

Natural co-infection of *Escherichia coli* and Infectious bronchitis virus in broilers and layers flocks

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ABSTRACT

The present study was designed to throw the light on isolation of Escherichia coli (E. coli) strains accompanied by Infectious bronchitis virus (IBV). Classification and serotyping of bacterial isolates of E.coli were done followed by their antimicrobial susceptibility testing, also isolation and identification of IBV was done. For that purpose, a total of 200 organs, swap and samples were aseptically collected from broilers and layers flocks suffered from respiratory symptoms and general illness from both private and governmental farms during the period from 2015 till the end of 2017. Results revealed that the prevalence of E. coli was 67.5% and IBV was 61.5% in the collected samples. The serological identification showed that the most predominant serotype for E. coli was O 158 and there was a high level of resistance of isolated strains against doxycycline and tetracycline.

Keywords: Broiler, E. coli, O158, IBV, co-infection.

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1. INTRODUCTION

Diseases of the respiratory tract are a significant component of the overall disease incidence in poultry. In many cases, respiratory disease observed in a flock may be a component of a multisystemic disease or it may be the predominant disease with lesser involvement of other organ systems. In some cases, such as infectious coryza or infectious laryngotracheitis, the disease may be limited to the respiratory system, at least initially. Various pathogens may initiate respiratory disease in poultry including a variety of viruses, bacteria and fungi. Environmental

factors may augment these pathogens to produce the clinically observed signs and lesions (Glisson, 1998).

Infectious bronchitis virus (IBV) is a single stranded positive sense, enveloped RNA virus (Lai and Cavanagh, 1997). The virus has been classified under the *Gammacoronavirus* genus in the family *Coronaviridae*, Order *Nidovirales*. Like other members of *coronavirus* family, the IBV genome is composed of structural and nonstructural proteins. Infectious bronchitis virus (IBV) is one of the major economically important poultry viral diseases distributed worldwide (OIE, 2013).

It affects both galliform and nongalliform birds. Its economic impact includes decreased egg production and poor egg quality in layers, stunted growth, poor carcass weight, and mortality in broiler chickens. (Bande *et al.*, 2016).

E. coli strains causing systemic disease in poultry (avian colibacillosis) are termed avian pathogenic E. coli (APEC). Colibacillosis is a disease of severe economic significance to all poultry producers worldwide and is characterized by a diverse array of lesions (Dziva and Stevens, 2008). These lesions varied between perihepatitis, airsacculitis and pericarditis, or other syndromes such as egg peritonitis, salpingitis, coligranuloma, cellulitis omphlitis, and osteomyelitis/ arthritis (Barnes and Gross, 1997).

Co-infection of *E. coli* with infectious bronchitis virus (IBV) may lead to a more complex outcome, usually associated with high morbidity and mortality. Similarly, infection with nephropathogenic IBV strains may result in pale, swollen, and mottled kidneys (Boroom and *et al.*, 2012).

Therefore, the present study was planned out to throw a light on the natural co-infection of isolated bacterial strains (*E. coli*) and IBV.

2. Materials and methods

2.1. Sample collection:

Birds showing symptoms of respiratory diseases and general illness were collected and submitted to the Reference Laboratory for Veterinary Quality Control on Poultry Production, Dokki, for investigating co-infection of respiratory viral infection (IBV) with *E.coli* infection in broilers and layers flocks. The birds were obtained from both private and governmental farms during the period from 2015 – 2017. A total of 200

broiler and layer flocks (200 samples) examined Organs and swabs were collected aseptically to prevent cross contamination. The collected organs were cultured within a time limit which did not exceed 24 hours from collection.

2.2. Bacterial isolation and identification: Preparation of samples:

The internal organs included liver, spleen, heart and other organs were prepared for isolation of E.coli according to (Quinn et al., 2002).

Microscopic examination:

Gram's stain was prepared and used as described by Cruickshank et al., (1975) for morphological study.

Biochemical Identification:

According to Quinn et al., (2002) including Indole reaction, Methyl red test, Voges Proskauer test, Citrate utilization test, Catales test, Sugar fermentation test, Oxidase test, Triple sugar iron and Christener's urea agar test.

Serological identification of E. coli isolates:

Typing of *E. coli* isolates was performed by the slide agglutination test using standard polyvalent and monovalent *E. coli* antisera according to Edwards and Ewing., (1972).

2.3. Determination of antimicrobial susceptibility profiles for E. coli isolates (CLSI, 2007):

The disk diffusion technique was applied according to (CLSI, 2007). The following antibiotics were used (Doxycycline, Sulphamethoxazole/trimethoprim,

Levofloxacin, Norfloxacin, Nitrofurantin, Ampicillin, Ciprofloxacin, Gentamycin, Tetracycline and Chloramphenicol) and interpreted according to (CLSI, 2007).

2.4. Viral identification:

Preparation of samples for IBV detection:

The collected organs (Trachea, lung and kidney) were washed in sterile saline, and then frozen at below-10°C. After thawing, the (10%) tissue homogenates w/v) were suspended in sterile saline (0.85% w/v)containing 100 IU/mL penicillin, 1.0 mg/ml streptomycin. By disrupting organs using sterile mortar and pestle, the homogenates were then centrifuged at 3000 rpm for 10 min, and the supernatant was further passed through 0.22 µm membrane filter for clarification. The supernatant was subjected to RNA extraction (OIE, 2013).

Detection of IBV directly from samples using RRT-PCR:

RNA extraction

RNA was extracted from the prepared suspension QIAamp Viral RNA Mini Kit(Qiagen, Valencia, Calif., USA,Cat. No. 52904.). The procedure was performed according to the company's instruction.

The extracted viral RNA was preserved at - 70°C until subjected for RRT-PCR.

Amplification of the extracted RNA by Real Time-RT-PCR (RRT-PCR)

Amplification was done using Quantitect RT-PCR(Oiagen, Inc. Valencia CA) in accordance with kit instruction In brief one reaction formed from 12.5 µl RT-PCR Master Mix, 6.6 µl Rnase free water, 0.5 µl for each primer and 0.125 µl probe. Oligonucleotide primers and probe used in real time PCR as previously described by Meir et al. (2010) in concentration of 50 pmol for primers and 30 pmol for probe, Briefly, foreword primer so AIBV-fr named 5' ATGCTCAACCTTGTCCCTAGCA 3' and reverse primer so named AIBV-as 5' TCAAACTGCGGATCATCACGT'3 and probe so named AIBV-TM (5'FAM-TTGGAAGTAGAGTGACGCCCAAACTTC A-TAMRA'3. Amplification conditions were Reverse transcription for 30 mins at 50 °C, primary denaturation for 15 min at 95 °C and then cycling steps 95 °C for 15 sec for 2nd

denaturation and then 60 °C for 45 sec for annealing and extension. that repeated for 40 cycles.

3. RESULTS

Bacterial isolation and serological identification.

The prevalence of *E.coli* infection reached 67.5% (135 out of 200 samples) many chickens showing symptoms of colisepticemia. The incidence of *E. coli* infection was 120 samples out of 160 samples with a percentage 75% in broiler and 15 out of 40 samples with a percentage 37.5% in layers Table (1).

Prevalence of E. coli strains in isolates from broilers and layer flocks.

Serotyping of 135 *E.coli* isolates was applied by slide agglutination test using polyvalent and monovalent O *E.coli* antisera. 14 different serotypes were identified among *E.coli* isolates and the most predominant serotypes were O158:k- ,O25:k11, O26:k60 ,O78:k80 and O44:k74 on the other hand 11.1% of isolates were untypable this showed in table (2)

Antimicrobial susceptibility test.

A high level of resistance of E.coli isolates was recorded to doxycycline (67.5%) and tetracycline (65%) followed by Nitrofurantin (64.2%), ampicillin (58.3%) finally Ciprofloxacin (6.7%) as shown in table (3).

Result of viral isolation.

Prevalence of IBV recovered from different flocks:

The result of virus detection by real time RT-PCR revealed that 103 out of 160 examined samples were positive for IBV with a percentage of 64.3% in broilers, while in layers 20 samples out of 40 were positive for IBV with a percentage of 50%. Meanwhile the total percentage of detection was 61.5% Table (4). Also studying occurrence of IBV and E. coli combined infection revealed that E. coli+ IBV was 20% as shown in table (5)

where E. coli alone prevalence percentage was 47.5% and IBV alone was 41.5%.

Type of flocks	Number of examined samples	Number of +ve samples	% of +ve samples
Broilers	160	120	75
Layers	40	15	37.5
Total	200	135	67.5

Table 1: Prevalence of E. coli recovered from broilers and layers flocks.

Table 2: Prevalence	of E. coli	strains ir	isolates	from	broilers	and laver fl	ocks.
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Serotype	Serotype number	Percentage of serotype		
O158:k-	18	13.3		
O25:k11	15	11.1		
O26:k60	11	8.1		
O78:k80	10	7.4		
O44:k74	9	6.7		
O114:K90	9	6.7		
O119:k69	8	6		
O103:K-	8	6		
O127:K63	7	5.2		
O125:K70	7	5.2		
O86:K-	6	4.4		
O91:K-	5	3.7		
O111:K58	4	3		
O8:K50	3	2.2		
Untypaple	15	11.1		

Table 3: Antibiogram profile of each antimicrobial against E. coli isolates from chicken.

Isolate	DO	TE	F	AMP	С	CN	LEV	SXT	NOR	CIP
Sensitive	•									
NO	. 10	13	14	19	34	28	50	43	65	103
%	8.3%	10.8%	11.6%	15.8%	28.3%	23.3%	41.7%	35.8%	54.2%	85.8%
Intermediate										
NC) 29	29	29	31	36	45	25	35	25	8
%	24.2%	24.2%	24.2%	25.8%	30%	37.5%	20.8%	29.2%	20.8%	6.7%
Resistant										
NO.	81	78	77	70	50	47	45	42	30	9
%	67.5%	65%	64.2%	58.4%	41.7%	39.2%	37.5%	35%	25%	7.5%

Doxycycline. (DO), Sulphamethoxazole/trimethoprim. (SXT), Levofloxacin.(LEV), Norfloxacin.(NOR), Nitrofurantin.(F), Ampicillin. (AMP), Ciprofloxacin. (CIP), Gentamycin.(CN), Tetracycline.(TE), Chloramphenicol.(C)

Table 4: Prevalence	of IB virus recovered	d from different flocks.

Type of flocks			+ve % of +ve samples
	examined samples	Samples	
Broilers	160	103	64.3
Layers	40	20	50
Total	200	123	61.5

Table 5: Prevalence of co-infection.

Type of infection	Number of +ve samples	% of +ve samples
E. coli alone	95	47.5
IBV alone	83	41.5
<i>E. coli</i> + IBV	40	20

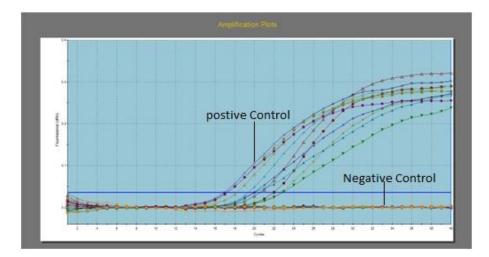


Fig.1. Amplification curves of the tested positive samples

4. DISCUSSION

In the present work it was recorded that the prevalence of isolation of E.coli reached 67.5% from the examined 200 samples showing symptoms of colisepticaemia. Almost similar percentages 67% in chickens were reported by Syuhada et al., (2013), and Stella et al., (2016) reported that From the 80 sampled birds, 48 (60%) E. coli was detected of them. While a higher incidence of 85.2% was reported by Wani et al., (2004), and Albarri et al., (2017) where it was (93.75%).

On the other side a lower incidence (20.5%) was recorded by Saidi *et al.*, (2013) and (43.1%) by Roshdy *et al.*, (2012).

The result of serotyping revealed that , the most commonly isolated serotypes were O158, O26 ,O78,O44 ,O114 ,O119 ,O103 ,O127, O125 ,O86, O91and O111. These result mostly similar to another study was conducted by Roshdy *et al.*, (2012). Also Abd El-Twab *et al.*, (2015) reported that the most commonly serogroups isolated from chickens were O44, O158, O125 and O103.

In the present study the susceptibility of *E.coli* isolated from broilers and layers

chicken showed a high sensitivity to Ciprofloxacin. This finding is similar to previous studies conducted by Guerra et al., (2003) and Hasan et al., (2011) which found that E. coli isolates were highly sensitive to ciprofloxacin. The high sensitivity ciprofloxacin might be because it is a broad spectrum antibiotic that is still relatively new and has limited use by poultry farmers. Unlike Omer *et al.*, (2010), who found that avian *E*. coli isolates were highly resistant to ciprofloxacin. The data of this study showed that there was a high level of resistance against doxycycline, tetracycline and ampicillin. This is go a hand with a several reports (Sharada et al., 2009) and (Zakeri and Kashefi, 2012). The presence of high resistance is probably due to the increased use of antibiotics as feed additives, for example, tetracyclines, bacitracin, and cloxacillin are widely used in poultry industries for growth promotion or prevention of diseases (Omer et al., 2010).

Concerning to viral isolation, a total of 123 samples out of 200 were positive for IBV detection (61.5% of the samples), where 64.3% and 50% of the samples were positive for broilers and layers respectively. Almost similar results were recorded by Roussan et al., (2008) and Zanaty. (2014). While a higher percentage reached to 88% recorded by Abdel-ElGhany et al., (2015). On the other hand a lower percentage was recorded by Mohamed and Ibrahim (2015) which was 14.28%. It was found that the percentage of co-infection occurrence was 20% which studied previously by Boroom and et al., (2012), who found that E. coli lead to more exacerbating effect to IBV, which may be due to E. coli provide the enzymes capable of cleaving the hemagglutinin of viruses like IBV enabling them to replicate and spread to a greater extent in that host (Bano et al., 2003)

It was concluded that co-infection between Infectious bronchitis virus and *E.coli* lead to a more complex outcome, than infection with IBV alone so we should give more attention to secondary bacterial infection specially *E.coli* and take the suitable preventive measures and precautions against them.

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