

Microbulbifer taiwanensis sp. nov., isolated from coastal soil.

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Details



Abstract

A Gram-negative, non-spore-forming rod (CC-LN1-12(T)) was isolated from coastal soil samples of Lutao Island (Green Island), Taiwan, and its taxonomic position was studied. 16S rRNA gene sequence analysis showed that isolate CC-LN1-12(T) was grouped into the Microbulbifer cluster, with the highest similarities to *Microbulbifer okinawensis* ABABA23(T) (97.9 %), *Microbulbifer maritimus* TF-17(T) (97.7 %) and *Microbulbifer donghaiensis* CN85(T) (97.7 %), similarities to all other species of the genus *Microbulbifer* were lower than 96.8 %. The polyamine pattern contained the major compounds spermidine and cadaverine. The fatty acid profile, comprising the major fatty acids iso-C(15 : 0), iso-C(17 : 1) ω 9c, C(18 : 1) ω 7c and iso-C(11 : 0) 3-OH as the major hydroxylated fatty acid, supported the affiliation of strain CC-LN1-12(T) to the genus *Microbulbifer*. DNA-DNA hybridizations between strain CC-LN1-12(T) and *Microbulbifer okinawensis* ABABA23(T), *M. donghaiensis* CN85(T) and *M. maritimus* JCM 12187(T) resulted in relatedness values of 21.5 % (14.3 %, reciprocal analysis), 35.9 % (48.5 %, reciprocal analysis) and 48.1 % (52.1 %, reciprocal analysis), respectively. From these data, as well as from physiological and biochemical tests, strain CC-LN1-12(T) could be clearly differentiated from the most closely

related species of the genus *Microbulbifer*. It is concluded that strain CC-LN1-12(T) represents a novel species, for which the name *Microbulbifer taiwanensis* sp. nov. is proposed. The type strain is CC-LN1-12(T) (= LMG 26125(T) = CCM 7856(T)).

Beteiligte Forschungseinheiten

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