

Species *Endomicrobium embiratermitis*

Etymology

[em.bi.ra.ter'mi.tis] N.L. **gen. n.** *embiratermitis*, of Embiratermes, referring to the termite host genus.

Nomenclatural type

[NCBI Assembly: GCA_009781315.1](#)^{TS}

Description

The species comprises only metagenome-assembled genomes. The species includes all bacteria with more than 95% average nucleotide identity (ANI) to the type genome. The GC content of the type genome is 39.6 mol% and the estimated genome size is 2.0 Mbp.

Classification

Bacteria » *Elusimicrobiota* » *Endomicrobiia* » *Endomicrobiales* » *Endomicrobiaceae* » *Endomicrobium* » *Endomicrobium embiratermitis*

References

Effective publication: Mies et al., 2024 [1]

Registry URL

<https://seqco.de/i:32903>

References

1. Mies et al. (2024). Genome reduction and horizontal gene transfer in the evolution of Endomicrobia—rise and fall of an intracellular symbiosis with termite gut flagellates. *mBio*. DOI:10.1128/mbio.00826-24