```
logo
NTDB id 291206 DEH83 RS06775 WP 125442754.1  .MGKRILLVENEKKLAQFISLELQNEGYRVDLFETGKDALTSARANQYDLFLLNFMLEDMTGTEFAEQLSLIKPASVIIV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1    .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDT.YIMM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDT.PIIM   79
consensus                                     **!*!!**!*!!*!!*!*!!!!**!!! *** **!***!* !  ***!!*!!**!!****!*!****!  *!**  !**

logo
NTDB id 291206 DEH83 RS06775 WP 125442754.1  LDNRDIILKYTEEIQRFAVSYMIEPFIVMDLVEKISAIFRGRDFIDHHCSQMKVSTSYRNLRIDVQNHTVYRGNEIISLT  159
NTDB id 471 HSISS4 RS01430 WP 002883757.1    MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDI.EAAKNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT  157
NTDB id 606 V4T04 RS01910 WP 003130756.1     MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEH.GHAV.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT  157
consensus                                    ***!!**** *******!**!***!!****!*****! !!* *  ***    *  ***!*!**!**!*!! !! **! !!

logo
NTDB id 291206 DEH83 RS06775 WP 125442754.1  RREYDLLATLMGS.HQALSREQLLERVWKYEGAVETNVVDVYIRYLRSKLDIAGQKSYIKTVRGVGYAMQE..  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