```
logo
NTDB id 91251 L6411 RS07745 WP 024532842.1  .MAKKILIAGKERNLSHFVSMELQKKDYLVDYVSTGKEALSLAHETDFDLILMSFQLSDMSSKELAKELLAMKPATVMIVAMDPA   84
NTDB id 471 HSISS4 RS01430 WP 002883757.1   .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYIMMMTARDS   84
NTDB id 606 V4T04 RS01910 WP 003130756.1    MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPIIMMTARDS   85
consensus                                    **!*!!!***!*!!**!!!*!!****! **  **!***!**! **!*!!!!****!******!****!  *!** *********

logo
NTDB id 91251 L6411 RS07745 WP 024532842.1  DVSQHGEEVLAYAVSYVVKPFVISDLVEQISAIFRGRDFIDNHCKQVHLHAAYRDLKVDFQNRTVTRGDELINLTRREYDLLATL  169
NTDB id 471 HSISS4 RS01430 WP 002883757.1   IM.DIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLTKREFDLLNTL  167
NTDB id 606 V4T04 RS01910 WP 003130756.1    TM.DRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLTRREYDLLLTL  167
consensus                                    * * ***** *!**!**!!!*!**!*****! !!* *   *     *  ***!!!**! *!!!! !!***! !!*!!*!!! !!

logo
NTDB id 91251 L6411 RS07745 WP 024532842.1  MNS.PEAVSREQLLERVWKYEAASETNVVDVYIRYLRGKLDLPNQESYIKTVRGVGYAMRD..   229
NTDB id 471 HSISS4 RS01430 WP 002883757.1   LSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.   229
NTDB id 606 V4T04 RS01910 WP 003130756.1    MQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK   230
consensus                                   **** ****!! !* *!!*!***!!!!!!!!!!!!!!*!*!*****!!!*!!!!*!!**!**
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X non conserved

X similar

X ≥ 50% conserved