

Klebsiella pneumoniae strain BSIKPN-9 chromosome, complete genome

GenBank: CP106912.1

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LOCUS CP106912 1500 bp DNA linear BCT 13-OCT-2022

DEFINITION Klebsiella pneumoniae strain BSIKPN-9 chromosome, complete genome.

ACCESSION [CP106912](#) REGION: 1948500..1949999

VERSION CP106912.1

DBLINK BioProject: [PRJNA862640](#)

BioSample: [SAMN29985365](#)

KEYWORDS .

SOURCE Klebsiella pneumoniae

ORGANISM [Klebsiella pneumoniae](#)

Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacterales; Enterobacteriaceae; Klebsiella/Raoultella group; Klebsiella.

REFERENCE 1 (bases 1 to 1500)

AUTHORS Zhang,N.

TITLE Clinical characteristics and evolution of 6 KPC-2 and NDM-5 co-producing Carbapenem-resistant Klebsiella pneumoniae strains

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 1500)

AUTHORS Zhang,N.

TITLE Direct Submission

JOURNAL Submitted (28-JUL-2022) Chinese People's Liberation Army Center for

Disease Control and Prevention, Chinese People's Liberation Army Center for Disease Control and Prevention, No. 20 Yard, East Street, Fengtai Road, Beijing 100000, Unknown or unspecified country

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 :: SPAdes v. 3.13.0

Genome Representation :: Full

Expected Final Version :: Yes

Genome Coverage :: 150X

Sequencing Technology :: PacBio

##Genome-Assembly-Data-END##

##Genome-Annotation-Data-START##


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          /locus_tag="OE131_09550"
CDS       complement(59..682)
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GO_process: GO:0006355 - regulation of DNA-templated
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ORIGIN

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