

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVM
.....
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVM

0
80

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

VYKTNRLARNTS DLLKIVEDLHKQNEFFSLSERMEVNTSSGK LMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN
.....
VYKTNRLARNTS DLLKIVEDLHKQNEFFSLSERMEVNTSSGK LMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN

0
160

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
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consensus

LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAK
.....
LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAK

0
240

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

YKDWSEKRRKGLNDKPVIAEGKHSPINQDLWDKVQMRKKQVSQKPQVHGKGTNLLTGTIHCPCGAPMAASNTNTLKD
.....
YKDWSEKRRKGLNDKPVIAEGKHSPINQDLWDKVQMRKKQVSQKPQVHGKGTNLLTGTIHCPCGAPMAASNTNTLKD

12
320

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

GTKKRIRYYSCFQFRNKGASVCHANSIRADQAEQFVAERIKETVQHPQIIKEVNSSST.....
GTKKRIRYYSCSNFRNKGSKVCSANSVRADVIEDVYMKQILEIVKSDKVIQRVVTHVNQENQVDGAALHHDIA YKQQQYD
!!!!!!!!!!!! *!!!!!! !!*!!!!*!!!! !*! * ! ! * * * ! ! * * *****

69
400

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

EVQIKLNNLIKTIEDNPDLTSVIRPSIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHI FHDIKHIEKSRLK
.....
EVQIKLNNLIKTIEDNPDLTSVIRPSIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHI FHDIKHIEKSRLK

69
480

logo

NTDB id 1045836 AB2Q74 RS01525 WP 014930816.1
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consensus

ALYLSVIDRIDIKKDG NHKKQFYVTLKLNNEIIKQLFNNKQLDEVHLSTSSFLPQTLYLTI
.....
ALYLSVIDRIDIKKDG NHKKQFYVTLKLNNEIIKQLFNNKQLDEVHLSTSSFLPQTLYLTI

69
542

 non conserved
 similar
 $\geq 50\%$ conserved