

## Family *Aquitaenariimonadaceae*

---

### Etymology

[A.qui.tae.na.ri.i.mo.na.da'ce.ae] N.L. fem. n. *Aquitaenariimonas*, referring to the type genus *Aquitaenariimonas*; -*aceae*, ending to denote a family; N.L. fem. pl. n. *Aquitaenariimonadaceae*, family of the genus *Aquitaenariimonas*

### Nomenclatural type

Genus *Aquitaenariimonas*

### Description

This family belongs to the order *Aquitaenariimonadales*, in the class *Gorgyraia*, phylum *Omnitrophota*. The description for this family is currently the same as for the genus *Aquitaenariimonas*, as no other representatives for this family is available. The type for the family is the genus *Aquitaenariimonas*.

### Classification

*Bacteria* » *Omnitrophota* » “*Gorgyraia*” » *Aquitaenariimonadales* » *Aquitaenariimonadaceae*

### References

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

### Registry URL

<https://seqco.de/i:23726>

## 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)