Micromonospora sagamiensis JCM 3310 DNA, complete genome

GenBank: AP023438.1

FASTA Graphics

Go to:

LOCUS AP023438 1947 bp DNA linear BCT 11-SEP-2020
DEFINITION Micromonospora sagamiensis JCM 3310 DNA, complete genome.
ACCESSION AP023438 REGION: 3031807..3033753
VERSION AP023438.1
DBLINK BioProject: PRJDB10510
BioSample: SAMD00245352
KEYWORDS .
SOURCE Micromonospora sagamiensis
ORGANISM Micromonospora sagamiensis
Bacteria; Bacillati; Actinomycetota; Actinomycetes;
Micromonosporales; Micromonosporaceae; Micromonospora.
REFERENCE 1
AUTHORS Wu,L. and Ma,J.
TITLE The Global Catalogue of Microorganisms (GCM) 10K type strain sequencing project: providing services to taxonomists for standard genome sequencing and annotation
JOURNAL Int. J. Syst. Evol. Microbiol. 69 (4), 895-898 (2019)
PUBMED 30832757
REFERENCE 2 (bases 1 to 1947)
AUTHORS Sun,Q. and Ohkuma,M.
TITLE Direct Submission
JOURNAL Submitted (10-SEP-2020) Contact:Qinglan Sun Institute of Microbiology, Chinese Academy of Sciences, WFCC-MIRCEN World Data Centre for Microorganisms (WDCM); No. 3, yard 1, Beichen W Rd, Ao Yun Cun, Chaoyang, Beijing, Beijing 1000101, China URL :http://www.wdcm.org/
COMMENT Genome sequencing and analysis were conducted under the Global Catalogue of Microorganisms (GCM) 10K type strain sequencing project (http://gctype.wdcm.org). The genomic data are also available at http://gctype.wdcm.org/sequencing/GCM10017556

Annotated by DFAST v.1.2.7 (https://dfast.nig.ac.jp).

##Genome-Assembly-Data-START##
Assembly Method :: Canu
Genome Coverage :: 62X
Sequencing Technology :: Illumina HiSeq X Ten; PacBio Sequel II
##Genome-Assembly-Data-END##
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