

Strain	Characteristics	Reference	Accession number
<i>Clostridium sordellii</i>			
ATCC9714	Reference, cause of oedema, contains pCS1-1	(1), (2)	LN679998
R28058	Isolated from infected crushed hand, Tet ^R	(2)	CEKZ00000000
7543-A	Unknown origin, pCS1-6	(3)	N/A
7508-A	Unknown origin, pCS1-7	(3)	N/A
S0804018	Equine intestinal disease, pCS1-5	This study	N/A
DLL5002 [<i>tcsLTT</i>]	ATCC9714(pDLL230), Erm ^R	(4)	N/A
DLL5078	DLL5002xR28058 transconjugant, R28058(pDLL230), Tet ^R , Erm ^R	This study	N/A
DLL5138 [<i>parBTT1</i>]	ATCC9714(pCS1-1 Ω <i>parB</i> ::TT), independent mutant 1, Erm ^R	This study	N/A
DLL5142 [<i>parBTT2</i>]	ATCC9714(pCS1-1 Ω <i>parB</i> ::TT), independent mutant 2, Erm ^R	This study	N/A
DLL5153	pCS1-1 free DLL5142, Str ^R , Rif ^R	This study	N/A
DLL5132 [<i>srtBTT1</i>]	ATCC9714(pCS1-1 Ω <i>srtB</i> ::TT), independent mutant 1, Erm ^R	This study	N/A
DLL5134 [<i>srtBTT2</i>]	ATCC9714(pCS1-1 Ω <i>srtB</i> ::TT), independent mutant 2, Erm ^R	This study	N/A
DLL5181 [<i>cstD4TT1</i>]	ATCC9714(pCS1-1 Ω <i>cstD4</i> ::TT), independent mutant 1, Erm ^R	This study	N/A
DLL5183 [<i>cstD4TT2</i>]	ATCC9714(pCS1-1 Ω <i>cstD4</i> ::TT), independent mutant 2, Erm ^R	This study	N/A
DLL5188 [<i>cstB4TT1</i>]	ATCC9714(pCS1-1 Ω <i>cstB4</i> ::TT), independent mutant 1, Erm ^R	This study	N/A
DLL5190 [<i>cstB4TT2</i>]	ATCC9714(pCS1-1 Ω <i>cstB4</i> ::TT), independent mutant 2, Erm ^R	This study	N/A
DLL5216 [<i>cstD4TT(cstD4⁺)</i>]	DLL5181(pDLL183), Erm ^R , Tm ^R	This study	N/A
DLL5217 [<i>cstD4TT(V)</i>]	DLL5181(pRPF185), Erm ^R , Tm ^R	This study	N/A
DLL5232 [<i>cstB4TT(cstB4⁺)</i>]	DLL5188(pDLL212), Erm ^R , Tm ^R	This study	N/A
DLL5230 [<i>cstB4TT(V)</i>]	DLL5188(pRPF185), Erm ^R , Tm ^R	This study	N/A
<i>C. difficile</i>			
P49	Unknown origin, contig 15 analysed	N/A	AVMN01000016
133	Human, node 23 analysed	N/A	NZ_MBGB01000023
20110869	Unknown origin, scaffold 00059 analysed	N/A	NZ_MTVT01000059

1. **Hall IC, Scott, J.P.** 1927. *Bacillus sordellii*, a Cause of Malignant Edema in Man. J Infect Dis **41**:329-335.
2. **Couchman EC, Browne HP, Dunn M, Lawley TD, Songer JG, Hall V, Petrovska L, Vidor C, Awad M, Lyras D, Fairweather NF.** 2015. *Clostridium sordellii* genome analysis reveals plasmid localized toxin genes encoded within pathogenicity loci. BMC Genomics **16**:392.
3. **Thiele TL, Stuber TP, Hauer PJ.** 2013. Detection of *Clostridium sordellii* strains expressing hemorrhagic toxin (TcsH) and implications for diagnostics and regulation of veterinary vaccines. Vaccine **31**:5082-5087.
4. **Carter GP, Awad MM, Hao Y, Thelen T, Bergin IL, Howarth PM, Seemann T, Rood JI, Aronoff DM, Lyras D.** 2011. TcsL is an essential virulence factor in *Clostridium sordellii* ATCC 9714. Infect Immun **79**:1025-1032.