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## Typing of Campylobacter jejuni isolated from poultry on the basis of flaA-RFLP by various

#### restriction enzymes

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#### **ABSTRACT**

RFLP analysis of the flagellin (*flaA*) gene was compared using three different restriction endonucleases *i.e Dde*I, *Hinf*I and *Dpn*II to determine the genetic diversity among 43 *Campylobacter jejuni* isolates of poultry origin from the same geographical area. *flaA* gene was amplified in all the isolates and RFLP analysis showed variations. *Dde*-based RFLP was found most efficient in discriminating *C. jejuni* isolates by generating 15 different *Dde*-RFLP patterns with discriminatory index (D.I) of 0.9258 whereas *Dpn*II produced seven *Dpn*-RFLP patterns (D.I.= 0.8427). While *Hinf*I enzyme produced only six *Hinf*-RFLP patterns (D.I.= 0.6977). The discrimination of *Dpn*-RFLP was comparable to discrimination given by *Dde*-RFLP analysis, which is generally used to study *flaA* gene RFLP.

KEYWORDS: C. jejuni, poultry, flaA RFLP, dpnII, ddeI

### 1. INTRODUCTION

Campylobacter species are the second most emerging bacterial zoonotic pathogen after Salmonella causing gastroenteritis (Silva et al., 2011; Epps et al., 2013). Of the many Campylobacter species identified, Campylobacter jejuni (C. jejuni) is the most predominant pathogen implicated in food borne infections followed by Campylobacter coli (Ccoli) (Coward et al., 2008; Biswas et al., 2011; Wieczorek and Osek, 2013; Bolton, 2015). Caeca of the chickens is frequently colonized by this organism and consumption of broiler meat and its products contaminated during production and processing is considered to be the most frequent source of infection (Pearson et al., 1993; Aydin et al., 2007; Wirz et al., 2010).

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