```
logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  MQQEHQPTFFFFDYETWGTSPAKDRPCQFAGVRTDMDFNIIGEPLVIYCQPPADYLPSPEAALITGITPQTAESQGLSEP   80
NTDB id 1141 GTF74 RS08270 WP 001183680.1      MQQEHQPTFFFFDYETWGVNPAKDRPSQFAGVRTDMDFNIIGEPLVIYCQLPNDYLPAPEAALITGITPQKANSQGLAEP   80
consensus                                      !!!!!!!!!!!!!!!!!!  !!!!!!*!!!!!!!!!!!!!!!!!!!!!!!*! !!!! !!!!!!!!!!!! ! !!!! !!

logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  EFIAKIHEQLATPNTISLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQNGNSRWDLLDVMRACYALRPEGVNWPENDEGFI  160
NTDB id 1141 GTF74 RS08270 WP 001183680.1      EFIARIHAELSTPNTINLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQHGNSRWDLLDVLRACHALRPEGIEWPENDDGFT  160
consensus                                      !!!!*!!  ! !!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!*!!!*!!!!!!* !!!!!*!!

logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  SFKLEHLSVANGIEHENAHDAMADVIATIEMAKKVKAAQPKLFDYFLSMRHKRKLNELIDIVNMTPLMHVSGMLGKECNY  240
NTDB id 1141 GTF74 RS08270 WP 001183680.1      SFKLEHLSVANGIEHSNAHDAMADVIATIELAKKVKAAQPKLFDYFFTMRHKRKLTELIDIVNQTPLMHVSGMLGRERHY  240
consensus                                      !!!!!!!!!!!!!!! !!!!!!!!!!!!!!*!!!!!!!!!!!!!!!! *!!!!!!! !!!!!!! !!!!!!!!!!!*!**!

logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  TSWIVPVAWHPTNQNAAICVDLAKDPQALLDLDTEQLNERLYTKRSELAPEELPVPVKLIHLNKCPILAPAKTLTAENAE  320
NTDB id 1141 GTF74 RS08270 WP 001183680.1      TSWIVPIAWHPTNQNAVIVVDLGKDPQPLLELDAETLQARLYTRYEDLAPDELPVPVKLVHLNKCPILAPAKTLTAENAQ  320
consensus                                      !!!!!!*!!!!!!!!!! !*!!!*!!!!*!!*!! ! !* !!!!*  *!!!*!!!!!!!!*!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  KIGIDREKCLENLALLRKHPEIREKLIGVYGIEREYEKSNDVDSMLYDGFFSPADKAAMNIIRETDPNNLSALDISFSDD  400
NTDB id 1141 GTF74 RS08270 WP 001183680.1      KIGIDREQCLAHLNVIRQHPEIREKLVTVFSQEREFNQESDVDSQLYAGFFSPADRAAMEIIRTTAPENLGSLAIRVADK  400
consensus                                      !!!!!!! !! *! **! !!!!!!!!* !*  !!!*    !!!! !! !!!!!!!*!!! !!! ! ! !!  ! !   !

logo
NTDB id 1094510 ACMV04 RS06775 WP 414932177.1  RIAPLLFRYRARHFPWTLDESEQQRWATHCRDYFESRIEDYMLNLENLVHEHESDQKKIAILKSIYQYVQKRVS  474
NTDB id 1141 GTF74 RS08270 WP 001183680.1      RIEPLLFRYRARHYPHTLTDTEQRRWAAHCRDYFERNLEGYMLNLENLVHEHQADPKKMAILKSVYHYVEKLAC  474
consensus                                      !! !!!!!!!!!!*!*!! **!! !!! !!!!!!!  *! !!!!!!!!!!!!  !*!!*!!!!!*!*!! !  *
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X non conserved

X similar

X ≥ 50% conserved