

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1104384 ACNM63 RS11870 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	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1104384 ACNM63 RS11870 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
NTDB id 1104384 ACNM63 RS11870 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRSDALRYGAVPIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 1104384 ACNM63 RS11870 WP 014418378.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI	399
NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPYCDPEEDPI	14
NTDB id 1104384 ACNM63 RS11870 WP 014418378.1	MSQYTICEQQTERRI	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	FDISPSRWQPNLLPKQLLRPYLWVPLTGNRLWLKMDPRNLTALFVRLMTRVQYEPVWITSDELTYLLNALS	471
NTDB id 1384 A4U84 RS08525 WP 010787006.1	PDPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLMKDPRNLTALDDVRLALKRKGLNVEVAPAVATEAAITKLI....	89
NTDB id 1104384 ACNM63 RS11870 WP 014418378.1	FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDELKYLNALS	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

 non conserved
 similar
 $\geq 50\%$ conserved