

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



80
80

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



160
160

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



240
240

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus



320
320

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus



400
400

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus



480
480

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus



560
560

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus

HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
!!

640
640

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus

LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
!!

720
720

logo

NTDB id 423313 G3R47 RS02165 WP 000156656.1
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consensus

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

742
742

- non conserved
- similar
- ≥ 50% conserved