



“Collinsella vaginalis” sp. nov., a new bacterial species cultivated from human female genital tract



Khoudia Diop ^a, Florence Bretelle ^{a,b}, Pierre-Edouard Fournier ^a, Didier Raoult ^{a,c}, Florence Fenollar ^{a,*}

^a Aix-Marseille Université, Institut hospitalo-universitaire Méditerranée-infection, URMITE, UM63, CNRS 7278, IRD 198, Inserm U1095, Faculté de médecine, 27 Boulevard Jean Moulin, 13385 Marseille cedex 05, France

^b Department of Gynecology and Obstetrics, Gynépole, Hôpital Nord, Assistance Publique-Hôpitaux de Marseille, AMU, Aix-Marseille Université, France

^c Special Infectious Agents Unit, King Fahd Medical Research Center, King Abdulaziz University, Jeddah, Saudi Arabia

ARTICLE INFO

Article history:

Received 14 September 2016

Revised 19 October 2016

Accepted 8 November 2016

Available online 15 November 2016

ABSTRACT

We present a brief description of “*Collinsella vaginalis*” strain P2666 (=CSUR P2666), a new bacterium that was cultivated from the vaginal sample of a 26-year-old woman affected from bacterial vaginosis.

© 2016 Elsevier Ltd. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Keywords:

“*Collinsella vaginalis*”

Culturomics

Vaginal flora

Bacterial vaginosis

Human microbiota

As part of the study of the human microbiota thanks to the approach of microbial culturomics [1], we analyzed the vaginal flora of women with bacterial vaginosis [2]. In May 2015, we cultivated from the vaginal swab of a 26 year-old French patient a bacterial strain that could not be identified using matrix-assisted laser desorption-ionization time-of-flight mass spectrometry (Microflex spectrometer, Bruker Daltonics, Leipzig, Germany) [3]. The study was authorized by the local ethics committee of the IFR48 (Marseille, France; agreement 09-022). The patient gave also her written consent.

The initial growth of strain Marseille P2666 was obtained at 37 °C under anaerobic conditions after 7 days of culture on CNA (Colistin and Naladixic Acid) agar (BD Diagnostics, Le Pont-de-Claix, France) after 15 days of pre-incubation in a blood culture bottle (BD Diagnostics) enriched with 4 ml of rumen that was filter-sterilized through at 0.2 µm pore filter (Thermo Fisher Scientific, Villebon-sur-Yvette, France) and 3 ml of sheep blood (bioMérieux, Marcy l'Etoile, France). Bacterial cells are rod-shaped Gram-positive, strictly anaerobic, non-motile, and non-spore-forming with a mean diameter of 0.4 µm and a mean length of 1.8 µm. After 2 days of incubation at 37 °C under anaerobic conditions on blood agar (bioMérieux), colonies are grey, circular, and

opaque with a diameter of 0.5–1.2 mm. Strain Marseille P2666 exhibited neither oxidase nor catalase activity.

The 16S rDNA sequence was obtained after amplification using the universal primer pair (fD1 and rP2) and a 3130-XL sequencer (Applied Biosciences, Saint-Aubin, France), as previously reported [4]. 16S rRNA gene sequence-based identification of strain Marseille P2666 exhibited 96.08% of identity with *Collinsella intestinalis* strain JCM 10643 (GenBank accession number NR_113165), the phylogenetically closest bacterium with a validly published name (Fig. 1). As this sequence was below the 98.7% threshold to define a new species [5], strain Marseille P2666 was considered as a new species within the *Collinsella* genus in the *Coriobacteriaceae* family. Created in 1999, the genus *Collinsella* contains currently 4 species [6]; all were isolated from human faeces.

Strain Marseille P2666 presents a 16S rRNA divergence around 3.8% with its phylogenetically closest species [7], we propose that strain Marseille P2666 may be the representative of a novel species named “*Collinsella vaginalis*” sp. nov. (va.gi.na'lis. L. n. vagina sheath, vagina; L. masc. suff. -alis suffix denoting pertaining to; N.L. masc. adj. vaginalis pertaining to vagina, of the vagina). Strain Marseille P2666^T is the type strain of the new species “*Collinsella vaginalis*” sp. nov.

MALDI-TOF MS spectrum accession number. The MALDI-TOF MS spectrum of *Collinsella vaginalis* is available at <http://www.mediterranee-infection.com/article.php?laref=256&titre=urms-database>.

* Corresponding author.

E-mail address: florence.fenollar@univ-amu.fr (F. Fenollar).

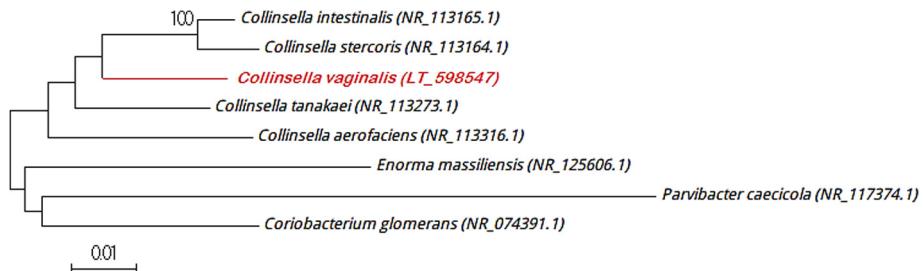


Fig. 1. Phylogenetic tree highlighting the position of “*Collinsella vaginalis*” strain Marseille P2666^T relative to other closest species. GenBank accession numbers are indicated after the name. Sequences were aligned using Muscle v3.8.31 with default parameters and phylogenetic inferences were obtained using neighbor-joining method with 500 bootstrap replicates, within MEGA6 software. Only bootstrap bigger than 95% are shown. The scale bar indicates a 1% nucleotide sequence divergence.

Nucleotide sequence accession number. The 16S rRNA gene sequence was deposited in EMBL-EBI under accession number LT598547.

Deposit in culture collection. Strain Marseille P2666 was deposited in the “Collection de Souches de l’Unité des Rickettsies” (CSUR, WDCM 875) under number P2666.

Acknowledgments

This research is funded by the Fondation Méditerranée Infection.

References

- [1] Lagier JC, Hugon P, Khelaifa S, Fournier PE, La Scola B, Raoult D. The rebirth of culture in microbiology through the example of culturomics to study human gut microbiota. Clin Microbiol Rev 2015;28:237–64.
- [2] Menard JP, Fenollar F, Henry M, Bretelle F, Raoult D. Molecular quantification of *Gardnerella vaginalis* and *Atopobium vaginae* loads to predict bacterial vaginosis. Clin Infect Dis 2008;20:33–43.
- [3] Seng P, Abat C, Rolain JM, Colson P, Lagier JC, Gouriet F, et al. Identification of rare pathogenic bacteria in a clinical microbiology laboratory: impact of matrix-assisted laser desorption ionization-time of flight mass spectrometry. J Clin Microbiol 2013;51:2182–94.
- [4] Drancourt M, Bollet C, Carlioz A, Martelin R, Gayral JP, Raoult D. 16S ribosomal DNA sequence analysis of a large collection of environmental and clinical unidentifiable bacterial isolates. J Clin Microbiol 2000;38:3623–30.
- [5] Stackebrandt E, Ebers J. Taxonomic parameters revisited: tarnished gold standards. Microbiol Today 2006;33:152–5.
- [6] Collinsella. <http://www.bacterio.net/collinsella.html>. [Accessed: 10-Sept-2016].
- [7] Yarza P, Richter M, Peplies J, Euzéby J, Amann R, Schleifer KH, et al. The All-Species Living Tree project: a 16S rRNA-based phylogenetic tree of all sequenced type strains. Syst Appl Microbiol 2008;31:241–50.