```
logo
NTDB id 66086 M3B RS01310 WP 011054187.1   MESIDKSKFRFVERDSEASEVIDTPAYSYWKSVFRQFFSKKSTVFMLVILVTVLMMSFIYPMFANYDFNDVSNINDFSKRYIWPN   85
NTDB id 477 HSISS4 RS06990 WP 021143810.1  MASIDKSKFQFVKRDDFASETIDAPAYSYWKSVMRQFFKKKSTTVMLGILIAIILMSFIYPMFSKFDFNDVSKVNDFSVRYVHPN   85
NTDB id 326 STU RS16120 WP 002946409.1     MASIDKSKFQFVKRDDFASETIDAPSYSYWKSVMRQFFKKKSTVVMLGILITIILMSFIYPMFSKFDFNDVSKVNDFSLRYVHPN   85
NTDB id 294 STER RS06920 WP 002946409.1    MASIDKSKFQFVKRDDFASETIDAPSYSYWKSVMRQFFKKKSTVVMLGILITIILMSFIYPMFSKFDFNDVSKVNDFSLRYVHPN   85
consensus                                  !*!!!!!!!*!!*!!**!!!*!!*! !!!!!!!*!!!!*!!!!**!!*!!*****!!!!!!!***!!!!!!**!!!!*!!**!!

logo
NTDB id 66086 M3B RS01310 WP 011054187.1   AEYWFGTDKNGQSLFDGVWYGARNSILISVIATLINITIGVVLGAIWGVSKVFDKVMIEIYNIISNIPSMLIIIVLTYSLGAGFW   170
NTDB id 477 HSISS4 RS06990 WP 021143810.1  AQYWFGTDSNGKSLFDSVWFGARNSILIAVIATFINVVIGLVVGAVWGISKTFDMIMMEIYNIISNIPALLVVIVLTYSLGAGFW   170
NTDB id 326 STU RS16120 WP 002946409.1     AQYWFGTDGNGKSLFDSVWFGARNSILIAVIATFLNVIIGLLVGAVWGISKTFDMIMMEIYNIISNIPSLLVVIVLTYSLGAGFW   170
NTDB id 294 STER RS06920 WP 002946409.1    AQYWFGTDGNGKSLFDSVWFGARNSILIAVIATFLNVIIGLLVGAVWGISKTFDMIMMEIYNIISNIPSLLVVIVLTYSLGAGFW   170
consensus                                  !*!!!!!!*!!*!!!!*!!*!!!!!!!!!*!!!!**!**!!***!!*!!*!!*!!**!*!!!!!!!!!!!**!**!!!!!!!!!!!!

logo
NTDB id 66086 M3B RS01310 WP 011054187.1   NLILAFCITGWIGVAYSIRVQILRYRDLEYNLASQTLGTPMYKIAVKNLLPQLVSVIMTMLSQMLPVYVSSEAFLSRFGIGLPTT   255
NTDB id 477 HSISS4 RS06990 WP 021143810.1  NMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRTLGTPTAKIVVKNIMPQLVSVIVTMASQLLPGFISYEAFLSYFGLGLPVT   255
NTDB id 326 STU RS16120 WP 002946409.1     NMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRNLGTPTAKIVIKNIMPQLVSVIVTMASQLLPGFISYEAFLSYFGLGLPVT   255
NTDB id 294 STER RS06920 WP 002946409.1    NMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRNLGTPTAKIVIKNIMPQLVSVIVTMASQLLPGFISYEAFLSYFGLGLPVT   255
consensus                                  !*!*!***!!!!!*!!*!!*!!*!!!!!!!!!!!* !!!!**!!**!!**!!!!!!!*!!*!!*!!***!*!!!!!*!!*!!!*!

logo
NTDB id 66086 M3B RS01310 WP 011054187.1   TPSLGRLIANYSSNLTTNAYLFWIPLVTLILVSLPLYIVGQNLADASDPRSHR   308
NTDB id 477 HSISS4 RS06990 WP 021143810.1  TPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
NTDB id 326 STU RS16120 WP 002946409.1     TPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
NTDB id 294 STER RS06920 WP 002946409.1    TPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
consensus                                  !!!!!!!!**!**!*!!!!!!!!!!*!!!!!!!!*!*!!!!!!!!!!!!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved