Species Endonucleibacter puteoserpentis^{TS}

Etymology

[pu.te.o.ser.pen'tis] **N.L. gen. masc. n.** puteoserpentis, of the deep-sea mussel *Bathymodiolus* puteoserpentis, based on the species epithet of the host

Nomenclatural type

NCBI Assembly: GCA 030674915.1 Ts

Description

A bacterium that invades the nuclei of deep-sea bathymodiolin mussels from hydrothermal vents and cold seeps. This organism was first discovered in *Bathymodiolus puteoserpentis* from the Logatchev hydrothermal vent field on the Mid-Atlantic Ridge. Fluorescence *in situ* hybridization and transmission electron microscopy analyses of the developmental cycle of *E. puteoserpentis* showed that the infection of a nucleus begins with a single rod-shaped bacterium which grows to an unseptated filament of up to 20 µm length and then divides repeatedly until the nucleus is filled with up to 80 000 bacteria. The greatly swollen nucleus destroys its host cell and the bacteria are released after the nuclear membrane bursts. Intriguingly, the only nuclei that were never infected by *E. puteoserpentis* were those of the gill bacteriocytes. These cells contain the symbiotic sulfur- and methane-oxidizing bacteria, suggesting that the mussel symbionts can protect their host nuclei against the parasite. *E. puteoserpentis* belongs to a monophyletic clade of *Gammaproteobacteria* associated with marine metazoans as diverse as sponges, corals, bivalves, gastropods, echinoderms, ascidians and fish.

Classification

Bacteria » Pseudomonadota » Gammaproteobacteria » Oceanospirillales » Endozoicomonadaceae » Endonucleibacter » Endonucleibacter puteoserpentis^{Ts}

References

Effective publication: Porras et al., 2024 [1] Assigned taxonomically: Porras et al., 2024 [1]

Registry URL

https://seqco.de/i:54836

References

1. Porras et al. (2024). An intranuclear bacterial parasite of deep-sea mussels expresses apoptosis inhibitors acquired from its host. *Nature Microbiology*. DOI:10.1038/s41564-024-01808-5