

Species *Kaerumiaba similis*

Etymology

[si.mi'lis] **L. fem. adj.** *similis*, A species related to but distinct from the existing species

Nomenclatural type

[NCBI Assembly: GCA_963744725.1](#) ^{Ts}

Description

Species established on the basis of ANI comparisons to GTDB R226 and 16S rRNA classifications to Silva 138.2. Formerly "uncultured bacterium" in Silva 138.2. Based on metabolic reconstruction, *Kaerumiaba similis* is inferred to encode the rod shape-determining protein MreB and peptidoglycan biosynthesis genes.

Classification

Bacteria » *Actinomycetota* » *Thermoleophilia* » *Solirubrobacterales* » *Maglebybacteriaceae* » *Kaerumiaba* » *Kaerumiaba similis*

References

Effective publication: Sereika et al., 2025 [1]

Registry URL

<https://seqco.de/i:53544>

References

1. Sereika et al. (2025). Genome-resolved long-read sequencing expands known microbial diversity across terrestrial habitats. *Nature Microbiology*. DOI:10.1038/s41564-025-02062-z