

## Occurrence of F42 colonization factor in *Escherichia coli* strains isolated from piglets with diarrhea<sup>1</sup>

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**ABSTRACT.** Penatti M.P.A., Silva A.S., Valadares G.F. & Leite D.S. 2005. **Occurrence of F42 colonization factor in *Escherichia coli* strains isolated from piglets with diarrhea.** *Pesquisa Veterinária Brasileira* 25(1):31-33. Depto Microbiologia e Imunologia, Instituto de Biologia, Unicamp, Campinas, SP 13081-970, Brazil. E-mail: domingos@unicamp.br

The objective of this study was to determine the presence of the colonization factor F42 in 168 strains of *Escherichia coli* isolated from diarrheic stools of newborn piglets. The presence of F42 in 12 (7.1%) strains was detected with the agglutination test. Through the Polymerase Chain Reaction (PCR) of F42 positive strains, gene encoding enterotoxins (ST-I, ST-II, LT-I and LT-II) were detected. The finding of ST-I/ST-II genes in 50% of the strains, ST-I (16%) and ST-II (25%) indicates a strong association of FC F42 with heat-stable enterotoxins (91%). In contrast, the thermolabile enterotoxin (LT-I and LT-II) genes were not detected. Serogroups of F42 positive strains were determined, serogroup O8 being the most prevalent (41,7%). Other serogroups, as there are O9, O11, O18, O32, O35, O98 and O101, were also identified. Thus, FC F42 was confirmed as an additional factor of virulence in the pathogenesis of porcine colibacillosis.

INDEX TERMS: *Escherichia coli*, F42, piglets, fimbriae, diarrhea, PCR.

**RESUMO.** [Prevalência do fator de colonização F42 em amostras de *Escherichia coli* isoladas de leitões com diarréia.] Este estudo determinou a presença do fator de colonização (FC) F42 em 168 amostras de *Escherichia coli* isoladas de suínos neonatos com diarréia. Ensaios de soroaglutinação determinaram a presença de F42 em 12 (7,1%) das amostras bacterianas. As amostras de *E. coli* F42 positivas foram estudadas através de PCR quanto à presença de genes das enterotoxinas (ST-I, ST-II, LT-I e LT-II). 50% das amostras apresentaram genes para ST-I/ST-II, 16% apresentaram genes para ST-I e 25% genes para ST-II. Não foram observados resultados positivos para as enterotoxinas termo-lábeis (LT-I e LT-II), indicando forte associação do FC F42 com as enterotoxinas termoestáveis (91%). O sorogrupo das amostras F42 positivas foi determinado, sendo o sorogrupo O8 prevalente (41,7%) e uma amostra dos sorogrupos O9, O11, O18, O32, O35, O98, e O101. Assim, o FC F42 foi confirmado como

um fator de virulência adicional na patogênese da diarréia neonatal suína.

TERMOS DE INDEXAÇÃO: *Escherichia coli*, F42, leitões, fimbria, diarréia, PCR.

### INTRODUCTION

*Escherichia coli*, which causes enterotoxic colibacillosis in suckling pigs, are referred to enterotoxigenic *E. coli* (ETEC) strains. They adhere to the microvilli of small intestinal epithelial cells and produce enterotoxins that act locally on the enterocytes (Nagy & Feket 1999). For pigs ETEC, the most common adhesins (also called colonization factors) are K88, K99 and 987P (Alexander 1994). In 1986, a new *E. coli* ETEC colonization factor (F42) isolated from feces of newborn piglet was described by Yano et al. (1986). Such strains showed mannose resistant haemagglutination (MRH) of chicken, guinea pig, sheep, horse and human erythrocytes. They also showed diffuse adherence into HeLa cells (human cervix carcinoma) and piglet enterocytes (Yano et al. 1986). Electron microscopy showed the presence of structures such as fimbriae when strains were cultured at 37°C but not at 18°C. The antiserum anti-F42 inhibited not only hemagglutination but also adherence to both HeLa cells and piglet intestinal epithelium, showing that the fimbriae were responsible for the adhesion.

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Later, Silveira et al. (1987) reported that F42, is associated with a non conjugative plasmid of 21.1 MDa coding for the heat-stable toxin (ST-I).

The purified F42 had a mass of 31 KDa and an isoelectric point of pH 3.2. Polyclonal antiserum anti-F42 obtained from the purified fimbriae, did not cross react with other haemagglutinating adhesins of pig origin (Leite et al. 1988). Castro et al. (1990) demonstrated that strains of F42 *E. coli* caused diarrhea in piglet when inoculated orally and could be detected by direct immunofluorescence in the small intestine of these animals after 42 hours.

Leite et al. (1997) showed that the hemagglutination caused by the F42 fimbriae was sensitive to the N-acetyl-galactosamine. In addition, potential receptors of F42 were shown through enzymatic treatment, to be a glycoprotein. They also verified, through Western Blotting that the reaction of F42 with membrane components varied with the source of the erythrocytes by different molecular structures.

In the present study, we report the detection of F42 among *E. coli* strains of piglets, their occurrence with the enterotoxins (ST-I and ST-II) genes and with serogroups.

## MATERIALS AND METHODS

***E. coli* strains and growth conditions.** A collection of 168 *E. coli* isolated from diarrheic feces of newborn piglets were studied. Standard strains used were: *E. coli* 567/7, O8:H<sup>+</sup>, F42<sup>+</sup> ST-I<sup>+</sup>; O101, F5<sup>+</sup>, ST-I<sup>+</sup>; 40T – LT-I<sup>+</sup>; P17 – ST-II<sup>+</sup>; PCLT-II – LT-II<sup>+</sup>; DH 5a e K12. For F42 detection the cultures were grown on Minimal Salt Medium (MM) (Davis & Mingioli 1950) containing 1.5% of Agar and 0.5% of glucose at 37° C, TSA (Trypticase Soy Agar, Difco) for serogroup test, and Luria Bertani broth (LB) for DNA amplification.

**F42 detection.** The *E. coli* strains were cultured overnight at 37° C on MM agar. The detection of F42 production was performed by slide agglutination test using specific anti-F42 rabbit antiserum (Leite et al. 1988).

**Polymerase Chain Reaction.** The strains were first screened in order to detect the presence of F42. Afterwards the positive strains in this reaction were subjected to PCR tests. Amplifications were made using 50 mg of purified DNA as template. PCR reactions were performed following the methodology described by Blanco et al. (1997) using specific primers for the genes that codify the toxins: LT-I, Blanco et al. (1997); LT-II, Pickett et al. (1986); ST-I and ST-II, Ojeniyi et al. (1994).

**Determination of the serogroups.** The serogrouping was determined using a microplate technique described by Guinée et al. (1972) and modified by Blanco et al. (1992).

## RESULTS

The slide agglutination test using rabbit polyclonal anti-serum anti-F42 identified 12 (7.1%) strains positive out of 168 isolates.

Among the 12 strains, 6 (50%) which had the genes for ST-I and ST-II, 3 (25%) were positive for ST-II, 2 (16%) for ST-I, and not any strain was positive for LT-I or LT-II in PCR. Only one was not toxigenic.

The serogroups from 12 F42<sup>+</sup> strains were determined: 5 reacted with anti-serum O8 (41.7%); antisera to O9, O11, O18, O32, O35, O98 and O101 reacted with a single isolate.

The combination of all results is shown in Table 1.

**Table 1. Enterotoxigenic genotypes and serogroups results of F42 *Escherichia coli* strains isolated of feces from diarrheic piglets**

ST-I	ST-II	Serogroup(s) (n° of strains)
+	+	O8 (3); O9(1); O11 (1); O32 (1).
+	-	O98 (1); O101 (1).
-	+	O8 (1); O18 (1); O35 (1).
-	-	O8 (1)

## DISCUSSION

In this study, the presence of the F42 colonization factor in *E. coli* strains from piglets with diarrhea, was identified. Through the slide agglutination test, the presence of F42 in 12 strains (7.1%) of the 168 strains was shown. These observations are in agreement with the previous report of Zerbini (1993) that showed the presence of F42 in 7.4% of the strains studied. The similarity of the FCs already described, the prevalence of which were observed in different studies, showed heterogeneous results in different countries. Our results suggest that FC F42 must be considered as a factor of virulence in porcine colibacillosis.

In our experiments, no amplification of primers for the thermo-labile enterotoxins (I or II) could be observed. These results are not surprising, as others reported that enterotoxin LT has not been detected, or that it has been detected in a small number of strains isolated from pigs (Woodward & Wray 1990, Flores-Abuxapqui et al. 1997, Osek 1998, Parma et al. 2000). Thus, the results let us conclude that probably FC F42 is not linked to the occurrence of this enterotoxin. On the other hand, the presence of the genes encoding ST-II and ST-I enterotoxins were found by us. The correlation between ST-I and ST-II was 50% of the total of the strains studied. In summary, all the results obtained, show the prevalence of genotypes profiles ST-I<sup>+</sup> ST-II<sup>+</sup> observed in 6 strains from serogroups O8 (3), O9 (1), O11 (1) and O32 (1). The genotype profile ST-I<sup>-</sup> ST-II<sup>+</sup> was found in 3 strains of the serogroups O8, O18 and O35, and the genotype profile ST-I<sup>+</sup> ST-II<sup>-</sup> was found in 2 strains of serogroups O98 and O101.

The correlation of ST-II enterotoxin with other colonization factors varies greatly and ranges from 0% to 78%; according to Söderlind et al. (1988) in 21%, Nagy et al. (1990) in 38%, Blanco et al. (1997) in 78%, Osek (1998) in 30%, Know et al. (1999) and according to Osek (1999) in 4.5%, but in non of the strains by Parma et al. (2000).

The serogroup O8 (41.7%) was predominant in our study, confirming the initial comment made by Yano et al. (1986). On the other hand, serogroups O9, O11, O18, O32, O35, O98 and O101 showed association with F42, what has not been described so far.

The combination of several factors was observed also for the colonization/enterotoxins in ETEC with neonatal diarrhea (Guinée & Jansen 1979, Gaastra & De Graaf 1982, Castro et al. 1984, Mainil et al. 1998). We demonstrated here the importance of the colonization factor F42 for the development of diarrhea in infections for ETEC, and the necessity of including this FC in the vaccine, in order to provide a major protection for the animals.

We think that more screening is needed, which includes variables such as age, race and breeding practices in distinct geographic regions to further clarify the distribution and association of this colonization factor.

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