```
logo
NTDB id 1120897 EL115 RS06910 WP 022526966.1  .MAKRILLVEKEKKLAQFIGLELQKEGYRVDLFETGKDALASARANQYDLFLLNFMLDDMTGTEFAAQLSVIKPASVIIV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1     .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDT.YIMM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDT.PIIM   79
consensus                                      **!*!!**!*!!*!!*!**!!!**!!! *** **!***!* !  ***!!*!!**!!****!*!****!  *!**  !**

logo
NTDB id 1120897 EL115 RS06910 WP 022526966.1  LDNRDNILQYTEEIQRFAVAYMVEPFIVTDLVEKISAIFRGRDFIDQHCSQMKVPTSYRNLRIDVQNHTVYRGEEVIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                     ***!!**** *******!**!***!!****!*****! !!* *   *     *  ***!*!**!**!*!! !!***! !!

logo
NTDB id 1120897 EL115 RS06910 WP 022526966.1  RREYDLLATLMGS.HQALSREQLLERVWKYEGAVETNVVDVYIRYLRSKIDIVGQKSYIKTVRGVGYVMQE..   229
NTDB id 471 HSISS4 RS01430 WP 002883757.1     KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1      RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                     *!!*!!! !!* ** ****!! !* *!!*!***!!!!!!!!!!!!! !!!* !**!!!*!!!!*!!!**!*
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X non conserved

X similar

X $\geq$ 50% conserved