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logo
NTDB id 545 H702 RS02385 WP 052071091.1      MIDIYVLEDNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLTNL.VEESFPQIFFLDIDIHGDNTKGIEVAKAIREKSRTAT   84
NTDB id 542 H702 RS02345 WP 039696227.1      MLDIFILEDNITQQFRIERQIETIMERNHWEYQRLEAFAASKDIIKKADGQGNHQVFFLDLEIKGDEKQGLDVARSIREKDPTAI   85
NTDB id 38281 SP670 RS02980 WP 001219134.1   .MRIFVLEDDFSQQTRIETTIEKLLKEHHITLSSFEVFGKPDQLLAEVHEKGAHQLFFLDIEIRNEEMKGLEVARKIRDRDPYAL   84
NTDB id 55 SMU RS08710 WP 002262114.1        MISIFVLEDDFLQQGRLETTIAAIMKEKNWSYKELTIFGKPQQLIDAIPEKGNHQIFFLDIEIKKEEKKGLEVANQIRQHNPSAV   85
consensus                                    ***!**!!! **!!*!*!**!* ** ******  **!*******  * *****!*!!!!**!******!**!!* !!*****!*

logo
IVFVTTHSEFMLLTYRYQVSALDFIDKTLNDEEFSSRIETCLYYANEYQTNSCSEESFYFESSKTQIRVPFKELLYEETSITYH
NTDB id 545 H702 RS02385 WP 052071091.1      IVFVTTHSEFMLLTYRAQVSALDFIDKTDKDDAIKDNLKKCLQKVFEDQ.GFNIPKEIFVFENNKTKIRIPLEDILYFETA.ESH   167
NTDB id 542 H702 RS02345 WP 039696227.1      IVFVTTHSEFMLLTYQSLVGAIDFIDKNMNDEAFGGRIELCLKEALKYQ.TGSFGENSYLFETAKARIRVAYDDILYFETSPTVH   169
NTDB id 38281 SP670 RS02980 WP 001219134.1   IVFVTTHSEFMPLSFRYQVSALDYIDKALSAEEFESRIETALLYANGQDS.KSLAEDCFYFKSKFAQFQYPFKEVYYLETSTRPH   168
NTDB id 55 SMU RS08710 WP 002262114.1        IVFVTTHSEFMPLTFQYQVSALDFIDKSLNPEEFSHRIESALYYAMENSQKNGQSEELFIFHSSETQFQVPFAEILYFETSSTAH   170
consensus                                    !!!!!!!!!!!*!** **!*!*!*!!!***** *  *****! ******   ********!** * *  *******!*!!*** !

logo
RVILYTKTGRIEFYGSITEVYKSDKRLLQCHRSFLINPNNVVRLDKKERLAYFRNGKSCLYSRYKVKKLRAYMRNLR
NTDB id 545 H702 RS02385 WP 052071091.1      RLKLITKTGQRHFYGSIKEVMQANQKLFRCHKSYLINLNNIVRLDKKEGLVYFDDDKACYVSRKYIKDLRSKMENLK...   244
NTDB id 542 H702 RS02345 WP 039696227.1      RVILHTKTGQTEFYGTIAEVAKSDKRLLKCHRSFVINVNNVTEFDKSTRTAYFENGSYCQVSRDKVKKLMAEMR......   243
NTDB id 38281 SP670 RS02980 WP 001219134.1   RVILYTKTDRLEFTASLEEVLKQEPRFLQCHRSFLINPANVVHLDKKEKLLFFPNGGSCLIARYKVREVSEAINNLH...   245
NTDB id 55 SMU RS08710 WP 002262114.1        KLCLYTYDERIEFYGSMTDIVKMDKRLFQCHRSFIVNPANITRIDRKKRLAYFRNNKSCLISRTKLTKLRAVIADQRRAK   250
consensus                                    ***!*!*** **!**** **** **** *!!*!***!* !* **!*******!*****!***! ******* * ***
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X non conserved

X similar

X ≥ 50% conserved