

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
85

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
170

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
255

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
340

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
425

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
510

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
595

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
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consensus

DNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKDNGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKE  
.....  
DNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKDNGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKE  
\*\*\*\*\*

43  
680

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
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consensus

IITSGDNKTALVYVDLEGNGTNNLIKKEVPGGKGLSSPTLVKDLDTGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGT  
.....  
IITSGDNKTALVYVDLEGNGTNNLIKKEVPGGKGLSSPTLVKDLDTGTVDIAYAGDRGGNMYRFDLSSQDPQQNSVRTIFEGT  
\*\*\*\*\*

43  
765

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
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consensus

KPITSAPAI SQLKDKRWVIFGTGSDLSEEDVDNMEEQYIYGFDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSD  
.....  
KPITSAPAI SQLKDKRWVIFGTGSDLSEEDVDNMEEQYIYGFDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTLFLTDYKRSD  
\*\*\*\*\*

43  
850

logo

NTDB id 21670 NMA RS14895 WP 306808038.1  
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consensus

GSGNKGWVVKLDGQQRVTVKPTVLRRTAFVTIHKYTGTDKCGAETAILGINTADGGKLTKKSARPIVPEANTAVAQYSGHKKGTN  
.....  
GSGNKGWVVKLDGQQRVTVKPTVLRRTAFVTIHKYTGTDKCGAETAILGINTADGGKLTKKSARPIVPEANTAVAQYSGHKKGTN  
\*\*\*\*\*

43  
935

logo

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consensus

GKSIPIGCMQKSNEIVCPNGYVYDKPVNRYLDEKKTDFSTTADGDAGGSGIDPAGKRSGKNRCSQKGVRTLLMNDLDSLDI  
.....  
GKSIPIGCMQKSNEIVCPNGYVYDKPVNRYLDEKKTDFSTTADGDAGGSGIDPAGKRSGKNRCSQKGVRTLLMNDLDSLDI  
\*\*\*\*\*

43  
1020

<0

logo

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NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TGPTCGMKRISWREVFY  
..... 43  
TGPTCGMKRISWREVFY\* 1037  
\*\*\*\*\*

non conserved  
 similar  
 ≥ 50% conserved