Citrobacter sp. CIP 55.13 genome assembly Citrobacter_pasteurii_55-13, contig contig_39, whole genome shotgun sequence

GenBank: CDHL01000055.1

FASTA Graphics

Go to:

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LOCUS       CDHL01000055          1662 bp    DNA     linear   BCT 04-DEC-2014
DEFINITION  Citrobacter sp. CIP 55.13 genome assembly
            Citrobacter_pasteurii_55-13, contig contig_39, whole genome shotgun
            sequence.
ACCESSION   CDHL01000055 REGION: complement(26930..28591)
VERSION     CDHL01000055.1
DBLINK      BioProject: PRJEB7595
            BioSample: SAMEA3099656
KEYWORDS    WGS.
SOURCE      Citrobacter pasteurii
  ORGANISM  Citrobacter pasteurii
            Bacteria; Pseudomonadati; Pseudomonadota; Gammaproteobacteria;
            Enterobacterales; Enterobacteriaceae; Citrobacter.
REFERENCE   1
  AUTHORS   Mornico,Damien.
  TITLE     Direct Submission
  JOURNAL   Submitted (12-NOV-2014) INSTITUT PASTEUR, Microbial Evolutionary
            Genomics, 25-28 RUE DU DOCTEUR ROUX 75015 PARIS, France
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