

Family *Vangianaceae*

Etymology

[van.gi.a.na'ce.ae] **N.L. fem.** *Vangiana*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Vangianaceae*, the Vangiana family

Nomenclatural type

Genus *Vangiana*

Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family GWA1-40-21 in GTDB R226. The description of the family is the same as for its type genus *Vangiana*.

Classification

Bacteria » *Patescibacteriota* » *Minisyncoccia* » “Zambryskiibacteriales” » *Vangianaceae*

References

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

Registry URL

<https://seqco.de/i:54720>

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)