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logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  MTKNSEQPSFFFFDYETWGTSPAKDRPSQFAGIRTDADFNIIGDPLVMYCRLPADYLPGAEAALITGITPQKAMSQGVSE   80
NTDB id 1141 GTF74 RS08270 WP 001183680.1      .MQQEHQPTFFFFDYETWGVNPAKDRPSQFAGVRTDMDFNIIGEPLVIYCQLPNDYLPAPEAALITGITPQKANSQGLAE   79
consensus                                      *  * *!!*!!!!!!!!!!  !!!!!!!!!!!*!!! !!!!!!*!!!*!! !! !!!!**!!!!!!!!!!!!! !!!* !

logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  PEFIGRIYEELARPNTTSLGYNSIRFDDEVTRYTCYRNFFDPYEWSWKNGNSRWDLLDVLRACHALRPEGIVWPENDEGF  160
NTDB id 1141 GTF74 RS08270 WP 001183680.1      PEFIARIHAELSTPNTINLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQHGNSRWDLLDVLRACHALRPEGIEWPENDDGF  159
consensus                                      !!!!*!!* !!  !!!  !!!!!!!!!!!!!!!!!!!!! !!! !!! *!!!!!!!!!!!!!!!!!!!!!! !!!!!*!!

logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  PSFKLEHLSKANGIEHDNAHDALADVIATIEMAKRVKAAQPKLFDYFYSMRNKRKLTELYDIVNMTPLMHVSGMLGRECQ  240
NTDB id 1141 GTF74 RS08270 WP 001183680.1      TSFKLEHLSVANGIEHSNAHDAMADVIATIELAKKVKAAQPKLFDYFFTMRHKRKLTELIDIVNQTPLMHVSGMLGRERH  239
consensus                                      *!!!!!!!! !!!!!! !!!!!*!!!!!!!!*!!*!!!!!!!!!!!!!**!!*!!!!!!!*!!!! !!!!!!!!!!!!!**

logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  YTSWIVPVAWHPTNQNAVITVDLAKDPTPLLELSAQELQTRLYTRREDLADDELPVPVKLVHLNKCPILAPAKTLTAENA  320
NTDB id 1141 GTF74 RS08270 WP 001183680.1      YTSWIVPIAWHPTNQNAVIVVDLGKDPQPLLELDAETLQARLYTRYEDLAPDELPVPVKLVHLNKCPILAPAKTLTAENA  319
consensus                                      !!!!!!!*!!!!!!!!!!! !!!*!!! !!!!! ! !! !!!!! !!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  KVIGINREQCLANLATLREHPAIREKLITLFSQEREFAKDDDVDTQLYDGFFSPADKAAMEIIRESNPEQLGSLDLQFSD  400
NTDB id 1141 GTF74 RS08270 WP 001183680.1      QKIGIDREQCLAHLNVIRQHPEIREKLVTVFSQEREFNQESDVDSQLYAGFFSPADRAAMEIIRTTAPENLGSLAIRVAD  399
consensus                                        !!! !!!!!!*!  *! !! !!!!!*!*!!!!!!!  * !!!*!!! !!!!!!!*!!!!!!! * !!*!!!! *   !

logo

NTDB id 1043102 ACEL84 RS10325 WP 261896273.1  KRIEPLLFRYRARHYPWTLSDTEQRRWANHCREYFENNIEAYMLNLENLVHEHESDAQKIAILKAVYRYVEQLAS  475
NTDB id 1141 GTF74 RS08270 WP 001183680.1      KRIEPLLFRYRARHYPHTLTDTEQRRWAAHCRDYFERNLEGYMLNLENLVHEHQADPKKMAILKSVYHYVEKLAC  474
consensus                                      !!!!!!!!!!!!!!!!!*!!*!!!!!!!! !!!*!!! !*!*!!!!!!!!!!!!  !* !*!!!! !!*!!! !!*
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X non conserved

X similar

X $\geq$ 50% conserved