

logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 399252 GI367 RS07790 WP 007612574.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
.....
.....

0
80
0
0

logo

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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 399252 GI367 RS07790 WP 007612574.1
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consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
.....
.....

0
160
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0

logo

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consensus

EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL
EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL
.....
.....

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240
0
0

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consensus

LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
.....
.....

0
320
0
0

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consensus

KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI
.....MSQYTYVESQTERI
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYIPYVDPEEDP.
.....
.....

14
399
0
0

* * *

logo

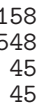
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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 399252 GI367 RS07790 WP 007612574.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKIWLAVDDENNITACEIFAFM.TH...KQIEPVVIASDEIKYLLINALS
.....PDPGAPLLLPEDLCCRYPGVFPHRLEGNRLVLLMKDPRNITLALDDVRLALKRKGLNVEVAPAVATEAAITKLT....
.....
.....

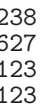
89
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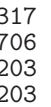
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consensus



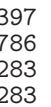
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consensus



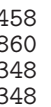
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consensus



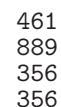
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consensus



 non conserved
 similar
 $\geq 50\%$ conserved