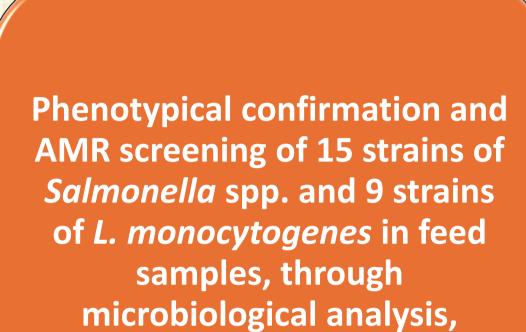
Phenotypic Characterization and Serotyping of Salmonella spp. and Listeria Monocytogenes Isolates From Feed Samples

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biochemical and serological

testing

\$\text{Strains of } Salmonella \text{ spp. were}

Strains of Salmonella spp. were susceptible to tetracycline, norfloxacin, ciprofloxacin, gentamicin, meropenem, while showed resistance to ampicillin, cefotaxime and ceftazidime, without any MDR being recorded

Solution L. monocytogenes strains were only susceptible to erythromycin and ampicillin, while one strain was MDR to the remaining antibiotics (tetracycline, penicillin, sulfamethoxazole, ciprofloxacin, meropenem)

\$\text{The recorded AMR of \$L\$.} \\ monocytogenes isolates was tetracycline, penicillin, sulfamethoxazole, ciprofloxacin, meropenem

Multiplex PCR was utilized to assign the isolated strains to the most prevalent and important public health-related Salmonella serotypes and L. monocytogenes PCR-serogroups

Salmonella isolates were classified into serotypes Thompson (60%), Typhimurium (6.7%) and Enteritidis (6.7%), whereas (26.6%) were identified as Salmonella spp.

\$\mathscr{G} L. monocytogenes isolates were classified into the PCR-serogroups IIa (44.5%), IIb (11.1%), IIc (11.1%) and Ivb (33.3%)

The results of the present study demonstrate the presence of important foodborne pathogenic bacteria with increased AMR to antibiotics caused at the primary production and at the farm level by the inappropriate use of pharmacological substances used to treat animal diseases, resulting in the potential detection of resistant bacterial strains of the pathogens to animal-originated food products



