

Halophibacterium profundimaris gen. nov. sp. nov., a novel short alkane-degrading bacterium isolated from deep sea water of Indian Ocean

Qiliang Lai, Yuanyuan Fu, Liping Wang, Fengqin Sun, Fuying Li and Zongze Shao*

(Key Laboratory of Marine Biogenetic Resources, Third Institute of Oceanography, State Oceanic Administration, People's Republic of China)

*Corresponding author: Zongze Shao. Tel: +86-592-2195390. Fax: +86-592-2085376. E-mail: shaozz@163.com)

Abstract: A taxonomic study was carried out on strain PC39^T, which was isolated from a PAHs-degrading consortium, enriched from a deep seawater sample collected from the Indian Ocean. The isolate was Gram-negative, short rods, no-mobile. Growth was observed at salinities from 1 to 36 ‰ and at temperatures from 4 to 40°C, and it was able to degrade Tween 80 but no gelatin. Phylogenetic analysis of the 16S rDNA sequence reveals that strain PC39^T forms a novel lineage close to the genus *Salinisphaera*, with the highest similarity to '*Salinisphaera sedimenticola*' T31B1^T (92.7 ‰), followed by *S. dokdonensis* CL-ES53^T (91.8 ‰), *Salinisphaera shabanensis* E1L3A^T (91.7 ‰) and *S. hydrothermalis* EPR70^T (91.2 ‰), all other species had sequence similarity below 89 ‰. The dominant fatty acids were C_{18:1}ω₉c (30.52 ‰), C_{18:0} (18.91 ‰), C_{19:0}ω₈c cyclo (13.57 ‰), Sum Feature 8 (C_{18:1}ω₇c/ω₆c) (10.35 ‰) C_{16:0} (8.28 ‰), Sum Feature 7 (C_{19:1}ω₆c and/or ECL 18.846 and/or C_{19:0}ω₁₀c cyclo) (8.01 ‰), and. The G+C content of the chromosomal DNA was 68.2 mol%. Phylogenetic analysis, as well as physiological and biochemical tests, showed that strain PC39^T was different from all members of the genus *Salinisphaera*. Strain PC39^T therefore represents a novel genus, for which the name *Halophibacterium profundimaris* gen. nov. sp. nov. is proposed (type strain PC39^T=CCTCC AB 2010013^T=LMG 25545^T=MCCC 1A03214^T).

key word: *Halophibacterium profundimaris*; alkane; deep sea