

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



40
85

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



40
170

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



40
255

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus



40
340

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus



40
425

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus



40
510

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus



40
595

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus

DNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKDNGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKE
.....
DNLNGQNRVFMFGAMGFGGRGAYALDLTKADGSDPTAVSLFDVKDNGNNGNRRVELGYTVGTPQIGKTHDGKYAAFLASGYATKE

40
680

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus

IITSGDNKTALVYVDLEGNGTNNLIKKIEVPGGKGLSSPTLVKDL DGTVDIAYAGDRGGNMYRFDLSSQDPQQWSVRTIFEGT
.....
IITSGDNKTALVYVDLEGNGTNNLIKKIEVPGGKGLSSPTLVKDL DGTVDIAYAGDRGGNMYRFDLSSQDPQQWSVRTIFEGT

40
765

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

KPITSAPAI SQLKDKRWVIFGTGSDLSEEDVDNMEEQYIYGFDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTFLFLTDYKRSD
.....
KPITSAPAI SQLKDKRWVIFGTGSDLSEEDVDNMEEQYIYGFDDDTATTGTVNFSDSGGGLLEQVLRDNDNKTFLFLTDYKRSD

40
850

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
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consensus

GSGNKGWVVKLDGQRVTVKPTVLRRTAFVTIHKYTGTDKCGAETAILGINTADGGKLTKKSARPIVPEANTAVAQYSGHKKGTN
.....
GSGNKGWVVKLDGQRVTVKPTVLRRTAFVTIHKYTGTDKCGAETAILGINTADGGKLTKKSARPIVPEANTAVAQYSGHKKGTN

40
935

logo

NTDB id 191838 QP84 RS14635 WP 306807940.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

GKSIPIGCMQKSNEIVCPNGYVYDKPVNVRYLDEKKTDFSTTADGDAGGSGIDPAGKRSKGNNRCFSQKGVRTLLMNDLDSLDI
.....
GKSIPIGCMQKSNEIVCPNGYVYDKPVNVRYLDEKKTDFSTTADGDAGGSGIDPAGKRSKGNNRCFSQKGVRTLLMNDLDSLDI

40
1020

<0

logo

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consensus

TGPTCGMKRISWREVFY
..... 40
TGPTCGMKRISWREVFY* 1037

X non conserved
X similar
X ≥ 50% conserved