

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



80
80

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



160
160

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



240
240

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus



320
320

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus



400
400

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus



480
480

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus



560
560

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus

HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
!!

640
640

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus

LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGP I P A P M E R K A G R Y R A H M V I L S A D R A R L H F Y L R Q W W A Q
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGP I P A P M E R K A G R Y R A H M V I L S A D R A R L H F Y L R Q W W A Q
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGP I P A P M E R K A G R Y R A H M V I L S A D R A R L H F Y L R Q W W A Q
!!

720
720

logo

NTDB id 361070 EA667 RS17110 WP 000156653.1
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consensus

LVHAPRQHQLRLSIDVDPQEF S
LVHAPRQHQLRLSIDVDPQEF S
LVHAPRQHQLRLSIDVDPQEF S
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

742
742

- non conserved
- similar
- ≥ 50% conserved