

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTITGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA	85
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTITGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA	0
NTDB id 45260 LL3 RS12715 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLGE	170
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLGE	0
NTDB id 45260 LL3 RS12715 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF	255
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF	0
NTDB id 45260 LL3 RS12715 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL	340
NTDB id 1384 A4U84 RS08525 WP 010787006.1	QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL	0
NTDB id 45260 LL3 RS12715 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPEEDPIFDISPSRWQPNLLPKOLLRRYLVWP	420
NTDB id 1384 A4U84 RS08525 WP 010787006.1	VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP.....PDPGAPLLLPEDLCRRYGVFP	39
NTDB id 45260 LL3 RS12715 WP 013352872.1	MSQYTICEQQTERIFDISPSRWQKNCEEKQLLLRYLAVP	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	* * * * *	

logo

NTDB id 1028 TT RS08215 WP 011173991.1	VRITGNRLVLLVKDPRNLTALFVRLMTRKGLNKETAIATAAEAAATKKLTNALSPQEPPFYCKSELSEYEQ	498
NTDB id 1384 A4U84 RS08525 WP 010787006.1	HRLEGNRLVLLMKDPRNLTALDDVRLALKRKGLNYEVAPAVATEAAATKLT.....ERFYGKAELSETAKEFAKKQAEVEEVP	109
NTDB id 45260 LL3 RS12715 WP 013352872.1	VQETEHKLWLAADDENNTACEIFAFM.TH....KQTEPVVIASDEKYLNALSPQQPIVEESLAFVEQ.....EQE	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	** * * * ** * * * * * * * * *	

