

## Species *Oestermarieana scandiae*<sup>Ts</sup>

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### Etymology

[scan.di'ae] **N.L. gen. n.** *scandiae*, A species found in Scandinavia

### Nomenclatural type

[NCBI Assembly: GCA\\_963737125.1](#)<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, *Oestermarieana scandiae* is inferred to encode the rod shape-determining protein MreB and peptidoglycan biosynthesis genes.

### Classification

*Bacteria* » *Chloroflexota* » *Oestermarieania* » *Oestermarieanales* » *Oestermarieaceae* » *Oestermarieana* » *Oestermarieana scandiae*<sup>Ts</sup>

### References

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

### Registry URL

<https://seqco.de/i:54595>

## 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)