



# Erratum: Large-Scale Genomic Analyses and Toxinotyping of *Clostridium perfringens* Implicated in Foodborne Outbreaks in France

Frontiers Production Office\*

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**Keywords:** foodborne outbreak, *Clostridium perfringens* enterotoxin, virulence gene profiles, virulence factors, real-time PCR, whole genome sequencing, coregenome SNP

## An Erratum on

### Large-Scale Genomic Analyses and Toxinotyping of *Clostridium perfringens* Implicated in Foodborne Outbreaks in France

by Mahamat Abdelrahim, A., Radomski, N., Delannoy, S., Djellal, S., Le Négrate, M., Hadjab, K., et al. (2019). *Front. Microbiol.* 10:777. doi: 10.3389/fmicb.2019.00777

Due to an editorial error, the Data Availability statement containing the accession numbers of the sequenced strains was erroneously omitted from the article.

The publisher apologizes for the mistake.

## OPEN ACCESS

### Approved by:

Frontiers Editorial Office,  
Frontiers Media SA, Switzerland

### \*Correspondence:

Frontiers Production Office  
production.office@frontiersin.org

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## DATA AVAILABILITY

Read sequences used in this study were submitted online on the NCBI network (<https://www.ncbi.nlm.nih.gov/biosample>) under the accession numbers: SAMN09721416, SAMN09721417, SAMN09721418, SAMN09721419, SAMN09721420, SAMN09721421, SAMN09721422, SAMN09721423, SAMN09721424, SAMN09721425, SAMN09721426, SAMN09721427, SAMN09721428, SAMN09721429, SAMN09721430, SAMN09721431, SAMN09721432, SAMN09721433, SAMN09721434, SAMN09721435, SAMN09721436, SAMN09721437, SAMN09721438, SAMN09721439, SAMN09721440, SAMN09721441, SAMN09721442, SAMN09721443, SAMN09721444, SAMN09721445, SAMN09721446, SAMN09721447, SAMN09721448, SAMN09721449, SAMN09721450, SAMN09721451, SAMN09721452, SAMN09721453, SAMN09721454, SAMN09721455, SAMN09721456, SAMN09721457, SAMN09721458, SAMN09721459, SAMN09721460, SAMN09721461, SAMN09721462, SAMN09721463, SAMN09721464, SAMN09721465, SAMN09721466, SAMN09721467, SAMN09721468, SAMN09721469, SAMN09721470, SAMN09721471, SAMN09721472, SAMN09721473.

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