

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



80
80

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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consensus



160
160

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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consensus



240
240

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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consensus



320
320

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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400
400

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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480
480

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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560
560

logo

NTDB id 1150279 HUW69 RS11115 WP 174566572.1
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consensus

TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSHHLLGKKTLRINQVSEQPVVWSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSHHLLGKKTLRINQVSEQPVVWSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSHHLLGKKTLRINQVSEQPVVWSADPQRLEQVLYNLIGNAIKY
!!

640
640

logo

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TSEKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDSRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDSRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDSRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
!!

720
720

logo

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GTTFSFTLPLASEEEEIAQARLTLPHFQAPEVLDSELPEQSTLPENEHGPLLVADEPVLNLRVLDSEFLRLEGYRVHTAQ
GTTFSFTLPLASEEEEIAQARLTLPHFQAPEVLDSELPEQSTLPENEHGPLLVADEPVLNLRVLDSEFLRLEGYRVHTAQ
GTTFSFTLPLASEEEEIAQARLTLPHFQAPEVLDSELPEQSTLPENEHGPLLVADEPVLNLRVLDSEFLRLEGYRVHTAQ
!!

800
800

logo

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DGHQVLEAIKREKPELVLLDIMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
!!

880
880

logo

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ARIVAHLTASKAELRRINAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGNKVRFANEAAARLFRRTPEQ
ARIVAHLTASKAELRRINAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGNKVRFANEAAARLFRRTPEQ
ARIVAHLTASKAELRRINAQLQKELKHRAMVEASLLETQGRILLEQLESAPAEILCVKEGNKVRFANEAAARLFRRTPEQ
!!

960
960

logo

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LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRPQGSGLQAMYIFNVGGSYNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRPQGSGLQAMYIFNVGGSYNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYGNIDVYVDDVRQHLSADVLRPQGSGLQAMYIFNVGGSYNAARIHNLETAVEALSSY
!!

1040
1040

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consensus

AFEGDKDKLQKLKELGGEFTRADKASGEYQSKQDLMRVAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDRLTQ
AFEGDKDKLQKLKELGGEFTRADKASGEYQSKQDLMRVAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDRLTQ
AFEGDKDKLQKLKELGGEFTRADKASGEYQSKQDLMRVAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDRLTQ
!!

1120
1120

