

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 118150 KHU1 RS10670 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTI GDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
MSVLTI GDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....
.....

85
0
0
0

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 118150 KHU1 RS10670 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG
EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG
.....
.....

170
0
0
0

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 118150 KHU1 RS10670 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF
LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF
.....
.....

255
0
0
0

logo

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NTDB id 118150 KHU1 RS10670 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVWLSDRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEVEVWLSDRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....
.....

340
0
0
0

logo

NTDB id 1028 TT RS08215 WP 011173991.1
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NTDB id 118150 KHU1 RS10670 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

VELGLARPEDVEEALQKQRRGGRLLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPIFDISPSRWQPNLLPKSLRRYLYWP
VELGLARPEDVEEALQKQRRGGRLLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPIFDISPSRWQPNLLPKSLRRYLYWP
.....
.....
MSQYITICEQQTERRIFDISPSRWQKNCEEKQLLLRYLAVP
.....
.....

420
39
0
0

* * * * *

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

YRLTGNRLWLVKDRNLTALFLYRIMTRVQYEPVWITSDELTYLLNALSPPEQRIYCKSELSEYEQEFAKKQAEVEVPE
YRLTGNRLWLVKDRNLTALFLYRIMTRVQYEPVWITSDELTYLLNALSPPEQRIYCKSELSEYEQEFAKKQAEVEVPE
.....
.....
HQEEEEKLVAMDENLTLCDFAFALKHKGLNKEIAIAAAEAATKKLLNALSPPEQRIYCKSELSEYEQEFAKKQAEVEVPE
.....
.....
VQETEHKLVAVDDENLTLACEIFAFM.TH...KQIEPVVIASDELKYLNLNALSPPEQRIYCKSELSEYEQEFAKKQAEVEVPE
.....
.....

498
109
0
0

** * * * ** * * * * * ** * * * *

logo

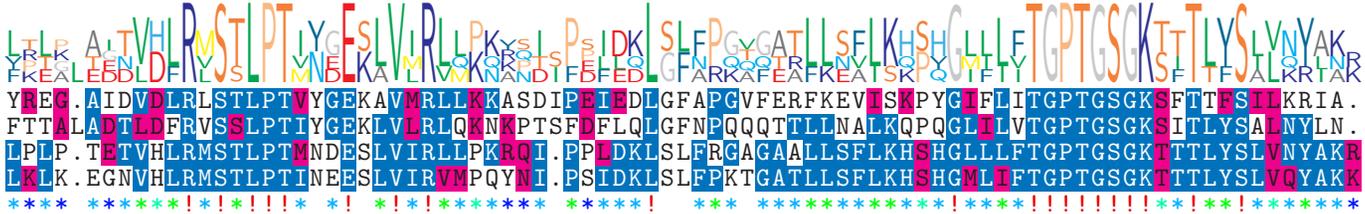
NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1384 A4U84 RS08525 WP 010787006.1
 NTDB id 118150 KHU1 RS10670 WP 012117985.1
 NTDB id 96 BSU 24730 NP 390353.1
 consensus



583
193
80
80

logo

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 NTDB id 1384 A4U84 RS08525 WP 010787006.1
 NTDB id 118150 KHU1 RS10670 WP 012117985.1
 NTDB id 96 BSU 24730 NP 390353.1
 consensus



666
277
163
163

logo

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 NTDB id 1384 A4U84 RS08525 WP 010787006.1
 NTDB id 118150 KHU1 RS10670 WP 012117985.1
 NTDB id 96 BSU 24730 NP 390353.1
 consensus



751
362
248
248

logo

NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1384 A4U84 RS08525 WP 010787006.1
 NTDB id 118150 KHU1 RS10670 WP 012117985.1
 NTDB id 96 BSU 24730 NP 390353.1
 consensus



832
432
325
325

logo

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



889
461
356
356

non conserved
 similar
 ≥ 50% conserved