

Molecular Detection of *Ureaplasma urealyticum* from Prostate Tissues using PCR-RFLP, Tehran, Iran

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KEY WORDS

Ureaplasma urealyticum
Ureaplasma parvum
PCR-RFLP
Prostatitis
Prostate tissue

ABSTRACT

Background: In most cases, prostatitis can be caused by a bacterial agent such as *Ureaplasma urealyticum*. Considering to the cumbersome of the culture method for the detection of *Ureaplasma* species in clinical samples such as prostate; PCR method that is faster and more appropriate than the cultivation methods, can be utilized for the detection of *U. urealyticum* and *U. parvum*. PCR-RFLP method can differentiate both biovars and assist in studies of the clinical diagnosis, epidemiology and pathology of this species in human. The aim of this study was to molecular detection of *U. urealyticum* in prostate tissue samples based on PCR- RFLP.

Methods: Two hundred prostate tissue samples were collected from patient suffering from prostatitis. The PCR assay was used to amplify a 559 bp fragment of 16S-23SRNA interspace region of *Ureaplasma*. After sequencing, PCR products from positive samples were digested with *TaqI* restriction enzyme.

Results: Seven cases (3.5%) out of 200 prostate tissue samples were positive for *U. urealyticum*. Results of PCR products sequencing demonstrated that all isolates were *U. parvum* biovar. PCR-RFLP results shown that there was not any differentiation in pattern of enzymatic digestion, in addition, all isolates were *U. parvum*, serovar 3.

Conclusion: *U. urealyticum* can be one of the causing agents of prostatitis. Using PCR-RFLP with specific primer and restriction enzyme is a rapid and cost-effect method for detection and differentiation of *Ureaplasma* from clinical samples.

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Introduction

Prostatitis is an inflammation of the prostate gland, which can be varied from an acute to chronic medical condition (1). It is mostly caused by a bacterial infection (2). In recent years,

the role of genital *Mycoplasma* especially *U. urealyticum* has been evaluated (3).

The genus *Ureaplasma* belongs to Mycoplasmataceae family and mollicutes class. Like other members of this class, *Ureaplasma* is lacking a cell wall and the smallest free-living

organisms known. *U. urealyticum* is considered as agent of nongonococcal urethritis (NGU), acquired arthritis and prostatitis in men (4). It is subdivided to *U. parvum*, including serotypes 1, 3, 6 and 14 and is designated as biovar 1. The other biovar of *U. urealyticum* include the remaining 10 of established serotypes and designated as biovar 2 (5).

Less than 60% DNA homology are between biovars (6). Using culture for detection of *Ureaplasma* species in patient samples such as prostate tissues is laborious, taking several days and requiring a high degree of technical skill (7). PCR method is specific, sensitive and provides results in short time. Due to its equivalent and fast performance than to culture, it can be replaced to conventional culture for detection of *U. urealyticum* and *U. parvum* (8, 9). In addition, culture alone does not differentiate *U. urealyticum* from *U. parvum*. In several molecular typing methods introduced for *U. urealyticum*, PCR-RFLP has some advantages such as easy interpretation, implementation and using for large quantities of samples (10, 11).

Both biovars can be differentiated by this method, which assists in studies of the clinical significance, epidemiology and pathology of this species in human. Ryo et al. examined the 16sRNA-23sRNA intergenic spacer regions of *U. urealyticum* and showed that this region also can be used for biovar identification (12). For the diagnosis of bacterial prostatitis, samples such as voided urine, urethral swabs, and expressed prostatic secretions can be used. The normal bacterial flora in the area of genitourinary causes contamination of samples and makes it difficult to interpret the results. To avoid contamination with flora, the appropriate sample is direct using of prostate tissue (1). Recently, a study was done especially on prostate samples for detection of *Ureaplasma* in cancer cases but there is not any report of the role of *U. urealyticum* in prostatitis in Iran (13). The aim of this study was to molecular detection of *U. urealyticum* in human

prostate tissue samples in prostatitis cases based on PCR-RFLP.

Materials and Methods

In this project, a cross sectional study was conducted and 200 prostate tissues were collected from prostatitis patients referred to Tehran hospitals during 2008 to 2010 for the preparation and examination. Only men with prostatitis were enrolled in this study and all of them subjected to prostate biopsy by a physician.

The prostatitis tissues were embedded with paraffin by pathologist. In the laboratory, using a microtome with disposable blades, 5-10 μ sections were prepared from blocks and transferred into sterile micro-tubes. To prevention cross contamination between the samples, the microtome blade and gloves were changed and the microtome were washed with 70% ethanol between samples. DNA was extracted from paraffin-embedded tissue samples using a special kit (manufactured by QiaGen Co). Briefly, 25 mg of cutting prostate tissue mixed with 20- μ g proteinase K and incubated at 56 °C until the tissue was completely lysed. Two hundred μ l of first buffer was added to the sample and incubated for 10 min at 70 °C. Then 200 μ l of ethanol were mixed with sample. Mixture was applied to the column and centrifuged at 600-x g for 1 min. Another Buffer was added in a sequence as protocol. At the end of process, DNA was collected in a clean micro-tube by Elution Buffer and stored at 4 °C for PCR or at -20 °C for long-term storage.

A PCR assay with primers UuF5'-(TGGAGTTAAGTCGTAACAAG)-3' and UuR5'-(CTGAGATGTTTCACTTCACC)-3' was used and amplified a 559-bp sequence. Primers for amplification of 16S-23SRNA inter-space region of *Ureaplasma* were synthesized by CinnaGen Company. The PCR reaction was

performed in a total volume of 30 µl. Each reaction contained 15 µl Master Mix (2X), 1.5 mM MgCl₂, 1 µg DNA template, 20 pmol of each reverse and forward primer, and sterile distilled water. PCR reaction mixture with *Ureaplasma* DNA and PCR reaction mixture without DNA were used as positive and negative control, respectively. The initial denaturation performed for 5 min at 95 °C. Total of 35 cycles was carried out as follows: denaturation for 30 sec at 94 °C, annealing for 60 sec at 56 °C and extension for 60 sec at 72 °C. The final extension was completed for 5 min at 72 °C. PCR product electrophoresed on 1.8% gel agarose- TBE buffer (Boric acid 27.5 gr, Tris base 54 gr, EDTA 20 ml (pH=8). For confirming of *U. urealyticum*, the PCR product was send for sequencing.

After sequencing, the BLAST program was used for alignment the sequence with other sequences available in database. The sequences were studied with Web Cutter software. Therefore, a list of restriction enzyme cutting sites was prepared. To identify polymorphism, the PCR product obtained from *Ureaplasma* DNA was digested with *EcoRI*, *AluI*, *TaqI*, *CacI*8 and *BbsI*. For enzymatic digestion, 15 µl reactions was prepared in the following ingredient; 6.3 µl sterile distilled water, 1.5 µl enzyme buffer, 0.2 µl restriction enzyme and 7 µl PCR product. After providing the reactions, micro-tubes containing all enzyme except *TaqI* were put in the water bath at 37 °C for 2 h and micro-tubes containing the enzyme *TaqI* were put at 65 °C for 2 h. After 2 h, products of RFLP were electrophoresed on 3% agarose gel. Finally, the size of the fragment resulted from enzymatic digestion of all positive PCR samples, was compared with each other and the biovars were identified.

Statistical analysis was conducted to determine how many samples were positive for *U. urealyticum*. Perspective analyses were performed and data with rounded-up numerical values (percentage) were documented.

Results

In 200 prostate tissue samples that their DNA was extracted and examined by PCR method, seven (3.5%) cases were positive for *U. urealyticum* and a fragment with 559 bp size was obtained (Fig. 1). Patients ranged in terms of age from 41 to 90 yr. Two patients were diagnosed with cancer which their samples contained *U. urealyticum*.

Obtaining results of PCR product sequencing and using of BLAST software shown that all of positive prostate samples contained *Parvum* biovar. *Urealyticum* biovar was not in samples.

The PCR products were confirmed by RFLP. The restriction enzyme selection was based on PCR product sequencing results. The PCR products of *U. urealyticum* were digested by *TaqI*, which yielded 227 and 332 bp fragments (Fig. 2).

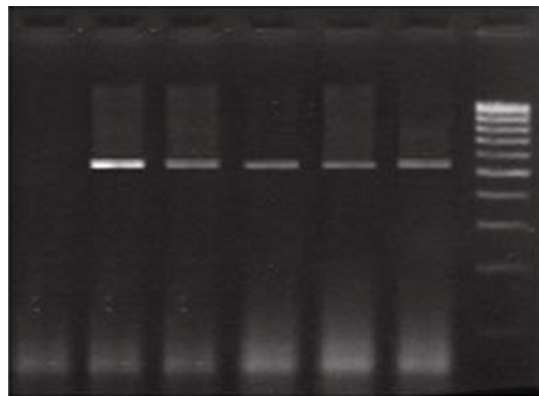
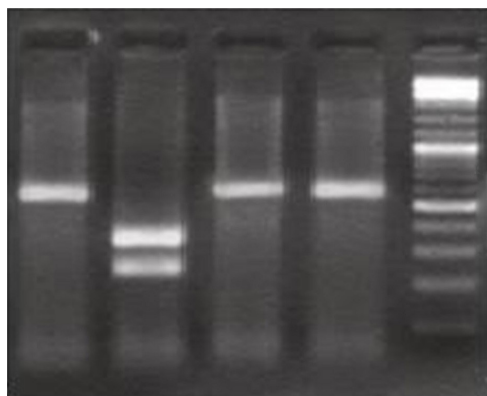


Fig. 1
Agarose gel electrophoresis of PCR amplified products. Lane 1. Negative control. Lane 2. Positive control. Lane 3,4,5,6. Positive samples containing of *Ureaplasma urealyticum* (559 bp band). Lane 7. Size marker (100bp DNA ladder)

Results of PCR-RFLP shown that there was not any differentiation in pattern of enzymatic digestion on PCR product and all isolates were from one type of *U. parvum* biovar belonging to serovar 3.

**Fig. 2**

Enzymatic digestion of PCR product obtained from positive samples. Lane 1. 559bp-long PCR product *Ureaplasma urealyticum* as positive control, Lane 2. PCR-RFLP pattern of *Ureaplasma urealyticum* after digestion with TaqI, specific enzyme that made two pieces of 227 bp and 332 bp; Lane 3, 4. PCR-RFLP pattern of *U. urealyticum* after digestion with Cac8I and BpsI (it is not digested); Lane 5. Size marker (100bp DNA ladder).

Discussion

Prostatitis is a common disorder of men and it is estimated that half of men suffer from symptoms at same stage of their live (14, 15). In many cases, urologists often find themselves, incapable to diagnosis and treat. Different bacteria causing the infection and the role of *Ureaplasma* as an etiologic agent of prostatitis are controversial (16). *Ureaplasma* one of the most common pathogens causing sexually transmitted disease that is important to determine its prevalence in men with prostatitis. Although *Ureaplasma* species have ability to visibly grow on the media but using molecular methods for identification of species and serovar for interpretation of pathogenicity is important (17, 18).

In this study, *U. urealyticum* was detected in 3.5% of prostate tissue samples using conventional PCR. In a study, multiplex PCR was used for detection of *U. urealyticum* in 92 specimens including prostate secretion and urine from patients. Amplifying a fragment of 16S rRNA gene by multiplex PCR method, detected *U. urealyticum*

in 4 (4.3%) samples (19). Lee et al. used multiplex PCR for detection of *U. urealyticum* and other bacteria in one-step in 96 urine samples. *U. urealyticum* was not detected in urine samples (20). However, present study differs from most of studies in type and size of sample and using of method. In these previous studies, urine samples and prostate secretions were used for detection of bacteria while we used prostate tissue for direct detection of *Ureaplasma* in prostate gland tissue.

PCR in the detection of pathogens in bacterial prostatitis is very effective. Because of many patients taking several antibiotics before an accurate diagnosis of bacterial, this will interfere with the growth of bacteria in vitro. Therefore, using PCR we can identify the causative agent directly and rapidly. Because the gene sequence obtained from PCR method has a specific restriction site for *Ureaplasma* species, using this method is possible to distinguish the different species.

In this study, RFLP method was used for typing to better understand the pathology, epidemiology and clinical significant of *Ureaplasma* species in human prostatitis. In the first step, result of PCR product sequencing showed that *U. parvum* (biovar 1) was predominant isolate and *U. urealyticum* (biovar 2) was not detected. In other hand, there was no difference between obtained patterns of TaqI enzyme digestion on PCR product from positive samples. These results state that all of *U. parvum* biovar belong to serovar 3. The findings are consistent with previous studies (21, 22). In a study, *U. parvum* was the most frequently occurring species (23). Eventually, *U. parvum* serovar 3 has a possible pathogenic role.

Conclusion

U. urealyticum could be one of the causing agents of prostatitis. Since all of biovars belong to *U. parvum*, therefore it can be said that the

causative agents of prostatitis in men are due to play a single biovar that makes it easier to follow up and treatment of infection origin. Using of PCR-RFLP with specific primer and restriction enzyme is a rapid and cost-effect method for detection and differentiation of *Ureaplasma* from clinical samples.

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Conflict of interest

The authors declare that there is no conflict of interests.

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