

Molecular Detection of Anaplasmosis in Crossbred Cattle

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ABSTRACT

Bovine anaplasmosis is an infectious, non-contagious, tick-borne disease of domestic as well as wild ruminants caused by obligate intraerythrocytic parasites of family Anaplasmataceae and genus *Anaplasma*. Fifty blood samples were collected from crossbred cattle in a dairy farm with a history of a drastic decrease in milk yield and fever in a few animals. Six (12%) samples showed characteristic *Anaplasma* organisms when stained with Giemsa's stain, whereas 19 (38%) samples were found positive for *A. marginale* by PCR assay. Low hemoglobin and total erythrocyte count were observed, whereas an increase in total leucocyte count was noted in *Anaplasma* affected animals compared to healthy animals at the same farm.

Keywords: Anaplasmosis, Cattle, Hematology, Molecular diagnosis, PCR.

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INTRODUCTION

Anaplasmosis is the most common cause of economic losses in the livestock industry throughout the world. Bovine anaplasmosis is a parasitic disease of cattle caused by the obligate intracellular (erythrocytic) rickettsia *Anaplasma marginale*, primarily occurring in tropical and subtropical regions of the world. Although cattle, sheep, goats, buffalo, and some wild ruminants can be infected with the erythrocytic *Anaplasma*, cattle have been more susceptible than buffaloes (Rajput *et al.*, 2005). Most commonly, *Rhipicephalus (Boophilus) microplus* (Asian blue tick) are biological vectors of *A. marginale*, but the pathogen is often transmitted mechanically to susceptible cattle by blood-contaminated mouthparts of biting flies or by needles and veterinary instruments that become contaminated with blood. In cattle aged > 2 years, *A. marginale* causes persistent fever, lethargy, icterus, weight loss, abortion, decreased milk yield, and death in more than 50% of untreated animals (Kocan *et al.*, 2003 and M'Ghirbi *et al.*, 2016)

The standard technique for diagnosing bovine anaplasmosis has traditionally been done by microscopic examination of Giemsa-stained blood smears revealing > 10⁶ infected erythrocytes/ml (Gale *et al.*, 1996). In the acute phase, inclusion bodies are easily detected in bovine erythrocytes (Maharana *et al.*, 2014) by microscopic examination. In contrast, this technique is not easily applicable for detecting chronic and sub-clinical infections. Such infected animals serve as a reservoir for spreading *A. marginale* (Eriks *et al.*, 1989). Polymerase Chain Reaction assays allow diagnosis of parasite at levels far below the detection limit of the frequently used parasitological techniques and have an advantage in separating clinical and subclinical forms of parasitic infection (Almeria *et al.*, 2001). In the present study, a PCR assay was used along with microscopy to detect anaplasmosis in crossbred cattle.

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MATERIALS AND METHODS

The present study was conducted on a dairy farm of district Ambala, Haryana, in the year 2019. Blood samples (n = 50) in EDTA vial were collected from crossbred cattle having fever in a few animals and a drastic decrease in milk yield. Blood smears were prepared on clean glass slides. Smears were fixed with methanol, and let to dry in the air, then stained with Giemsa's stain and examined under a microscope (X1000) with immersion oil for the presence of haemoparasites infections. The haemoparasites were identified to species level as per morphological characters as described by Soulsby (1982). Various hematological parameters such as hemoglobin (Hb), total erythrocyte count (TEC), and total leucocyte count (TLC) were compared between healthy (n=8) and affected animals (n = 19). The blood samples were kept at 4°C and further analyzed by PCR. DNA was extracted from the blood samples using a DNA extraction

kit by Qiagen. *Anaplasma marginale* specific primers for the gene *msp4* (Forward primer 5'-CTGAAGGGGAGTAATGGG-3' and Reverse primer 5'-GGTAATAGCTGCCAGAGATTCC-3') were used for diagnosis and amplification of 344 bp gene product (Torina *et al.*, 2012). The amplified PCR products were separated by electrophoresis on 1% agarose gel and visualized under UV transilluminator for detection of 344 bp amplified product in positive cases. A sample with heavy parasitemia of *A. marginale* was used as a positive control for the standardization of PCR assay. A no-template reaction was also used as a negative control for all the PCR reactions.

RESULTS AND DISCUSSION

Out of 50 blood samples, only six (12%) samples showed characteristic rickettsial organisms when stained with Giemsa's stain as well as PCR (Fig. 1). The PCR product of positive control yielded a specific band of 344 bp on gel electrophoresis while there was no band in the negative control. In addition, 13 more samples were positive for *A. marginale* by PCR. Overall, 19 (38%) samples were found positive for *A. marginale* by PCR. Gel electrophoresis of PCR products from these 19 samples showed a specific fragment of 344 bp (Fig. 2). The blood samples found negative by PCR were also found negative for *A. marginale* on blood smear examination. These samples belonged to animals whose production performance was poor and gradually decreased. Various studies revealed that anaplasmosis is prevalent in the cattle population in Haryana. Ganguly *et al.* (2017) observed *T. annulata* (33.32%), followed by *B. bigemina* (4.76%),

A. marginale (1.67%), and mixed infection of these parasites (1.58%) in crossbred cattle in Haryana. Bhanot (2020) also reported haemoparasitic infection in crossbred cattle and found theileriosis (44.6%), followed by anaplasmosis (3.4%) and babesiosis (2.6%) to be the major disease problems.

Recent molecular technology has gained a lot of importance in the diagnosis of haemoparasitic infection. Conventional parasitological techniques like microscopic examination of Giemsa stained thin blood smear is always a standard gold test for diagnosing anaplasmosis in clinically suspected animals, while in subclinical and chronic infections Giemsa stained thin blood smear is not suitable (Eriks *et al.*, 1989; Ashuma *et al.*, 2013). The 2 X 2 contingency table for blood smear examination and PCR is shown in Table 1. The result revealed that the PCR assay is 100% sensitive (true positive (TP)/true positive (TP) + false negative (FN) × 100) and 70.4% specific (true negative (TN)/true negative (TN) + false positive (FP) × 100) when compared with microscopy based detection method. This further asserts its usefulness in rapid

Table 1: 2 x 2 contingency table for blood smear examination and PCR

	Blood Smear Examination		Total
	Positive	Negative	
<i>PCR</i>			
Positive	6 ^a	13 ^b	19
Negative	0 ^c	31 ^d	31
Total	6	44	50

^a True positive, ^b False positive, ^c False negative, ^d True negative
Sensitivity of PCR = $a / (a+c) \times 100 = 100\%$; Specificity of PCR = $d / (b+d) \times 100 = 70.4\%$

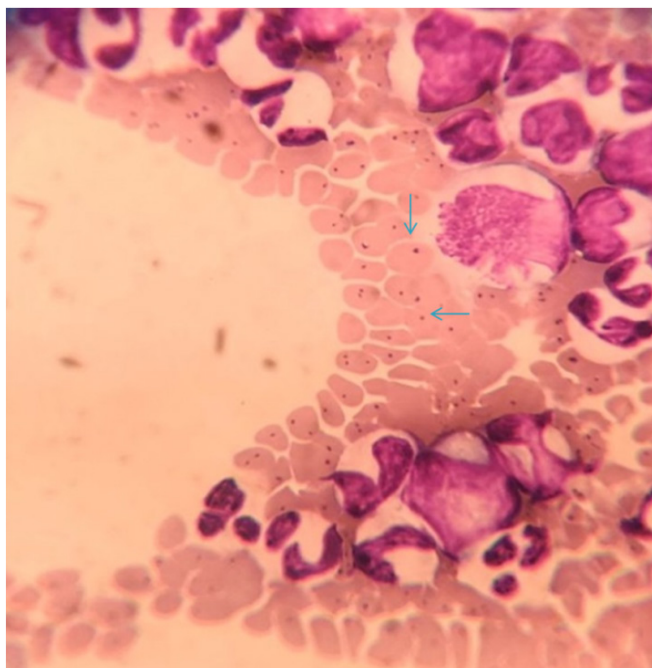


Fig. 1: *Anaplasma* organisms on the margin of RBC by microscopic examination (X1000) blood smears

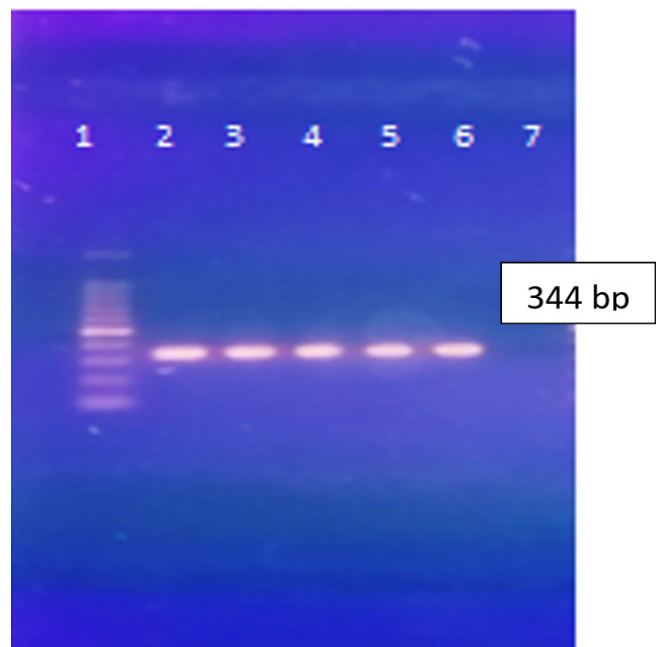


Fig. 2: Agarose gel electrophoresis of amplified 344 bp product by PCR using *msp4* specific primers for *A. marginale* Lane 1: DNA marker (100bp DNA Ladder), Lane 2: Positive Control, Lane 3-6: Sample tested, Lane 7: Negative Control

diagnosis of *A. marginale* with high specificity and sensitivity especially in carrier animals/subclinical infections with very low parasitemia. In the case of a carrier state with low levels of parasitemia, *Babesia* and *Anaplasma* organisms may not be found on microscopic examination (Atif *et al.*, 2012b). For confirmatory diagnosis, PCR is used for detection even in case of a low level of parasitemia (Ge *et al.*, 1995; Torina *et al.*, 2008). Bisen *et al.*, 2021 also reported molecular detection by msp5 nested-PCR as highly sensitive and reliable for detecting carrier cattle for *Anaplasma* infection.

In the present study, hematological parameters (Hb, TEC $\times 10^6$ and TLC $\times 10^3$) were compared between healthy and affected animals. The results of the study revealed Low Hb (5.6 ± 0.25) and TEC (4.02 ± 0.275) levels with increase in TLC (9.4 ± 0.412) count as compared to healthy animals (Hb- 10.25 ± 0.72 ; TEC -7.02 ± 0.58 ; TLC- 6.01 ± 0.35). The values for different cells of leucocyte (Mean \pm SE) are presented in Table 2 reveals that neutrophil and monocyte count in the infected cattle were low whereas lymphocytes and eosinophil count were high as compared to healthy animals. The major sign of anaplasmosis infection is extravascular hemolytic anemia in cattle (Ajayi *et al.*, 1978; Kuttler, 1984; Atif *et al.*, 2012a). A high risk of anemia could be due to phagocytosis of infected erythrocytes by the bone marrow cells and spleen (Jain, 1993). Normally there is simultaneous hemolysis of intravascular and extravascular RBCs (Riond *et al.*, 2007). Destruction of the non-parasitized RBCs also starts in addition to the breakdown of parasitized RBCs due to immune-mediated autolysis. The disease also adversely affects the milk production of cattle. This alterations in haematological parameters is in agreement with Ganguly *et al.*, (2018) who observed significantly decreased levels of TEC, Hb and PCV in animals infected with *Anaplasma* than healthy control animals. Ashuma *et al.* (2013) reported that haemogram of the *Anaplasma* infected animals revealed anaemia as indicated by significant decrease ($P < 0.05$) in RBC, Hb and PCV in both the infected groups positive by GSTBS and PCR as compared to the healthy control group.

It can be deduced that Anaplasmosis is associated with alteration in haemogram which may results in production loss. Although microscopy is widely accepted and cost

effective technique for the diagnosis of haemoprotozoans and haemoreticidal organisms, the technique is not highly sensitive. Further, subclinical infection of anaplasmosis can be better detected by molecular methods. Appropriate tick control strategies are also to be adopted for containing the infection at reduced level in the herd in endemic areas.

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Table 2: Haematological parameters of *Anaplasma* positive and healthy animals

Parameter	Positive (N=19) (Mean \pm SE)	Healthy control (N=8) (Mean \pm SE)
Hb(g/dl)	5.6 \pm 0.25	10.25 \pm 0.72
TLC($\times 10^3$)	9.4 \pm 0.41	6.01 \pm 0.35
TEC ($\times 10^6$)	4.02 \pm 0.27	7.02 \pm 0.58
Neutrophil	38.15 \pm 1.9	52.15 \pm 3.6
Lymphocyte	58.84 \pm 1.7	46 \pm 3.8
Monocyte	0.63 \pm 0.16	0.75 \pm 0.16
Eosinophil	1.89 \pm 0.24	0.5 \pm 0.19

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