```
logo
NTDB id 284723 SK637 RS08035 WP 033689345.1  .MGKRILLLEKERNLAHFLSLELQKEQYRVDQVEEGQKALSMALQTDYDLILLNARLGDMTAQDFADKLSRTKPASVIMV   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 284723 SK637 RS08035 WP 033689345.1  LDHREELQDQIETIQRFAVSYIYKPVIIENLVARISAIFRGRDFIDQHCSQMKVPTSYRNLRMDVEHHTVYRGEEMIALT   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 284723 SK637 RS08035 WP 033689345.1  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