

logo

```
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1042625 ACEF29 RS13070 WP 053573196.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
```

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 80
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 0
..... 0
..... 0
..... 0

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consensus
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ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTKSAFLYALAKHYPELGLPVPPPPSG 160
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTKSAFLYALAKHYPELGLPVPPPPSG 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**I**L**VEL**GLAR**P**ED**VEE**AL**Q**K**QRR**GGG**R**LE**D**TL**VQ**SG**K**LR**P**EAL**AQ**AVAT**ML**GG**P**CD**PEED**PI
 KEAL**EVQ**KGL**P**RAK**PL**GE**I**L**VEL**GLAR**P**ED**VEE**AL**Q**K**QRR**GGG**R**LE**D**TL**VQ**SG**K**LR**P**EAL**AQ**AVAT**QL**GY**P**Y**PV**D**P**EE**D**P
MS**Q**Y**T**IC**E**Q**Q**T**E**R

399
 14
 0
 0

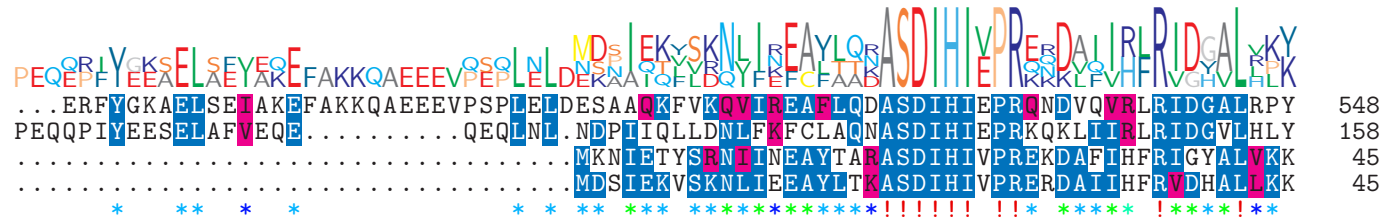
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FDISPSRWQPNLLPEKQLRRYLYVPHLEENRLWLVMKDERNLTALCELYRFLMTRKGLNKYEAPVVIATSPDELTLYLNALS
...PDPGAPLLLPEDLCRRYGVFPHRLEGNRIVLLMKDPRNITLALDDVRLALKRKGLNYPEVAPATAEAAITKKLTI... 471
FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDENNLTACIFAFM.TH...KQIEPVVIASDELKYLNLALS 89
... 0
... 0

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 non conserved
 similar
 $\geq 50\%$ conserved