```
logo
NTDB id 1139388 DQM69 RS02755 WP 172456394.1  .MAKRILLIEGEKSLSRLVSLKLQERGYQTVVSQDGKSGLKTAMEKDFELIFIDLKLPDQDGLELMQQLRGHKSVPVVLM   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1     .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMM   79
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMM   80
consensus                                      **!*!!**!*!!*!*!*!!!*!***!! **** **!**!!* !**!!**!!**!!*!!**!!*!* **!* *!******!

logo
NTDB id 1139388 DQM69 RS02755 WP 172456394.1  VGREDVATLLPKTDNLADDYLIKPFAVEELLDKVDETFQRKDFKRRKQSTSQTTAYQDLILDEANRSVIRKGESIALTKR   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     TARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKNAPAKASTYRDLKLDVQNRTVVRGDEAIPLTKR   159
NTDB id 606 V4T04 RS01910 WP 003130756.1      TARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV.ERAENTSFRDLVIDKTNRTVHRGKKVIDLTRR   159
consensus                                     **!**********! *!!!!*!!!!*!!!!******!*! *   ***  *  ****!!**!  !!*!*!* * ! !!*!

logo
NTDB id 1139388 DQM69 RS02755 WP 172456394.1  EFALLSTLLRHINMVMTREELLDYVWSYSDGTIETNVVDVYVRYLRGKIDIPGQESYIQTVRGLGYVIRER.   230
NTDB id 471 HSISS4 RS01430 WP 002883757.1     EFDLLNTLLSNMNQVMTREELLLQVWKYDDA.IETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1      EYDLLLTLMQNVGDVVTREHLVSQVWGYEEG.TETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                     !**!! !!* *** !*!!!*!* *!! !*** *!!!!!!!!!*!!!!*!!!**!**!!!!!!!!*!!!*!!*
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X non conserved

X similar

X $\geq$ 50% conserved