

## Order *Moltrupihabिताntales*

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### Etymology

[mol.tru.pi.ha.bi.tan.ta'les] **N.L. masc.** *Moltrupihabitans*, the type genus of the order; **L. fem. pl. suff.** *-ales*, ending to denote an order; **N.L. fem. pl. n.** *Moltrupihabिताntales*, the *Moltrupihabitans* order

### Nomenclatural type

Genus *Moltrupihabitans*

### Description

Order established on the basis of phylogenomic comparisons to GTDB R226. Formerly known as order UBA12157 in GTDB R226. The description of the order is the same as for the family *Moltrupihabिताntaceae*. The type genus is *Moltrupihabitans*.

### Classification

*Bacteria* » *Patescibacteriota* » *Moltrupihabिताntia* » *Moltrupihabिताntales*

### References

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

### Registry URL

<https://seqco.de/i:54568>

## 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)