

logo

	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHRTLKGSGRMIGAKSSA	
NTDB id 372 SMU RS05280 WP 002262198.1		0
NTDB id 1419 SGL RS12550 WP 010873252.1		0
NTDB id 1143 A1552VC RS01965 WP 001883436.1		0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 1046 HON27 RS03130 WP 168726941.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHRTLKGSGRMIGAKSSA	80
NTDB id 55045 CYAN10605 RS14695 WP 015220734.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDNFKQKRAHTLDFRPLILLGQQQLQQQSSLEPALEELLQLS	
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

logo

	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	
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NTDB id 55045 CYAN10605 RS14695 WP 015220734.1		0
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consensus		

logo

	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQ	
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consensus		

logo

	HGDRTDPLMP SHGLV SQLQLD VSEL L DAE LDFEKGIRNEFPDYLERLSEQADLLLQH THSQAMLGLHEYTSQLKESYEV	
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consensus		

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	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAQQRVG VVKQHQSI L EELKLYTQYTPDISNDLPQQDSTSF EPIYNI EP	
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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	EPEVVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEELEGG..IDTDLNIWVGEQENFAALNN	
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consensus		

logo

	LMRYLHTL...KGGANMVQATYLG LIAHELES IYERLIQKQLVVTSD LIDFIRLVQDDLADR....LQIM....REQQL	
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consensus		

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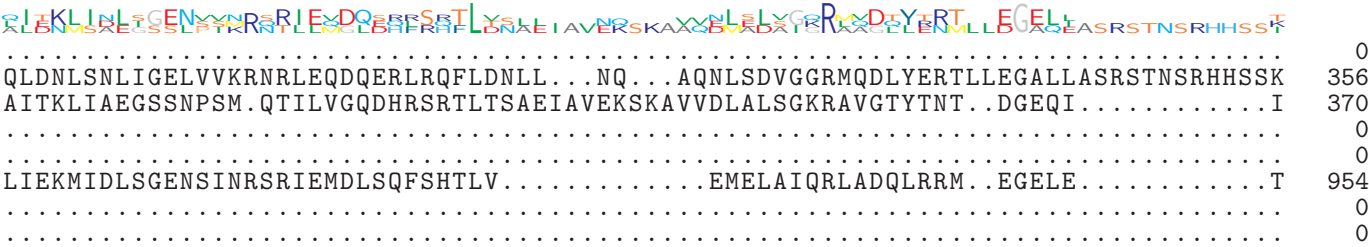
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consensus
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consensus
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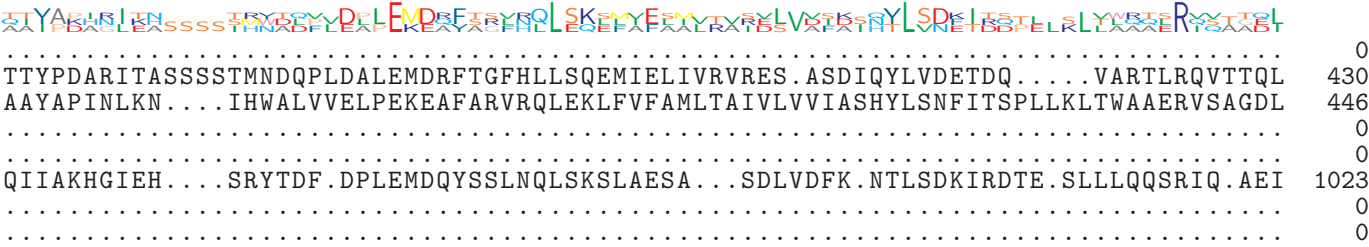
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consensus



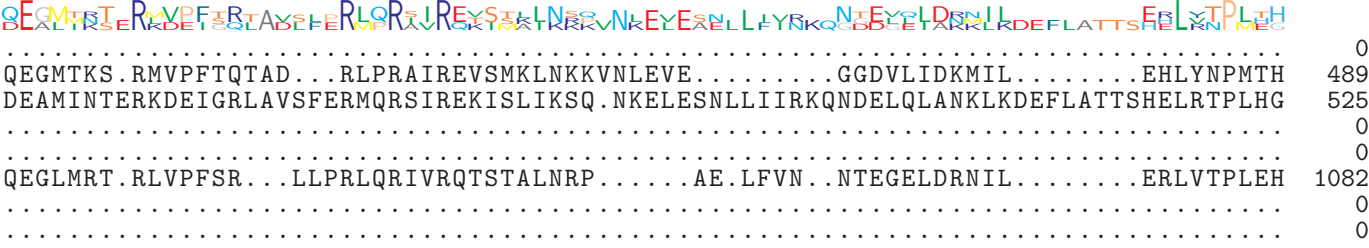
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consensus



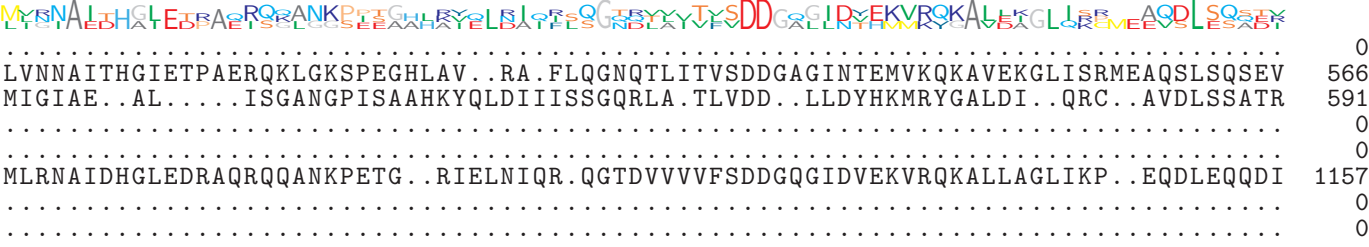
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consensus



logo

NTDB id 372 SMU RS05280 WP 002262198.1
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NTDB id 1143 A1552VC RS01965 WP 001883436.1
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NTDB id 1417 SGL RS04200 WP 014407071.1
consensus



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NTDB id 1417 SGL RS04200 WP 014407071.1		121
consensus		

- ⓧ non conserved
- ⓧ similar
- ⓧ ≥ 50% conserved