

Genus *Vangiana*

Etymology

[van.gi.a'na] **Danish n.** *Vang*, Danish parish; **N.L. fem. n.** *Vangiana*, Microbe named after the Danish parish of Vang, near which the genome of the type species was recovered.

Nomenclatural type

Species *Vangiana lacustris*^{Ts}

Description

Genus established on the basis of phylogenomic comparisons to GTDB R226 and 16S rRNA classifications to Silva 138.2. Formerly known as genus Bog-1340 in GTDB R226. The description of the genus is the same as for its type species *Vangiana lacustris*.

Classification

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

References

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

Registry URL

<https://seqco.de/i:54684>

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