```
logo
NTDB id 1142329 DQN71 RS08420 WP 015605580.1  .MAKRILLVENEKNLARFVTLELQKEAYLLDVVETGSQGLALARENDYDLLLLNYDLQDMTASEFAKQLSLLKPASVIIV   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 1142329 DQN71 RS08420 WP 015605580.1  LASREEIKEHVDEIQRFAVSYTIKPFIISDLVEKVSDIFRGRDFIDQHCSKLKTPTAYRNLRVDLKNHIVYRGEEVIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                     ***!***** !******!**!*!!!*!**!****** !!* *   *     *  ***!*!**!* !**! !!***! !!

logo
NTDB id 1142329 DQN71 RS08420 WP 015605580.1  RREYDLLVTLMSNQ.KVMSREQLLERVWKYEGATETNVVDVYIRYLRSKIDIAGQTSYIKTVRGIGYAMQE..   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