

logo

NTDB id 146751 ABA10 RS12345 WP 015483431.1
NTDB id 96 BSU 24730 NP 390353.1
NTDB id 1028 TT RS08215 WP 011173991.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
.....
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
80

logo

NTDB id 146751 ABA10 RS12345 WP 015483431.1
NTDB id 96 BSU 24730 NP 390353.1
NTDB id 1028 TT RS08215 WP 011173991.1
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNPLDLSLEEVEDLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
.....
ALLPAEKAKELKAIPFALDEEAGVVRVAFNPLDLSLEEVEDLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
160

logo

NTDB id 146751 ABA10 RS12345 WP 015483431.1
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NTDB id 1028 TT RS08215 WP 011173991.1
consensus

EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL
.....
EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL
240

logo

NTDB id 146751 ABA10 RS12345 WP 015483431.1
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consensus

LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
.....
LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
320

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NTDB id 146751 ABA10 RS12345 WP 015483431.1
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consensus

KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP
.....
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP
400

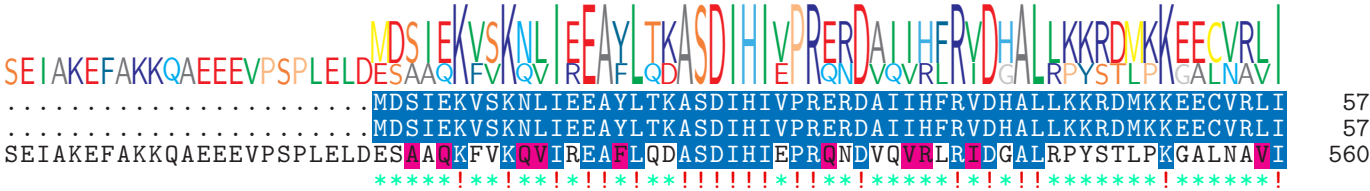
logo

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consensus

DPGAPLLLPEDLCRRYGVPFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNVEVAPAVATEAAITKLIERFYGKAEL
.....
DPGAPLLLPEDLCRRYGVPFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNVEVAPAVATEAAITKLIERFYGKAEL
480

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consensus



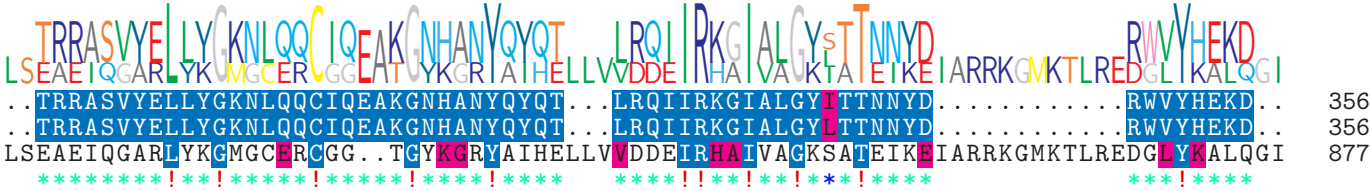
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consensus



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NTDB id 1028 TT RS08215 WP 011173991.1
consensus



 non conserved
 similar
 ≥ 50% conserved