

## Family *Vindingisomataceae*

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### Etymology

[vin.din.gi.so.ma.ta'ce.ae] **N.L. neut.** *Vindingisoma*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Vindingisomataceae*, the *Vindingisoma* family

### Nomenclatural type

Genus *Vindingisoma*

### Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family F1-60-MAGs149 in GTDB R226. The description of the family is the same as for its type species *Vindingisoma humicum*.

### Classification

*Bacteria* » *Actinomycetota* » *Rubrobacteria* » *Gaiellales* » *Vindingisomataceae*

### References

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

### Registry URL

<https://seqco.de/i:54003>

## 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)