Mycoplana subbaraonis

Submitted by Van Lill, Melandre

Species Mycoplana subbaraonis

Etymology

[sub.ba.ra.o'nis] **N.L. gen. masc. n.** *subbaraonis*, of Subba Rao, named after Professor N. S. Subba Rao, an eminent microbiologist who significantly contributed to the knowledge of Rhizobium biofertilizers in India

Nomenclatural type

NCBI Assembly: GCF 900220975.1 Ts

Reference Strain

Strain sc|0039684: JC85 = DSM 24765 = KCTC 23614

Description

Forms yellow coloured colonies on nutrient agar. Cells are rod-shaped, $0.8-1.2~\mu m$ in diameter. Gram-stain-negative. Oxidase- and catalase-positive. Growth mode is chemoorganoheterotrophic (respiration/fermentation). Good growth occurs after 48 h of incubation on nutrient agar at 30 °C. Growth occurs between pH 6.0–7.5 (optimum 7.0). NaCl is not required and tolerated up to 3 % (w/v). Utilises a number of carbon sources for growth. Good carbon sources for growth are d-fructose, α -d-glucose, d-sorbitol, glycerol, malate, succinate and glutamate. Major fatty acid is C18: $1\omega7c$ with minor amounts of C19: 0 cyclo $\omega8c$, C16: 0, C12: 0 and C19: $1\omega6c$. Q-10 is the major quinone.

The type strain is JC85T (= DSM 24765T = KCTC 23614T). The type strain was isolated as an endolith from a beach sand sample collected from Diamond Harbour, Kolkata, West Bengal, India. The DNA G+C content of the type strain is 67.7 mol%.

Classification

Bacteria » Pseudomonadota » Alphaproteobacteria » Hyphomicrobiales » Rhizobiaceae » Mycoplana » Mycoplana subbaraonis

References

Effective publication: Hördt et al., 2020 [1]

Original (not valid) publication: Ramana et al., 2013 [2]

Registry URL

https://seqco.de/i:50861

References

- 1. Hördt et al. (2020). Analysis of 1,000+ Type-Strain Genomes Substantially Improves Taxonomic Classification of Alphaproteobacteria. *Frontiers in Microbiology*. DOI:10.3389/fmicb.2020.00468
- 2. Ramana et al. (2013). Rhizobium subbaraonis sp. nov., an endolithic bacterium isolated from beach sand. *International Journal of Systematic and Evolutionary Microbiology*. DOI:10.1099/ijs.0.041442-0

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:igm2z748** submitted by **Van Lill, Melandre** and including 1 new name has been successfully validated.

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