CORRECTION

Correction to: Description of *Mediterraneibacter massiliensis*, gen. nov., sp. nov., a new genus isolated from the gut microbiota of an obese patient and reclassification of *Ruminococcus faecis*, *Ruminococcus lactaris*, *Ruminococcus torques*, *Ruminococcus gnavus* and *Clostridium glycyrrhizinilyticum* as *Mediterraneibacter faecis* comb. nov., *Mediterraneibacter lactaris* comb. nov., *Mediterraneibacter torques* comb. nov., *Mediterraneibacter gnavus* comb. nov. and *Mediterraneibacter glycyrrhizinilyticus* comb. nov.

Amadou Hamidou Togo · Awa Diop · Fadi Bittar · Marie Maraninchi · René Valero · Nicholas Armstrong · Grégory Dubourg · Noémie Labas · Magali Richez · Jeremy Delerce · Anthony Levasseur · Pierre-Edouard Fournier · Didier Raoult · Matthieu Million

Published online: 28 September 2018
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Correction to: Antonie van Leeuwenhoek
<https://doi.org/10.1007/s10482-018-1104-y>

Subsequent to the publication of the above article, it has been noticed that the designation of the type strain is not correct. The strain referred to throughout the article as strain AT7^T should be designated as strain

Marseille-P2086^T (= CSUR P2086^T = DSM 100837^T). The corrected protologue for the species *Mediterraneibacter massiliensis*, represented by strain Marseille-P2086^T as type strain, is given below.

The original article can be found online at
<https://doi.org/10.1007/s10482-018-1104-y>.

A. H. Togo · N. Armstrong · G. Dubourg · M. Richez · J. Delerce · A. Levasseur · D. Raoult · M. Million (✉)
Aix Marseille Univ, IRD, MEPHI, IHU-Méditerranée Infection, 19-21 Boulevard Jean Moulin, 13005 Marseille, France
e-mail: matthieumillion@gmail.com

A. Diop · F. Bittar · N. Labas · P.-E. Fournier
Aix Marseille Univ, IRD, VITROME, IHU-Méditerranée Infection, 19-21 Boulevard Jean Moulin, 13005 Marseille, France

M. Maraninchi · R. Valero
NORT “Nutrition, Obesity and Risk of Thrombosis”, INSERM1062, INRA1260, Aix Marseille Université, 13385 Marseille, France

R. Valero
CHU Hôpital de la Conception, Service Nutrition, Maladies Métaboliques et Endocrinologie, APHM, 13385 Marseille, France

Description of *Mediterraneibacter massiliensis* sp. nov.

Mediterraneibacter massiliensis (mas.si.li.en'sis. L. masc. adj. *massiliensis*, of Massilia, the Latin name for Marseille).

In addition to the characteristics in the genus description, cells are coccobacillary-shaped, with a width ranging from 0.2 to 0.4 μm and a length ranging from 1 to 1.4 μm . Colonies are translucent with a diameter of 0.5–1 mm on 5% sheep blood Columbia agar. Oxidase negative. Optimum growth temperature is 37 °C under anaerobic conditions and pH tolerance ranges from 6.5 to 8.5. The major fatty acids are C_{16:0} and C_{18:1n9}. The major end product of carbohydrate

metabolism also includes isohexanoic acid and isobutanoic acid. The draft genome of the type strain is 3,069,882 bp long with a DNA G + C content of 42.4%.

The type strain Marseille-P2086^T has been deposited in the CSUR and DSMZ culture collections under numbers CSUR P2086 and DSM 100837, respectively. The type strain was isolated from the stool sample of a 37-year-old obese French woman. The draft genome and 16S rRNA sequences of the type strain have been deposited in EMBL-EBI under accession numbers FAVJ000000000 and LN881607, respectively.