

logo

NTDB id 1063213 ACG4I6 RS10935 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



80
80

logo

NTDB id 1063213 ACG4I6 RS10935 WP 001883436.1
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consensus



160
160

logo

NTDB id 1063213 ACG4I6 RS10935 WP 001883436.1
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consensus



240
240

logo

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consensus



320
320

logo

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400
400

logo

NTDB id 1063213 ACG4I6 RS10935 WP 001883436.1
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480
480

logo

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560
560

logo

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consensus

TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
!!

640
640

logo

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TSEGVIVISATYIDDKVRVQVVDGQGI PAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGVIVISATYIDDKVRVQVVDGQGI PAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGVIVISATYIDDKVRVQVVDGQGI PAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
!!

720
720

logo

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GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDLDFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDLDFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPIIIIVADDEPVNLRVLDLDFLRLEGYRVHTAQ
!!

800
800

logo

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DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
!!

880
880

logo

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ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
!!

960
960

logo

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LKRSNAEELIAPKFLNIEQEHYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
!!

1040
1040

logo

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AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
!!

1120
1120

logo

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TRTLDKYLRIETLPKTPRWRTVLNSLDYILEHCKEAGPERTHIEMQRDQLKLLTSE
TRTLDKYLRIETLPKTPRWRTVLNSLDYILEHCKEAGPERTHIEMQRDQLKLLTSE
!!

1177
1177

- non conserved
- similar
- ≥ 50% conserved