```
logo
NTDB id 169989 AXE83 RS02970 WP 060955366.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDQVEEGQKALSMALQTDYDLILLNAHLGDMTAQDFADKLSRTKPAAVIMV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1    .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYI.MM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPI.IM   79
consensus                                     **!*!!**!*!*!!!*!*!!!!**!*! **  **!* *!**!* *!*!!!!!***!*********** **!** * **

logo
NTDB id 169989 AXE83 RS02970 WP 060955366.1  LDHREELQDQLETIQRFAVSYIYKPIIIDDLVSRISAIFRGRDFIDQHCSQMKVPTAYRNLRMDVEHHTVYRGDEMIALT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1    MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAAK.NAPAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHAV..ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                    ***!*****! *******!**!! !!**!**!**!**! !!* *   *     *  ***!*!**!* **!! !!***! !!

logo
NTDB id 169989 AXE83 RS02970 WP 060955366.1  RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKIDVKGQKSYIKTVRGVGYTMQE..   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 $\geq$ 50% conserved