

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
NTDB id 1384 A4U84 RS08525 WP 010787006.1  
NTDB id 116719 AW02 RS12365 WP 003153083.1  
NTDB id 96 BSU 24730 NP 390353.1  
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA  
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA  
.....

85  
0  
0  
0

logo

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NTDB id 116719 AW02 RS12365 WP 003153083.1  
NTDB id 96 BSU 24730 NP 390353.1  
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTLGLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE  
 EKAKELKAIPFALDEEAGVVRVAFNLPLDTLGLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE  
 .....  
 .....

170  
0  
0  
0

logo

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consensus

LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAAALLLLRSDALRYGAVPIGF  
 LLLQKGWISREALVEEALVEQEKTDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAAALLLLRSDALRYGAVPIGF  
 .....  
 .....

255  
0  
0  
0

logo

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consensus

QNGEVEVVLSDPRHKEAVAQLLNRPARFYLLALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL  
QNGEVEVVLSDPRHKEAVAQLLNRPARFYLLALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL  
.....

340  
0  
0  
0

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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPEEDPIFDISDPGAKLCCEEDCLRLRYLYP  
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP...PDPGAPLLLPEDLCRRYGVFP  
.....MSQYTICEQQTERIFDISPSRWQKNCEEKQLLLRYLAVP

420  
39  
0

\* \* \* \* \*

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NTDB id 96 BSU 24730 NP 390353.1  
consensus

YRLTISNRLLVLYKDPNRNITALELYRIMLTRKGLNKETAI A AA EAATKKLTNALSP EQEPFEEAAEAKFAKKQAE E EVPE  
 HRLEGNRLVLLMKDPRNITALLDVRLLAKRKGLN Y E VAPAVATEAAITKLI.....ERFYGKAELSEIAKEFAKKQAE E EVPS  
 VQETEHKLWLAVDDENNITACEIFAFM.TH...KQIEPVIASDELKYLIINALSP EQQPIY E EELAFVEQE.....QE

498  
109  
0

\* \* \* \*

