```
logo
NTDB id 1139100 DQM54 RS02480 WP 111723335.1  .MAKRILLVENEKNLARFIMLELQKESYQVDLAETGQEGLALAKENDYDLLLLNNQFADGTAFAFAQQLSLLKPVAVIIV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1     .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDT.YIMM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDT.PIIM   79
consensus                                      **!*!!**!*!!!!!!!**!!!**!*! *******!**!!*!! **!*!!*!!********!**!*****!  *!**  !**

logo
NTDB id 1139100 DQM54 RS02480 WP 111723335.1  MVSREEIADSAEEIQQFAVSYMVKPFIISDLVDKISAIFRGRDFIDQKCSQLKVPTAYRNLRVDLKNHTVYRGDEIIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKN.APAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV..ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                     !**!****! ***** *!**!**!!!*!**!*****! !!* *   **    *  ***!*!**!* !*!! !!***! !!

logo
NTDB id 1139100 DQM54 RS02480 WP 111723335.1  RREYDLLVTLMSS.NKVMSREQLLDRVWKYESASETNIVDVYIRYLRSKIDLPGQESYIKTVRGIGYSMQG..   229
NTDB id 471 HSISS4 RS01430 WP 002883757.1     KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1      RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                     *!!*!!!*!!*****  !**!!  !*  *!!*!***!!!*!!!!!!!!!  !!!**!**!!!*!!!!*!!*****
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X non conserved

X similar

X $\geq$ 50% conserved