

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



45
80

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



45
160

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



45
240

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
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consensus



45
320

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
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consensus



45
400

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



45
480

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus



45
560

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

QNT**EST**LAKDV**RT**FAEKGYV**GDRY**GVDGGFVLRKVDNLNGQNRV**FMFGAM**FGG**R**GAYALDLTKADGSDPTAVSLFDV**KD**
.....
QNT**EST**LAKDV**RT**FAEKGYV**GDRY**GVDGGFVLRKVDNLNGQNRV**FMFGAM**FGG**R**GAYALDLTKADGSDPTAVSLFDV**KD**

45
640

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GT**TNNLIKKIEVPGGK**GL**S**SP**
.....
NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GT**TNNLIKKIEVPGGK**GL**S**SP**

45
720

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDPQ**QW**SVRTIFEGTKPITSAPAI**SQL**KDKR**WV**IFGTGS**DL**SEEDVD**NME**
.....
TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDPQ**QW**SVRTIFEGTKPITSAPAI**SQL**KDKR**WV**IFGTGS**DL**SEEDVD**NME**

45
800

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

EQYI**YGI**FDDDTATTG**TVN**FSDSGG**LL**EQVLR**RD**N**DN**KT**LF**LTDYKRSDGSG**NK**GWV**KL**KD**Q**RV**TV**KPT**V**LR**TA**FV
.....
EQYI**YGI**FDDDTATTG**TVN**FSDSGG**LL**EQVLR**RD**N**DN**KT**LF**LTDYKRSDGSG**NK**GWV**KL**KD**Q**RV**TV**KPT**V**LR**TA**FV

45
880

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

TIHKYT**G**TDK**CGA**ETAILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**Y**SGHKK**G**TNGKS**I**PI**G**CMQ**K**SNEI**V**CP**NG**Y**V**YDK
.....
TIHKYT**G**TDK**CGA**ETAILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**Y**SGHKK**G**TNGKS**I**PI**G**CMQ**K**SNEI**V**CP**NG**Y**V**YDK

45
960

<0

logo

NTDB id 301747 DRA66 RS11370 WP 192887912.1
NTDB id 1090 CAA90909.1 1..3114()
consensus

PV**N**VRYLDEK**K**TDGF**ST**TADGDAGG**SG**IDPAGK**R**SGK**NN**R**C**FSQ**K**GV**RT**LL**M**NDL**D**SLDITG**P**TC**G**M**K**RI**S**W**R**EV**F**Y
.....
PV**N**VRYLDEK**K**TDGF**ST**TADGDAGG**SG**IDPAGK**R**SGK**NN**R**C**FSQ**K**GV**RT**LL**M**NDL**D**SLDITG**P**TC**G**M**K**RI**S**W**R**EV**F**Y*

45
1037

non conserved
 similar
 ≥ 50% conserved