# The colonization of broilers with Campylobacter

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Poultry is a major reservoir of Campylobacter jejuni, which is currently the leading cause of acute bacterial diarrhoea in western countries. This results in substantial accumulated economic loss due clinical costs and lost working hours. In developed countries reinforcement of hygienic practices and consumer education have so far been inadequate to significantly decrease numbers of human cases of campylobacteriosis. The control of poultry associated infection in humans may also depend upon the control of Campylobacter colonization in broiler flocks. However, sources of C. jejuni and the transmission routes through which broilers become colonized are not clearly defined. A better understanding of C. jejuni epidemiology is urgently needed. This review summarizes current theories on the epidemiology of C. jejuni broiler colonization.

**Keywords:** Campylobacter; colonization; epidemiology; broilers.

## Introduction

Each species of Campylobacter has a favoured reservoir, Campylobacter enteritis is regarded as a food-borne disease because the natural habitat of Campylobacter species is the intestines of both domestic animals, e.g. dogs (Workman et al., 2005), livestock, e.g. pigs and poultry and wild animals, e.g. migratory birds (Griffiths and Park, 1990; Thomas et al., 1999a). C. jejuni is currently the leading cause of human foodborne gastroenteritis in developed countries and infection and can be followed by severe clinical complications, such as Guillain-Barré syndrome (GBS) (Boyd et al., 2005). Most campylobacteriosis cases are sporadic and a number of foods have been implicated as risk factors for sporadic Campylobacter infection, including barbecued meat, raw milk, bird pecked milk, bottled mineral water and poultry (ACMSF, 2004). Evidence for sporadic sources of human Campylobacter infections is mostly indirect (Corry and Atabay, 2001). Several studies have demonstrated the consumption of undercooked poultry, to be associated with human illness (Studahl and Andersson 2000), and it has been suggested that between 20% and

© World's Poultry Science Association 2005 World's Poultry Science Journal, Vol. 61, December 2005 Received for publication June 12, 2005 Accepted für publication June 27, 2005 40% of sporadic disease might be due to the consumption of chicken (ACMSF, 2004). While not all strains isolated from poultry are the same as those from humans, several studies, where discriminatory typing methods were used, show a close correspondence between the strains of campylobacters found in human infections and those found in chickens (Pearson *et al.*, 1993). However, consumption or occupational contact with livestock or their faeces can also actually be protective and is associated with a significant decrease in the risk of becoming ill with *Campylobacter* (Adak *et al.*, 1995). Controlling *Campylobacter* carriage in the poultry reservoir might have a measurably beneficial effect on human disease incidence (ACMSF, 2004). Despite steadily improving molecular biology identification techniques, *e.g. flu* typing; by PCR-restriction fragment length polymorphism analysis of the *flaA* and *flaB* genes, (Newell *et al.*, 2001), the epidemiology of colonization remains unclear. There is an urgent need for further research in this area to help to minimize incidences of consumer *Campylobacter* infection.

## Campylobacter and poultry

The ubiquitous nature of *Campylobacter* and its optimal growth in the gastrointestinal tract of poultry ensures that poultry frequently become heavily infected (Evans, 1992). Isolation of campylobacters from poultry carcasses range from 50-60% in the UK, Canada, Switzerland, the USA and up to 94% in Australia (Altekruse *et al.*, 1998; Shapton and Shapton, 1998; Wittwer *et al.*, 2005). In 2001, an extensive Food Safety Authority (FSA) (*Table* 1) survey found the UK national average of *Campylobacter* contamination to be 50% in finished raw chicken meat (FSA, 2001), and C. *jejuni* can also survive routine poultry processing techniques (Evans, 1992).

|                   | England | Wales | Scotland | N. Ireland |
|-------------------|---------|-------|----------|------------|
| Fresh             | 52%     | 47%   |          | 89%        |
| Frozen            | 31%     | 29%   | 35%      | 41%        |
| Number of samples | 2,481   | 800   | 800      | 800        |

Campylobacters in living broilers are found in highest numbers in the large intestine, caecum and cloaca (Shane, 2000; Corry and Atabay, 2001). When colonizing intestines, C. jejuni expresses several virulence factors, e.g. chernotaxis (Takata et al., 1992; van Vliet and Ketley, 2001). The organism is located in the mucosal film (Shane, 2000; Corry and Atabay, 2001) as it is attracted to L-fucose, a terminal sugar of the glycoprotein constituent of mucin (Shane, 2000). Heat shock proteins (HSPs) are associated with the thermal stress response of bacteria and are important virulence factors e.g. DnaJ mutants are unable to colonize chickens (Konkel et al., 1998; van Vliet and Ketley 2001). Campylobacter penetrate the mucous layer covering the intestinal cells using their polar flagella and 'cork-screw' motion (Szymanski et al., 1995). Adhesion and invasion are dependent on both motility and flagellar expression, as C. jejuni mutants with decreased motility due to paralysed flagella show reduced adhesion and no invasion, indicating that while flagella are involved in adherence, other adhesions are involved in subsequent internalization (Yao et al., 1994; van Vliet and Ketley, 2001). The outer-membrane adhesion proteins CadF and PEBI are involved in adherence and invasion. The PEBI protein is encoded by the peb1A locus, with significantly decreased adherence, invasion of HeLa cells and reduced colonization of mouse models associated with peb1A mutants

(Konkel et al., 1997; Pei et al., 1998; Ziprin et al.. 1999; van Vliet and Ketley, 2001). CadF mutants are unable to bind fibronectin and colonize newly hatched leghorn chickens (Konkel et al., 1997; Pei et al., 1998; Ziprin et al., 1999; van Vliet and Kelley, 2001). The major outer membrane constituents. lipo-oligosaccharide (LOS) and lipopolysaccharides (LPS) are involved in serum resistance, endotoxicity and adhesion (Jin et al., 2001; van Vliet and Ketley, 2001). The superoxide dimutase (SOD) protein SoaB is the main component of the C. jejuni superoxide stress defence and sodB mutants show significantly decreased intracellular survival in human embryonic intestinal (INT-407) cells (Pesci et al., 1994; Purdy and Park, 1994).

Numbers in the region of 10<sup>5</sup>-10<sup>9</sup> colony forming units (CFUs) per g of intestinal contents have been commonly observed (Corry and Atabay, 2001; Newell, 2001). Poultry faecal samples may contain high concentrations of C. jejuni cells, e.g. 107 CFUs per g (Pearson et al., 1993). Organisms may also be recovered from throughout the gastrointestinal tract, as well as the spleen and liver, indicating some systemic infection has occurred (Newell, 2001). Rapid and cross-contamination of poultry carcasses occurs during processing (Pearson et al., 1996; Corry and Atabay, 2001). Experimental challenge of chicks of all ages, appears to induce an identical outcome to that observed during natural infection, where C. jejuni act as a commensal in this host because colonization is both extensive and asymptomatic (Newell, 2001). C. jejuni can be pathogenic for young birds, but is generally not so for adult birds (Pearson et al. 1996). Newly hatched chicks can become easily colonized, without morbidity, by single oral dosage or cloacal inoculation, or by repeated low level inoculation through drinking water (Evans, 1992; Altekruse et al., 1998). For example, 1-d-old chicks orally challenged with as few as 30 CFUs of fresh isolates of C. jejuni can achieve caecal colonization with levels of up to 1 x 1010 CFUs per g of caecal contents within 3 d (Newell, 2001). The level and extensiveness of colonization in chick models varies according to the genetic lineage of the birds, the challenge strain and the degree of laboratory adaptation of the strain (Newell, 2001). Specific flocks that become infected, show a rapid rate of intra-house transmission and a high isolation rate from caecal swabs, water and litter (Hafez et al., 2001). Within colonized flocks, the organism usually spreads so rapidly that the proportion of birds colonized is often close to 100% (Pearson et al., 1993; Corry and Atabay, 2001). Coprophagy may partly explain the rapid transmission of the organism between birds, because it has been demonstrated that the virulence of invasive C. jejuni isolates is enhanced by passage through chicks (Evans, 1992). Once excreted into the environment, the organism does not multiply, due to its relatively high minimal growth temperature of >30°C (van de Giessen et al., 1996).

Most commercial chickens are colonized with C. *jejuni*, without clinical signs, around 3 weeks of age (Evans, 1992; Fields and Swerdlow, 1999; Boyd *et al.*, 2005). The reasons for this delay are unclear, but may include maternal antibodies in young chicks, as most broiler-breeder flocks are *Campylobacter-positive* and anti *-Campylobacter* antibodies may be present in egg yolks, and the presence in young birds of bacterial floras antagonistic to *Campylobacter* spp. (ACMSF, 2004). In the UK, individual broiler flocks are colonized by a limited number of subtypes of C. *jejuni* (Newell *et al.*, 2001). In some, but not all cases, the same subtypes, isolated from the ceca, contaminate the end product as observed in carcass washes (Newell *et al.*, 2001).

Following prolonged environmental stresses, *e.g.* low nutrient conditions in water, certain strains of C. *jejuni* can enter a dormant coccal form, where it is no longer possible to culture the organism using conventional media (Jones *et al.*, 1991). After campylobacters are no longer culturable, electron microscopy has demonstrated that portions of cocci do not show signs of degeneration, which is also consistent with continued viability, i.e. a viable, but not culturable state (VBNC) (Jones *et al.*, 1991). The

animal gut remains the only natural site of multiplication for thermophilic campylobacters (Jones *et al.*. 1991). The epidemiological significance of VBNC *C. jejuni is* based upon the potential of the form to resuscitate and give rise to infection (Thomas *et al.*, 1999b). However, only a limited proportion of isolates produce the VBNC form (Tholozan *et al.*, 1999), suggesting that this form is not significant in transmission to poultry flocks and it is not a universal survival mechanism.

#### Vertical transmission

The theory of vertical transmission is a controversial issue. Campvlobacter is present in the reproductive tracts and semen of poultry which could lead to vertical transmission of Campylobacter from the hen to the chick (Cole et al., 2004a; Cole et al., 2004b). However, most evidence suggests that the control of infection in breeder flocks appears to have little importance, since most researchers have found no evidence that campylobacters are transmitted vertically, or in the egg (Shane, 1992; Corry and Atabay, 2001). Low C. jejuni isolation rates from naturally or experimentally infected eggs (Shane et al., 1986; Pearson et al., 1996; Sahin et al., 2003), lack of egg penetration (Doyle, 1984; Shanker et al., 1986; Evans, 1992), no multiplication inside eggs, except in yolks and poor survival on eggs shells (Corry and Atabay, 2001) collectively suggest that vertical transmission of C. jejuni through eggs is a rare event and does not play a major role in the introduction of Campvlobacter to chicken flocks. Shane (1992) found that faecal shedders of C. jejuni did not produce infected eggs. The theory of vertical transmission may be supported by the fact that the same C. jejuni strains may be present in both breeder flocks and their progeny (Cox et al., 2002). However, this could equally be due to low-level horizontal transmission of a persistent source or reservoir external to broiler house environments, e.g. cattle (On et al., 1998; Shreeve et al., 2002). Therefore, control measures should be directed at limiting the sources of infection for broilers and reducing the rate of transmission within flocks (van de Giessen et al., 1996; Evans, 1992).

## **Horizontal transmission**

Confinement of poultry flocks is critical because many warm-blooded animals and insects may serve as vectors (Altekruse et al., 1998) and broilers readily pick up C. je juni from the environment, so there may be many potential sources of infection (Evans, 1992; Pearson et al., 1993). Flock positivity is generally higher (up to 100%) in organic and free-range flocks compared to intensively reared flocks, presumably reflecting the level of environmental exposure of such birds, as well as the increased age of the birds at slaughter (Newell and Fearnly, 2003). An elevated risk of broiler Campvlobacter infection is associated with the presence of other farm animals, e.g. cattle and sheep (Annanm-Prah and Janc 1988; Cardinale et al., 2004). Cross-contamination of Campylobacter between sheds by contaminated air, dust, insects or litter, are possibilities (Pearson et al., 1993; Whyte et al., 2001). Lack of hygiene barriers, uncemented poultry-house floors (Cardinale et al., 2004) dividing flocks into batches for staggered slaughter and down periods of less than 14 d all significantly increase Campvlobacter prevalence in broilers (Hald et al., 2000; Hald et al., 2001). Transportation of broilers prior to processing significantly increases Campylobacter colonization of the ceca and carcass contamination (Stern et al., 1995; Hald et al., 2001). This is probably because trucks, forklifts, pallets, crates, drivers' and catchers' boots are all potential sources of C. jejuni for broilers (Ramabu et al., 2004).

Wild birds can acquire Campylobacter by feeding on raw sewage and rubbish, and can

spread these agents to humans directly or by contaminating commercial poultry operations (Reed et al., 2003). The lesser mealworm, Alphitobius diaperinus (Panzer), is a carrier of Campylobacter in poultry facilities and could acquire and harbour Campylobacter from an environmental source (Strother et al., 2005). The lesser mealworm is capable of passing viable bacteria to chickens that consume the beetle, and should be included in attempts to reduce Campylobacter prominence in poultry facilities (Strother et al., 2005).

The presence of Campylobacter in the intestinal tract implicates ingestion of a contaminated source, e.g. feeds and water (Montorse et al., 1985). Neither feeds nor fresh litter seem to be likely sources of campylobacters (Pearson et al., 1993). The moisture content of commercial feed is about 8-10% (Montorse et al., 1985). Feed is dried and pelleted, may be pasteurized and is air blown into silos (Pearson et al., 1993). C. *jejuni* is very sensitive to dehydration and dies rapidly in aerosols and the litter used on farms is generally wood shavings, which are dry and resinous, being mainly softwood and normally comes directly from sawmills (Pearson et al., 1993). Spent wood shaving litter is extremely inhibitory to the growth of salmonellas (Pearson et al., 1993). Its effect on campylobacters is less well understood, but experimental work has shown that litter artificially contaminated with campylobacters can infect chickens, under laboratory conditions (Pearson et al., 1993).

## **Drinking water**

Drinking water has sometimes been found to be a significant source of infection, or a very significant risk factor (Kapperud et al., 1993; Pearson et al., 1993). There is evidence that low (10-10<sup>2</sup> per litre) numbers of campylobacters in water supplies can colonize poultry flocks (Shapton and Shapton, 1998). Although the levels of chlorination normally used for potable water would usually be considered lethal to planktonic C. jejuni, water-borne protozoa, such Tetrahymena pyrifonnis and Acanthamoeba castellanii, have strong potential to act as protective reservoirs for C. jejuni in the drinking water systems of intensively reared broilers (Snelling et al., 2005). In-vitro, experimental co-cultivation of C. jejuni with such protozoa can significantly reduce the susceptibility of the bacteria to chlorine (King et al., 1988) as well as to the industrial disinfectant Virudene (p>0.05) and can significantly (p>0.05) prolong the viability of Campylobacter at 25°C, the temperature at which intensively reared broilers are maintained (Snelling et al., 2005). These factors may partially explain observations by Stern et al. (2002) that chlorination of broiler drinking water had no effect on the C. jejuni colonization of broilers (Newell et al., 2003). Although no link between resistance to disinfectants and antibiotics has been observed (Avrain et al., 2003), water-borne protozoa could putatively act as protective delivery vehicles for Campylobacter (Snelling et al., 2005) with antiobiotic resistance genes, which could then be exchanged between strains (Wittwer et al., 2005).

Despite these new findings, overall evidence suggests that C. jejuni-contaminated water constitutes a low risk of colonization for broiler flocks (Newell and Fearnly, 2003). Further research is required to establish the role of water line and reservoir contamination in flock colonization and the value of water sanitizers as control measures (Newell and Feamley, 2003).

#### Conclusion

In several on-farm studies where horizontal transmission was suspected, and where epidemiological typing of *Campylobacter* isolates were performed, the exact source of

Campylobacter infection in broilers were not established (Jacobs-Reitsma et al., 1995; van de Giessen et al., 1996). In theory, problems like these should be addressed with rapidly evolving molecular biology methods to provide more sensitive Campylobacter detection methods and typing methods with more accuracy. However, because Campylobacter spp. are ubiquitous in the environment, cases are mostly sporadic, outbreaks are rare and due to the lack of global protocol standardization, source tracing can be difficult (Wassenaar and Newell, 2000). It is extremely difficult to identify the most suitable Campylobacter genotyping method, e.g. pulsed-field gel electrophoresis (PFGE), that fulfils all of the requirements for molecular epidemiological investigations (Wassenaar and Newell, 2000). However, these issues are now starting to be addressed by international consortia, e.g. CAMPYNET (Wassenaar and Newell, 2000) and PulseNet, to try to attain standard methods for strain comparison. This should help to give a greater insight into both the population structure and distribution of Campylobacter over the next few years.

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