

Genus *Allingihabitans*

Etymology

[al.lin.gi.ha.bi'tans] **Danish n.** *Oester Alling*, Danish parish; **L. masc. pres. part.** *habitans*, inhabiting; **N.L. masc. n.** *Allingihabitans*, Microbe arbitrarily named after the Danish parish of Oester Alling, near which the genome of the type species was recovered.

Nomenclatural type

Species *Allingihabitans sediminis*^{Ts}

Description

Genus established on the basis of phylogenomic comparisons to GTDB R226 and 16S rRNA classifications to Silva 138.2. Formerly known as genus DAHUXN01 in GTDB R226. The description of the genus is the same as for its type species *Allingihabitans sediminis*.

Classification

Bacteria » *Acidobacteriota* » *Terriglobia* » *Acidiferrales* » "Acidiferraceae" » *Allingihabitans*

References

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

Registry URL

<https://seqco.de/i:54798>

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