

## Family *Christiansfeldiaceae*

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### Etymology

[chris.ti.ans.fel.di.a'ce.æ] **N.L. neut.** *Christiansfeldium*, the type genus of the family; **L. fem. pl. suff.** *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Christiansfeldiaceae*, the Christiansfeldium family

### Nomenclatural type

Genus *Christiansfeldium*

### Description

Family established on the basis of phylogenomic comparisons to GTDB R226. The description of the family is the same as for its type genus *Christiansfeldium*.

### Classification

*Bacteria* » "Bacteroidota\_A" » *Skagenicola* » *Skagenicolales* » *Christiansfeldiaceae*

### References

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

### Registry URL

<https://seqco.de/i:54613>

## 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