

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1058693 ACFXAA RS19175 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

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

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NTDB id 1058693 ACFXAA RS19175 WP 003153083.1
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consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLD~~TL~~SLEEVEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVP~~PP~~PSG 160
 ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLD~~TL~~SLEEVEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVP~~PP~~PSG
 0
 0
 0

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EGQKDLKLGELLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL 240
EGQKDLKLGELLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL 0
..... 0
..... 0
..... 0

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[illegible]

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KEAL**EVQ**KGL**P**RA**K**PL**G**E**I**L**V**ELGLAR**P**ED**V**EEAL**Q****K**QRRGGGR**L**ED**T**LV**Q**S**G**KLR**P**EAL**A**Q**A**V**A**T**M**L**G****P****I**C**D**PE**E**D**P****I** 399
 KEAL**EVQ**KGL**P**RA**K**PL**G**E**I**L**V**ELGLAR**P**ED**V**EEAL**Q****K**QRRGGGR**L**ED**T**LV**Q**S**G**KLR**P**EAL**A**Q**A**V**A**T**M**L**G****P****I**C**D**PE**E**D**P****I** 14
MS**Q****T****I**C**E****Q****T****T****E****R**I 0
 0
 0

logo

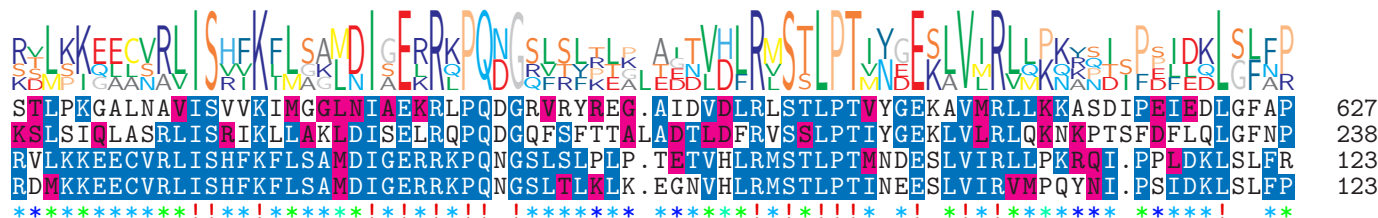
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FDISPSRWQAKNCEEKQLLRRYLYYPVRLTEENRLVLVMDERNITALELYRMLTRKGLNKYEYEPVVIASDELKYLLNALS
 ...PDPGAPLLLLPEDLCRRYGVPFHRLEGNRIIVLLMKDPRIILALDDVRLALKRKGLNVEYAPAVATEAAITKKLI... 471
 FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLLWLAVDDENNITACEIFAFM.TH...KQIEPVVIASDELKYLLNALS 89
 0
 0
 0

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 non conserved
 similar
 $\geq 50\%$ conserved