Streptomyces smaragdinus sp. nov., isolated from the gut of the fungus growing-termite Macrotermes natalensis.

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Projects

Investigation of secondary metabolites from insect-associated microbes and their contribution to insect homeostasis and defense

Details

Metabolomic and transcriptomic analysis of the defensive role of Actinobacteria within the fungus-growing termite system

Details

Abstract

The taxonomic position of a novel aerobic, Gram-positive actinobacteria, designated strain RB5T, was determined using a polyphasic approach. The strain, isolated from the gut of the fungus-farming termite Macrotermes natalensis, showed morphological, physiological and chemotaxonomic properties typical of the genus Streptomyces. Based on 16S rRNA gene sequence analysis, the closest phylogenetic neighbour of RB5T was Streptomyces polyrhachis DSM 42102T (98.87 %). DNA-DNA hybridization experiments between strain RB5T and S. polyrhachis DSM 42102T resulted in a value of 27.4 % (26.8 %). The cell wall of strain RB5T contained ll-diaminopimelic acid as the diagnostic amino acid. Mycolic acids and diagnostic sugars in whole-cell hydrolysates were not detected. The strain produced the following major phospholipids: diphosphatidylglycerol, phosphatidylethanolamine, phosphatidylinositol, phosphatidylinositol-mannoside and phosphatidylserine. The menaquinone profile showed hexa- and octahydrogenated menaquinones containing nine isoprene units [MK-9(H6) and MK-9(H8)]. The strain exhibited a fatty acid profile containing the following major fatty acids: 12-methyltridecanoic acid (iso-C14 : 0) 12-methyltetradecanoic acid (anteiso-C15 : 0), 13-methyltetradecanoic acid (iso-C15 : 0) and 14-methylpentadecanoic acid (iso-C16 : 0). Here, we propose a novel species of the genus Streptomyces - Streptomyces smaragdinus with the type strain RB5T (=VKM Ac-2839T=NRRL B65539T).

Identifier

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