

---

## Species *Aalborgicola bjergmarkensis*

---

### Etymology

[bjerg.mar.ken'sis] **N.L. masc. adj.** *bjergmarkensis*, of the city of Bjergmarken, from where the MAG was obtained

### Nomenclatural type

[NCBI Assembly: GCA\\_016708725.1](#) <sup>Ts</sup>

### Description

Bacterium found in activated sludge.

Genome-wide gene annotation suggested the potential for full glycolysis, pentose phosphate pathway, citric acid cycle and glyoxylate pathway.

The *gtsABC*+*malK* transporters of glucose uptake was also predicted.

Potential to reduce nitrate with narGHI. Lastly it also had nitrous oxide reductase, nosZ predicted.

### Classification

*Bacteria* » *Pseudomonadota* » *Betaproteobacteria* » *Burkholderiales* » *Burkholderiaceae* » *Aalborgicola* » *Aalborgicola bjergmarkensis*

### References

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

### Registry URL

<https://seqco.de/i:44112>

## References

1. Petersen et al. (2025). Diversity and physiology of abundant Rhodoferrax species in global wastewater treatment systems. *Systematic and Applied Microbiology*.  
[DOI:10.1016/j.syapm.2024.126574](https://doi.org/10.1016/j.syapm.2024.126574)