```
logo
NTDB id 30594 SPCG RS01430 WP 000900660.1  MLGILTFILVFGIIVVVHEFGHFYFAKKSGILVREFAIGMGPKIFAHIGKDGTAYTIRILPLGGYVRMAGWGDDTTEIKTGTPVS   85
NTDB id 291 STER RS01205 WP 011680685.1    MKAIITFLLIFCVIVVFHEFGHFFFAKRSGILVREFAIGMGPKIFAHTGKDGTVYTIRILPLGGYVRMAGWGEDTTEIKTGSPAS   85
NTDB id 323 STU RS10650 WP 011225396.1     MKAIITFLLIFCVIVVFHEFGHFFFAKRSGILVREFAIGMGPKIFAHTGKDGTVYTIRILPLGGYVRMAGWGEDTTEIKTGSPAS   85
consensus                                  !**!*!!*!*!**!!!*!!!!!!*!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!*!!!!!!!!!!!!!!!!!!*!!!!!!!!!*!*!
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logo
NTDB id 30594 SPCG RS01430 WP 000900660.1  LTLADDGKVKRINLSGKKLDQTALPMQVTQFDFEDKLFIKGLVLEEEKTFAVDHDATVVEADGTEVRIAPLDVQYQNATIWGKLI   170
NTDB id 291 STER RS01205 WP 011680685.1    LTLGKDGKVRRINLSDRQVDQTALPMNVTAYDLEDKLTITGLVLGETKTYEVDHDATIVEEDGTELRIAPKDVQYQNASIWGRLI   170
NTDB id 323 STU RS10650 WP 011225396.1     LTLGKDGKVRRINLSDRQVDQTALPMNVTAYDLEDKLTITGLVLGETKTYEVDHDATIVEEDGTELRIAPKDVQYQNASIWGRLI   170
consensus                                  !!!**!!!!*!!!!!****!!!!!!!*!!**!*!!!!*!*!!!!*!*!!**!!!!!!*!!*!!!!*!!!!*!!!!!!!*!!!*!!
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logo
NTDB id 30594 SPCG RS01430 WP 000900660.1  TNFAGPMNNFILGVVVFWVLIFMQGGVRDVDTNQFHIMPQGALAKVGVPETAQITKIGSHEVSNWESLIQAVETETKDKT.APTL   254
NTDB id 291 STER RS01205 WP 011680685.1    TNFAGPMNNFILGVLVFIILAFVQGGVQDTSTNLIQVANGGAAQVSGLKTGDAIVAINKDKVTDWDSLKEALRENTQKFSKGDSL   255
NTDB id 323 STU RS10650 WP 011225396.1     TNFAGPMNNFILGVLVFIILAFVQGGVQDTSTNLIQVANGGAAQVSGLKTGDAIVAINKDKVTDWDSLKEALRENTQKFSKGDSL   255
consensus                                  !!!!!!!!!!!!!!!*!!**!*!*!!!!!*!**!!*******!!****!******!**!****!**!*!!**!****!********!
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logo
NTDB id 30594 SPCG RS01430 WP 000900660.1  DVTI.SEKGSDKQVTVTPEDSQGRYLLGVQPGVKSDFLSMFVGGFTTAADSALRILSALKNLIFQPDLNKLGGPVAIFKASSDAA   338
NTDB id 291 STER RS01205 WP 011680685.1    SVTVKRSNGQEETISVKPQESQGSYFLGVSPVLKTGLKDKIFGGFQMAWEGATAILATLKGLITNFSLNKLGGPVAMFQMSAQAS   340
NTDB id 323 STU RS10650 WP 011225396.1     SVTVKRSNGQEETISVKPQKSQGSYFLGVSPVLKTGLKDKIFGGFQMAWEGATAILATLKGLITNFSLNKLGGPVAMFQMSAQAS   340
consensus                                  *!!*****!******!*!**!!!*!*!!!!*!**!********!!!**!***!**!!**!!*!!****!!!!!!!!!*!**!**!*
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logo
NTDB id 30594 SPCG RS01430 WP 000900660.1  KNGIENILYFLAMISINIGIFNLIPIPALDGGKIVLNILEAIRRKPLKQEIETYVTLAGVVIMVVLMIAVTWNDIMRLFFR   419
NTDB id 291 STER RS01205 WP 011680685.1    ESGLISILDLMGMLSINLGIFNLIPIPALDGGKIVMNIIEAIRRKPLNQEIESYITLAGVAVMVVLMIAVTWNDIMRAFF.   420
NTDB id 323 STU RS10650 WP 011225396.1     ESGLISILDLMGMLSINLGIFNLIPIPALDGGKIVMNIIEAIRRKPLNQEIESYITLAGVAVMVVLMIAVTWNDIMRAFF.   420
consensus                                  **!***!!****!*!!!*!!!!!!!!!!!!!!!!!!!
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