Detection of *Toxoplasma gondii* DNA in Sheep and Goat Milk in Northwest of Iran by PCR-RFLP

Mousa Tavassoli 1,*, Bijan Esmaeilnejad 1, Farnaz Malekifard 1, Ali Soleimanzadeh 2, Mahdi Dilmaghani 3

1Department of Pathobiology, Faculty of Veterinary Medicine, Urmia University, Urmia, IR Iran  
2Department of Clinical Sciences, Faculty of Veterinary Medicine, Urmia University, Urmia, IR Iran  
3Department of Microbiology, Faculty of Veterinary Medicine, Urmia University, Urmia, IR Iran

*Corresponding author: Mousa Tavassoli, Department of Pathobiology, Faculty of Veterinary Medicine, Urmia University, P.O. Box 1177, Urmia, IR Iran. Tel: +98-4412972654, Fax: +98-4412701508, E-mail: mtavassoli2000@yahoo.com.

Received: September 17, 2012; Revised: January 19, 2013; Accepted: January 27, 2013

**Background:** Toxoplasmosis is a widespread disease in humans and many other species of warm-blooded animals. Among livestock animals, sheep and goat are more widely infected by *Toxoplasma gondii*. This parasite is a major cause of abortion, with significant economic losses for sheep and goat breeders.

**Objectives:** The polymerase chain reaction (PCR) method was employed to detect of the *T. gondii* DNA in the milk of sheep and goats based on its B1 gene.

**Materials and methods:** A total of 625 milk samples were collected from 345 sheep and 280 goats from randomly selected flocks of Northwest of Iran.

**Results:** Of 625 examined milk samples, 19 animals (3.04%) yielded a specific *T. gondii* B1 fragment (529 bp), of which *T. gondii* was detected in 16 (4.63%) sheep milk samples and 3 (1.07%) goat milk samples. Restriction fragment length polymorphism (RFLP) analysis of the PCR products of *T. gondii* with AluI restriction enzyme produced only one distinct pattern among all positive samples, which indicates that one RFLP profile of *T. gondii* exists in the study area.

**Conclusions:** Presence of *T. gondii* DNA in the milk of sheep and goats raises the possibility that this parasite is transmitted through consumption of raw milk. Since sheep and goats are important milk sources in Iran, there is a high risk of contamination through milk from these hosts due to their susceptibility to infection. Further studies are required on milk producing animals to implement effective control strategies against toxoplasmosis.

**Keywords:** *Toxoplasma gondii*; Milk; Sheep; Goat; Iran

**Implication for health policy/practice/research/medical education:** Presence of *Toxoplasma gondii* DNA in the milk of sheep and goats raises the possibility that this parasite can be transmitted through consumption of sheep and goats raw milk.
eral domestic animals (11-13). Thus, due to the presence of tachyzoites in the milk, consumption of unpasteurized goat, sheep or cow milk and its products is a high-risk action (11). Occurrence of clinical toxoplasmosis in human has been attributed to consumption of unpasteurized goat milk (5, 14-16). Recently, due to the lack of studies involving the role of sheep milk as a potential source of toxoplasmosis transmission and based on the phylogenetic affinity between sheep and goats, several studies were carried out on ovine milk to determine the role of ovine milk as a potential source of toxoplasmosis transmission (17, 18).

2. Objectives

Cattle, sheep and goat are the most important meat and milk producing animals in Iran (19) and *T. gondii* is one of the most prevalent protozoan parasites in this country (20). Due to the economic importance of the disease as well as the paucity of published reports about the *T. gondii* presence in the milk of sheep and goats and also a health concern due to neonatal complications in Iran, it is necessary to investigate the prevalence of *T. gondii* infection in the milk of producing animals especially sheep and goats which supply the main sources of milk for local consumption in North West of Iran.

The aim of the present study was to detect of *T. gondii* DNA in the milk of infected sheep and goats. Moreover, PCR-restriction fragment length polymorphism (RFLP) was used for identification of the B1 gene polymorphism among *T. gondii* parasites obtained from sheep and goats milk samples of North West of Iran.

3. Materials and Methods

3.1. Sampling

625 small ruminants (including: 345 sheep and 280 goats) were randomly selected from various regions of North-West of Iran, from March to September 2011. Milk samples were taken manually by milking the teats which had previously been disinfected with iodine alcohol and use of gloves during manipulation. The samples were kept under refrigeration and transferred in sterile microtubes to the laboratory of Parasitology, Faculty of Veterinary Medicine, Urmia University, for molecular analysis.

3.2. PCR Amplification

Total DNA was extracted from each sheep and goat milk sample using a Genomic DNA purification kit (Fermentas, Germany), adjusted to a total volume of 200 µL in TE buffer and stored at -20°C until the usage time. Amplification of the B1 gene of *T. gondii* was performed by sensitive and previously reported species-specific primers used to amplify a fragment of 529 bp (21). The sequences of primers were as follows:

TOX4 (CGCTGCAGGAGGACAGCAAAGTGT)

TOX5 (CGCTGCAGACAGTGACATGGATT)

The PCR reaction was performed in a 50 µL total reaction volume containing 5 µL of 10X PCR buffer, 2 mM MgCl₂, 250 µM of each of the four deoxynucleotide triphosphates, 1.25 U Taq DNA polymerase (Fermentas, Germany), 50 pmol of each primer and 5 µL of the extracted DNA. The positive control for *T. gondii* was kindly provided by Razi Vaccine and Serum Research Institute (Tehran branch, Iran). Sterile water was served as negative control.

Cycling condition was 94°C for 7 minutes, followed by 33 cycles of 1 minute at 94°C, 1 minute at 55°C and 1 minute at 72°C with a final step at 72°C for 10 minutes. Five microliters of each DNA sample were used as the template. PCR products were analyzed by 2% agarose gel electrophoresis followed by ethidium bromide staining and photography.

3.3. RFLP of PCR Products

The amplified products were digested with AluI restriction enzyme (Fermentas, Germany) as described by the manufacturer, and analyzed using 2% agarose gel. Each digestion reaction was set up in a 20 µL volume containing 2 µL of the 10 X reaction buffer, 10 µL of PCR products and 10 U of the restriction enzyme. The digestion mixture was incubated at 37°C for 2 hours.

4. Results

4.1. Identification of *T. gondii* in the Milk Samples

All obtained milk samples were examined by PCR. The results showed that 16 (4.63%) sheep and 3 (1.07%) goats milk samples were infected (Figure 1).

![PCR-Amplified Products Using *T. gondii* Specific Primers](image)

Lane 1, 50 bp Ladder (Fermentas, Germany); lane 2, positive control; lane 3, negative control, lanes 4 - 5, sheep milk Samples; lanes 6 - 8, goats milk Samples
4.2. RFLP of *T. gondii* B1 Gene Amplified Region Isolated From Animals

A single RFLP profile, expected three fragments (364 bp, 87 bp and 78 bp), was yielded from PCR positive milk samples using AluI restriction enzyme (Figure 2).

![Figure 2. RFLP Patterns of Amplified 529 bp PCR Products of *T. gondii* Generated Using AluI](image)

Lane 1, 50 bp ladder (Fermentas, Germany); lane 2, undigested PCR product, lanes 3 - 4, digested PCR products of sheep milk; lanes 5 - 6, digested PCR products of goats milk

5. Discussion

Sheep and goats are economically important in many countries, because of production of milk and meat (22). Consumption of unpasteurized goat milk is a source of infection for humans (5, 15, 16). Chiari and Neves had reported the excretion of tachyzoit in the milk of naturally infected goats (15). During an acute infection in goats, tachyzoit of *T. gondii* may be excreted in the milk and become a possible source of human infection (5, 15). In addition, physiologic decrease in the peripartum immunity may lead to reactivation of *T. gondii* cyst and excretion of parasite tachyzoites in the milk (18). Tachyzoites are generally not considered as important sources of oral transmission of *T. gondii* because they are sensitive to proteolytic enzymes of the milk and are thought to be immediately destroyed by the gastric juice (11, 13). Presumably, a part of tachyzoites is excreted in the milk which is not destroyed by the gastric juice due to its rapid passage through this digestive compartment (23).

Although pasteurization will kill *T. gondii* in milk, unpasteurized raw milk and goat cheeses made out of it can be among the sources of *T. gondii* infection (7). Hiramoto et al. has shown that consumption of unpasteurized milk or fresh cheese is the major source of contamination by this parasite in rural areas (24). Based on the similarity between sheep and goats, ovine milk can be a potential source of infection to humans (18). There are several reports on PCR method usage for detection of *T. gondii* DNA in the milk of sheep and goats. Camossi et al. identified *T. gondii* DNA in seven milk samples of 20 naturally infected sheep by PCR. They demonstrated the presence of *T. gondii* DNA in the milk (18). In another study by Fusco et al. the transmission potential of ovine milk and dairy products was presented. They detected the parasite DNA in 3.4% of total analyzed milk samples of sheep by PCR (17).

In this study, *T. gondii* DNA was found in 16 (4.63%) and 3 (1.07%) milk samples from 345 and 280 naturally infected sheep and goats, respectively. Lower infection rates in goats compared to those in sheep may be attributed to the differences in susceptibility to *T. gondii* and the feeding habits of the animals (25). The presence of *T. gondii* DNA, detected by molecular analysis, raises the possibility that this parasite is transmitted through consumption of raw milk and its non-pasteurized derivatives which are highly-concerned result for the public health (18).

RFLP is a technique in which organism may be differentiated by analysis of the pattern derived from the cleavage of its DNA. Diversity of the generated patterns can be used to differentiate species and even strains from each other. PCR-RFLP has been used to identify the genotypes of *T. gondii* isolates (26). There are three major genotypes (type I, type II, and type III) of *T. gondii*. These genotypes differ in their pathogenicity and prevalence. Type III is more common in animals than in human toxoplasmosis (27). Zia-Ali et al. found four isolates of *T. gondii* from adult sheep in Iran, two of which were Type II and two were Type III. Type II was the predominant lineage of the strains isolated from sheep (20).

To date, No Type I isolate of *T. gondii* has been found in sheep (4). In a study on *T. gondii* infection in domestic animals of Urmia by PCR - RFLP, Tavassoli et al. showed that sheep and horse were infected with the same strain of *T. gondii* (28). As PCR-RFLP genotyping is simple and cost-effective compared to DNA sequencing, it is considered as a rapid and practical tool for identification and estimation of genetic diversity in *T. gondii* (29). In this study, the PCR-RFLP results indicated that a strain of *T. gondii* exists in the study area, and there is a need for more investigations regarding the distribution of different strains in other parts of the country.

In conclusion, our findings demonstrated the presence of *T. gondii* DNA in the milk of sheep and goats. Presence of *T. gondii* DNA raises the possibility that this parasite is transmitted through consumption of raw sheep and goats’ milk. Since sheep and goats are the most important milk sources in Iran, there may be a high risk of contamination through milk from these hosts due to their...
susceptibility to the infection. Therefore, further investigations are necessary to collect more information on the prevalence of T. gondii DNA in milk-producing animals to apply effective control strategies against toxoplasmosis.

Acknowledgements
Authors would like to thank Mr. A. Kazemnia for his technical assistance. We would like to sincerely thank the members of the Faculty of Veterinary Medicine and Urmia University Research Council for the approval and support of this research.

Authors’ Contribution
Study concept and design: Tavassoli, Esmaeilnejad, Malekifard. Analysis and interpretation of data: Tavassoli, Soleimanzadeh, Dilmaghani. Drafting of the manuscript: Tavassoli, Malekifard. Analysis and interpretation of data: Tavassoli, Esmaeilnejad, Malekifard. Funding/Support of this research.

Urmia University Research Council.

References