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NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS	LAEVLVDMGLLSERRIAQTI	EDRFGIPLVELHRVEIPPKVKALLPA	85
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS	LAEVLVDMGLLSERRIAQTI	EDRFGIPLVELHRVEIPPKVKALLPA	0
NTDB id 64205 U722 RS11760 WP 003153083.1				0
NTDB id 96 BSU 24730 NP 390353.1				0
consensus				

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NTDB id 1028 TT RS08215 WP 011173991.1	EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPPSGEGQKDLKLGE	170
NTDB id 1384 A4U84 RS08525 WP 010787006.1		0
NTDB id 64205 U722 RS11760 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF	255
NTDB id 1384 A4U84 RS08525 WP 010787006.1		0
NTDB id 64205 U722 RS11760 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	QNQGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL	340
NTDB id 1384 A4U84 RS08525 WP 010787006.1		0
NTDB id 64205 U722 RS11760 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP.....PDPGAPLLLPEDLCRRYGVFP	420
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSQYITICEQQTIRIFDISPSRWQKNCEEKQLLLRYLAVP	39
NTDB id 64205 U722 RS11760 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*.....*	

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NTDB id 1028 TT RS08215 WP 011173991.1	HRLEGNRLVLLVLDPRNLTALFDVRLALKKRGLNYEYAPAVATEAAATKLLI.....ERFYGKAELSELAKELFAKKQAEVEVPS	498
NTDB id 1384 A4U84 RS08525 WP 010787006.1	VQETEHKLWLAADDENNTACEIFAFM.TH....KQIEPVVIASDELKYLLNALSPEQQPIVEESLAFVEQE.....QE	109
NTDB id 64205 U722 RS11760 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*** * * * * ** * * * * * * * * * * * * * * *	

