

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



80  
80

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



160  
160

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



240  
240

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



320  
320

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



400  
400

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



480  
480

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



560  
560

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

TLVDDLLDYHKMRYGALDIQRCVDLSSATRLVLELSHLLGKKTLRIINQVSEQPVVVSADPQRLEQVLYNLIGNAIKY  
TLVDDLLDYHKMRYGALDIQRCVDLSSATRLVLELSHLLGKKTLRIINQVSEQPVVVSADPQRLEQVLYNLIGNAIKY  
!!

640  
640

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

TSEGKIVISATYIDDKVRVQVDTGGGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV  
TSEGKIVISATYIDDKVRVQVDTGGGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV  
TSEGKIVISATYIDDKVRVQVDTGGGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV  
!!

720  
720

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

GTTFSFTLPLASEEEIAQARLTLPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDSEFLRLEGYRVHTAQ  
GTTFSFTLPLASEEEIAQARLTLPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDSEFLRLEGYRVHTAQ  
GTTFSFTLPLASEEEIAQARLTLPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDSEFLRLEGYRVHTAQ  
!!

800  
800

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

DGHQVLEAIKREKPELVLLDIMPPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA  
DGHQVLEAIKREKPELVLLDIMPPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA  
DGHQVLEAIKREKPELVLLDIMPPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA  
!!

880  
880

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

ARIVAHLTASKAELRRIENAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGKRVFANEAAARLFRRTPEQ  
ARIVAHLTASKAELRRIENAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGKRVFANEAAARLFRRTPEQ  
ARIVAHLTASKAELRRIENAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGKRVFANEAAARLFRRTPEQ  
!!

960  
960

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLLETAVEALSSY  
LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLLETAVEALSSY  
LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLLETAVEALSSY  
!!

1040  
1040

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus

AFEGDKDKLQKLKELGGFTRLADKASGEYQSKQDLMRVAVLVEAMTSALNYWERSVSGSKFTFAEQSGLWRVYLDLDRSTLQ  
AFEGDKDKLQKLKELGGFTRLADKASGEYQSKQDLMRVAVLVEAMTSALNYWERSVSGSKFTFAEQSGLWRVYLDLDRSTLQ  
AFEGDKDKLQKLKELGGFTRLADKASGEYQSKQDLMRVAVLVEAMTSALNYWERSVSGSKFTFAEQSGLWRVYLDLDRSTLQ  
!!

1120  
1120

logo

NTDB id 85913 K6J79 RS00825 WP 001883436.1  
NTDB id 1143 A1552VC RS01965 WP 001883436.1  
consensus



1177  
1177

- non conserved
- similar
- ≥ 50% conserved