

TITLE: IDENTIFICATION OF LISTERIA SPP. BY MALDI-TOF MASS SPECTROMETRY IN FECES AND FOOD.

AUTHORS: CAMPOS, T. M.¹; REIS, E. M. F.¹; PEREIRA, R. C. L.¹; BOTELHO, L. A. B.²; HOFER, E.¹; HOFER, C. B.³; VALLIM, D. C.¹

INSTITUTION: ¹.LABZOO/IOC/FIOCRUZ - LABORATÓRIO DE ZONÓSES BACTERIANAS/ INSTITUTO OSWALDO CRUZ, RIO DE JANEIRO, RJ (AVENIDA BRASIL, 4365, PAVILHÃO ROCHA LIMA, 3º ANDAR, RIO DE JANEIRO – RJ, BRAZIL).

².LIN/IMPPG/UFRJ – LABORATÓRIO DE INVESTIGAÇÃO EM MICROBIOLOGIA MÉDICA/INSTITUTO DE MICROBIOLOGIA PROF. PAULO DE GÓES, RIO DE JANEIRO, RJ (AV. CARLOS CHAGAS FILHO, 373, BL.I, SALA i2 059, CIDADE UNIVERSITÁRIA, RIO DE JANEIRO – RJ, BRAZIL).

³.UNIVERSIDADE FEDERAL DO RIO DE JANEIRO, HUCFF/IPPMG (AV BRIG TROMPOWSKY, S/ NO., ILHA DO FUNDÃO, 21044-020- CIDADE UNIVERSITÁRIA, RJ – BRASIL)

ABSTRACT:

Listeria monocytogenes is a food-borne pathogen agent of human listeriosis, an opportunistic infection that primarily infects pregnant, neonates and immunologically compromised individuals. Rapid and accurate discrimination among *Listeria* strains is essential for appropriate therapeutic and timely intervention for infection control. Matrix Associated Laser Desorption Ionization–Time of Flight Mass Spectrometry (MALDI-TOF MS) is a rapid and simple microbial identification method, which is a promise for identification of *Listeria* spp. In the study, the performance of MALDI TOF MS in detection and identification of *Listeria* spp. directly from the broth for enrichment of fecal material and food samples was evaluated. In order to do that, 34 standard strains of different *Listeria* species belonging to the Collection of *Listeria* (CLIST/LABZOO-FIOCRUZ), as well as the broth for enrichment of 12 samples of pregnant feces and 15 samples of *sashimi* and *temaki*, all positive for *Listeria*, were analyzed by MALDI-TOF MS. Following a short cell extraction procedure with ethanol, formic acid and acetonitrile, the cell material from the broth was deposited on a sample target, dried, overlaid with a matrix necessary for the MALDI process and analyzed. The method was also confirmed using the conventional methodology and PCR with 23S rDNA and *hly* primers in the detection and identification of this pathogen, respectively. This study showed that 97% (n=33) of the standard strains were identified correctly. Except for one *L. innocua* that was identified only on the genus level, all other isolates were identified in both. Results revealed that was possible to detect *Listeria* in 40% (n=6) of the food enrichment broths, among them 27% (n=4) were identified at the genus and 13% (n=2) at the genus and species. From the fecal broth it was possible to identify the presence of *Listeria* in 25% (n=3), being 17% (n=2) identified at the level of genus and species and 8% (n=1) at the genus. The remaining 58% (n=7) of the total broth of fecal material weren't analyzed because it didn't remain viable until the end of the study. This was probably due to the long storage time, even under refrigeration (4°C), since these samples were collected in previous studies. This study indicates the possibility of identifying *Listeria* in samples contaminated directly from the enrichment stage, which can contribute to the reduction of costs of laboratory tests and provide a faster diagnosis of listeriosis.

Keywords: *Listeria* spp., MALDI-TOF MS, Enrichment broth, *Listeria monocytogenes*, Diagnosis.

Development Agency: CAPES, CNPq

