

Molecular Detection of *Chlamydomphila Abortus* In Aborted Fetal Tissues by Using Polymerase Chain Reaction (PCR) In Tabriz, Northwest of Iran

Mahsa Alem^{1*}, Reza Asadpour², Raziallah Jafari Joozani², Katayoon Nofouzi¹

¹ Department of Pathobiology, School of Veterinary Medicine, Tabriz University, Tabriz, Iran

² Department of Clinical Science, School of Veterinary Medicine, Tabriz University, Tabriz, Iran

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Abstract

Enzootic abortion of ewes (EAE) induced by *Chlamydomphila abortus* (formerly *Chlamydia psittaci* serotype1) is a major cause of reproductive failures in most sheep producing countries. In this study, the abortion prevalence of *Chlamydomphila abortus* (*C. abortus*) in sheep abortion cases are evaluated by polymerase chain reaction (PCR) and objectives are considered Chlamydiosis incidence of abortion in sheep in the northwest region of Iran. For this purpose, the contents of fetal tissue from 50 aborted fetuses were homogenized and DNA extracted from them and then PCR is done using specific primers for *Chlamydomphila* (CHOMP191, CHOMP371). The study showed that about 26% of the total sample is contained *Chlamydomphila*. This study confirms that *C. abortus* as an important cause of ovine abortion in the northwest of Iran and showed the PCR in tissue pools of aborted fetuses is a useful method for rapid detection of *Chlamydomphila abortus* Ovis infections.

Keywords: *Chlamydomphila abortus*, Molecular detection, Polymerase Chain Reaction (PCR)

Introduction

Ovine enzootic abortion (OEA) induced by *Chlamydomphila abortus* (*C. abortus*) (formerly *Chlamydia psittaci* serotype 1) is a major cause of reproductive failures in most sheep producing countries (Aitken et al., 1993). In an epidemic of OEA, up to 30% of ewes abort in the last three weeks of gestation or give premature birth to weak or dead lambs. After the abortion, the ewes develop a protective immunity and, in endemically infected flocks, 5-10% of the ewes abort annually (Rodlakis et al., 1998; Aitken, 2000).

Despite the economically important losses due to late-term abortions and weak lambs, the prevalence of *C. abortus* is not well known in most Asian countries, including Iran. Diagnosis of OEA, by examination of fetal tissues following postmortem examination, can be achieved by a variety of methods.

All of these methods are based on direct antigen detection and include modified Ziehl-Neelsen (MZN) staining of placental smears and direct fluorescent-antibody staining of *Chlamydiae* in frozen cryostat sections of the placenta or fetal tissues. Several enzyme-linked immunosorbent assays (ELISAs) have been proposed for testing for

C. abortus antibodies in sheep (Anderson et al., 1995; Longbottom et al., 2002).

However, the majority of sheep have preexisting background levels of antibody acquired from natural widespread infection by strains of *C. pecorum*. PCR appears to be an ideal alternate means of *Chlamydiaceae* detection because it offers advantages in sensitivity and reduced processing time over the conventional serological techniques. Furthermore, PCR does not rely on the presence and maintenance of viable organisms and poses less risk to laboratory staff than culture through eggs or cells. There have been many reports of PCR for the detection of *Chlamydomphila* nucleic acids in specific animals or from particular tissues (Hewinson et al., 1997; Laroucau et al., 2001; Messmer et al., 1997; Bomhardvon et al., 2003), but at the time of the manuscript developing there are limited publications on the impact of a pool of samples in preparation for molecular detection of *Chlamydomphila abortus* in aborted fetuses of ewes.

The aim of the present study was to determine the possible role of *Chlamydomphila abortus* ovis as an etiological agent of ovine abortion by PCR in a pool of samples of varying tissues of aborted fetuses.

Corresponding authors E-mail:

*Mahsa.Alem@gmail.com

Materials and Methods

Sampling Methods

During breeding season of 2011-2012, 50 abomasal aborted ovine fetus samples were collected from 10 flocks in Tabriz, northwest of Iran. Placentas were collected, macroscopically assessed for EAE lesions and representative cotyledons removed for molecular analysis. Following macroscopic examination, brain, skeletal muscle, liver, spleen, and abomasum were removed from each fetus. However, the condition of some of the fetuses was such that not all tissues could be collected. Fetal tissues and placental samples were then washed with PBS containing 1000 units/ml of penicillin (Pharmacia-Upjohn) and 1000 units/ml of streptomycin sulfate (Bristol-Myers Squibb) and then different tissues from each fetus were mixed in equal proportion and pulverized under liquid nitrogen. Approximately 200 mg of the sample powder stored as pool samples at -20°C until required for DNA extraction.

DNA Extraction of Tissues

Total DNA was extracted from approximately 200 mg powdered pool tissues using a commercial kit (Aquaprep DNA Tissue kit Bioneer, S. Korea) according to manufacturer's instructions. DNA concentration was measured at 260 and 280 nm (Biophotometer plus, Eppendorf, Germany). Electrophoresis of each DNA sample on 0.5% agarose gel in 1X Tris/Borate/EDTA (TBE) buffer was done to check the integrity of the DNA. A 60 μl aliquot of total DNA was produced from each sample and stored at -20°C until required for PCR analysis.

Polymerase Chain Reaction

DNA extract (5 μl) was used to amplify a 630-bp fragment of ompA gene of Chlamydomphila, using primers CHOMP191 (5'-GCI YTI TGG GAR TGY GGI TGY GCI AC-3') and CHOMP371 (5'-TTA GAA ICK GAA TTG IGC RTT IAY GTG IGC IGC-3'), as described by Kaltenbock et al. (1997) and modified by Sachse and Hotzel (2005). One μl of the amplicon was used as template for the second-round amplification, using the primer set CHOMP218 (sense) (5'-GTA ATT TCI AGC CCA GCA CAA TTY GTG-3')/ CHOMP336 (antisense) (5'-CCR CAA GMT TTT CTR GAY TTC AWY TTG TTR AT-3'), that yields a 400 bp product for *C. abortus*. Amplification reactions were carried out in a final volume of 25 μl , containing 100 ng of DNA, 0.5 μM of each primer, 2.5 μl 10X PCR buffer, 1.5 mM MgCl_2 , 0.2 mM dNTPs and 1 unit of Taq DNA

polymerase. The following cycles were applied: initial denaturation step at 95°C for 5 minutes followed by 35 cycles: denaturation at 94°C for 30 seconds, primer annealing at 50°C for 30 seconds, PCR products synthesis at 72°C for 30 seconds and final synthesis step at 72°C for 5 minutes. PCR products were recognized by electrophoresis on 1.5% agarose gel, stained with ethidium bromide and images were obtained in UVIdoc gel documentation systems (UK). Positive controls (DNA from references strains; ATCC 25116 and ATCC 2215), samples controls (DNA from abomasal fluids and lungs without etiological diagnostic), and none template control (distilled water) were included in each PCR run. PCR products with the molecular size of 440bp were considered indicative for identification as *C. abortus*.

Results

Genomic DNA was successfully extracted from aborted ovine fetus tissues using the DNA extraction kit. The PCR products of the primers specific for ompA gene (CTU-F and CTL-R) revealed the 630bp DNA fragment. Chlamydomphila was isolated in 13 out of 50 cases of ovine abortion (26%). The positive control showed the expected amplification product specific for Chlamydomphila (630 bp) and *C. abortus* (440bp). PCR products obtained from these samples and *C. abortus* reference strains with the molecular length of 440bp (according to the new taxonomy: Chlamydia and Chlamydomphila), are shown in Fig 1.

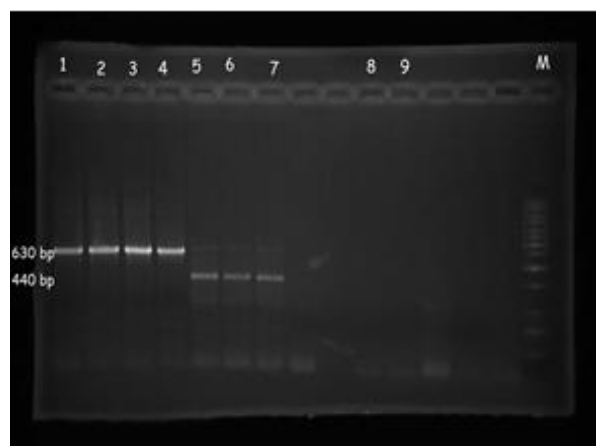


Figure 1. Ethidium bromide-stained 1.5% agarose gel electrophoresis of PCR products amplified using the mentioned primer. M, 100bp DNA ladder; lane 1, positive control sample of Chlamydomphila; lane 2-4, Chlamydomphila positive sample; lane 5, positive control sample of *C. abortus* ovis; lane 6-7, *C. abortus* ovis positive sample; lane 8, NTC (negative template control); lane 9, negative control (distilled water).

Discussion

OEA is one of the most important diseases affecting sheep flocks and because of abortion, weak lambs and a decrease in milk production can have negative economic effects. In previously reported serosurveys, the less-sensitive and -specific CFT had been used. Positive CFT titers can be caused by *C. abortus* as well as by *C. pecorum*.

According to the World Organisation for Animal Health (OIE) (2004), the most commonly used method for serodiagnosis of animal chlamydiosis the complement fixation test (CFT). However, the technique is laborious, has limited sensitivity and often impaired by cross-reactions between chlamydial species (McCauley et al., 2007). Recently developed serodiagnostic tests are mainly based on two main cross-reactive antigens present in all chlamydial species, lipopolysaccharide and the major outer membrane protein (MOMP) and thus, are not species-specific for diagnosing animals infected by OEA.

Other more specific tests need to be developed and evaluated in a European context, such as those based on specific monoclonal antibodies (Salti-Montesanto et al., 1997) and recombinant protein fragments (Sheehy et al., 1996). According to antigen detection, cultivation in cell culture is still regarded as the standard. While this time-consuming method is only applicable to cultivable strains, many strains are difficult to grow and not all laboratories have the facilities and expertise to culture. The possibilities for diagnostic detection of chlamydiae have considerably improved following the introduction of DNA-based methods, particularly the PCR, which permits direct identification from clinical specimens and differentiation of species. A number of tests have been published in the literature (Anderson et al., 1996; Kaltenbock et al., 1997; Messmer et al., 1997; Longbottom et al., 2001), but none of them has been validated in veterinary laboratories so far. One of the main difficulties in evaluating new tests for detection of *C. abortus* is a lack of a completely reliable method for comparative purposes.

Isolation, which was at one time used as the standard procedure has proved to have insufficient sensitivity (Wood et al., 1992; Domeika et al., 1994). Contamination of test samples with other bacteria, inadequate transport conditions, autolysis, antibiotic treatment, and other toxic factors may all interfere with the success of isolation (Sanderson et al., 1989). A further obstacle to isolation is the possible presence of chlamydiae in samples in the form of latent infection (the so-called “cryptic” form), which is well known in infections of the human uterine tube

and joints (Sachse et al., 2005).

Methods based on nucleic acid and antigen detection are recommended for diagnostic uses. Some authors have found the former (Thiele et al., 1992) and others the latter (Domeika et al., 1994; Trevejo et al., 1999), to be more sensitive in the detection of Chlamydia. In research by Lenzko et al. (2011), PCR and DNA microarray testing revealed the presence of chlamydiae in 78% of studied flocks. In sheep and goats, when *C. abortus* is present in the fetal membranes in large numbers, the two methods probably have almost identical sensitivity. In conclusion, our results showed that *C. abortus* is one of the important factors in abortions of sheep that are unknown and the vaccine has been applied to the control of abortions by this infection. Also, this study indicates that Chlamydia infection in the abomasal is an important factor for abortion in Iranian ovine.

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