```
logo
NTDB id 1139888 DQM93 RS01835 WP 000518011.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDLVEEGQKALSMALQTDYDLILLNVNLGDMMAQDFAEKLSRTKPASVIMI   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 1139888 DQM93 RS01835 WP 000518011.1  LDHWEDLQEELEVVQRFAVSYIYKPVLIENLVARISAIFRGRDFIDQHCSLMKVPRTYRNLRIDVEHHTVYRGEEMIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1     T.ARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAK.NAPAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1      T.ARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV..ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                     * ******* *******!**!! !!**!!*!*!!**! !!* *   *     *  ***!*!**!* **!! !!***! !!

logo
NTDB id 1139888 DQM93 RS01835 WP 000518011.1  RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTMQE..   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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