

## Species *Oerdingivens similis*<sup>Ts</sup>

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### Etymology

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

### Nomenclatural type

[NCBI Assembly: GCA\\_963720695.1](#)<sup>Ts</sup>

### Description

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

### Classification

*Bacteria* » *Actinomycetota* » *Acidimicrobiia* » *Acidimicrobiales* » *Ilumatobacteraceae* » *Oerdingivens* » *Oerdingivens similis*<sup>Ts</sup>

### References

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

### Registry URL

<https://seqco.de/i:53242>

## 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)