

## “*Collinsella bouchedurhonensis*” sp. nov., identified in human stool sample

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### Abstract

We report the main characteristics of “*Collinsella bouchedurhonensis*” strain Marseille-P3296, which was isolated from a stool sample of a healthy 50-year-old pygmy (Baka) woman.

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**Keywords:** “*Collinsella bouchedurhonensis*”, culturomics, emerging bacteria, gut microbiota, human microbiota

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We obtained approval for the study from the ethics committee of the Institut Fédératif de Recherches 48 (no. 09-022) before we initiated the project. Samples were collected in Congo for analysis as part of the project describing the human microbiome by culturomics [1].

First, a stool sample from a 50-year-old pygmy (Baka) woman was diluted with 1 mL phosphate-buffered saline and incubated in a blood culture bottle containing an extra 5 mL of filtered rumen and sheep’s blood. The culture bottle was incubated at 37°C for 30 days under anaerobic conditions. At day 10, a “*Collinsella bouchedurhonensis*” colony was isolated on 5% sheep’s blood–enriched Columbia agar (bioMérieux, Marcy l’Étoile, France). Strain Marseille-P3296 identification by matrix-assisted desorption ionization–time of flight mass spectrometry (MALDI-TOF MS) using a Microflex spectrometer (Bruker Daltonics, Leipzig, Germany) failed [2]. Then, 16S rRNA gene sequencing was performed for further analysis using a 3130-XL sequencer (Applied Biosciences, Saint Aubin, France) with fDI-rP2 primers (Eurogentec, Seraing, Belgium) as previously described [3]. Strain Marseille-P3296 exhibited a 96.19% sequence identity with *Collinsella aerofaciens* type strain JCM

10188 (AB 011816), the phylogenetically closest species with standing in nomenclature (Fig. 1). Thus, strain Marseille-P3296 can be classified as a new species within *Collinsella* genus [4].

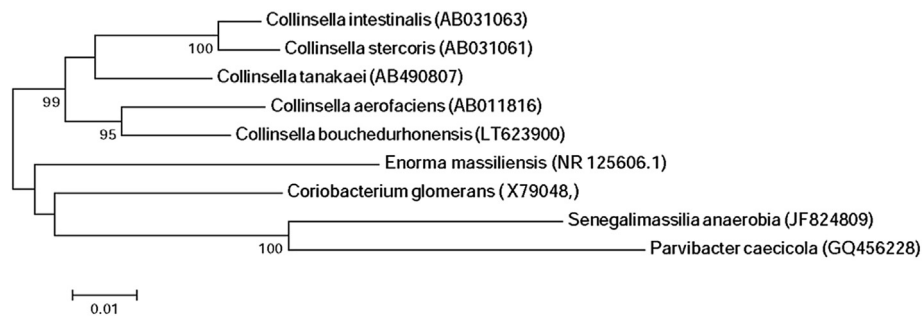
Because the 16S rRNA gene sequence of strain Marseille-P3296 diverges by more than 1.3% from the 16S rRNA gene sequence of its phylogenetically closest species with standing in nomenclature [5], we suggest the discovery of the new species “*Collinsella bouchedurhonensis*” (*bou.ches.du.rho.nen’sis*, N.L. fem. adj. *bouchedurhonensis*, pertaining to Bouches du Rhône, the name of the French territory where strain Marseille-P3296 was isolated). Strain Marseille-P3296 is the type strain of the new species “*Collinsella bouchedurhonensis*.” Colonies were smooth with a mean diameter of 0.1 to 0.5 mm. Bacterial cells were Gram-positive bacilli, were catalase and oxidase negative and had a mean diameter of 2.6 µm.

### MALDI-TOF MS spectrum accession number

The MALDI-TOF MS spectrum of “*C. bouchedurhonensis*” is available online (<http://www.mediterranee-infection.com/article.php?laref=256&titre=urms-database>).

### Nucleotide sequence accession number

The 16S rRNA gene sequence was deposited in GenBank under accession number LT623900.



**FIG. 1.** Phylogenetic tree showing branching of “*Collinsella bouchesdurhonensis*” strain Marseille-P3296 among other phylogenetically close neighbours. Sequence alignment was performed by CLUSTALW tool; then MEGA software was used for phylogenetic tree generation by neighbour-joining method. Five hundred bootstraps were performed; scores of at least 90% are shown on nodes. Scale bar indicates 2% nucleotide sequence divergence.

### Deposit in a culture collection

Strain Marseille-P3296 was deposited in the Collection de Souches de l'Unité des Rickettsies (CSUR, WDCM 875) under number P3296.

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### Conflict of Interest

None declared.

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