```
logo
NTDB id 61868 SIR RS11515 WP 021002458.1   .MGKRILLVENEKKIAQFISLELQNEGYRVDLFETGKDALTSARANQYDLFLLNFMLEDMTGTEFAEQLSLIKPASVIIVLDNRD   84
NTDB id 471 HSISS4 RS01430 WP 002883757.1  .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDT.YIMMMTARD   83
NTDB id 606 V4T04 RS01910 WP 003130756.1   MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDT.PIIMMTARD   84
consensus                                   **!*!!**!*!!**!*!*!!!!**!!! *** **!***!* !  ***!!*!!**!!****!*!****!  *!**  !*****!!

logo
NTDB id 61868 SIR RS11515 WP 021002458.1   IILKYTEEIQRFAVSYMIEPFIVMDLVGKISAIFRGRDFIDHHCSQMKVSTSYRNLRIDVQNHTVYRGNEIISLTRREYDLLATL   169
NTDB id 471 HSISS4 RS01430 WP 002883757.1  SIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDI.EAAKNAPAKASTYRDLKLDVQNRTVVRGDEAIPLTKREFDLLNTL   167
NTDB id 606 V4T04 RS01910 WP 003130756.1   STMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEH.GHAV.ERAENTSFRDLVIDKTNRTVHRGKKVIDLTRREYDLLLTL   167
consensus                                  **** *******!**!***!!****!*****! !!* *  ***    *  ***!*!**!**!*!! !! **! !!*!!*!!! !!

logo
NTDB id 61868 SIR RS11515 WP 021002458.1   MGS.HQALSREQLLERVWKYEGAVETNVVDVYIRYLRSKLDTAGQKSYIKTVRGVGYAMQE..   229
NTDB id 471 HSISS4 RS01430 WP 002883757.1  LSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1   MQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                  * ** ****!! !* *!!*!***!!!!!!!!!!!!! !*!* !**!!!*!!!!*!!***!*
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X non conserved

X similar

X $\geq$ 50% conserved