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logo
NTDB id 1081722 ACK2I7 RS06160 WP 057489790.1  .MAKRILLVEKEKKLAQFIGLELQKEGYRVDLFETGKDALASARANQYDLFLLNFMLDDMTGTEFAAQLSVIKLASVIIV   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 1081722 ACK2I7 RS06160 WP 057489790.1  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 1081722 ACK2I7 RS06160 WP 057489790.1  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 ≥ 50% conserved