



logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

QNT**EST**LAKDV**RT**FAEKGYV**GDRY**GVDGGFVLRKVDNLNGQNRV**FMFGAM**FGG**RGAY**ALDLTKADGSDPTAVSLFDV**KD**  
.....  
QNT**EST**LAKDV**RT**FAEKGYV**GDRY**GVDGGFVLRKVDNLNGQNRV**FMFGAM**FGG**RGAY**ALDLTKADGSDPTAVSLFDV**KD**  
\*\*\*\*\*

41  
640

logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GTNN**L**IKK**IEVPGGK**GGL**S**SP**  
.....  
NGNNGN**NR**VELGYTVG**TPQ**IGK**THD**GKYAAFLASGYATKEIITSGDNKTALYVYDLEGN**GTNN**L**IKK**IEVPGGK**GGL**S**SP**  
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41  
720

logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDP**QQW**SVRTIFEGTKPITSAPAI**SQL**KDKR**WV**IFGTGS**DL**SEEDVD**NME**  
.....  
TLVDKDL**DGT**VDIAYAGDRGG**NMY**RFDLSSQDP**QQW**SVRTIFEGTKPITSAPAI**SQL**KDKR**WV**IFGTGS**DL**SEEDVD**NME**  
\*\*\*\*\*

41  
800

logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

EQYI**YGI**FDDDTATTG**TVN**FSDSGG**LL**EQVLR**RDND**NKTLFLTDYKRSDGSG**NKG**W**VKL**KD**GQR**VT**VKPT**W**LRT**AFV  
.....  
EQYI**YGI**FDDDTATTG**TVN**FSDSGG**LL**EQVLR**RDND**NKTLFLTDYKRSDGSG**NKG**W**VKL**KD**GQR**VT**VKPT**W**LRT**AFV  
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41  
880

logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

TIHKYT**GTD**KCGA**ET**AILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**YSG**HKKGT**NG**KS**IP**IG**MQ**KS**NE**IVCP**NG**YVYDK  
.....  
TIHKYT**GTD**KCGA**ET**AILG**INT**ADGG**KL**TKKSAR**P**IVPEANTAVAQ**YSG**HKKGT**NG**KS**IP**IG**MQ**KS**NE**IVCP**NG**YVYDK  
\*\*\*\*\*

41  
960

<0

logo

NTDB id 305764 DV144 RS14220 WP 275895500.1  
NTDB id 1090 CAA90909.1 1..3114( )  
consensus

PV**NV**RYLDEK**KT**DGF**STT**ADGDAG**GS**GID**PAG**KRS**GK**NN**RC**FSQ**KG**V**RTL**LL**MND**LSLDIT**GP**TC**GM**KRIS**W**RE**VFY**  
.....  
PV**NV**RYLDEK**KT**DGF**STT**ADGDAG**GS**GID**PAG**KRS**GK**NN**RC**FSQ**KG**V**RTL**LL**MND**LSLDIT**GP**TC**GM**KRIS**W**RE**VFY**\*  
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41  
1037

non conserved  
 similar  
 ≥ 50% conserved