## Listeria spp IDENTIFICATION by PCR

MOURA, G. F.1, CASTRO, V. S1; FIGUEIREDO, E.E.S.1,2

<sup>1</sup>Universidade Federal de Mato Grosso, Programa de Pós Graduação em Ciência Animal, Cuiabá-MT. Brasil.

<sup>2</sup>Universidade Federal de Mato Grosso, Faculdade de Nutrição Cuiabá-MT, Brasil.

The genus Listeria spp. comprises fifteen species and six subspecies. Among these, L. monocytogenes, L. ivanovii and L. seeligeri can be associated with diseases in humans, and, specifically, L. monocytogenes can cause severe food-borne illness, where the manifestation can vary from a simple cold to meningitis, encephalitis, septicemia and abortion. The detection of Listeria spp is usually performed based on classical culture and biochemical identification, methods which are laborious and slow, requiring around ten days to complete, of which approximately two days are intended for selective enrichment only. Targeting a specific and quick test as a screening to replace this step, the Polymerase Chain Reaction (PCR) technique has proven efficient, reducing the assay time and cost of analysis. The present study aimed to perform a PCR standardization for the rapid and specific identification of Listeria spp. Primers designed for the 16S-23S region, previously described in the literature, IGS 1 (5' -GGCCTATAGCTCAGCTGGTTA-3') and IGS 2 (5' -GCTGAGCTAAGGCCCCGTAAA-3'), were selected and used for the internal standardization of optimal Listeria spp.-specific PCR conditions. To optimize the amplification conditions, reference and environmental strains of Listeria monocytogenes (ATCC 19117), Listeria inoccua, Escherichia coli (ATCC 43895), Staphylococcus aureus and Salmonella thyphi (ATCC 10749) after growth were subjected to DNA extraction with the aid of a commercial kit Blood and Tissue (Qiagen®) and quantification by fluorimetry using Qubit (Invitrogen). The conditions for the amplification of a 200 bp fragment were established and the analytical specificity and sensitivity were determined. The PCR was standardized using 66 ng target DNA, 100 mM dNTPs, 100 mM of each primer, 2 mM MgCland 1U Taq DNA polymerase. The amplification cycle adopted the following parameters: initial denaturation 95 ° C for 10 min, 30 cycles of 95 ° C for 1 min, 61 ° C for 1 min, 72 ° C for 1 min and final extension at 72 ° C for 7 min. The resulting PCR products were analyzed by gel electrophoresis on 1.5% ultrapure agarose gels stained red. The PCR reaction showed sensitivity and specificity of 100% for the tested conditions, detecting only members of the Listeria spp. genus. This PCR assay is efficient, therefore, in reducing the assay time and cost analysis, and could be used as a screening step to identify suspected Listeria spp. colonies.

Keywords: listeriosis, 16S-23S region, molecular diagnostics

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