Bacillus cereus ATCC 14579 strain P1 chromosome00001, whole genome shotgun sequence

NCBI Reference Sequence: NZ_JBJJSR010000001.1

FASTA Graphics

Go to:

LOCUS NZ_JBJJSR010000001 1221 bp DNA linear CON 08-JAN-2025

DEFINITION Bacillus cereus ATCC 14579 strain P1 chromosome00001, whole genome shotgun sequence.

ACCESSION NZ_JBJJSR010000001 REGION: complement(3064201..3065421)

VERSION NZ_JBJJSR010000001.1

DBLINK BioProject: PRJNA224116
BioSample: SAMN44839476
Assembly: GCF_046524075.1

KEYWORDS WGS; RefSeq.

SOURCE Bacillus cereus ATCC 14579

ORGANISM Bacillus cereus ATCC 14579
Bacteria; Bacillati; Bacillota; Bacilli; Bacillales; Bacillaceae; Bacillus; Bacillus cereus group.

REFERENCE 1 (bases 1 to 1221)

AUTHORS Harrison,S., Sparklin,R., Verratti,K. and Sozhamannan,S.

TITLE Draft Genome Assemblies of Ciprofloxacin Resistant Derivatives of Bacillus cereus

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1221)

AUTHORS Harrison,S., Sparklin,R., Verratti,K. and Sozhamannan,S.

TITLE Draft Genome Assemblies of Ciprofloxacin Resistant Derivatives of Bacillus cereus

JOURNAL Unpublished

REFERENCE 3 (bases 1 to 1221)

AUTHORS Harrison,S. and Sparklin,R.

TITLE Direct Submission

JOURNAL Submitted (24-NOV-2024) AOS QLA, Johns Hopkins Applied Physics Lab, 11100 Johns Hopkins Rd, Laurel, MD 20723, USA

REFERENCE 4 (bases 1 to 1221)

AUTHORS Harrison,S. and Sparklin,R.

TITLE Direct Submission

JOURNAL Submitted (24-NOV-2024) AOS QLA, Johns Hopkins Applied Physics Lab, 11100 Johns Hopkins Rd, Laurel, MD 20723, USA

COMMENT REFSEQ INFORMATION: The reference sequence is identical to JBJJSR010000001.1.
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/

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            Expected Final Version :: Yes
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            Annotation Software revision     :: 6.9
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            Genes (coding)                  :: 5,333
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            Genes (RNA)                     :: 150
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            complete rRNAs                  :: 13, 13, 13 (5S, 16S, 23S)
            tRNAs                           :: 106
            ncRNAs                          :: 5
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ORIGIN

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