

## Species *Xiheicaenimonas phototrophica*<sup>Ts</sup>

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### Etymology

[pho.to'tro.phi.ca] **Gr. n.** *phos*, photos light; **Gr. adj.** *trophikos*, nursing, tending or feeding; **N.L. fem. adj.** *phototrophica*, referring to the likely ability to use light for energy generation

### Nomenclatural type

[NCBI Assembly: GCA\\_016793725.1](https://ncbi.nlm.nih.gov/assembly/GCA_016793725.1)<sup>Ts</sup>

### Description

The nomenclatural type for the species is the genomic assembly LLY-WYZ-6\_1 (GCA\_016793725.1). Genome is predicted to 9.75 Mb in 939 scaffolds. The GC content is 69.60%. Genome has complete bacteriochlorophyll synthesis pathways, and encodes reaction center proteins and other key enzymes, suggesting potential phototrophic lifestyle. Genome for this species originated from activated sludge.

### Classification

*Bacteria* » *Myxococcota* » *Myxococcia* » *Myxococcales* » *Myxococcaceae* » *Xiheicaenimonas* » *Xiheicaenimonas phototrophica*<sup>Ts</sup>

### References

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

### Registry URL

<https://seqco.de/i:32844>

## 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)