

logo

	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	
NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDLLGVGDVEPAITNPG	
NTDB id 1419 SGL RS12550 WP 010873252.1	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDLLGVGDVEPAITNPG	160
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	HSFAGADRVTPLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	
NTDB id 1419 SGL RS12550 WP 010873252.1	HSFAGADRVTPLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	240
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLIGELVVKRNRLEQDQERLRQFLDNLLNQAQ	
NTDB id 1419 SGL RS12550 WP 010873252.1	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLIGELVVKRNRLEQDQERLRQFLDNLLNQAQ	320
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHSSKTTYPDARITASSSSTMNDQPLDALEMDFRFTGFHLLSQEMI	
NTDB id 1419 SGL RS12550 WP 010873252.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHSSKTTYPDARITASSSSTMNDQPLDALEMDFRFTGFHLLSQEMI	400
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	
NTDB id 1419 SGL RS12550 WP 010873252.1	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	480
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	EHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQS	
NTDB id 1419 SGL RS12550 WP 010873252.1	EHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQS	560
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVGKGTTFITIRLPLTSLICKALCCLSDRARI	
NTDB id 1419 SGL RS12550 WP 010873252.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVGKGTTFITIRLPLTSLICKALCCLSDRARI	640
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGSVYGGKQEEEDVVRVVI IARGSSGNLL	
NTDB id 1419 SGL RS12550 WP 010873252.1	AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGSVYGGKQEEEDVVRVVI IARGSSGNLL	720
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTVNEVHMDP	
NTDB id 1419 SGL RS12550 WP 010873252.1	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTVNEVHMDP	800
NTDB id 374 SMU RS06885 WP 002262930.1	2
NTDB id 413 AAK55817.1 141..845()	2
NTDB id 1047 HON27 RS03110 WP 000389061.1MEDAFQNL	8
NTDB id 411815 GSQ19 RS07740 WP 010995103.1	2
NTDB id 1417 SGL RS04200 WP 014407071.1	2
consensus	*

