

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
NTDB id 1072 ABD1 RS01805 WP 000156662.1  
consensus



80  
80

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
NTDB id 1072 ABD1 RS01805 WP 000156662.1  
consensus



160  
160

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus



240  
240

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus



320  
320

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus



400  
400

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus



480  
480

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus



560  
560

logo

NTDB id 359496 EA719 RS16690 WP 000156646.1  
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consensus

HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL  
HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL  
!!

640  
640

logo

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consensus

LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMVLSADRARLHFYLRQWVAQ  
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMVLSADRARLHFYLRQWVAQ  
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMVLSADRARLHFYLRQWVAQ  
!!

720  
720

logo

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consensus

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

742  
742

- non conserved
- similar
- ≥ 50% conserved