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logo

NTDB id 68334 SNAG RS01675 WP 033629311.1 MSTIGKEKFQFVKRDDFASETIDAPAYSYWGSVFRQFLKKKSTVIMLGILVAIILMSFIYPMFSNFDFNDVSKVNDFSARFIKPN  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 68334 SNAG RS01675 WP 033629311.1 AEHWFGTDSNGKSLFDGVWFGARNSILISVIATFINLVIGVIVGGIWGISKSVDRVMMEVYNIISNIPSLLIVIVLTYSIGAGFW  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 68334 SNAG RS01675 WP 033629311.1 NLIFAMSVTTWIGIAYMIRIQIMRYRDLEYNLASQTLGTPTFKIIVKNIMPQLVSVIVSTMTLMLPSFISYEAFLSRFGLGLPVT  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 68334 SNAG RS01675 WP 033629311.1 VPSLGRLISDYSQNVTTNAYLFWIPLTTLILVSLSLFVVGQNLADASDPRTHR  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