

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 994828 AAVC70 RS02710 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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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 994828 AAVC70 RS02710 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALVEEQEKTCDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLGELLLQKGWISREALVEEQEKTCDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
NTDB id 994828 AAVC70 RS02710 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 994828 AAVC70 RS02710 WP 013352872.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPPEEDPI	399
NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGIPICDPPEEDPI	14
NTDB id 994828 AAVC70 RS02710 WP 013352872.1	MSQYTICQQTETRI	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	0

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NTDB id 1028 TT RS08215 WP 011173991.1	FDISPSRWQPNLLEKQLLRPYLYYDYLTCNRLWLVLVDENNTALELYRIMTRKGLNKEITAIAAAEAATKKLTNALS	471
NTDB id 1384 A4U84 RS08525 WP 010787006.1	PDPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLMKDPRNLTALDDVRLALKRKGLNYEAPAVATEAATKLT....	89
NTDB id 994828 AAVC70 RS02710 WP 013352872.1	FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNTACIFAFM.TH....KQIEPVVIASDELKYLTNALS	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	0

 non conserved
 similar
 $\geq 50\%$ conserved