

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 63175 U471 RS11465 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA 85
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA 0
..... 0
..... 0

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 63175 U471 RS11465 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG 170
EKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLG 0
..... 0
..... 0

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consensus

LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF 255
LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF 0
..... 0
..... 0

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consensus

QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLEGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL 340
QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLEGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL 0
..... 0
..... 0

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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPEEDPIFDISPSRWQPNLLPKOLLRRYLVYP 420
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPEEDPIFDISPSRWQPNLLPKOLLRRYLVYP 39
..... MSQYITICEQQTERRIFDISPSRWQKNCEEKQLLLRYLAVP 0
..... 0
* * * * * ** *

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consensus

VRITGNRLVLLVLDPRNITALEYVRLMTRVYQYEDVWITSDELTYLITNALSPQEPPFYCKSELSEYEQEFAKKQAEVEEVS 498
HRLEGNRTVLLMKDPRNITALEYVRLMTRVYQYEDVWITSDELTYLITNALSPQEPPFYCKSELSEYEQEFAKKQAEVEEVS 109
VQETEHKLVLAWDDENNTACEIFAFM.TH...KQTEPVVIASDEIKYLLNALSPQEPPFYCKSELSEYEQEFAKKQAEVEEVS 0
..... 0
** * * * ** * * * * * * * * *

logo



logo



logo



logo



logo



non conserved
 similar
 ≥ 50% conserved