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logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  MHQDNQPTFFFFDYETWGTNPAKDRPSQFAGVRTDENFNIIGEPLVMYCQPPADYLPSPEAALITGITPQKTVQEGLSEP   80
NTDB id 1141 GTF74 RS08270 WP 001183680.1    MQQEHQPTFFFFDYETWGVNPAKDRPSQFAGVRTDMDFNIIGEPLVIYCQLPNDYLPAPEAALITGITPQKANSQGLAEP   80
consensus                                    !*!**!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!  !!!!!!!!!*!!!*! !!!! !!!!!!!!!!!!!    !! !!

logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  EFIAKIHAELARPNTTSLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQNGNSRWDLLDVLRACHALRPEGVEWPENEDGFT  160
NTDB id 1141 GTF74 RS08270 WP 001183680.1    EFIARIHAELSTPNTINLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQHGNSRWDLLDVLRACHALRPEGIEWPENDDGFT  160
consensus                                    !!!!*!!!!!  !!!  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!*!!!!

logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  SFKLEHLSVANGIEHSNAHDAMADVVATIEMAKKVKAAQPKLFDYFFSMRQKRKLNDLVDIVNMTPLMHVSGMLGRECQY  240
NTDB id 1141 GTF74 RS08270 WP 001183680.1    SFKLEHLSVANGIEHSNAHDAMADVIATIELAKKVKAAQPKLFDYFFTMRHKRKLTELIDIVNQTPLMHVSGMLGRERHY  240
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!*!!!!!!!!!!!!!!!!!*!!*!!!!  *!*!!!!  !!!!!!!!!!!!!**!

logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  TSWIVPVAWHPTNNNAVITIDLAKDPQPLLELSAEELHERLYTKREDLG.ELLPVPVKLVHLNKCPILAPAKTLTAENAE  319
NTDB id 1141 GTF74 RS08270 WP 001183680.1    TSWIVPIAWHPTNQNAVIVVDLGKDPQPLLELDAETLQARLYTRYEDLAPDELPVPVKLVHLNKCPILAPAKTLTAENAQ  320
consensus                                    !!!!!!*!!!!!!*!!!! *!!*!!!!!*!!! !! !* !!!!* !!!*** !!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  SIGIDRQQCLANLALLRQHPEIREKLISLFSVERKFDKSDDVDTQLYDGFFSPADRAAMDIIRETDPNNLAALDIEFDDK  399
NTDB id 1141 GTF74 RS08270 WP 001183680.1    KIGIDREQCLAHLNVIRQHPEIREKLVTVFSQEREFNQESDVDSQLYAGFFSPADRAAMEIIRTTAPENLGSLAIRVADK  400
consensus                                     !!!!! !!!!*! **!!!!!!!!!!***!! !! !    !!!*!!! !!!!!!!!!!!*!!! ! ! !!* ! !   !!

logo

NTDB id 377931 FR729 RS04515 WP 167415153.1  RIKPLLFRYRARNFPGTLDEQEQRRWALHCREVFESQIEEYMLNLENLVHEHESDEKKIAILKSVYRYVESLAS  473
NTDB id 1141 GTF74 RS08270 WP 001183680.1    RIEPLLFRYRARHYPHTLTDTEQRRWAAHCRDYFERNLEGYMLNLENLVHEHQADPKKMAILKSVYHYVEKLAC  474
consensus                                    !! !!!!!!!!!**!*!! * !!!!!! !!!* !! **! !!!!!!!!!!!!   !*!!*!!!!!!!*!!! !!*
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- X non conserved
- X similar
- X ≥ 50% conserved