

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
80

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
160

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
240

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
320

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
400

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
480

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus



43  
560

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
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**  
\*\*\*\*\*

43  
640

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GT**TNNLIK**KE**IEVPGGK**GGL**S**SP**  
.....  
NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GT**TNNLIK**KE**IEVPGGK**GGL**S**SP**  
\*\*\*\*\*

43  
720

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDP**QQ**WSV**RT**IFEGTKPITSAPAI**SQL**KDKR**W**IFGTGS**DL**SEEDVD**NME**  
.....  
TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDP**QQ**WSV**RT**IFEGTKPITSAPAI**SQL**KDKR**W**IFGTGS**DL**SEEDVD**NME**  
\*\*\*\*\*

43  
800

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

EQYI**Y**GIFDDDTATTG**TV**NFSDSGG**LL**EQVLR**RD**N**DN**KT**LF**LTDYKRSDGSG**NK**GWV**KL**KD**GQ**RV**TV**KPT**W**L**R**TA**FV**  
.....  
EQYI**Y**GIFDDDTATTG**TV**NFSDSGG**LL**EQVLR**RD**N**DN**KT**LF**LTDYKRSDGSG**NK**GWV**KL**KD**GQ**RV**TV**KPT**W**L**R**TA**FV**  
\*\*\*\*\*

43  
880

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TI**H**KYT**G**TDK**CGA**ETAILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**Y**SGH**KK**GT**NG**KS**I**PI**G**CM**Q**KS**NE**IVCP**NG**Y**VY**DK  
.....  
TI**H**KYT**G**TDK**CGA**ETAILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**Y**SGH**KK**GT**NG**KS**I**PI**G**CM**Q**KS**NE**IVCP**NG**Y**VY**DK  
\*\*\*\*\*

43  
960

<0

logo

NTDB id 301652 DRA75 RS11425 WP 306807169.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

P**V**N**V**RYLDEK**KT**DGF**ST**TADGDAGG**SG**ID**P**AG**K**RS**G**K**NN**R**C**FS**Q**K**G**V**R**T**LL**M**N**D**L**D**S**L**D**IT**G**PT**CG**M**K**R**I**SW**R**EV**F**Y  
.....  
P**V**N**V**RYLDEK**KT**DGF**ST**TADGDAGG**SG**ID**P**AG**K**RS**G**K**NN**R**C**FS**Q**K**G**V**R**T**LL**M**N**D**L**D**S**L**D**IT**G**PT**CG**M**K**R**I**SW**R**EV**F**Y\*  
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43  
1037

non conserved  
 similar  
 ≥ 50% conserved