

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



80
80

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



160
160

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



240
240

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



320
320

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



400
400

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



480
480

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus



560
560

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
TLVDDLLDYHKMRYGALDIQRCAVDLSSATRLVLELSSHLLGKKTLLRIINQVSEQPVMVSADPQRLEQVLYNLIGNAIKY
!!

640
640

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
TSEGKIVISATYIDDKVRVQVDTGQGIPAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLVSSQPMV
!!

720
720

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPLLLVADDEPVNLRVLDLDFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPLLLVADDEPVNLRVLDLDFLRLEGYRVHTAQ
GTTFSFTLPLASEEEIAQARLTELPHFQAPEVLDSELPEQSNLPENEHGPLLLVADDEPVNLRVLDLDFLRLEGYRVHTAQ
!!

800
800

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
DGHQVLEAIKREKPELVLLDIMMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEAGANDYLSKPFNKQELA
!!

880
880

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVFANEAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVFANEAAAARLFRRTPEQ
ARIVAHLTASKAELRRIENAQLQKELKHRAVVEASLLETQGRILLEQLESAPAEAILCVKEGKRVFANEAAAARLFRRTPEQ
!!

960
960

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSDVLRQLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSDVLRQLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
LKRSNAEELIAPKFLNIEQEHCYCGNIDVYVDDVRQHLSDVLRQLPQGSGLQAMYIFNVGGSVNAARIHNLETAVEALSSY
!!

1040
1040

logo

NTDB id 1062974 ACG4I2 RS11140 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
AFEGDKDKLQKLKELGGEFTRLADKASGEYQSKQDLMRAVLVEAMTSALNYWERVSGQSKFTFAEQSGLWRVYLDLDRSTLQ
!!

1120
1120

