

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 45335 BAXH7 RS12250 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....

85
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logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 45335 BAXH7 RS12250 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFLNPLD~~TL~~SLEEVEDLTGLWEPYQTTKSAFLYALAKHYPELGLPVP~~PP~~PPPSGEGQDKLKLGE
 EKAKELKAIPFALDEEAGVVRVAFLNPLD~~TL~~SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVP~~PP~~PPPSGEGQDKLKLGE

170
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logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 45335 BAXH7 RS12250 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALVEEALVEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF
 LLLQKGWISREALVEEALVEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF

255
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0

logo

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NTDB id 1028 TT RS08215 WP 011173991.1
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NTDB id 45335 BAXH7 RS12250 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
```

QNGEEVWVLSDPRHKEAQAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEEVWVLSDPRHKEAQAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

340
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0
0

logo

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NTDB id 45335 BAXH7 RS12250 WP 013352872.1
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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSCKLRPEALAQAVATMLGPTCDPEEPIDFISPSRWQNLLPKQLRLRYLYXP
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSCKLRPEALAQAVATMLGYPYVDPEEDP...PDPGAPLLLLPEDLCRRYGVFP
MSQYTICQQTERIFDISPSRWQKNCEEKQLLLRYLAVP

* * *

420
39
0
0

logo

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NTDB id 45335 BAXH7 RS12250 WP 013352872.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

[illegible]

498
109
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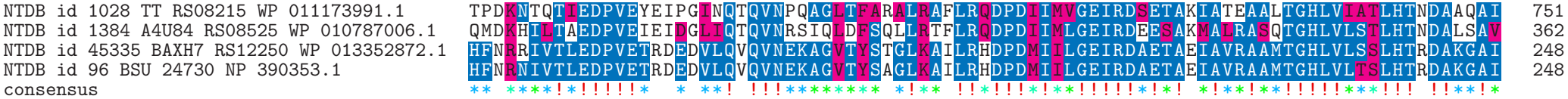
logo



logo



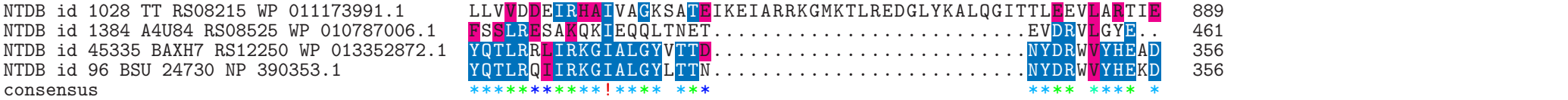
logo



logo



logo



⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved