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logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  MDNNRAPSPADNREQKPTFFVFDYETWGTSPAKDRPSQFAGIRTDEDFNIIGEPVVLYCKLPSDYLPNPEAALITGITPQ   80
NTDB id 1141 GTF74 RS08270 WP 001183680.1      ..........MQQEHQPTFFFDYETWGVNPAKDRPSQFAGVRTDMDFNIIGEPLVIYCQLPNDYLPAPEAALITGITPQ   70
consensus                                      ********** * !* !!!! !!!!!!!   !!!!!!!!!!!*!!! !!!!!!!!*!*!! !! !!!! !!!!!!!!!!!!

logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  KAQQEGLIESEFIARIHQELSTPNTTSFGYNNIRFDDEVTRYTCYRNFIDPYAWSWQNGNSRWDLLDVLRACHALRPEGI  160
NTDB id 1141 GTF74 RS08270 WP 001183680.1      KANSQGLAEPEFIARIHAELSTPNTINLGYNSIRFDDEVTRYTCYRNFIDPYAWSWQHGNSRWDLLDVLRACHALRPEGI  150
consensus                                      !!*  !! !*!!!!!!! !!!!!!!   !!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  EWPENDDGFTSFKLEHLSVANGIEHANAHDALADVIATIEMAKKVKTAQPKLFDYFLSMRHKRKLNGLVDIVNMTPLMHV  240
NTDB id 1141 GTF74 RS08270 WP 001183680.1      EWPENDDGFTSFKLEHLSVANGIEHSNAHDAMADVIATIELAKKVKAAQPKLFDYFFTMRHKRKLTELIDIVNQTPLMHV  230
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!*!!!!!!!!*!!!!! !!!!!!!!! *!!!!!!!   !*!!!! !!!!!!

logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  SGMLGRECQYTSWVVPIAWHPTNNNAVIVADLAKDPSPLLELDAQQLAERLYTKRDQL.GDKSPVPLKLIHLNKCPILAP  319
NTDB id 1141 GTF74 RS08270 WP 001183680.1      SGMLGRERHYTSWIVPIAWHPTNQNAVIVVDLGKDPQPLLELDAETLQARLYTRYEDLAPDELPVPVKLVHLNKCPILAP  310
consensus                                      !!!!!!!**!!!!*!!!!!!!!!*!!!!! !!*!!! !!!!!!!   !   !!!!* * !**!   !!!*!!*!!!!!!!!!!

logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  AKTLTADNAAHIGIDRQQCLDNLAQLRKHAELVNKVVDLYSNQPDFAADDDVDSQLYQGFFSSADKAAMSVVRTTQPENL  399
NTDB id 1141 GTF74 RS08270 WP 001183680.1      AKTLTAENAQKIGIDREQCLAHLNVIRQHPEIREKLVTVFSQEREFNQESDVDSQLYAGFFSPADRAAMEIIRTTAPENL  390
consensus                                      !!!!!!*!! *!!!!! !!! *!  *! !*!*  !*! **!* **!  * !!!!!!! !!!!*!!*!!! **!!! !!!!

logo

NTDB id 1027397 AB0763 RS07270 WP 306100083.1  EGLDIQFSDPRLTPLLFRYRARNYPASLNQQEQEKWAEHCREHFTHALPDHLARIEQLVQQHHDDENKLAILKSVYRYTE  479
NTDB id 1141 GTF74 RS08270 WP 001183680.1      GSLAIRVADKRIEPLLFRYRARHYPHTLTDTEQRRWAAHCRDYFERNLEGYMLNLENLVHEHQADPKKMAILKSVYHYVE  470
consensus                                        ! !   !*!* !!!!!!!!!*!!**!   !! *!! !!!**! * !* **  *!*!!* !* !* !*!!!!!!!*! !

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NTDB id 1027397 AB0763 RS07270 WP 306100083.1  SIAS   483
NTDB id 1141 GTF74 RS08270 WP 001183680.1      KLAC   474
consensus                                       *!*
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X non conserved
X similar
X $\geq$50% conserved