```
logo
NTDB id 156788 SanJ4206 RS07220 WP 003029931.1  .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 156788 SanJ4206 RS07220 WP 003029931.1  LDHRDNILQYTEEIQRFAVAYMVEPFIVTDLVEKISAIFRGRDFIDQHCSQMKVPTSYRNLRIDVQNHTVYRGEEVIALT   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 156788 SanJ4206 RS07220 WP 003029931.1  RREYDLLATLMGS.HQALSREQLLERVWKYEGAVETNVVDVYIRYLRSKIDIVGQKSYIKTVRGVGYAMQE..   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 ≥ 50% conserved