Presence and Antibiotic Resistance of *Listeria monocytogenes* in Raw Milk and Dairy Products

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Abstract

In this study, the presence of *Listeria monocytogenes* in raw milk and milk products produced from raw milk by traditional method, such as cheese and butter using cultural and molecular techniques and the antibiotic resistances were investigated. Isolation was performed using the FDA technique. Microbact 12L test kit was used for biochemical identification. Genotyping of isolates was done by traditional PCR method. Antibiotic resistance of isolates was determined by standard disk diffusion method. *Listeria* was detected in 16 (5.3%) of the total 300 food samples according to the biochemical identification results. The highest prevalence of *Listeria* sp. was in raw milk samples (10%), followed by cheese samples (4%) and butter samples (2%). All 36 *Listeria* sp. isolates which were obtained from biochemical identification were tested for *L. monocytogenes* by PCR, and 15 (41.6%) of them were confirmed as *L. monocytogenes*. According to the antibiotic resistance test results of *L. monocytogenes* strains, four (26.7%) isolates were resistant to at least one antibiotic while only one (6.7%) had multiple antibiotic resistance. The highest resistance was found against trimethoprim-sulfamethoxazole. It has been determined that *L. monocytogenes* isolates had the same degree resistance against penicillin G, meropenem, amikacin and vancomycin. In conclusion, the presence of *L. monocytogenes* which has been detected in dairy products made from raw milk and in raw milk itself may constitute a potential risk for public health.

Keywords: Milk, Dairy product, *L. monocytogenes*, *Listeria* spp., Antibiotic resistance

Çiğ Süt ve Süt Ürünlerinde *Listeria monocytogenes* Varlığı ve Antibiyotik Direnci

Öz


Anahtar sözcükler: Süt, Süt ürünleri, *L. monocytogenes*, *Listeria* spp., Antibiyotik direnci

INTRODUCTION

Listeria genus is a Gram-positive, spore-free and facultative anaerobic [1]. There are ten known pathogenic and non-pathogenic species. These include *Listeria monocytogenes*, *Listeria innocua*, *Listeria welshimeri*, *Listeria grayi*, *Listeria

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seeligeri, Listeria ivanovii, Listeria marthii, Listeria rocourtiae, Listeria weihenstephanensis and Listeria fleischmannii. Among these, L. monocytogenes is the primary pathogenic species for humans and causes serious diseases [1,2].

Listeria monocytogenes is an important pathogen originating from food, which has emerged in the second half of the 20th century. Listeria species are commonly found in raw and unprocessed food products. Listeriosis has high mortality and morbidity rate, and is caused by various food products including soft cheese, deli meats and vegetable products [3]. L. monocytogenes can be found in a wide spectrum of raw and processed products. Listeria spp. contamination has been detected in milk and dairy products, beef, pork, meat and meat products such as fermented sausage, fresh products like radish and cabbage and in seafood and fish [4]. This bacterial infection poses a serious threat for vulnerable individuals including immunocompromised patients, newborns, elderly people and pregnant women [5]. Generally, non-invasive gastrointestinal disease is seen in individuals with strong immune system. However, invasive listeriosis infections cause septicemia, meningitis infections and abortion in pregnant women in immunocompromised individuals [6].

Antibiotic resistance is a significant problem worldwide as many forms of resistance spread rapidly. A minimum of 2 million people are infected with serious diseases caused by resistant bacteria each year in the U.S. and at least 23,000 people die as a direct result of infections caused by these antibiotic-resistant bacteria [7]. Currently, the treatment for human listeriosis is ampicillin or penicillin G combined with gentamicin. Trimethoprim-sulfamethoxazole, vancomycin and erythromycin are second choice to treat listeriosis in pregnant women [8].

In a study conducted in Turkey by Kevenk and Terzi Gülel [9], the presence of L. monocytogenes in a total of 210 raw milk and milk samples was investigated, and in five (5%) milk, one (5%) white cheese, three (30%) kuymak cheese, three (30%) cokkelek, two (20%) farm cheese samples Listeria monocytogenes were positive while in kashar cheese, butter and ice cream Listeria monocytogenes were not detectible. Sağun et al. [10] has reported that six of the raw milk samples (2.40%) were Listeria spp. positive, and three (1.20%) of them were L. monocytogenes, one (0.40%) was L. innocua and one (0.40%) was found to be L. welshimeri positive. While 13 of the herb cheese samples (5.11%) were Listeria spp. positive, 10 (3.93%) of these isolated strains were also L. monocytogenes, one (0.39%) was L. ivanovii, one (0.39%) was L. innocua and one (0.39%) was L. welshimeri positive. Soyutemiz et al. [11] and Erol and Şireli [12] detected 3% and 5% L. monocytogenes from the raw milk, respectively. In another study, Arslan and Özdemir [13] have reported that they detected 9.2% of L. monocytogenes in white cheese samples.

It is estimated that 1600 people suffer from L. monocytogenes and 1.500 of them are hospitalized and 260 deaths are reported every year in the United States [14]. The incidence of listeriosis has been reported as 633 person among total population in Turkey of 68 million for 2003 [15]. Majority of listeriosis epidemics have been reported to occur due to the consumption of contaminating milk and dairy products [10,16]. About half of the reported listeriosis outbreaks in Europe have been associated with dairy products and the cause of outbreaks is mostly attributed to consumption of raw or unpasteurized milk products. Outbreaks from non-pasteurized soft cheeses in Switzerland between 1983 and 1987, from unpasteurized milks in Austria in 1986, and from Brie-type cheese in 1995 which was produced from unpasteurized milk in France put forth the risks of soft cheese consumption [17].

In this study, the presence of L. monocytogenes in cheese and butter produced from raw milk by traditional methods and its resistance against some antibiotics was determined.

MATERIAL and METHODS

Sampling

In this study, a total of 300 samples, including 100 samples of raw milk, 100 white cheese and 100 butter samples were obtained from different retail stores in Kars between October 2012 and March 2013. White cheese and butter samples are produced from raw milk in the traditional way which are offered for sale without any packaging. The samples were put in sterile bags and delivered to the laboratory as soon as possible under cold chain and were analyzed immediately.

Isolation of Listeria Species

For the isolation of Listeria species from the samples, the FDA (Food and Drug Administration) method as recommended by Hitchins [18] was used. According to this method, 25 g/25 mL sample was homogenized in 225 mL Buffered Listeria Enrichment Broth (BLEB, Oxoid, CM 897 Basingstoke, UK) medium and incubated at 30°C for 4 h. Subsequently, selective supplement (Listeria Selective Enrichment Supplement, SR0141) was added and incubation was completed to a total of 48 h. This pre-enrichment culture was inoculated to Listeria Selective Agar Base (LSA, Oxoid CM0856, Basingstoke, UK) medium consisting of Listeria Selective Supplement (Oxoid SR0140) by streak plate technique. The LSA plates were incubated at 37°C for 48-72 h. Blackish green brown colonies 2-3 mm in diameter surrounded by a black zone and having a sunken center which grew in the medium following the incubation were evaluated as suspected Listeria colonies. Listeria suspected colonies were inoculated by the streak plate method on Tryptone Soya Yeast Extract Agar (TSYEa) prepared as 30 g/L CASO Broth (Merck 1.05459), 6 g/L yeast extract (Merck 1.03753) and 15 g/L agar (Merck 1.01613) for
purification and identification procedures, and incubated at 30-37°C for 24 h.

Identification of Listeria Species

For the identification of the isolated strains, Gram-staining, oxidase (Bactident Oxidase, Merck) and catalase tests were applied to colonies growing in TSYEA. The Microbact™ 12L (MB1128A, Oxoid, Basingstoke, UK) test kit was used in order to identify Gram-positive bacil and coccobacil, oxidase negative and catalase positive colonies [19] on the basis of the species [20]. Each colony was incubated at 37°C for 24 h in a nutrient broth followed by transfer to each well in the 100 µL strain test kit. After incubation for 4-24 h, test results were recorded in the assessment form. The reaction results obtained were compared with the table provided in the kit. The results were assessed in the computer-aided identification program of Microbact 12L. As a positive control, L. monocytogenes ATCC 7644 strain was used in biochemical tests.

Genetic Identification of the Isolates

DNA extraction kit (QIAGEN QIamp DNA mini kit-51304) was used for the genomic DNA extraction from the isolates. The extraction procedure was performed in accordance with the recommendations of the manufacturing company. For the amplification of the extracted DNAs, the method recommended by Aznar and Alarcon [21] was used in a modified way. The target gene for L. monocytogenes was selected as hylA, and LM1: CCTAAGACGCCAATCGAA and LM2: AAGCGCTTGCAACTGCTC primers were used [22]. For PCR, master mix PCR buffer (10 mmol/L, Tris-HCl, pH 8.8, 1.5 mmol/L MgCl2, 50 mmol/L KCI, 0.1% Triton X-100) was prepared in a total volume of 25 µL with 1 µmol/L of each primer (Methabion, Germany), 100 µmol/L of each dNTP (Sigma), 1 U of Taq DNA Polymerase (Sigma D1806) and 5 ng DNA/µL. Thermal conditions for PCR were applied in TProfessional Basic Thermocycler (Biometra, Model TProfessional Basic Gradient) device as initial denaturation at 94°C for 5 min, 30 cycles of denaturation at 94°C for 30 sec, annealing at 50°C for 1 min, synthesis (extension) at 72°C for 1 min and final extension at 72°C for 5 min. 0.5 X TBE (Tris-Borate-EDTA) buffer at 1% agarose gel was used in order to observe the PCR products. L. monocytogenes (ATCC 7644) was used as positive control and the hlyA gene was visualized at 702 bp under UV illumination.

Antibiotic Resistance of the Isolates

Antimicrobial susceptibility tests of the isolates were carried out by disc diffusion method according to the Clinical and Laboratory Standards Institute [23,24]. After the isolates were activated, inoculation onto a TSA medium was performed. A suspension was prepared directly in physiological saline solution from single colonies. The density of the suspension was adjusted in a turbidity equivalent to the McFarland 0.5 standard. Inoculation was conducted on Mueller-Hinton agar containing 5% sheep blood by streaking the sterile swab over the surface. Four antibiotic discs were placed on each plate. The plates were incubated at 35°C for 24 h. After the incubation, inhibition zones on the plates were measured and recorded as resistant, intermediate and susceptible. Staphylococcus aureus ATCC 25293 was used as a quality control standard. Antimicrobial susceptibility test results were evaluated according to the intervals recommended by CLSI for Staphylococcus spp and by EUCAST for L. monocytogenes [23-25]. The following antibiotics were used for antimicrobial susceptibility tests: amikacin (30 µg), ampicillin (10 µg), cefotaxime (30 µg), chloramphenicol (30 µg), ciprofloxacin (5 µg), erythromycin (15 µg), gentamycin (10 µg), meropenem (10 µg), penicillin G (10 U), rifampin (5 µg), streptomycin (10 µg), tetracycline (30 µg), trimethoprim/sulfamethoxazole (1.25/23.75 µg), vancomycin (30 µg).

RESULTS

Isolation and Identification of Listeria spp.

Thirty-six Listeria spp. were isolated from a total of 300 foods having milk and milk products. As a results of the biochemical tests on these isolates using the Microbact 12 L, 22 (61.1%) L. monocytogenes, four (11.1%) L. seeligeri,

<table>
<thead>
<tr>
<th>Type of Sample</th>
<th>No. Listeria spp. Positive Samples</th>
<th>No. Listeria spp. Positive Isolates</th>
<th>Microbact 12L</th>
<th>PCR</th>
</tr>
</thead>
<tbody>
<tr>
<td>Raw milk</td>
<td>10 (10%)</td>
<td>26</td>
<td>16</td>
<td>4</td>
</tr>
<tr>
<td>White cheese</td>
<td>4 (4%)</td>
<td>8</td>
<td>5</td>
<td>2</td>
</tr>
<tr>
<td>Butter</td>
<td>2 (2%)</td>
<td>2</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>Total</td>
<td>16 (5.3%)</td>
<td>36</td>
<td>22</td>
<td>4</td>
</tr>
</tbody>
</table>

<table>
<thead>
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<tr>
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<td>10 (10%)</td>
<td>16</td>
<td>4</td>
</tr>
<tr>
<td>White cheese</td>
<td>4 (4%)</td>
<td>5</td>
<td>2</td>
</tr>
<tr>
<td>Butter</td>
<td>2 (2%)</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>Total</td>
<td>16 (5.3%)</td>
<td>36</td>
<td>4</td>
</tr>
</tbody>
</table>

Table 1. Identification and verification of Listeria spp isolates obtained from raw milk and dairy products
eight (22.2%) L. ivanovii, one (2.7%) L. welshimeri and one (2.7%) was identified as L. innocua. When distribution of the isolates in term of foods was examined, 16 of the 300 samples (5.3%) were found to be positive for Listeria spp. Listeria spp. were isolated the highest from raw milk samples (10%), followed by cheese (4%) and butter samples (2%). Identification of Listeria spp. isolates obtained from raw milk and dairy products was shown (Table 1).

Identification of Listeria monocytogenes by Molecular Techniques

A total of 36 Listeria spp. isolates identified by Microbat 12L were tested genetically for L. monocytogenes by PCR. Fig. 1 shows the band obtained by PCR. Of the 36 isolates examined, 15 (41.6%) were identified as L. monocytogenes. According to the PCR results, 11 (3.7%) out of total 300 foodstuff L. monocytogenes were detected. The test results of L. monocytogenes using PCR were given (Table 1).

Antibiotic Resistance

Antimicrobial resistance profiles of 15 isolates confirmed by PCR as L. monocytogenes were examined. Of the 15 L. monocytogenes positive isolates, four (26.7%) of them were resistant to only one antibiotic while one (6.7%) were resistant to more than one antibiotic, and 10 (66.6%) were not resistant to any antibiotics. The highest resistance was against trimethoprim-sulfamethoxazole. It has been determined that L. monocytogenes isolates had the same degree resistance against penicillin G, meropenem, amikacin and vancomycin. The antimicrobial resistance of L. monocytogenes isolates was shown in (Table 2).

DISCUSSION

It has been demonstrated that milk and dairy products are responsible for approximately 50% of listeriosis epidemics and various sporadic cases in Europe [3,17,26,27]. In this study,

<table>
<thead>
<tr>
<th>Antibacterial Agent</th>
<th>Listeria monocytogenes (%)</th>
<th>R</th>
<th>I</th>
<th>S</th>
</tr>
</thead>
<tbody>
<tr>
<td>Amikacin</td>
<td>1 (6.7)</td>
<td>2 (13.3)</td>
<td>12 (80)</td>
<td></td>
</tr>
<tr>
<td>Ampicillin</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>Cefotaxime</td>
<td>0 (0)</td>
<td>1 (6.7)</td>
<td>14 (93.3)</td>
<td></td>
</tr>
<tr>
<td>Chloramphenicol</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>Ciprofloxacin</td>
<td>0 (0)</td>
<td>1 (6.7)</td>
<td>14 (93.3)</td>
<td></td>
</tr>
<tr>
<td>Erythromycin</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>Gentamycin</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>Meropenem</td>
<td>1 (6.7)</td>
<td>0 (0)</td>
<td>14 (93.3)</td>
<td></td>
</tr>
<tr>
<td>Penicillin G</td>
<td>1 (6.7)</td>
<td>0 (0)</td>
<td>14 (93.3)</td>
<td></td>
</tr>
<tr>
<td>Rifampin</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>Streptomycin</td>
<td>0 (0)</td>
<td>3 (20)</td>
<td>12 (80)</td>
<td></td>
</tr>
<tr>
<td>Ticarcycline</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>15 (100)</td>
<td></td>
</tr>
<tr>
<td>SXT</td>
<td>4 (26.7)</td>
<td>0 (0)</td>
<td>11 (73.3)</td>
<td></td>
</tr>
<tr>
<td>Vancomycin</td>
<td>1 (6.7)</td>
<td>0 (0)</td>
<td>14 (93.3)</td>
<td></td>
</tr>
</tbody>
</table>

* SXT: Trimethoprim-Sulfamethoxazole R: Resistant I: Intermediate S: Susceptible; a Results were evaluated for Staphylococcus spp. according to the intervals recommended by CLSI 2012, 2014; b Results were evaluated for L. monocytogenes according to the intervals recommended by EUCAST 2015

Fig 1. Amplification products obtained through PCR. Lane M: 100-bp DNA ladder, Lane 1-3: Listeria monocytogenes PCR amplification products, Lane 4: Positive control for Listeria monocytogenes, Lane 5: DNA free PCR mixture for Listeria monocytogenes negative control

L. monocytogenes was found to be positive in 6% of the raw milk samples. In partial consistency with our findings, Keven and Terzi Gülel [9], Erol and Şireli [12] and Jamali et al.[1] reported that they isolated L. monocytogenes from raw milk at the rates of 5%, 5% and 5.4% respectively. In contrast, Gebretsadik et al.[28] and Usman et al.[29] reported isolation of L. monocytogenes from raw milk at higher rates than our results revealed, i.e. at 13% and 8.3% respectively. On the other hand, Sağun et al.[18], Soyutemiz et al.[31], Rahimi et al.[30] Aygün and Pehlivanlar [30], Mahmoodi [31], Shamloo et al.[32]. Ünlü et al.[33] and Osman et al.[34] reported isolation of L. monocytogenes at the rates of 1.2%, 3%,...
1.1%, 0%, 1.7-3.3%, 4.39%, 4% and 2% respectively. These results are lower than our findings.

Aygün and Pehlivanlar [30] and Mahmodii [31] reported isolation of *L. monocytogenes* from white cheese at the rates of 2.35% and 3.3-6.7% respectively (at two different production sites). Our study also determined that 3% of the white cheese samples were *L. monocytogenes* positive which is a finding close to those of the above-mentioned researchers. On the other hand, Keven and Terzi Güle [9], Arslan and Özdemir [13], Rahimi et al. [16], Akkaya and Alişarlı [33], and Gülmez and Güven [30] reported isolation of *L. monocytogenes* at the rates of 5%, 9.2%, 15%, 6% and 5% respectively. These findings are higher than ours. On the other hand, Shamloo et al. [12] reported no isolation of *L. monocytogenes* in traditionally manufactured cheese samples. Karadal and Yıldırım [37] reported isolation of *L. monocytogenes* at a rate of 1% from both white cheese and tulum cheese made from raw milk by using traditional methods. These findings are lower than ours. It is observed that there are differences between the findings obtained in our study and the findings reported by other researchers. The reasons behind such difference may be the contamination level of the milk used in cheese production, environmental and seasonal differences, the differences in the traditional cheese production techniques and secondary contamination.

In our study, it was determined that 2% of the butter samples were contaminated with *L. monocytogenes*. Rahimi et al. [18] reported isolation of *L. monocytogenes* in 4% of the traditional butter samples. This result was higher than our findings. As for the findings of other researchers, Keven and Terzi Güle [9], Aygün and Pehlivanlar [30], Shamloo et al. [12] and Rahimi et al. [38] reported that none of the butter samples were identified with *L. monocytogenes*. Our findings are higher than those of the above-mentioned researchers. This is considered to have resulted from the differences in the butter production techniques and especially lack of pasteurisation process. Other reasons may be cross contamination of the samples, inattentiveness to cold storage conditions during storage and marketing, and poor hygiene.

The number of *Listeria spp.* found in food products may vary depending on the type of food tested. It has been reported that the detection of *L. monocytogenes* in complex food matrices such as cheese may be more difficult than milk [30]. Thus, the present study revealed the highest level of *L. monocytogenes* contamination in raw milk followed by white cheese and butter. It is observed that various studies have obtained different results in similar products. This may be due to the isolation and identification methods used.

In general, *L. monocytogenes* is susceptible to those antimicrobials that are effective against Gram-positive bacteria whereas antimicrobial resistance against these bacteria has been reported in isolates obtained from food producing animals, food processing environments and food [40]. In this study, it has been detected that *L. monocytogenes* isolates are resistant to trimethoprim-sulfamethoxazole, penicillin G, meropenem, amikacin and vancomycin. The highest resistance has been found against trimethoprim-sulfamethoxazole. In parallel to our findings, Jamali et al. [11], Usman et al. [29], Osman et al. [34] and Harakeh et al. [19] reported resistance to trimethoprim-sulfamethoxazole. On the other hand, Karadal and Yıldırım [37] and Mackiw et al. [40] indicated no resistance to trimethoprim-sulfamethoxazole. Many researchers [1,8,13,16,29,34,38,39] reported resistance to penicillin G. In parallel to these results, our study also determined resistance to penicillin G. On the other hand, Karadal and Yıldırım [37] and Conter et al. [34] reported no resistance to penicillin G. Aydin et al. [34] reported that *L. monocytogenes* strains of food origin are resistant to meropenem, penicillin G and trimethoprim-sulfamethoxazole. In parallel to these findings, this study also determined resistance to the three antibiotics indicated. In this study, resistance to amikacin has been detected. Some researchers reported resistance to amikacin in *L. monocytogenes* isolates of food origin in parallel to our findings [40]. In our study, *L. monocytogenes* isolate with vancomycine resistance has been found. Our findings are consistent with the findings of certain researchers [9,33,41]. However, some researchers [1,16,34,38] reported that there was no vancomycine resistance. *L. monocytogenes* isolates can develop resistance by acquiring mobile genetic components such as mobilizable plasmids and conjugative transposons. Further, mutational events in chromosomal genes may play a role in giving *Listeria spp.* resistance [39].

Many variables including the number and rate of saprophytic microorganisms available in the environment, geographic regions and infected food affect genetic variety and the resistance of *L. monocytogenes* serotypes to antibiotics which in turn lead to different research results [44]. The present study also involves results which are different from those of some other researchers.

In conclusion, this study has revealed the presence of *L. monocytogenes* in raw milk and dairy products made with raw milk in the region. It has been concluded that this may constitute a potential risk to public health and that there is a need for enhancing the safety of milk and dairy products. Implementing good food and good manufacturing practices is extremely important for consumer health. It is essential to take the necessary hygienic measures during the entire process starting from the production of milk and dairy products until their consumption. In order to provide product safety, it is primarily recommended that pasteurisation conditions are fulfilled, cross contamination is prevented, cold storage conditions are complied with and especially unpackaged products are kept under inspection.
CONFLICT OF INTEREST

The authors declare no conflict of interest.

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