

logo

	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	
NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
consensus		

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	AELDALFTDEDAPTTAIDPFESSWLEIDQMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDLLGVGDVEPAITNPGL	
NTDB id 1419 SGL RS12550 WP 010873252.1	AELDALFTDEDAPTTAIDPFESSWLEIDQMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDLLGVGDVEPAITNPGL	160
NTDB id 413 AAK55817.1 141..845( )	.....	0
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NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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	HSFAGADRVPTLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	
NTDB id 1419 SGL RS12550 WP 010873252.1	HSFAGADRVPTLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	240
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLIGELVVKRNRLEQDQERLRQFLDNLLNQAQ	
NTDB id 1419 SGL RS12550 WP 010873252.1	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLSNLIGELVVKRNRLEQDQERLRQFLDNLLNQAQ	320
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSKTTYPDARITASSSSTMNDQPLDALEMDFRFTGFHLLSQEMIELIV	
NTDB id 1419 SGL RS12550 WP 010873252.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSKTTYPDARITASSSSTMNDQPLDALEMDFRFTGFHLLSQEMIELIV	400
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
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NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIIL	480
NTDB id 413 AAK55817.1 141..845( )	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIIL	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	EHLYNPMTHLVNNAITHGIEPTAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEASQ	560
NTDB id 413 AAK55817.1 141..845( )	EHLYNPMTHLVNNAITHGIEPTAERQKLGKSPGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEASQ	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVGKGTFTTIRLPLTLSICKALCCLSDRARI	640
NTDB id 413 AAK55817.1 141..845( )	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVGKGTFTTIRLPLTLSICKALCCLSDRARI	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	AFPMDGVEDTRDYLDPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGSVYGGKQEEDEVVRV I IARGSSGNLL	720
NTDB id 413 AAK55817.1 141..845( )	AFPMDGVEDTRDYLDPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGSVYGGKQEEDEVVRV I IARGSSGNLL	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTMEDAFMDL	800
NTDB id 413 AAK55817.1 141..845( )	ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSI MPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTVNEVHMDP	2
NTDB id 374 SMU RS06885 WP 002262930.1	.....	2
NTDB id 1060296 ACFFRD RS11280 WP 069073698.1	.....	2
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	2
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....MEDAFQNL	8
consensus		

