

Bacillus cereus strain FDAARGOS_802 chromosome, complete genome

GenBank: CP053965.1

[FASTA Graphics](#)

LOCUS CP053965 1320 bp DNA linear BCT 04-JUN-2020
DEFINITION Bacillus cereus strain FDAARGOS_802 chromosome, complete genome.
ACCESSION [CP053965](#) REGION: 2647276..2648595
VERSION CP053965.1
DBLINK BioProject: [PRJNA231221](#)
BioSample: [SAMN11056517](#)
KEYWORDS .
SOURCE Bacillus cereus
ORGANISM [Bacillus cereus](#)
Bacteria; Firmicutes; Bacilli; Bacillales; Bacillaceae; Bacillus;
Bacillus cereus group.
REFERENCE 1 (bases 1 to 1320)
AUTHORS Nelson,B., Plummer,A., Tallon,L., Sadzewicz,L., Zhao,X.,
Vavikolanu,K., Mehta,A., Aluvathingal,J., Nadendla,S., Myers,T.,
Yan,Y. and Sichtig,H.
TITLE FDA dAtabase for Regulatory Grade micrObial Sequences (FDA-ARGOS):
Supporting development and validation of Infectious Disease Dx
tests
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 1320)
AUTHORS Nelson,B., Plummer,A., Tallon,L., Sadzewicz,L., Zhao,X.,
Vavikolanu,K., Mehta,A., Aluvathingal,J., Nadendla,S., Myers,T.,
Yan,Y. and Sichtig,H.
TITLE Direct Submission
JOURNAL Submitted (26-MAY-2020) Center for Devices and Radiological Health,
US Food and Drug Administration, 10903 New Hampshire Avenue, Silver
Spring, MD 20993-0002, USA
COMMENT The annotation was added by the NCBI Prokaryotic Genome Annotation
Pipeline (PGAP). Information about PGAP can be found here:
https://www.ncbi.nlm.nih.gov/genome/annotation_prok/

##Genome-Assembly-Data-START##
Assembly Method :: Canu v. 1.4, SPAdes v. 3.11.1
Genome Coverage :: 995.39x
Sequencing Technology :: Pacbio; Illumina
##Genome-Assembly-Data-END##

##Genome-Annotation-Data-START##
Annotation Provider :: NCBI
Annotation Date :: 05/28/2020 21:50:58
Annotation Pipeline :: NCBI Prokaryotic Genome

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ORIGIN

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