**Original Article**

**Isolation of Listeria monocytogenes from Meat and Dairy Products**

Ashraf Haj Hosseini1, *Anousheh Sharifan*, Akram Tabatabae2

1Department of Food Science and Technology, Science and Research Branch, Islamic Azad University, Tehran, Iran; 2Department of Biology, East Tehran Branch, Islamic Azad University, Tehran, Iran.

Received Feb 06, 2016; accepted May 03, 2016

**Introduction:** This study was intended to determine the presence and distribution of *Listeria monocytogenes* in various meat and dairy products from Qazvin Province by culture followed by biochemical and morphological assays. The identity of the isolates was further obtained by amplification of *pfA* gene in bacteria isolates. This gene is a transcriptional activator of virulence gene expression within the pathogenic *L. monocytogenes*. **Method:** In a cross-sectional design, a total of 182 different food samples were collected from different areas in Qazvin, Iran. Bacterial isolates were obtained by the cold enrichment method. DNA extraction from the pelleted cells was conducted and then *pfA* gene was amplified by conventional PCR. **Results:** As many as 37 (20.3%) food samples were positive for *Listeria* spp. including 21 (56.8%) *L. monocytogenes*, 7 (18.9%) *Listeria innocua*, 4 (10.8%) *Listeria welshimeri*, 3 (8.1%), *Listeria seeligeri*, and 2 (5.4%) *Listeria grayi*. None of the isolated specimen was *Listeria ivanovii*. The *PfA* gene was amplified in all *L. monocytogenes* specimen. Moreover, PCR assay had high sensitivity and specificity for bacterial identification. **Conclusion:** To sum up, presence of *L. monocytogenes* in food samples was confirmed in this region, it was more frequent in milk specimen. In addition to common culture techniques, PCR assay showed higher sensitivity and specificity for *L. monocytogenes* detection in contaminated foods. *J Med Microbiol Infec Dis*, 2014, 2 (4): 159-162.

**Keywords:** *Listeria monocytogenes*, *PfA*, PCR, Milk, Meat.

**INTRODUCTION**

*Listeria* spp. are Gram-positive and facultative anaerobic bacteria, with a low G+C content in their genetic material. The genus comprises six species, namely *Listeria monocytogenes*, *Listeria ivanovii*, *Listeria seeligeri*, *Listeria innocua*, *Listeria welshimeri*, and *Listeria grayi*. The only pathogenic species of this genus is *L. monocytogenes*, although there have been few reports on the capability of causing disease by *L. seeligeri*, *L. ivanovii* and *L. innocua* [1].

*L. monocytogenes* is an intracellular pathogen, it is referred as a main cause of human food-borne infections across the globe [2]. Food-borne listeriosis induced by *L. monocytogenes* is a relatively rare, yet a serious condition with a greater mortality (20-30%) than other food-borne microbial pathogens, such as *Salmonella* spp. [3]. While *L. monocytogenes* commonly cause a mild gastroenteritis in healthy adults, it may be associated with high severity in susceptible individuals. Basically, *L. monocytogenes* frequently affects immunocompromised individuals, pregnant women, and elderly people. The signs and symptoms of *L. monocytogenes* infection range from flu-like illness to meningitis and septicemia and it may lead to spontaneous abortion or listeriosis of the newborn [3].

*Listeria* species are tolerant to harsh circumstances including low pH, low temperature and high salt [4, 5]. Accordingly, they are able to grow in various soil environments, sewages, silages, water, effluents and various food items. Concurrent with globalization and growing popularity of manufactured ready-to-eat foods, much attention has been currently drawn to *L. monocytogenes*.

On the other hand, the huge amounts of reports on contamination in food industry products by pathogenic bacteria have raised a great public concern. *Listeria* species, *L. monocytogenes* in particular, have been recognized in different food sources such as meat and chicken [6]. The presence of *Listeria* spp. in meat is a serious threat for food safety since the organism can grow in both raw and cooked meat during refrigerated storage [7]. There have been some reports on successful isolation of the *Listeria* spp. across the globe. However, no thorough records on outbreaks of human listeriosis are available [8]. Over the past 25 years, listeriosis has been a health-threatening food-related disease leading to considerable number of deaths. There has been plenty of evidence regarding this issue that outbreaks of human listeriosis have been associated dairy products namely milk [9] and soft cheese [10]. Given the growing concern over *L. monocytogenes* infection as well as its potential adverse effects on the human’s health, it seems a priority to conduct research on contamination of various food sources with this pathogenic agent in different parts of the country, including Qazvin Province. Therefore, the aims of this study were: (i) to specify the prevalence of *L. monocytogenes* in three different food categories; (ii) and to specify the performance of two pairs of primers for identifying *L. monocytogenes*.

**Correspondence:** Anousheh Sharifan
Department of Food Science and Technology, Science and Research Branch, Islamic Azad University, Daneshgah Blvd, Hesarak, Tehran, Iran, 1477893855.

**Email:** anousheh.sharifan123@yahoo.com

**Tel:** +98 (21) 44867154 **Fax:** +98 (21) 66460700

http://jommid.pasteur.ac.ir
MATERIAL AND METHODS

Different samples were randomly selected from dairy products (raw milk, soft cheese, butter and yogurt, n=84) as well as raw and processed meat (beef, chicken, sausages, chicken nugget and burgers, n=98). They were directly transferred to the laboratory using an ice box. The International Standards Organization (ISO) 11290 method with a two-stage enrichment process, which inhibits competitive organisms, was utilized for isolation and identification of L. monocytogenes [11]. The samples were thawed overnight in refrigerator and then were aseptically and carefully separated from their casings and grounded completely by a sterile mortar. The isolation was initially carried out by the use of Listeria selective enrichment broth (LSEB) to increase the growth of Listeria spp. The LSEB base contained trypticase soy broth besides 0.6% yeast extract supplemented with Listeria selective supplement (Hi Media laboratories) including acriflavine-HCL (15 mg L⁻¹), nalidixic acid (40 mg L⁻¹), and cycloheximide (50 mg L⁻¹). Amounts of 10 g of meat samples were aseptically mixed with LSEB (90 mL). For dairy products, 25 g of samples were aseptically mixed with 225 mL Brain Heart Infusion (BHI). All samples were incubated at 4°C for 7 to 20 days. A loopful of the homogenate was used to streak onto Listeria selective agar base (BHI and LSEB agar Media laboratories) followed by incubation at 37°C for 24 h. Identification of suspected colonies of Listeria was performed by various assays including Gram staining, catalase reaction, umbrella-shaped motility pattern, hemolysis on sheep blood agar, fermentation of mannitol, rhamnose, xylose, glucose and maltose, and MR-VP and CAMP tests, in accordance with Bergey’s Manual of Systematic Bacteriology [12] as summarized in table 1. Cooling enrichment methods were performed for Listeria spp. isolation during four months as well.

DNA extraction and PCR assay. The obtained culture (10 mL) was cultivated to mid-log growth phase using Trypsophas Soy (TYS) broth for every Listeria isolates. Of which, 1 mL of cells were pelleted via centrifugation at 13.000×g for 5 min. The pellets were subsequently resuspended and centrifuged again at 12.500×g for 15 min. DNA was then extracted from the pelleted cells using the Wizard genomic DNA purification kit (Promega, USA) and stored at -20°C until used. Two primers, LIS-F and LIS-R, were used (Table 2) to amplify the prfA gene of L. monocytogenes as described by others [13]. The final 25 μL PCR reactions contained 2 μl of extracted DNA, 12.5 μl GoTaq Green Master Mix (Promega, USA), forward primer and reverse primers, 500 μM each, and Ultra-Pure DNase/RNase-Free distilled Water (Gibco, USA). The amplifications were performed in a thermocycler (Techne, Cyclogene) with pre-incubation at 95°C for 30 s, denaturation at 95°C for 30 s, annealing at 54°C for 30 s, extension at 72°C for 30 s, as well as final elongation at 72°C for 5 min.

PCR products were resolved through 1% agarose gels and stained using ethidium bromide. Detection was performed via a short-wavelength ultraviolet light source and photographed by an EDVOTEK Gel documentation system. The method adopted by Jami et al. was utilized to measure the sensitivity and specificity of the primers. Briefly, serial dilutions based on the overnight culture of L. monocytogenes in BHI broth were enumerated applying surface plating technique. As for the specificity of the primers, DNA extracted from various bacterial colonies was taken into account as the template for the m-PCR assay [14].

Table 1. Main laboratory tests for the differentiation of Listeria spp.

<table>
<thead>
<tr>
<th>Species</th>
<th>Hemolysis</th>
<th>Phosphatidylinositol phosphatase</th>
<th>Acid production</th>
<th>Surface motility</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>L. monocytogenes</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>Liss 1983</td>
</tr>
<tr>
<td>L. ivanovi</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td></td>
</tr>
<tr>
<td>L. seeligeri</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td></td>
</tr>
<tr>
<td>L. innocua</td>
<td>-</td>
<td>-</td>
<td>V</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>L. welshimeri</td>
<td>-</td>
<td>-</td>
<td>V</td>
<td>+</td>
<td></td>
</tr>
<tr>
<td>L. grayi</td>
<td>-</td>
<td>-</td>
<td>V</td>
<td>-</td>
<td></td>
</tr>
</tbody>
</table>

Rapid Method (PCR)

Table 2. Oligonucleotide sequences used for identification of L. monocytogenes by PCR

<table>
<thead>
<tr>
<th>Target gene</th>
<th>Primer sequence (5‘ - 3’)</th>
<th>Amplified fragment length</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>prfA gene</td>
<td>LIS-F: TCA TCG ACG GCA ACC TCG G</td>
<td>217 bp</td>
<td>[13]</td>
</tr>
<tr>
<td></td>
<td>LIS-R: TGA GCA ACG TAT CCT CCA GAG T</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Table 3. Isolation rate of Listeria spp. in different food and human sample

<table>
<thead>
<tr>
<th>Type odd samples</th>
<th>No. of analyzed sample</th>
<th>Positive sample</th>
<th>%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Meat products</td>
<td>98</td>
<td>12</td>
<td>32.4</td>
</tr>
<tr>
<td>Milk products</td>
<td>84</td>
<td>25</td>
<td>67.6</td>
</tr>
<tr>
<td>Total</td>
<td>182</td>
<td>37</td>
<td>100</td>
</tr>
</tbody>
</table>
RESULTS

Tables 3 and 4 presented data for the analysis of milk and meat products. As many as 37 (20.3%) food samples were positive for Listeria spp. including 21 (56.8%) L. monocytogenes, 7 (18.9%) L. innocua, 4 (10.8%) L. welshimeri, 3 (8.1%), L. seeligeri, and 2 (5.4%) L. grayi. None of the isolated specimen was L. ivanovii. The PrfA gene in L. monocytogenes was detected in all samples. Meat and chicken products were frequently contaminated with L. monocytogenes. The performance of the primers was confirmed by measuring their sensitivity and specificity. The findings indicated that these primers possess high sensitivity (96.9%) and specificity (100%) for the target sequence and merely for Listeria spp. (data not shown).

**DISCUSSION**

The recurring occurrence of L. monocytogenes in food items including meat or dairy products may create formidable challenges on the part of customers. This is claimed that human infections foundationally arise from consuming contaminated foods which in turn, are associated with serious, if not life-threatening conditions of listeriosis [7, 14]. It has been indicated that the food borne transmission of L. monocytogenes is one of the pivotal causes behind such cases [10, 15].

Molla et al. revealed that raw meat products possessed a high level of contamination (50.6%) with Listeria spp. [16]. Jamali et al. reported that prevalence of L. monocytogenes in ready-to-eat food samples was 45 out of 396 [17]. It is assumed that the slaughter methods (evisceration), food handling practice, and food processing increase the probability of microbial contamination of Listeria spp. in meat products. Moreover, Listeria spp. widely spread out in the environment [13]. Panahi Dorcheh et al. found that the prevalence of Listeria spp. in meat, liver, and feces of quail were 15, 5, and 30%, respectively. More to the point, heart and kidney demonstrated to be free of Listeria spp. In their study, L. innocua showed the high frequency, even more than Listeria spp. [18]. El Marnissi et al. exhibited 5.90% as an overall prevalence of L. monocytogenes in raw milk [15]. Boubendir et al. observed a comparable occurrence in bovine raw milk from the North Eastern Algeria [19]. Jami et al. reported a lower contamination rate for milk samples in Mashhad, Iran [14]. Therefore, it was concluded that meat products have been presented to be at high risk for L. monocytogenes contamination rather than dairy ones.

Furthermore, the current disparity of the prevalence depends on geographic location, sampling technique, detection procedure, types of media utilized, and hygienic production [14, 20].

L. innocua has shown to grow more rapidly than other pathogenic species in enrichment broth media, and it is more likely to overgrow L. monocytogenes. L. innocua holds the similar ecological niche with high incidence. In other words, L. innocua is capable of being contaminated by L. monocytogenes [21]. Lower records were found in China, where the prevalence rate was 0.23% to 1.2% [22]. As for common sources of L. monocytogenes in raw milk, following parameters are influential: milking processing, transportation, storage methods, infected cows, and silage quality [14, 23].

Akin to the study by Jami et al., our study corroborated the sensitivity and the specificity of the applied primers for identification of L. monocytogenes [14]. Therefore, it is recommended that application of common culture techniques along PCR assay can considerably enhance the chance of detecting L. monocytogenes dairy and meat products.

This study revealed the presence of L. monocytogenes and other Listeria spp. in various dairy and meat products collected from different areas of Qazvin province, Iran. Moreover, the combination of culture methods and PCR analysis can afford to confirm whether the presumptive isolates were L. monocytogenes.
ACKNOWLEDGEMENT
This study was supported by Shahd Ar Caspian Company.

CONFLICT OF INTEREST
The authors declare that there are no conflicts of interest associated with this manuscript.

REFERENCES