

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



80
80

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



160
160

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



240
240

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



320
320

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



400
400

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



480
480

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus



560
560

logo

NTDB id 1089387 ACLRDL RS04330 WP 175295302.1
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consensus

TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVVVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVVVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVVVSADPQRLEQVLYNLIGNAIKY
!!

640
640

logo

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TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
!!

720
720

logo

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GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLD_sELPEQSNLPENEHGPIIIYADDEPVNLRVLD_sFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLD_sELPEQSNLPENEHGPIIIYADDEPVNLRVLD_sFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLD_sELPEQSNLPENEHGPIIIYADDEPVNLRVLD_sFLRLEGYRVHTAQ
!!

800
800

logo

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DGHQVLEAIKREKPELVLLDIMPMGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMPMGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMPMGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
!!

880
880

logo

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ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVRFANAAAARLFRRTPEQ
!!

960
960

logo

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LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSADVLRLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
!!

1040
1040

logo

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AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLD_sRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLD_sRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLD_sRSTLQ
!!

1120
1120

logo

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TRTLDKYLRIETLPKTPRWRTVLNSLDYILEHCKEAGPERTHIEMQRDQLQKLLTSE
TRTLDKYLRIETLPKTPRWRTVLNSLDYILEHCKEAGPERTHIEMQRDQLQKLLTSE
!!

1177
1177

non conserved
 similar
 ≥ 50% conserved