

## ANTIBIOGRAM PROFILE AND BIOFILM FORMING POTENTIAL OF PSEUDOMONAS SPECIES ISOLATED FROM VARIOUS CLINICAL SPECIMENS

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### ABSTRACT

**Objective:** The present study aimed at finding the resistance pattern of *Pseudomonas aeruginosa* and other *Pseudomonas* species isolated from various clinical specimens in the laboratory.

**Methods:** A total of 150 isolates of different species of *Pseudomonas* obtained from various clinical specimens processed at the Microbiology laboratory of Kasturba Medical College, Manipal Academy of Higher Education, were taken for this study. Antibiotic susceptibility testing was performed by Kirby-Bauer disc diffusion method and interpreted according to the CLSI guidelines. Biofilm assay was performed by modified O'Toole and Kolter method. The results were analyzed using SPSS 17.0 and Student's unpaired t-test, Kruskal-Wallis, Mann-Whitney, ANOVA, and Chi-square test.  $p < 0.05$  was considered statistically significant.

**Results:** Increased resistance was observed by *P. aeruginosa* to cefotaxime, cotrimoxazole, levofloxacin, ofloxacin, and ticarcillin clavulanate. There was also a good correlation between antibiotic resistance to aztreonam, netilmicin, and ceftazidime and biofilm production. Results of the present study, therefore, demonstrated the occurrence of resistance to various antipseudomonal agents among the biofilm-producing *P. aeruginosa* isolates.

**Conclusion:** The present study may help in assessing the seriousness of drug resistance caused by biofilm formation in *P. aeruginosa* and devise strategies through antibiotic policies to minimize such problems.

**Keywords:** Biofilm, *Pseudomonas aeruginosa*, Antibiogram, Antipseudomonal agents.

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### INTRODUCTION

*Pseudomonas* is a large group of aerobic, non-spore forming Gram-negative motile rods, which are pervasive in nature. *Pseudomonas aeruginosa* is a major opportunistic pathogen responsible for acute and chronic infections mainly in hospital settings, especially in patients with compromised host defense mechanism and also with serious underlying disease conditions [1]. According to the CDC, approximately 8% of all health care-associated infections reported to the National Healthcare Safety Network are caused by *P. aeruginosa*.

The survival of *Pseudomonas* within the host in the initial stages of infection is aided by the secretion of various toxins and virulence factors including pyocyanin, proteases, and elastases [2]. *P. aeruginosa* possesses various intrinsic and acquired mechanisms of drug resistance [3]. Risk factors associated with the emergence of drug-resistant strains include previous antipseudomonal drug treatment and prolonged use of antibiotics. Besides, prolonged hospital stay and increased susceptibility of patients to secondary bacteremia lead to the acquisition of resistant strains [4].

*P. aeruginosa* forms microcolonies enclosed in extracellular polymeric substances (EPSs) termed as biofilms [5]. Biofilm formation leads to persistent and chronic infection by resisting the action of antimicrobial agents [6]. It has the ability to resist the suppression of the organism by various physical and chemical treatments [5]. Biofilm forms the major response mechanism to external stress factors by inducing many additional phenotypic alterations such as loss of motility, reduced growth rate, and altered susceptibility to host response [6,7].

The present study aimed at correlating biofilm production by *P. aeruginosa* and other species of *Pseudomonas* isolated from various clinical samples with their antibiogram pattern.

### METHODS

#### Collection and Identification of bacterial isolates

A total of 150 isolates of *Pseudomonas* (*P. aeruginosa* -75 and other *Pseudomonas* species-75) obtained from various clinical specimens processed at the Microbiology laboratory of Kasturba Medical College, Manipal Academy of Higher Education, were taken for this study (with a 95% confidence level and 80% power, the sample size came up to 75 each). The Institutional Ethics Committee clearance was obtained for the study. The isolates were identified by standard biochemical methods [8] or by VITEK 2 system. Antibiotic susceptibility testing was performed by Kirby-Bauer disc diffusion method and interpreted according to the CLSI guidelines [9]. For isolates identified by VITEK 2 system, sensitivity was recorded from it.

#### Biofilm assay

This was done by modified O'Toole and Kolter method [10-13]. The bacterial colony was inoculated in brain heart infusion (BHI) broth and incubated at 37°C for 18 h. It was then diluted with fresh BHI broth, and the turbidity was adjusted to 0.5 McFarland standard. 200 µl of the suspension was dispensed into microtiter plate wells in duplicate. The plate was incubated at 37°C for 24 h. The contents were aspirated and washed with (phosphate-buffered saline (pH 7.4) following which 100 µl bouin fixative was added. The plate was incubated at 25°C for 10 min. The contents were discarded, and the wells were stained with 1% crystal violet. After 1 min, the excess stain was rinsed off by placing the plate under running tap water. Then, 33% glacial acetic acid was added to each well and optical densities of stained adherent bacterial films were read with Micro ELISA plate reader at 570 nm. Mean reading from two wells were calculated. *P. aeruginosa* ATCC 27853 was included as control.

Statistical analysis: All experiments were performed in duplicate. Statistical analysis was performed using SPSS 17.0 for Windows (SPSS, Inc., Chicago, IL, USA) and using Student's unpaired t-test, Kruskal-Wallis, Mann-Whitney, ANOVA, and Chi-square test.  $p < 0.05$  was considered statistically significant.

**RESULTS**

Of 75 isolates of *P. aeruginosa*, 64 (35.2%) were isolated from males, while 11 (14.7%) were isolated from females. With regard to the other *Pseudomonas* spp., of 75 isolates, 54 (72.0%) were from males and 21 (28.0%) were from females. As shown in Fig. 1, the highest rate of isolation was from swabs and exudate material from wound infections (21.3%). Similarly, isolation rate of other species of *Pseudomonas* also accounted to 24.0% as shown in Fig. 2.

**Antibiotic resistance of the isolates**

Among the *P. aeruginosa* strains, 80% were resistant to cefotaxime, 75% to cotrimoxazole, and 68.2% to ticarcillin/clavulanic acid. All the isolates were sensitive to colistin and polymyxin-B (Fig. 3). Resistance to levofloxacin was 62.5%. Similarly, the other species of *Pseudomonas* also showed maximum resistance to cefotaxime (75%) and cotrimoxazole (75%), while levofloxacin was third in the list (66.7%) as shown in Fig. 4.

**Resistance to aminoglycosides**

Among the aminoglycosides, the percentage resistance of *P. aeruginosa* revealed 38.4% resistance to amikacin, 41.3% resistance to gentamicin, and also a percentage as high as 46.2% for netilmicin. However, the percentage of resistance to amikacin was lower in other species of *Pseudomonas*, being 20.8% for amikacin, while it was 36.2% for gentamicin and 12.5% for netilmicin.

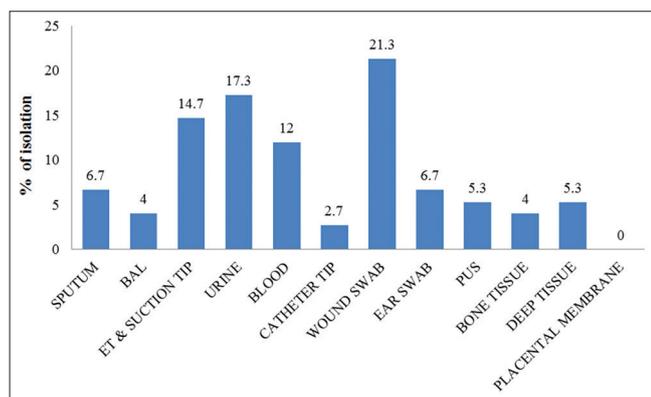


Fig. 1: The percentage isolation of *Pseudomonas aeruginosa* from various specimens. BAL: Bronchoalveolar lavage

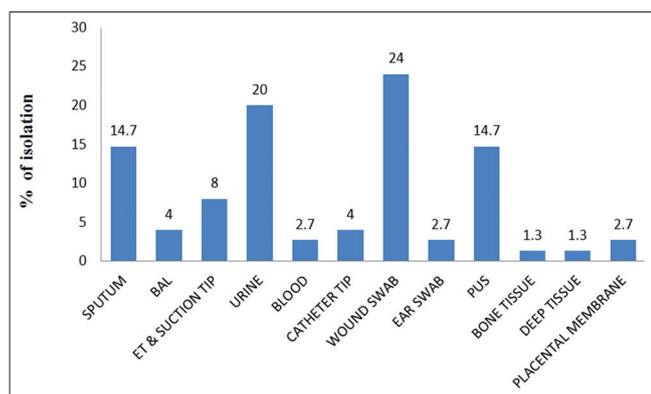


Fig. 2: The percentage isolation of *Pseudomonas* spp. from various specimens. BAL: Bronchoalveolar lavage

**Carbapenem resistance**

*P. aeruginosa* strains exhibited 24.7% resistance to imipenem while it was 19.2% for meropenem which was almost on par with other species of *Pseudomonas*, and the resistance to imipenem and meropenem being 27% and 21.1%, respectively.

**Resistance to antipseudomonal penicillins**

Most of the strains were sensitive to piperacillin and piperacillin-tazobactam. Resistance to piperacillin and piperacillin-tazobactam by *P. aeruginosa* was 16.7% and 12.1%, respectively, while for other species, the percentage of resistance to piperacillin and piperacillin-tazobactam was 19.6% and 17.8%, respectively. However, 68.2% resistance was shown by *P. aeruginosa* against ticarcillin-clavulanic acid, a carboxypenicillin.

**Resistance to quinolones**

Both *P. aeruginosa* and the other species showed high level of resistance to the quinolones with higher resistance to levofloxacin (62.5% and 66.7%, respectively), while for ciprofloxacin, the resistance exhibited by *P. aeruginosa* and the other species was 45.8% and 34.2%, respectively.

**Resistance to minocycline**

There was minimal resistance to minocycline, a tetracycline group of drug by *P. aeruginosa* strains, while other species showed 100% susceptibility to this drug.

**Resistance to cephalosporins**

The isolates also exhibited resistance to ceftazidime, a third-generation antipseudomonal drug (28.4% by *P. aeruginosa* and 23.2% by the other

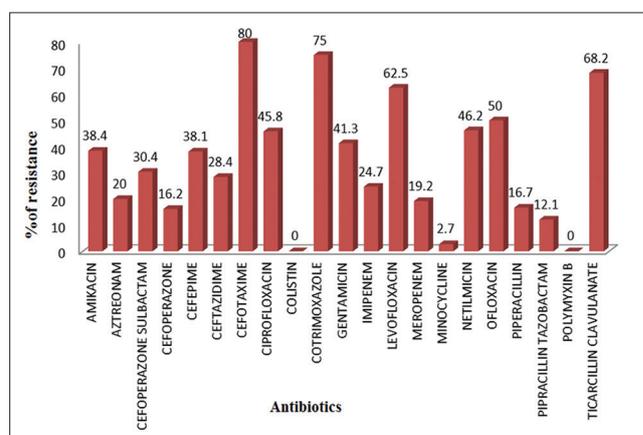


Fig. 3: The resistance pattern of *Pseudomonas aeruginosa* to various antibiotics

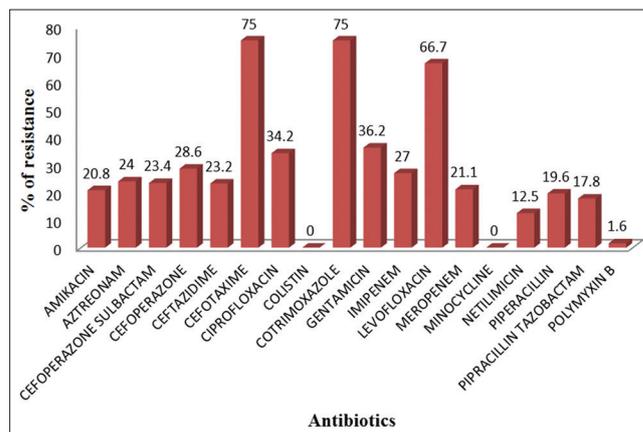


Fig. 4: The resistance pattern of *Pseudomonas* spp. to various antibiotics

species). There was a very high percentage of resistance to cefotaxime by *P. aeruginosa* and the other species of *Pseudomonas* (80% and 75%, respectively). The percentage resistance of *P. aeruginosa* and the other species of *Pseudomonas* to cefoperazone-sulbactam was 30.4% and 23.4%, respectively, while the resistance to cefoperazone was 16.2% and 28.6%, respectively.

**Resistance to monobactams**

There was 20% resistance to aztreonam, a monobactam antibiotic by *P. aeruginosa*, while 24% resistance was shown by other species of *Pseudomonas*.

**Results of biofilm formation by the isolates**

OD<sub>570</sub> of biofilm produced by the isolates is depicted in Tables 1 and 2, and the comparison between the biofilm produced by *P. aeruginosa* and the other species of *Pseudomonas* is shown in Fig. 5 and Table 3.

**Aminoglycosides**

Among the three aminoglycosides, gentamicin- and netilmicin-resistant strains of *P. aeruginosa* showed more biofilm production while amikacin sensitive strains produced more biofilm. In case of other *Pseudomonas* spp., more biofilm production was shown by aminoglycoside-sensitive strains.

**Carbapenems**

*P. aeruginosa* which was resistant to imipenem produced more amount of biofilm, whereas in other species production, it was more in case of sensitive isolates. With regard to meropenem, intermediately susceptible strains produced more biofilm in case of both *P. aeruginosa* and *Pseudomonas* spp.

**Antipseudomonal penicillins**

Biofilm production was more by piperacillin susceptible strains of both *P. aeruginosa* and *Pseudomonas* spp. The strains which were intermediately susceptible to piperacillin-tazobactam isolates showed more biofilm in case of *P. aeruginosa*. In other species, piperacillin-tazobactam resistant strains produced more biofilm.

**Quinolones**

*P. aeruginosa* which was resistant to ciprofloxacin produced more amount of biofilm, whereas in *Pseudomonas* spp. strains that were intermediately susceptible to ciprofloxacin produced more biofilm. *P. aeruginosa* sensitive to levofloxacin showed more biofilm production, while in case of other species, levofloxacin-resistant ones produced more biofilm.

**Cephalosporins**

Isolates of both *P. aeruginosa* and *Pseudomonas* spp. which were resistant to cefoperazone sulbactam produced more biofilm. In case of ceftazidime, more biofilm production was showed by *P. aeruginosa* intermediately susceptible isolates and ceftazidime-resistant isolates of other *Pseudomonas* spp. Isolates of both *P. aeruginosa* and *Pseudomonas* spp. which were resistant to cefotaxime and cefoperazone produced more biofilm.

**Monobactams**

*P. aeruginosa* which was resistant to aztreonam produced more amount of biofilm, whereas in *Pseudomonas* spp., more biofilm production was in sensitive strains.

Minimum inhibitory concentration (MIC) of certain antibiotics (ceftazidime, ciprofloxacin, and piperacillin-tazobactam) against biofilms also demonstrated that when MIC values increase, there was a significant increase in biofilm.

**DISCUSSION**

Antibiotic resistance is a major problem in *P. aeruginosa*. The organism exhibits intrinsic resistance to several beta-lactam antibiotics and may also acquire additional resistance mechanisms either due to mutational events or due to the acquisition of transferable genetic elements [14].

*P. aeruginosa* also shows intrinsic resistance by the expression of chromosomally encoded inducible AmpC beta-lactamase and also by several important efflux pump systems that export antibiotics, biocides, dyes, detergents, metabolic inhibitors, organic solvents, and molecules involved in bacterial cell-to-cell communication [15].

Carbapenem resistance mechanisms have emerged under the pressure of carbapenem use in clinical settings and may be classified as enzymatic, which include carbapenemases, aminoglycoside-modifying enzymes, and 16S rRNA methylases or nonenzymatic involving decreased transcription of OprD gene and overproduction of MexAB-OprM efflux system [16]. Carbapenem resistance, however,

**Table 1: OD<sub>570</sub> of biofilm produced by *P. aeruginosa* isolated from various clinical specimens**

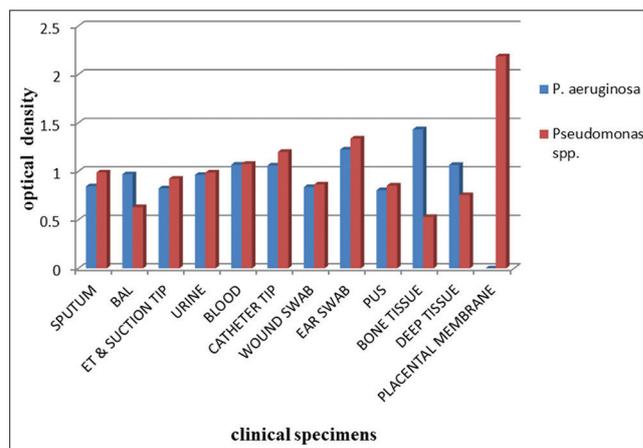
Clinical specimen	Mean number of isolates±SD	OD <sub>570</sub> ±SD
Sputum	5±1.023	0.847
Bronchoalveolar lavage	3±0.135	0.971
Et and suction tip	11±0.473	0.825
Urine	13±0.484	0.965
Blood	9±0.512	1.069
Catheter tip	2±0.520	1.062
Wound swab	16±0.439	0.839
Ear swab	5±0.554	1.226
Pus	4±0.446	0.807
Bone tissue	3±0.690	1.434
Deep tissue	4±0.323	1.067
Placental membrane	0	0

*P. aeruginosa: Pseudomonas aeruginosa*

**Table 2: OD<sub>570</sub> of biofilm produced by *Pseudomonas* spp. isolated from various clinical specimens**

Clinical specimen	Mean number of isolates±SD	OD <sub>570</sub> ±SD
Sputum	11±0.524	0.989
Bal	3±0.393	0.631
Et and suction tip	6±0.517	0.925
Urine	15±0.387	0.988
Blood	2±0.517	1.077
Catheter tip	3±0.581	1.202
Wound swab	18±0.480	0.866
Ear swab	2±0.941	1.339
Pus	11±0.499	0.854
Bone tissue	1±	0.531
Deep tissue	1±	0.756
Placental membrane	2±0.684	2.187

*P. aeruginosa: Pseudomonas aeruginosa*



**Fig. 5: Comparison of biofilm production in *Pseudomonas aeruginosa* and *Pseudomonas* spp.**

Table 3: Correlation between antibiotic sensitivity pattern and biofilm production

Antimicrobial agents	Organism	Mean OD <sub>570</sub> isolates±SD Sensitive	Mean OD <sub>570</sub> isolates±SD Resistant	Mean OD <sub>570</sub> isolates±SD Intermediate	p value
Amikacin	<i>P. aeruginosa</i>	0.975±0.471	0.966±0.58		0.941
	<i>Pseudomonas</i> spp.	1.03±0.553	0.758±0.347	0.807±0.089	0.166
Ceftazidime	<i>P. aeruginosa</i>	0.905±0.458	0.923±0.551	1.491±0.492	0.015
	<i>Pseudomonas</i> spp.	0.970±0.507	1.067±0.600	0.694±1	0.698
Cefoperazone sulbactam	<i>P. aeruginosa</i>	0.941±0.466	1.131±0.614	0.708±0.410	0.242
	<i>Pseudomonas</i> spp.	0.983±0.492	1.005±0.592	0.871±0.603	0.901
Ciprofloxacin	<i>P. aeruginosa</i>	0.901±0.445	1.055±0.586		0.209
	<i>Pseudomonas</i> spp.	0.968±0.490	0.932±0.539	2.08±1	0.093
Gentamicin	<i>P. aeruginosa</i>	0.888±0.454	1.059±0.588	0.873±0.191	0.367
	<i>Pseudomonas</i> spp.	1.046±0.546	0.852±0.466	0.951±1	0.337
Imipenem	<i>P. aeruginosa</i>	0.886±0.433	1.194±0.682		0.027
	<i>Pseudomonas</i> spp.	1.014±0.573	0.832±0.347	1.014±0.099	0.408
Meropenem	<i>P. aeruginosa</i>	0.959±0.508	0.864±0.489	1.442±1	0.507
	<i>Pseudomonas</i> spp.	1.004±0.550	0.783±0.331	1.396±0.968	0.183
Piperacillin	<i>P. aeruginosa</i>	0.891±0.454	0.972±0.432		0.647
	<i>Pseudomonas</i> spp.	0.945±0.494	1.05±0.764		0.569
Piperacillin-tazobactam	<i>P. aeruginosa</i>	0.961±0.524	0.784±0.398	1.466±1	0.395
	<i>Pseudomonas</i> spp.	0.962±0.482	1.044±0.677	0.335±1	0.617
Netilmicin	<i>P. aeruginosa</i>	0.805±0.190	1.289±0.529		0.044
	<i>Pseudomonas</i> spp.	1.080±0.491	0.717±0.094		0.328
Cotrimoxazole	<i>P. aeruginosa</i>	0.709±0.050	1.175±0.951		-
	<i>Pseudomonas</i> spp.	1.340±1	0.878±0.161		-
Aztreonam	<i>P. aeruginosa</i>	0.776±0.337	1.613±0.783		0.003
	<i>Pseudomonas</i> spp.	1.007±0.582	0.874±0.437		0.614
Cefotaxime	<i>P. aeruginosa</i>		1.11±0.674	0.744±1	-
	<i>Pseudomonas</i> spp.	0.738±1		0.934±0.441	-
Cefoperazone	<i>P. aeruginosa</i>	0.891±0.443	1.097±0.681	0.392±0.081	0.211
	<i>Pseudomonas</i> spp.	0.907±0.460	1.059±0.681	0.896±1	0.709
Levofloxacin	<i>P. aeruginosa</i>	1.473±0.508	1.251±0.688	0.771±0.084	0.451
	<i>Pseudomonas</i> spp.	0.531±1	0.845±0.213		0.996

*P. aeruginosa*: *Pseudomonas aeruginosa*

develops frequently due to the concomitant presence of more than one mechanism [17,18].

The present study showed a higher level of imipenem resistance as compared to meropenem resistance. This could also be due to the frequent use of imipenem which could change the permeability of the outer membrane or modify the target sites of the organism when given with piperacillin, thus resulting in carbapenem resistance [19].

Most of the isolates were sensitive to aztreonam, the only monobactam possibly effective against carbapenem-hydrolyzing strains of *P. aeruginosa* which is often used in patients who are penicillin allergic or who cannot tolerate aminoglycosides [20].

Previous studies have shown growing resistance to ciprofloxacin and cefotaxime by *P. aeruginosa* [21-24]. This was quite similar to the resistance pattern that was observed in our study with resistance percentage to cefotaxime being as high as 80%, while for ciprofloxacin, it was 45.8%. Ceftazidime was shown a high level of resistance in previous study [1], but the present study shows a decrease in resistance pattern.

Quinolones are generally preferred by the physicians as an empirical therapy mainly because of the easy availability in oral forms and reasonable cost [25-27].

However, emerging quinolone-resistant strains of *P. aeruginosa* have surfaced owing to the changes in target enzymes of the bacteria and active efflux pumps generated to prevent the entry of the drugs. The present study also found very high resistance to levofloxacin and ciprofloxacin by *P. aeruginosa*.

The present study also showed a high level of resistance to ticarcillin-clavulanic acid, a carboxypenicillin. Studies have demonstrated that clavulanate may induce AmpC expression in *P. aeruginosa* [28]. Significant induction was shown to occur with pharmacokinetically

relevant concentrations of clavulanate. Besides, the induction of AmpC by clavulanate was shown to significantly antagonize or substantially diminish the antibacterial activity of ticarcillin. The study also suggested that, in the selection of an antipseudomonal  $\beta$ -lactam for the treatment of *P. aeruginosa* infections, the combination of ticarcillin-clavulanate should be avoided, especially with immunocompromised patients, for whom bacterial killing is required to ensure clinical success.

It was interesting to note the correlation between antibiotic resistance and biofilm production. Among the aminoglycosides, netilmicin resistance and biofilm production demonstrated good correlation ( $p=0.044$ ). Aminoglycosides which constitute a vital component of antipseudomonal therapy have been showing resistance in the recent past. Hence, it is important to remember that prolonged therapy may lead to persisters by forming biofilms [29].

Imipenem-resistant *P. aeruginosa* strains produced more amount of biofilm. Imipenem exposure may lead to an expression of alginate which is known to provide the biofilm bacteria with a protected environment [30].

Aztreonam, the monobactam commonly preferred for treating *Pseudomonas* infections in patients allergic to penicillins or aminoglycosides, showed only 20% resistance, but the resistant strains were strong biofilm producers ( $p=0.003$ ). It is possible that resistance may lead to persister cells in this case and this needs to be studied further.

Inhaled aztreonam has been used in combination with tigecycline in treatment of cystic fibrosis, and the study has also shown enhancement of antibiofilm activity. However, some strains have shown tolerance to aztreonam-tigecycline combination [31]. Ps1EPS production appears to relate to antibiotic tolerance in some of the strains. In addition to this, other mechanisms such as lower bacterial outer membrane permeability, antibiotic efflux pump, and beta-lactamase resistance may play a role in *P. aeruginosa* tolerance to aztreonam [32,33]. In a particular study, even same antibiotics showed significant quorum sensing inhibitory activity

against different test organisms, and in clinical isolates, similar antibiotics exhibited varied effects in their effective concentration [34,35].

Our study revealed that isolates of both *P. aeruginosa* and *Pseudomonas* spp. which were resistant to cefotaxime and cefoperazone produced more biofilm. A particular study has shown the ability of cefuroxime to dislodge the biofilm formation in *Pseudomonas* by 65.57%.

Increase in biofilm with an increase in MIC of antibiotics seen in the present study is on par with other studies reported [10,13,14].

## CONCLUSION

Results of the present study demonstrated the occurrence of resistance to various antipseudomonal agents among the *P. aeruginosa* isolates as well as in other species of *Pseudomonas*. While certain drugs such as piperacillin-tazobactam, carbapenems, and amikacin remain the mainstay of the treatment of pseudomonal infections, it is alarming to note an increase in resistance levels to these antibiotics. Besides this, the organisms also produce biofilms which serve as barriers to effective therapy. Regular antimicrobial susceptibility monitoring would help and guide the physicians in prescribing the right combinations of antimicrobial to limit and prevent the emergence of multidrug-resistant strains of *P. aeruginosa*. Antibiotics should be used judiciously and at the optimum concentration so as to inhibit biofilm formation and eradicate persister cells. Studies have postulated that combination of a biofilm inhibitor with a conventional antibiotic to control biofilms thereby permits the drug to reach the cells trapped inside the biofilm. As this is a hospital-based epidemiological data, the present study will help in implementation of better patient management and infection control strategies.

## CONFLICTS OF INTEREST

All authors have none to declare.

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