# 'Clostridium mediterraneense,' a new bacterial species isolated from the human gut

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#### Abstract

We report the main characteristics of 'Clostridium mediterraneense' sp. nov., strain Marseille-P2434<sup>T</sup> (CSUR P2434), a new species within the genus Clostridium. This strain was isolated from the gut microbiota of a 66-year-old diabetic patient in Marseille, France. © 2017 The Authors. Published by Elsevier Ltd.

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In November 2015, a 66-year-old diabetic patient was hospitalized in Timone Hospital, Marseille, France. He was diagnosed a malignant haemopathy. As part of the culturomics study [1], while the patient was asymptomatic, we isolated from his stool sample a bacterial strain that could not be identified by our systematic matrix-assisted laser desorption/ionization time-offlight mass spectrometry (MALDI-TOF MS) screening on a MicroFlex spectrometer (Bruker Daltonics, Bremen, Germany) [2]. The study was approved by the Institut Fédératif de Recherche 48 (agreement 09-022, Marseille, France), and the patient's consent was obtained.

The culture of strain Marseille-P2434<sup>T</sup> was achieved by preincubation of the stool specimen in an anaerobic blood culture vial (Becton Dickinson, Le Pont-de-Claix, France) at 37° C supplemented of 5% sheep's blood and 5 mL filter-sterilized rumen for 5 days. Isolated colonies were obtained by subculture on 5% sheep's blood–enriched Columbia agar (bio-Mérieux, Marcy l'Etoile, France) at 37°C in anaerobic atmosphere using AnaeroGen (bioMérieux) after 48 hours. The

colonies were circular, beige and nonhaemolytic with a diameter of about 1.0 mm after 3 days of incubation on 5% sheep's blood agar. The strain Marseille-P2434<sup>T</sup> is a strict anaerobic, motile and Gram-positive bacterium, rod shaped (0.6–0.8  $\mu$ m × 3.5–5.5  $\mu$ m) and spore forming. Catalase and oxidase were negative. A molecular identification was performed by sequencing the complete 16S rRNA gene using a 3130-XL sequencer (Applied Biosciences, Saint Aubin, France) and the fDI-rP2 primers as previously described [3]. The obtained sequence was 95.2% similar to the 16S rRNA gene sequence of Clostridium perfringens ATCC 13124 (GenBank accession no. CP000246), the phylogenetically closest species with standing in nomenclature (Fig. 1).

According to the 16S rRNA gene sequence similarity for species demarcation of prokaryotes [4], we propose that strain Marseille-P2434<sup>T</sup> is representative of a new species within the *Clostridium* genus, for which we propose the name *'Clostridium mediterraneense*' sp. nov. (me.di.ter.rane.ense, L. fem. adj. *mediterraneense*, for the Mediterranean Sea, in front of which the city of Marseille is located, where strain Marseille-P2434<sup>T</sup> was isolated).

# **MALDI-TOF MS** spectrum

The MALDI-TOF MS spectrum of 'C. mediterraneense' is available online (http://www.mediterraneeinfection.com/article.php? laref=256&titre=urms-database).

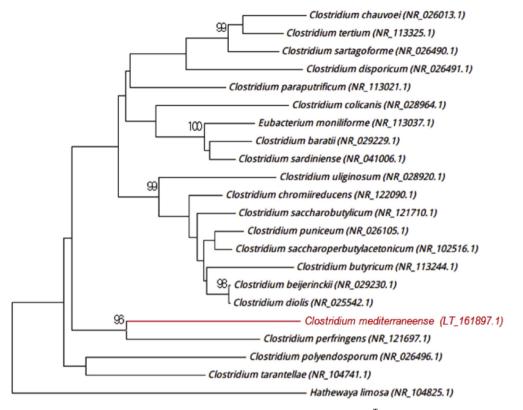


FIG. 1. Phylogenetic tree highlighting position of '*Clostridium mediterraneense*' strain Marseille-P2434<sup>T</sup> (red) relative to other phylogenetically close members of family *Clostridiaceae*. Numbers at nodes are percentages of bootstrap values obtained by repeating analysis to generate majority consensus tree 500 times. Only values >95% are displayed. Scale bar represents 2% nucleotide sequence divergence.

### Nucleotide sequence accession number

The 16S rRNA gene sequence was deposited in GenBank under accession number LT161897.

# Deposit in a culture collection

Strain Marseille-P2434<sup>T</sup> was deposited in the Collection de Souches de l'Unité des Rickettsies (CSUR, WDCM 875) under number P2434.

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# **Conflict of Interest**

None declared.

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