```
logo
NTDB id 298496 DK877 RS08560 WP 002942700.1  .MAKKILIAGKERNLSHFVSMELQKKDYLVDYASTGKEAMSLAHETDFDLILMSFQLSDMSSKELAAELLAIKPATVMIV   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 298496 DK877 RS08560 WP 002942700.1  VVEPIEVSQYGEEVLSYAVSYVVKPFVISDLVEQISAIFRGRDFIDNNCKQVHMHAAYRDLKVDFQNRTVTRGDELINLT   159
NTDB id 471 HSISS4 RS01430 WP 002883757.1    TARDSIM.DIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT   157
NTDB id 606 V4T04 RS01910 WP 003130756.1     TARDSTM.DRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT   157
consensus                                    ***** * * ***** *!**!**!!!*!**!*****! !!* *   *     *  ***!!!**! *!!!! !!***! !!

logo
NTDB id 298496 DK877 RS08560 WP 002942700.1  RREYDLLATLMNS.PEPVSREQLLERVWKYEAASETNVVDVYIRYLRGKLDLPHQDSYIKTVRGVGYAMRD..   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