

## Family *Velesiimonadaceae*

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### Etymology

[Ve.le.si.i.mo.na.da'ce.ae] **N.L. fem. n.** *Velesiimonas*, referring to the type genus *Velesiimonas*; *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Velesiimonadaceae*, the *Velesiimonas* family

### Nomenclatural type

Genus *Velesiimonas*

### Description

This family belongs to the order *Velesiimonadales*. As a monotypic family, the description of the family is currently the same as for the genus *Velesiimonas*, the nomenclatural type for the family.

### Classification

*Incertae sedis* (Bacteria) » *Velesiimonadales* » *Velesiimonadaceae*

### References

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

### Registry URL

<https://seqco.de/i:23738>

## 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)