGYVIREK....   229
consensus                                     !******!*!!!*!* *!!*!********!!!**!*!!*!!** **** !**!*!!*!!***
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X non conserved

X similar

X ≥ 50% conserved