

## Family *Ghiorseimicrobiaceae*

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### Etymology

[Ghi.or.sei.mi.cro.bi.a'ce.ae] **N.L. neut. n.** *Ghiorseimicrobium*, referring to the type genus *Ghiorseimicrobium*; -*aceae*, ending to denote a family; **N.L. fem. pl. n.** *Ghiorseimicrobiaceae*, the *Ghiorseimicrobium* family

### Nomenclatural type

Genus *Ghiorseimicrobium*

### Description

This family belongs to the order *Ghiorseimicrobiales*, class *Velamenicoccia*, phylum *Omnitrophota*. The description for this monotypic family is the same as for its nomenclatural type, the genus *Ghiorseimicrobium*.

### Classification

*Bacteria* » *Omnitrophota* » “*Velamenicoccia*” » *Ghiorseimicrobiales* » *Ghiorseimicrobiaceae*

### References

Effective publication: Seymour et al., 2023 [1]

### Registry URL

<https://seqco.de/i:23761>

## References

1. Seymour et al. (2023). Hyperactive nanobacteria with host-dependent traits pervade Omnitrophota. *Nature Microbiology*. [DOI:10.1038/s41564-022-01319-1](https://doi.org/10.1038/s41564-022-01319-1)