

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGEPTVIQGEFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 1047 HON27 RS03110 WP 000389061.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEGEPTVIQGEFGSPSSGEEDPPVRMTLDFNSPE	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	AELDALFTDEDAPTTAIDPFESSSWLEIDQVMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLGVDVEPAITNPGL	160
NTDB id 1047 HON27 RS03110 WP 000389061.1	AELDALFTDEDAPTTAIDPFESSSWLEIDQVMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLGVDVEPAITNPGL	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	HSFAGADRVTPLTNEVLLFAAIDDL DGLINAPVLDEEFSFTELET LIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEA	240
NTDB id 1047 HON27 RS03110 WP 000389061.1	HSFAGADRVTPLTNEVLLFAAIDDL DGLINAPVLDEEFSFTELET LIDTPAPKSFATISHPFRLPVAMEEDEDEFKDLEA	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LLEQADSS LGGGIAGMGGARLPARPRQVKSR SFEQ TMRVP I KQLDNLSNL IGELVVKRNRLEQDQERLRQFLDNLLNQAQ	320
NTDB id 1047 HON27 RS03110 WP 000389061.1	LLEQADSS LGGGIAGMGGARLPARPRQVKSR SFEQ TMRVP I KQLDNLSNL IGELVVKRNRLEQDQERLRQFLDNLLNQAQ	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSKTTYPDARITASSSSTMNDQPLDALEM DRFTGFHLLSQEMI ELIV	400
NTDB id 1047 HON27 RS03110 WP 000389061.1	NLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSKTTYPDARITASSSSTMNDQPLDALEM DRFTGFHLLSQEMI ELIV	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	RVRESASDIQYLVD ETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	480
NTDB id 1047 HON27 RS03110 WP 000389061.1	RVRESASDIQYLVD ETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMIL	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1		0
NTDB id 1417 SGL RS04200 WP 014407071.1		0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	EHLYNPMTHLVNNAITHGIIETPAERQKLCKSP	560
NTDB id 1047 HON27 RS03110 WP 000389061.1	EHLYNPMTHLVNNAITHGIIETPAERQKLCKSP	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	EHLYNPMTHLVNNAITHGIIETPAERQKLCKSP	0
NTDB id 1417 SGL RS04200 WP 014407071.1	EHLYNPMTHLVNNAITHGIIETPAERQKLCKSP	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDV	640
NTDB id 1047 HON27 RS03110 WP 000389061.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDV	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDV	0
NTDB id 1417 SGL RS04200 WP 014407071.1	LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDV	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	AFPM DGVEDTRDYLPKDIVTDEDGQKCVKYGKE	720
NTDB id 1047 HON27 RS03110 WP 000389061.1	AFPM DGVEDTRDYLPKDIVTDEDGQKCVKYGKE	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	AFPM DGVEDTRDYLPKDIVTDEDGQKCVKYGKE	0
NTDB id 1417 SGL RS04200 WP 014407071.1	AFPM DGVEDTRDYLPKDIVTDEDGQKCVKYGKE	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATV	800
NTDB id 1047 HON27 RS03110 WP 000389061.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATV	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATV	0
NTDB id 1417 SGL RS04200 WP 014407071.1	ALQIDQVLGEEEEIVIKQIEGPIPKPPGIAGATV	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQE	880
NTDB id 1047 HON27 RS03110 WP 000389061.1	TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQE	0
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQE	0
NTDB id 1417 SGL RS04200 WP 014407071.1	TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQE	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAAN	960
NTDB id 1047 HON27 RS03110 WP 000389061.1	TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAAN	6
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1	TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAAN	0
NTDB id 1417 SGL RS04200 WP 014407071.1	TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAAN	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1
NTDB id 1047 HON27 RS03110 WP 000389061.1
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1
NTDB id 1417 SGL RS04200 WP 014407071.1
consensus
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logo

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NTDB id 1419 SGL RS12550 WP 010873252.1
NTDB id 1047 HON27 RS03110 WP 000389061.1
NTDB id 54773 OSCIL6304 RS03815 WP 015147160.1
NTDB id 1417 SGL RS04200 WP 014407071.1
consensus
```



-  non conserved
-  similar
-  $\geq 50\%$ conserved