

logo

NTDB id 339869 ES968 RS12565 WP 129134318.1
NTDB id 96 BSU 24730 NP 390353.1
NTDB id 1028 TT RS08215 WP 011173991.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
.....
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK

0
0
80

logo

NTDB id 339869 ES968 RS12565 WP 129134318.1
NTDB id 96 BSU 24730 NP 390353.1
NTDB id 1028 TT RS08215 WP 011173991.1
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDtlsleevedLTGLWEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
.....
ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDtlsleevedLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG

0
0
160

logo

NTDB id 339869 ES968 RS12565 WP 129134318.1
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NTDB id 1028 TT RS08215 WP 011173991.1
consensus

EGQKDLKLGELLLQKGWISREALEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL
.....
EGQKDLKLGELLLQKGWISREALEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL

0
0
240

logo

NTDB id 339869 ES968 RS12565 WP 129134318.1
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LRSDALRYGAVPIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
.....
LRSDALRYGAVPIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL

0
0
320

logo

NTDB id 339869 ES968 RS12565 WP 129134318.1
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KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP
.....
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP

0
0
400

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NTDB id 339869 ES968 RS12565 WP 129134318.1
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DPGAPLLLPEDLCRRYGVPFHRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNVEVAPAVATEAAITKLIERFYGKAEL
.....
DPGAPLLLPEDLCRRYGVPFHRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNVEVAPAVATEAAITKLIERFYGKAEL

0
0
480

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consensus



- X non conserved
- X similar
- X >50% conserved