

## Species *Sjoerringivita contigua*<sup>Ts</sup>

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### Etymology

[con.ti'gu.a] **L. fem. adj.** *contigua*, Close to, referring to a species related to but distinct from the existing species

### Nomenclatural type

[NCBI Assembly: GCA\\_963713925.1](#)<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, *Sjoerringivita contigua* is inferred to encode the rod shape-determining protein MreB and peptidoglycan biosynthesis genes.

### Classification

*Bacteria* » “Babelota” » “Babeliae” » “Babeliales” » “Vermiphilaceae” » *Sjoerringivita* » *Sjoerringivita contigua*<sup>Ts</sup>

### References

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

### Registry URL

<https://seqco.de/i:52966>

## 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)