```
logo
NTDB id 1155622 QNH80 RS05890 WP 012027645.1  .MAKKILIAGKERNLSHFVSMELQKKDYLVDYASTGKEAMSLAHETDFDLILMSFQLSDMSSKELATELLAIKPATVMIV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1     .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMM   79
NTDB id 606 V4T04 RS01910 WP 003130756.1      MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMM   80
consensus                                      **!*!!!***!*!!**!!!*!!****! ** ***!*****!  **!*!!!!****!******!****!   *!**  ****

logo
NTDB id 1155622 QNH80 RS05890 WP 012027645.1  VVEPTEVSQYGEEVLSYAVSYVVKPFVISDLVEQISAIFRGRDFIDNNCKQVHMHAAYRDLKVDFQNRTVTRGDELINLT  159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     TARDSIM.DIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT  157
NTDB id 606 V4T04 RS01910 WP 003130756.1      TARDSTM.DRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT  157
consensus                                     ***** * * ***** *!**!**!!!*!**!*****! !!* *   *     *  ***!!!*!* *!!!! !!***! !!

logo
NTDB id 1155622 QNH80 RS05890 WP 012027645.1  RREYDLLATLMNS.PEPVSREQLLERVWKYEAASETNVVDVYIRYLRGKLDLPHQDSYIKTVRGVGYAMRD..  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  ≥ 50% conserved
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