

## Species *Houyibacterium oceanicum*<sup>Ts</sup>

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### Etymology

[o.ce.a'ni.cum] **N.L. fem. adj.** *oceanicum*, of or pertaining to the ocean

### Nomenclatural type

[NCBI Assembly: GCA\\_031800675.1](#)<sup>Ts</sup>

### Description

The nomenclatural type for the species is the genomic assembly LLY-WYZ-15\_3 (GCA\_031800675.1). Genome is predicted to 9.41 Mb in 214 scaffolds. The GC content is 73.94%. Genome has complete bacteriochlorophyll synthesis pathways, and encodes reaction center proteins and other key enzymes, suggesting potential phototrophic lifestyle. Genome also has PR, suggesting the potential capacity using both proton-pumping and bacteriochlorophyll-based photosystems. Genomic assemblies for this species originated from seawater.

### Classification

*Bacteria* » *Myxococcota* » *Polyangia* » *Polyangiales* » *Houyibacteriaceae* » *Houyibacterium* » *Houyibacterium oceanicum*<sup>Ts</sup>

### References

Effective publication: Li et al., 2023 [1]

### Registry URL

<https://seqco.de/i:31318>

## References

1. Li et al. (2023). Globally distributed Myxococcota with photosynthesis gene clusters illuminate the origin and evolution of a potentially chimeric lifestyle. *Nature Communications*.  
[DOI:10.1038/s41467-023-42193-7](https://doi.org/10.1038/s41467-023-42193-7)