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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1079070 ACJJTF RS12060 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
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MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 80
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 0
..... 0
..... 0
..... 0

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1079070 ACJJTF RS12060 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
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ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTKSAFLYALAKHYPELGLPVPPPPSG 160
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTKSAFLYALAKHYPELGLPVPPPPSG 0
..... 0
..... 0
..... 0

```
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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1079070 ACJJTF RS12060 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
```

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

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consensus

LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 320
 LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 0
 0
 0
 0

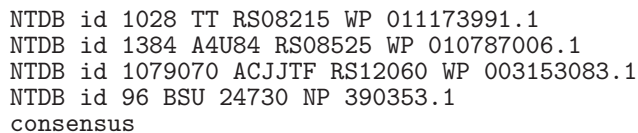
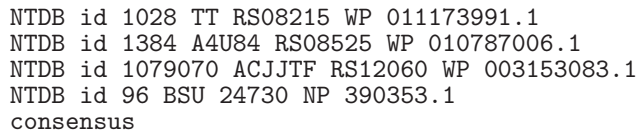
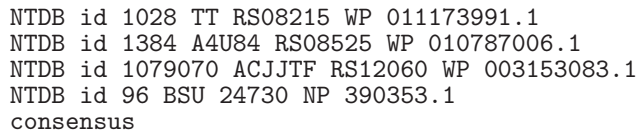
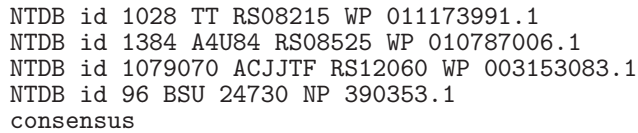
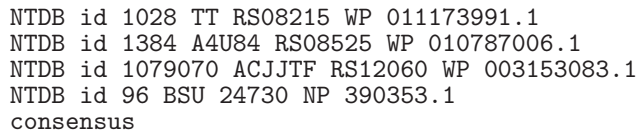
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consensus

KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPTCDPEEDPI
 KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP. 399
MSQYTICQQTTERI 14
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consensus

FDISPSRWQPNLLKQLRLRYLYVPHVLTCHNLWLVLVKDERNLTLALELYRMLTRKGLNKYEIPVWITSDEITYLINALS
PDPGAPLLLPEDLCRRYGVFPHRLGNRLVLLMKDPRNITLALDDVRLALKRKGLNYEVPAAVDEAAAITKLI.... 471
 FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDEIKYLIINALS 89
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NTDB id 96 BSU 24730 NP 390353.1
consensus



 non conserved
 similar
 $\geq 50\%$ conserved