

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQGEFGSPSSGEEDPPVRMTLDFNSPEAELEDA	85
NTDB id 413 AAK55817.1 141..845()	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQGEFGSPSSGEEDPPVRMTLDFNSPEAELEDA	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	LFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGHLHSFAGADRVT	170
NTDB id 413 AAK55817.1 141..845()	LFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGHLHSFAGADRVT	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	PLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEALLEQADSSLGGGIAG	255
NTDB id 413 AAK55817.1 141..845()	PLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEALLEQADSSLGGGIAG	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	MGGARLPARPRQVKSRSFQETMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEG	340
NTDB id 413 AAK55817.1 141..845()	MGGARLPARPRQVKSRSFQETMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEG	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	ALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMDRFTGFHLLSQEMI ELIVRVRESASDIQYLVDETDQVARTLRQ	425
NTDB id 413 AAK55817.1 141..845()	ALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMDRFTGFHLLSQEMI ELIVRVRESASDIQYLVDETDQVARTLRQ	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	VTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMILEHLYNPMTHLVNNAITHGIIETPAERQKLGK	510
NTDB id 413 AAK55817.1 141..845()	VTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMILEHLYNPMTHLVNNAITHGIIETPAERQKLGK	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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	SPEGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVR	595
NTDB id 413 AAK55817.1 141..845()	SPEGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVR	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	TSLIDIRGAINIESTVGKGTTFTIRLPLTLSICKALCCLSDRARIAPFMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLY	680
NTDB id 413 AAK55817.1 141..845()	TSLIDIRGAINIESTVGKGTTFTIRLPLTLSICKALCCLSDRARIAPFMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLY	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	PLKDLLNYNRQLSRGSVYGGKQEEDVVRVIIARGSSGNLLALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADV	765
NTDB id 413 AAK55817.1 141..845()	PLKDLLNYNRQLSRGSVYGGKQEEDVVRVIIARGSSGNLLALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADV	0
NTDB id 374 SMU RS06885 WP 002262930.1		0
NTDB id 271729 NPM RS24175 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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NTDB id 1419 SGL RS12550 WP 010873252.1	LELIEIAQGRLRTDSSGGLWRKTLTPTVNEVHMDPTVLIVDDSIIVRELISLSFKKAGFRVEQARDGQEAWDQIKSGLPCDLVFC	850
NTDB id 413 AAK55817.1 141..845()	LELIEIAQGRLRTDSSGGLWRKTLTPTVNEVHMDPTVLIVDDSIIVRELISLSFKKAGFRVEQARDGQEAWDQIKSGLPCDLVFC	51
NTDB id 374 SMU RS06885 WP 002262930.1		51
NTDB id 271729 NPM RS24175 WP 069073698.1		51
NTDB id 1417 SGL RS04200 WP 014407071.1		51
NTDB id 1047 HON27 RS03110 WP 000389061.1	MEDAFQNLKVMVIDDSKTIIRTAETLLQREGCEVITAVDGFLEALSKEIAEA.NPDLIVFV	57
consensus	MEDAFQNLKVMVIDDSKTIIRTAETLLQREGCEVITAVDGFLEALSKEIAEA.NPDLIVFV	

