

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQGEFGSPSSGEEDPPVRMTLDFNSPEAELEDA	85
NTDB id 374 SMU RS06885 WP 002262930.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGETVIQGEFGSPSSGEEDPPVRMTLDFNSPEAELEDA	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGHLHSFAGADRVT	170
NTDB id 374 SMU RS06885 WP 002262930.1	LFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGHLHSFAGADRVT	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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logo

NTDB id 1419 SGL RS12550 WP 010873252.1	PLTNEVLLFAAIDDDLGLINAPVLDEEFSFTELETIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEALLEQADSSLGGGIAG	255
NTDB id 374 SMU RS06885 WP 002262930.1	PLTNEVLLFAAIDDDLGLINAPVLDEEFSFTELETIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEALLEQADSSLGGGIAG	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	MGGARLPARPRQVKSRSEFQTMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEG	340
NTDB id 374 SMU RS06885 WP 002262930.1	MGGARLPARPRQVKSRSEFQTMRVPIKQLDNLSNLI GELVVKRNRLEQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEG	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	ALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMDRFTGFHLLSQEMI E L I V R V R E S A S D I Q Y L V D E T D Q V A R T L R Q	425
NTDB id 374 SMU RS06885 WP 002262930.1	ALLASRSTNSRHHSSKTTYPDARITASSSSTMNDQPLDALEMDRFTGFHLLSQEMI E L I V R V R E S A S D I Q Y L V D E T D Q V A R T L R Q	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	VTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMI LEHLYNPMTHLVNNAITHG IETPAERQKLGK	510
NTDB id 374 SMU RS06885 WP 002262930.1	VTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMI LEHLYNPMTHLVNNAITHG IETPAERQKLGK	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	SPEGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVR	595
NTDB id 374 SMU RS06885 WP 002262930.1	SPEGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVR	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	TSLIDIRGAINIESTVGKGTTFITIRLPLTLSICKALCCLSDRARI AFPMDGVEDTRDYLPKD IVTDEDGQKCVKYGKEALLLPLY	680
NTDB id 374 SMU RS06885 WP 002262930.1	TSLIDIRGAINIESTVGKGTTFITIRLPLTLSICKALCCLSDRARI AFPMDGVEDTRDYLPKD IVTDEDGQKCVKYGKEALLLPLY	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	PLKDLLNYNRQLSRGSVYGKQEEDVVRV I IARGSSGNLLALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADV	765
NTDB id 374 SMU RS06885 WP 002262930.1	PLKDLLNYNRQLSRGSVYGKQEEDVVRV I IARGSSGNLLALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADV	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 24693 AVA RS02570 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
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NTDB id 1419 SGL RS12550 WP 010873252.1	LELIEIAQGRLRRTDSSGGLWRKTLTPTVNEVHMDPTVLIVDDSI TRELISLFFKKAGFRVEQARDGQEAWDQIKSGLPCDLVFC	850
NTDB id 374 SMU RS06885 WP 002262930.1	LELIEIAQGRLRRTDSSGGLWRKTLTPTVNEVHMDPTVLIVDDSI TRELISLFFKKAGFRVEQARDGQEAWDQIKSGLPCDLVFC	51
NTDB id 413 AAK55817.1 141..845()		51
NTDB id 1047 HON27 RS03110 WP 000389061.1	MEDAFQNLKVMVIDDSKTI RR TAETLLQREGCEVITAVDGFEALSKIAEA.NPDIIVFV	57
NTDB id 24693 AVA RS02570 WP 010995103.1	MSTVLIVEDSLAQREMITDLLKASGLTVTHATDGLEALEAIKEE.PPDLVVL	51
NTDB id 1417 SGL RS04200 WP 014407071.1	MNAVLVEDSSSQREMISSGILKDHGQVTTIACDGVEALEKIQNF.SPDLVVL	51
consensus		

