

Family *Oesbyanaceae*

Etymology

[oes.by.a.na'ce.ae] **N.L. fem.** *Oesbyana*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Oesbyanaceae*, the Oesbyana family

Nomenclatural type

Genus *Oesbyana*

Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family Gp7-AA8 in GTDB R226. The description of the family is the same as for its type genus *Oesbyana*.

Classification

Bacteria » *Acidobacteriota* » *Thermoanaerobaculia* » *Oesbyanales* » *Oesbyanaceae*

References

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

Registry URL

<https://seqco.de/i:54434>

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)