

Detection of *Babesia ovis* in small ruminants by using microscopic and molecular techniques

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The present study was planned to detect *Babesia ovis* (*B. ovis*) in naturally infected sheep and goat herds of Faisalabad, Toba Tek Singh and Jhang districts of Punjab, Pakistan through microscopy and molecular diagnostic techniques *i.e.*, PCR and nPCR. For this purpose, a total of 892 blood samples were collected *i.e.*, sheep (n=544), goats (n=348) in both seasons (summer + winter). Initially, the samples were screened via the blood film and the blood smear examinations revealed parasite merozoites as tear drop in pair identified as babesia. The overall prevalence in summer based upon microscopic analysis was 22.05% (60/272) in sheep and 16.05% (28/174) in goat, respectively. In winter the prevalence was 7.34% (20/272) and 4.59% (8/174) in sheep and goat, respectively. In summer the results based upon PCR were 29.7% (81/272) and 23.56% (41/174) and n-PCR were 31.98% (87/272) and 27.01% (47/174) in sheep and goats respectively. On the other hand, in winter results showed that 12.13% (33/272), 9.77% (17/174) and 13.60% (37/272), 13.21% (23/174) samples positive for *B. ovis* through PCR and nPCR respectively. The results have shown that sensitivity of NPCR is more as compared to conventional PCR. The product size of *B. ovis* through PCR and NPCR was 186 base pair (bp) and 549 bp. Furthermore, the prevalence based upon different risk factors *i.e.*, age, sex, housing, floor system, presence or absence of ticks was determined. Statistical analysis of different risk factors indicated that, increased age, non-cemented floor, closed housing, weak body condition and heavy infestation of ticks increase the incidence of disease by many folds.

Keywords: *Babesia ovis*, PCR, NPCR, prevalence, small ruminants.

INTRODUCTION

Livestock sector plays an important role in Pakistan's economy and share of livestock in national GDP is 11.6% along with it contributes about 55.1% of agriculture value added during 2011-2012. Most of the rural economy of developing countries is dependent directly or indirectly on livestock (Abdel-Rahman *et al.*, 2020; Bukhari *et al.*, 2020; Rahat *et al.*, 2020; Štrbac *et al.*, 2021). However, livestock production systems have always been threatened with huge burden of infectious diseases (Anees *et al.*, 2020; Muhammad *et al.*, 2020). Among livestock, rearing of sheep and goats has much importance in rural areas especially for non-agricultural poor families (Shahzad *et al.*, 2013). Total number of heads of sheep and goats in Pakistan assessed as 28.24 M and 49.14 Million, respectively (Shahzad *et al.*, 2013) and they yielded 0.040 and 0.915 billion liters of milk for human consumption, 0.629 Million tons of meat and 0.043 Million tons of wool during 2011-2012. Despite of huge number of animals in Pakistan production is very low, reasons behind the low

production are many few of them are poor breeding, poor management, improper housing and increased infestation of ticks, which lead to increased prevalence of hemoparasitic diseases. Pakistan being in the sub-tropical region in the world has environment, which favors the growth of ticks, which play an important role in the transmission of such diseases in small ruminants such as babesiosis. Ovine babesiosis develops in RBCs of peripheral blood of small ruminants (Abed *et al.*, 2017).

Three species of genus *Babesia* namely *B. motasi*, *B. crassa* and *B. ovis* are responsible for disease in small ruminants. *Babesia motasi* and *B. crassa* are considered to be less pathogenic and infection is mostly subclinical, while *B. ovis* is extremely pathogenic and instigates severity of infection along with 30-50% mortality rate in sheep and goats particularly in aged animals (Mathieu *et al.*, 2018; Hurtado *et al.*, 2018). The main clinical signs of babesiosis in sheep and goats include high fever, anemia, hemoglobinuria, depression and loss of appetite, as a consequence death may occurred (Rodastits, 2007; AbouLaila *et al.*, 2021). Because of

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Table 1. Primer sets used for *Babesia ovis* DNA detection.

Parasite	Assay	Primer name	Primer sequence	Size	Reference
<i>B. ovis</i>	PCR	Bov-F	TGGGCAGGACCTTGTTGTTCT	549bp	(Esmaeilnejad <i>et al.</i> , 2014)
		Bov-R	CCGCGTAGCGCCGGCTAAATA		
	nPCR	iBov-F	AAGAATTTACCTATGACAG	186bp	(Haghi <i>et al.</i> , 2013)
		iBov-R	GTCTGCGCGCGCCTTTGCG		

zoonotic nature some of *Babesia* species cause diseases in human population so increased emergence of such tick-borne parasites are considered to be a major threat to public health (Fakhar *et al.*, 2012; Zheng *et al.*, 2018; Young *et al.*, 2019). Different investigators have reported the occurrence of babesiosis in different parts of country, which were geographically different from the study like PCR based prevalence of babesiosis in sheep was 50% (Iqbal *et al.*, 2011) and Durrani *et al.*, (2012) documented other hemoparasite incidence that was 35%. However, before the present study few data is available on molecular prevalence of babesiosis in small ruminants through conventional and nested PCR in Faisalabad, Pakistan.

Traditionally, parasites are detected by clinical signs and microscopy (Hussain *et al.*, 2017; Karatepe *et al.*, 2019). These methods are insufficient to diagnose the carrier animals with low parasitemia. The disadvantages of conventional diagnostic methods can be improved by using modern molecular diagnostic approaches (Wang *et al.*, 2018).

Keeping in view the benefits of molecular techniques like PCR and nested PCR (nPCR) in terms of high specificity and sensitivity, the current study was planned to detect *Babesia ovis* in naturally infected sheep and goat herds of Faisalabad, Toba Tek Singh and Jhang districts of Punjab, Pakistan. Before that no such kind of work has been reported in purposed areas.

MATERIALS AND METHODS

Sample collection and experimental ethics: A total of 892 blood samples in both seasons (summer + winter) from ovine were collected *i.e.*, sheep (n=544), goats (n=348) with a history of high fever or presence of ticks on the body residing in Faisalabad, Toba Tek Singh and Jhang districts in Punjab, Pakistan. Samples were taken in tube free from any kind of anticoagulant. Thin blood smears were formed and fixed with absolute methanol and stained with Giemsa, all the laboratory precautions were followed. The presence of piroplasm was determined by examining the blood smears under light microscope with 100X magnification. Piroplasm were identified by morphological characteristics.

DNA Isolation: *Babesia* genomic DNA was extracted from suspected sheep and goat blood as depicted beforehand (Almeria, 2001). Phenol chloroform isoamyl liquor technique was utilized for separating DNA. Erythrocytes were slicked in lysis cradle, at that point 20 ml proteinase K (10 mg/ml) was added, and the incubation for 2 hours at 56°C was done

to process the proteins. At that point, phenol chloroform isoamyl liquor was added and centrifuged at 13400 rpm for 15 min. Upper fluid layer was transferred to another smaller scale eppendorf containing ethanol 96% (2.5 ml volumes of the sample) and centrifuged at 13400 rpm for 15 min. Subsequent to washing with ethanol 70%, at long last removed DNA air-dried, disintegrated in TE cushion (10mM Tris-HCl pH 8, 0.1mM EDTA) at 55° C, and kept up at - 4° C for 24 hours at that point kept at - 20° C until use.

PCR Analysis: So as to amplification of DNA parts of *B. ovis*, the PCR and nPCR methods were utilized with explicit ground works previously revealed for *Babesia* spp. (Table 1) got from flanking part of hyper variable locale of 18SrRNA. The PCR was acted in an all-out volume of 25 µl including 5µl of DNA master mix, 1 x PCR cradle, 0.1 U Taq polymerase, 0.5 µl of each primer (P1/P2, 20 mM), 125 µM of each of dNTPS (Fermentas) and 1.5 mM MgCl₂ in a programmed DNA Thermocycler (Bio-Rad); with the accompanying conditions: 5 min at 95°C to denature twofold strand DNA, 38 patterns of 45s at 94°C, 45s at 56°, 45s at 72°C and a last expansion step for 10 min (Esmaeilnejad *et al.*, 2014). The intensified items will be settled by 1.5% agarose gel electrophoresis and recolored with ethidium bromide for visual location by bright transillumination. Tests introducing noticeable groups of around 549 base sets (PCR) thought about positive for *B. ovis*, PCR positive groups were of 549 bp for *B. ovis* (Fig. 5).

nPCR Analysis: Recently portrayed conventions were utilized for the nPCR measures for location by explicit enhancement of DNA pieces of *B. ovis*. Experiment was performed in a volume of 25 µl: 12.5 µl of Master Mix, 5.5 µl of ultrapure water, 10 pM of each primer (Table 1) and 5 µl of DNA fragment. The nPCR thermocycling conditions for *B. ovis* were: 1 min at 94 °C, 1 min at 62 °C and 1 min at 72 °C. The nPCR conditions were indistinguishable with the exception of the tempering advance, which was at 69°C for 30 seconds (Haghi *et al.*, 2013). Decontaminated tests of *B. ovis* was submitted to DNA extraction and utilized as positive control. DNA tests were viewed as positive if in nPCR explicit groups will be imagined on 1.5% agarose gel *B. ovis* 186 bp (Fig. 4).

RESULTS

In this study a total of 892 blood samples in both seasons (summer + winter) from ovine were collected *i.e.*, sheep (n=544), goats (n=348) residing in Faisalabad, Toba Tek Singh and Jhang districts in Punjab, Pakistan. First of all, the samples were screened via the blood film in both seasons (summer & winter) and the blood smears examination revealed parasite as tear drop in pair identified as *Babesia* and the overall microscopic prevalence in summer was 22.05% (60/272) and 16.05% (28/174) in sheep and goat, respectively and in winter was 7.34% (20/272) and 4.59% (8/174) in sheep and goat, respectively.

Furthermore, the prevalence based upon different risk factors *i.e.*, age, sex, housing, floor system, presence or absence of ticks was determined. Statistical analysis of different risk factors (Table 2) (Fig. 2) indicated that, age ($P = 0.026$), sex ($P = 0.048$), floor ($P = 0.033$), housing ($P = 0.047$), body condition ($P = 0.033$), tick burden ($P = 0.036$) increased age, female, non-cemented floor, closed housing, weak body

condition and heavy infestation of ticks increase the incidence of disease by many folds.

The district wise prevalence of babesiosis in goats and sheep was also vary in both seasons *i.e.* in summer in goats the prevalence was 15.38% (12/78) in Faisalabad, 12.50% (6/48) in Toba Tek Singh and 20.83% (10/48) in Jhang, respectively, while in sheep prevalence was 10.90% (12/110), 20.00% (16/80) and 39.02% (32/82) in Faisalabad, Toba Tek Singh and Jhang respectively. While in winter the prevalence of *Babesia ovis* in goats was 5.12% (4/78) in Faisalabad, 2.08% (1/48) in Toba Tek Singh and 6.26% (3/48) in Jhang respectively, while in sheep *B. ovis* prevalence in Faisalabad was 5.66% (03/53), 3.44% (03/87) in Toba Tek Singh and 10.60% (14/142) in Jhang, respectively (Table 3) (Fig. 3).

The samples were further analyzed through the conventional PCR and nPCR. In summer the results based upon PCR were 29.7% (81/272) and 23.56% (41/174) and nested PCR were 31.98% (87/272) and 27.01% (47/174) in sheep and goats

Table 2. Prevalence of *Babesia ovis* in ovine in relation to different risk factors in both seasons.

Risk factors	Variables	Summer			Winter		
		Prevalence %	Chi-square	P-Value	Prevalence %	Chi-square	P-Value
Age	Young	22/214(10.28)	34.22	0.038	07/220(3.18)	38.35	0.026
	Adult	66/232(28.44)			21/216(9.72)		
Sex	Female	76/338(22.48)	16.43	0.026	23/306(7.51)	17.22	0.048
	Male	12/108(11.11)			05/140(3.57)		
Feeding System	Ground	53/210(25.23)	18.33	0.053	20/229(8.73)	21.42	0.053
	Trough	35/236(14.83)			08/217(3.68)		
Floor	Cemented	20/203(9.85)	17.43	0.045	07/232(3.01)	14.32	0.033
	Non-cemented	68/243(27.98)			21/214(9.81)		
Housing	Open	23/213(10.79)	28.33	0.028	04/226(1.76)	28.21	0.047
	Closed	63/233(27.03)			24/220(10.9)		
Herd size	>20 Animals	58/231(25.10)	19.33	0.046	21/209(10.0)	21.44	0.056
	<20 Animals	28/215(13.02)			07/237(2.95)		
Body conditions	Good	14/209(6.69)	11.31	0.062	04/206(1.94)	13.43	0.033
	Poor	64/237(27.00)			24/240(10.0)		
Ticks	Present	55/203(27.09)	22.53	0.057	22/224(9.82)	27.43	0.036
	Absent	33/243(13.58)			06/222(2.70)		
Dog Association	Present	56/213(26.29)	20.43	0.035	19/226(8.40)	32.45	0.065
	Absent	32/233(13.73)			09/220(4.09)		

Table 3. District wise prevalence of *Babesia ovis* in small ruminants in summer and winter season.

Districts	Specie	Summer			Winter		
		Prevalence %	Chi-square	P-Value	Prevalence %	Chi-square	P-Value
Faisalabad	Goat	12/78(15.38)	12.53	0.039	04/78(5.12)	15.64	0.042
	Sheep	12/110(10.90)			03/53(5.66)		
Toba Tek Singh	Goat	06/48(12.50)	16.36	0.042	01/48(2.08)	20.53	0.049
	Sheep	16/80(20.00)			03/87(3.44)		
Jhang	Goat	10/48(20.83)	21.53	0.050	03/48(6.25)	19.43	0.027
	Sheep	32/82(39.02)			14/132(10.60)		
Total	Goat	28/174(16.09)	18.55	0.030	08/174(4.59)	14.74	0.064
	Sheep	60/272(22.05)			20/272(7.35)		

respectively. On the other hand, in winter results showed that 12.13% (33/272), 9.77% (17/174) and 13.60% (37/272), 13.21% (23/174) samples positive for *B. ovis* through PCR and nPCR respectively. The results have shown that sensitivity of nPCR is more as compared to conventional PCR. The product size of *B. ovis* through PCR and nPCR was 186 bp and 549 bp (Fig. 4 and Fig. 5)

DISCUSSION

Tick borne diseases (TBDs) are major threat to animals particularly in small ruminants common in temperate zone in tropical and sub-tropical countries (Misgana *et al.*, 2017; Zeb *et al.*, 2020). Most of them are zoonotic in nature and leads to severe economic losses to a country in terms of decreased production and trade allegations (Shah *et al.*, 2017; Jelcic *et al.*, 2020). Babesiosis is one of them, babesia was first discovered by Babes (1888) in the cattle populations of Romania that caused high mortalities; previously it was considered as bacteria. The prevalence of babesiosis in different parts of the world ranges from 0% to 100% (Jorgensen *et al.*, 1992; Noaman, 2013).

Babesiosis is characterized by fever, decreased production, poor quality wool, anemia, hemoglobinuria and birth of weak offspring. Sheep and goats are prone to babesiosis and the disease endemic in different areas of Pakistan (Shahabuddin *et al.*, 2006; Rashid *et al.*, 2010; Iqbal *et al.*, 2011; Durrani *et al.*, 2011; Durrani *et al.*, 2012; Naz *et al.*, 2012). Babesiosis is commonly diagnosed by microscopic examination (Reddy *et al.*, 2016; Haq *et al.*, 2017; Vannier *et al.*, 2020) but this practice is useful in acute cases where the parasitemia is high. Drawback of this technique is that subclinical infections and carrier animals can not be detected by this method. Another assay which is indirect fluorescent antibody assay (IFAT) also used previously but this method has also some limitations like lack of skilled persons in reading the microscopic slides of specimen (Persichetti *et al.*, 2017). It is need of time to develop a more sensitive tool and accurate treatment of disease to decrease the economic losses because of the babesiosis. Different researchers were agreed with the fact that PCR and nPCR are more sensitive and specific tool for detection of hemoparasites without showing any apparent signs which mostly occurs in carrier animals (Talkhan *et al.*, 2010; Shah Nawaz *et al.*, 2011; Ziapour *et al.*, 2011). In the past sheep and goats were investigated separately either by PCR or nPCR, but the current study firstly, encompasses the collectively investigation of sheep and goat by two molecular techniques.

In the current study, the overall prevalence in summer based upon microscopic analysis was 22.05% (60/272) in sheep and 16.05% (28/174) in goats. In winter the prevalence was 7.34% (20/272) in sheep and 4.59% (8/174) in goat respectively. The prevalence reported on PCR and nPCR based upon molecular techniques was 29.7% (81/272) and 23.56% (41/174), 31.98% (87/272) & 27.01% (47/174) in sheep and goats respectively in summer. On the other hand, in winter results showed that

12.13% (33/272), 9.77% (17/174) and 13.60% (37/272), 13.21% (23/174) samples positive for *B. ovis* through PCR and n-PCR respectively. Similar findings have been reported by Iqbal *et al.*, (2012) in southern Punjab, Pakistan. They reported 50% samples positive through molecular techniques i.e., PCR and 3% detected positive via microscopic analysis. Similarly, 18% blood samples were reported positive through PCR as compared to microscopic 3% (Zulifqar *et al.*, 2012). Further risk factors analysis of data revealed that, adult animals were more prone to babesiosis (28.44%) ($P = 0.026$) as compared to young ones. This is because of adult animals are active grazers and when they go out for grazing, they are more at risk to ticks secondly immunity decreases with increase in age, these findings were in accordance with previous literature (Niazi *et al.*, 2008). Females were more infected with babesiosis (22.48%) ($P = 0.048$) as compared to males. High prevalence in females might be due to the fact that females were reared for longer period of time, and female are more prone to physiological stress like milk production, breeding, parturition and post parturition as compared to male (Alim *et al.*, 2011; Kabir *et al.*, 2011). High occurrence of disease was associated with non-cemented floor system (27.98%) ($P = 0.033$) as compared to cemented floor system, because in non-cemented there is no proper drainage system, cleanliness of shed along with some cracks in the floor which harbor the parasites that why the eggs of ticks were not properly removed that leads to the incidence of babesiosis in animals. Similar results have been reported by Roy *et al.*, (2018). Disease was found to be more prevalent in closed housing system (27.03%) ($P = 0.047$) as compared to open housing system. This is because of zoonotic nature of disease was more frequent in intensive farming (Tolkacz *et al.*, 2017). Weak animals were more infected with babesiosis (27%) ($P = 0.033$), as compared to healthy ones (Ward *et al.*, 2018). It was also noted that animals with high tick infestations were more suffered with *Babesia* infection (27.09%) ($P = 0.036$) as compared to less tick burden. This finding was similar to Surez *et al.*, (2017). The transmission of disease is due to ticks because they act as an intermediate host (Ybanez *et al.*, 2019). According to current study presence of dogs ($P = 0.065$) and herd size ($P = 0.056$) has no effect on the prevalence of disease (Theodoropoulos *et al.*, (2006).

Conclusion: *Babesia ovis* is much prevalent in sheep and goat herds that leads to huge economic losses in terms of poor growth rate, poor quality wool and increased treatment cost after disease outbreak. Commonly smear method is used for detection of blood parasite i.e., babesia, this is not useful for detection of carrier animals or very low parasitemia. Therefore, current study was conducted to investigate the babesiosis in ovine and caprine species in Faisalabad Division. PCR for the detection of *B. ovis* is specific and sensitive. The test is suitable for tracing carrier animal and provide a qualitative and validated measure that is useful in epidemiological surveys, follow ups for drug treatment and

ticks control programs in endemic areas to enhance the livestock productivity.

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Conflict of interest: There is no conflict of interest

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