```
logo
NTDB id 1141807 DQK99 RS02140 WP 000518008.1  .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 1141807 DQK99 RS02140 WP 000518008.1  LDHWEDLQEELEVVQRFAVSYIYKPVFIENLVARISAIFRGRDFIDQHCSLMKVPRTYRNLRIDVEHHTVYRGEEMIALT  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 1141807 DQK99 RS02140 WP 000518008.1  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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