```
logo
NTDB id 293567 DLJ51 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 293567 DLJ51 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 293567 DLJ51 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                                    !**!! !!* *** !*!!!*!* *!! !*** *!!!!!!!!!*!!!!*!!!**!*!!!!!!!!!*!!!*!!*
```

X non conserved

X similar

X $\geq$ 50% conserved