

## Species *Hemmediaba affinis*

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### Etymology

[af.fi'nis] **L. fem. adj.** *affinis*, A species related to but distinct from the existing species

### Nomenclatural type

[NCBI Assembly: GCA\\_963807005.1](#) <sup>Ts</sup>

### 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, *Hemmediaba affinis* is inferred to encode the rod shape-determining protein MreB and peptidoglycan biosynthesis genes.

### Classification

*Bacteria* » *Actinomycetota* » *Actinomycetes* » *Hemmediabales* » *Hemmediabaceae* »  
*Hemmediaba* » *Hemmediaba affinis*

### References

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

### Registry URL

<https://seqco.de/i:54093>

## 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](https://doi.org/10.1038/s41564-025-02062-z)