

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
NTDB id 1384 A4U84 RS08525 WP 010787006.1  
NTDB id 258212 BVQ RS13060 WP 053573196.1  
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 258212 BVQ RS13060 WP 053573196.1  
NTDB id 96 BSU 24730 NP 390353.1  
consensus

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

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consensus

LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF 255  
LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPASAALLLLRSDALRYGAVPIGF 0  
..... 0  
..... 0

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consensus

QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRNGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL 340  
QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRNGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL 0  
..... 0  
..... 0

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VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPTCDPEEDPIFDISPSRWQPNLLPKOLLRRYLVYP 420  
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPTCDPEEDPIFDISPSRWQPNLLPKOLLRRYLVYP 39  
..... MSQYITICEQQTERRIFDISPSRWQKNCEEKQLLLRYLAVP 0  
..... 0  
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consensus

VRITGNRLVLLVLDPRNITALEIVRIMTRVYVEDVWITSDELTYLITNALSPEQEPFLYCKSELSEYEQEFAKKQAEVEEVS 498  
HRLEGNRTVLLMKDPRNITALEIVRIMTRVYVEDVWITSDELTYLITNALSPEQEPFLYCKSELSEYEQEFAKKQAEVEEVS 109  
VQETEHKLVLAWDDENNTACEIFAFM.TH...KQTEPVVIASDEIKYLLNALSPEQQPIVEESLAFVEQE...QE 0  
..... 0  
\*\* \* \* \* \*\* \* \* \* \* \* \* \* \* \* \* \* \* \*

