

# Supplementary Materials: A Highly Specific Holin-Mediated Mechanism Facilitates the Secretion of Lethal Toxin TcsL in *Paeniclostridium sordellii*

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**Table S1.** Bacterial isolates and plasmids used in this study.

Strain/ plasmid	Characteristics	Reference
<i>P. sordellii</i>		
ATCC9714	Bovine, myonecrosis, contains pCS1-3, <i>tcsH tcsL</i> <sup>+</sup>	[1]
DLL5002 [ <i>tcs</i> LT <sub>T</sub> ]	ATCC9714Ω <i>tcsL</i> ::TT, Em <sup>R</sup>	[2]
DLL5143 [pCS1-1·]	ATCC9714 cured of pCS1-1, <i>tcsL</i> <sup>-</sup>	This study, derived from [3].
DLL5036 [ <i>tcs</i> ETT1]	ATCC9714Ω <i>tcsE</i> ::TT, independent mutant 1, Em <sup>R</sup>	This study
DLL5037 [ <i>tcs</i> ETT2]	ATCC9714Ω <i>tcsE</i> ::TT, independent mutant 2, Em <sup>R</sup>	This study
DLL5159 [ <i>tcs</i> ETT1( <i>tcsE</i> <sup>+</sup> )]	DLL5036(pDLL101), Em <sup>R</sup> , Tm <sup>R</sup>	This study
DLL5240 [ <i>tcs</i> ETT1(V)]	DLL5036(pRPF185), Em <sup>R</sup> , Tm <sup>R</sup>	This study
<i>E. coli</i>		
MC1061/ λ <i>cI</i> <sub>857</sub> <i>Sam</i> 7	Lysogenic MC1061 carrying λ <i>cI</i> <sub>857</sub> <i>Sam</i> 7, does not encode a functional holin	[4]
MC1061/ λ <i>Cm</i> <sup>R</sup> Δ(SR)	Lysogenic MC1061 carrying λ <i>Cm</i> <sup>R</sup> Δ(SR), does not encode a functional holin or endolysin	[5]
DH5α	F- φ80lacZΔM15 Δ( <i>lacZYA-argF</i> )U169 <i>recA1 endA1 hsdR17(rK-, mK+) phoA supE44 λ-thi-1 gyrA96 relA1</i> , used for subcloning	Thermo Fisher Scientific
HB101(pVS520)	<i>recA123, Res<sup>-</sup>, Mod<sup>-</sup>, Str<sup>R</sup></i> containing pVS520, Tc <sup>R</sup> , used as conjugation donor for <i>P. sordellii</i>	[6]
<b>Plasmids</b>		
pBRQ(ΔRBS)	pJN4 derivate; λS gene with a deletion in its ribosome binding site (RBS) under the control of the late transcription regulatory (LTR) elements of phage λ	[7]
pJN5	pBR322 derivative; carries S105 under the control of the LTR elements of phage λ	[8]
pRG32	pBRQ(Δrbs) carrying <i>tcdE</i> with its own RBS	[7]
pDIA6884	pBRQ(Δrbs) carrying <i>tcsE</i> with its own RBS	This study
pDIA6883	pBRQ(Δrbs) carrying <i>tpeE</i> with its own RBS	This study
pDIA6885	pBRQ(Δrbs) carrying <i>tcnE</i> with its own RBS	This study
pDLL46	Clostridial TargeTron vector, contains RP4 and Tn916 <i>oriTs</i> and <i>lacZα</i> within retargeting region for blue white screening	[3]
pVS520	Tra <sup>+</sup> , Mob <sup>+</sup> , RP1 derivative	[6]
pRPF185	Clostridial tetracycline inducible expression vector	[9]
pDLL101	pRPF185 carrying <i>tcsE</i> with its predicted RBS under a tetracycline inducible promoter	This study

**Table S2.** Oligonucleotide primers used in PCR.

Primer	Sequence (5'–3')	Use
OBD778	GCTCTAGATGAATATAACAATATCTTTTATCAAAA	Amplify <i>tcsE</i> and predicted RBS (+) for cloning into pBRQ(ΔRBS)
OBD779	CCCAAGCTTCATTATTTTATCTATCCTCAATTTTAC	Amplify <i>tcsE</i> and predicted RBS (-) for cloning into pBRQ(ΔRBS)
OBD776	GCTCTAGACTGGATTCAAGAACTGTTAAAATTATG	Amplify <i>tpeE</i> and predicted RBS (+) for cloning into pBRQ(ΔRBS)
OBD777	CCCAAGCTTCTAATTATTATTCAATTATTTTATGTCTTC	Amplify <i>tpeE</i> and predicted RBS (-) for cloning into pBRQ(ΔRBS)
OBD780	GCTCTAGAACGGATAAACAGAAATTTAACAC	Amplify <i>tcnE</i> and predicted RBS (+) for cloning into pBRQ(ΔRBS)
OBD781	CCCAAGCTTGCAAGACTATTGTCCTGCTC	Amplify <i>tcnE</i> and predicted RBS (-) for cloning into pBRQ(ΔRBS)
JRP3867	CGAAATTAGAAACTTGCAGTCAGTAAC	TargeTron EBS universal
DLP965	AAAAAAAGCTTATAATTATCCTAAATATCCAAGCTGTGCCAGATAGGGTG	<i>tcsE</i> TargeTron IBS, 161/162 antisense
DLP966	CAGATTGTACAAATGTGGTGATAACAGATAAGTCCAAGCTATTAACTTAC- CTTCTTTGT	<i>tcsE</i> TargeTron EBS1d, 161/162 antisense
DLP967	TGAACGCAAGTTCTAATTCCGGTTATATTCCGATAGAGGAAAGTGTCT	<i>tcsE</i> TargeTron EBS2, 161/162 antisense
DLP525	AAAGAGCTCTAGGAGGCATTATGAATATAAC	Amplify <i>tcsE</i> and predicted RBS (+) for complementation - introduces SacI site, screening of mutants, generation of probe
DLP526	AAAGGATCCATTTTCATTATTTTATCTATCCTC	Amplify <i>tcsE</i> and predicted RBS (-) for complementation - introduces BamHI site, screening of mutants, generation of probe
DLP968	AAGAACTCAGCGAAACAAATGAC	Internal <i>tcsL</i> (+), RT-ddPCR
DLP969	TTACTAAACTGGTATCCCTGCTG	Internal <i>tcsL</i> (-), RT-ddPCR
DLP970	GAAGCACAAGGACCATGTACAG	Internal <i>P. sordellii rpoA</i> (+), RT-ddPCR
DLP971	CTTCCTTATTACAGATATTCCATG	Internal <i>P. sordellii rpoA</i> (-), RT-ddPCR

(+) forward primer, (-) reverse primer.

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