```
logo
NTDB id 210642 BTR42 RS07530 WP 061459706.1   MTSIDKEQFEFVTLDSAASETIDAPTYSYWKSVFKQFFSRKSTFFMLAILIAIILMSFVYPLFVHCDFSDVSDINDFSKR   80
NTDB id 477 HSISS4 RS06990 WP 021143810.1     MASIDKSKFQFVKRDDFASETIDAPAYSYWKSVMRQFFKKKSTTVMLGILIAIILMSFIYPMFSKFDFNDVSKVNDFSVR   80
NTDB id 326 STU RS16120 WP 002946409.1        MASIDKSKFQFVKRDDFASETIDAPSYSYWKSVMRQFFKKKSTVVMLGILITIILMSFIYPMFSKFDFNDVSKVNDFSLR   80
NTDB id 294 STER RS06920 WP 002946409.1       MASIDKSKFQFVKRDDFASETIDAPSYSYWKSVMRQFFKKKSTVVMLGILITIILMSFIYPMFSKFDFNDVSKVNDFSLR   80
consensus                                     !*!!!!**!*!!**!**!!!!!!!*!!!!!!!**!!!**!!!**!!*!!!! !!!!!!*!!*!***!!*!!!**!!!!*!

logo
NTDB id 210642 BTR42 RS07530 WP 061459706.1   YVWPNSHYWFGTDKDGRSLFDCVWYGARSSILISVIATVLNMIIGVVIGGIWGISKTVDAIMIEVYNVISNIPQLLIVMV   160
NTDB id 477 HSISS4 RS06990 WP 021143810.1     YVHPNAQYWFGTDSNGKSLFDSVWFGARNSILIAVIATFINVVIGLVVGAVWGISKTFDMIMMEIYNIISNIPALLVVIV   160
NTDB id 326 STU RS16120 WP 002946409.1        YVHPNAQYWFGTDGNGKSLFDSVWFGARNSILIAVIATFLNVIIGLLVGAVWGISKTFDMIMMEIYNIISNIPSLLVVIV   160
NTDB id 294 STER RS06920 WP 002946409.1       YVHPNAQYWFGTDGNGKSLFDSVWFGARNSILIAVIATFLNVIIGLLVGAVWGISKTFDMIMMEIYNIISNIPSLLVVIV   160
consensus                                     !!*!!**!!!!!!**!*!!!!*!!*!!!*!!!!*!!!!**!**!!***!**!!!!!!*!*!!*!*!!*!!!!!*!!*!*!

logo
NTDB id 210642 BTR42 RS07530 WP 061459706.1   LTYSLGAGFWNMILAFCITGWIGIAYSIRVQILRYRDLEYNLASQILGTSKVKIIVKNLLPQLISVIVTMVSQLLPSYIS   240
NTDB id 477 HSISS4 RS06990 WP 021143810.1     LTYSLGAGFWNMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRTLGTPTAKIVVKNIMPQLVSVIVTMASQLLPGFIS   240
NTDB id 326 STU RS16120 WP 002946409.1        LTYSLGAGFWNMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRNLGTPTAKIVIKNIMPQLVSVIVTMASQLLPGFIS   240
NTDB id 294 STER RS06920 WP 002946409.1       LTYSLGAGFWNMIFAMTVTGWIGIAYTIRIQIMRYRDLEYNLASRNLGTPTAKIVIKNIMPQLVSVIVTMASQLLPGFIS   240
consensus                                     !!!!!!!!!!!!!!*!***!!!!!!!!*!!*!!*!!!!!!!!!!!!**!!!***!**!!**!!!*!!!!!!*!!!!!**!!

logo
NTDB id 210642 BTR42 RS07530 WP 061459706.1   YEAFLSYFGIGLPITDPSLGRLISNYSSNLTSNAYLFWIPLTILVLVSLPLYIVGQNLADASDPRTHR   308
NTDB id 477 HSISS4 RS06990 WP 021143810.1     YEAFLSYFGLGLPVTTPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
NTDB id 326 STU RS16120 WP 002946409.1        YEAFLSYFGLGLPVTTPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
NTDB id 294 STER RS06920 WP 002946409.1       YEAFLSYFGLGLPVTTPSLGRLISDYAQNVTVNAYLFWIPLTTLILVSLALFIVGQNLADASDPRTHR   308
consensus                                     !!!!!!!!!*!!!*!*!!!!!!!!*!**!*!*!!!!!!!!!!*!*!!!!*!*!!!!!!!!!!!!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved