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Research Article

Detection of Brucella spp. in Dairy Products by Real-Time PCR

Elham Moslemi,¹ Mohammad Mehdi Soltandalal,^{2,3} Mohammad Reza Beheshtizadeh,¹ Afsoon Taghavi,⁴ Hamidreza Kheiri Manjili,⁵ Reza Mahmoudi Lamouki,⁶ and Amir Izadi^{7,8,*}

²Food Microbiology Research Center, Tehran University of Medical Sciences, Tehran, Iran

³Division of Microbiology, Department of Pathobiology, School of Public Health, Tehran University of Medical Sciences, Tehran, Iran

⁴Research Cancer Center, Shahid Beheshti University of Medical Sciences, Tehran, Iran

⁵Assistant Professor, Department of Pharmaceutical Nanotechnology, School of Pharmacy, Zanjan University of Medical Sciences, Zanjan, Iran

⁶Department of Medical Genetic, Faculty of Medicine, Shahid Beheshti University of Medical Sciences, Tehran, Iran

⁷Banej Exir Research Center, Tehran, Iran

⁸Young Researcher Club, East Tehran Branch, Islamic Azad University, Tehran, Iran

^{*}Corresponding author: Amir Izadi, Banej Exir Research Center, Tehran, Iran. Tel: +98-9126057392, E-mail: amir_izad@yahoo.com

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Abstract

Background and Objectives: *Brucella* is an intracellular gram-negative bacterium that can infect many kinds of mammals like humans, sheep, cattle, etc. Brucellosis is a contagious occupational disease caused by *Brucella* spp. that affects individuals who have close contact with infected animals. The clinical features of Brucellosis are not disease-specific and almost every organ can be affected. This zoonotic disease is a great health concern and economically important in many countries, such as Iran. The aim of this study was to detect *Brucella* spp. in pasteurized and non-pasteurized dairy products.

Methods: In this study, 208 samples, including goat, sheep, and cow raw and pasteurized milk as well as pasteurized and non-pasteurized cheese, were collected in Tehran province. The DNA was extracted, and then the real-time PCR was used for detection of the *Brucella* spp. gene.

Results: The prevalence of *Brucella* spp. contamination in the dairy products was: 45.5% in goat's raw milk, 39.1% in non-pasteurized cheese, 27.3% in sheep's raw milk, 26.3% in cow's raw milk, 25% in pasteurized cheese, and 14.7% in pasteurized milk.

Conclusions: Rapid and exact detection of pathogens in dairy products is the most significant factor to prevent foodborne diseases. In addition, the real-time PCR assay is sensitive and specific enough to detect a low number of *Brucella* spp. in dairy products.

Keywords: Real-Time PCR, Brucellosis, Pasteurization, Dairy Products, Brucella spp.

1. Background

Non-pasteurized dairy products are one of the most significant hosts of many foodborne pathogens including *Enterobacteriaceae* family, *Brucella* spp., and *Campylobacter* spp. (1). The genus of *Brucella* comprises facultative Gram-negative bacteria that are able to infect a vast variety of mammalians such as humans and chattel. Brucellosis is a contagious occupational disease caused by *Brucella* spp. that affects individuals who have close contact with infected animals. The most common way of Brucellosis transmission is ingestion of non-pasteurized milk and dairy products obtained from infected animals (2, 3).

Brucella spp. are accumulated in mammary glands and supra-mammary lymph nodes of infected animals and therefore the milk of these animals will be a source of pathogens (4). The clinical features of Brucellosis are not disease-specific; but almost every organ can be affected (5, 6). Infection with *Brucella* can cause a variety of problems in animals; for example, in females, it could be the cause of abortion and in males, usually orchitis and epididymitis are seen. Depending on conditions such as proper temperature, pH, and humidity, *Brucella* can remain in a contaminated environment for several months (7). Several epidemiological studies have reported a high frequency of brucellosis in endemic countries such as Saudi Arabia (19%), Iran (20%), Peru (8%), and Azerbaijan (10%) (8).

Four out of six major identified species of *Brucella* are human pathogens. Human brucellosis is mostly caused by *B. melitensis* while *B. abortus* is the second cause of human brucellosis, mostly infecting cattle, buffalos, elks, yaks, and camels; *B. canis* is the other cause of human brucellosis and *B. suis* that infects domestic pigs and rodents is the last one (9). Because *Brucella* can easily be transmitted as aerosols, it was used in the former U.S. biological weapons program (10).

¹Department of Biology, East Tehran Branch, Islamic Azad University, Tehran, Iran

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Currently, diagnostic methods for detection of Brucella spp. rely on serological, microbiological, and molecular techniques. Serological techniques are standard for the epidemiological surveillance of brucellosis (11). The most common method for Brucella detection in milk and milk products is MRT (milk ring test) that has low sensitivity and accuracy (12). Molecular detection methods have been widely used for Brucella diagnosis in the last decades (13). Real-time PCR, which has less hazard and high sensitivity, has been developed for Brucella detection (14). Real-time PCR does not require extensive manipulation that minimizes the risk of contamination (15). Several nucleic acid sequences for Brucella spp. have been used to be amplified by PCR technique like16S rRNA, 16S-23S intragenic spacer region, omp2, and bcsp31 (16). A real-time Light-Cycler PCR (LC-PCR) assay that is based on the use of SYBR Green I DNAbinding fluorophore dye was developed by a clinical laboratory to simplify the molecular diagnosis of brucellosis (17).

Despite the decreased incidence of Brucellosis, in Iran and many endemic countries, it remains as an important public health.

Furthermore, a survey displayed that nearly 7.4% of cows in Iran were infected with *Brucella* spp. (18). Approximately, 500000 cases of human brucellosis globally are reported to the world health organization annually (19). Therefore, it seems that the contaminated dairy products are one of the most common causes of brucellosis. Therefore, the aim of this study was to detect *Brucella* spp. in different dairy products.

2. Methods

2.1. Preparation of Samples

In this case study, 208 different samples including 57 samples of cow raw milk, 34 samples of pasteurized milk (from different companies), 28 samples of pasteurized cheese (from different companies), 23 samples of unpasteurized cheese, 33 samples of goat raw milk, and 33 samples of sheep raw milk were collected in the province of Tehran. The samples were collected from 2014 to 2016 and stored at -20°C. For DNA extraction, 100 μ L of each sample were obtained and DNA was extracted from samples with QIAamp DNA Mini Kit (Qiagen, USA), according to the protocols).

2.2. Real-Time PCR Technique

The real-time PCR assays were optimized and applied to all samples by the 7500 real-time PCR system (Applied Biosystems).

2.3. Real-Time PCR Reactions

The Taqman/ROX qPCR Master Mix (2X) (Applied Biosystems) was used. Each reaction mixture contained 100 ng of template DNA, 12.5 μ L Master Mix (Applied Biosystems), 1 μ L of each F/R (Forward/ Reverse) primers and probe (10 mM), and 9.5 μ L nuclease-free water in a final volume of 25 μ L per reaction. All reactions had a positive control that contained *Brucella* spp. DNA with exact concentration and a negative control that contained dilute water instead of DNA.

The mixture was subjected to the following PCR conditions: primary denaturation temperature at 95°C for 10 minutes to activate AmpliTaq Gold polymerase, followed by 40 cycles of denaturation at 95°C for 15 seconds, and annealing at 59°C for 1 minute. To minimize Ct values for experimental variability, the threshold cycle, in which the fluorescence signal raised significantly above background in the exponential phase of the amplification, was specified by the second derivative maximum method.

2.4. Template Preparation of the Standard Curve

The PCR amplification of the Brucella gene fragment was performed by using a universal primer. The reaction mixture contained Brucella DNA (1 - 20 ng), 1X PCR Buffer, 1.5 mM MgCl₂, 0.2 mM dNTP, 0.4 μ M each F and R primers, and 2U Tag DNA polymerase (5 U/ μ L; Fermentas, USA) in a final volume of 25 μ L. Amplification included a primary denaturation at 95°C for 3 minutes, followed by 25 cycles of denaturation at 94°C for 30 seconds, annealing at 54°C for 30 seconds, and extension at 72°C for 1 minutes. The PCR Amplicon (766 bp) was cloned into a pTZ57R vector and transformed in E. coli JM107 cells with T/A cloning Kit, (Fermentas, cat #: k1213) following the manufacturer's instructions. DNA sequencing confirmed the cloning. The absorbance of the DNA solution was measured at 260 nm. The standard curve was drawn by the Fit Point's analysis method that included the 7500 real-time PCR system (Applied Biosystems). The concentration log of a dilution series of the standard or reference template DNA (Brucella spp.) was plotted versus the cycle number in which the fluorescent signal increased up to the background or threshold (Ct value). The slope of the standard curve, which was provided for each detected approach, was put into the following equation to determine the reaction's efficiency: efficiency = $10^{-(1/sl^{\circ}pe)}$ (20).

2.5. DNA Sequence Analysis and Design of the Primers and Probe

The target gene for the designed probe and primers set was *Brucella spp*. gene (Accession No. HE603359). The Primer Express Software provided by Applied Biosystems was used for designing primers. All primers were supplied by Bioneer (Korea). The sequences of each primer are listed in Table 1.

2.6. Sensitivity and Specificity Determination of Real-Time PCR Assays

For sensitivity determination of real-time PCR, different dilutions of bacterial DNA from 1 million up to 10 particles were provided. The DNA of mice, humans, *Salmonella*, *Shigella*, *Saccharomycescerevisiae*, and *Escherichiacoli* were used to verify the PCR specificity.

3. Results

3.1. Testing of Samples for Brucella by Real-Time PCR

The results of real-time PCR show the following prevalence of contamination for *Brucella* spp.: 45.5% in goat raw milk, 39.1% in non-pasteurized cheese, 27.3% in sheep raw milk, 26.3% in cow raw milk, 25% in pasteurized cheese, and 14.7% in pasteurized milk. The highest and lowest prevalence rates of *Brucella* spp. were observed in goat raw milk and pasteurized milk, respectively. Most of the contaminated samples were collected from companies without any standard certification. In the milk and cheese produced in rural areas with no observation of sanitation organizations, a high number of *Brucella* spp. was detected.

The results of all samples are described in Table 2.

3.2. Sensitivity and Specificity of Real-Time PCR

The real-time PCR after optimization had high specificity by showing no reactions to infectious agents except *Brucella* spp. The real-time sensitivity was 10 particles.

The threshold cycle values are plotted against the *Brucella* spp. input copy number.

4. Discussion

It seems that people who are living far from *Brucella* endemic regions are at low risk of infection; however, recently some reports indicated that brucellosis is increasing in non-endemic areas as in endemic regions. In addition, it seems that brucellosis could be easily transferred from rural to urban regions.

It is usually done by transferring raw milk and dairy products that are infected by *Brucella* spp. from far and near distances (2).

B. abortus can spread in food and water. Under conditions of high humidity, low temperatures, and no sunlight, these organisms can stay viable for a long time in the water, aborted fetuses, wool, feces, hay, clothes, and equipment. *Brucella* species can withstand drying, particularly when organic substances are present in the soil. In low temperature, survival is longer, especially when it is below freezing (21). Brucellosis affects many organs and tissues as a systemic infection (6). The highest significant incidence of brucellosis bacteremia occurs in spring and summer while the lowest occurrence is in winter. This is because of the consumption of unpasteurized milk products as a result of more travel to rural areas at these times (22). As mentioned by WHO, nowadays, brucellosis is the most common infection in the world with 500,000 infected cases each year (6, 23).

Endemic countries suffer from the lack of productivity and its adverse effects on human health (24). Several different factors like socioeconomic factors and some cultural habits cause various prevalence rates of brucellosis in distinct areas all over the world (25).

The prevalence of human brucellosis in different parts of Iran varied from 1.5 to 107.5 per 100,000 people in 2003. The highest levels of infection appeared in Hamadan with 107.5, Kurdistan with 83.5, Western Azarbaijan with 71.4, and Zanjan with 67.1 per 100,000 people (26).

Clearly, the prevalence of Brucella contamination varies according to the sensitivity of the used methods. Although isolation and phenotyping of Brucella are timeconsuming and unsafe and need well-trained staff, they are still the gold standard for diagnosis of Brucella spp. (27). Today by using molecular detection techniques like PCR, the detection of brucellosis is significantly increasing (28, 29). Indeed, several studies have shown that agents such as lipids, enzymes, polysaccharides, proteins, and Ca2+ in high concentration that are present in dairy products can play the role of PCR inhibitor by interfering with nucleic acid degradation or with the amplification activity of polymerase (30). Unlike the circulating bacteria and DNA, antibodies against Brucella antigen remain in the blood for a long time, making sometimes PCR results negative while ELISA results are positive.

In the study performed by Lindahl-Rajala et al. in 2017, it was stated that 10.3% of non-pasteurized cow milk samples were infected with *Brucella* spp. in Tajikistan. They also declared that since the consumption of non-pasteurized milk is common in this area, this problem has caused anxiety and prevalence of the diseases (30). Similar to this study, they also used real-time PCR technique and stated that traditional detection methods are not sufficient for detection of this bacterium.

Probert et al. in 2004 established the multiplex PCR test for the detection of *Brucella* spp., *B. abortus* and *B. melitensis*, in a single test (31). In Iran, some limited studies have been performed in the field of detecting *Brucella* spp. by using real-time PCR on dairy products. In the study by Majid Yaran et al. in 2016, the prevalence of *B. melitensis*

able 1. Primers Sequences Used for Detection of Brucella spp.					
Methods	Forward Primer (5' - 3')	Reverse Primer (5' - 3')	Fragment Size, bp		
Real Time PCR	CATATCGTTGCGCGTAAGGA	GAAACGCGCTTGCCTTTC	64		
Probe	FAM CAAACATCAAATCGGTCGCGGACC MGB				
Universal primer	TGCCCGGTCTCGTAGCGACG	TCTGCGCCGGGATGCAGC	766		



Cable 2. PCR Results of Brucella spp. Detection in Dairy Products					
Sample	No.	Real-time PCR			
		Positive	%		
Raw cow milk	57	15	26.3		
Pasteurized milk	34	5	14.7		
Pasteurized cheese	28	7	25		
Traditional cheese	23	9	39.1		
Raw goat milk	33	15	45.5		
Raw sheep milk	33	9	27.3		



and *B. abortus* in raw milk and dairy products were evaluated by using real time PCR (3). In spite of the differences in the obtained results, in both studies, the necessity of using exact and sensitive molecular techniques for detection of *Brucella* spp. in dairy products was emphasized.

The assessment of real-time PCR technique for detection of *B. melitensis* in non-pasteurized milk was done by Wareth et al. in 2014. They notified that non-pasteurized dairy products are important sources for the prevalence of brucellosis and real-time PCR is qualified and efficient for detection of this pathogen (2).

As mentioned earlier, one of the most important ways for brucellosis infection transmission is consumption of infected dairy products. The results of this study showed that high percentages of non- pasteurized dairy products including milk and traditional cheese are infected with *Brucella* spp.

In spite of this fact, the consumption of nonpasteurized dairy products in many places still makes a great concern for the disease prevalence. In addition, since the results showed 14.7% of pasteurized milk and 25% of pasteurized cheese samples were infected with *Brucella* spp., it seems the pasteurization methods are not effective for destruction of this pathogen. As the dairy products are controlled before being distributed, it seems that commercial quality control of these products is not sufficient and exact. Therefore, reducing the possibility of being infected by this pathogen by using accurate molecular detection techniques like real-time PCR should be considered (3, 32).

In order to decrease the venture of *Brucella* infection due to the ingestion of contaminated dairy products, food

safety management systems, which guarantee the sanitary quality of the products, have to control and improve the production of dairy products. Intransitive training of dairy makers and consumers should be provided and the consumers should be notified of serious health risks due to unpasteurized milk and dairy products (33).

Finally, for exact detection of bacteria and evaluation of the amount of pathogen, it is essential to use sensitive and specific methods such as real-time PCR to detect *Brucella* spp. in dairy products.

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