```
logo
NTDB id 668119 MP387 RS07565 WP 242746013.1  .MVKRILLLEKERNLAHFLSLELQKEQYRVDLVEEGQKALSMALQTDYDLILLNARLGDMTAQDFAERLSRTKPASVIMV   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 668119 MP387 RS07565 WP 242746013.1  LDHREELQDQIETIQRFAVSYIYKPVIIDNLVARISAIFRGRDFIDQHCSQMKVPTSYRNLRMDVEHHTVYRGEEMIALT  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 668119 MP387 RS07565 WP 242746013.1  RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTMQE..  229
NTDB id 471 HSISS4 RS01430 WP 002883757.1    KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK.  229
NTDB id 606 V4T04 RS01910 WP 003130756.1     RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK  230
consensus                                    *!!*!!! !!* **  !*!!! !* *!!*!*****!!*!!!!!!!!! !*!! !**!!!*!!!!*!!***!*
```

X non conserved

X similar

X $\geq$ 50% conserved