





# INTERNATIONAL JOURNAL OF SYSTEMATIC AND EVOLUTIONARY MICROBIOLOGY

Volume 73, Issue 11

## Paenibacillus polygoni sp. nov., an endophytic bacterium isolated from Polygonum lapathifolium L. in wetland 8

Yong Long<sup>1</sup>, Yansheng Li<sup>1</sup>, Guanghua Wang<sup>1</sup>, Jian Jin<sup>1</sup>, Mengfan Mao<sup>1</sup>, Lizheng Gao<sup>1</sup>, Guihua Liu<sup>2</sup>, Guoquan Fan<sup>3</sup>, Zhenhua Yu<sup>1</sup>

View Affiliations

Published: 28 November 2023

#### Most read this month

A taxonomic note on the genus *Lactobacillus*: Description of 23 novel genera, emended description of the genus *Lactobacillus* Beijerinck 1901, and union of *Lactobacillaceae* and *Leuconostocaceae* 

Jinshui Zheng, Stijn Wittouck, Elisa Salvetti, Charles M.A.P. Franz, Hugh M.B. Harris, Paola Mattarelli, Paul W. O'Toole, Bruno Pot, Peter Vandamme, Jens Walter, Koichi Watanabe, Sander Wuyts, Giovanna E. Felis, Michael G. Gänzle and Sarah Lebeer

pp.: 2782-2858 (77)

#### Valid publication of the names of forty-two phyla of prokaryotes

Aharon Oren and George M. Garrity

#### International Code of Nomenclature of Prokaryotes. Prokaryotic Code (2022 Revision)

Aharon Oren, David R. Arahal, Markus Göker, Edward R. B. Moore, Ramon Rossello-Mora and Iain C. Sutcliffe Editors:

Robust demarcation of 17 distinct *Bacillus* species clades, proposed as novel *Bacillaceae* genera, by phylogenomics and comparative genomic analyses: description of *Robertmurraya kyonggiensis* sp. nov. and proposal for an emended genus *Bacillus* limiting it only to the members of the Subtilis and Cereus clades of species

Radhey S. Gupta, Sudip Patel, Navneet Saini and Shu Chen pp.: 5753–5798 (46)

Valid publication of names of two domains and seven kingdoms of prokaryotes

Markus Göker and Aharon Oren

第1页 共3页 2024-2-20 10: 35

A Gram-stain-positive, aerobic, rod-shaped, non-motile, yellowish and glossy strain, C31<sup>T</sup>, was isolated from a wetland plant *Polygonum lapathifolium* L. located south of Poyang Lake, Jiangxi Province, PR China. Phylogenetic analysis based on 16S rRNA gene sequences indicated that strain C31<sup>T</sup> showed similarity values of lower than 97.0% to other type species belonging to the genus *Paenibacillus*. The genomic average nucleotide identity values between strain C31<sup>T</sup> and its reference type species ranged from 68.9–70.9% and the digital DNA–DNA hybridization values were lower than 27.8%. The genomic DNA G+C content of strain C31<sup>T</sup> was 41.9 mol%. The optimal growth temperature, pH and NaCl concentration were 37°C, pH 7 and 0.5%, respectively. The major cellular fatty acids (>5.0%) of strain C31<sup>T</sup> were anteiso-C<sub>15:0</sub> (73.7%), anteiso-C<sub>17:0</sub> (8.4%) and iso-C<sub>15:0</sub> (5.2%). The polar lipids of strain C31<sup>T</sup> were diphosphatidylglycerol, phosphatidylglycerol, phosphatidylethanolamine and unidentified phospholipids. The respiratory quinone was MK-7. Based on these phylogenetic and phenotypic characterizations, strain C31<sup>T</sup> represents a novel species within the genus *Paenibacillus*. Therefore, the proposed name for this newly identified species is *Paenibacillus polygoni* sp. nov. and the type strain is C31<sup>T</sup> (=CCTCC AB 2022349<sup>T</sup>=KCTC 43565<sup>T</sup>).

Keyword(s): 16S rRNA gene, ANI, Paenibacillus, Polygonum lapathifolium and polyphasic taxonomy

#### **Funding**

This study was supported by the:

- National Key R&D Program of China (Award 2021YFD1500803)
  - Principle Award Recipient: YanshengLi
- Jilin Province-CAS Special Program for High-Tech Industrialization in Science and Technology Cooperation (Award 2023SYHZ0046)
  - Principle Award Recipient: ZhenhuaYu
- Strategic Priority Research Program of the Chinese Academy of Sciences (Award XDA28100200)
  - o Principle Award Recipient: YanshengLi

© 2023 The Authors

### 

#### olimination Introducing EzBioCloud: a taxonomically united database of 16S rRNA gene sequences and wholegenome assemblies

Seok-Hwan Yoon, Sung-Min Ha, Soonjae Kwon, Jeongmin Lim, Yeseul Kim, Hyungseok Seo and Jongsik Chun

ODNA-DNA hybridization values and their relationship to whole-genome sequence similarities

Johan Goris, Konstantinos T. Konstantinidis, Joel A. Klappenbach, Tom Coenye, Peter Vandamme and James M Tiedje

A taxonomic note on the genus *Lactobacillus*: Description of 23 novel genera, emended description of the genus *Lactobacillus* Beijerinck 1901, and union of *Lactobacillaceae* and *Leuconostocaceae* 

Jinshui Zheng, Stijn Wittouck, Elisa Salvetti, Charles M.A.P. Franz, Hugh M.B. Harris, Paola Mattarelli, Paul W. O'Toole, Bruno Pot, Peter Vandamme, Jens Walter, Koichi Watanabe, Sander Wuyts, Giovanna E. Felis, Michael G. Gänzle and Sarah Lebeer

第2页 共3页 2024-2-20 10: 35

Paeni bacillus polygoni sp. nov., an endophytic bacte... <a href="https://www.microbiologyresearch.org/content/journal">https://www.microbiologyresearch.org/content/journal</a>...

Proposed minimal standards for the use of genome data for the taxonomy of prokaryotes

Jongsik Chun, Aharon Oren, Antonio Ventosa, Henrik Christensen, David Ruiz Arahal, Milton S. da Costa,

Alejandro P. Rooney, Hana Yi, Xue-Wei Xu, Sofie De Meyer and Martha E. Trujillo



**3 OrthoANI: An improved algorithm and software for calculating average nucleotide identity** Imchang Lee, Yeong Ouk Kim, Sang-Cheol Park and Jongsik Chun



+ More

第3页 共3页 2024-2-20 10:35