

## Species *Endomicrobiellum guadaloupense*

### Etymology

[gu.a.da.lou.pen'se] N.L. neut. adj. *guadaloupense*, of Guadalupe, referring to the geographic origin of the host genus.

### Nomenclatural type

[NCBI Assembly: GCA\\_031278895.1](#) <sup>Ts</sup>

### 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 32.5 mol% and the estimated genome size is 1.3 Mbp.

### Classification

*Bacteria* » *Elusimicrobiota* » *Endomicrobia* » *Endomicroiales* » *Endomicrobiaceae* » *Endomicrobiellum* » *Endomicrobiellum guadaloupense*

### References

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

### Registry URL

<https://seqco.de/i:32924>

## 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)