

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
NTDB id 1384 A4U84 RS08525 WP 010787006.1  
NTDB id 1020205 AB0R72 RS13495 WP 003153083.1  
NTDB id 96 BSU 24730 NP 390353.1  
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 80  
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 0  
..... 0  
..... 0  
..... 0

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NTDB id 1020205 AB0R72 RS13495 WP 003153083.1  
NTDB id 96 BSU 24730 NP 390353.1  
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 160  
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 0  
..... 0  
..... 0  
..... 0

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EGQKDLKLGELLLQKGWISREALVEEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL 240  
EGQKDLKLGELLLQKGWISREALVEEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL 0  
..... 0  
..... 0  
..... 0

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LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 320  
 LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 0  
 ..... 0  
 ..... 0  
 ..... 0

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KEAL**EVQ**KGL**P**RAK**PL**GE**IL**VELGLAR**P**ED**VEE**AL**Q**K**Q**RRGGGR**L**ED**T**LV**Q**SG**K**LR**P**EAL**A**Q**A**V**A**T**M**LG**P**CD**P**EE**P**I  
 KEAL**EVQ**KGL**P**RAK**PL**GE**IL**VELGLAR**P**ED**VEE**AL**Q**K**Q**RRGGGR**L**ED**T**LV**Q**SG**K**LR**P**EAL**A**Q**A**V**A**T**Q**LG**P**Y**P**V**D**EE**D**P  
 .....MS**Q****Y****T****I****C****E****Q****T****T****E****R**I  
 .....  
 .....

399  
 14  
 0  
 0

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[illegible]

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 non conserved  
 similar  
  $\geq 50\%$  conserved