

logo

	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWQSENIAATLTEIRRHFHHTLKGSGRMIGAKSSA	
NTDB id 1046 HON27 RS03130 WP 168726941.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWQSENIAATLTEIRRHFHHTLKGSGRMIGAKSSA	80
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFQKQKRAHTLDFRPLILLGQQLQQQSLLEPALEELLQLS	
NTDB id 1046 HON27 RS03130 WP 168726941.1	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFQKQKRAHTLDFRPLILLGQQLQQQSLLEPALEELLQLS	160
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	
NTDB id 1046 HON27 RS03130 WP 168726941.1	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	240
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVVRDSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQ	
NTDB id 1046 HON27 RS03130 WP 168726941.1	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVVRDSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQ	320
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	HGDRTPPLMPHGLVLSQLLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	
NTDB id 1046 HON27 RS03130 WP 168726941.1	HGDRTPPLMPHGLVLSQLLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	400
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVVKQHQSI LLEELKLYTQYTPDISNDLPQQDSTSFEP IYNIEP	
NTDB id 1046 HON27 RS03130 WP 168726941.1	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVVKQHQSI LLEELKLYTQYTPDISNDLPQQDSTSFEP IYNIEP	480
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	EPEVWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM	
NTDB id 1046 HON27 RS03130 WP 168726941.1	EPEVVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM	560
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	RYLHTLKGGANMVQATYLGILIAHELESYERLIQKQLVWTSDLIDFIRLVQDDLADRLQIMREQQLDYAAPYITINALKRA	
NTDB id 1046 HON27 RS03130 WP 168726941.1	RYLHTLKGGANMVQATYLGILIAHELESYERLIQKQLVWTSDLIDFIRLVQDDLADRLQIMREQQLDYAAPYITINALKRA	640
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	GQNSNFQPLPVVEAFDAESEVFSEKQVISEIIDEIPVELEPALAELETHHDQVFDTAVTELATPVEMITTVTSQENEVA	
NTDB id 1046 HON27 RS03130 WP 168726941.1	GQNSNFQPLPVVEAFDAESEVFSEKQVISEIIDEIPVELEPALAELETHHDQVFDTAVTELATPVEMITTVTSQENEVA	720
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	ANEQDIEAVVEQTFLEEATELLEMAESLLKQWFEQRTNRSILLQLQRAVHSLKGGARMVGLAEVQAIAYQLENAFEQFAL	
NTDB id 1046 HON27 RS03130 WP 168726941.1	ANEQDIEAVVEQTFLEEATELLEMAESLLKQWFEQRTNRSILLQLQRAVHSLKGGARMVGLAEVQAIAYQLENAFEQFAL	800
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	HHFNSNIYDHLLLESAIAWLKDAIFNHNYQHFDGLQQSLENIQFFETTIIQIPTKLTRADLFSSEPVMTFIQGDGTEPPPM	
NTDB id 1046 HON27 RS03130 WP 168726941.1	HHFNSNIYDHLLLESAIAWLKDAIFNHNYQHFDGLQQSLENIQFFETTIIQIPTKLTRADLFSSEPVMTFIQGDGTEPPPM	880
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	GAWEQTERLDQNNEMIRVSADLIEKMIDLSGENSINRSRIEMDLSQFSHTLVEMELAIQRLADQLRRMEGELETQIIAKH	
NTDB id 1046 HON27 RS03130 WP 168726941.1	GAWEQTERLDQNNEMIRVSADLIEKMIDLSGENSINRSRIEMDLSQFSHTLVEMELAIQRLADQLRRMEGELETQIIAKH	960
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 1132934 NMG88 RS14025 WP 026795306.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

