Magnimaribacter mokuoloeensis gen. nov. sp. nov.

Submitted by Freel, Kelle

Order Magnimaribacterales

Etymology

[Mag.ni.ma.ri.bac.ter.a'les] **N.L. masc. n.** *Magnimaribacter,* referring to the type genus Magnimaribacter; *-ales,* ending to denote an order; **N.L. fem. pl. n.** *Magnimaribacterales,* the Magnimaribacter order

Nomenclatural type

Genus Magnimaribacter

Description

This order was named following the isolation of a SAR86 bacterium from surface seawater of the tropical Pacific Ocean collected from Kāne'ohe Bay, O'ahu, Hawai'i. It is the first isolated member of the SAR86 clade of gammaproteobacteria, which we have named the Magnimaribacterales. The Magnimaribacterales are frequently one of the most abundant groups of marine bacteria found across the global ocean.

Classification

Bacteria » Pseudomonadota » Gammaproteobacteria » Magnimaribacterales

References

Effective publication: Ramfelt et al., 2024 [1]

Registry URL

https://seqco.de/i:44153

Family Magnimaribacteraceae

Etymology

[Mag.ni.ma.ri.bac.ter.a'ce.ae] **N.L. masc. n.** *Magnimaribacter,* referring to the type genus Magnimaribacter; *-aceae,* ending to denote a family; **N.L. fem. pl. n.** *Magnimaribacteraceae,* the Magnimaribacter family

Nomenclatural type

Genus *Magnimaribacter*

Description

This family was established following the isolation of a SAR86 bacterium from surface seawater of the tropical Pacific Ocean collected from Kāne'ohe Bay, O'ahu, Hawai'i. It is the first isolated member of the SAR86 clade of gammaproteobacteria, which we have named the Magnimaribacterales. The Magnimaribacterales are frequently one of the most abundant groups of marine bacteria found across the global ocean.

Classification

Bacteria » Pseudomonadota » Gammaproteobacteria » Magnimaribacterales » Magnimaribacteraceae

References

Effective publication: Ramfelt et al., 2024 [1]

Registry URL

https://segco.de/i:44054

Genus Magnimaribacter

Etymology

[Mag.ni.ma.ri.bac'ter] L. masc. adj. magnus, great in importance or number; L. fem. n. mare, sea; N.L. masc. n. bacter, a rod; N.L. masc. n. Magnimaribacter, an abundant microbe in the global oceans

Nomenclatural type

Species Magnimaribacter mokuoloeensis^{Ts}

Description

This type isolate of this genus was isolated from surface seawater of the tropical Pacific Ocean collected from Kāne'ohe Bay, O'ahu, Hawai'i. It is the first isolated member of the SAR86 clade of gammaproteobacteria, the order of which we have named the Magnimaribacterales. The Magnimaribacterales are frequently one of the most abundant groups of marine bacteria found across the global ocean. This isolate opens an unprecedented window toward understanding the genomic and metabolic traits that contribute to the success of this lineage.

Classification

Bacteria » Pseudomonadota » Gammaproteobacteria » Magnimaribacterales » Magnimaribacteraceae » Magnimaribacter

References

Effective publication: Ramfelt et al., 2024 [1]

Registry URL

https://seqco.de/i:44052

Species Magnimaribacter mokuoloeensis^{Ts}

Etymology

[mo.ku.o.loe.en'sis] **N.L. masc. adj.** *mokuoloeensis*, Named after Moku o Lo'e, an island in Kāne'ohe Bay, O'ahu, where the organism was discovered from

Nomenclatural type

NCBI Assembly: GCA_041860225.1 Ts

Reference Strain

Strain sci0038783: Strain HIMB1674

Description

This bacterium was isolated from surface seawater of the tropical Pacific Ocean collected from Kāne'ohe Bay, O'ahu, Hawai'i. It is the first isolated member of the SAR86 clade of gammaproteobacteria, which we have named the Magnimaribacterales. The Magnimaribacterales are frequently one of the most abundant groups of marine bacteria found across the global ocean. This isolate opens an unprecedented window toward understanding the genomic and metabolic traits that contribute to the success of this lineage.

Classification

Bacteria » Pseudomonadota » Gammaproteobacteria » Magnimaribacterales » Magnimaribacteraceae » Magnimaribacter » Magnimaribacter mokuoloeensis^{Ts}

References

Effective publication: Ramfelt et al., 2024 [1]

Registry URL

https://seqco.de/i:44051

References

1. Ramfelt et al. (2024). Isolate-anchored comparisons reveal evolutionary and functional differentiation across SAR86 marine bacteria. *The ISME Journal*. DOI:10.1093/ismejo/wrae227

Register List Certificate of Validation

On behalf of the *Committee on the Systematics of Prokaryotes Described from Sequence Data* (SeqCode Committee), we hereby certify that the Register List **seqco.de/r:pj-wgdq1** submitted by **Freel, Kelle** and including 4 new names has been successfully validated.

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