

MOLECULAR STUDIES ON SOME VIRULENCE FACTORS OF *PSEUDOMONAS AERUGINOSA* ISOLATED FROM CHICKENS AS A BIOFILM FORMING BACTERIA

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ABSTRACT

This study was aimed to isolate and identify *Pseudomonas aeruginosa* (*P. aeruginosa*) from 150 diseased broiler chickens in addition to 50 environmental swabs from water pipes and tanks to examine their susceptibility against some commonly used antimicrobial agents, in addition to detection of some virulence genes using Polymerase Chain Reaction (PCR) and evaluation of its ability to form biofilm *in vitro*. Clinically the affected chickens were subjected to postmortem examination (P.M) then samples from internal organs such as (liver, heart, lung, spleen, and intestine); tracheal and environmental swabs were collected and subjected to bacteriological examination and identification. Twenty *P. aeruginosa* isolates were recovered from the diseased chickens and environmental swabs with an incidence of (10%). Sixteen isolates were isolated from internal organs of 150 diseased chickens with an incidence of (10.66%); meanwhile 4 isolates were recoded from 50 water pipes and tanks swabs with a percentage of (8%). Antimicrobial sensitivity testing showed highly sensitivity to amikacin and colistin sulphate with percentages of (90%) for each of them. Doxycycline showed resistance with a percentage of (75%). PCR technique was a good tool for testing three virulence genes; *pslA*, *pelA* and *fliC* genes; the three genes were detected in of the examined samples with a percentage of (100%). A significant relationship between the existence of three virulence genes in the isolated *P. aeruginosa* and ability of biofilm formation (Slime producing ability) was reported in this study.

Key words: *Pseudomonas aeruginosa*, virulence genes, biofilm, chicken

INTRODUCTION

P. aeruginosa is a serious poultry pathogen and zoonotic bacterial agent causing nosocomial infections (Elsayed *et al.*, 2016); it's associated with heavy economic losses in broiler farms (Devriese *et al.*, 1975) causing significant morbidity and mortality (Hancock and Speert, 2000).

P. aeruginosa is a motile Gram negative rod shaped bacteria belonged to family *Pseudomonadaceae* (Blanc *et al.*, 1998); grows readily on common bacteriologic media and producing a water-soluble green pigment composed of fluorescein and pyocyanin (Barnes, 2003). It's a ubiquitous organism usually associated with soil, drinking water, and humid environments (Mena and Gerba, 2009).

Birds at any age may be infected; young birds are most susceptible. Severely stressed or immunodeficient birds and concurrent infections with viruses and other bacteria enhance susceptibility to *Pseudomonas* infection (Stipkovits *et al.*, 1993). *P. aeruginosa* produce a variety of toxins and enzymes that may contribute to its pathogenicity (Lin *et al.*, 1993), birds infected with *P. aeruginosa* suffer from septicemia, respiratory infections, sinusitis, keratoconjunctivitis and embryonic death rates in hatcheries (Hai-ping, 2009). It also produces cheesy deposits in serous surfaces lining air sacs and peritoneal cavity in addition to congestion of internal organs, perihepatitis and pericarditis (Riad, 1994).

Biofilms are intricate bacterial communities found attached to living or abiotic surface and surrounded by a bacterially produced extracellular matrix composed of exopolysaccharides, DNA and proteins (Whitchurch *et al.*, 2002); Alginate, Psl and Pel are three exopolysaccharides that considered the main components in biofilm matrix (Al-Wrafy *et al.*, 2016), these polysaccharides are produced by *P.*

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aeruginosa and determine its ability to form biofilm structure (Ghafoor *et al.*, 2011).

pslA and *pelA* genes play an important role in formation of carbohydrate-rich structure of biofilm matrix (Cotton *et al.*, 2009). *pel* genes were identified in *P. aeruginosa* (PA14 strain) and its required to produce a glucose-rich matrix material that involved in a thick pellicle formation (Vasseur *et al.*, 2005).

Flagella mediate swimming and swarming motility of *P. aeruginosa* which act as an adhesin and produce cell-to-surface interactions then enable the bacteria to swim and overcome repulsive electrostatic forces that present between the cell surface and the surface of materials forming film (O'Toole and Kolter, 1998). The *fliC* gene as a key gene involved in flagella production, has an important role in the encoding of subunit protein, flagellin (FliC) (Wolfgang *et al.*, 2004).

Several factors are required for biofilm development, including attachment via adhesive protein, cell aggregation via proteins, extracellular DNA, polysaccharides and cell motility (Kjelleberg and Givskov, 2007). The ability of *P. aeruginosa* to metabolize a variety of nutrients enables them to survive in a variety of habitats by developing biofilms (Meliani and Bensoltane, 2015). Within biofilms microorganisms are generally well protected against influence of antibiotics, disinfectants (Hoiby *et al.*, 2010) and host immune system (Jensen *et al.*, 2010). The ability of *P. aeruginosa* to form biofilm renders its ineffective clearance by immune defense systems and antibiotherapy (Rasamiravaka *et al.*, 2015), that considered as a protective mode that allows bacteria to survive and colonize in hostile environments (Yang *et al.*, 2011).

The aim of this study was to isolate and identify *P. aeruginosa* from diseased broiler chickens and environmental swabs collected from water pipes and tanks. Also examine the susceptibility of this microbe against some common usable antimicrobial agents in addition to detection of some virulence genes using PCR technique and evaluation of its ability to form biofilm in vitro.

MATERIALS AND METHODS

1. Samples collection

One hundred and fifty diseased broiler chickens (Cobb breed) were collected from 40 farms in Dakahlia Governorate at 30 days of age. The collected birds suffered from depression, anorexia with lameness, swollen hock joint, diarrhea and

necrotic foci in (liver, spleen and kidney). The mortality rate ranged from (4%) to (10%) Samples from internal organs such as (liver, heart, lung, spleen, intestine and tracheal swabs were collected aseptically and pooled together for each bird. A total of 50 swabs from water pipes and tanks in chicken farms were also collected aseptically. All samples were labeled and transported directly in ice box to reference laboratory for veterinary Quality Control on poultry production (Dakahlia branch- Gamasa lab.) for further examinations.

2. *P. aeruginosa* isolation and identification

The collected samples were cultured in nutrient broth tubes and incubated at 37°C for 24 hours then loopfull from broth was streaked on nutrient agar, MacConkey and blood agar plates then incubated at 37 °C for 24 hours aerobically. The suspected colonies were transferred to nutrient agar slant and stored in refrigerator at 4 °C for further identifications (colony morphology, pigment production and biochemical reactions which were preformed following the methods described by MacFadden, (2000).

3. Antimicrobial susceptibility testing according to Finegold and Martin (1982).

Antimicrobial susceptibility testing was done by agar disc diffusion method using Muller Hinton agar plates on isolates that confirmed to be *P. aeruginosa*. The used antibiotics were doxycycline (30 µg), streptomycin (10 µg), amikacin (30 µg), gentamycin (10 µg), cefotaxime (30 µg), ceftazidime (30 µg) and colistin sulphate (10µg). The antibiotics were categorized into resistant, intermediate and susceptible categories according to Clinical Laboratory Standards resistance according to (CLSI, 2016).

4. Detection of *pslA*, *pelA* and *fliC* genes in the isolated *P. aeruginosa*

4.1. DNA extraction

DNA was extracted from samples using QIAamp DNA Mini kit (Qiagen, Germany, GmbH) with modifications from the manufacturer's recommendations. Briefly, 200 µl of the sample suspension was incubated with 10 µl of proteinase K and 200 µl of lysis buffer at 56°C for 10 min. After incubation, 200 µl of 100% ethanol was added to the lysate. The sample was then washed and centrifuged following the manufacturer's recommendations. Nucleic acid was eluted with 100 µl of elution buffer.

4.2. Oligonucleotide primers that used were provided from Metabion (Germany) listed in table (1).

Table (1): Primers sequences, target genes, amplicon sizes and cycling conditions.

Gene	Primer sequence	Seg. (bp)	P. dent.	Amplification 35 cycles			Final ext.	References
				Sec. dent.	Annealing	Extens.		
<i>pslA</i>	F:5'TCCCTACCTC AGCAGCAAGC-3'	656	94°C 5 min.	94°C 30 sec.	60°C 40 sec.	72°C 45 sec.	72°C 10 min.	Ghadaksaz <i>et al.</i> , (2015)
	R:5'- TGTTGTAGCCGTA GCGTTTCTG-3'							
<i>pelA</i>	F:5'- CATACCTTCAGCC ATCCGTTCTTC-3'	786	94°C 5 min.	94°C 30 sec.	60°C 40 sec.	72°C 45 sec.	72°C 10 min.	
	R:5'- CGCATTGCGCCGA CTCAG-3'							
<i>fliC</i>	F:5'- TGAACGTGGCTA CCAAGAACG -3'	180	94°C 5 min.	94°C 30 sec.	56.2°C 30 sec.	72°C 30 sec.	72°C 7 min.	
	R:5'- TCTGCAGTTGCTT CACTTCGC -3'							

Seg. (bp)= amplified segment& P. dent.= primary denaturation& Sec. dent.= secondary denaturation & Extens= extension& Final ext.= final extension

4.3. PCR amplification

Primers were utilized in a 25- µl reaction containing 12.5 µl of EmeraldAmp Max PCR Master Mix (Takara, Japan), 1 µl of each primer of 20 pmol concentrations, 4.5 µl of water, and 6 µl of DNA template. The reaction was performed in an Applied biosystem 2720 thermal cycler.

4.4. Analysis of PCR products

The products of PCR were separated by electrophoresis on 1.5% agarose gel (Applichem, Germany, GmbH) in 1x TBE buffer at room temperature using gradients of 5V/cm (1.5 gm agarose was prepared in 100 ml TBE buffer and heated in microwave then allowed to cool at 70°C, then 0.5 µg/ml ethidium bromide was added and mixed thoroughly. The warm agarose was poured directly in gel casting apparatus with desired comb in apposition and left at room temperature for polymerization). For gel analysis, 20 µl of the products was loaded in each gel slot. A gelpilot 100 bp DNA Ladder (Qiagen, Germany, GmbH) and generuler 100 bp ladder (Fermentas, Germany) were used to determine the fragment sizes. The gel was photographed by a gel documentation system (Alpha

Innotech, Biometra) and the data was analyzed through computer software.

5. Detection of biofilm formation in the isolated *P. aeruginosa* using qualitative tube method

According to Christensen *et al.* (1982) and Maram, (2011) with some modifications; a loopful of test organisms was inoculated in 10 mL of trypticase soy broth with 1% glucose in test tubes. The tubes were incubated at 37°C for 24 hours. One ml from the inoculated broth was transferred into another tube containing 4 ml trypticase soy broth with 1% glucose, one tube used as a control negative (not inoculated) and another tube was inoculated with *P. aeruginosa* (positive control). All test tubes were incubated at 37°C for 5 days. After incubation, tubes were decanted and washed with phosphate buffer saline (pH 7.3) and dried. Tubes were then stained with crystal violet (0.1%). Excess stain was washed with deionized water. Tubes were dried in inverted position. The results of tube method were compared with the control positive strain. Biofilm formation was considered positive when a visible film lined the wall and the bottom of the tube. The experiment was performed in triplicate and repeated three times.

RESULTS

1- Cultural, morphological, staining characteristics and biochemical reactions of the isolated *P. aeruginosa*

The cultural characteristics of *P. aeruginosa* isolates were large irregular translucent colonies with a greenish diffusible pigment with fruity smell. The isolated micro-organism was Gram negative, motile, rod shape and produce β hemolysis on blood agar. Biochemical reactions of the examined *P. aeruginosa* were positive for oxidase, catalase, citrate utilization and urease tests meanwhile negative for indole, methyl red and Voges – proskauer tests.

2 - Incidence of *P. aeruginosa* among the examined samples

A total of 20 *P. aeruginosa* isolates were isolated from 200 samples (150 diseased chickens and 50 water pipes and tanks swabs) with an incidence of (10 %). Sixteen isolates were recorded from internal organs of 150 diseased chickens with an incidence of (10.66%) (16 isolates/150 samples of internal organs); meanwhile 4 isolates were recoded from 50 water pipes and tanks swabs with a percentage of (8 %).

3 - Antimicrobial Susceptibility pattern of the isolated *P. aeruginosa*

The results of antimicrobial susceptibility testing in table (2) revealed that *P. aeruginosa* isolates were highly sensitive to amikacin, colistin sulphate and gentamycin with percentages of (90%), (90%), (90%), (70%) and (65%) respectively. Doxycycline, ceftazidime and streptomycin showed resistance with a percentages of (75%), (65%), (60%) and (50%) respectively.

Table (2): Antimicrobial Susceptibility pattern of the most frequently isolated *P. aeruginosa* isolates.

Antimicrobial agent	<i>P. aeruginosa</i> (20 isolates)		
	Sensitive NO. (%)	Intermediate NO. (%)	Resistant NO. (%)
Amikacin	18 (90%)	1 (5%)	1 (5%)
Cefotaxime	3 (15%)	12 (60%)	5 (25%)
Ceftazidime	2 (10%)	6 (30%)	12 (60%)
Colistin sulphate	18 (90%)	0 (0%)	2 (10%)
Doxycycline	2 (10%)	3 (15%)	15 (75%)
Gentamycin	13 (65%)	4 (20%)	3 (15%)
Streptomycin	7 (35%)	3 (15%)	10 (50%)

*Percentage calculated by dividing number of *Pseudomonas aeruginosa* isolates showed resistance or sensitivity on total number of *Pseudomonas aeruginosa* isolates

4 - Detection of *pslA*, *pelA* and *fliC* genes in the examined *P. aeruginosa* isolates

From figures (1, 2 and 3), *pslA*, *pelA* and *fliC* genes were detected in all of the examined samples with percentage of (100%).

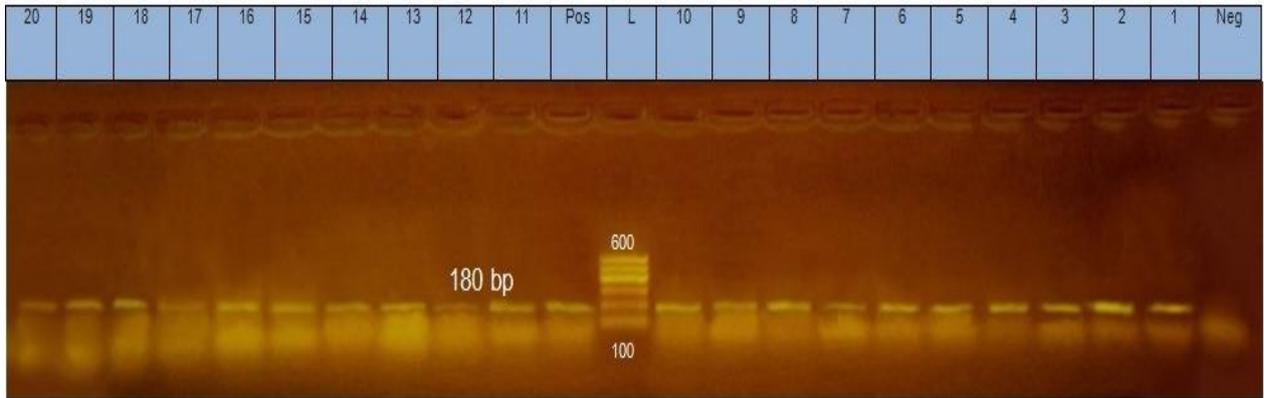


Figure (1): Agarose gel electrophoresis of PCR products for *P. aeruginosa* isolates to detect *fliC* gene in genomic DNA. Lane L: 100-600 bp DNA ladder. Neg: Negative control, Pos: Positive control. Lane: 1 to 20 were positive samples

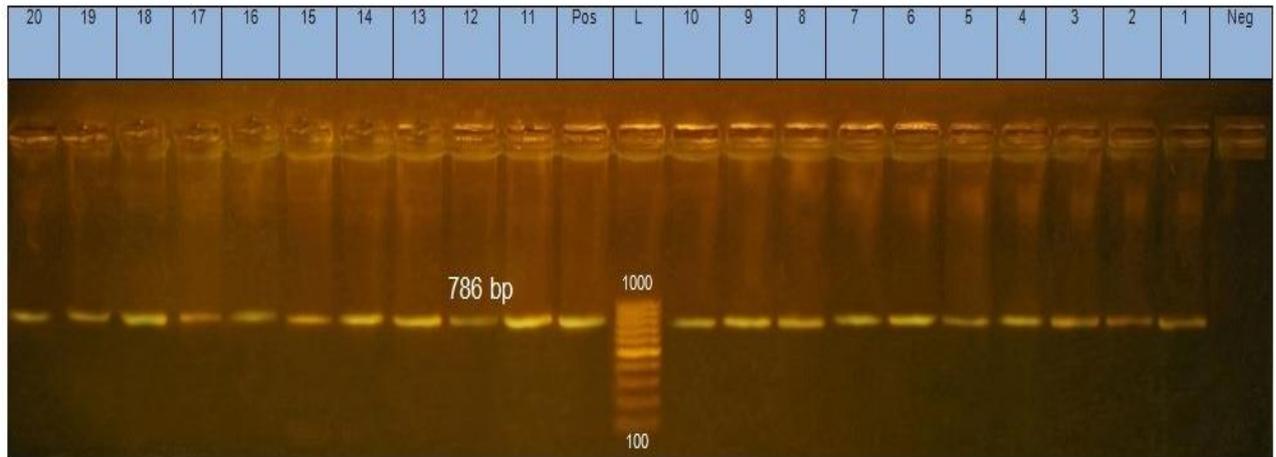


Figure (2): Agarose gel electrophoresis of PCR products for *P. aeruginosa* isolates to detect *pelA* gene in genomic DNA. Lane L: 100-1000 bp DNA ladder. Neg: Negative control, Pos: Positive control. Lane: 1 to 20 were positive samples

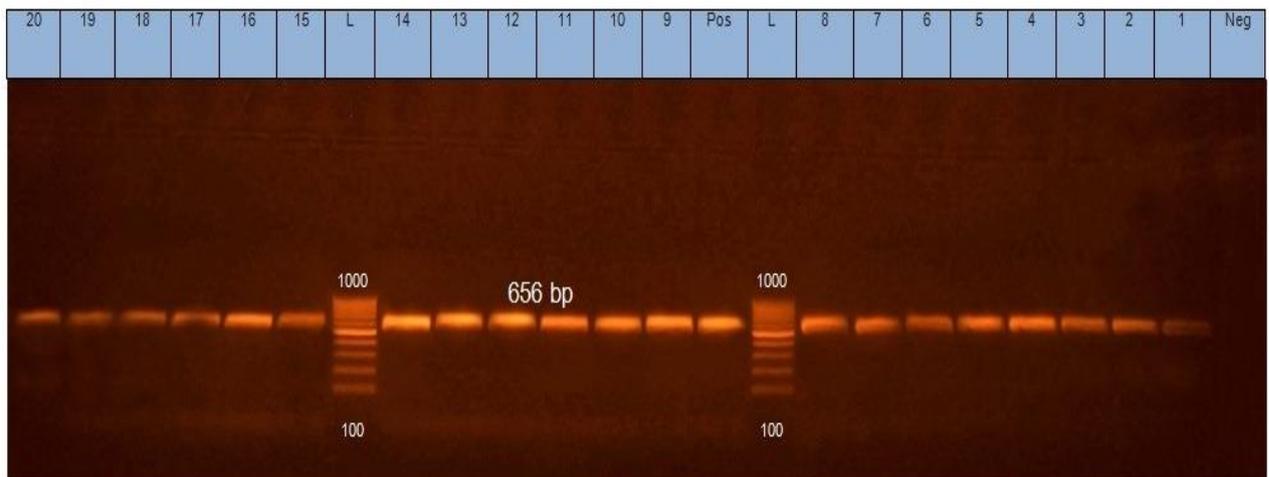


Figure (3): Agarose gel electrophoresis of PCR products for *P. aeruginosa* isolates to detect *psIA* gene in genomic DNA. Lane L: 100-1000 bp DNA ladder. Neg: Negative control, Pos: Positive control. Lane: 1 to 20 were positive samples

5-Biofilm formation in the examined *P. aeruginosa* isolates

A total of 20 *P. aeruginosa* isolates were examined for biofilm formation and all of them were totally

positive with a visible film lined the wall and the bottom of the test tubes. The experiment was performed in triplicate and repeated three times.

6-Correlation between biofilm formation, virulence factors and antibiotic resistance in the examined *P. aeruginosa* isolates

All of the isolated *P. aeruginosa* in this study were found to produce biofilm by using a qualitative tube method. Some virulence factors that have an important role in biofilm formation such as presence of exopolysaccharide and flagella were examined using PCR technique. The percentages of (*pslA* and *pelA*) genes were (100%) of the examined strains. The results in this study explain the significant role of such genes in facilitating the attachment of *P. aeruginosa* to inert or living surfaces and enclosed in an extracellular polysaccharide matrixes forming biofilm.

Flagellum plays an extremely important role in production and assembly of biofilm formation. *fliC* gene that encodes for presence of flagella was detected in this study with a percentage of (100%).

In the current study *pelA* and *pslA* genes that encodes for these exopolysaccharides were detected and this explains the occurrence of resistance against some of the used antibiotics.

DISCUSSION

In this study *P. aeruginosa* was isolated from internal organs of diseased chickens with a percentage of (10.66%) which was nearly coordinated with Sidhom, (2007) and Omran, (2012) who isolated it from chickens with a percentage of (13.34%) and (12%) respectively.

Results in this study varied to some extent from Abd El- Tawab *et al.* (2014) who reported *P. aeruginosa* with a percentage of (6.25%) from 224 samples collected from diseased chickens; Younes *et al.* (1990) who isolated *P. aeruginosa* from 20 dead chickens with percentage of 4.9%. several authors reported a higher incidence of isolation than this study such as Farhan, (2006) who isolated *P. aeruginosa* from infected chickens with an incidence of (20.5%); Kamel *et al.* (2011) who isolated *P. aeruginosa* from poultry respiratory tract with a percentage of (31%).

The results of antimicrobial susceptibility testing in table (2) revealed that *P. aeruginosa* isolates were highly sensitive to amikacin, ciprofloxacin, colistin sulphate, norfloxacin and gentamycin with percentages of (90%), (90%), (90%), (70%) and (65%) respectively. Doxycycline, penicillin, ceftazidime and streptomycin showed resistance with a percentages of (75%), (65%), (60%) and (50%) respectively.

A study performed by Abd El-Gawad *et al.* (1998) reported the same results of the present study; that *P. aeruginosa* isolates from chickens were highly sensitive to colistin sulphate and amikacin. The obtained results agreed with Hassan, (2013) who said that *P. aeruginosa* isolated from chickens showed high sensitivity to amikacin. The results in this study were nearly coordinated with to Abd El-Tawab *et al.* (2014) who reported that *P.aeruginosa* isolates were sensitive to colistin sulphate and norfloxacin; Kurkure *et al.* (2001) who stated that *P. aeruginosa* isolated from broiler were sensitive to gentamycin and ciprofloxacin in a percentage (88.57%) and (62.85%) respectively.

Unlike to our results that reported a higher sensitivity to amikacin and colistin sulphate; a study conducted by El Gohary, (2004) who reported that norfloxacin and ciprofloxacin showed a higher effect of sensitivity on *P. aeruginosa* while amikacin showed a moderate effect.

pslA, *pelA* and *fliC* genes were detected in all of the examined samples with percentage of (100%). These results varied greatly from that recorded by Ghadaksaz *et al.* (2015) who reported *pslA*, *pelA* and *fliC* genes with percentages of (45.2%), (83.7%) and (70.2%) respectively in *P. aeruginosa* isolated from clinical samples.

The results in this study reported that *P. aeruginosa* has the ability to form biofilm and this is also mentioned by several authors such as Price and Ahearn, (1988); Hoiby *et al.* (2010); Bjarnsholt, (2013); Meliani and Bensoltane, (2015) and Rasamiravaka *et al.* (2015).

Some virulence factors of *P. aeruginosa* that have an important role in biofilm formation such as presence of exopolysaccharide and flagella were examined using PCR technique. The percentages of (*pslA* and *pelA*) genes were (100%) of the examined strains. The results in this study explain the significant role of such genes in facilitating the attachment of *P. aeruginosa* to inert or living surfaces and enclosed in an extracellular polysaccharide matrixes forming biofilm. Some authors such as Cotton *et al.* (2009) explained the important role of *pslA* and *pelA* genes in formation of carbohydrate rich structure of biofilm matrix, any mutations in these genes cause deficiency in the biofilm formation ability.

Flagellum plays an extremely important role in production and assembly of biofilm formation. *fliC* gene that encodes for presence of flagella was detected in this study with percentage of (100%). This *fliC* gene is one of the key genes in flagella production that have a role in subunit protein encoding, flagellin (*fliC*) (Feldman *et al.*, 1998), the

activities of such flagella and type IV pili play a role in biofilm formation (Pratt and Kolter, 1999). In this study there is a significant relationship between the existence of *fliC* gene in the isolated *P. aeruginosa* and the important role that this gene plays in biofilm formation. The same findings were reported by Ghadaksaz *et al.* (2015) who reported also such significant relationship.

Biofilms are inherently resistant to antibiotics due to their failure to fully diffuse through the biofilm, but some antibiotics are more effective in penetrating and killing cells than others (Stewart, 2001). This explains the variation in antibiotic susceptibility of the used antibiotics in this study, slow diffusion of antibiotics allowing them to be deactivated in the outer layers of the biofilm leading to antibiotic resistance. However, a study performed by Lederberg, (2000) mentioned that antibiotics fail to kill the bacterium in the biofilm even after antibiotic therapy is completed.

The exopolysaccharides (Psl and Pel) form the biofilm matrix of *P. aeruginosa* which protect the bacterial cell from antibiotics and the immune system (Al-Wrafy *et al.*, 2016). In the current study *pelA* and *pslA* genes that encodes for these exopolysaccharides were detected and this explains the occurrence of resistance against some of the used antibiotics. Other mechanisms of resistance may also exist including antibiotic modifying enzymes, limited membrane permeability for the antibiotics, acquisition of chromosomally or plasmid encoded antibiotic resistance genes, mutations (Potron *et al.*, 2015).

In this study the resistance of *P. aeruginosa* to the used aminoglycosides antibiotics is explained by Stewart, (2001) who mentioned that oxygen is often completely consumed by the surface layers of biofilm and creating anaerobic conditions in the less exposed biofilm regions; these anaerobic conditions are often unaffected by aminoglycoside antibiotics. So, the accumulation of waste products produce pH differences, the acidity prevents the antibiotic from killing all of bacterial in biofilm.

CONCLUSION

P. aeruginosa is an opportunistic pathogenic bacterium responsible for serious problems in poultry farms. It considered as a good example of environment associated bacteria. Beside its natural resistance to many antimicrobial agents and conventional disinfectants, it has ability to form biofilm. There is increasing evidence that biofilm-mediated infection facilitates the development of chronic infectious diseases and recurrent infections due to the inherently resistance of biofilm to antimicrobial agents. Non microbicidal strategies

should be applied to struggle biofilm formation such as avoiding microbial attachment to a surface, disrupting biofilm development to enhance the penetration of antimicrobials. Alternative methods of treatment should be applied such as phage therapy. Poultry farms should take all stringent measures of management to overcome any possible sources of *pseudomonas* infection.

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دراسات جزيئية على بعض عوامل الضراوة للسيدوموناس ايريجنوزا المعزولة من الدجاج كبيكتيريا مكونة للبيوفيلم

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تهدف هذه الدراسة الي عزل وتوصيف ميكروب السيدوموناس ايريجنوزا من عدد ١٥٠ طائر من دجاج التسمين المريض بالاضافة الي ٥٠ مسحة من مواسير المياه والخزانات ومن ثم فحص حساسيتها لمضادات الميكروبات الشائعة الاستخدام وايضا فحص بعض عوامل الضراوة باستخدام اختبار تفاعل البلمرة المتسلسل وتقييم قابليتها لتكوين البيوفيلم. تم اجراء التشريح المرضي للدجاج محل الدراسة واخذ عينات لاجراء الفحص البكتيريولوجي من الاعضاء الداخلية مثل (الكبد- القلب- الرئتين- الطحال- الامعاء) وكذلك مسحات من القصبة الهوائية وخزانات ومواسير المياه. تم تسجيل ٢٠ معزولة من السيدوموناس ايريجنوزا بنسبة (10%) . تم عزل 16 معزولة من من الدجاج المريض بنسبة (10.66%) بينما تم عزل 4 معزولات من مواسير المياه والخزانات بنسبة (8%). باجراء اختبارات الحساسية وجد ان المعزولات محل الدراسة حساسة للاميكاسين- الكولستين سلفات بنسب (٩٠%) لكل منهما بينما اظهرت المعزولات مقاومة للدوكسي سيكلين بنسبه (٧٥%). تم الكشف عن بعض عوامل الضراوة باستخدام اختبار تفاعل انزيم البلمرة المتسلسل حيث تواجدت جينات (*fliC*, *pelA*, *pslA*) بنسبة (١٠٠%). تبين في هذه الدراسة وجود علاقة ذات اهمية بين تواجد عوامل الضراوة في السيدوموناس ايريجنوزا وقابليتها لتكوين البيوفيلم.