

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 374 SMU RS06885 WP 002262930.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	0
NTDB id 413 AAK55817.1 141..845()		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
NTDB id 67600 NOS3756 RS25195 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	AELDALFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGL	160
NTDB id 374 SMU RS06885 WP 002262930.1	AELDALFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLQDLLGVGDVEPAITNPGL	0
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NTDB id 67600 NOS3756 RS25195 WP 010995103.1		0
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consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	HSFAGADRVTPLTNEVLLFAAIDDL DGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEA	240
NTDB id 374 SMU RS06885 WP 002262930.1	HSFAGADRVTPLTNEVLLFAAIDDL DGLINAPVLDEEFSFTELETLIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEA	0
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NTDB id 67600 NOS3756 RS25195 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	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSEFQTMRVPIKQLDNLSNLIGELVVKRNRLREQDQERLRQFLDNLLNQAQ	320
NTDB id 374 SMU RS06885 WP 002262930.1	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSEFQTMRVPIKQLDNLSNLIGELVVKRNRLREQDQERLRQFLDNLLNQAQ	0
NTDB id 413 AAK55817.1 141..845()		0
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NTDB id 67600 NOS3756 RS25195 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	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSSSTMNDQPLDALEMDRFTGFHLLSQEMI E L I V	400
NTDB id 374 SMU RS06885 WP 002262930.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSSSTMNDQPLDALEMDRFTGFHLLSQEMI E L I V	0
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NTDB id 67600 NOS3756 RS25195 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	RVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	480
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consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	EHLYNPMTHLVNNAITHGIIETPAERQKLCKSPEGHLAVRAFLQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEAQS	560
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NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAINIESTVCKGTTFTIRLPLTLSICKALCCLSDRARI	640
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NTDB id 1047 HON27 RS03110 WP 000389061.1		0
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consensus		

logo

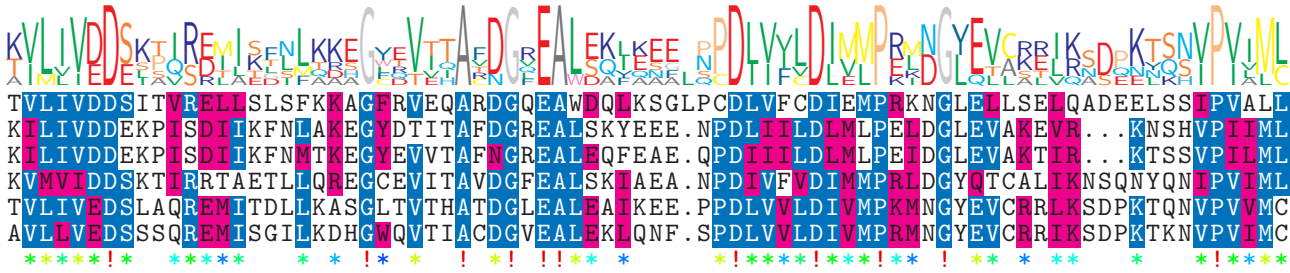
NTDB id 1419 SGL RS12550 WP 010873252.1	AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLLPLYPLKDLLNYNRQLSRGGSVYGGKQEEDVVRVIIARGSSGNLL	720
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NTDB id 67600 NOS3756 RS25195 WP 010995103.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTMEDIK	800
NTDB id 374 SMU RS06885 WP 002262930.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATVRGDGSIMPIADVLELIEIAQGRRLRTDSSGGLWRKTLTPTVNEVHMDP	2
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consensus	*	

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NTDB id 1419 SGL RS12550 WP 010873252.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 413 AAK55817.1 141..845()
NTDB id 1047 HON27 RS03110 WP 000389061.1
NTDB id 67600 NOS3756 RS25195 WP 010995103.1
NTDB id 1417 SGL RS04200 WP 014407071.1
consensus



880
78
78
87
81
81

logo

NTDB id 1419 SGL RS12550 WP 010873252.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 413 AAK55817.1 141..845()
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NTDB id 1417 SGL RS04200 WP 014407071.1
consensus



960
157
156
127
121
121

logo

NTDB id 1419 SGL RS12550 WP 010873252.1
NTDB id 374 SMU RS06885 WP 002262930.1
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NTDB id 1417 SGL RS04200 WP 014407071.1
consensus

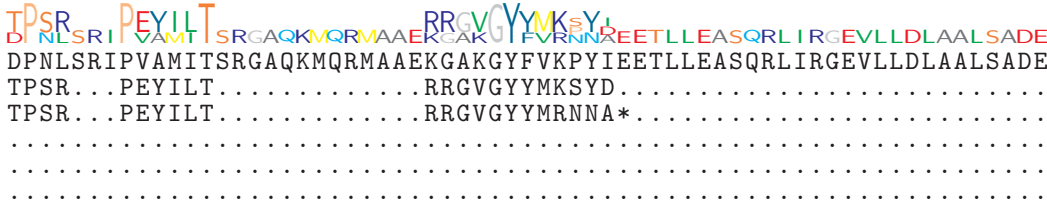


1030
213
212
127
121
121

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logo

NTDB id 1419 SGL RS12550 WP 010873252.1
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NTDB id 413 AAK55817.1 141..845()
NTDB id 1047 HON27 RS03110 WP 000389061.1
NTDB id 67600 NOS3756 RS25195 WP 010995103.1
NTDB id 1417 SGL RS04200 WP 014407071.1
consensus



1095
235
234
127
121
121

☒ non conserved
☒ similar
☒ ≥ 50% conserved