

Family *Faaborgimorphaceae*

Etymology

[fa.a.bor.gi.mor.pha'ce.ae] **N.L. fem.** *Faaborgimorpha*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Faaborgimorphaceae*, the *Faaborgimorpha* family

Nomenclatural type

Genus *Faaborgimorpha*

Description

Family established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as family Bog-793 in GTDB R226. The description of the family is the same as for its type species *Faaborgimorpha humica*.

Classification

Bacteria » *Actinomycetota* » *Acidimicrobiia* » *Acidimicrobiales* » *Faaborgimorphaceae*

References

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

Registry URL

<https://seqco.de/i:54079>

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)