

Species *Endomicrobiellum mastotermitis*

Etymology

[mas.to.ter'mi.tis] **N.L. gen. n.** *mastotermitis*, of Mastotermes, referring to the termite host genus.

Nomenclatural type

[NCBI Assembly: GCA_031272445.1](#) ^{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 31.0 mol% and the estimated genome size is 0.9 Mbp.

Classification

Bacteria » *Elusimicrobiota* » *Endomicrobia* » *Endomicrobiales* » *Endomicrobiaceae* » *Endomicrobiellum* » *Endomicrobiellum mastotermitis*

References

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

Registry URL

<https://seqco.de/i:32914>

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](https://doi.org/10.1128/mbio.00826-24)