

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
80

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
160

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
240

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
320

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
400

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
480

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



40  
560

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

QNTTESTLAKDVRTFAEKGYVGDRYGVDDGGFVLRKVDNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKD  
.....  
QNTTESTLAKDVRTFAEKGYVGDRYGVDDGGFVLRKVDNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKD  
\*\*\*\*\*

40  
640

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
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consensus

NGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKEIITSGDNKTALVYDLEGNGTNNLIKKIEVPGGKGGGLSSP  
.....  
NGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKEIITSGDNKTALVYDLEGNGTNNLIKKIEVPGGKGGGLSSP  
\*\*\*\*\*

40  
720

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TLVDKDL DGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGTKPITSAPAI SQLKDKRWIFGTGSDLSEEDVDNME  
.....  
TLVDKDL DGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGTKPITSAPAI SQLKDKRWIFGTGSDLSEEDVDNME  
\*\*\*\*\*

40  
800

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

EQYIYGI FDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSDGSGNKGWVVKLDGQRVTVKPTVWLRTAFV  
.....  
EQYIYGI FDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSDGSGNKGWVVKLDGQRVTVKPTVWLRTAFV  
\*\*\*\*\*

40  
880

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TIHKYTGTDKCGAETA I LGINTADGCKLTTKSARPIVPEANTAVAQYSGHKKGTNGKSIP I GCMQKSNEIVCPNGYVYDK  
.....  
TIHKYTGTDKCGAETA I LGINTADGCKLTTKSARPIVPEANTAVAQYSGHKKGTNGKSIP I GCMQKSNEIVCPNGYVYDK  
\*\*\*\*\*

40  
960

<0

logo

NTDB id 1147896 CKV73 RS15175 WP 308805568.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

PVNVRYLDEKKT DGFSTTADGDAGGSGIDPACRKSCKNNRCFSQKGVRTLLMNDLDSLDITGPTCGMKRISWREVFY  
.....  
PVNVRYLDEKKT DGFSTTADGDAGGSGIDPACRKSCKNNRCFSQKGVRTLLMNDLDSLDITGPTCGMKRISWREVFY\*  
\*\*\*\*\*

40  
1037

non conserved  
 similar  
 ≥ 50% conserved