```
logo
NTDB id 267538 C0971 RS16315 WP 248892339.1  MKAINFDTKTLDQILEKMIDTVVNSKDEIFRISEQCRKDYETLTDELKDVKQMVLQTIAEGDRLEVQARLARKRLSEVSM   80
NTDB id 130 BSU 35500 NP 391430.1            MNKTKMDSKVLDSILMKMLKTVDGSKDEVFQIGEQSRQQYEQLVEELKQIKQQVYEVIELGDKLEVQTRHARNRLSEVSR   80
consensus                                    !     !*! !! !! !!* !!   !!!!*! ! !!*!  !! ! *!!! *!! !   !  !!*!!!! !*!! !!!!!!

logo
NTDB id 267538 C0971 RS16315 WP 248892339.1  HFKDYTEEEVRDAYEKAHNLQMDLTMNRQLEKQLRQRRDEIERRLAGLSETIERADYLISQITIVLNYLMSDLKQVGEVL  160
NTDB id 130 BSU 35500 NP 391430.1            NFHRFSEEEIRNAYEKAHKLQVELTMIQQREKQLRERRDDLERRLLGLQEIIERSESLVSQITVVLNYLNQDLREVGLLL  160
consensus                                    *!* **!!!*! !!!!!! !!**!!!  ! !!!!! !!!**!!!! !! ! !!! * !*!!!!*!!!!!  !!* !! *!

logo
NTDB id 267538 C0971 RS16315 WP 248892339.1  ENAKQKQDFGLKIIEAQEEERKRLSREIHDGPAQMLANVMMRSDLIERIYKERSAEEAIDEIRNLKRMVRSALYEVRRII  240
NTDB id 130 BSU 35500 NP 391430.1            ADAQAKQDFGLRIIEAQEEERKRVSREIHDGPAQMLANVMMRSELIERIFRDRGAEDGFQEIKNLRQNVRNALYEVRRII  240
consensus                                      !  !!!!!!*!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!***! !!**  !!*!!*  !! !!!!!!!!!

logo
NTDB id 267538 C0971 RS16315 WP 248892339.1  YDLRPMALDDLGLVPTLRKYLQTIEEYHNRTKIYFFNIGE..DKRLPAKYEVALFRLVQESVQNALKHADASEIQVKLEI  318
NTDB id 130 BSU 35500 NP 391430.1            YDLRPMALDDLGLIPTLRKYLYTTEEYNGKVKIHFQCIGETEDQRLAPQFEVALFRLAQEAVSNALKHSESEEITVKVEI  320
consensus                                    !!!!!!!!!!!!!*!!!!!!! ! !!!* * !!*! *!!!**! !!** *!!!!!!! !! ! !!!!! *  !! !!*!!

logo
NTDB id 267538 C0971 RS16315 WP 248892339.1  NKSHVNVVIKDNGKGFDTNE...KKPESFGILGMKERVDLLEGEISIHSKIGAGTVIIIQVPLK.   379
NTDB id 130 BSU 35500 NP 391430.1            TKDFVILMIKDNGKGFDLKEAKEKKNKSFGLLGMKERVDLLEGTMTIDSKIGLGTFIMIKVPLSL   385
consensus                                     ! *! **!!!!!!!!!  !***!!* !!!*!!!!!!!!!!!! **!*!!!! !! !*! !!! *
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X non conserved

X similar

X $\geq$ 50% conserved