

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 45187 BAMTA208 RS12225 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

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

logo

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NTDB id 45187 BAMTA208 RS12225 WP 013352872.1
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consensus
```

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

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EGQKDLKLGELLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLL	240
EGQKDLKLGELLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLL	0
.....	0
.....	0
.....	0

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[illegible]

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KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAAVATMLGYPYDPEEDP. 399
 KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAAVATMLGYPYDPEEDP. 14
MSQYTICQQTERTI 0
 0
 *

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FDISPSRWQAKNCEEKQLLRRYLYYPVRLTEENRLVLVMDERNITALELYRFLMTRKGLNKYEYEPVVIATSPDELTLYLNALS
 ...PDPGAPLLLLPEDLCRRYGVPFHRLEGNRIIVLLMKDPRIILALDDVRLALKRKGLNVEYAPAVATEAAITKKLI... 471
 FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLLWLAVDDENNITACEIFAFM.TH...KQIEPVVIASDELKYLNLALS 89
 0
 0
 0

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 non conserved
 similar
 $\geq 50\%$ conserved