

**Supplemental Table S1:** Primers used for PCR and sequencing

Gene	Function	Primer	PCR and sequencing primers (5'-3') forward/reverse	GenBank accession number	Positions	Amplicon size (bp)
<i>cTPs</i>	CTP synthase	P1967 P1968	TGGTTGAATTCCTATGCTTCTT ATAACAGCGGCTTCACTTGG	ABDQ01000000	982- 1545	564
<i>glpK</i>	Glycerol kinase	P1965 P1966	TCAATGCTTCCAGAAGTTAAACC GCAAGACCTGCAAGATAAGCA	ABDQ01000000	637- 1349	713
<i>rpoB</i>	RNA polymerase B subunit	P1985 P1986	CCTAAGGCTGAACATGATGG TTCCTGCCATCTTATCTCCAA	ACSJ01000000	2080- 2821	742
<i>gyrA</i>	DNA gyrase A subunit	P1987 P1988	TTTGATGGAGAAGAAAAGAACC TTCTTCTTGTTATAACTTCTTCTTGGGA	ACSJ01000000	442- 1087	646
<i>dnaK</i>	Chaperone protein (hsp70)	P1989 P1990	GCATCACTTGCATATGGTCTTG TTGAACTGCTGCACCCATAG	ACSJ01000000	451- 1050	600
Botulinolysin <sup>1</sup>	Pore-forming toxin	P2006 P2007	TGCAAATAGRAATCTTRTAGAAAAYAA CATTCCCAWGCAAGRCCWG	NZ_ACSJ01000007	405- 1475	1071
Novyilysin	Pore-forming toxin	P2004 P2005	AATCCAGCATATCCGATTTCA CACCATTCCCATGCAAGAC	NZ_ACSJ01000007	1141- 1478	338
C2 toxin component I	Actin ADP-ribosylating toxin	P2012 P2013	GAAGAAAAACGTTGGTTTACGA TTGGCTTTACAGATAGAGCAGAA	AB465554	73- 871	799
C2 toxin component II	Actin ADP-ribosylating toxin	P2014 P2015	AAAAGCTGTAGCATGGGATGA GGATAGGTTCCACCAGGATTT	AB465554	621- 1238	618

<sup>1</sup>Botunolysin gene is noted "tetanolysin O" in the genome annotation of NZ\_ACSJ01000007.

**Supplemental Table S2.** Genome sequences and strains used in this study. Additional strains were *C. botulinum* type C 468, *C. novyi* A ATCC17861<sup>T</sup> (AIP10062<sup>T</sup>), *C. novyi* B AIP212.86, *C. haemolyticum* ATCC9650<sup>T</sup> (AIP10052<sup>T</sup>).

	GenBank genome sequence accession number (sequence status) where available
AIP981.10	This study
<i>Clostridium botulinum</i> A ATCC 3502	AM412317 (complete)
<i>Clostridium botulinum</i> C str. Eklund	ABDQ01000000 (assembly)
<i>C. botulinum</i> type C 468	This study
<i>Clostridium botulinum</i> C/D mosaic str. BKT028387	AESB01000174 (assembly)
<i>Clostridium botulinum</i> C/D mosaic str. BKT015925	CP002410 (complete)
<i>Clostridium botulinum</i> C/D mosaic str. V891	AESCA00000000 (draft)
<i>Clostridium botulinum</i> C/D mosaic AIP26.09	This study
<i>Clostridium botulinum</i> C/D mosaic str. 6813	This study
<i>Clostridium botulinum</i> D/C mosaic str. OFD05	This study
<i>Clostridium botulinum</i> D/C mosaic str. OFD09	This study
<i>Clostridium botulinum</i> D str. 1873	AC SJ01000000 (assembly)
<i>C. haemolyticum</i> AIP10052 <sup>T</sup> (=ATCC9650 <sup>T</sup> )	This study
<i>C. novyi</i> A AIP10062 <sup>T</sup> (=ATCC17861 <sup>T</sup> )	This study
<i>Clostridium novyi</i> B AIP212.86	This study
<i>Clostridium novyi</i> NT	CP000382 (complete)
<i>Clostridium perfringens</i> str. 13	BA000016 (complete)