```
logo
NTDB id 518 SMSK321 RS11040 WP 000413917.1   .MEKMKKQIRVLLLFTIFGISSCSCLSLSTWALMLNQNCFYYLVGMLLLASIGSGINYYSGQRNQSYSVFK.EDKQQFLYSLLLL   83
NTDB id 490 SM12261 RS08340 WP 000970273.1   MMEKMKKQIRVLLLLTVFGLSSCSCLSLSTWALMLNQNCFYYLVGMLLLASIGAGLNYYRGQGNQHYSVFK.EDKQQFLYSLLLL   84
NTDB id 142 SP RS09775 WP 000839908.1        .....MKSMRILFLLALIQISLSSCFLWKECILSFKQSTAFFIGSMVFVSGICAGVNYLYTRKQEVHSVLASKKSVKLFYSMLLL   80
NTDB id 178 SPR RS08850 WP 000839908.1       .....MKSMRILFLLALIQISLSSCFLWKECILSFKQSTAFFIGSMVFVSGICAGVNYLYTRKQEVHSVLASKKSVKLFYSMLLL   80
NTDB id 212 SPD RS09290 WP 000839908.1       .....MKSMRILFLLALIQISLSSCFLWKECILSFKQSTAFFIGSMVFVSGICAGVNYLYTRKQEVHSVLASKKSVKLFYSMLLL   80
consensus                                    *!**!*!*!******!**!!*******!***!*******!*****!**!*!!**********!!**********!!*!!!

logo
NTDB id 518 SMSK321 RS11040 WP 000413917.1   VNILGLALIVIDHILAGNASVEGFVEAFLPSFFFLFGIDLLVFLPIQKYVNGLKKILGKKKICFISGLAILIFLRNPWMLLSMTF   168
NTDB id 490 SM12261 RS08340 WP 000970273.1   VNILGLALIVIEHILAGNTSVEGFVEAFLPSFFFLFGIDLLVFLPIQKYVNGVKKFLGKKEICFISGLAILIFLRNPWMLLSMTF   169
NTDB id 142 SP RS09775 WP 000839908.1        INLLGAVLVLSDNLFIKNTLQQELVDFLLPSFFFLFGLDLLIFLPLKKYVRDFLAMLDRKKTVLVTILATLLFLRNPMTIVSLLI   165
NTDB id 178 SPR RS08850 WP 000839908.1       INLLGAVLVLSDNLFIKNTLQQELVDFLLPSFFFLFGLDLLIFLPLKKYVRDFLAMLDRKKTVLVTILATLLFLRNPMTIVSLLI   165
NTDB id 212 SPD RS09290 WP 000839908.1       INLLGAVLVLSDNLFIKNTLQQELVDFLLPSFFFLFGLDLLIFLPLKKYVRDFLAMLDRKKTVLVTILATLLFLRNPMTIVSLLI   165
consensus                                    *!*!!**!*********!******!***!!!!!!!!!!*!!!*!!!**!!!******!**!*******!!*!*!!!!!****!***

logo
NTDB id 518 SMSK321 RS11040 WP 000413917.1   YITLGMLFLGLLVPKVIRWEVSFYSHLMRDILLVVFLLFLF   209
NTDB id 490 SM12261 RS08340 WP 000970273.1   YIALGMLFLGLLVPKVIRWEVSFYSHLMRDILLVVFLLFLF   210
NTDB id 142 SP RS09775 WP 000839908.1        YIGLGLFFAAYLVPNSVKKEVSFYGHIFRDLVLVIVTLIFF   206
NTDB id 178 SPR RS08850 WP 000839908.1       YIGLGLFFAAYLVPNSVKKEVSFYGHIFRDLVLVIVTLIFF   206
NTDB id 212 SPD RS09290 WP 000839908.1       YIGLGLFFAAYLVPNSVKKEVSFYGHIFRDLVLVIVTLIFF   206
consensus                                    !!*!!**!***!!!*****!!!!!*!**!!**!!***!**!
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X non conserved

X similar

X ≥ 50% conserved