```
logo
NTDB id 405055 GOM46 RS02115 WP 235083846.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDQVEEGQKALSMALQTDYDLILLNAHLGDMTAQDFADKLSRTKPASVIMI   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1    .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMM.   78
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIM.   79
consensus                                     **!*!!**!*!*!!!*!*!!!!**!*! **  **!* *!**!* *!*!!!!!***!***********! **!** **!

logo
NTDB id 405055 GOM46 RS02115 WP 235083846.1  LDHREELQGQLETIQRFAVSYIYKPVIIDDLVNRIAAIFRGRDFIDQHCSQMKVPTSYRNLRMDLENHTVYRGDEMIALT   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 405055 GOM46 RS02115 WP 235083846.1  RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNVVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTMQE..   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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