```
logo
NTDB id 241572 CHF41 RS04245 WP 119876135.1  .MVKKILIAGRERNLSHFVSMELQKKDYLVDYTSTGKEALQLAHETDFDLILMSFRLSDMSSQELAAELVNIKPTTVMIV   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 241572 CHF41 RS04245 WP 119876135.1  VVDSEDAAIYGAEIQRYAVSYIIKPFVISDLVDQVTSIFRGRDYIDEHCKQVKLQAAYRDLKIDFQNRTVTRGPELINLT  159
NTDB id 471 HSISS4 RS01430 WP 002883757.1    TARDS.IMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT  157
NTDB id 606 V4T04 RS01910 WP 003130756.1     TARDS.TMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT  157
consensus                                    *****  ** *!*****!**!!*!!!*!**!****** !!* *   *     * ****!!!**! *!!!! !! **! !!

logo
NTDB id 241572 CHF41 RS04245 WP 119876135.1  RREYDLLATLMNS.QETLTREQLLDRVWKYEATAETNVVDVYIRYLRGKLDLPGQESYIKTVRGVGYAMREE.  230
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