Title: THE INFLUENCE OF THREALOSE METABOLISM ON TYPE I FIMBRIA EXPRESSION IN EXTRAINTESTINAL *E. COLI* MT78

AUTHORS: KLEMBERG, VS¹; PAVANELO, DB¹, HORN, F¹.

INSTITUITION: ¹UNIVERSIDADE FEDERAL DO RIO GRANDE DO SUL-UFRGS, PORTO ALEGRE/RS. (AVENIDA BENTO GONÇALVES, 9500, CAMPUS DO VALE. PRÉDIO 43433. LABORATÓRIO 103. CEP: 91501-970).

Extraintestinal pathogenic *Escherichia coli* (ExPEC) are responsible for a number of infections outside the gastrointestinal tract, including urinary tract infections and neonatal meningitis in humans, and infections in farmed animals. In poultry, avian colibacillosis - as ExPEC infections are collectively called - causes great economic losses worldwide. After screening a library of 1710 mutants of the ExPEC MT78 strain generated by signature-tagged mutagenesis, one of the attenuated mutants had the *treA* gene affected. This mutant showed a decrease in type I fimbriae expression, a reduction of 35% in adhesion and of 65% in invasion of avian fibroblasts (CEC-32 cell line) compared to the wild type strain. The enzyme trehalase, encoded by the *treA* gene, degrades the disaccharide trehalose in the periplasm; otsA and otsB genes encode the cytoplasmic enzymes of trehalose biosynthesis. In E. coli K-12, the pathways of trehalose biosynthesis and degradation are involved in the response to osmotic stress. Thus, under osmotic stress, K-12 accumulates trehalose by increasing the expression of *treA*, otsA and otsB. In order to elucidate whether the excess of trehalose in the periplasm is affecting type I fimbriae expression, we generated mutants MT78 Δ otsAB and MT78 Δ *treAotsAB* by the non-polar mutation lambda red technique. Afterwards, we performed yeast agglutination tests to observe type I fimbriae expression in these mutants. Bacterial cultures were grown both statically for 24 hours and under shaking up to the mid-log phase (OD 600 nm ~ 0.6) in LB medium. An initial suspension of approximately 10¹¹ CFU in PBS was serially diluted 1:2 in microtiter plates. Equal volumes of a 1.5% yeast solution were added to the wells. After 30 minutes on ice, the agglutination was visually monitored and the most diluted titer with agglutination was determined. Strain DM34, a type I fimbria null mutant of MT78 was used as a negative control. The most diluted yeast agglutination titers were: 2^8 for wild-type strain, 2^6 for MT78 Δ otsAB and 2^3 for MT78 Δ treAotsAB. Our results indicate that excess of trehalose in the bacterial periplasm is not responsible for the observed reduction in type I fimbriae expression; this reduction may be explained by other mechanism yet to be elucidated.

Key words: E.coli MT78, otsA, otsB, treA, type I fimbriae.

Development Agency: CAPES