```
logo
NTDB id 293154 DK181 RS02865 WP 002959089.1  .MAKRILLIEGEKSLSRLVSLKLQERGYQTVVSQDGKSGLKTAMEKDFELIFIDLKLPDQDGLELMQELRGHKSVPVVLM   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 293154 DK181 RS02865 WP 002959089.1  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 293154 DK181 RS02865 WP 002959089.1  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