Register list for Ryujinia shimokita gen. nov. sp. nov. and their lineage

Submitted by Rinke, Chris

Phylum Ryujiniota

Etymology

[Ry.u.ji.ni.o'ta] **N.L. fem. n.** *Ryujinia*, referring to the type genus Ryujinia; *-ota*, ending to denote a phylum; **N.L. neut. pl. n.** *Ryujiniota*, the Ryujinia phylum

Nomenclatural type

Genus Ryujinia

Description

Ryujiniota phyl. nov. has a global distribution and was mainly present in public shotgun metagenome datasets labelled as marine sediment metagenome, peat metagenome, sediment metagenome, soil metagenome and groundwater metagenome. The highest relative abundances of Ryujiniota were found in Costa Rica margin subsurface and east pacific subsurface metagenomes with 15.75% and 15.33%, respectively.

Classification

Bacteria » Ryujiniota

References

Effective publication: Sun et al., 2025 [1] Assigned taxonomically: Sun et al., 2025 [1]

Registry URL

https://seqco.de/i:49691

Class Ryujiniia

Etymology

[Ry.u.ji.ni'i.a] **N.L. fem. n.** *Ryujinia*, referring to the type genus Ryujinia; *-ia*, ending to denote a class; **N.L. neut. pl. n.** *Ryujiniia*, the Ryujinia class

Nomenclatural type

Genus Ryujinia

Description

The description is the same as for the type genus Ryujinia

Classification

Bacteria » Ryujiniota » Ryujiniia

References

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

Registry URL

https://seqco.de/i:49693

Order Ryujiniales

Etymology

[Ry.u.ji.ni.a'les] **N.L. fem. n.** *Ryujinia*, referring to the type genus Ryujinia; *-ales*, ending to denote an order; **N.L. fem. pl. n.** *Ryujiniales*, the Ryujinia order

Nomenclatural type

Genus Ryujinia

Description

The description is the same as for the type genus Ryujinia

Classification

Bacteria » Ryujiniota » Ryujiniia » Ryujiniales

References

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

Registry URL

https://segco.de/i:49694

Family Ryujiniaceae

Etymology

[Ry.u.ji.ni.a'ce.ae] **N.L. fem. n.** *Ryujinia*, referring to the type genus Ryujinia; *-aceae*, ending to denote a family; **N.L. fem. pl. n.** *Ryujiniaceae*, the Ryujinia family

Nomenclatural type

Genus Ryujinia

Description

The description is the same as for the type species is Ryujinia shimokita sp. nov

Classification

Bacteria » Ryujiniota » Ryujiniia » Ryujiniales » Ryujiniaceae

References

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

Registry URL

https://seqco.de/i:49695

Genus Ryujinia

Etymology

[Ry.u.ji'ni.a] **N.L. fem. n.** Ry.u.ji'ni.a, named after Ryūjin (\square , lit. 'Dragon God') the protector deity of the sea in Japanese mythology; **N.L. fem. n.** Ryujinia, named after Ryūjin (\square , lit. 'Dragon God') the protector deity of the sea in Japanese mythology

Nomenclatural type

Species Ryujinia shimokita^{Ts}

Description

The description is the same as for the type species is Ryujinia shimokita sp. nov

Classification

Bacteria » Ryujiniota » Ryujiniia » Ryujiniales » Ryujiniaceae » Ryujinia

References

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

Registry URL

https://seqco.de/i:49696

Species Ryujinia shimokita^{Ts}

Etymology

[shi.mo.ki'ta] **N.L. fem. n.** *shimokita*, named after the sampling site, the international waters off the Shimokita Peninsula in Japan

Nomenclatural type

INSDC Nucleotide: BAAGKY000000000.1 Ts

Description

This uncultured species is represented by the genome 'SP_28H5_5', whose genome size is 2.0 MB, with the presence of 23S, 16S and 5S rRNA genes. *R. shimokita* is inferred to have a chemolithotrophic lifestyle, since this species encodes a nearly complete Wood-Ljungdahl (WL) pathway. It also has the potential for a heterotrophic lifestyle with substrate-level phosphorylation through glycolysis, while it lacks most genes for the citrate cycle and beta-oxidation. The type genome does not encode the ability for the oxidoreduction of nitrogen or sulphur compounds.

Classification

Bacteria » Ryujiniota » Ryujinia » Ryujiniales » Ryujiniaceae » Ryujinia » Ryujinia shimokita^{Ts}

References

Effective publication: Sun et al., 2025 [1] Assigned taxonomically: Sun et al., 2025 [1]

Registry URL

https://segco.de/i:49692

References

 Sun et al. (2025). Metagenomic insights into taxonomic and functional patterns in shallow coastal and deep subseafloor sediments in the Western Pacific. *Microbial Genomics*. DOI:10.1099/mgen.0.001351

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:zf1yjhk0** submitted by **Rinke, Chris** and including 6 new names has been successfully validated.

Date of Priority: 2025-09-17 07:39 UTC **DOI:** 10.57973/seqcode.r:zf1yjhk0

