

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 260177 CXR48 RS11880 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWVEPYQTtksaflyalakhyPELGLPVPPPPSG	160
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDtlsleevedLTGLWVEPYQTtksaflyalakhyPELGLPVPPPPSG	0
NTDB id 260177 CXR48 RS11880 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLgelllQKGWISREALEEALVEQEKtGdLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL	240
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLgelllQKGWISREALEEALVEQEKtGdLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL	0
NTDB id 260177 CXR48 RS11880 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAQAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAQAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 260177 CXR48 RS11880 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		0

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NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDtlvqsgklrPEALAQAVATMLGIPtyvesoter	399
NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDtlvqsgklrPEALAQAVATMLGIPtyvesoter	14
NTDB id 260177 CXR48 RS11880 WP 003153083.1	MSQYtICEQqTERI	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	0

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NTDB id 1028 TT RS08215 WP 011173991.1	FDISPSRWQPNLLPKQLRRYLVYPVBLTGRLWLVKDPRLNLTALFIVRMTRVQYEPVWITSDELTYLLNALS	471
NTDB id 1384 A4U84 RS08525 WP 010787006.1	PDPGAPLLLLPEDLCRRYGVPFHRLEGNRIVLLMKDPRNLTALDDVRLALKRKGLNVEVAPAVATEAAITKLLI....	89
NTDB id 260177 CXR48 RS11880 WP 003153083.1	FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFM.TH....KQIEPVVIASDELKYLLNALS	0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	0

 non conserved
 similar
 $\geq 50\%$ conserved