

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 114849 V529 RS12515 WP 038459186.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTI GDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
MSVLTI GDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
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85
0
0
0

logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 114849 V529 RS12515 WP 038459186.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG
EKAKELKAIPFALDEEAGVVRVAFNLPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG
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170
0
0
0

logo

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NTDB id 114849 V529 RS12515 WP 038459186.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREAL EEA LVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDP SAALLLLRSDALRYGAVPIGF
LLLQKGWISREAL EEA LVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDP SAALLLLRSDALRYGAVPIGF
.....
.....

255
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVWLS D PRHKEAVAQLLN RPARFYLA LPQAW EELFR RAYPQKNR LGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEVEVWLS D PRHKEAVAQLLN RPARFYLA LPQAW EELFR RAYPQKNR LGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
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.....

340
0
0
0

logo

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NTDB id 114849 V529 RS12515 WP 038459186.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQS GKLRPEALAQAVATMLGIPYCDPEEDPIFDISPSRWQPNLLPKSLRRYLYVP
VELGLARPEDVEEALQKQRRGGGRLEDTLVQS GKLRPEALAQAVATMLGIPYCDPEEDPIFDISPSRWQPNLLPKSLRRYLYVP
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420
39
0
0

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logo

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NTDB id 114849 V529 RS12515 WP 038459186.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

YRLTGNRLWLVKDRNLTAL ELYRIMTR YQYEPVWITSDELTYLLNALSPEQEPFEEAEAEAKLFAKKQAE E E VPE
YRLTGNRLWLVKDRNLTAL ELYRIMTR YQYEPVWITSDELTYLLNALSPEQEPFEEAEAEAKLFAKKQAE E E VPE
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498
109
0
0

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