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Data Article

Draft genome sequence data and analysis of *Brachybacterium* sp. strain EE-P12 isolated from a laboratory-scale anaerobic reactor



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ABSTRACT

The species of the genus *Brachybacterium* belonging to the family *Dermabacteraceae* within the phylum *Actinobacteria* are gram-positive, facultatively anaerobic or aerobic, nonmotile and non-sporeforming bacteria. Cells of *Brachybacterium* spp. vary in shape from coccoid forms (stationary phase) to rods (exponential phase). *Brachybacterium* species can be isolated from numerous sources such as poultry deep litter, human gut, soil, food products. Here we describe the draft genome sequence of *Brachybacterium* sp. EE-P12 that was isolated from a laboratory-scale anaerobic digester. The genome sequencing generated 3,964,988 bp, with a G+C content of 72.2%. This draft genome data has been deposited at DDBJ/ENA/GenBank under the accession number QXCP00000000 (<https://www.ncbi.nlm.nih.gov/nuccore/QXCP00000000>).

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