

logo

	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFHRTLKGSGRMIGAKSSA	
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NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1046 HON27 RS03130 WP 168726941.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFHRTLKGSGRMIGAKSSA	80
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NTDB id 1132922 NMG85 RS19415 WP 214560400.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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	ELAWTVEDTLNLRVINQSLQLTPTIQSYVQLVFKFYFLKLVDNFKQKRAHTLDFRPLILLGQQQLQQQSSLEPALEELLQLS	
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTLMEEAEHLATIHQFLDQELHQYDSYNALIRALHTLR	
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consensus		

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	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVVRDSLELLSRYSSSEQLEARLLQFNQSWDAYVEQ	
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NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 413 AAK55817.1 141..845()		0
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	HGDRTDPLMP SHGLVSQLLQLDVSELDDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVKQHQSI LLEELKLYTQYTPDISNDLPQQDSTSFEPIYNIEP	0
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NTDB id 1046 HON27 RS03130 WP 168726941.1	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVKQHQSI LLEELKLYTQYTPDISNDLPQQDSTSFEPIYNIEP	480
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	EPEVVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGI DTDLNIWVGEQENFAALNNLM	0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	RYLHTLKGGANMVQATYLG LIAHELES IYERLI QKQLVVTSDLI DFI RLVQDDLADRLQIMREQQLDYAAPYTINALKRA	0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	GQNSNFQPLPVVEAFDAESEVFSEQKVI SEI I I DEIPVELEPALAELETHHDQVFD TAVTELATPVEMI TTVTSQENEVA	0
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consensus		

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NTDB id 413 AAK55817.1 141..845()		0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	HHFNSNIYDHLLES AIAWLKDAIFNHNYQHFDGLQQSLENI QFFETT I QI I PTKLTRADLFSSEPVM TFI QGDGTEPPPPM	0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	GAWEQTERLDQNNEMI RVSADLI EKMI DLSGENS I NRSRI EMDLSQFSHTLVEMELAI QRLADQLRRMEGELETQ I I AKH	0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	G I E H S R Y T D F D P L E M D Q Y S S L N Q L S K S L A E S A S D L V D F K N T L S D K I R D T E S L L L Q Q S R I Q A E I Q E G L M R T R L V P F S R L L P	0
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	R L Q R I V R Q T S T A L N R P A E L F V N N T E G E L D R N I L E R L V T P L E H M L R N A I D H G L E D R A Q R Q Q A N K P E T G R I E L N I Q R Q G T D V	0
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consensus		

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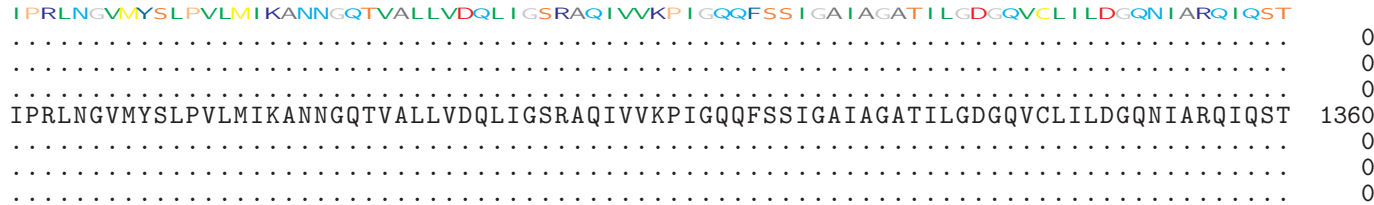
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consensus		

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NTDB id 372 SMU RS05280 WP 002262198.1	V Y G Q G T T F T I R V P T T V A V S D A L M V K V A D Q Q F A I P L A Q I D R I I R V S P A S L E Q Y F E S P Q E L F E Y E N K R Y P L R Y L S E F V G N Q P	0
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consensus		

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consensus



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consensus



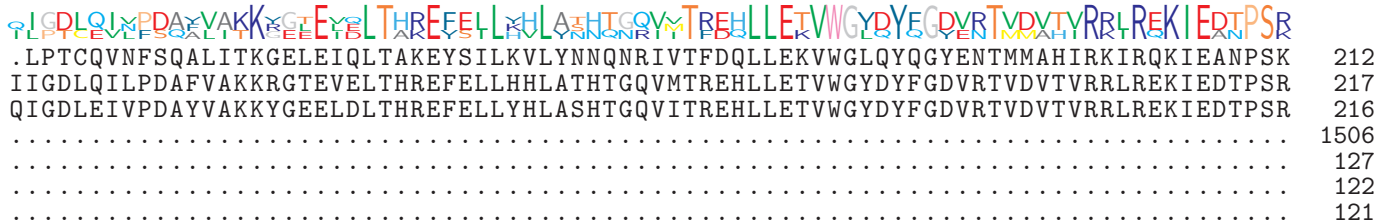
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consensus



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consensus



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NTDB id 1132922 NMG85 RS19415 WP 214560400.1			122
NTDB id 1417 SGL RS04200 WP 014407071.1			121
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved