```
logo
NTDB id 1170227 ACLV7Z RS09005 WP 218774094.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDQVEEGQKALSMALQTDYDLILLNARLGDMTAQDFADKLSRTKPSSVIMV   79
NTDB id 471 HSISS4 RS01430 WP 002883757.1      .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYI.MM   78
NTDB id 606 V4T04 RS01910 WP 003130756.1       MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPI.IM   79
consensus                                       **!*!!*!*!*!!!*!*!!!!**!*! **  **!* *!**!* *!*!!!!***!*********** **!** * **

logo
NTDB id 1170227 ACLV7Z RS09005 WP 218774094.1  LDHREELQDQLETIQRFAVSYIYKPVIIDNLVARISAIFRGRDFIDQHCSQMKVPTSYRNLRMDVEHHTVYRGEEMIALT  159
NTDB id 471 HSISS4 RS01430 WP 002883757.1      MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT  157
NTDB id 606 V4T04 RS01910 WP 003130756.1       MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT  157
consensus                                      ***!****** *******!**!! !!**!**!*!!**! !!* *   *     *  ***!*!**!* **!! !!***! !!

logo
NTDB id 1170227 ACLV7Z RS09005 WP 218774094.1  RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTMQE..  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 ≥ 50% conserved