

logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1468 ASU2 RS05090 WP 014991734.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 29949 BACAU RS12020 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....

0
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85
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0

logo

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NTDB id 1468 ASU2 RS05090 WP 014991734.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 29949 BACAU RS12020 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE
.....
EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE
.....

0
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170
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logo

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NTDB id 29949 BACAU RS12020 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLLLRSDALRYGAVPIGIF
.....
LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLLLRSDALRYGAVPIGIF
.....

0
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logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1
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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 29949 BACAU RS12020 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....
QNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

0
0
340
0
0

logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1
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NTDB id 1028 TT RS08215 WP 011173991.1
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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYVEVEEETDPAELLWQKNCLEKSLLLRYLAVP
.....
MSQYTTICEQQTERIFDISPSRWQKNCEEKQLLLRYLAVP
.....
MQYSVCDMNSQQVFDISESLWQKNCDEKHIILRYLAIP
.....
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPPDPGAPL.....LLPEDLCRRYGVFP
.....

39
38
420
0
0

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