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logo

NTDB id 627 AAM18091.1 1453..2205( )          .MKVLILEDVIQHQVRLERTLDEISKELNIPI.SYKTTGKEREFKEYIENDEVNQLYFLDIDIHGVEKKGFEVAQFIRRH   78
NTDB id 545 H702 RS02385 WP 052071091.1       MIDIYVLEDNVNQQFRIEKMIGDLLETNQWETRHFDLFSEPDCLTNL.VEESFPQIFFLDIDIHGDNTKGIEVAKAIREK   79
NTDB id 542 H702 RS02345 WP 039696227.1       MLDIFILEDNITQQFRIERQIETIMERNHWEYQRLEAFAASKDIIKKADGQGNHQVFFLDLEIKGDEKQGLDVARSIREK   80
NTDB id 105870 AABA08 RS02340 WP 001219133.1  .MRIFVLEDDFSQQTRIETTIEKLLKEHHITLNSFEVFGKPDQLLAEVHEKGAHQLFFLDIEIRNEEMKGLEVARKIRDR   79
NTDB id 55 SMU RS08710 WP 002262114.1         MISIFVLEDDFLQQGRLETTIAAIMKEKNWSYKELTIFGKPQQLIDAIPEKGNHQIFFLDIEIKKEEKKGLEVANQIRQH   80
consensus                                     ** ***!!! * *! !*!*** **** **    *******  **  * * * *!**!!!**!******!**!!* !!**

logo

NTDB id 627 AAM18091.1 1453..2205( )          NPYAIIVFITSRSEFATLTYKYQVSALDFVDKDINDELFKKRIEQSILYTKSMLLENKDVVDYFDYNYKGNDLKIPYHDI  158
NTDB id 545 H702 RS02385 WP 052071091.1       SRTATIVFVTTHSEFMLLTYRAQVSALDFIDKTDKDDAIKDNLKKCLQKVFEDQ.GFNIPKEIFVFENNKTKIRIPLEDI  158
NTDB id 542 H702 RS02345 WP 039696227.1       DPTAIIVFVTTHSEFMLLTYQSLVGAIDFIDKNMNDEAFGGRIELCLKEALKYQ.TGSFGENSYLFETAKARIRVAYDDI  159
NTDB id 105870 AABA08 RS02340 WP 001219133.1  DPYALIVFVTTHSEFMPLSFRYQVSALDYIDKALSAEEFESRIETALLYANSQDS.KSLAEDCFYFKSKFAQFQYPFKEV  158
NTDB id 55 SMU RS08710 WP 002262114.1         NPSAVIVFVTTHSEFMPLTFQYQVSALDFIDKSLNPEEFSHRIESALYYAMENSQKNGQSEELFIFHSSETQFQVPFAEI  160
consensus                                      **!*!!!*!**!!!* !******!*!*!**!! **** *  ***  * **          ** *** *    ***** **

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NTDB id 627 AAM18091.1 1453..2205( )          LYIETTGVSHKLRIIGKNFAKEFYGTMADIQEKDKDTQRFYSAHKSFLVNVGNIREIDRKNLEVVFYEDHRCPITRLKVR  238
NTDB id 545 H702 RS02385 WP 052071091.1       LYFETA.ESHRLKLITKTGQRHFYGSIKEVMQA...NQKLFRCHKSYLINLNNIVRLDKKEGLVYFDDDKACYVSRKYIK  234
NTDB id 542 H702 RS02345 WP 039696227.1       LYFETSPTVHRVILHTKTGQTEFYGTIAEVAKS...DKRLLKCHRSFVINVNNVTEFDKSTRTAYFENGSYCQVSRDKVK  236
NTDB id 105870 AABA08 RS02340 WP 001219133.1  YYLETSPRPHRVILYTKTDRLEFTASLEEVFKQ...EPRLLQCHRSFLINPANVVHLDKKEKLLFFPNGGSCLIARYKVR  235
NTDB id 55 SMU RS08710 WP 002262114.1         LYFETSSTAHKLCLYTYDERIEFYGSMTDIVKM...DKRLFQCHRSFIVNPANITRIDRKKRLAYFRNNKSCLISRTKLT  237
consensus                                     *!*!!*   !** * ***   *!**** ** *    * *** *!*!***!**!*  *!** ****! *   ! **! ***

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NTDB id 627 AAM18091.1 1453..2205( )          KLRDILEKKSKK*   250
NTDB id 545 H702 RS02385 WP 052071091.1       DLRSKMENLK...   244
NTDB id 542 H702 RS02345 WP 039696227.1       KLMAEMR......   243
NTDB id 105870 AABA08 RS02340 WP 001219133.1  EVSEAINKLH...   245
NTDB id 55 SMU RS08710 WP 002262114.1         KLRAVIADQRRAK   250
consensus                                     ***  *
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X non conserved

X similar

X ≥50% conserved