

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 205990 BMJ37 RS12245 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 205990 BMJ37 RS12245 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	240
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
NTDB id 205990 BMJ37 RS12245 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 205990 BMJ37 RS12245 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI	399
NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI	14
NTDB id 205990 BMJ37 RS12245 WP 014418378.1	MSQYTICEQQTTERI	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	

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NTDB id 1028 TT RS08215 WP 011173991.1	FDISPSRWQPNLLPKQLRRYLVYPVBLTGRLWLVKDPRLNTALEIVRIMTRVQYEPVWITSDELTYLLNALS	471
NTDB id 1384 A4U84 RS08525 WP 010787006.1	PDPGAPLLLLPEDLCRRYGVPFHRLEGNRIVLLMKDPRNIALALDDVRLALKRKGLNVEVAPAVATEAAITKLLI....	89
NTDB id 205990 BMJ37 RS12245 WP 014418378.1	FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDELKYLLNALS	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	

 non conserved
 similar
 $\geq 50\%$ conserved