

High diversity of chickpea Mesorhizobium species isolated in a Portuguese agricultural region



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Author(s): Laranjo M, Machado J, Young JPW, Oliveira S

Source: FEMS MICROBIOLOGY ECOLOGY **Volume:** 48 **Issue:** 1 **Pages:** 101-107 **Published:** APR 1 2004

Times Cited: 13 **References:** 46 [Citation Map](#)

Abstract: Chickpea rhizobia isolated from Portuguese soils were assigned to the genus Mesorhizobium by 16S-rDNA sequencing. High species diversity was found within populations of an agricultural region in the south of Portugal. Besides the expected Mesorhizobium ciceri and M. mediterraneum, some isolates were close to M. loti or M. tianshanense and some formed a clade that may represent a new species. A new PCR-based approach, named direct amplified polymorphic DNA (DAPD) analysis, supported the 16S-based phylogeny. This suggests that this method could be used as a molecular tool to assess genetic relationships. Evaluation of genetic diversity by 16S-rDNA sequence, DAPD and protein profiles showed different levels of heterogeneity in natural populations. 2004 Federation of European Microbiological Societies. Published by Elsevier B.V. All rights reserved.

Document Type: Article

Language: English

Author Keywords: Mesorhizobium; 16S-rDNA; DAPD; chickpea; diversity; microbial

KeyWords Plus: CICER-ARIETINUM-L; POLYMERASE-CHAIN-REACTION; SP-NOV; GENETIC DIVERSITY; RHIZOBIUM-MEDITERRANEUM; RNA GENE; STRAINS; SOILS; LOTI; TIANSHANENSE

Reprint Address: Oliveira, S (reprint author), Univ Evora, ICAM, Lab Microbiol Solo, Evora, Portugal

Addresses:

1. Univ Evora, ICAM, Lab Microbiol Solo, Evora, Portugal
2. Univ Evora, Dept Biol, P-7002554 Evora, Portugal
3. Inst Nacl Saude Dr Ricardo Jorge, Bacteriol Lab, Lisbon, Portugal
4. Univ York, Dept Biol, York YO10 5DD, N Yorkshire England

E-mail Addresses: ismo@uevora.pt

Publisher: ELSEVIER SCIENCE BV, PO BOX 211, 1000 AE AMSTERDAM, NETHERLANDS