```
logo
NTDB id 1059199 ACFKHJ RS02695 WP 239604567.1  .MAKKILIAGKERNLSHFVSMELQKKEYIVDYASTGKEALSLAHETDFDLVLMSFQLSDMSSKELAKELLAIKPATVMIV   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 1059199 ACFKHJ RS02695 WP 239604567.1  VVEPTEVADHGEDVLTYAVFYVVKPFVISDLVEQISDIFRGRDFIETNCKQVHLHAAYRDLKVDFQNRTVTRGDELINLT  159
NTDB id 471 HSISS4 RS01430 WP 002883757.1      TAR.DSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEA.AKNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT  157
NTDB id 606 V4T04 RS01910 WP 003130756.1       TAR.DSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGH.AV.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT  157
consensus                                      *** ****! ***** *!**!**!!!*!**!****** !!* *    *    *  ***!!!**! *!!!! !!***! !!

logo
NTDB id 1059199 ACFKHJ RS02695 WP 239604567.1  RREYDLLATLMNS.PEPVSREQLLERVWKYEAASETNVVDVYIRYLRGKLDVPGQASYIRTVRGFGYTMRD..  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