

logo

	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLGVGDVEPAITNPGL	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLGVGDVEPAITNPGL	160
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	HSFAGADRVTPLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	HSFAGADRVTPLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	240
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQ	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQ	320
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMVDRFTGFHLLSQEMI E L I V	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMVDRFTGFHLLSQEMI E L I V	400
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	480
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	EHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQS	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	EHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQS	560
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVCKGTTFTIRLPLTLSICKALCCLSDRARI	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVCKGTTFTIRLPLTLSICKALCCLSDRARI	640
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGVSYGKQEEDVVRVVIARGSSGNLL	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGVSYGKQEEDVVRVVIARGSSGNLL	720
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MP IADVLELIEIAQGRRLTDSSGGLWRKTLTPTVNEVHMDP	
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MP IADVLELIEIAQGRRLTDSSGGLWRKTLTPTVNEVHMDP	800
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 35865 RIV7116 RS03775 WP 015116941.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		



