

Species *Avelusimicrobium stercoris*

Etymology

[ster'co.ris] **L. gen. n.** *stercoris*, of excrement, referring to the isolation source

Nomenclatural type

[NCBI Assembly: GCA_002432025.1](#) ^{Ts}

Description

The species comprises only metagenome-assembled genomes. The species includes all bacteria with more than 95% average nucleotide identity (ANI) to the type genome. The GC content of the type genome is 49% and the estimated genome size is 1.7 Mbp.

Classification

Bacteria » *Elusimicrobiota* » *Elusimicrobia* » *Elusimicrobiales* » *Elusimicrobiaceae* »
Avelusimicrobium » *Avelusimicrobium stercoris*

References

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

Registry URL

<https://seqco.de/i:49142>

References

1. Mies et al. (2025). Comparative genomics of Elusimicrobiaceae (phylum Elusimicrobiota) and description of the isolates *Elusimicrobium simillimum* sp. nov., *Elusimicrobium posterum* sp. nov., and *Parelusimicrobium proximum* gen. nov. sp. nov. *Systematic and Applied Microbiology*. [DOI:10.1016/j.syapm.2025.126606](https://doi.org/10.1016/j.syapm.2025.126606)