

## Family *Ravnsnaesimenaceae*

---

### Etymology

[ravn.snae.si.me.na'ce.ae] **N.L. neut.** *Ravnsnaesimen*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Ravnsnaesimenaceae*, the Ravnsnaesimen family

### Nomenclatural type

Genus *Ravnsnaesimen*

### Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family SZUA-252 in GTDB R226. The description of the family is the same as for its type genus *Ravnsnaesimen*.

### Classification

*Bacteria* » "Eisenbacteria" » *Ravnsnaesimenia* » *Ravnsnaesiminales* » *Ravnsnaesimenaceae*

### References

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

### Registry URL

<https://seqco.de/i:54620>

## 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)