```
logo
NTDB id 1067662 ACFSN5 RS02610 WP 165329206.1  .MAKRILLIEGEKSLSRLVSLKLQEKGYQAFVSQNGRSGLKMALERDFELIFIDFKLSDRDGLKLMQDIKEVKKVPIVLM   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 1067662 ACFSN5 RS02610 WP 165329206.1  VNREDAEN.LPKTDGLTDDYLIKPFAVEELLDKVDEIFQRKGLKRKP.PKSNQTAYQDLIMDEASRSVIRRGEAIALTKR   157
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 1067662 ACFSN5 RS02610 WP 165329206.1  EFALLSTLLRHINTVMTREELLDYVWSYSDGTVETNVVDVYVRYLRGKIDIPGHESYIQTVRGLGYVIRER.   228
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