

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



80
80

logo

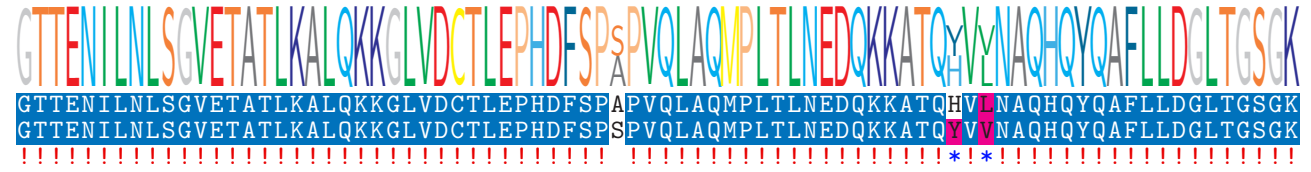
NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



160
160

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



240
240

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



320
320

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



400
400

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



480
480

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



560
560

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
HVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGRAGRGEHKGHVYLQTLRPDHP LLTTLIEKDYRAVAKQTLAERKVAL
!!

640
640

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
LPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVDIWGPI PAMP ERKAGRYRAHVMILSADRARLHFYLRQWVAQ
!!

720
720

logo

NTDB id 409198 GQ096 RS17060 WP 032030927.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

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

742
742

- non conserved
- similar
- ≥ 50% conserved