



# Prevalence of Zoonotic Species of *Campylobacter* in Broiler Chicken and Humans in Assiut Governorate, Egypt



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## Abstract

The lack of a national *campylobacteriosis* surveillance system in Egypt warrants the need for periodical disease evaluation. This study aimed to determine *campylobacteriosis* prevalence in broiler chicken farms and slaughterhouses as well as in humans at Assiut Governorate, Egypt. Biochemical and molecular reactions have been employed to determine *Campylobacteriosis* prevalence. Additionally, the predisposing factors for human *campylobacteriosis* were assessed. The overall prevalence of *Campylobacter* species detected by biochemical reactions and multiplex PCR assay was found to be 23.51% and 22.46%, respectively. Prevalence rates of 16.83%, 24% and 27.55% were recorded for the incorporated samples from broiler farms, slaughterhouses and humans, respectively. None of the analyzed demographic determinants (age, gender and residence) significantly affected *Campylobacteriosis* prevalence in humans. Interestingly, mixed infection was the predominate finding among positive samples and none was confirmed to have *C. jejuni* alone. Additionally, *Arcobacter* was recovered either independently or coexisted with *Campylobacter* in poultry samples; nevertheless, zero detected from humans. This study confirms that broilers could represent an important public health threat to Assiut Governorate residents. Accordingly, measures on how to minimize the contamination level at farms, slaughterhouses and during culinary practices should be disseminated to farm workers, slaughterhouse men and consumers.

**Keywords:** *Campylobacteriosis*; Multiplex PCR; Zoonosis; Poultry; Egypt

## Introduction

Among the zoonotic *Campylobacter* species, e.g. *C. jejuni*, *C. coli*, *C. lari* and *C. upsaliensis*, the former two species are responsible for the vast majority of the human food borne infections, accounting for 90% and 5-10% of cases [1], respectively. Generally, *Campylobacter* is the most common bacterium inducing gastroenteritis in human beings globally and can be resulting in a fatal consequence in very young children, geriatric populations and immune compromised patients [2]. Being harbor *Campylobacter* in their intestinal flora, poultry and subsequently their products are considered a common and main source of the bacterium to human [3]. This mainly occurs as a result of extensive contamination of broiler carcasses and different organs with the gut contents following mechanical evisceration of birds [4]. Despite the public health burden associated with *campylobacteriosis* and the well-known role of poultry in its transmission cycle, the disease trend is still difficult to be accurately estimated in developing countries given the lack of a national surveillance system for many zoonotic diseases [5]. The

situation is aggravated with the presence of funding constraints in such countries, hence, targeted epidemiological studies could be the only resort to fill the gap. Therefore, this study was undertaken to reevaluate the prevalence of *C. jejuni* and *C. coli* in poultry in farms and slaughterhouses at Assiut Governorate, Egypt. Also, the prevalence and risk factors associated with *campylobacteriosis* caused by the two species of interest in patients attending Assiut University Hospital were studied.

## Materials and Methods

### Samples

A total of 285 samples, including 205 poultry samples and 80 human samples were recruited to this study. The two hundred and five poultry samples were collected from four poultry farms (101 cloacal swabs) and two poultry slaughterhouses (104; 26 each for intestinal swab, liver, neck and wing) located at Assiut Governorate, Egypt. However, human samples included 60 children and 20 adults

were collected from patients attending Assiut University Hospital. The age, gender, residence and clinical summary of the sampled humans were included and analyzed.

### Isolation and biochemical identification of *Campylobacter*

Samples were enriched in Bolton selective enrichment broth (Oxoid) for 24 hours at 10% Co<sub>2</sub> under 42 °C and were then cultured on modified *Campylobacter* selective agar base Cefoperazone Charcoal Desoxycholate Agar, mCCDA (Oxoid) at 42 °C for 48 hr under microaerophilic conditions. Typical colonies were biochemically identified utilizing catalase, oxidase, Gram's stain, hippurate test, sensitivity to nalidixic acid and cephalothin [3].

### Molecular confirmation of *Campylobacter* isolates

DNA was extracted from overnight cultures of *Campylobacter* using Patho Gene-spin DNA/RNA Extraction Kit (Rx Biosciences, USA). Three pairs of specific primers were chosen [6] and purchased from metabion international AG, Germany. They were selected for amplification of a 650bp of *Campylobacter* 23s rRNA gene (F: tataccggtaaggagtgtggag and R: atcaattaaccttcgagaccg), a 323bp of *C.jejuni* hipo gene (F: acttcttattgcttgctgc and R: gccacaacaagtaaagaagc) and a 126bp of *C.coli glyA* gene (F:

### Results

**Table 1:** Identification of *Campylobacter* spp. by biochemical & molecular techniques

Source & type of sample		No. Examined	Biochemical Tests			Multiplex PCR Assay			
			<i>Campylobacter</i> species	<i>C. jejuni</i>	<i>C. coli</i>	<i>Campylobacter</i> species	<i>C. jejuni</i>	<i>C. coli</i>	Mixed Infection
			Positive No. (%)	Positive No. (%)	Positive No. (%)	Positive No. (%)	Positive No. (%)	Positive No. (%)	Positive No. (%)
Poultry farms	Cloacal swab	101	20 (19.80%)	11(10.89)	9 (8.91)	17 (16.83)	0	4 (3.96)	13 (12.87)
Poultry slaughterhouses	Intestinal swab	26	5 (19.23)	4 (15.38)	1 (3.85)	5 (19.23)	0	0	5 (19.23)
	Liver	26	5 (19.23)	3 (11.54)	2 (7.69)	5 (19.23)	0	0	5 (19.23)
	Neck	26	7 (26.92)	3 (11.54)	4 (15.38)	7 (26.92)	0	2 (7.69)	5 (19.23)
	Wing	26	8 (30.77)	4 (15.38)	4 (15.38)	8 (30.77)	0	2 (7.69)	6 (23.08)
Assiut Uni. Hospital	Human stool	80	22 (27.5)	16 (20)	6 (7.5)	22 (27.5)	0	4 (5)	18 (22.5)
Total		285	67 (23.51)	41 (14.39)	26 (9.12)	64 (22.46)	0	12 (4.21)	52 (18.25)

The overall prevalence of *Campylobacter* species detected by biochemical reactions was found to be 23.51% (67 of 285). Only three isolates were found to be negative by multiplex assay. None of the confirmed isolates were found to be *C.jejuni*; nevertheless, *C.coli* was confirmed in 12 isolates and the rest of isolates were matching a mixed infection of *C. jejuni* and *C. coli* (Table 1). *Arcobacter*

gtaaaccacaaagcttatcgtg and R: tccagcaatgtgtgcaatg). The assay was performed using Qiagen Multiplex PRR Kit in Veriti Thermal Cycler (applied Biosystems) and cycling conditions were applied as recently published [7]. Gel electrophoresis was conducted for 1 hour at 100 volt and products were visualized by UV transilluminator and photographed by the gel documentation system (Biometra, Göttingen, Germany).

### Statistical analysis

Statistical differences were determined using Fisher's exact test and odds ratio in the Graphpad prism 5 (Graph Pad Software, Inc., La Jolla, CA, USA). P value ≤ 0.05 was considered significant.

### Ethical statement

Informed written consent was obtained from each adult participant and parental consent was attained prior to the participation of children in this study. All participants were informed that the obtained information will be strictly confidential. Animal blood samples were collected following agreement of the owners and animals were handled according to the Assiut University regulatory rules for animal research. The animal and human work were approved by the ethics committee of Assiut University, Egypt.

prevalence in farms and slaughterhouses was also reported in this study. *Arcobacter* alone or combined with *Campylobacter* was found in 30.24% and 6.83%, respectively, out of 205 examined poultry samples. Unlike poultry, we failed to isolate *Arcobacter* from the examined human samples (Table S1).

**Table S1:** Co-existence of *Arcobacter* and *Campylobacter* in poultry samples

Source of Sample	No. Examined	Arcobacter Only		Campylobacter + Arcobacter	
		No. (%)	Odds ratio (95% CI)	No. (%)	Odds ratio (95% CI)
Farms					
A	23	12	11.45 (2.166 to 60.58)	4 (17.39)	1.47 (0.3276 to 6.629)
B	23	2	Ref.	0 (0)	0.134 (0.006892 to 2.635)
C	23	10 (43.48)	8.07 (1.522 to 42.85)	0 (0)	0.134 (0.006892 to 2.635)
D	32	28	73.50 (12.28 to 440.1)	4 (12.5)	Ref.
Total	101	52 (51.48)	P < 0.0001	8	P = 0.0525
Slaughter houses					
A	52	2	0.22 (0.04433 to 1.092)	0 (0)	0.06813 (0.003733 to 1.243)
B	52	8	4.54 (0.9159 to 22.56)	6 (11.54)	14.68 (0.8043 to 267.8)
Total	104	10 (9.62)	P = 0.0923	6	P = 0.0268
Overall total	205	62 (30.24)	-	14 (6.83)	-

Fisher's exact test	Arcobacter only	Campylobacter + Arcobacter
	Odds ratio (95% CI)	Odds ratio (95% CI)
Farms	9.98 (4.665 to 21.33)	1.40 (0.4695 to 4.205)
Slaughterhouses	0.10 (0.04688 to 0.2143)	0.71 (0.2378 to 2.130)
P value	< 0.0001	0.59

**Table 2:** Distribution of *Campylobacter species* in different poultry farms and slaughterhouses

Source & type of samples		No. examined	<i>Campylobacter species</i>		<i>C.coli</i>	Mixed Infection
			No. (%)	Odds ratio (95%, CI)		
Farms						
A	Cloacal swab	23	6 (26.09)	7.76 (0.85-70.79)	1 (4.34)	5 (21.74)
B		23	6 (26.09)	7.76 (0.85-70.79)	2 (8.69)	4 (17.39)
C		23	1 (4.34)	Ref.	1 (4.34)	0 (0.00)
D		32	4 (12.)	3.14 (0.33-30.18)	0 (0.00)	4 (12.5)
Total		101	17 (16.83)		4 (3.96)	13 (12.87)
Slaughterhouses						
	Intestinal swab	13	2 (15.38)	Ref.	0 (0.00)	2 (15.38)
A	Liver	13	3 (23.07)	1.650 (0.2269 to 12.00)	0 (0.00)	3 (23.07)
	Neck	13	4 (30.77)	2.44 (0.3609 to 16.55)	1 (7.69)	3 (23.07)
	Wing	13	2 (15.38)	1.000 (0.1187 to 8.425)	1 (7.69)	1 (7.69)
	Total	52	11 (21.15)	0.7282 (0.2946 to 1.800)	2 (3.84)	9 (17.30)
	Intestinal swab	13	3 (23.08)	1.650 (0.2269 to 12.00)	0 (0.00)	3 (23.08)
B	Liver	13	2 (15.38)	Ref.	0 (0.00)	2 (15.38)

	Neck	13	3 (23.08)	1.650 (0.2269 to 12.00)	1 (7.69)	2 (15.38)
	Wing	13	6 (46.15)	4.714 (0.7337 to 30.29)	1 (7.69)	5 (38.46)
	Total	52	14 (26.92)	1.373 (0.5556 to 3.394)	2 (3.84)	12 (23.07)
Overall Total		104	25 (24)	-	4 (3.84)	21 (20.19)

**Table 3:** Simultaneous detection of *Campylobacter* species in different examined parts of positive chicken carcasses (n=15) a: Mixed infection, b: *C. coli*

Carcass	No of <i>Campylobacter</i> spp Isolates	Type of Isolates		Origin of Isolates			
		<i>C.coli</i>	Mixed infection	Intestine	Liver	Neck	Wing
1	1	-	1	-	-	a	-
2	1	-	1	-	-	-	a
3	1	-	1	-	a	-	-
4	4	1	3	a	a	a	b
5	2	-	2	a	a	-	-
6	1	-	1	-	-	a	-
7	1	1	-	-	-	b	-
8	2	1	1	-	-	a	b
9	2	1	1	-	-	b	a
10	2	-	2	-	-	a	a
11	1	-	1	-	a	-	-
12	2	-	2	a	a	-	-
13	1	-	1	-	-	-	a
14	2	-	2	a	-	-	a
15	2	-	2	a	-	-	a
Total	25	4	21	5	5	7	8

The prevalence rates of *Campylobacter* spp. in sampled chicken from farms and slaughterhouses were 16.83% and 24%, respectively. Farm A similar to farm B had the highest prevalence rate and slaughterhouse B had a slightly higher prevalence than slaughterhouse A (Table 2). All sampled chicken carcasses, but no. 7 had a mixed infection at least in one organ. However, neither the intestinal swabs nor livers sampled from slaughterhouses had *C. coli* contamination. Additionally, two wings and two livers were found to have the species of interest (Table 3).

(Table 4) revealed that the prevalence of *Campylobacter* infection was higher in adults (30%) than children (26.66%) and none of the analyzed variables had significant impact on getting *campylobacteriosis* in humans. As illustrated in Table S2, fever was documented in 87.5% of infected children, however, all had diarrhea and abdominal pain. Additionally, blood was seen in 25% of stool samples of infected children and surprisingly, our study reported the presence of an un expectedly high percentage (87.5%) of infected children manifesting emesis.

**Table 4:** Prevalence and risk factors of *Campylobacter* infection in humans

Variable	No. examined	<i>Campylobacter</i> species		<i>C. coli</i>	Mixed infection
		No. (%)	Odds ratio (95% CI)	No. (%)	No. (%)
Children					
Age (Year)					
≤ 1	33	10 (30.3)	1.478 (0.4263 to 5.126)	2 (6.06)	8 (24.24)
1.1-2	22	5 (22.72)	Ref.	2 (9.09)	3 (13.64)
>2	4	1 (25)	1.133 (0.09551 to 13.45)	0 (0.00)	1 (25)

Gender					
Male	33	10 (30.30)	1.522 (0.4711 to 4.916)	1 (3.03)	9 (27.27)
Female	27	6 (22.22)	0.6571 (0.2034 to 2.123)	3 (11.11)	3 (11.11)
District					
Manfalut	20	2 (10)	Ref.	0 (0.00)	2 (10)
Dayrut	8	2 (25)	3 (0.3434 to 26.21)	1 (12.5)	1 (12.5)
Abnub	8	5 (62.5)	15 (1.939 to 116)	1 (12.5)	4 (50)
Sodfa	7	2 (28.57)	3.6 (0.4002 to 32.38)	0 (0.00)	2 (28.57)
El Badari	6	2 (33.33)	4.5 (0.4791 to 42.27)	2 (33.33)	0 (0.00)
Others	11	3 (27.27)	3.37 (0.4688 to 24.3)	0 (0.00)	3 (27.27)
Total	60	16 (26.66)	0.85 (0.2783 to 2.587)	4 (6.66)	12 (20)
Adults					
Age (Year)					
18-39	14	6 (42.86)	9.941 (0.4693 to 210.6)	0 (0.00)	6 (42.86)
≥ 40	6	0 (0.00)	0.1006 (0.004749 to 2.131)	0 (0.00)	0 (0.00)
Gender					
Male	12	4 (33.33)	1.500 (0.2028 to 11.09)	0 (0.00)	4 (33.33)
Female	8	2 (25)	0.67 (0.09 to 4.93)	0 (0.00)	2 (25)
Diarrhea					
Yes	8	4 (50)	5 (0.6397 to 39.08)	0 (0.00)	4 (50)
No	12	2 (16.66)	0.20 (0.02559 to 1.563)	0 (0.00)	2 (16.66)
Total	20	6 (30)	1.179 (0.3866 to 3.593)	0 (0.00)	6 (30)

**Table S2:** Clinical symptoms on the *Campylobacter*-positive children

Symptom	No. of patients (Total no. 16)	%
Fever	14	87.5
Vomiting	14	87.5
Blood in stool	4	25
Diarrhea	16	100
Abdominal pain	16	100

## Discussion

Being of a low price relative to red meat and fish, poultry meat is a popular protein source for all Egyptians. However, contaminated poultry and poultry products can disseminate many zoonoses to humans, primarily including *campylobacteriosis*, thus representing a serious public health threat [8]. This threat is underestimated in developing countries owing to the lack of a national surveillance system for a such type of foodborne zoonosis [5]. Accordingly, periodical evaluation of the foodborne *campylobacteriosis* has to be implemented.

The present study combined both biochemical and multiplex PCR assays for identification of *Campylobacter* in broiler chicken and

humans in Assiut Governorate, Egypt. Multiplex PCR assay confirmed that 64 isolates of the 67 presumptively identified *Campylobacter* colonies were true positive. In contrary to biochemical reactions which does not typically detect mixed infection of *Campylobacter*, the current multiplex assay could reveal that up to 18.25% (52 of 285) were having a mixed infection of both *C. jejuni* and *C. coli*, the rest of the samples were found to be *C. coli* alone (4.21%, 12 of 285); never the less, none of them confirmed to have *C. jejuni* alone. Unlike other studies that showed the predominance of *C. jejuni* [9] or *C. coli* [10], our study stated that mixed infection was superior in all types of examined samples. Although there was an in significant difference in using either biochemical reactions or multiplex PCR assay in identifying *Campylobacter* species (95.52% compatibility), but multiplex assay surpassed biochemical reactions in detecting mixed infection, indicating the importance of such advanced assay in precise *Campylobacter* identification. Accordingly, a combination of both assays would be an excellent match for setting an efficient and cost effective diagnostic strategy for *campylobacteriosis*, thus facilitating rapid monitoring of such important foodborne zoonosis.

In this study, 3 of the 41 biochemically identified *C. jejuni* were found to be negative by multiplex PCR assay. Also, 5 isolates of them were confirmed to be *C. coli*. Such false negative results produced by hippurate test could be due to the existence or even production



of amino acids or peptides in the culture media [11]. Only 7 of 26 *C. coli* were confirmed by multiplex PCR while the rest was found to be a mixed infection. A reliable hippurate test result depends mainly on adjusting cell suspension turbidity which subsequently affects the color production and intensity [12]. This can be straight forward in testing reference strains, but in mixed infection the process is difficult to be controlled. For instance, upon employing the hippurate test, we have recorded a color change that was strong in some isolates and weak in others. Strong color-related isolates were suspected as *C. jejuni* from the beginning and mostly were further confirmed to be a mixed infection as noted above. However, those of weak color or colorless test results (along with the antimicrobial testing) were suspected as *C. coli*. Most of these isolates were also merged to mixed infection. The color weakness or the lack of color even with the presence of *C. jejuni* could be attributed to the variability in the *C. jejuni* concentration in the used inocula compared to other *Campylobacter* species or related species or the presence of amino acids or peptides as noted earlier.

The appearance of unfavorable growth at the initial *Campylobacter* isolation and identification steps followed in the present study was challenging. Such growth was in the form of translucent, beige to off-white, small colonies unlike the typical colonies of *Campylobacter* on mCCDA plates incubated at 37 °C. Our first thought was that the growth might be *Campylobacter*-related bacteria able to grow on the selective agar medium, mCCDA as reported elsewhere [13]. The colonies were found to be Gram negative with spiral shaped rods, hippurate negative, grown in air at 25 °C and 37 °C and not at 42 °C that typically matches the *Arcobacter* spp. identification scheme [3]. Taking the advantage of the inability of *Arcobacter* to grow above 37 °C unlike *Campylobacter*, we have completed our isolation protocol at 42 °C and this change was the possible reason behind the failure of further isolation of *Arcobacter* from those handled by the new protocol i.e. some samples derived from slaughterhouses and all human samples.

Despite the insignificant difference between farms, OR showed that both farms A and B had a higher *Campylobacter* contamination (OR for each= 7.76 (95% CI, 0.85–70.79) compared to other farms. Compared to our study, a higher percentage, 38.1% of *Campylobacter* contamination were reported by [14]. Mixed infection was the predominate finding in the examined cloacal swabs (12.87%, 13 of 101). This was nearly similar to that (12.4%) reported in Ecuador [15]. Conversely, a higher percentage of mixed infection (22%) was encountered by [16] and a lower percentage, 7% was detected by [17], respectively. *C. coli* was isolated from 3.96% of the examined cloacal swabs. Lower percentage (1.79%) was isolated by [18] and a much higher percentage, 33.33%, was detected by [19], respectively.

The variation in the *Campylobacter* recovery rate from different farms included in this study could be as a result of the difference in handling and management practices applied in different farms that affected the transmission pattern of *Campylobacter* from the environment to the broilers [20]. This study showed a low prevalence rate of *Campylobacter* in farms because sampling was

conducted between March and May. Such sampling period falls in the months (January and May) which previously found to have the lowest cumulative incidence of *Campylobacter* [21].

The overall prevalence in slaughterhouses was 24% accounted for 21.19% and 26.92% for slaughterhouse A and B, respectively. No statistical difference was observed between the two studied slaughterhouses, but slaughterhouse B showed a slightly higher odds of contamination compared to slaughterhouse A. Chicken neck had the highest contamination ratio (30.77%, OR= 2.44 (95% CI, 0.3609 to 16.55)) in the sampled chicken derived from slaughterhouse A, however, wing had greatest prevalence and odds of contamination in the other slaughterhouse (46.15, OR= 4.714 (95% CI, 0.7337 to 30.29)). Similarly, [22] reported that wing, neck and legs had the highest rate of contamination with *Campylobacter*. The high isolation rate of *Campylobacter* in chicken wings recovered in this study could be due to imperfect scalding, post scalding contamination, or due to the combination of both [23]. Also, feathers could be played an essential role in such a high contamination following getting the bacteria during transport, plucking process or during their mechanical removal in the slaughterhouses [24].

Owing to its lower price compared to beef liver, chicken liver became a popular for many people in the Egyptian cuisine. Unfortunately, *Campylobacter* enteritis associated with its consumption has been reported mainly as a result of post slaughtering contamination and additionally following systemic infection of live birds [25]. *Campylobacter* could be isolated in a 19.23% (5 of 26) of examined samples, thus chicken liver could represent an important public health threat and measures on how to minimize contamination should be disseminated to both slaughterhouse men and consumers. A much lower percentage (4%) was detected recently in Giza, Egypt [7] however, a higher isolation rate (24%) was documented elsewhere [26]. Contamination level detected in this study varied from carcass to another as well as between the organs of the same carcass. Accordingly, strict hygienic measures have to be applied and awareness regarding the risk of environmental contamination should be promoted to slaughterhouse men along with postmortem examination of poultry.

On the other hand, the overall prevalence of *Campylobacteriosis* in humans was 27.5%. This result was higher than a 8.1% [27]. The much higher percentage detected in our study could be attributed to inclusion of stool samples primarily obtained from gastroenteritis-infected individuals rather than investigating the disease in the general population. Sampled children showed a higher prevalence of *Campylobacter* spp. (26.66%) compared to a previous study that showed an isolation rate accounted for 9.6% [28]. Similarly, our study had a higher prevalence of *C. coli* (6.66%) and mixed infection (20%) in children than those reported by [29] which accounted for 1.11% and 2.22%, respectively. Such high prevalence noticed in our study may be attributed to; firstly, most of the sampled humans were originating from villages which usually does not adopt the basic hygienic standards and precautions in

contact and handling of live poultry. Secondly, over 41% (33 of 88) of the sampled humans were primarily risky infants of an age  $\leq$  1 year. Infants were documented to be at a higher risk of getting *campylobacteriosis* based on their impaired immunity, especially in developing countries [5]. Finally, sampling was achieved between May and August and those months were previously shown to have a substantial increase in the *Campylobacter* isolation rate in children compared to other months of the year [30].

Children are usually prone to get *campylobacteriosis* more likely than adults as mentioned in several reports and the bacterium is one of the most frequently isolated bacteria from stools of infants with diarrhea in developing countries [5]. For instance, it was found that children were more commonly infected with *Campylobacter* (19%) than adult males (8%) and females (7%) [31]. Never the less, our study stated a higher prevalence in adults (30%) than children (26.66%). This result came in accordance with [32] who noticed that prevalence of *campylobacter* in adult was also higher than that in children, however, he reported much lower prevalence rate accounted for 9.5% and 3.6%, respectively. Our finding is statistically insignificant and also, the odds ratio revealed that adults are at slightly higher risk (OR= 1.179 [95% CI, 0.3866 to 3.593]) of acquiring the disease compared to children. Regarding the gender of the examined children, a 30.03% and 22.22% of males and females were found to be infected with *Campylobacter*, respectively. The result was insignificant, but odds ratio indicated that males are more likely to contract *campylobacteriosis* (OR=1.522 [95% CI, 0.4711 to 4.916]) than females. Similarly, [33] reported that the *Campylobacter* infection rate was significantly higher among males than females.

The impact of age on acquiring *campylobacteriosis* by children was also assessed in the present study. Among the three studied age groups,  $\leq$  1 year aged children had the highest prevalence rate (33%). A typically similar rate (32.6%) has been previously reported in Cairo [34]. Concerning the residence of sampled children, insignificant variation between different districts was encountered, however, the OR revealed that not all districts had the same *Campylobacter* distribution; Abnub was the highest (OR= 15 [95% CI, 1.939 to 116]) followed by El Badari (OR= 4.5 [95% CI, 0.4791 to 42.27]) and Sodfa (OR= 3.6 [95% CI, 0.4002 to 32.38]).

This difference is difficult to be reasoned, but it can be related to eating customs and preferences and close proximity to birds as well as the incomparable number of children examined from each district. Additionally, it is worthy to mention that most of the children sampled from Abnub were primary of  $\leq$  1 year that was recorded to have the highest risk of acquiring *campylobacteriosis*. These factors could individually or collectively contribute to alleviate such clear discrepancy between different districts. Fever was recorded in 87.5% of infected children, however, all patients had diarrhea and abdominal pain. This results were assured by [35]. The presence of blood in 25% of stool samples of infected children may be due to the delayed hospitalization of sick children. Blood in stools of *Campylobacter*-infected children appears within the first day of disease onset and usually lasts for three days [36]. Another reason

behind this relatively lower percentage of bloody stool compared to other symptoms in patients could be the early life production of IgG antibodies against *Campylobacter* in developing countries [37]. Although vomiting is a less frequent sign of *Campylobacteriosis* and only occurs in approximately 20% of patients [38], our study showed a completely contradicting finding. Surprisingly, our study reported the presence of an unexpected percentage (87.5%) of infected children manifesting emesis.

This finding has never been reported elsewhere and needs explanation. The presence of other gastrointestinal infection [39] and the small number of infected patients (no=16) used for setting the present estimate could be possible reasons. Interestingly, the impact of gender on acquiring *Campylobacter* infection in adults was nearly similar to that in children participated in the present study. *Campylobacter* was found to infect adult males more likely (OR=1.500 [9% CI, 0.2028 to 11.09]) than adult females (OR= 0.67 [95% CI, 0.09 to 4.93]). This difference was also found insignificant and the increased odds of exposure of males versus females could be attributed to the immune response. Under rural Egyptian setting, females get involved in practicing traditional aviculture as well as kitchen responsibilities early in their life thus they are exposed to *Campylobacter* repeatedly than males and as a result, females may be more likely develop specific antibodies to the bacteria than males.

*Campylobacteriosis* prevalence appears to wane with age as reported in our study. The prevalence of *Campylobacter* infection was decreased from 42.86% in the age group of 18-39 years old to 0% in  $\geq$ 40 years old. This result indicated that the disease is not a problem in old ages. This reinforces the impression that excessive exposure of individuals to *Campylobacter* over the life could result in the development of protective immunity. Similarly, [40] reported that 36.5% of all *Campylobacter* isolates was isolated from adults aged 20-39 years.

The prevalence of *Campylobacter* in diarrheic adults (50%) was higher than that of non- diarrheic (16.66%). Statistically, the difference between both groups was not significant, however, diarrheic individuals were found to be infected more likely with *Campylobacter* compared to non-diarrheic.

## Conclusion

The present study reports the detection of mixed infection of *Campylobacter* species in different poultry farms and slaughterhouses. Co-existence of *Arcobacter* with *Campylobacter* in the examined poultry samples is a striking feature for this study. This confirms that broilers could represent an important public health threat to the people residing in Assiut Governorate. Accordingly, measures on how to minimize the contamination level at farms, slaughterhouses and during culinary practices should be disseminated to farm workers, slaughterhouse men and consumers. Furthermore, a combination of both biochemical and multiplex assays would be an excellent match for setting an efficient and cost effective diagnostic strategy for *campylobacteriosis*, thus facilitating rapid monitoring of such important foodborne zoonosis.

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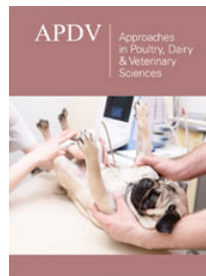
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