

Antibiotic Effect on *Pseudomonas Aeruginosa* Isolated From Patient In Hila City

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Abstract

In this study, a total of 400 clinical samples were collected from patients different hospitals in hilla city((13.51%) Margin teaching hospital, (27.03%)Al-Hilla teaching hospital,(43.24%) Babylon hospital for maternity and pediatric and (16.22%)Chest Disease Center) , (37) isolates were identified as to *P. aeruginosa*, from 9 (18%) isolates from burn specimen, 7 (14%) isolates from otitis , 5 (10%)isolates from UTI , 4 (8%) isolates from wound , 3(7.5%) was isolated from blood specimen where as 6 (6%)was isolated sputum and only 3 (5%)isolates were detected from skin. The period of collection was extended from January 2019 to June 2019. Also antibiotic susceptibility were preformed using disk employed diffusion and agar utilizing dilution processes. Furthermore , the resistance of isolate to a variety of antibiotics has been investigated and found that these isolate have been resistance to more than one antibiotic : 37(100%) resistance to Cloxacilin , 37(100%) to Amoxicillin /Clavulanic acid, 37(100%) to cefotaxime, 36(97%) to ceftizoxone, 34(92%) to Ceftazidime, 14(37%) to Levofloxacin, 18(49%) to Gentamicin, , 3(8%) to Norfloxacin and 1(3%) to Azetronam.

Keywords: *Pseudomonas aeruginosa*, Different sources, Antibiotic susceptibility, Resistant, Gram-negative bacteria.

Introduction

Pseudomonas aeruginosa is a Gram-negative bacterium, an aerobe rod belongs to the Pseudomonadaceae bacterial family and so taxonomic classification relies on preserved substances, including 16srRNA which is a range of members of the *Pseudomonas* genus, divided into eight classes¹⁷. *Pseudomonas aeruginosa* infection allowed the incorporation of the constituents of the outer membrane, namely secretary toxins like Lipopolysaccharides (LPs), general endotoxin, which is a protein, toxoids, mutant reacting toxins or nontoxic substances, Pilli, and Flagella^{23,35}. *P. aeruginosa* allows β -Lactamases; enzymes to deactivate the antibiotics by hydrolyzing the peptide attachment of the Ring β -Lactam. *P.aeruginosa* is capable of producing numerous β -lactamases, such as extended-spectrum β -lactamases (ESBL), chromosomal cephalosporinase (AmpC), and metallo- β -lactamases (MBL). Different kinds of MBLs were previously identified and brought on integrated products. Earlier

researches have shown that high levels of morbidity and mortality are connected with MBL-generating *P.aeruginosa* in bloodstream diseases. the ampC gene mutation generates such resistance which unlike other β -lactamases^{1,9,10,13}.

Material and Methods

1- Patients:

A total of 37 *Pseudomonas aeruginosa* isolates were collected from different clinical specimens in Babylon province during the period from January to June 2019. These specimens were collected from 400 specimens obtained from inpatients suffering from different infections by taking swabs from Burn, wound, ear, UTI, blood, urine ,sputum and skin. MacConkey and Nutrient agars were utilized in aerobic conditions with a temperature at 42°C for 24 to 48hrs. Later, the bacteria were characterized using traditional biochemical tests depending on protocols from MacFaddin, (2000).

2- Antibiotic susceptibility tests:

The susceptibility of *P. aeruginosa* isolates were determined by disk diffusion method (Kirby–Bauer standardized disk method).

The Mueller–Hinton medium was employed for this test. The medium was cooled to 45–50°C and with a sterile wire loop, the 4–5 pure colonies were transferred to a tube containing broth at 5ml of BHI 37°C-incubated until its turbidity standard. This usually required at least 4–6 hours incubation. The cells density was compared with McFarland standard tube No. 0.5. Sterile swab made of wood and cotton was standardized-suspension-dipped streaking it onto the Mueller–Hinton medium dry surface with three different directions to obtain an even distribution of the inoculums. Using a flat and even surface, the plates were let to stay without distribution

for 3 to 5mins for better dryness. The disks were inserted on the surface using gentle pressing with a sterile forceps within 15mins and incubated for 18hrs at 37°C with inverted position. The inhibition zones later were ruler-measured (millimeter) using CLSI (2012) as guidelines for deciding the sensitivity or resistance of the bacteria to the antibiotics used.

Result and Discussion

1) Isolation of *Pseudomonas aeruginosa*

During the period of study, 400 hospital-patient-based specimens were collected in hilla city, only 37 (9.25%) isolates from *P. aeruginosa* bacterium, 9 (18%) isolates burn specimen, 7 (14%) from otitis, 5 (10%) isolates, 4 (8%) isolates from wound; 3 (7.5%) was isolated from blood specimen whereas 6 (6%) was belong to sputum and only 3 (5%) isolates were detected from skin. The outcomes were identified, table (1).

Table (1): *P. aeruginosa* isolate frequency from different clinical specimens.

Specimen source	No. of Specimen	No. (%) of <i>P.aeruginosa</i>
Burn	50	9 (18%)
Wound	50	4(8%)
Sputum	100	6(6%)
UTI	50	5(10%)
Otitis	50	7(14%)
Blood	40	3(7.5%)
Skin	60	3(5%)
Total	400	37(9.25%)

The second Gram negative HAP bacterium isolated from health care centers is *P. aeruginosa* with leveled up incidence and fatality rates^{22,27} with highly risky illnesses, such as septic burns, in patients with low defense of the immune system^{11,34}.

The *P. aeruginosa* incidence findings, here, assured 9.25 % (37/400) of the hospital samples with lower rates

than that, 39.1%, uncovered by Okon et al. ,2009 from Nigerian patients with infected wounds and, 25.5%, revealed by Ndip et al., (2005) in Cameron, in Al-Sulaimania City Hospital, Iraq, at 17.85% , and in Egypt at 18.6%. Sample size and geographical distribution may have induced variations in the collected findings^{30,14,15}.

P. aeruginosa is well-revealed as HAP antimicrobial and chemical resistant gene carrying as unveiled of those genetically transferred materials such as via plasmids Nordmann (1993).

2. Properties of *P. aeruginosa*

Smooth and irregular looking colonies on nutrient agar with aromatic sweetish odor. *Pseudomonas aeruginosa* appear on *Pseudomonas* base agar Shiny, opaque, shiny, convex smooth, greenish-yellow colony Medium turned light blue.as shown in figure (1), a wide collection of the bacterium isolates unveiled colonies with green-blue fluorescent color with pyocyanin of medium diffusion in nature ²⁵



Figure (1) *Pseudomonas aeruginosa* on *Pseudomonas* base agar .

β -hemolysin was recorded to be released by certain isolates on blood agar. The growing on MacConkey agar was the dominant feature of all isolates; however, a limited number claimed positive lactose fermentation properties. Positive oxidase with Gram negative non-fermenting bacterial rods was uncovered for the survey isolates ⁸

3. Antibiotic susceptibility test of *P. aeruginosa*:

Disk diffusion method:

The results of presents study showed resistant and sensitive isolates, so the effects of different antibiotics on *P. aeruginosa* isolates were investigated, and the results were similar to locally and worldly studies and the drug susceptibility patterns of *P. aeruginosa* isolates

were varied as shown in table (2) .

Table (2): *P. aeruginosa* based susceptibility-to-antibiotics profile

Antibiotic	Sensitive No.(%)	Resistance No.(%)
Cloxacilin	0(0%)	37(100%)
Amoxicilin / Clavulan acid	0(0%)	37(100%)
Cefotaxime	0(0%)	37(100%)
Ceftazidime	3(8%)	34(92%)
Ceftriaxone	1(2%)	36(97%)
Azetronam	36(97%)	1(3%)
Gentamicin	19(51%)	18(49%)
Levofloxacin	23(62%)	14(37%)
Norfloxacin	34(91%)	3(8%)

As shown above antibiotic susceptibility to *P.aeruginosa* results showed 97% resistance to ceftizoxone (30 μ g), 63% to Norfloxacin , 100% to Cloxacilin , 100% to Amoxicilin /Clavulan acid, 37% to Levofloxacin, 49% to Gentamicin (10 μ g), 92% to Ceftazidime (30 μ g) , 100 % to cefotaxime (30 μ g) and 71% to Azetronam , the results of presents study in line with results of were locally studies conducted by Al-Maamori (2011), Al-saffar (2005), Al-Gibouri (2006) and Zeki (2006). In presents study *P. aeruginosa* isolates showed resistant to aminoglycoside antibiotics, gentamicin (49%) to be disagreed with Lee et al in (2007) who was detected aminoglycoside-resistant in (47%) of aeruginosal reported variants and Alammary, (2013) declared resistant to aminoglycoside antibiotics, gentamicin (45%) in Hilla teaching hospital because the appearance of aminoglycoside resistant as acquired ⁶.

Resistance of *P. aeruginosa* to antibiotics of aminoglycosidic nature shows geographical variation as highest in Southern Europe with Greece, 49.8%, and the United Kingdom, 96.6%, of non-susceptibility to gentamicin ³⁶

The greatest antibiotic resistance rates were observed in ceftizoxime (100%), Cloxacilin (100%), Amoxicilin /Clavulan acid (100%) followed by Ceftriaxone (97%) which was agreement with other studies in ceftizoxime (99.2%), Cloxacilin (91%) and Amoxicilin /Clavulan acid (94%) (Amini and Mobasseri, 2017) and Kerman (95%) ceftizoxime, Cloxacilin (88%) and Amoxicilin /Clavulan acid (95%) (Moosavian and Rahimzadeh, 2015) This can make ceftizoxime, Cloxacilin and Amoxicilin /Clavulan acid a bad choice for further prescription against *P. aeruginosa*. (ICA, 2016). Show in Figure (2).

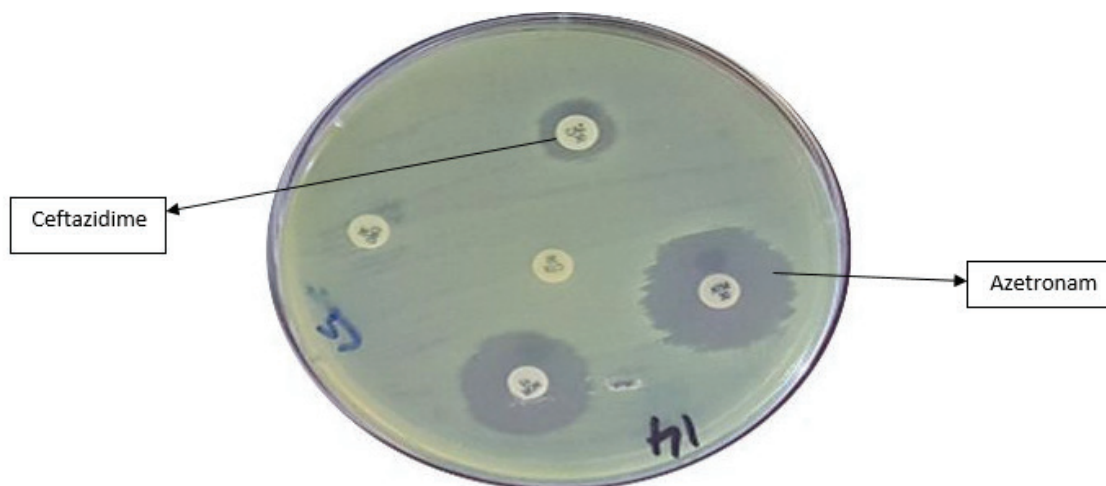


Figure (2) Antibiotics on *Pseudomonas aeruginosa*

Strains of *P. aeruginosa* have evolved increased resistance rates against in ceftizoxime (100%), Cloxacilin (100%), Amoxicilin /Clavulan acid (100%) and Ceftriaxone (97%) (Sofianou et al., 1997 and Snyderman, 2010), The super bacterial resistance may be due the incidence of genetic modifications via mutations and gene moving between strains of *P. auroginosa*.^{7, 32}

The result of present study was disagreement with Babaeekhou et al., (2018) who showed 78.9% resistance to ceftizoxime, 68.1% to Amoxicilin /Clavulan acid, 67.3% to Ceftriaxone 67.3% to carbenicillin, 65.2% to Cloxacilin 63.7% to ceftazidime 63.04% to gentamicin, 53.6%.

Azetronam and Norfloxacin had been recorded the lowest rate of resistance in the present study (3%) and (8%) respectively. It shows relatively low frequencies in other studies, the current findings from this work in line with Saderi and Owlia (2015) findings in Tehran (10.4%) resistance to Azetronam and (13.6%) resistance to Norfloxacin.

Levofloxacin had been recorded resistance rate in present study (37%) was agreement with study concluded by Peymani et al., (2015) in Iran showed resistance of *P.*

aeruginosa to levofloxacin (33.7%).

Geographical and time criteria may apply critical alterations in the status of resistance to those drugs. However, MDR of the bacterium is a featured property narrowing the antibiotic options available for managing related diseases. De novo occurrence of elevated rates of MDR was ensured via drug exposure with circulating of MDR strains between individuals.^{2, 18, 21}

Conclusion

Pseudomonas aeruginosa recovering from patients' infected burns constitute a serious therapeutic problem. *Pseudomonas aeruginosa* isolates were resistant to Cloxacilin, Amoxicilin /Clavulanic acid and Cefotaxime, whereas *Pseudomonas aeruginosa* isolates were susceptible to Azetronam.

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Conflict of Interest: None to declare.

Ethical Clearance: All experimental protocols were approved under the College of Science for women and all experiments were carried out in accordance with

approved guidelines.

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