

# Low prevalence of Cutibacterium acnes in prostatic tissue biopsies in a French hospital

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## Anaerobe Version 2 2 Low prevalence of *Cutibacterium acnes* in prostatic tissue biopsies in a French hospital 4 A-L. Bidaud<sup>1</sup>, G. Karam<sup>2</sup>, C. Kandel-Aznar<sup>3</sup>, L. Ruffier d'Epenoux<sup>1,4</sup>, A. Guillouzouic<sup>1</sup>, 6 P. Bémer<sup>1</sup>, A-G. Lerov<sup>1</sup>, S. Corvec<sup>1,4</sup> 8 <sup>1</sup> Service de Bactériologie-Hygiène Hospitalière, CHU Nantes, Nantes Université, France <sup>2</sup> Service d'Urologie et transplantation, CHU Nantes, Nantes Université, France 10 <sup>3</sup> Service d'Anatomo-Pathologie, CHU Nantes, Nantes Université, France 12 <sup>4</sup> CRCINA, INSERM U1232, Nantes Université, Nantes, France 14 Running title: Cutibacterium acnes and prostatic tissue 16 Keywords: Cutibacterium acnes, prostate, phylotypes, inflammation, chronicity 18 20 22 24 26 **Correspondence:** Stéphane Corvec 28 Institut de Biologie des Hôpitaux de Nantes, Service de Bactériologie-Hygiène hospitalière CHU de Nantes 9 quai Moncousu, 44093 Nantes Cedex 01, France 30 Email: stephane.corvec@chu-nantes.fr Tel: +33.2.40.08.39.55 32 Fax: +33.2.40.08.39.28

34	Abstract: 53 words
	We evaluated the Cutibacterium acnes prevalence in prostatic biopsies and characterized the
36	strains at a molecular level. 18 out of 36 biopsies (50%) were sterile after seven days in
	culture. C. acnes was observed in only two biopsies. Its prevalence was low (5.6%). Finally,
38	the molecular characterization revealed diverse clusters including phylotypes IA1, IB and II.
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#### Note: 941 words

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Prostatic adenocarcinoma remains one of the most frequent tumors for male patients. The 62 persistence of microorganisms leading to a chronic inflammation of the prostate gland has been suspected to be responsible for the carcinoma process [1]. Viruses or bacteria such as 64 Cutibacterium acnes have been reported for their remanence in prostatic tissues, especially in Asia and Northen Europe [2,3]. C. acnes prevalence in prostatic samples has been reported 66 from 23% to 95%, depending on the method used [4]. However, the relationship between C. acnes detection and prostatic cancer or chronic prostatitis remains controversial. Currently, no 68 study has been conducted in France. The aim of this monocenter prospective study was to assess *C. acnes*' prevalence within prostatic biopsies. 70 During a four-month period at Nantes University hospital, 36 patients were included prospectively during their urology consultation. These patients were admitted for prostatic 72 biopsies as part of the adenocarcinoma diagnosis associated with prostatitis or not. The tissue sampling, using a transrectal approach with a 400 mg ofloxacine antibioprophylaxis, was 74 realized with a biopsy sterile gun. The first needle was meant for the bacteriology analysis while the 12 other cartography prostatic biopsies were used for histological analysis. Prostatic 76 biopsies were placed in sterile conditions, using physiological water. After grinding with glass beads for 15min, 50 µL of the sample were seeded on a Schaedler agar plate incubated 78 in anaerobic condition at 37°C for 7 days. A first look was performed at day 2 and another one at day 7 (D7). At each time, suspect colonies were identified using MALDI-TOF mass spectrometry as previously described [5]. For each C. acnes strains detected, a molecular 80 typing was performed [6].

Patients' median age was 69.5 and the median PSA level was 7.5 ng/mL (N <4,50 ng/mL).

Prostatic adenocarcinoma was diagnosed for 23 patients (64%) with a Gleason score ranging

from 6 to 10.

Regarding the microbiological analysis, 18 out of 36 biopsies (50%) were sterile at D7 in culture including 7/13 (54%) in the non-cancer group and 11/23 (48%) in the adenocarcinoma group (**Figure 1**). The 18 remaining samples were tested positive, mostly with a polymorphic flora, including *Bifidobacterium sp.* (n=4) or *Mobiluncus curtisii* (n=3).

A positive *C. acnes* culture (usually a few colonies) was observed in only two biopsies representing a prevalence of only 5.6%. One patient (non-cancer group) had two different types of *C. acnes* (hemolytic and non-hemolytic) associated with *Escherichia coli* and *Streptococcus salivarius* group. The other patient (adenocarcinoma group) had a *C. acnes* associated with *Collinsella aerofaciens* and *Finegoldia magna*, both present in the gastrointestinal tract. Regarding these three strains' molecular types, three different lineages were found. For the first patient, a phylotype IB, a SLST-type K16 belonging to clonal complex CC36 (hemolytic) and a phylotype II, SLST-type K16 belonging to CC53 (non-hemolytic), generally detected in prosthetic joint infections [7,8]. For the second patient, a phylotype IA<sub>1</sub>, SLST-type K1 belonging to CC18, usually found on the skin [9]. Interestingly, *Cutibacterium avidum* was also found in the three samples either alone (adenoma patient) and with *C. aerofaciens* and *Mobiluncus curtisii* (non-cancer patient). At last, no microorganism or mixed flora was found in patients presenting a prostatitis diagnosis, regardless of adenocarcinoma's presence.

The role of *C. acnes* in the inflammatory stimulation prostatitis and the carcinogenesis process remains under debate. Compared to the existing literature and despite crushing prostatic biopsies, the prevalence of *C. acnes* remains surprisingly low in this work series. In the review of Capoor et al, the detectable *C. acnes* prevalence varied within cancer tissue

samples significantly from 23 to 95% depending on the method used [4]. This is a key finding. Indeed, it seems really difficult to compare microbiological culture methods, universal 16S rRNA amplification, specific C. acnes-target PCR, Next Generation Sequencing approach, histologic observations, in situ fluorescence microscopy or antibody methods to detect viable or alive C. acnes. These methods allow to distinguish infection, colonization and skin contamination [4]. Moreover, 16S rRNA PCR is probably not the best molecular target to assess C. acnes presence [10,11]. Therefore, notwithstanding our expertise in orthopedic-device related infections and skin proinflammatory disease [8,9], as far as our ability to detect C. acnes strains in prostatic specimens is concerned, we question a possible difference regarding the ecology of the prostatic biopsies analyzed. Indeed, we detected for almost 50% of the samples, a polymorphic flora. This seemingly implies that we would have found a higher positivity rate with a higher microbial diversity using other methods, including molecular and sequencing ones as previously reported [12]. If certain C. acnes strains may likely play a role in either chronic inflammation of prostatic tissue or prostatic tumor [4], we did not find a specific tropism of a phylotype or a specific lineage as stressed by Mak et al. [13]. A different ecology hypothesis across ethnicities has been previously suggested concerning viruses detected in prostatic biopsies, [2] but the differences are still to be confirmed by a large scale, cross-center prospective study. A major limitation of this study is its small sample size at a single location. However, grinding should have helped detecting the intracellular C. acnes bacteria [14]. Another limit ties to the fact that only one biopsy per patient was analyzed. A cartography, using different prostatic biopsies, would have allowed to collect more accurate data. Conducting further studies in different European countries using the same methodology would also help. As a reminder, this methodology entails (i) analyzing the potential differences in C. acnes

frequency in prostatic tissues, (ii) characterizing at a molecular level the strains involved and

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132 determining a potential specific tropism and finally (iii) better understanding the correlation between *C. acnes* presence and either prostatic inflammation or cancer development.

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136 Conflict of interest: nothing to declare

#### **Ethical approval** 138

The human biological samples collected during the treatment of patients undergoing prostate biopsies were stored in the microbiology laboratory of the Nantes University Hospital. These

samples integrated into the collections of human biological samples attached to the

"Infectiology" research program declared on 05/09/2011 under the n° DC-2011-1399 and in 142 the following amending declarations (DC-2012-1555; DC-2013-1832; DC2014-2206 and DC-

144 2017-2987 currently pending) at the Ministry of Research and having obtained a favorable decision from the CPP Ouest IV on 07/04/2015.

Data collection followed the principles outlined in the Declaration of Helsinki; all patients 146 had signed an informed consent agreement to deliver their own anonymous information for

148 future studies.

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Fig1. Flow chart of the study, polymorphic flora including microorganims from the gut flora:

E. coli, E. faecium, Collinsella aerofaciens, Bifidobacterium spp., Mobiluncus curtisii and Finegoldia magna

