



Detecting Some Virulent Bacterial Causes of Omphalitis in Broiler Chickens

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Abstract

OMPHALITIS is the main cause of mortality and low performance in broiler chicks. Detection of some virulent bacterial causes of yolk sac infection in broilers was the aim of this study. This investigation included 45 examined broiler flocks in Delta region of Egypt, aged 1-8 days old and suffered from high mortality, depression, diarrhea and low vitality. Bacterial isolation revealed 39 *E. coli* isolates (86.6%), 7 *Salmonella* isolates (15.5%), 6 *Pseudomonas aeruginosa* isolates (13.3%) and 3 coagulase-negative *Staphylococcus* isolates (6.6%). The polymerase chain reaction test used to detect virulent genes in these isolates revealed that all 4 examined *E. coli* isolates had *iroN*, *hlyF* and *ompT* genes. The *invA*, *stn* and *fimA* genes were detected in 4 examined *S. Enteritidis* isolates. *Staphylococcus lentus* isolate was positive to *sea* and *tsst* genes and negative to *etb* gene, and *S. saprophyticus* isolate was positive to *sea*, *tsst* and *etb* genes while *S. xylosus* was positive to *etb* gene and negative to *sea* and *tsst* genes. *tox A*, *phzM* and *oprL* genes were identified in 4 tested *P. aeruginosa* isolates. Antimicrobial sensitivity testing revealed that all 4 tested *E. coli* isolates were resistant to ampicillin while doxycycline and imipenem showed sensitivity. While the 4 examined *S. Enteritidis* isolates were resistant to ampicillin but showed sensitivity to gentamycin, amikacin and imipenem. The coagulase-negative *Staphylococcus* isolates were resistant to clindamycin but sensitive to doxycycline, ciprofloxacin and gentamycin. *P. aeruginosa* isolates (n=4) were resistant to imipenem and were sensitive to amikacin, cefepime and ciprofloxacin.

Keywords: Omphalitis, Yolk sac infection, Bacteria, Virulence genes, Multi drug resistant.

Introduction

Ensuring the health and productivity of poultry is therefore essential for sustaining this industry, particularly during the critical early stages of a chick's life. Embryonic development of chickens depends heavily on nutrients derived from the yolk, which serves as the primary reservoir of nourishment. The yolk sac, analogous to the mammalian placenta, facilitates the absorption, digestion, and transfer of these nutrients to the developing embryo. However, the period following hatching, especially the first week, is fraught with

challenges. This stage accounts for 30-50% of overall mortality in broiler chicks, with omphalitis emerging as a leading cause of death along-side other conditions such as avian encephalomyelitis, brooder pneumonia, and pullorum disease [1].

Omphalitis, also known as yolk sac infection (YSI) or mushy chick disease, is a non-contagious condition often linked to unhealed navels which may be caused by infectious and/or non-infectious agents. This disease is economically significant, as it not only causes increased mortality but also leads to poor weight gain, stunted growth, and inferior carcass quality in surviving chicks [2]. The prevalence of

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YSI varies between 5% and 30%, with higher incidence rates reported during colder seasons such as winter and autumn [3].

The unabsorbed yolk sac in young chicks provides an ideal medium for bacterial growth such as *Escherichia coli*, *Salmonella* spp., and *Staphylococcus* spp. which responsible for omphalitis in broiler and layer chicks [4]. Infection typically occurs when the navel fails to close properly after the yolk sac is drawn into the abdominal cavity, allowing environmental bacteria to invade the chick tissues. Poor hygiene on breeding farms and improper hatchery management exacerbates the problem [5].

The degree of the organism's virulence is based on certain virulence factors that may have acquired. These factors may be toxins or surface structures that enable the bacterium to adhere to vector cells or to attack the immune system. According to *E. coli* the five essential virulence genes carried by plasmid, hlyF (hemolysin), ompT (outer membrane protease), iroN (siderophore), iss (serum survival), and iutA (iron transport), are considered markers for APEC from other avian fecal *E. coli* [6]. All strains and serovars of *Salmonella* enterica were found to carry the *Salmonella* enterotoxin determinant (stn) gene which has been proposed to be a putative causative agent of diarrhea [7]. *Salmonella* adhesion to the host cell is an initial and most important step in the pathogenesis of salmonellosis. Fimbriae which encoded by (fimA, agfA, sefA and sefD) genes are one of the most important tools of bacterial adhesion to host cell [8]. In addition, Toxic shock syndrome (TSS) caused by *Staphylococcus* is an acute, life-threatening illness characterized by the sudden onset of high fever, hypotension, and a widespread erythematous rash. If untreated, it can progress rapidly to multi-organ dysfunction and potentially fatal shock [9]. *P. aeruginosa* also has got massive arsenal of virulence supplies such as lipopolysaccharide, elastase, alkaline proteases, pyocyanin, pyoverdine, hemolysins, phospholipase C and rhamnolipids [10]. The outer membrane proteins of *P. aeruginosa* (OprI and OprL) play important roles in the interaction of the bacterium with the environment as well as the inherent resistance of *P. aeruginosa* to antibiotics And pyocyanin produced by *P. aeruginosa* is a key virulence factor that causes damages to airway and lung in hosts. Conversion of phenazine-1-carboxylic acid to pyocyanin involves an extra metabolic pathway which contains two enzymes encoded, by phzM and phzS [11].

While antimicrobial therapy is commonly used to treat omphalitis, the overuse of antibiotics in both human and veterinary medicine has led to the emergence of antibiotic-resistant pathogens. This growing resistance is a global concern, with predictions suggesting that by 2050, antibiotic-resistant infections could result in 10 million deaths

annually [12]. The persistent release of antibiotics into the environment due to extensive use in the veterinary sector exacerbates this issue, underlining the need for alternative strategies to combat bacterial infections.

This research aimed to unveil some bacterial causes of omphalitis in broiler baby chicks, focusing on their incidence, virulence genes and antimicrobial sensitivity.

Material and Methods

Sample collection for bacterial isolation and identification

Samples from liver and yolk sacs (n=10 per flock) in diseased broiler chicken flocks (n= 45) in Delta region, Egypt from May till October 2023. The examined flocks aged 1 - 8 days old were suffering from high mortality (3-12%) in the first 10 days of age, retarded growth, depression, huddling together, profuse watery diarrhea (pasty vent) and lameness. Samples were weighed and suspended in nutrient broth (as 1:10 dilution) and incubated at 37 C for 20 hours. Loops-full of suspension were streaked onto MacConkey agar (Oxoid, Ltd, UK), Baird parker agar (Oxoid, Ltd, UK), Cetrimide agar (Himedia, India) and incubated at 37 C for 24 hours. Further culture of suspected colonies on selective media such as eosin methylene blue agar (EMB, Oxoid, Ltd, UK), Rappaport-Vassiliadis, Xylose Lysine Deoxycholate agar media (XLD, Oxoid, Ltd, UK) were applied, and the suspected colonies were stained by Gram's stain and examined microscopically. Biochemical identification tests (indole test, methyl red Test, Voges – Proskauer reaction, citrate utilization test, urea hydrolysis test, hydrogen sulfide production test, oxidase test and catalase test) were carried out on recovered suspected colonies according to Quinn et al. [13]. This work followed the guidelines of the Ethics Committee of Damanhur University and was approved by the committee decision No. DMU/VetMed-2025/027.

Serological, virulence genes identification and antimicrobial sensitivity test

These tests was applied on four *E. coli*, four *Salmonella*, three *Staphylococcus* and four *P. aeruginosa* isolates recovered from cases which showed severe postmortem lesions with the highest mortality rates.

Serotyping of E. coli

Four isolates were selected for serological identification according to using rapid diagnostic *E. coli* antisera sets for (O) and (H) antigens (Denka Seiken Co., Japan) for diagnosis of the Enteropathogenic types and pathotype categorization was according to Bahgat et al. [14].

Serotyping of *Salmonella* isolates

Also, 4 *Salmonella* isolates were selected for serological identification by using *Salmonella* antisera of (Sifin Company, Germany) and interpretation of results according to white Kauffman scheme 2007 [15].

Identification of *Staphylococcus* isolates

VITEK2 COMPACT identification system (bioMérieux, France) was used for definite identification of 3 obtained *Staphylococcus* isolates directly from suspected colonies grew on Baird parker agar media.

Identification of *P. aeruginosa*

The identification of *P. aeruginosa* was dependent on colony morphology on cetrimide agar which is specific to *P. aeruginosa* then application of biochemical tests.

Virulence genes identification by using PCR test

DNA was extracted from isolates by using QIAamp DNA mini kit (Qiagen N.V, Germany) and the Oligonucleotide primers used in cPCR as shown in Table 1. Preparation of PCR Master Mix was used according to Emerald Amp GT PCR master mix (Takara, Japan) by adding 12.5 µl of Emerald Amp GT PCR master mix (2x premix), 5.5 µl of PCR grade water, 1 µl of forward primer (20 pmol), 1 µl of reverse primer (20 pmol) and 5 µl of template DNA. Cycling conditions of the two primers during PCR are shown in Table 1. Agarose gel electrophoreses (1%) applied according to Sambrook et al. [25] with some modifications. Each uniplex PCR product, negative and positive controls (20 µl) were loaded to the gel. The gel was examined under UV cabinet and photographed by a gel documentation system.

Antimicrobial sensitivity test

The antimicrobial sensitivity test was carried out according to the guidelines of Clinical and Laboratory Standards Institute [26] using Mueller-Hinton agar (Oxoid, Ltd, UK) for *E. coli*, *Salmonella* and *P. aeruginosa* isolates, and nutrient agar (Oxoid, Ltd, UK) for coagulase-negative *Staphylococcus* isolates. The following antimicrobial discs: ampicillin (10 µg), cefepime (30 µg), cefotaxime (30 µg), ceftazidime (30 µg), gentamycin (10 µg), amikacin (30 µg), ciprofloxacin (5 µg), imipenem (10 µg), trimethoprim-sulfamethoxazole (1.25/23.75 µg), doxycycline (30 µg), erythromycin (15 µg), linezolid (30 µg) and clindamycin (2 µg) were used. Criteria for the interpretation of the zone diameter for sensitive, sensitive dose dependent, intermediate and resistance was done and recorded as CLSI guidelines, 2020 as shown in Tables 2, 3, 4.

Results

Postmortem examinations

Internally, the examined chicks showed retained putrefactive yolk sac, hyperemia of visceral blood vessels, inflammation and edema around the navel, discoloration of yolk sac (greenish yellow or dark brown) and in severe cases distension of chick's abdomen, congestion of internal organ and fibrinous airsacculitis was found.

Isolation and identification of bacterial isolates

Bacteriological examination of diseased chicks with omphalitis revealed that 43 out of 45 broiler flocks (95.5%) were bacteriologically positive. Single bacterial isolation was in 35 flocks (77.8 %), but mixed bacterial infection was in 8 flocks (17.8 %). Identification of bacterial isolates revealed that out of 45 examined samples *E. coli* 39 isolates (86.6%), *Salmonella* 7 isolates (15.5%), coagulase-negative *Staphylococcus* 3 isolates (6.6%) and *Pseudomonas aeruginosa* 6 isolates (13.3%). Morphology of growing colonies on selective media (MacConkey, EMB, XLD, Baird parker and Cetrimide agar media) was shown in Table 5, and biochemical test results were shown in Table 6.

Serotyping of *E. coli* isolates

Serological identification of *E. coli* isolates revealed suspected enteropathogenic *E. coli* serotypes O119: H6 and O146: H21, enterohemorrhagic *E. coli* O55: H7 and O128: H2 as enterotoxigenic *E. coli*.

Serotyping of *Salmonella* isolates

Serological identification of (O) and (H) antigen of 4 *Salmonella* isolates revealed *S. Enteritidis* O (1, 9, 12) and H (g, m).

Identification of *Staphylococcus* isolates

Three *Staphylococcus* isolates were identified by using VITEK 2 Compact technique into *Staphylococcus xylosus*, *Staphylococcus saprophyticus* and *Staphylococcus lentus*.

Seasonality

This Study was performed in 6 months from May to October and Its noticed that all *Pseudomonas aeruginosa* isolates recovered during October and November (Fig. 1).

PCR for virulence genes identification of bacterial isolates

E. coli isolates

Four *E. coli* isolates had *iroN*, *hlyF* and *ompT* virulence genes (Fig. 2).

Salmonella isolates

Four *Salmonella* isolates had the *invA*, *stn* and *fimA* virulence genes (Fig. 3 and 4).

Staphylococcus species isolates

Three *Staphylococcus* isolates examined by PCR for virulence genes of *sea*, *tsst* and *etb*. Sample no. (1) (*Staphylococcus lentus*) was positive to *sea* and *tsst* genes and negative to *etb* gene, sample no. (2) (*Staphylococcus saprophyticus*) was positive to *sea*, *tsst* and *etb* genes while sample no. (3) (*Staphylococcus xylosus*) was positive to *etb* gene and negative to *sea* and *tsst* genes (Fig. 5).

P. aeruginosa isolates

The four examined *P. aeruginosa* isolates had these virulence genes *toxA*, *phzM* and *oprL* (Fig. 6).

Antimicrobial Sensitivity tests

The in-vitro antimicrobial sensitivity test for different bacterial isolates of *E. coli*, *S. Enteritidis*, *Staphylococcus* species and *P. aeruginosa* are presented in Table 7.

Discussion

The present study was carried out for the isolation of bacteria associated with yolk sac infection (omphalitis) in broiler chicks from 45 flocks of 1 - 8 days old suffered from high mortality, depression, pasty vent, watery chalky diarrhea and decrease in vitality. Postmortem examination of sick chicks showed retained yolk sac, hyperemia of visceral blood vessels, inflammation and edema around the navel, discoloration of yolk sac (greenish yellow or dark brown) and distension of chick's abdomen in all cases. In addition, in severe cases no (1, 3, 8, 13, 15, 22, 27, 34, 36, 37, 39, 42 and 43) there were putrefactive odor of yolk sac, enteritis, severe congestion of internal organs and fibrinous airsacculitis.

Bacteriological examination of chicks with omphalitis from 45 broiler flock revealed that 43 flock (95.5%) of examined samples were positive for bacterial isolation. *E. coli* was the most predominant bacteria causing omphalitis in 86.6%, which was very close to results of Shaheen et al. [27] with a rate of 87.5%. Virulence factors of *E. coli* may be toxins or surface structures that enable the bacterium to adhere to vector cells or to attack the immune system. Detection of the *iroN*, *hlyF* and *ompT* genes using PCR indicated that all the four examined *E. coli* isolates are virulent. The *iroN* gene was widely prevalent among APEC strains, consistent with findings from previous studies by Pourhossein et al. [28], reporting prevalence rate of 78.69%. Similarly, the *hlyF* gene was identified in 100% of cases in the study by Sedeek et al. [29]. The prevalence of the *ompT* gene was reported as 86.9% by Mohamed et al. [30]. The *iroN*, *hlyF*, and *ompT* genes are primarily located on the ColV plasmid, which is known to harbor numerous antimicrobial resistance genes [17]. The presence of these plasmid-associated virulence genes in the isolates examined in this study

may explain their resistance to multiple antibiotics, reflecting the co-selection and dissemination of both virulence and resistance determinants through horizontal gene transfer.

Antimicrobial therapy has played a crucial role in controlling bacterial infections in poultry, which can lead to significant health issues and economic losses. However, the widespread use of antimicrobials has also contributed to the emergence of resistant strains, making it challenging to control infections effectively. This resistance can lead to cross-resistance issues, especially concerning human enteric pathogens, as some antimicrobials used in veterinary settings overlap with those used in human medicine.

Antimicrobial sensitivity test was performed on *E. coli*, *Salmonella* and *P. aeruginosa* isolates which cause severe omphalitis in addition to three recovered coagulase-negative *Staphylococcus* species. Multidrug resistance in *E. coli* has become a worrying issue that is observed worldwide in humans and veterinary medicine. As it has a great ability to gain antibacterial resistance genes, mostly through horizontal gene transmission [31]. In the present *E. coli* isolates were completely resistant to ampicillin, showed also 75% resistance to cefotaxime and gentamycin and 50% to amikacin, ciprofloxacin and trimethoprim-sulfamethoxazole but showed no resistance to doxycycline and imipenem.

Salmonella Enteritidis was the second pathogen responsible for omphalitis with a percentage of 15.5% and this rate of infection was constituted with results obtained by Kaboudi et al. [32] who reported *Salmonella* isolation rate from omphalitis at 10.6%. Also, *Salmonella* infection was the second bacteria causing omphalitis but with high rate of infection as 62.5% as reported by Shaheen et al. [27]. The main infection route with *Salmonella* species was through an unhealed navel, but transmission can also occur via the bloodstream or through yolk contamination, leading to infection in chicks [3]. Based on serological identification of (O) and (H) antigen, 4 tested *Salmonella* isolates were *S. Enteritidis*. The cPCR results indicated the presence of *invA*, *stn* and *fimA* virulence genes in these *S. Enteritidis* isolates. The *invA* and *stn* genes are considered main genes and widely spread in *Salmonella* isolates as it was detected with 100 % in the studies of Shen et al. [33].

In addition, the *fimA* gene was also detected in all isolates in the previous studies of Lozano-Villegas and Rondón-Barragán [34].

Epidemiological studies indicate that multidrug resistant *Salmonella* serotypes are more virulent than others, as reflected by increased severity and more prolonged disease in patients infected by these strains [35]. Antimicrobial sensitivity test of *Salmonella* isolates showed complete resistance to Ampicillin, high resistance (75%) to cefotaxime and

doxycycline, 25% resistant to ciprofloxacin, trimethoprim-sulfamethoxazole and cefepime, and *Salmonella* isolates were completely sensitive to gentamycin, amikacin and imipenem.

Staphylococcus infection rate was 6.6%. However, coagulase-positive *Staphylococcus* is more common and more virulent than coagulase-negative *Staphylococcus* species, all 3 recovered *Staphylococcus* isolates were coagulase-negative (*S. lentus*, *S. saprophyticus* and *S. xylosus*).

Similarly, Nasrin et al. [36] isolated coagulase-negative staphylococci with percentages (24% in 1-3 days old chicks and 28.6% in 4-7 days old chicks). Our data revealed that there is a complete correlation between coagulase-negative *Staphylococcus* (CoNS) infection and *Salmonella* infection as all three recovered CoNS isolates are mixed infection with *Salmonella* isolates, and that may be due to activation of a complex immune response by *Salmonella* infection, which not only combats the invading pathogen but also disrupts the accurate balance of the gut microbiota. In the past, CoNS have been considered as a non-pathogenic microorganism as most of these species can establish a commensal relationship with humans and animals [37]. But CoNS have gained high importance over the past decade, as they have been involved in different infections in humans and animals [38]. Historically, toxic shock syndrome (TSS) has been associated with coagulase-positive *Staphylococcus* species. However, in the present study, the *tsst* gene was identified in two CoNS isolates (*Staphylococcus lentus* and *Staphylococcus saprophyticus*) but was not detected in *Staphylococcus xylosus*. Similarly, the presence of the *tsst* gene in CoNS strains has been documented in studies by Nasaj et al. [39] with reported rates of 25%. Enterotoxin genes were widely distributed among CoNS strains. For instance, Nasaj et al. [39] reported that CoNS isolates harbored enterotoxin-encoding genes with rate of 43% respectively. In addition, the present study found that the *sea* gene was identified in two CoNS isolates (*Staphylococcus lentus* and *Staphylococcus saprophyticus*) but was absent in *Staphylococcus xylosus*. Exfoliative toxins (ETs) play a significant role in cell-to-cell adhesion in the superficial epidermis, leading to the formation of blisters. These toxins contain a critical peptide bond that blocks their active sites, rendering them enzymatically inactive in their native state [40]. Although the *etb* (exfoliative toxin B) gene is rarely detected in CoNS isolates. In the present study, it was detected in two isolates (*Staphylococcus saprophyticus* and *Staphylococcus xylosus*) but was absent in the *Staphylococcus lentus* isolate. Coagulase-negative staphylococci are an important reservoir of antimicrobial-resistant genes as almost all CoNS had a multidrug-resistant profile [41]. Presence of *tsst*, *sea* and *etb* genes in the genome of CoNS isolates

may refer to indirectly increased their possibility of antibiotic resistance, as these virulent genes and antibiotic resistance genes are often found together due to their association with mobile genetic elements. This facilitates co-selection under antibiotic pressure, especially in clinical environments. In the present study CoNS isolates were completely resistant to clindamycin and highly resistant to trimethoprim-sulfamethoxazole, linezolid and erythromycin with 66.6%. No resistance was reported to doxycycline ciprofloxacin and gentamycin.

Pseudomonas aeruginosa causes high mortality in young birds due to omphalitis and yolk sac infections acquired in the hatchery, giving a mortality following yolk sac inoculation reached 100% mortality in experimentally inoculated chickens [42]. In the present study, the incidence of *P. aeruginosa* infection was 13.3% and this result was nearly constitute with results obtained by EL-Sawah et al. [43] who recovered *P. aeruginosa* with a percentage of 11%, Hassan [44] isolated *P. aeruginosa* with a percentage of 20% from one day old broiler chicks Although the bacterial isolation was carried out during the period from May to October, all *P. aeruginosa* isolates were recovered only during September and October and that may be due to the high humidity resulting from using cooling pads during summer months in breeder houses [45]. PCR test identified the presence of *Tox A*, *phzM* and *oprL* virulence genes in the four examined *P. aeruginosa* isolates recovered from diseased chicks. The *tox A* gene is recognized as a prevalent marker in *P. aeruginosa*, with studies by Algammal et al. [46] reporting that all tested isolates were positive for this gene. The *phzM* gene was detected in *P. aeruginosa* isolates from various clinical sources, with rates of 53.7% as reported by Ali et al. [47]. Additionally, the *oprL* gene is widely distributed in the *P. aeruginosa* genome, as it was reported in 100% of isolates in the studies of Algammal et al. [46].

Pseudomonas is naturally resistant bacteria to most antibiotics and able to quickly develop resistance to the commonly used ones [48], The multidrug resistant *P. aeruginosa* developed by several mechanisms such as multi-drug resistance efflux pumps, production of β -lactamase enzymes, biofilm formation and aminoglycoside modifying enzymes [49]. In the present study antimicrobial sensitivity test of *P. aeruginosa* isolates showed complete resistance towards imipenem, 75% resistant to ceftazidime, 25% resistance to gentamycin, and complete sensitive to amikacin, cefepime and ciprofloxacin.

Finally, we can conclude that the most common bacterial causes of omphalitis in chicks cause high mortality during the first 8 days of age are *E. coli*, *Salmonella*, staphylococci and *P. aeruginosa*. All these isolates have different virulence genes and

showed multidrug resistant (MDA) to different antimicrobial drugs.

Conclusion

The current study highlighted the role of virulent and multidrug resistant (MDR) *E. coli*, *Salmonella Enteritidis*, *Pseudomonas aeruginosa*, and coagulase-negative *Staphylococcus* spp., in omphalitis and early mortality in broiler chicks in the Delta region of Egypt. Ongoing monitoring of bacterial profiles and resistance patterns is crucial for developing appropriate treatment strategies and reducing economic losses in poultry production.

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Disclosure statement

No potential conflict of interest was reported by the author(s).

TABLE 1. Oligonucleotide primer sequences (Metabion, Germany).

Bacteria	Target gene	Sequence	Amplified product	Reference
<i>E. coli</i>	<i>iroN</i>	ATC CTC TGG TCG CTA ACT G CTG CAC TGG AAG AAC TGT TCT	847 bp	[16]
	<i>ompT</i>	TCATCCCGGAAGCCTCCCTCACTACTAT TAGCGTTTGCTGCACTGGCTTCTGATAC	496 bp	[17]
	<i>hlyF</i>	GGCCACAGTCGTTTAGGGTGCTTACC GGCGGTTTAGGCATTCCGATACTCAG	450 bp	
<i>Salmonella</i>	<i>fimA</i>	CCT TTC TCC ATC GTC CTG AA TGG TGT TAT CTG CCT GAC CA	85 bp	[18]
	<i>invA</i>	GTGAAATTATCGCCACGTTTCGGGCAA TCATCGCACCGTCAAAGGAACC	284 bp	[19]
	<i>Stn</i>	TTG TGT CGC TAT CAC TGG CAA CC ATT CGT AAC CCG CTC TCG TCC	617 bp	[20]
<i>Pseudomonas</i>	<i>toxA</i>	GACAACGCCCTCAGCATCACCAGC CGCTGGCCCATTCGCTCCAGCGCT	396 bp	[21]
	<i>oprL</i>	ATG GAA ATG CTG AAA TTC GGC CTT CTT CAG CTC GAC GCG ACG	504 bp	[22]
	<i>phzM</i>	ATGGAGAGCGGGATCGACAG ATGCGGGTTTCCATCGGCAG	875 bp	[23]
<i>Staphylococcus</i>	<i>sea</i>	GGTTATCAATGTGCGGGTGG CGGCACTTTTTTCTCTTCGG	102 bp	[24]
	<i>etb</i>	ACAAGCAAAGAATACAGCG GTTTTGGCTGCTTCTCTTG	226 bp	
	<i>tsst</i>	ACCCCTGTTCCCTTATCATC TTTTAGTATTTGTAACGCC	326 bp	

TABLE 2. Zone diameter interpretative standards for *E. coli* and *Salmonella* species

Antimicrobial drug	Susceptible	Susceptible dose dependent	Intermediate	Resistant
Ampicillin	≥ 17	—	14-16	≤ 13
Gentamycin	≥ 15	—	13-14	≤ 12
Amikacin	≥ 17	—	15-16	≤ 14
Cefepime	≥ 25	19-24	—	≤ 18
Cefotaxime	≥ 26	—	23-25	≤ 22
Doxycycline	≥ 14	—	11-13	≤ 10
Ciprofloxacin	≥ 31	—	21-30	≤ 20
Imipenem	≥ 23	—	20-22	≤ 19
Trimethoprim-Sulfamethoxazole	≥ 16	—	11-15	≤ 10

TABLE 3. Zone diameter interpretative standards for coagulase-negative *Staphylococcus*

Antimicrobial drug	Susceptible	Intermediate	Resistant
Erythromycin	≥ 23	14-22	≤ 13
Trimethoprim -Sulfamethoxazole	≥ 16	11-15	≤ 10
Linezolid	≥ 21	—	≤ 20
Clindamycin	≥ 21	15-20	≤ 14
Doxycycline	≥ 16	13-15	≤ 12
Ciprofloxacin	≥ 21	16-20	≤ 15
Gentamycin	≥ 15	3-14	≤ 12

TABLE 4. Zone diameter interpretative standards for *Pseudomonas aeruginosa*

Antimicrobial drug	Susceptible	Intermediate	Resistant
Ceftazidime	≥ 18	15-17	≤ 14
Ciprofloxacin	≥ 25	19-24	≤ 18
Cefepime	≥ 18	15-17	≤ 14
Amikacin	≥ 17	15-16	≤ 14
Gentamycin	≥ 15	13-14	≤ 12
Imipenem	≥ 19	16-18	≤ 15

TABLE 5. Morphology of colonies growth on specific media.

Bacteria Species	MacConkey agar	EMB Agar	XLD agar	Baird parker agar	Cetrimide agar
<i>E. coli</i>	Dry Pink colonies	Convex black colonies with green sheen	Large, flat, yellow colonies	Large brown colonies	No growth
<i>Salmonella Species</i>	Colorless colonies	-	Pink colonies with black center	No growth	No growth
<i>Pseudomonas aeruginosa</i>	Colorless colonies	-	Red flat colonies	No growth	Blue-green colonies
<i>Staphylococcus species</i>	No growth	-	No growth	Black colonies without opaque zone around	No growth

EMB agar: eosin methylene blue, XLD: xylose lysine deoxycholate agar

TABLE 6. biochemical tests results of suspected isolates recovered from specific media

Biochemical test	<i>E. coli</i>	<i>Salmonella species</i>	<i>Pseudomonas aeruginosa</i>	<i>Staphylococcus species</i>
Gram stain	-	-	-	+
Indole	+	-	-	NT
Methyl red	+	+	-	NT
Voges Proskauer	-	-	-	NT
Citrate utilization	-	-	+	NT
Urease	-	-	-	NT
Oxidase	-	-	+	-
Catalase	+	+	+	+
Coagulase	NT	NT	NT	-
TSI	A/A, gas +, H ₂ S -	K/A, gas +, H ₂ S +	K/K, gas -, H ₂ S -	NT

NT: Not Tested; A/A: Acid slant/acid butt; K/A: Alkaline slant/acid butt; K/K: Alkaline slant/alkaline butt; “+” positive reaction, “-” negative reaction

TABLE 7. Antimicrobial sensitivity test.

Bacteria isolates	<i>E. coli</i>				<i>S. Enteritidis</i>				<i>Staphylococcus spp.</i>			<i>P. aeruginosa</i>			
Sample no.	8	15	42	43	1	3	13	27	27	22	13	34	36	37	39
Ampicillin	R	R	R	R	R	R	R	R	-	-	-	-	-	-	-
Gentamycin	S	R	R	R	S	S	S	S	S	S	S	-	-	-	-
Amikacin	S	S	R	R	S	S	S	S	-	-	-	S	S	S	S
Cefepime	S	SDD	S	SDD	SDD	R	S	SDD	-	-	-	S	S	S	S
Cefotaxime	I	R	R	R	I	R	R	R	-	-	-	-	-	-	-
Ciprofloxacin	S	R	R	S	R	S	S	S	S	I	S	I	I	I	S
Imipenem	S	S	S	S	S	S	S	S	-	-	-	R	R	R	R
Trimethoprim-Sulphamethoxazole	S	R	R	S	R	S	S	S	R	R	S	-	-	-	-
Doxycycline	S	S	S	S	R	R	R	S	S	S	S	-	-	-	-
Ceftazidime	-	-	-	-	-	-	-	-	-	-	-	R	R	R	S
Erythromycin	-	-	-	-	-	-	-	-	I	R	I	-	-	-	-
Linezolid	-	-	-	-	-	-	-	-	R	R	S	-	-	-	-
Clindamycin	-	-	-	-	-	-	-	-	R	R	R	-	-	-	-

R, resistant; S, sensitive; I, intermediate; SDD, sensitive-dose dependent; (-) not tested

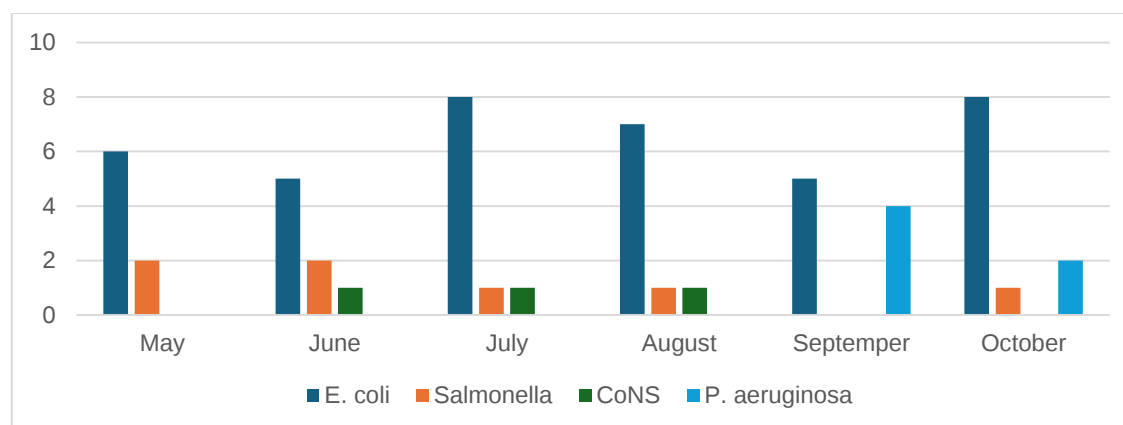


Fig. 1. *E. coli*, *Salmonella*, coagulase-negative *Staphylococcus* and *Pseudomonas aeruginosa* isolates number recovered in months from May to October



Fig. 2. Agarose gel electrophoresis (1%) stained with ethidium bromide showing PCR amplification of *hlyF*, *ompT*, and *iron* genes in *E. coli* isolates. L: 10 band DNA ladder from 100 to 1000bp; Lanes N: negative control; Lanes P: positive control; Lanes E1-E4: examined *E. coli* isolates showing positive amplification for the tested genes. Amplified products appeared at 450 bp for *hlyF*, 496 bp for *ompT*, and 847 bp for *iron* gene

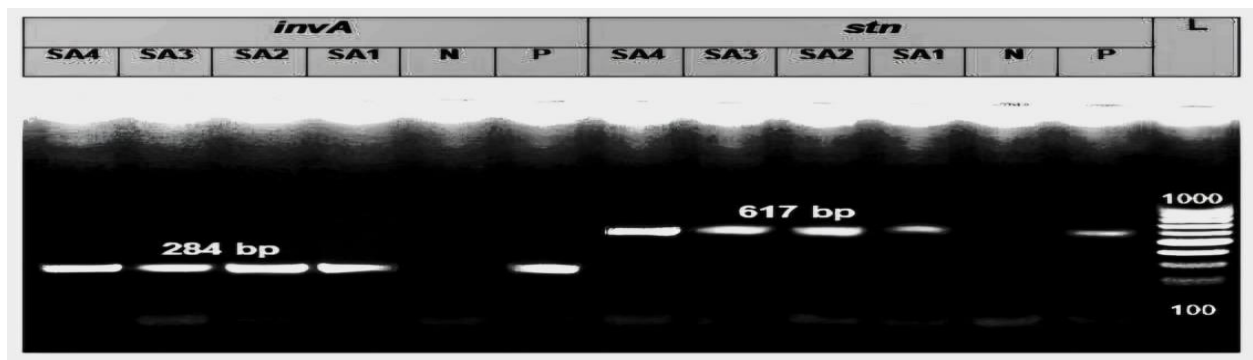


Fig. 3. Agarose gel electrophoresis (1%) stained with ethidium bromide showing PCR amplification of *invA* and *stn* genes in *Salmonella* isolates L: 10 band DNA ladder from 100 to 1000bp; Lanes N: negative control; Lanes P: positive control; Lanes SA1-SA4: examined *Salmonella* isolates showing positive amplification for the tested genes. Amplified products appeared at 284 bp for *invA*, 617 bp for *stn*



Fig. 4. Agarose gel electrophoresis (1%) stained with ethidium bromide showing PCR amplification of *fimA* gene in *Salmonella* isolates L: 17 band DNA ladder from 50 to 1500 bp; Lanes N: negative control; Lanes P: positive control; Lanes SA1-SA4: examined *Salmonella* isolates showing positive amplification for the tested gene. Amplified products appeared at 85 bp for *fimA* gene.

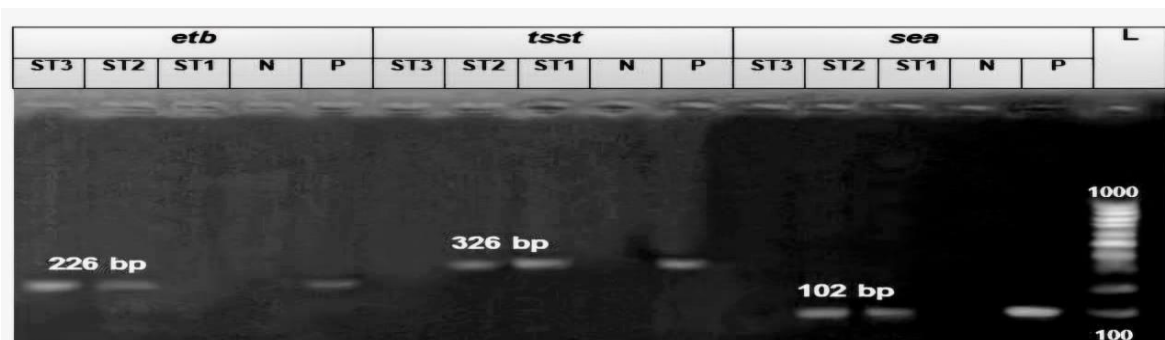


Fig. 5. Agarose gel electrophoresis (1%) stained with ethidium bromide showing PCR amplification of *etb*, *tsst* and *sea* genes in *Staphylococcus* isolates L: 10 band DNA ladder from 100 to 1000bp; Lanes N: negative control; Lanes P: positive control; Lanes ST1; ST2 and ST3: examined *Staphylococcus* isolates showing positive amplification for the tested genes. Amplified products appeared at 226 bp for *etb*, 326 bp for *tsst*, and 102 bp for *sea* gene

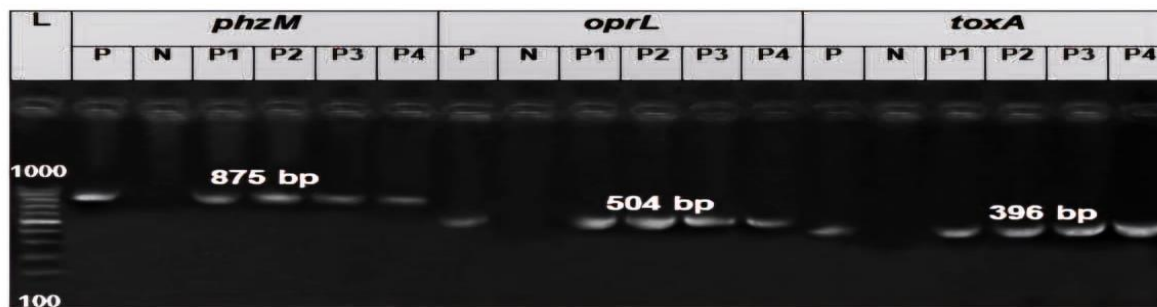


Fig 6. Agarose gel electrophoresis (1%) stained with ethidium bromide showing PCR amplification of *phzM*, *oprL* and *toxA* genes in *Pseudomonas aeruginosa* isolates. L: 10 band DNA ladder from 100 to 1000bp; Lanes N: negative control; Lanes P: positive control; Lanes P1-P4: examined *Pseudomonas aeruginosa* isolates showing positive amplification for the tested genes. Amplified products appeared at 875 bp for *phzM*, 504 bp for *oprL*, and 396 bp for *toxA* gene

References

- Rai, M. F., Khan, S. A., Asim Aslam, A. A. and Khalid Saeed, K. S. Effect of yolk sac infection in chicken. *Avian and Poultry Biology Reviews*, **16**(2), 87–93(2005).
- Oliveira, G. D. S., Pires, P. G. D. S., McManus, C., de Jesus, L. M., Santos, P. H. G. D. S. and Dos Santos, V. M. Plant extract in the control of poultry omphalitis. *Pathogens*, **13**(6), 438(2024).
- Khan, K. A., Khan, S. A., Aslam, A., Rabbani, M. and Tipu, M. Y. Factors contributing to yolk retention in poultry: A review. *Pakistan Vet. J.*, **24**, 46-50 (2004).
- Rezaee, M. S., Liebhart, D., Hess, C., Hess, M. and Paudel, S. Bacterial infection in chicken embryos and consequences of yolk sac constitution for embryo survival. *Vet. Pathol.*, **58**(1), 71-79(2021).
- Gordon, R. F. and Jordon, F. T. N. (1982). "Poultry Dis.", 2nd ed. Bailliere, Tindall, London, UK, pp: 6062.
- De Oliveira, A. L., Rocha, D. A., Finkler, F., de Moraes, L. B., Barbieri, N. L., Pavanelo, D. B., Winkler, C., Grassotti, T. T., de Brito, K. C. T., de Brito, B. G. and Horn, F. Prevalence of ColV Plasmid-Linked Genes and in Vivo Pathogenicity of Avian Strains of *Escherichia coli*. *Foodborne Pathogens and Disease*, **12**(8), 679–685 (2015).
- Mohamed, M. Y. I. and Habib, I. Virulence gene landscapes of *Salmonella* in Eastern and Southern Africa. *Frontiers in Microbiol.*, **16**, 1631550(2025).
- Rehman, T., Yin, L., Latif, M. B., Chen, J., Wang, K., Geng, Y., ... & Ouyang, P. (2019). Adhesive mechanism of different *Salmonella* fimbrial adhesins. *Microbial Pathogenesis*, **137**, 103748.
- Atchade, E., De Tymowski, C., Grall, N., Tanaka, S. and Montravers, P. Toxic shock syndrome: a literature review. *Antibiotics*, **13**(1), 96 (2024).
- Sachdeva, C., Satyamoorthy, K. and Murali, T. S. *Pseudomonas aeruginosa*: metabolic allies and adversaries in the world of polymicrobial infections. *Critical Reviews in Microbiol.*, **51**(4), 619-638(2025).
- Wang, K., Kai, L., Zhang, K., Hao, M., Yu, Y., Xu, X., Yu, Z., Chen, L., Chi, X. and Ge, Y. Overexpression of *phzM* contributes to much more production of pyocyanin converted from phenazine-1-carboxylic acid in the absence of RpoS in *Pseudomonas aeruginosa*. *Archives of Microbiology*, **202**(6), 1507–1515(2020).
- Castro-Vargas, R. E., Herrera-Sánchez, M. P., Rodríguez-Hernández, R. and Rondón-Barragán, I. S. Antibiotic resistance in *Salmonella* spp. isolated from poultry: A global overview. *Vet. World*, **13**(10), 2070–2084(2020).
- Quinn, P. J., Markey, B. K., Carter, M. E. and Carter, G. R. (2002). "Veterinary microbiology and microbial disease". Oxford, UK: Blackwell Science Ltd.
- Bahgat, O. T., Rizk, D. E., Kenawy, H. I. and Barwa, R. Prevalence of *E. coli* pathotypes: a comparative study between clinical and environmental isolates. *Egyptian Journal of Medical Microbiology*, **32**(3), 59-69(2023).
- Grimont, P. A. and Weill, F. X. Antigenic formulae of the *Salmonella* serovars. *WHO collaborating centre for reference and research on Salmonella*, **9**, 1-166(2007)..
- Ewers, C., Li, G., Wilking, H., Kießling, S., Alt, K., Antão, E. M. and Wieler, L. H. Avian pathogenic, uropathogenic, and newborn meningitis-causing *Escherichia coli*: how closely related are they. *International J. Med. Microbiol.*, **297**(3), 163-176 (2007).
- Johnson, T. J., Wannemuehler, Y., Doetkott, C., Johnson, S. J., Rosenberger, S. C. and Nolan, L. K. Identification of minimal predictors of avian pathogenic *Escherichia coli* virulence for use as a rapid diagnostic tool. *J. Clinical Microbiol.*, **46**(12), 3987-3996(2008)..
- Cohen, H. J., Mechanda, S. M. and Lin, W. PCR amplification of the *fimA* gene sequence of *Salmonella typhimurium*, a specific method for detection of *Salmonella* spp. *Applied and environmental microbiol.*, **62**(12), 4303-4308 (1996).
- Oliveira, S. D., Rodenbusch, C. R., Ce, M. C., Rocha, S. L. S. and Canal, C. W. Evaluation of selective and non-selective enrichment PCR procedures for *Salmonella* detection. *Letters in Applied Microbiol.*, **36**(4), 217-221(2003)..
- Murugkar, H. V., Rahman, H. and Dutta, P. K. Distribution of virulence genes in *Salmonella* serovars isolated from man & animals. *Indian J. Med. Res.*, **117**, 66-70(2003).

21. Matar, G. M., Ramlawi, F., Hijazi, N., Khneisser, I. and Abdelnoor, A. M. Transcription levels of *Pseudomonas aeruginosa* exotoxin A gene and severity of symptoms in patients with otitis externa. *Current Microbiol.*, **45**, 350-354(2002)..
22. Xu, J., Moore, J. E., Murphy, P. G., Millar, B. C. and Elborn, J. S. Early detection of *Pseudomonas aeruginosa* comparison of conventional versus molecular (PCR) detection directly from adult patients with cystic fibrosis (CF). *Annals C. Microbiol. and Antimicrobials*, **3**, 1-5(2004).
23. Finnan, S., Morrissey, J. P., O'gara, F. and Boyd, E. F. Genome diversity of *Pseudomonas aeruginosa* isolates from cystic fibrosis patients and the hospital environment. *J. C. Microbiol.*, **42**(12), 5783-5792(2004).
24. Mehrotra, M., Wang, G. and Johnson, W. M. Multiplex PCR for detection of genes for *Staphylococcus aureus* enterotoxins, exfoliative toxins, toxic shock syndrome toxin 1, and methicillin resistance. *J. C. Microbiol.*, **38**(3), 1032-1035(2000).
25. Sambrook, J., Fritsch, E. F. and Maniatis, T. (1989). "Molecular cloning: A laboratory manual" New York, NY. Cold Spring Harbor Laboratory Press.
26. Clinical and Laboratory Standards Institute CLSI. (2020). Performance standards for antimicrobial susceptibility testing (30th ed.; CLSI supplement M100). Clinical and Laboratory Standards Institute, 950 West Valley Road, Suite 2500, Wayne, Pennsylvania 19087, USA.
27. Shaheen, R., El-Abasy, M., El-Sharkawy, H. and Ismail, M. M. Prevalence, molecular characterization, and antimicrobial resistance among *Escherichia coli*, *Salmonella* spp., and *Staphylococcus aureus* strains isolated from Egyptian broiler chicken flocks with omphalitis. *Open Vet. J.*, **14**(1), 284–291(2024).
28. Pourhossein, Z., Asadpour, L., Habibollahi, H. and Shafighi, S. T. Antibacterial Resistance and Virulence Potential of Avian Colibacillosis-Causing *Escherichia coli* Isolates. *Infection Epidemiology and Microbiol.*, **8**(4), 307-315(2022)..
29. Sedeek, D. M., Rady, M. M., Fedawy, H. S. and Rabie, N. S. Molecular epidemiology and sequencing of avian pathogenic *Escherichia coli* APEC in Egypt. *Adv. Anim. Vet. Sci.*, **8**(5), 499-505 (2020).
30. Mohamed, L., Ge, Z., Yuehua, L., Yubin, G., Rachid, K., Mustapha, O. and Karine, O. Virulence traits of avian pathogenic (APEC) and fecal (AFEC) *E. coli* isolated from broiler chickens in Algeria. *Tropical Animal Health and Production*, **50**, 547-553(2018).
31. Poirel, L., Madec, J.-Y., Lupo, A., Schink, A. K., Kieffer, N., Nordmann, P. and Schwarz, S. Antimicrobial Resistance in *Escherichia coli*. *Microbiol. Spectrum*, **6**(4), 1128 (2018).
32. Kaboudi, K., Mamlouk, A., Romdhane, R. B., Khayech, M. and Bouzouaia, M. Gross pathology and bacteriological study of the yolk sac infections (omphalitis) in broiler chicks, North East Tunisia. *Revue Marocaine des Sciences Agronomiques et Vétérinaires*, **9**(3), 390-395 (2021).
33. Shen, X., Yin, L., Zhang, A., Zhao, R., Yin, D., Wang, J. and Liu, Y. Prevalence and characterization of *Salmonella* isolated from chickens in Anhui, China. *Pathogens*, **12**(3), 465 (2023).
34. Lozano-Villegas, K. J. and Rondón-Barragán, I. S. Virulence and Antimicrobial-Resistant Gene Profiles of *Salmonella* spp. Isolates from Chicken Carcasses Markets in Ibagué City, Colombia. *International J. Microbiol.*, **2024**(1), 4674138 (2024).
35. Eng, S. K., Pusparajah, P., Ab Mutalib, N. S., Ser, H. L., Chan, K. G. and Lee, L. H. *Salmonella*: A review on pathogenesis, epidemiology and antibiotic resistance. *Frontiers in Life Science*, **8**(3), 284–293 (2015).
36. Nasrin, S., Islam, M., Khatun, M., Akhter, L. and Sultana, S. Characterization of bacteria associated with omphalitis in chicks. *Bangladesh Veterinarian*, **29**(2), 63–68(2013).
37. Otto, M. *Staphylococcus* colonization of the skin and antimicrobial peptides. *Expert Review of Dermatology*. **5**(2), 183–195(2010).
38. Elbehiry, A. and Marzouk, E. *Staphylococci* in Livestock: Molecular Epidemiology, Antimicrobial Resistance, and Translational Strategies for One Health Protection. *Vet. Sci.*, **12**(8), 757(2025).
39. Nasaj, M., Saeidi, Z., Tahmasebi, H., Dehbashi, S. and Arabestani, M. R. Prevalence and distribution of resistance and enterotoxins/enterotoxin-like genes in different clinical isolates of coagulase-negative *Staphylococcus*. *European J. Med. Res.*, **25**, 1-11(2020).
40. Wiśniewski, P., Gajewska, J., Zadernowska, A. and Chajęcka-Wierzchowska, W. Identification of the enterotoxigenic potential of *Staphylococcus* spp. from raw milk and raw milk cheeses. *Toxins*, **16**(1), 17(2023)..
41. Silva, V., Caniça, M., Ferreira, E., Vieira-Pinto, M., Saraiva, C., Pereira, J. E., Capelo, J. L., Igrejas, G. and Poeta, P. Multidrug-Resistant Methicillin-Resistant Coagulase-Negative staphylococci in Healthy Poultry Slaughtered for Human Consumption. *Antibiotics*, **11**(3), 365(2022).
42. Lin, M. Y., Cheng, M. C., Huang, K. J. and Tsai, W. C. Classification, pathogenicity, and drug susceptibility of hemolytic gram-negative bacteria isolated from sick or dead chickens. *Avian Dis.*, 6-9 (1993).
43. El-Sawah, A. A., Hussien Dahshan, A. M., Nasef, S. A., El-Nahass, E.-S. and I, N. A. Characterization of *E. coli* and *Salmonella* spp. isolates associated with omphalitis in baby chicks. *J. Vet. Med. Res.*, **2016**(1), 61–70(2016).
44. Hassan, H.M. Characterization of *Pseudomonas aeruginosa* isolated from different pathological lesions in chickens. *M. V. Sc. Thesis*, (Microbiol.), Fac. Vet. Med., Beni-Suef Univ., (2013)

45. Elsayed, M. M., Abd, A. A., Tawab, E., Elhofy, F. I., and Rizk, A. M. Matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS) technique for identification of *Pseudomonas aeruginosa* isolated from broilers and humans. *Benha Vet. Med. J.*, **45**(2),1685 (2023).
46. Algammal, A. M., Eidaros, N. H., Alfifi, K. J., Alatawy, M., Al-Harbi, A. I., Alanazi, Y. F. and El-Tarabili, R. M. Opr 1 gene sequencing, resistance patterns, virulence genes, quorum sensing and antibiotic resistance genes of xdr *Pseudomonas aeruginosa* isolated from broiler chickens. *Infection and Drug Resistance*, 853-867(2023).
47. Ali, M. D. Y. and Abdulrahman, Z. F. Molecular identification, susceptibility pattern, and detection of some virulence genes in *Pseudomonas aeruginosa* isolated from burn patients. *Plant Archives*, **20**(1), 2573-2580 (2020).
48. Laborda, P., Hernando-Amado, S., Martínez, J. L. and Sanz-García, F. Antibiotic Resistance in *Pseudomonas*. *Advances in Experimental Medicine and Biology*, **1386**, 117–143(2022).
49. Zhao, Y., Xu, H., Wang, H., Wang, P. and Chen, S. Multidrug resistance in *Pseudomonas aeruginosa*: genetic control mechanisms and therapeutic advances. *Molecular Biomed.*, **5**(1), 62(2024).

الكشف عن بعض المسببات البكتيرية شديدة الضراوة لالتهاب كيس المح في دجاج التسمين

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الملخص

يُعد التهاب كيس المح من الأسباب الرئيسية للنفوق وضعف الأداء في كتاكيت التسمين. هدفت هذه الدراسة إلى الكشف عن بعض الأسباب البكتيرية المرتبطة بعدوى كيس المح في الكتاكيت. شملت الدراسة فحص 45 قطيعة من دواجن التسمين في منطقة الدلتا بمصر، تراوحت أعمارها بين 1 إلى 8 أيام، وكانت تعاني من معدلات نفوق مرتفعة، وخمول، وإسهال، وضعف في الحيوية. أسفر العزل البكتيري عن تحديد 39 معزولة من الإشريكية القولونية (86.6%)، و 7 معزولات من السالمونيلا (15.5%)، و 6 معزولات من السودوموناس (13.3%)، و 3 معزولات من المكورات العنقودية سالبة التلخثر (6.6%). أظهرت نتائج اختبار تفاعل البلمرة المتسلسل (PCR) للكشف عن الجينات الفاتكة أن جميع معزولات الإشريكية القولونية المختبرة احتوت على جينات *iroN* و *ompT* و *hlyF*، في حين احتوت معزولات السالمونيلا على جينات *invA* و *stn* و *fimA* كما أظهرت معزولات المكورات العنقودية نتائج متفاوتة من حيث وجود الجينات *sea* و *tsst* و *etb*. وتم الكشف عن جينات *toxA* و *phzM* و *oprL* في معزولات السودوموناس كما أظهرت اختبارات الحساسية للمضادات الحيوية مقاومة تامة لمعزولات الإشريكية القولونية تجاه الأمبيسلين، بينما كانت حساسة تمامًا للذوكسي سيكلين والإيمبيبينيم. كما أظهرت معزولات السالمونيلا مقاومة تامة للأمبيسلين، لكنها كانت حساسة للجنتاميسين والأميكاسين والإيمبيبينيم. أما معزولات المكورات العنقودية سالبة التلخثر فكانت مقاومة تمامًا للكليندامايسين، لكنها حساسة للذوكسي سيكلين، السيبروفلوكساسين، والجنتاميسين. وبيّنت معزولات السودوموناس مقاومة تامة للإيمبيبينيم، وحساسية كاملة تجاه الأميكاسين والسييفيم والسيبروفلوكساسين.

الكلمات الدالة: التهاب كيس المح، جينات الضراوة، بكتيريا متعددة المقاومة للمضادات الحيوية.