

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

QNTTESTLAKDVRTFAEKGYVGDRYGVDDGGFVLRKVDNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKD
.....
QNTTESTLAKDVRTFAEKGYVGDRYGVDDGGFVLRKVDNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKD

42
640

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

NGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKEIITSGDNKTALVYVDLEGNGTNNLIKKIEVPGGKGGGLSSP
.....
NGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKEIITSGDNKTALVYVDLEGNGTNNLIKKIEVPGGKGGGLSSP

42
720

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

TLVDKDLDGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGTKPITSAPAI SQLKDKRWIFGTGSDLSEEDVDNME
.....
TLVDKDLDGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGTKPITSAPAI SQLKDKRWIFGTGSDLSEEDVDNME

42
800

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

EQYIYGI FDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSDGSGNKGWVVKLDGQRVTVKPTVWLRTAFV
.....
EQYIYGI FDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSDGSGNKGWVVKLDGQRVTVKPTVWLRTAFV

42
880

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

TIHKYTGTDKCGAETAALGINTADGGKLTTKSARPIVPEANTAVAQYSGHKKGTNGKSIPIGCMQKSNEIVCPNGYVYDK
.....
TIHKYTGTDKCGAETAALGINTADGGKLTTKSARPIVPEANTAVAQYSGHKKGTNGKSIPIGCMQKSNEIVCPNGYVYDK

42
960

<0

logo

NTDB id 1144566 C7R97 RS14310 WP 306473064.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

PVNVRYLDEKKTDFSTTADGDAGGSGIDPACRKSCKNNRCFSQKGVRTLLMNDLDSLDITGPTCGMKRISWREVFY
.....
PVNVRYLDEKKTDFSTTADGDAGGSGIDPACRKSCKNNRCFSQKGVRTLLMNDLDSLDITGPTCGMKRISWREVFY*

42
1037

non conserved
 similar
 ≥ 50% conserved