```
logo
NTDB id 291432 DG474 RS08000 WP 255778025.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDLVEEGQKALSMALQMDYDLILLNARLGDMTAQDFAERLSRTKPASVIMV   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 291432 DG474 RS08000 WP 255778025.1  LDHREELQDQIETIQRFAVSYIYKPVIIDQLVARISAIFRGRDFIDQHCSQMKVPTSYRNLRMDVEHHTVYRGEEMIALT  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 291432 DG474 RS08000 WP 255778025.1  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 $\geq$ 50% conserved