

Family *Maglebybacteriaceae*

Etymology

[mag.le.by.bac.te.ri.a'ce.ae] **N.L. neut.** *Maglebybacterium*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Maglebybacteriaceae*, the *Maglebybacterium* family

Nomenclatural type

Genus *Maglebybacterium*

Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family 70-9 in GTDB R226. The description of the family is the same as for its type species *Maglebybacterium contiguum*.

Classification

Bacteria » *Actinomycetota* » *Thermoleophilia* » *Solirubrobacterales* » *Maglebybacteriaceae*

References

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

Registry URL

<https://seqco.de/i:53534>

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