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سورة الإسراء الآية رقم ٨٥

**SHEDDING AND COLONIZATION OF CAMPYLOBACTER COLI
IN BROILERS FROM DAY OF HATCH TO SLAUGHTER**

A Thesis

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تواجد ميكروب الكامبيلو باكتر كولاي فى بدارى التسمين وخروجه منها من يوم الفقس حتى الذبح

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I-INTRODUCTION

Campylobacter coli and *Campylobacter jejuni* have been recognized since that date 1970s as important agents of gastrointestinal infections through out the world, in the United states these infections affected approximately 1% of the population each year (**Tauxe et al., 1992**).

In many industrialized countries the incidence of Campylobacteriosis exceeds Salmonellosis. *Campylobacter* bacteria are transmitted to human mainly through food especially poultry meat products (**Rivoal et al., 2005**).

Campylobacters are carried in the intestinal tract of a wide variety of wild and domestic animals especially birds. They can establish a temporary symptomatic carrier state as well as illness in human. Consumption of food and water contaminated with untreated animal or human wastes accounts for 70 % of *Campylobacter* related illness as each year.

The food includes unpasteurized milk, meats, poultry, shellfish, fruits and vegetables (**Abeyta and Kaysner, 1987 and Abeyta, 1998**).

Once **Campylobacter** colonization occurs in birds within a flock, ceacal can colonized within 72 hours (**Newell , et al . 1983**) , reach high numbers in ceacal contents as high as 10^9 cfu in experimentally challenged birds (**Wassenaar, et al . 1993**). This level may also increased during evisceration of carcasses , washing and processing due to contamination by personnel (**Oosterom , et al . 1983**) . Several serotypes were identified in one flock , some of these were not found again on further examination of the same flocks during the growing period (**Pokamunski . et al 1986**) . However the horizontal transmission is generally the most significant cause of broiler flock (**Pattison , 2001**) varied consider with season and region (**Hofshagen , 2005**) .

Campylobacter is one of the most common causes of bacterial gastroenteritis in humans world and yet is still a poorly understood bacterial pathogen, little is known about the prevalence of different virulence factors and the ability to produce toxin among *Campylobacter* isolates obtained from different sources (**Bang et al., 2003**).

The presence of **Campylobacter** Spp. on broiler carcasses and in closed tank water in a commercial poultry processing facility was monitored at monthly intervals from July to December (**Hinton et al., 2004**).

Species within the genus **Campylobacter** have emerged over the last three decades as significant clinical pathogens particularly of human public health concern where the majority of acute bacterial; enteritis in the western world is due to those organisms of particularly concern ,the species *C. jejuni* and *C. coli* which are responsible for most of these gastrointestinal related infections. Although these organisms have already emerged as causative agents of zoonoses. (**Moore et al., 2005**). Yet , there is no medium available to allow growth of *C. jejuni* and inhibit *C. coli* or vice versa (**Martin et al ., 2002**)

Enteritis due to **Campylobacter** is the most common cause of acute bacterial diarrhoea worldwide. In most cases infection occurs as a result of consuming contaminated water or food especially raw meat of fowls. (Seyed et al., 2005).

There are no validated serological tests developed for the identification of infected mammals or birds (Cawthraw et al ., 1994). Due to some technical difficulties such as long incubation time uncertainly in results and some atypical strains which are difficult to culture , the detection of campylobacters is more complicated and genome based detection methods like PCR have gained a preminent importance to compare the results with other data (seyed et al ., 2005)

The present work was designed to cover the following points.

- 1-Incidence of Campylobacter coli infection in broiler chickens from day of hatch to slaughter.
- 2-Identification of isolates of Campylobacter coli by bacteriological methods.
- 3 _ Comparison of PCR and conventional laboratory methods for diagnosis of Enteropathogenic Campylobacter spp. in Fowl faeces .

II-REVIEW OF LITERATURE

II-1.Incidence of Campylobacter:-

Oosterom et al. (1983) believed that, the source of bacterial contamination of poultry meat is essentially with intestine or gut content which may come in contact with carcasses already in the broiler house and during transport and slaughter either directly or indirectly through a vehicle such as transport and processing equipment. High levels of bacterial cross contamination may occur especially during defeathering and water chilling with intestinal contamination apparently being the only source. However, these level may also increased during evisceration of carcasses washing and processing due to contamination by personnel.

Milakovic-Novak et al. (1984) isolated *C. Jejuni* from the small intestines of 12 (6.6 %) out of 180 and the large intestines 12 (2.2 %) out of the 558 broilers that had died and from the cloaca of 28 (24 %) of 122 healthy laying hens and 47 (40 % of 118 hens with clocitis (vent gleet). Of the 29 isolates that were types 8 were *C. jejuni* biotype 1 and 21 were *C. coli*.

Neill et al. (1984) showed that *C. Jejuni* was not usually present in intestines of broiler chicks less than two weeks old but they were present in most older birds.

Pokamanski et al. (1985) examined six flocks for *Campylobacter* from hatching to slaughter. A total of 1530 cloacal swabs and ceacum samples were collected from 768 chicks at one-day and 1, 4 and 8 weeks old *Campylobacter* was not isolated from 360 chicks (healthy condemned unhatched) at hatching only one flock was negative all samples of 172 isolates 147 were *C. Jejuni* and 25 were *C. coli*.

Pokamanski et al. (1986) examined 1440 samples. From 720 broilers (From 5 flocks) for *C. jejuni*. *Campylobacter* species were not isolated from day old chicks and were only isolated from one week old chicks in one flock. In three flocks *Campylobacter* species were isolated from all chicks sampled at 4 weeks of age. In the 4th flock all chicks sampled were negative until 8 weeks of age when all were positive. The 5th flock remained negative through the 8 weeks of its life eleven broiler flocks were examined only at slaughter, 24 caecal samples were examined from each flock. In three flocks *Campylobacter* was not isolated in one flock one broiler chick was positive and in 7 flocks (From 58 to 100 %) of sampled broiler chickens were positive of the 146 isolates typed, 123 were *C. Jejuni* and 23 were *C. coli*. The authors concluded that in some flocks , several serotypes were identified, some of these were not found again on further were identified, some of these were not found again on further examinations of the same flocks during the growing period.

Varga et al. (1986) isolated 62 *Campylobacter* strains from hens with hepatitis from the cecum of clinically healthy hen and from the surface of slaughtered chickens. About 5 % had diarrhea but the mean mortality rate did not exceed 2.3 %. Sixty strains were identified as *C. Jejuni* one as *C. coli*. Most *C. jejuni* strains had typical growth and biochemical characteristics.

Acevedo et al. (1987) cultured samples of intestine, gall bladder and liver from 161 chickens of different ages, 85 with diarrhea and 75 apparently healthy yielded *Campylobacter* from 43 (26 %). Most isolates were from the small intestine and cecum very few from liver and gall bladder. *C. jejuni* was isolated more often than *Campylobacter coli* and was of the same serotypes as those known to occur in man. Isolations were obtained from chickens older than 3 weeks but not from younger ones. There was no relationship between the presence of *Campylobacter* and diarrhoea.

Hoope and Ehram (1987) examined a large poultry farming organization over 18 months for *C. jejuni* and *C. coli*. Incidence of infection ranged 4 to 78 %. In 4 of 34 broiler units (12 %). *Campylobacter* shedders were found in 6 of 153 flocks (4 %) first isolations occurring between the 34th and 42nd of fattening 16 % of 625. Skin samples from birds prepared for sale were positive. It was concluded that *Campylobacter* was introduced into broiler flocks as a result of inadequate protection and that the slaughter process was responsible for contamination of poultry.

Torre Manas et al. (1987) studied the incidence of *Campylobacter* in the cloacal contents of broiler chickens. The study revealed three species, *C. jejuni* (37.24 %) *C. coli* (32.28 %) and thermophilic nalidixic acid resistance, *Campylobacter* (10.93%).

Kaino et al. (1988) showed that the intestines of poultry are easily colonized with *C. coli* and *C. jejuni*. Day old chickens can be colonized with a few 35 organisms. Most chickens in commercial operation are colonized by 4 weeks (**Humphrey et al., 1993**)

Kapperud et al. (1993) indicated that, vertical transmission from breeder flocks to progeny has been suggested in one study but is not widely accepted (**Pearson et al., 1996**).

Jacobs-Reitsma et al. (1995) found that, the vertical transmission of *Campylobacter* to flocks via contaminated egg breeder hens are usually colonized by multiple strains of *C. jejuni*. The recovered from various segments of the reproduction tract.

Stern et al. (1995) investigated the colonization of the ceca and the contamination of chicken carcasses by *Campylobacter* species. Samples were taken on the farm and after transport and handling. In the first set of experiments, 20 chickens obtained from each of 10 broiler farms were collected from houses containing 6 to 7 week old birds. Half of the birds were slaughtered at the farm, the other birds were slaughtered at the farm, the other half was transported (10 birds per chicken coop) to a holding facility and killed within 16 to 18 h. Caeca from birds in 9 of the 10 farms. Samples were positive for *Campylobacter* species. Colonization level ranged from 10 (4.11) to 10 (2.28) CFU *Campylobacter* species / g caecal matter except on one farm, where the organism was not isolated. The mean count on the farm was 10 (9.44) CFU *Campylobacter* species /g caecal material and after transport the mean was 10 (6.15) CFU/g.

Berndtson et al. (1996) showed that, the prevalence of flock positively is also depend on flock size and the system. Flock positively is generally higher up to 100. In organic and free compared to intensively reared flocks. This presumably reflects the level of such birds as well as the increased age of the birds at slaughter.

Cawthraw et al. (1996) explained that, *C. jejuni* close relative *C. coli*. In chickens *C. jejuni* colonizes the mucus overlying the epithelial cells primarily in the ceca and the small intestine but may also recovered from elsewhere in the gut and from the spleen and liver. Experimentally the dose of viable *C. jejuni* required to colonize chicks and chickens can be as low as 40 CFU. However this dose and the kinetics of colonization may be dependent on both the bacterial strain and the chickens strain. Once colonization is established *Campylobacter*s can rapidly reach extremely higher numbers in the caecal contents, as high as 10^9 CFU experimentally challenged birds (**Wassenarr et al., 1993**), although this level may be lower in naturally colonized birds.

Chuma et al. (1997) found the incidence of *C. jejuni* and *C. coli* in broiler farms was 33.9% (19/56) *C. jejuni* positive flock accounted for 20.0 % (17/85) and *C. coli* positive ones was 4.7 % (4/85).

Gregory et al. (1997) found that, *Campylobacter* most establish at transient low level of infection but more typically colonization of the broilers occurs between days 14 and 49 of the grow out period with almost complete. Flock involvement by day 49. Whether the broiler house was new or had previously housed broilers influenced neither the time nor rate of colonization of the flock. Although various species of wild life around the grow-out houses were found to be positive for *Campylobacter*. There was no correlation between this and the onset of flock colonization.

In Denmark, **Nielsen et al. (1997)** examined 929 cloacal swabs between (1995 – 1996) and found that 36 % were positive for thermophilic *Campylobacter*.

Willis and Murray (1997) examined seasonal variation of the level of *Campylobacter* contamination of broiler chickens . *Campylobacter* have been reported to exhibit a cyclical pattern of contamination of poultry. We release the level of contamination consistently increased and decreased depending on the season of the year.

Andrews (1998) found that *Campylobacter* species are intestinal commensals of many different wild animals, domestic livestock and pets to include cow, cattle, poultry (Turkeys and chickens), sheep , dogs and cats. The organism regularly shed in the feces of these animals and consequently has associated with inadequately pasteurized milk and improperly handled, cooked meat.

Byrd et al. (1998) evaluated the effect of pre-slaughter feed withdrawal on the incidence of *Campylobacter* isolation from the crop was determined immediately before and after feed withdrawal in 7 weeks old broiler chickens obtained from each of nine separate broiler house. Ceca were collected from broilers in six of the same flocks for comparison with the crop samples. Feed withdrawal caused a significant ($P < 0.025$) increase in *Campylobacter* positive crop samples in seven of the nine houses sampled. A total number of *Campylobacter* positive crops increased significantly ($P < 0.001$) from 90/360 (25 %) before feed removal to 224 / 359 (62.4%) after the feed withdrawal period.

Achen et al. (1998) mentioned that at 43 days old only 37.5 % of birds were shedding *C. jejuni* in their faeces, and enumeration of *C. jejuni* in the crop, jejunum and ceacum on day

43 revealed the cecum was the major colonization site and out of 24 birds carried *C. jejuni* in their intestinal tract.

Colin et al. (1999) examined the colonization of the ceca and organ invasion by different isolates of *C. jejuni* were investigated in day-of-hatch leghorn chicks. This model of *Campylobacter* colonization of the ceca demonstrates that (**Hirn and Aho, 1988**) day-of hatch birds don't naturally contain cecal *Campylobacter* ceca can be colonized with *C. jejuni* by oral gavage and not by cloacal inoculation. *C. jejuni* can be recovered from the ceca up until at least 7 days post-inoculation (**Berndtson et al., 1996**) ceacal colonization when as little as 10^2 colony forming units is orally inoculated into chicks (**Demming et al., 1987**). Different *C. jejuni* isolates via both in their ability to colonize the ceca and in their ability to invade the liver.

Pattison (2001) considered that the horizontal transmission is generally the most significant cause of broiler flocks. *Campylobacter* are ubiquitous in the environment and could the house by a number of vehicles including human activity associated with management. However, this has yet to be proven by reproductive observation genotyping of strains in the environment with subsequently result in flock.

Richard et al. (2001) examined ten genotypically strains of *Campylobacter* strains *coli* were isolated from a swine production facility. These porcine isolates were then orally inoculated into day-of hatching horn chicks and were excellent colonizers of the chick cecum. *C. coli* recovered from inoculated chickens were genotypically identical to the challenge strain. The absence of the host specificity suggests a possible movement of strains among swine, field animals, birds and poultry houses.

Stern et al. (2001) found the proportion of broiler flocks colonize with *Campylobacter* varies, this variation may reflect at least in different sampling and indicated that nearly 90% of flock. In Europe this prevalence varies from 18 to 90 % with the Northern most countries lower figure than Southern most countries. In Sweden for National survey indicates the flock positively is less than 100 %. The reason for this is unknown. Zootechnical parameters including the climatic conditions and distance between farms may all influence flock pre possible that. The poultry industries in these northern European countries are there fore exploit hewer facilities and more closely regulated thus elsewhere is seen whether similar trends occur in other regions of the world.

Ringoir and Korolik (2002) detected the colonization and shedding in poultry has not been fully accepted that colonization in chickens persists at least for the life span of a cycle reared birds. This is usually less than 47 days , however in experimentally persistence of colonization may vary among *Campylobacter* strain.

Newell and Fearnley (2003) indicated the epidemiological investigations of commercial flocks the flock *Campylobacter* colonization is age dependent. Newly hatched chicks appear to be free. Europe this negatively persists until at least 10 days of age become infected only 2 to 3 weeks after the placement of chicks into a broiler colonized chicks into a broiler colonized. Chickens usually show no absorbable clinical symptoms of infected chicks are exposed to high doses under experimental condition.

In Europe (**Takkinen and Ammon, 2003**) studied the prevalence of *Campylobacter* species in poultry and cattle has been shown to be *C. jejuni* in swine, *C. coli* and in dog *C. upsalinensis*. Most human infections (90 – 95 %) are due to *C. jejuni*. Broiler flocks are after contaminated with *C. jejuni* and *C. coli* there is strong evidence for frequent cross contamination during slaughtering and product processing.

Hofshagen (2005) examined 10803 flocks from 562 broiler farms were tested altogether 521 (4.8 %) of the flocks were identified as positive for *Campylobacter* Spp., primarily *C. Jejuni*. The positive flocks originated from 257 (25.7%) of the farm . During the period 2002 to 2004 there was a large and steady reduction in flock prevalence. From 6.3 % in 2002 to 3.3 in 2004. the proportion of flocks positive for *Campylobacter* Spp. varied consider with season and region. The proportion of farms producing flocks positive for *Campylobacter* Spp. each year reduced substantially from 28.4 % in 2002 to 17.8 % in 2004.

Keller et al. (2005) examined samples for *C. coli* and *C. jejuni* were derived prior to slaughter from 100 randomly selected flocks (Five birds per flock) raised on three different farm types. The observed flock prevalence was 54 %., in total with 50 % for conventional and 69 % for free – range farms. Birds held on farms with a confined roaming area had the lowest prevalence of 37 % *Campylobacter* isolates were characterized by amplified fragment length Polymorphism (AFLP).

Ring et al. (2005) examined ten conventional and four extensive out door broiler flocks distributed over nine farms were investigated twice per week during a 35 – 58 days rearing period to observe the dynamics of *Campylobacter* Spp. Spread within these flocks. Spread within these flocks. A total of 4112 samples were collected 157 (3.8 %) of these samples were *Campylobacter* positive with all *C. jejuni*. The positive samples were distributed over three conventional and two extensive out door flocks on five farms. These five positive flock farms were colonized the fifth to the seventh week of age and remained colonized until slaughter. The data revealed to a single source from environment may have been responsible for the colonization of each flock.

Tesfaye et al. (2005) determined the prevalence of thermophilic *Campylobacter* Spp. in various food animals in Jimma Zone South West Ethiopia by studied urban and rural farm animal setting in Jimma collected. Fecal specimens from 485 various food animal. Cattle n = 205, Poultry n = 1991, Pigs n = 18, Sheep n = 71 and culture using standard method. The results *Campylobacter* Spp. were isolated from 192 (39.8 %) out of 485 fecal specimens. The highest isolation rate was recorded among chickens (68.1 %), followed by pigs (50.0 %), sheep (38. %) and cattle (12.7 %). Among the 192 thermophilic *Campylobacter* isolated 135 (70.3 %) were identified to be *C. jejuni*, 51 (26.6 %) were *C. coli* and 6 (3.1 %), *C. lari* *C. jejuni* was the most prevalent species in chickens (80.8 %), followed by sheep (59.3 %) and cattle (53.8 %). All isolates found in pigs were identified to be *C. coli* (100 %).

Wesley et al. (2005) said that, the *Campylobacter* is a major bacterial foodborne pathogen with consume contaminated poultry regarded as a major risk factor for human infections. The stress is feed withdrawal catching-crating transport and holding at the abattoir prior to slaughter been shown to alter the levels of *Campylobacter*. They reported

that 6 – 95 % of market *Campylobacter* as determined by cloacal swabs of birds both prior to transport at the abattoir the population of *C. coli* increased after transport the five trials. They also noted that the population of *C. coli* were more diverse than those of based on DNA profiles.

Rasschaert (2006) examined the relation between internal carriage and surface contamination with thermophilic *Campylobacter* species in broiler chickens. Samples were collected from 39 flocks in three Belgian poultry slaughter houses from each flock crop swabs before slaughtering and intestines and neck skin during slaughter were collected isolates were identified as *C. jejuni* (90 %), *C. coli* (8.37%) and *C. lari* (2.2 %). Seventy two percent of the flocks arriving at the abattoir were colonized with *Campylobacter*s after slaughter 79 % of the flocks had contaminated neck skin. Four of these flocks were initially free of *Campylobacter*. These four flocks may have no contaminated carcasses after logistic slaughtering.

Hansson (2007) analyzed and identified sources and risk factors for colonization of *Campylobacter* Spp. in broilers at both farm level and 5 slaughter groups with a low within group prevalence were identified split slaughter was confirmed as a risk factor and contamination of carcasses was shown to occur both during transport and during slaughter process. Environmental *Campylobacter* load was comparable on high and low incidence. Forms indicating that hygiene regime are of greater importance than environmental load.

Sarah and Laura (2008) detected the prevalence of infection in broiler breeder flocks has been found to be as high as 80 % but *Campylobacter*s are isolated from hatcheries or newly hatched chicks at the prevalence reported in broiler flock varies according to age and isolation technique or reason. The prevalence reported in broiler flocks also varies between countries Sweden, Finland and Norway report relatively low rates infection (5 – 20 %) whereas the UK, other European countries and the U.S.A appear to have higher levels of infection with up to 90 % of broiler flocks infected. *C. jejuni* is the most frequently isolated species from poultry but occasionally *C. coli* and *C. lari* are found.

2-Pathology and pathogenicity of *Campylobacter* in broiler chickens:

Newell et al. (1983) found the poultry is to be colonized primarily with *C. jejuni* (65 – 95 %) less often with *C. coli* and rarely with other species. Colonization rates in broiler chickens are age related. Most flocks are negative until 2 – 3 weekly of age. Once *Campylobacter* colonization occurs in a broiler flock, transmission via coprophagy, is extremely rapid and up to 100 % of birds within a flock can become colonized within 72 hours. Samples from liver birds, destined for the food chain, should therefore be taken as close to slaughter as possible the majority of birds shed large numbers of organisms ($> 10^6$ colony-forming units/g faeces). *Campylobacter*s can be isolated from fresh faeces / caecal droppings or cloacal swabs.

Establishment of *C. jejuni* in the intestinal tract results in extensive fecal excretion (**Welkos, 1984**). Found that about 83 % of the birds excreted organisms within 83 % of the bird's excreted organisms within 24 h of challenge. However, all birds did not excrete organisms, formally on all sampling days a cyclical or intermittent fecal shedding pattern was observed similar observation were made in the study of (**Achen et al., 1998**). The