```
logo
NTDB id 247936 CO687 RS00630 WP 012000010.1  .MAKRILLVENEKNLARFIMLELQKESYQVDLAETGQEGLALAKENDYDLLLLNNQFADGTAFAFAQQLSLLKPAAVIIV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1    .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKD.TYIMM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKD.TPIIM   79
consensus                                     **!*!!**!*!!!!!!!**!!!**!*! *******!**!!**! **!*!!*!!******!**!*****!  *!* * !**

logo
NTDB id 247936 CO687 RS00630 WP 012000010.1  MVSREEIADSAEEIQQFAVSYMVKPFIISDLVDKISAIFRGRDFIDQKCSQLKVPTAYRNLRVDLKNHTVYRGEEIIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1    MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAKN.APAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV..ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                    !**!****! ***** *!**!**!!!*!**!*****! !!* *   **     *  ***!*!**!* !*!! !!***! !!

logo
NTDB id 247936 CO687 RS00630 WP 012000010.1  RREYDLLVTLMSS.NKVMSREQLLDRVWKYESASETNIVDVYIRYLRSKIDLPGQESYIKTVRGIGYSMQG..   229
NTDB id 471 HSISS4 RS01430 WP 002883757.1    KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1     RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                    *!!*!!!*!!***** !**!! !* *!!*!***!!!*!!!!!!!!! !!!**!**!!!*!!!!*!!*****
```

X non conserved

X similar

X $\geq$ 50% conserved