

Genomic Detection of *Brucella Abortus* in Milk obtained from Farms in Lorestan Province using PCR Method

Rahim Tahmasebi, Amin Jaydari* , Nemat shams , Heidar Rahimi 

Department of Pathobiology, Faculty of Veterinary Medicine, Lorestan University, Khorram Abad, Iran

ABSTRACT

Background and Aim: Brucellosis is a zoonotic disease which has become endemic in Iran. Contaminated milk with *Brucella* bacteria is the main way of transmission of this disease in humans. Therefore, the aim of this study was to investigate the prevalence of *Brucella spp.* and *B. abortus* in raw milk samples collected from farms belongs to six geographical areas of Lorestan province.

Materials and Methods: In the present study, 100 raw milk samples of cows which were less than 4 years, 4 to 6 years and over 6 years old were randomly collected. The isolates were identified by PCR method using specific bcsp31 and IS711 primers for *Brucella spp.* and *B. abortus*, respectively.

Results & Conclusion: The results showed that 26 of the samples were infected with *Brucella* bacteria, of which 19 samples (73%) were *B. abortus*. Most of the brucellosis infection in cows belonged to cows less than 4 years (31.4%) and 4 to 6 years (31.4%) categories, 11 samples in both groups, so that in cows older than 6 years this rate was 13% (4 samples). The east of the province with 12 samples and the northwest of the province with one sample showed the highest and lowest levels of *Brucella* infection, respectively. The findings demonstrated that there is a significant difference between the frequency of *Brucella* contamination in milk in the eastern regions of Lorestan province with other regions ($p < 0.05$). It should be noted that the possibility of transmitting brucellosis to humans through consumption of contaminated raw milk and dairy products in Lorestan province is high.

Keywords: *B. abortus*, zoonosis, milk, cattle, PCR

Received: 2022/04/14;

Accepted: 2022/06/26;

Published Online: 2022/08/08

Corresponding Information:

Amin Jaydari, Department of Pathobiology, Faculty of Veterinary Medicine, Lorestan University, Khorram Abad, Iran Email: jaydari.a@lu.ac.ir



Copyright © 2022, This is an original open-access article distributed under the terms of the Creative Commons Attribution-noncommercial 4.0 International License which permits copy and redistribution of the material just in noncommercial usages with proper citation.

Use your device to scan and read the article online



Tahmasebi R, Jaydari A, shams N, Rahimi H. Genomic Detection of *Brucella Abortus* in Milk obtained from Farms in Lorestan Province using PCR Method. Iran J Med Microbiol. 2022; 16 (5):479-84.

Download citation: [BibTeX](#) | [RIS](#) | [EndNote](#) | [Medlars](#) | [ProCite](#) | [Reference Manager](#) | [RefWorks](#)

Send citation to:  [Mendeley](#)  [Zotero](#)  [RefWorks](#)

1. Introduction

Brucellosis is a zoonotic disease with a bacterial agent that is important from both health and economic aspects. Transmission of brucellosis to humans is through consumption of raw milk and contaminated dairy products or direct contact with fluids or tissues of infected animals (1). Despite control and prevention programs such as vaccination and slaughter of infected animals, Iran is one of the pandemic with high prevalence of this disease (2). Lorestan province, with about 6.5 million livestock units, accommodates 5.5% of the country's livestock population and has played an important role in the

production of livestock products, which is now one of the centers with a high prevalence of Brucellosis in the country (3). The eradication of this disease depends not only on the prevention of new cases of the disease but also on the timely diagnosis of the disease in humans and animals.

Serological and bacterial culture methods, which are the most common tests for diagnosing the *Brucella* infection, have major drawbacks. Disadvantages of the bacterial culture method include time consuming, facilitates the risk of acquiring laboratory infection

and the need for biosafety Level 3 (BSL-3) laboratories. In addition, serological methods are less specific and may provide false results. Also, the clinical similarity of the symptoms of brucellosis with many other infectious and non-infectious diseases, has led to the use of molecular methods with high sensitivity and accuracy to diagnose the strain of *Brucella* (2, 4). Therefore, due to the importance of cow's milk as the main source of dairy production and the important role of milk in the possible transmission of brucellosis to humans, the present study were addressed to evaluate the prevalence of *Brucella spp.* and *B. abortus* in cow milk obtained from various regions of Lorestan province using the PCR method.

2. Materials and Methods

Sample collection

In this study, a total of 100 raw milk samples obtained from industrial and traditional farms in 6 regions of north (Noor Abad), south (Pole Dokhtar), west (Aligoudarz, Azna and Doroud), east (Kouhdasht), northwest (Boroujerd and Alashtar) and center (Khorram Abad) of Lorestan province (Figure 1) were randomly collected from February to June 2020. 50 ml of milk was collected from each cow after disinfecting the nipples with 70% alcohol in sterile. Then the age of the animals was recorded and finally, the samples under aseptic conditions were transferred to the laboratory of veterinary laboratory, Lorestan University on the icebox.

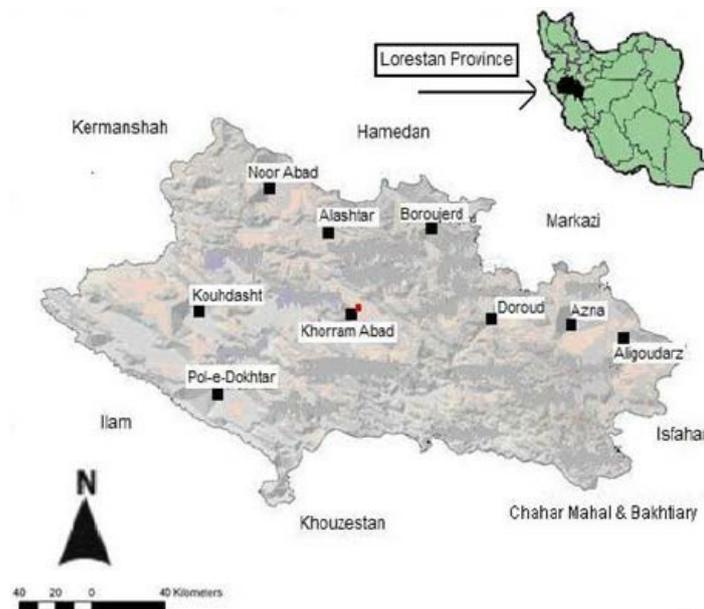


Figure 1. Geographical map of Lorestan province.

DNA extraction

Genomic DNA Extraction was done using 10 ml of each milk sample. Samples were centrifuged at 6000 rpm for 15 minutes. Then, 200 μ l of the fatty top layer was transferred to the 1.5 ml tube of each sample and using a DNA extraction kit (Gene All South Korea), the DNA extraction process was performed according to the manufacturer's protocol. The quality and concentration of the extracted DNA were evaluated by spectrophotometry with a wavelength of 260 to 280 nm, and the extracted DNA was stored in a -20 freezer for further study.

Identification of *Brucella spp.* and *B. abortus* by PCR:

To identify *Brucella spp.* and *B. abortus*, specific primers bcsp31 (31 kDa outer membrane protein) and IS711 prepared by Takapo Zist Tehran (Iran) were

used, respectively (Table 1). PCR amplification was performed using PCR master kit (Ampliqon Taq DNA Polymerase Master Mix RED 1.25 mL, Ampliqon Denmark) with 25 μ L mixtures containing 12.5 μ L of 2X master mix, 0.5 μ L of each primer, and 5 μ L of the extracted DNA. For positive control represented by genomic DNA isolated from vaccine strain RB51 (Razi Vaccine and Serum Institute - Iran) and for the negative control, sterile water was added instead of nucleic acids. Further, the amplification was conducted by Bio-Rad thermocycler (Model T- 100, USA) under the following conditions: A for *Brucella spp.* and B for *B. abortus*.

Table 1. PCR primers used for *Brucella spp.* and *B. abortus* detection.

Target gene	Primer sequence	Amplified Product size	References
<i>bcs p31</i>	(F) 5'-TGG CTC GGT TGC CAATAT CAA-3'	223 (bp)	(5)
	(R) 5'- CGC GCT TGC CTT TCA GGT CTG-3'		
<i>IS711</i>	(F) 5'-GACGAACGGAATTTTCCAATCCC-3'	498 (bp)	(6)
	(R) 5'-TGCCGATCACTTAAGGGCCTTCAT-3'		

A: The initial step of 95°C for 5 min, followed by 40 cycles of 90°C for 1 min, 60°C for 30 Second, 72°C for 2 min, and finally, 72°C for 7 min.

B: The initial step of 95°C for 5 min, followed by 35 cycles of 95°C for 75 Second, 55.5°C for 1 min, 72°C for 2 min, and finally, 72°C for 10 min.

The PCR products were separated in a 1.5% (w/v) agarose gel (Merck, Germany) containing 2.5 µg/mL gel stain (Sina Gene, Iran). Electrophoresis was performed in 0.5x Tris/Borate/EDTA (TBE) buffer for one hour at 100 V. The resulting PCR products were visualized under a UV transilluminator (E-Box, Iran) and the 100 bp DNA ladder (Smobio, Taiwan) plus was used as the molecular size marker.

Statistical analysis

Data analysis was performed using Chi-square Tests. For this purpose, SPSS software version 22 (IBM, USA) was used to perform this statistical test and the difference was considered significant with $p < 0.05$.

3. Results & Discussion

100 milk samples were investigated using PCR method. Based on the results 26 (26%) milk samples were positive for *Brucella spp.* and 19(73%) cases of them were positive for *B. abortus*, respectively ([Figure 2](#)).

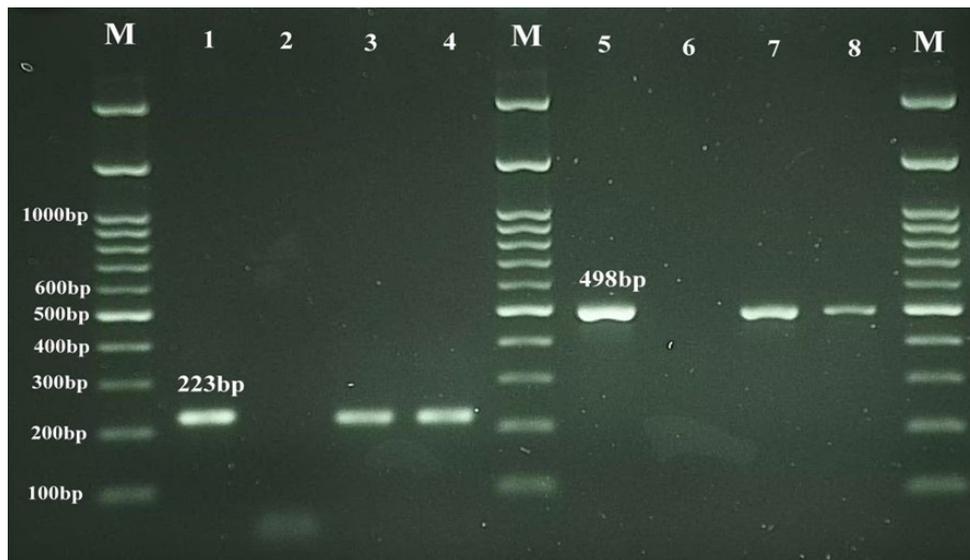


Figure 2. PCR assay for the detection of *Brucella .spp* and *B. abortus* in raw milk samples. Lane M: Standard DNA marker (100bp DNA ladder); Lane 1: Positive control for *Brucella .spp* (RB51 vaccine strain); Lane 2: Negative control; Lanes 3, 4: Positive samples for *Brucella .spp*; Lane 5: Positive control for *B. abortus* (RB51 vaccine strain); Lane 6: Negative control; Lanes 7, 8: Positive samples for *B. abortus*.

The highest level of contamination with *Brucella* was observed in the eastern region of Lorestan province. Out of 24 milk samples, 12 samples (50%) were positive. While in the central, northern, northwestern, southern and western regions of the province, this rate was 2 samples were determined (14.2%), 3 samples (20%), 1 sample (7, 6), 2 samples (14.2 %) and 6 samples (30%), respectively. In addition, the prevalence of *B. abortus* in the east of

the province with 9 samples (75%) was identified more than other areas. The results of this study show that there is a significant difference between the frequency of milk contamination in the eastern regions of Lorestan province with other regions ($p < 0.05$). Therefore, it can be concluded that the geographical area significantly effects on the rate of *Brucella* infection in Lorestan province ($p < 0.05$) ([Table 2](#)).

Table 2. Prevalence of *Brucella spp.* and *B. abortus* detection in different geographical areas.

Area	<i>Brucella spp.</i>	<i>B. abortus</i>	Negative	Total
East	12	9	12	24
Center	2	2	12	14
North	3	2	12	15
North West	1	1	12	13
South	2	1	12	14
West	6	4	14	20
Total	26	19	74	100

In this study, three age groups (less than 4 years, 4 to 6 years and over 6 years) were selected. From 35 milk samples which were belonged to less than 4 years old group, 11 samples (31.4%) and of 35 milk samples which were taken from the 4 to 6 years group 11 samples (31.4%) were positive for *Brucella* infection. Also, out of 30 samples from the over 6 years old group, 4 samples (13.3%) were infected with *Brucella*. The results of this study showed that there was no significant relationship between the presence of *Brucella* and the age of cattle.

In cattle populations in industrial units, testing and slaughter programs are implemented and comprehensive vaccination of cattle and calves with RB51 and S19 vaccine is also done, so the basis of prevention of brucellosis is its control in animal populations. Despite extensive efforts to combat the disease, prevalence of brucellosis is not only high among the livestock population, but also the occurrence of the disease in the human population is considered as an infectious disease with a higher prevalence than other infectious diseases in most provinces of the country (3,7-11). Awareness of the number of positive cases and the incidence of disease in humans and livestock in the provinces, epidemiological characteristics of infected areas in the two sectors of health and veterinary through the exchange of information in recent years has been one of the main activities in combating the disease which requires more extensive studies in this field.

In a study conducted in 2012 in Kurdistan province by Shafiei et al, using PCR method on 60 samples of raw cow's milk, they found that 20 samples (33.33%) were infected with *Brucella* bacteria, of which nine samples (45%) were detected to be *B. abortus*. The author attributes the high prevalence of brucellosis to the proximity of Kurdistan province to countries such as Iraq and Turkey and the entry of non-native strains into Kurdistan province and the possible spread of *B. abortus* vaccine in milk after vaccination in the studied areas (12). In the study by Khalili et al, which was performed to evaluate the contamination of raw milk with *Brucella* by PCR method in Kerman, 8.3% of milk samples were positive for the genome of *Brucella spp.* (13). In another similar study in 2017, in Lorestan

province by Shams et al. The prevalence of *Brucella* in 120 samples of milk tanks was reported to be 10% (14). In the study by Shakrian et al. (2012), in Isfahan and Chaharmahal Bakhtiari provinces, which aimed to investigate the contamination of raw cow milk samples with *B. abortus* by PCR method, only 1% of the samples were infected with *B. abortus*. These results are not in accordance with the results of the present study, which may be due to differences in sampling method, number of samples, geographical factors of the study areas, vaccination and measures to control this disease among the livestock of these two provinces. The study of geographical distribution of this disease shows that Chaharmahal and Bakhtiari province are classified in the group with moderate infection and Isfahan province in the group with low pollution, while Lorestan province is in the group with high prevalence (15). In the study by Entezari and Sepahvand, (2014) it has been shown that climatic conditions and geographical environment can be one of the factors affecting the prevalence of the brucellosis disease in susceptible areas (16). In confirmation of this report, the present study also showed that there is a significant difference between the frequency of bovine milk contamination in the eastern regions of Lorestan province with other regions ($p < 0.05$). Also, this study showed the highest prevalence of brucellosis after the eastern region of the province (Aligudarz, Azna and Dorud cities) in the western part of the province (Hennpoldakhtar and Kuhdasht cities). Due to the summer and winter location of the regions of the province, as well as the traffic route of nomadic herds, having suitable livestock pastures, has caused the highest prevalence of brucellosis in these areas. In a comparison of Khalili study in Kerman province and Shams in Lorestan province on samples with the same conditions obtained from milk reservoirs, it was found that the prevalence of brucellosis in Lorestan province is higher than Kerman province. Therefore, it can be concluded that the prevalence of brucellosis is lower in Kerman, Isfahan, Chaharmahal and etc. provinces, than in Lorestan province (13-15) and in parallel, the higher prevalence of this disease in Kurdistan province (12) can be due to the effects of geographical area on the rate of *Brucella* infection and excretion through milk.

In addition, this argument is confirmed by comparing the prevalence of *Brucella* in the present study with studies conducted abroad. As, The prevalence of *Brucella spp.* and *B. abortus* in raw milk in developing countries such as Sudan is reported to be 22.4% and 40% (17), Kenya 18.9% and 65.5%, respectively (18). Also in Iraq, the prevalence of *Brucella* has been reported from 8.4% to 56% by serological examination (19). While in most European countries brucellosis has been eradicated or has a very low prevalence (20, 21). So far, no study has examined the relationship between the age of the animal and the prevalence of brucellosis. The present study showed that there is no significant relationship between the prevalence of brucellosis and the age of the animal. However, *Brucella* is more prevalent in cows under 6 years of age than in cattle over 6 years of age. This seems to be due to the vaccine received over many years and the safety of older animals.

4. Conclusion

The findings of the present study show that the prevalence of brucellosis is still high in Lorestan province and more comprehensive planning, and

policies should be done to prevent and eradicate this disease, especially in the eastern and western parts of the province. In addition, due to the high livestock population in these areas and the main role of dairy products in the diet of the people, especially the cheese preparation in rural areas from raw milk, necessary training should be provided to detect the presence of *Brucella* bacteria in cow's milk and to investigate the possibility of contracting Brucellosis in case of consumption of raw milk and contaminated dairy products by health and veterinary organs.

Acknowledgment

The authors would like to thank Mr. Ali Karimpor and Veterinary personnel of Noor Abad and Kuhdasht counties for their aid in sample collection.

Funding

This research is financially supported by Lorestan University

Conflict of Interest

The authors declare that they have no conflict of interest.

Reference

1. Taleski V, Zerva L, Kantardjiev T, Cvetnic Z, Erski-Biljic M, Nikolovski B, et al. An overview of the epidemiology and epizootology of brucellosis in selected countries of Central and Southeast Europe. *Vet Microbiol.* 2002;90(1-4):147-55. [DOI:10.1016/S0378-1135(02)00250-X]
2. Dadar M, Alamian S, Behroozikhah AM, Yazdani F, Kalantari A, Etemadi A, et al. Molecular identification of *Brucella* species and biovars associated with animal and human infection in Iran. *Vet Res forum an Int QJ.* 2019;10(4):315-21.
3. Golshani M, Buozari S. A review of Brucellosis in Iran: Epidemiology, Risk Factors, Diagnosis, Control, and Prevention. *Iran Biomed J.* 2017;21(6):349-59.
4. Rahimi H, Tukmechi A, Rashidian E. Use of touch-down polymerase chain reaction to enhance the sensitivity of *Brucella melitensis* detection in raw milk. *Anim Biotechnol.* 2022;33(1):104-9. [DOI:10.1080/10495398.2020.1777149] [PMID]
5. Baily GG, Krahn JB, Drasar BS, Stoker NG. Detection of *Brucella melitensis* and *Brucella abortus* by DNA amplification. *J Trop Med Hyg.* 1992;95(4): 271-5.
6. Bricker BJ, Halling SM. Differentiation of *Brucella abortus* bv. 1, 2, and 4, *Brucella melitensis*, *Brucella ovis*, and *Brucella suis* bv. 1 by PCR. *J Clin Microbiol.* 1994;32(11):2660-6. [PMID] [PMCID] [DOI:10.1128/jcm.32.11.2660-2666.1994]
7. Kashfi SM, Hatamian N, Rakhshani T. Epidemiological Study of the Brucellosis in Iran, Andimeshk, 2001-2016. *J Heal Sci Surveill Syst.* 2018;6(1):23-8.
8. Behroozikhah AM, Bagheri Nejad R, Amiri K, Bahonar AR. Identification at Biovar Level of *Brucella* Isolates Causing Abortion in Small Ruminants of Iran. Wong H-C, editor. *J Pathog.* 2012;2012:357235. [DOI:10.1155/2012/357235] [PMID] [PMCID]
9. Pishva E, Salehi R, Hoseini A, Kargar A, Taba FE, Hajiyan M, et al. Molecular typing of *Brucella* species isolates from Human and livestock bloods in Isfahan province. *Adv Biomed Res.* 2015;4:104.
10. Ashrafganjooyi SH, Saedadeli N, Alamian S, Khalili M, Shirazi Z. Isolation and biotyping of *Brucella spp.* from sheep and goats raw milk in southeastern Iran. *Trop Biomed.* 2017;34(3):507-11.
11. Azadi Chegeni S, Jaydari A. Seroprevalence of Malta Fever in Veterinary Staff of Lorestan Province Using Indirect ELISA Method in 2018-2019. *Yafteh.* 2020 ;22(2):36-42.
12. Abdali F, Hosseinzadeh S, Berizi E, Pourmontaseri

- M. Prevalence of *Brucella* species in unpasteurized dairy products consumed in Shiraz province using PCR assay. *Mol Biol Res Commun.* 2020;9(3):117-21.
13. Khalili M, Aflatoonian MR, Aliabadi FS, Abshenas J. *Brucella* contamination in raw milk by polymerase chain reaction. *Tehran Univ Med J.* 2016;74(7):517-21.
 14. Shams N, Jaidari A, Etemadfar L. Molecular Detection of *Brucella abortus* and *Brucella melitensis* in Raw and Unpasteurized Bulk Cow Milk Tanks of Traditional Domestic Dairy Sale Centres in Khorramabad. *Iran J Med Microbiol.* 2017;11(4):13-20.
 15. Shakerian A. Study of contamination rate in raw milk and its traditional products with *Brucella abortus*, and *Brucella mellitensis* in Isfahan and Chaharmahal and Bakhtiari Provinces. *J Shahrekord Univ Med Sci.* 2015;17(1):16-23.
 16. Entezari M, Sepahvand S. Investigating Geographical Factors Affecting the Prevalence of Brucellosis in the Lorestan Province, Iran. *J Isfahan Med Sch.* 2014; 32(283): 569-79.
 17. Abdalla A, Hamid ME. Comparison of conventional and non-conventional techniques for the diagnosis of bovine brucellosis in Sudan. *Trop Anim Health Prod.* 2012;44(6): 1151-5. [[DOI:10.1007/s11250-011-0051-7](https://doi.org/10.1007/s11250-011-0051-7)] [[PMID](#)]
 18. Muendo EN, Mbatha PM, Macharia J, Abdoel TH, Janszen P V, Pastoor R, et al. Infection of cattle in Kenya with *Brucella abortus* biovar 3 and *Brucella melitensis* biovar 1 genotypes. *Trop Anim Health Prod.* 2012;44(1):17-20. [[DOI:10.1007/s11250-011-9899-9](https://doi.org/10.1007/s11250-011-9899-9)] [[PMID](#)]
 19. Dahl MO. Brucellosis in food-producing animals in Mosul, Iraq: A systematic review and meta-analysis. *PLoS One.* 2020;15(7):e0235862-e0235862. [[PMCID](#)] [[DOI:10.1371/journal.pone.0235862](https://doi.org/10.1371/journal.pone.0235862)] [[PMID](#)]
 20. Jansen W, Linard C, Noll M, Nöckler K, Al Dahouk S. *Brucella*-positive raw milk cheese sold on the inner European market: A public health threat due to illegal import? *Food Control.* 2019; 100:130-7. [[DOI:10.1016/j.foodcont.2019.01.022](https://doi.org/10.1016/j.foodcont.2019.01.022)]
 21. Authority EFS, Control EC for DP and. The European Union: summary report on trends and sources of zoonoses, zoonotic agents and food-borne outbreaks in 2016. EFSA. 2017 ;15(12):e05077. [[DOI:10.2903/j.efsa.2017.5077](https://doi.org/10.2903/j.efsa.2017.5077)] [[PMID](#)]