

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVM

0
80

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
VYKTNRLARNTSDLLKIVEDLHKQNEFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN

0
160

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAK

0
240

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
YKDWSEKRRKGLNDKPVIAEGKHSPINQDLWDKVQMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKD
*****!!!!!!

12
320

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
GTTKRIRYYSCFQFRNKGASVCHANSIRADQAEQFVAERIKETVQHPQIIKEVNSSST.....
GTTKRIRYYSCSNFRNKGSKVCSANSVRADVIEDVVMKQILEIVKSDKVIQRVVTHVNQENQVDGAALHHDIA YKQQQYD
!!!!!!!!!!!! *!!!!!! !!*!!!!*!!!! !*! * ! ! * * * ! ! * * *****

69
400

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
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consensus

.....
EVQIKLNNLIKTIEDNPDLTSVIRPSIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLK

69
480

logo

NTDB id 1047759 AB2Q12 RS01525 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

.....
ALYLSVIDRIDIKKDGNHKKQFYVTLKLNNEIIKQLFNNKQLDEVHLSTSSLFLPQTLYLTI

69
542

 non conserved
 similar
 $\geq 50\%$ conserved