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Original Article

Study on ITS1 Gene of Iranian *Trichomonas vaginalis* by Molecular Methods

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Abstract

Background: Trichomoniasis is a worldwide protozoan parasitic disease and metronidazole is a choice drug for its treatment. Because of disease importance in public health and its controversial ideas about the prevalence of drug resistance, this study was carried out.

Methods: Fifty-two suspected vaginal samples were collected from 2006 to 2007 in Gynecology Maryam Hospital, Tehran, Iran. All isolates were examined by microscopic, culture and PCR techniques. The PCR products were analyzed by RFLP and CSGE methods and two suspected samples were sequenced.

Results: *Trichomonas vaginalis* was identified from all 52 samples. Of 52 isolates, 45 samples were successfully cultured and amplified by PCR except one. Seven were positive only by PCR. Finally, ITS1 fragment was successfully amplified in 51 of 52. CSGE analysis and PCR products digestion by MspI followed by sequencing showed nucleotide mutation at position 209 (C209T) of the ITS1 fragment in two (3.9%) of them.

Conclusion: The results showed mutation in ITS1 fragment of *T. vaginalis* in two (3.9%) of Iranian isolates which may be related to metronidazole resistance.

Keywords: *Trichomonas vaginalis*, Mutation, ITS1 fragment

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Introduction

Trichomoniasis is the most widespread non-viral sexually transmitted disease caused by *T. vaginalis* (1).

The clinical disease in women ranges from asymptomatic to symptomatic such as severe urethritis, vulvo-vaginitis, cervicitis (2), birth of preterm or low-birth-weight infant and possibility of development of cervical cancer are associated complication in affected women (3,4). The infected women are more susceptible to other transmitted diseases including HIV (5). One of the most significant clinical features of *T. vaginalis* is resistance to choice drug, metronidazole (6-8) which was recognized as an effective treatment with cure rate of approximately 95% (9).

Metronidazole resistance has been reported since 1962 (10) and some studies have been reported from 2.5% - 5% (11, 12). More than 10% of infected patients in the United States does not respond to this drug (13), and some clinical isolates have showed decreasing susceptibility to metronidazole (14). Rasoloson et al. showed mechanism of in vitro development of resistance to metronidazole in *T. vaginalis* (15). Meri et al. reported the first three cases of metronidazole resistant *T. vaginalis* from Finland (16). Resistance testing was performed for 78 of 467 isolates of *T. vaginalis*, 4 ones were resistance to difference concentrations of metronidazole (17). Clinical metronidazole resistance is defined by failure infection clearance by standard treatment (8). Internal transcribed spacers (ITS1 and 2) are nonfunctional fragments in ribosomal RNA between 18s rRNA and 28s rRNA and found in all eukaryotic organisms which are transcribed, but not translated. Some investigators proposed that it has an important role in using metronidazole as choice treatment (8). Identification of the molecular methods to detect drug resistance

may lead to faster results for clinical treatments (18).

This study was designed for the first time in Iran for mutation detection in ITS1 gene of *T. vaginalis* by molecular methods.

Materials and Methods

Sampling

As a descriptive study on fifty two vaginal samples were collected from both treated and untreated women suspected to trichomoniasis that referred to Gynecology Department of Maryam Hospital in Tehran from 2006 to 2007. Wet mounts examinations were done for all samples and all were cultured in 50 ml of TYM medium (19).

DNA extraction

Cultures were centrifuged at 430 ×g for 20 min at 4°C and pellets were washed once with PBS (pH 7.4). Total nucleic acid was extracted with the Prim Prep Genomic DNA Isolation Kit (GeNet Bio company, Chungcheongnam-Do, Korea), according to the manufacturer's manual.

PCR amplification

The ITS1 fragment was amplified by designed primers based on *T. vaginalis* ribosomal DNA (rDNA) gene (ACCESSION AF466750). TVITS F: 5'- ACA CCG CCC GTC GCT CCT AC -3' and TVITSR: 5' AAT TTG CAT TCA AAG ATT AAC- 3'. These primers were amplified 313 bp of *T. vaginalis* ITS1. The PCR reaction contained 500 ng DNA, 12.5 µl master mixes (Ampliqon, California, US) and 20 pmol of each of the forward and reverse primers in 25 µl of final volume. PCR reactions were carried out in 30 cycles, denaturation at 94°C for 30 sec, annealing at 52°C for 30 sec and extension at 72°C for 30 sec. The reaction was incubated for 5 min. at 94°C and 72°C as pre

denaturation and post extension steps, respectively. The PCR products were subjected to electrophoresis using 2% agarose gel, stained by ethidium bromide and visualized under a UV transilluminator.

PCR product analysis

All PCR products were analyzed by CSGE method (20) and RFLP using digestion by MspI restriction enzyme. The CSGE is a universal mutation detection system was developed for study into rapid and non-radioactive heteroduplex based detection for mutation screening. PCR products were mixed by control PCR product, subjected to 5 min at 98 °C and 30 min at 68 °C, mixed by five µl of loading buffer (30% glycerol, 0.25% bromophenol blue and 0.25% xylen cyanol FF), and subjected to electrophoresis on 10% CSGE gel. The PCR products were digested with MspI restriction enzyme for 3h at 37°C in the appropriate buffer and restriction patterns were seen on 3% agarose gel.

Results

After investigating *T. vaginalis* isolates by three methods including direct smear, culture in TYI and PCR, 44 isolates were positive in both culture and PCR but seven were

positive only by PCR. The ITS1 fragment was amplified by PCR in 51 of 52 DNA extractions. Figure 2 showed 2% agarose gel electrophoresis of amplified 313 bp as PCR product.

All the 51 PCR products of *T. vaginalis* isolates were electrophoresed on CSGE gel and two PCR products were suspected. The suspected samples were purified and sequenced by dideoxy chain termination method. Our results demonstrated mutation in ITS1 fragment in two cases (3.9%) that were symptomatic and none of them used metronidazole before examination. Both sequences were aligned in GenBank by Blast software and demonstrated mutation related to aligned genes in GenBank.

The all-51 amplified PCR products were digested by MspI restriction enzyme and electrophoresed on 3% agarose gel. Figure 3 showed RFLP pattern of mutant wiled samples on agarose gel. Sequence of mutated samples submitted to GenBank at accession numbers EU308571 and EU308572. The results showed mutation at position 209 (C209T) of the ITS1 region in which the thymidine was replaced by cytosine in accession number EU308572.

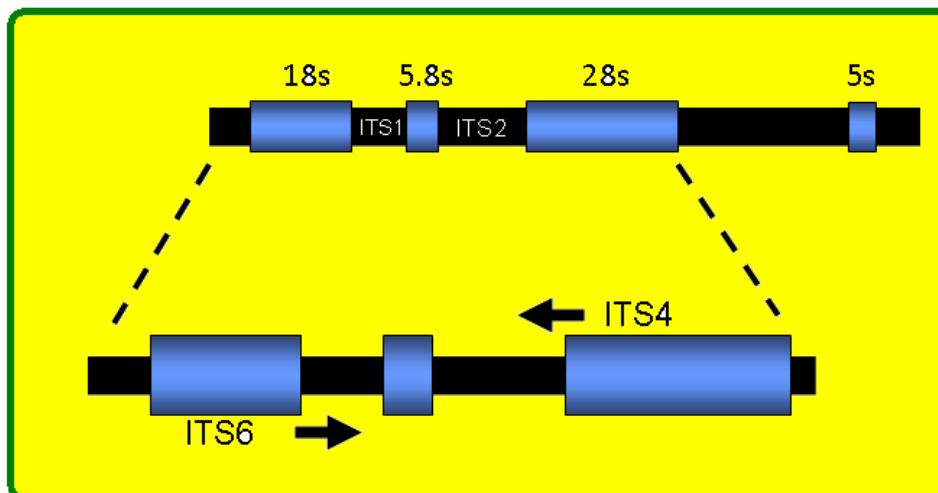


Fig. 1: The map position of ITS1 and ITS2 in eukaryotic genome

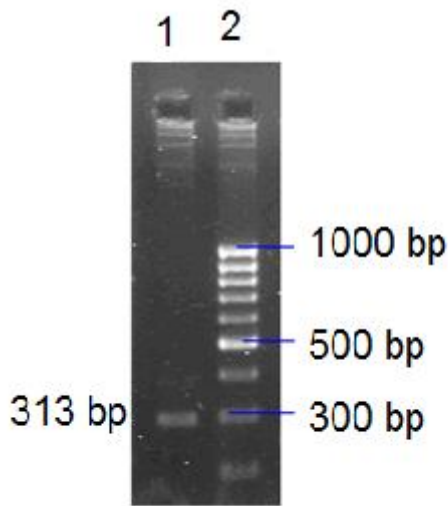


Fig. 2: Electrophoresis of PCR product on agarose

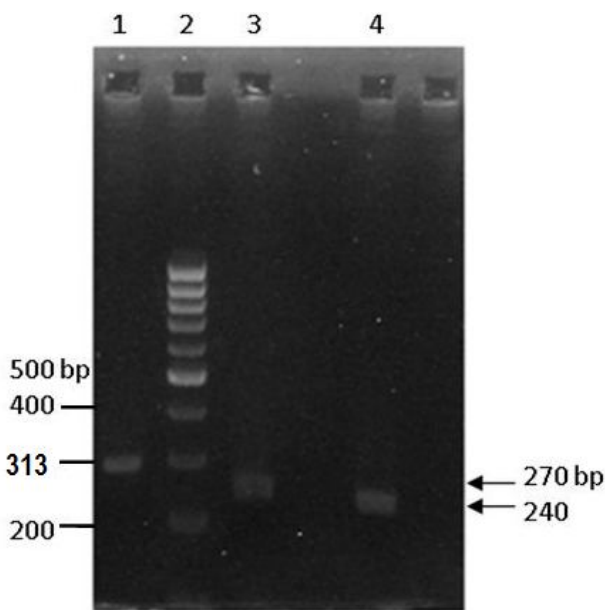


Fig. 3: Agarose gel electrophoresis of RFLP pattern. Lane 1. PCR Product (313bp). Lane 2. 100 bp DNA Ladder marker. Lane 3. MspI digested wild type PCR Product (270 bp). Lane 4. MspI digested mutant type PCR Product (240 bp)

Discussion

The purpose of this study was detection mutations in ITS1 of *T. vaginalis* isolated from Iranian patients. Although correlation between metronidazole resistance and ITS1 mutations have been reported among clinical isolates of *T. vaginalis* (8), but others reported that correlation between the ITS1 sequence variations and drug resistance was not observed and may be related to *Mycoplasma hominis* (21). Previous studies reported relationship between ITS1 mutations and drug resistance, but they showed different results. The mechanism of metronidazole resistance in *T. vaginalis* from treatment failure is not clear (22). Additional studies are needed to adapt the differences between these studies. Internal transcribe spacer (ITS) has been considered as an appropriate candidate gene for determination of metronidazole susceptibility on *T. vaginalis*, the sequence of this gene have conserved among related species (21).

Despite its high prevalence, the genetic variability and drug resistance features of this parasite are poorly understood. The mutations in genomic level (like ferredoxin mutation) are the important cause of drug resistance phenomenon (23). Butler and Xioa in two separate studies showed 51% and 14% resistance to metronidazole *in vitro* respectively (18, 21). In other study Upcroft et al., using PCR, showed 17.4% resistance to metronidazole (24). Due to different methods, our results were different from mentioned studies.

This suggests that single mutation on ITS1 gene may be a marker for identify resistance that could lead to control and treatment strategies to avoid from prevent of their distribution. Clearly, further studies with larger samples from different endemic regions are needed to explain this contradiction.

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