```
logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  MHQDNQPTFFFFDYETWGTNPAKDRPSQFAGVRTDENFNIIGEPLVMYCQPPADYLPSPEAALITGIPPQKAVQDGLSEP   80
NTDB id 1141 GTF74 RS08270 WP 001183680.1     MQQEHQPTFFFFDYETWGVNPAKDRPSQFAGVRTDMDFNIIGEPLVIYCQLPNDYLPAPEAALITGITPQKANSQGLAEP   80
consensus                                     !*!**!!!!!!!!!!!!! !!!!!!!!!!!!!!!!  !!!!!!!!!*!!!*! !!!! !!!!!!!!!*!!!!   !! !!

logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  EFIAKIHAELSKPNTTSLGYNSIRFDDEVTRYACYRNFIDPYAWSWQNGNSRWDLLDVMRACHALRPEGIEWPENEDGFT  160
NTDB id 1141 GTF74 RS08270 WP 001183680.1     EFIARIHAELSTPNTINLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQHGNSRWDLLDVLRACHALRPEGIEWPENDDGFT  160
consensus                                     !!!!*!!!!!! !!!  !!!!!!!!!!!!!!! !!!!!!!!!!!!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!*!!!!

logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  SFKLEHLSVRNGIEHSNAHDAMADVIATIEMAKKVKAAQPKLFEYFFSMRQKRKLNELVDIVNMTPLMHVSGMLGRECQY  240
NTDB id 1141 GTF74 RS08270 WP 001183680.1     SFKLEHLSVANGIEHSNAHDAMADVIATIELAKKVKAAQPKLFDYFFTMRHKRKLTELIDIVNQTPLMHVSGMLGRERHY  240
consensus                                     !!!!!!!!! !!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!*!!!!*!!*!!!! !!*!!!! !!!!!!!!!!!!!**!

logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  TSWIVPVAWHPTNNNAVITIDLAKDPQPILDLSAEELHERLYTKREELG.DLLPVPVKLVHLNKCPILAPAKTLTAENAE  319
NTDB id 1141 GTF74 RS08270 WP 001183680.1     TSWIVPIAWHPTNQNAVIVVDLGKDPQPLLELDAETLQARLYTRYEDLAPDELPVPVKLVHLNKCPILAPAKTLTAENAQ  320
consensus                                     !!!!!!*!!!!!!*!!!! *!!*!!!!!*!*! !! !* !!!!* !*!**! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  SIGIDRQQCLTNLALLRQHPEIREKLISLFAVERKFEKDDDVDTQLYDGFFSPADRTAMDIIRETDPNNLAALDIEFDDK  399
NTDB id 1141 GTF74 RS08270 WP 001183680.1     KIGIDREQCLAHLNVIRQHPEIREKLVTVFSQEREFNQESDVDSQLYAGFFSPADRAAMEIIRTTAPENLGSLAIRVADK  400
consensus                                      !!!!! !!! *! **!!!!!!!!!!!***!  !! !  * !!!*!!! !!!!!!!! !!*!!! ! ! !!* ! !   !!

logo
NTDB id 1112430 KHN79 RS06230 WP 182007974.1  RIKPLLFRYRARNFPGTLDEQEQRRWALHCREVFESQIEEYMMNLENLVHEHESDEKKIALLKSVYRYVESLAS  473
NTDB id 1141 GTF74 RS08270 WP 001183680.1     RIEPLLFRYRARHYPHTLTDTEQRRWAAHCRDYFERNLEGYMLNLENLVHEHQADPKKMAILKSVYHYVEKLAC  474
consensus                                     !! !!!!!!!!!**!*!! * !!!!!! !!!* !! **! !!*!!!!!!!!!  !*!!*!*!!!!!*!!! !!*
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X non conserved

X similar

X ≥ 50% conserved