

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 69519 BVS141 RS12050 WP 007408320.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 80
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 0
..... 0
..... 0

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consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 160
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 0
..... 0
..... 0

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EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL 240
EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL 0
..... 0
..... 0

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LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 320
LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 0
..... 0
..... 0

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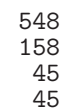
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI 399
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPPYDPEEDP. 14
.....MSQYTICEQQTTERI 0
..... 0
* * *

logo

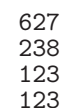
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FDISPSRWQPNLLPKQLRRYLVYPVBLTGRLWLKVDPRNLTALFVRLMTRVQYEPVWITSDELTYLLNALS 471
....PDPGAPLLLPEDLCCRYPGVFPHRLEGNRIVLLMKDPRNLTALDDVRLALKRKGLNVEVAPAVATEAAITKLLI.... 89
FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDELKYLNNALS 0
..... 0
* * * * * * * * * * * * * * * * * * * *

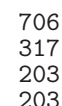
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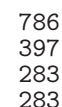
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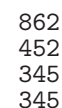
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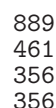
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 non conserved
 similar
 $\geq 50\%$ conserved