



Intrinsic Risk Factors for Bovine Tropical Theileriosis in Indian Cattle: A Meta-Analysis

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Received: 26 March, 2021

Revised: 19 April, 2021

Accepted: 25 April, 2021

ABSTRACT

Meta-analysis of systematically selected studies published during 2015-2020 on the prevalence of bovine tropical theileriosis from Indian cattle was done. It was found that age, sex and breed of cattle are important risk factors to get infected with *Theileria* sp. Adult cattle are found at higher risk of getting infected with BTT than the younger stocks with an odds ratio of 2.062, however as regards sex of the cattle is concerned cows are little more susceptible to BTT than bullocks / bulls with an odds ratio of 1.632. The breed factor of cattle has shown that exotic/ crossbred cattle are more prone to contract BTT with an odds ratio of 2.113. The meta-analysis showed no publication bias.

HIGHLIGHTS

- Age, sex, and breed of cattle are major risk factors for Theileriosis in Indian cattle.
- Very few studies recorded those risk factors.
- More studies on epidemiological aspect of bovine tropical Theileriosis are needed.

Keywords: *Theileria*, cattle, risk factors, MetaXL

In dairy animals vector borne haemoprotozoan diseases are of prime importance considering their impact on production and substantial economic costs ultimately upsetting the dairy enterprises. With significant morbidity and mortality in its ruminant hosts, theileriosis is a major impediment in dairy industry globally (Demessie and Derso, 2015). Bovine tropical Theileriosis (BTT) is a tick borne haemoprotozoan disease caused by an apicomplexan parasite *Theileria* spp., the obligate intracellular protozoan parasite which infects domestic and wild ruminants. The ticks of Ixodidae family are the vectors of this protozoa and they have a complex life cycle both in vertebrate and invertebrate hosts. Amongst several species of *Theileria* genus that infect cattle, *T. parva* and *T. annulata* are most pathogenic and prevalent species. BTT is very lethal and debilitating hemoprotozoan disease infecting cattle which is caused by *T. annulata* and transmitted by *Hyalomma* sp ticks (Mans *et al.*, 2015). The disease is clinically characterized by high fever and enlargement

of superficial lymph nodes. As per earlier estimate about 33 million Indian cattle are at risk of BTT (Minjauw and McLeod, 2003) and annual economic losses due to this disease approximately amounts to ₹ 8426 crores in India (Narladkar, 2015).

As the sub-tropical climatic conditions of Indian subcontinent are extremely conducive for ticks and therefore for tick borne disease, BTT is a major disease dairy animals are exposed to in this region including India. Several earlier studies from across India reported prevalence of BTT of different extent attributed to various risk factors *viz.* age, sex, breed, season, management practices *etc.* (Gul *et al.*, 2015). Various factors such as management practices, sex, health status, nutritional

How to cite this article: Bhangale, G.N., Prakash, B.K.C. and Dhabale, A.R. (2021). Intrinsic risk factors for bovine tropical theileriosis in Indian cattle: A meta-analysis. *J. Anim. Res.*, 11(3): 497-503.

Source of Support: None; **Conflict of Interest:** None



deficits, breed, herd size, agro-climate (humidity, temperature) directly affects the livestock health and pose them to get this disease (Ghosh and Nagar 2014, Ayadi *et al.*, 2016). Considering its importance epidemiological studies have since long being conducted in India to estimate its spread so as to formulate effective control strategies to minimize the losses incurred by BTT.

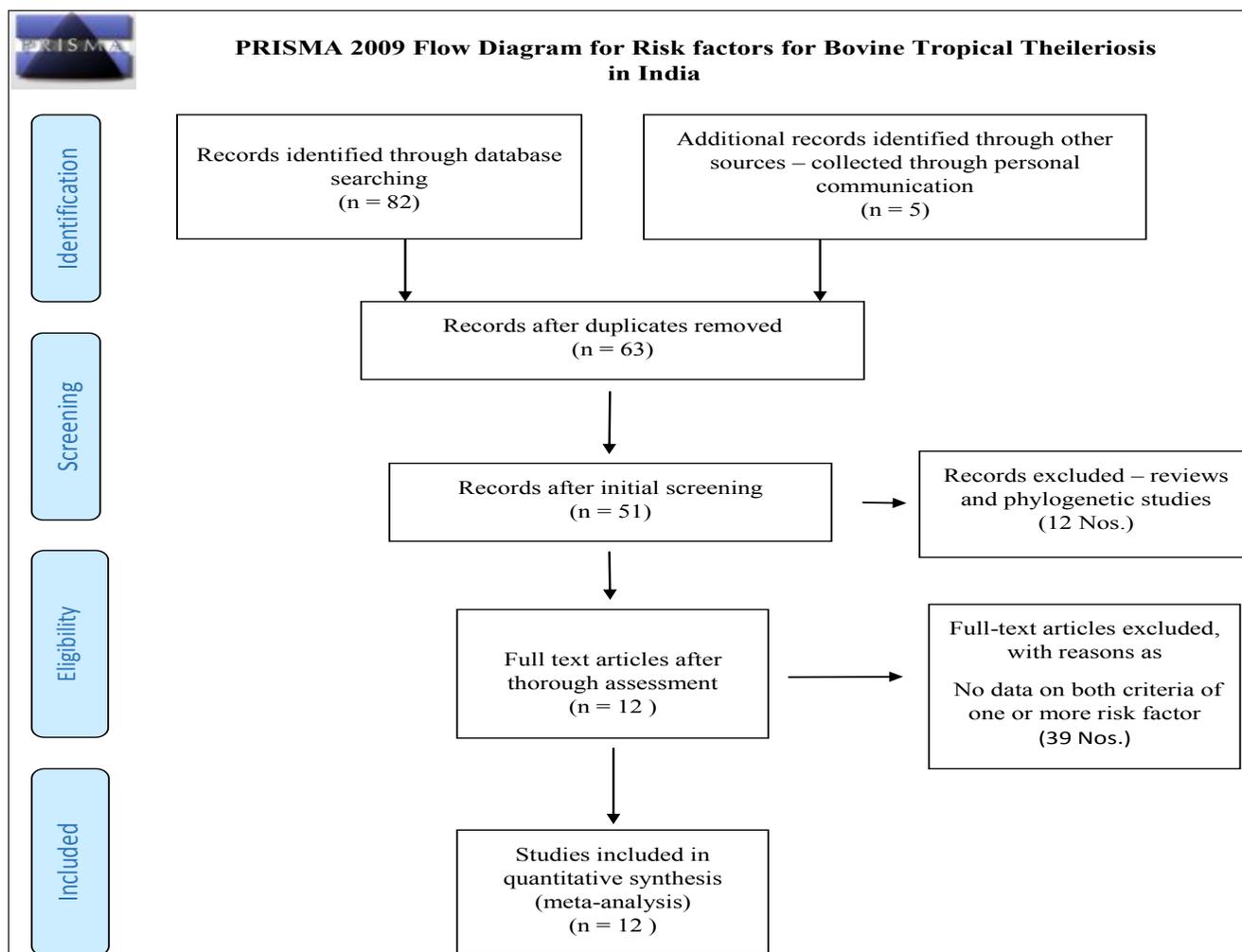
The present study has detailed on animal bio-markers as the risk factors for BTT in India though meta-analysis approach based on earlier published reports.

MATERIALS AND METHODS

Present study is based on meta-analysis of earlier reports on risk factors for BTT in India. This meta-analysis was

conducted according to criteria laid down under PRISMA Guidelines (Moher *et al.*, 2009). The studies were selected for this meta-analysis were spanned over years 2015-2020. The studies were selected through a systematic search from online free databases viz. GoogleScholar, PubMed, IndianJournals. The keys used for this search were Theileria, Theileriosis, Bovine, Cattle, India with combinations by using Boolean words *i.e.* AND, OR. the criteria for selection of studies were such that the study to be selected should have reported prevalence of Theileriosis in cattle based various risk factors including age, sex and breed as the biomarkers of the cattle. The search and strategic selection based on the aforementioned criteria yielded 12 studies for final meta-analysis.

The requisite data on the BTT prevalence estimates based



Moher *et al.* (2009). www.prisma-statement.org

on risk factors were extracted from the selected studies and entered in an MS-excel sheet. Depending on the extent of heterogeneity, the random effects model and fixed effects model was employed for this meta-analysis. When heterogeneity was above 75%, random effects model was employed otherwise fixed effect model was used.

Risk factors for BTT as age, sex, breed was considered for this analysis and accordingly data was categorized. Studies with exact and essentially with both the number of animals tested and the number animals found positive for at least one of these factors was included in this analysis. Individual spreadsheet for data on each of the risk factors was prepared and subjected to meta-analysis. Meta-analysis was performed on MetaXL add-in with the help the command NumOR and results were presented through forest plot. Heterogeneity and publication bias were assessed and presented accordingly.

Table 1: Characteristics of the included studies on risk factor wise prevalence of BTT in India

Risk Factor – Age	Adult		Young	
	Total	Positive	Total	Positive
Kundave <i>et al.</i> , 2015	74	59	15	7
Tuli <i>et al.</i> , 2015	840	243	241	33
Agrawal <i>et al.</i> , 2016	83	26	55	2
Naik <i>et al.</i> , 2016	115	28	35	7
Dadhich <i>et al.</i> , 2017	210	41	90	27
Brahma <i>et al.</i> , 2018	314	189	249	86
Debbarma <i>et al.</i> , 2020	163	49	147	22
Farooq <i>et al.</i> , 2019	921	57	762	16
Khawale <i>et al.</i> , 2019	60	13	7	2
Selim <i>et al.</i> , 2020	140	81	86	43
Risk Factor – Sex	Female		Male	
	Total	Positive	Total	Positive
Tuli <i>et al.</i> , 2015	1053	322	225	52
Agrawal <i>et al.</i> , 2016	104	26	34	2
Naik <i>et al.</i> , 2016	110	28	40	7
Maharana <i>et al.</i> , 2016	42	14	76	4
Debbarma <i>et al.</i> , 2020	254	63	56	8
Brahmbhatt <i>et al.</i> , 2019	48	10	16	3
Farooq <i>et al.</i> , 2019	1492	67	191	6
Khawale <i>et al.</i> , 2019	62	13	5	2
Selim <i>et al.</i> , 2020	215	118	11	6
Risk Factor – Breed	Crossbred / Exotic Breed		Indigenous Breed	
	Total	Positive	Total	Positive
Tuli <i>et al.</i> , 2015	972	315	56	11

Agrawal <i>et al.</i> , 2016	125	27	13	1
Naik <i>et al.</i> , 2016	97	27	53	8
Farooq <i>et al.</i> , 2019	1635	72	48	1
Khawale <i>et al.</i> , 2019	65	15	2	0
Selim <i>et al.</i> , 2020	220	122	6	2

RESULTS AND DISCUSSION

Through the systematic search and following selection criteria amongst 87 studies reviewed from databases 12 were eligible for this meta-analysis (Fig 1). The characteristics of the selected studies and details of each are given in Table 1. Meta-analysis results are also presented in Table 2 wherein the log odds ratio for each of risk factor along with heterogeneity data and publication bias accounts are given. Publication bias was presented through LFK Index and Doi plot and depicted through Figures 5, 6, 7.

Table 2: Results of meta-analysis of risk factors of BTT in India

Risk Factor	No of studies found eligible	Pooled OR (95% CI)	Cochrane's Q	I ² (%)	p value	Tau squared value	LFK index
Age	10	2.062 (1.352 – 3.145)	38.729	76.76	0.00	0.31	0.36
Sex	09	1.632 (1.266 – 2.104)	13.923	42.54	0.08	0.15	0.27
Breed	06	2.113 (1.315 – 3.396)	0.304	0.00	0.99	0.00	0.75

Age as a risk factor

Results of this meta-analysis revealed that age plays an important role in development of Theileriosis in cattle. Adult cattle are found twice at risk for BTT than the younger stocks with an odds ratio of 2.062 (95% CI 1.352 – 3.145) which was highly significant (Q= 38.729; p=0.0) with substantial heterogeneity (I² = 476.76%). As regards age factor in the distribution of BTT these findings correspond to earlier reports by Panda *et al.* (2011); Kumar (2012); Agrawal (2016); Kolte *et al.* (2017); Ghosh *et al.* (2020). However, several other studies contradicted these findings (Salih *et al.*, 2013; Velusamy *et al.*, 2014). Since adult cattle particularly cows are usually subjected to

milk production, the stress due to lactation and pregnancy will be a major predisposing factor for getting BTT and other infectious disease. Additionally, higher exposure of vectors to adult cattle due to open grazing is also an important contributing factor. On the other hand, young stock which are generally raised on intensive feeding and confined housing are less prone to attack by vectors and thereby vector borne diseases (Bhutto *et al.*, 2010). Even earlier reports justifying the age related resistance in young stock to most of the tick borne protozoan and rickettsial diseases including *T.annulata* (Dumanli *et al.*, 2005).

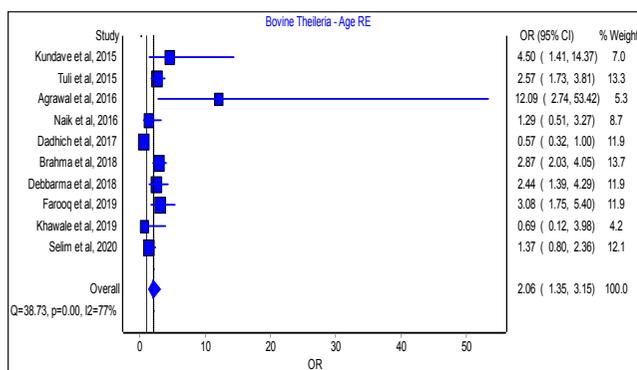


Fig. 2: Forest plot depicting age as a risk factor of BTT in Indian cattle

Sex as a risk factor

Sex of cattle was found to have moderate impact to make the animal susceptible for BTT. It was estimated that cows are little more susceptible to BTT than bullocks / bulls with an odds ratio of 1.632 (95% CI 1.266 – 2.104). However, with lower heterogeneity between studies ($I^2 = 42.54\%$) this estimate was non-significant ($Q = 13.923$; $p=0.08$). These findings also corroborates with earlier reports by Zaman *et al.* (2020). However, the rationale for such distribution may be attributed to certain intrinsic factors such as hormones may have a role to play (Kumar *et al.*, 2017). As males are either culled or utilized for drought purposes as against cows which are raised longer for milk production and breeding are under persistent physiological stress due to pregnancy and/or lactation. This stress along with suppression of immunity may predispose those cows to higher risk of getting infected with BTT (Kamani *et al.*, 2010, Ghosh *et al.*, 2018).

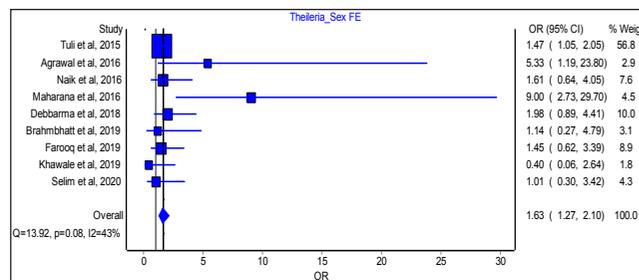


Fig. 3: Forest plot depicting sex as a risk factor of BTT in Indian cattle

Breed as a risk factor

As being reported since long, exotic / crossbred cattle are more prone to contract BTT with an odds ratio of 2.113 (95% CI = 1.315 – 3.396) which was even supported by several earlier evidences based on molecular studies (Kumar *et al.*, 2017). Similarly, the estimates of current meta-analysis also seconded this, however there is absence of heterogeneity ($I^2 = 0\%$) between the studies under this meta-analysis. The results on the basis of Q statistics (0.304; $p= 0.99$) are non-significant. Yet the current meta-analysis revealed that exotic and cross bred cattle are two fold at risk of getting infected with BTT than those cattle of domestic breeds. Similar trends were reported earlier by Salih *et al.*, (2013); Ghosh *et al.*, (2018); Gharbi *et al.*, (2020). The observations of current study have strongly justified the earlier report by Kolte *et al.*, (2017) wherein they have validated that germplasm of the cattle was crucial for the infection of Theileriosis and found that carrier crossbred cattle attracts more maintenance costs to the dairy enterprises. Furthermore, earlier reports also pointed out low case fatality and infection rates in local breeds while the higher morbidity and mortality in exotic and crossbred cattle (Jensen *et al.*, 2007).

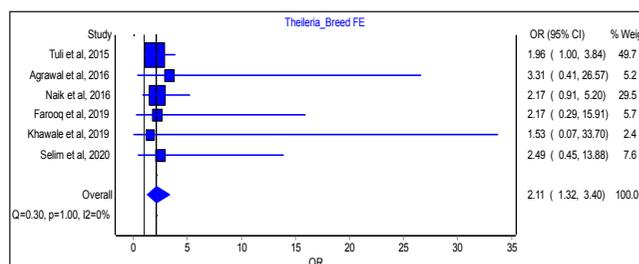


Fig. 4: Forest plot depicting breed as a risk factor of BTT in Indian cattle

Since the 95% confidence intervals of odds ratios for all the factors under this study are more than 1 and did not include 1 (*i.e.* OR \neq 1), the effects size *i.e.* pooled ORs are significant at 5% level of significance.

Publication bias

The meta-analysis conducted for individual risk factor with selected set of studies for each of them has also tested for publication bias and presented through LFK Index and Doi plot individually. The LFK indices for each of them are in the range of 0 – 1 which signifies no publication bias (Table 2). Also visual observation of Doi plots also seconded the absence of publication bias between the studies included for this analysis (Figs. 2, 3, 4). No publication bias in meta-analysis of risk factors of BTT in Indian cattle was found for individual analysis conducted (Barendregt and Doi, 2011 and Kanamori *et al.*, 2018).

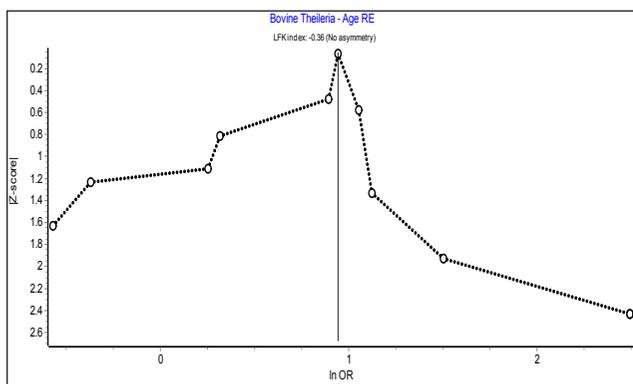


Fig. 5: Doi plot and LFK index for publication bias in meta-analysis of age as risk factor for BTT

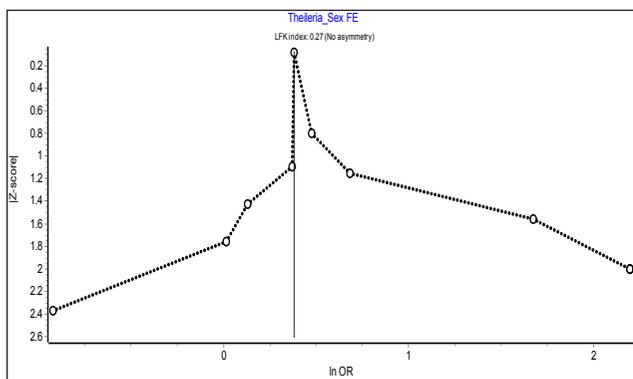


Fig. 6: Doi plot and LFK index for publication bias in meta-analysis of sex as risk factor for BTT

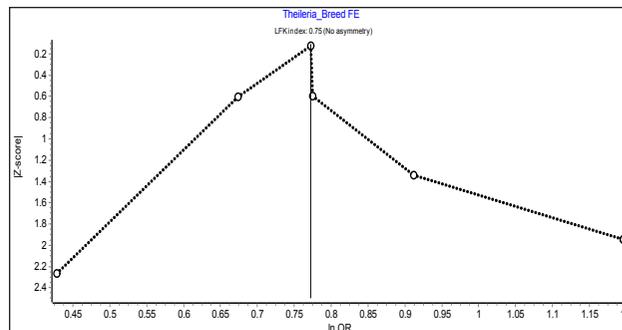


Fig. 7: Doi plot and LFK index for publication bias in meta-analysis of breed as risk factor for BTT

CONCLUSION

It is herewith concluded that among the variables *viz.* age, sex, breed of cattle considered to be potential risk factors for BTT in Indian cattle, age and sex are found significant association with the occurrence of Theileriosis. On the other hand, even if breed of cattle has shown non-significant relation with the BTT incidence, its importance as a risk factor based on evidence from previous studies cannot be overruled. Though meta-analysis gives pooled estimates, the evidence obtained through it depends primarily on the qualities of primary studies included in it. To overcome this, further studies with appropriate sampling from each of the risk groups need to be undertaken to build a concrete evidence for better understanding of epidemiology of BTT.

ACKNOWLEDGMENTS

Authors express gratitude to the authorities of COVAS, Parbhani and MAFSU, Nagpur for providing necessary facilities for this study.

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