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logo
NTDB id 38737 KE3 RS00795 WP 020916081.1   MLGIITFIIVFGILVIVHEFGHLYFAKKAGILVREFSIGMGPKIFSHFDKEGTAYTFRILPLGGYVRMAGWGDDTTEIKTGTPAS   85
NTDB id 323 STU RS10650 WP 011225396.1     MKAIITFLLIFCVIVVFHEFGHFFFAKRSGILVREFAIGMGPKIFAHTGKDGTVYTIRILPLGGYVRMAGWGEDTTEIKTGSPAS   85
NTDB id 291 STER RS01205 WP 011680685.1    MKAIITFLLIFCVIVVFHEFGHFFFAKRSGILVREFAIGMGPKIFAHTGKDGTVYTIRILPLGGYVRMAGWGEDTTEIKTGSPAS   85
consensus                                  !**!!!!***!***!**!!!!!**!!!**!!!!!!!*!!!!!!!!!*!**!*!!*!!*!!!!!!!!!!!!!!!!*!!!!!!!!*!!!

logo
NTDB id 38737 KE3 RS00795 WP 020916081.1   LSLNKDGLVTRINLSHKQLDNTALPMNVTSYDLEDKLEITGLVLDEQKTYTVHHDATIVEEDGTEIRIAPLDVQYQNATVWGRLM   170
NTDB id 323 STU RS10650 WP 011225396.1     LTLGKDGKVRRINLSDRQVDQTALPMNVTAYDLEDKLTITGLVLGETKTYEVDHDATIVEEDGTELRIAPKDVQYQNASIWGRLI   170
NTDB id 291 STER RS01205 WP 011680685.1    LTLGKDGKVRRINLSDRQVDQTALPMNVTAYDLEDKLTITGLVLGETKTYEVDHDATIVEEDGTELRIAPKDVQYQNASIWGRLI   170
consensus                                  !*!*!!!*!*!!!!!**!*!*!!!!!!!!*!!!!!!!*!!!!!!*!*!!!*!*!!!!!!!!!!!!*!!!!*!!!!!!!**!!!!*

logo
NTDB id 38737 KE3 RS00795 WP 020916081.1   TNFAGPMNNFILGTLAFILLVFMQGGVPNPSTNAVRVTDGGAMQAAGVKDGDKVLAVGDYKVSNWSELTEAVAKSTKGISKDTTI   255
NTDB id 323 STU RS10650 WP 011225396.1     TNFAGPMNNFILGVLVFIILAFVQGGVQDTSTNLIQVANGGAAQVSGLKTGDAIVAINKDKVTDWDSLKEALRENTQKFSKGDSL   255
NTDB id 291 STER RS01205 WP 011680685.1    TNFAGPMNNFILGVLVFIILAFVQGGVQDTSTNLIQVANGGAAQVSGLKTGDAIVAINKDKVTDWDSLKEALRENTQKFSKGDSL   255
consensus                                  !!!!!!!!!!!!!*!*!!*!*!*!!!!***!!!***!**!!!*!**!*!*!!***!****!!**!**!*!!****!***!!****

logo
NTDB id 38737 KE3 RS00795 WP 020916081.1   PVTVKDASGKTKTLDIKPVKNHGSYLIGVSSALKTGFWDKITGGFQMSWQSAMLILNALKGIVSNFSLDKLGGPVAMYQASSQAA   340
NTDB id 323 STU RS10650 WP 011225396.1     SVTVKRSNGQEETISVKPQKSQGSYFLGVSPVLKTGLKDKIFGGFQMAWEGATAILATLKGLITNFSLNKLGGPVAMFQMSAQAS   340
NTDB id 291 STER RS01205 WP 011680685.1    SVTVKRSNGQEETISVKPQESQGSYFLGVSPVLKTGLKDKIFGGFQMAWEGATAILATLKGLITNFSLNKLGGPVAMFQMSAQAS   340
consensus                                  *!!!!***!***!***!!****!!!**!!!**!!!!**!!!*!!!!!*!**!**!!**!!!***!!!!*!!!!!!!!*!*!*!!*

logo
NTDB id 38737 KE3 RS00795 WP 020916081.1   SYGLTSVINLLALLSINLGIVNLIPIPALDGGKILMNLIEIVRRKPLKQETETYVTLVGVVIMIILMIAVTWNDIMRVFF   420
NTDB id 323 STU RS10650 WP 011225396.1     ESGLISILDLMGMLSINLGIFNLIPIPALDGGKIVMNIIEAIRRKPLNQEIESYITLAGVAVMVVLMIAVTWNDIMRAFF   420
NTDB id 291 STER RS01205 WP 011680685.1    ESGLISILDLMGMLSINLGIFNLIPIPALDGGKIVMNIIEAIRRKPLNQEIESYITLAGVAVMVVLMIAVTWNDIMRAFF   420
consensus                                  **!!*!***!***!!!!!!!*!!!!!!!!!!!!!!*!!*!!**!!!!!*!!*!*!*!!*!!**!**!!!!!!!!!!!!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved