```
logo
NTDB id 977115 MGAS37857 RS08215 WP 010921892.1  MESIDKSKFRFVERDSEASEVIDTPAYSYWKSVFRQFFSKKSTVFMLVILVTVLMMSFIYPMFANYDFNDVSNINDFSKR   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 977115 MGAS37857 RS08215 WP 010921892.1  YIWPNAEYWFGTDKNGQSLFDGVWYGARNSILISVIATLINITIGWVLGAIWGVSKAFDKVMIEIYNIISNIPSMLIIIV  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 977115 MGAS37857 RS08215 WP 010921892.1  LTYSLGAGFWNLILAFCITGWIGVAYSIRVQILRYRDLEYNLASQTLGTPMYKIAVKNLLPQLVSVIMTMLSQMLPVYVS  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 977115 MGAS37857 RS08215 WP 010921892.1  SEAFLSFFGIGLPTTTPSLGRFIANYSSNLTTNAYLFWIPLVTLILVSLPLYIVGQNLADASDPRSHR  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