TITLE: OVERVIEW OF VIRULENCE GENES OF BIOFILM-FORMING E. coli

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ABSTRACT:

Regarding the importance of the virulence factors of E. coli in the dairy environment, the association of the biofilm with its persistence in different places stands out. Thus, the objective of this work was to evaluate the biofilm production in E. coli strains related to the milk production environment. A total of 444 samples were collected within the dairy environment. The identification of E. coli was performed by biochemical tests and MALDI TOF MS. The microplate test was used to evaluate the biofilm production. The bacterial DNA was extracted by thermal lysis to evaluate the *fim*H, *csg*A and *flu* genes by PCR. A total of 41.44% (63/152) of E. coli strains phenotypically produced the biofilm. The fimH gene was highly expressed in the studied strains. The strains presented a total percentage of 92.1% (140/152) for the fimH gene being found in the feces 96.8% (91/94) followed by 93.3% (14/15) from the water, 82, 8% (29/35) of the milk and 75% (6/8) of the milking environment. In relation to the csgA gene, the strains had 88.8% (135/152) of which 94.6% (89/94) were isolated in feces, 93.3% (14/15) in water, 77.1% (27/35) in milk and 62.5% (5/8) in the milking environment. The total percentage of the presence of the flu gene in the studied strains was 29.6% (45/152), being 41.4% (39/94) found in the feces, 25% (2/8) in the milking environment, 20% (3/15) in the water and 2.8% (1/35) in the milk. It can be concluded that the high expression of the genes in fecal strains may have occurred due to the location of *E. coli* in the outermost layer of mucin in the colon, this glycoprotein can stimulate the formation of biofilm, together with the secreted IgA, favoring the E. coli biofilm in the intestinal mucosa through type 1 fimbria. This could be the contributing factor for the dispersion of this microorganism in the evaluated production environment.

Keywords: enterobacteria, milk environment, virulence

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