

A prospective study to analyse antibiotic susceptibility pattern of *Pseudomonas aeruginosa* in a tertiary care hospital

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ABSTRACT

Background: *Pseudomonas aeruginosa*, a gram-negative pathogen is commonly associated with nosocomial infections. Infections caused by *P. aeruginosa* can range from superficial skin infections to fulminant sepsis. Antimicrobial resistance is an area of prime concern in pseudomonal infections. The objective of the study was to evaluate and analyse the antibiotic susceptibility pattern of *P. aeruginosa* at a tertiary care hospital in South India.

Methods: The study was carried out at Kasturba Medical College and Hospital, Manipal, India from January 2011 to December 2011. Ten different types of specimens were collected from patients who were culture positive for *Pseudomonas aeruginosa*. Antibiotic susceptibility was confirmed by disk diffusion technique on Muller-Hinton medium and was performed according to the Clinical Laboratory Standard Institute (CLSI) guidelines.

Results: Out of 200 samples of *P. aeruginosa* 69.5% and 30.5% were from male and female patients respectively. Majority of the specimen from which *P. aeruginosa* was isolated consisted of sputum, pus and urine. Among the antimicrobial drugs tested, organism was most sensitive to carbapenems (77.5%), piperacillin-tazobactam (77%) and cefoperazone-sulbactam (72%). Resistance rates were high for fluoroquinolones (FQs) (43.5%), gentamicin (40.5%), tobramycin (40.5%), ticarcillin-clavulanic acid (39%) and aztreonam (38%) when compared to cefepime (31.5%), ceftazidime (32.5%), netilmicin (34.5%) and amikacin (35%).

Conclusions: Carbapenems and piperacillin-tazobactam were the best antipseudomonal agents with highest sensitivity to *P. aeruginosa*. FQs, gentamicin and tobramycin were the least effective drugs against *P. aeruginosa* as monotherapy.

Keywords: *P. aeruginosa*, Antibiotic susceptibility, Antipseudomonal agents, Nosocomial infections

INTRODUCTION

Pseudomonas aeruginosa, a gram-negative pathogen is commonly associated with nosocomial infections. It is involved in wide variety of human infections, ranging from superficial skin infections, acute and chronic lung infections to fulminant sepsis. It is distinguished as an opportunistic pathogen causing infection in patients with defective physical, phagocytic and immunological defense mechanisms.¹

Historically, it was considered a major burn wound pathogen, an agent of bacteremia in neutropenics and the most common pathogen in cystic fibrosis patients. However, these interesting associations have undergone considerable changes. Now *P. aeruginosa* is the 2nd most common cause of nosocomial pneumonia (17%), 3rd important etiology for urinary tract infection (7%), 4th most common cause of surgical site infection, 5th most common isolate (9%) overall from all sites and 7th most frequently isolated pathogen from the bloodstream.²

Antimicrobial resistance (AMR) to broad spectrum antibiotics is an area of prime concern in pseudomonal infections.^{1,3} It is the most widespread multidrug-resistant (MDR) gram-negative pathogen causing pneumonia in hospitalized patients. It not only limits therapeutic options but also affects clinical outcome by increasing morbidity and mortality.⁴ Therefore, a study was conducted to determine the current antibiotic sensitivity and resistance rates which would help in laying down current recommendations for empirical antibiotic regimens for treating *P. aeruginosa* infections and also to minimize progression of MDR.

The objective of the study was to analyse antibiotic susceptibility patterns of *P. aeruginosa* in study population.

METHODS

The study was conducted in Kasturba Hospital, Manipal, Karnataka, India over a period of one year (January to December 2011). Study subjects included were patients aged above 18 years of either sex who got admitted to the hospital during study period with positive growth for *P. aeruginosa*. Subjects were explained about the study and written informed consent was obtained. Cases from burns ward and OPD were excluded from study.

A total of 200 patients with positive culture for *P. aeruginosa* were included in the study after considering inclusion and exclusion criteria. Ethical clearance was obtained from institutional ethical committee before the start of study.

Pre-designed proforma was used to collect demographic details, laboratory data and treatment information of the patient. Microbiological data including specimen, site, presence of any associated organism, antibacterial sensitivity and resistance patterns of *P. aeruginosa* was noted.

Microbiology

Culture examination was carried out using blood agar, nutrient agar and MacConkey's agar, followed by study of colony morphology, pigment production, positive oxidase reaction and oxidase in oxidation fermentation medium.⁵ Antibiotic susceptibility was confirmed by disk diffusion technique on Muller-Hinton medium and was performed according to the Clinical Laboratory Standard Institute (CLSI) guidelines. Paper disks were impregnated with antibiotics. These were commercially procured from span diagnostics.

Antibiotics which were tested for sensitivity are ceftazidime, cefoperazone-sulbactam, cefepime or ceftiprome, piperacillin, piperacillin-tazobactam, ticarcillin-clavulanic acid, aztreonam, carbapenem, gentamicin, tobramycin, amikacin, netilmicin and ciprofloxacin or levofloxacin.

Cultures were incubated overnight at 37°C. Diameter of the zone of inhibition was measured and compared to that of standard strain and results were interpreted as sensitive and resistant, based on CLSI guidelines.⁶

Statistical analysis

Analysis was primarily descriptive. Data thus obtained was entered in excel sheet. Analysis was done using SPSS version 17.0 as percentages and proportions.

RESULTS

Out of 200 cases analysed in our study, there were 141 (70.5%) males and 59 (29.5%) females and majority of them were aged 40 and above (Table 1). Various specimens were studied including pus-wound swab, sputum, urine, blood, etc. *P. aeruginosa* was most commonly isolated from sputum (31.5%) followed by pus-wound swab (25.5%) and urine (22.5%) (Table 2).

Table 1: Age and sex distribution of study subjects.

Age group (years)	Males (%)	Females (%)	Total (%)
18-40	23	21	44
41-60	57	22	79
>60	59	18	76
Total	139	61	200

Table 2: Microbiological specimens with positive culture for *P. aeruginosa*.

Specimen	Total no. of cases	Percentage of cases
Pus and wound swab	51	25.5%
Sputum	63	31.5%
Urine	45	22.5%
Blood	25	12.5%
Others*	16	8.0%

*ear swab, tracheal swab, pleural fluid, endotracheal aspirate, bronchoalveolar lavage and intercostal drainage

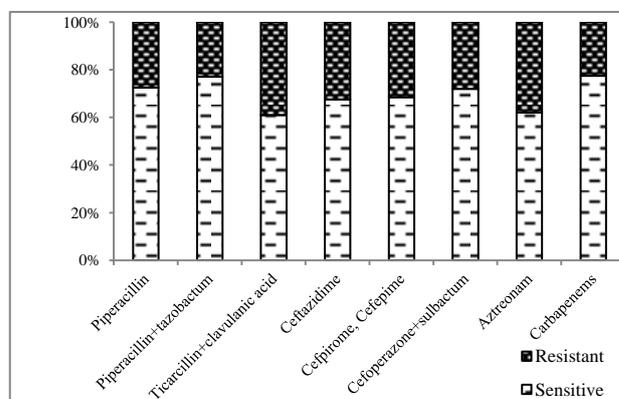


Figure 1: In vitro susceptibilities of *P. aeruginosa* to anti-pseudomonal beta-lactam antibiotics.

Susceptibility to anti-pseudomonal β -lactam antibiotics in decreasing order of frequency was as follows: carbapenems (77.5%), piperacillin-tazobactam (77%), piperacillin (72.5%), cefoperazone-sulbactam (72%), cefpirome or cefepime (68.5%), ceftazidime (67.5%), aztreonam (62%) and ticarcillin-clavulanic acid (61%) (Figure 1).

Percentage of susceptible isolates for aminoglycoside group of antibiotics was 65.5% (netilmicin), 65% (amikacin), 59.5% (tobramycin) and 59.5% (gentamicin). The fluoroquinolones (FQs) tested were ciprofloxacin and levofloxacin, 56.5% of isolates were susceptible to FQs (Figure 2).

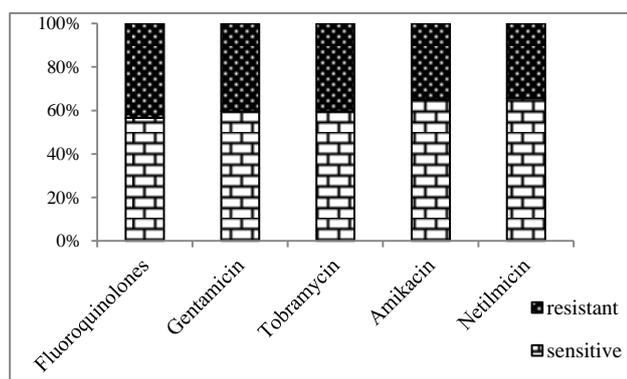


Figure 2: In vitro susceptibilities of *P. aeruginosa* to fluoroquinolones and aminoglycosides.

DISCUSSION

Out of 200 cases analysed in our study *P. aeruginosa* infection was more in patients with age above 40 years with a male preponderance (69.5%), while females constituted only 30.5% of cases. This was consistent with a similar study done in India.⁷ Highest number of *P. aeruginosa* was isolated from sputum making the organism the most common cause of respiratory tract infection amongst all hospital acquired infections (HAIs). It is amongst the most common causes of ventilator associated pneumonia and carries the highest mortality among HAIs. It also harbours in damaged bronchi of bronchiectasis patients leading to acute exacerbations.⁴

Among the β -lactam antibiotics tested, carbapenems and piperacillin-tazobactam showed least resistance rates. These antibiotics have broader spectrum of activity than most β -lactam antibiotics, therefore were most effective against treating *P. aeruginosa* infections. Concurrent administration of β -lactamase inhibitor tazobactam enhanced antipseudomonal sensitivity of an extended spectrum penicillin like piperacillin when compared to piperacillin alone. Sensitivity rates of cefepime and cefpirome were slightly superior to ceftazidime due to their non-susceptibility to inducible β -lactamases. Cefoperazone-sulbactam was superior among all cephalosporins.

We observed that aminoglycosides and FQs were less effective when compared to most of β -lactam antibiotics. Netilmicin and amikacin were most effective among the aminoglycosides. Among all the drugs tested FQs showed highest resistance rates followed by gentamicin and tobramycin, hence these drugs should be avoided as monotherapy especially in MDR *P. aeruginosa* infections.

Multidrug resistance phenotypes are slowly increasing in prevalence among *P. aeruginosa* isolates.⁸⁻¹¹ However, comparison between studies is often difficult, because definitions of multidrug resistance have not been uniform.^{9,10,12} The reasons for increasing nosocomial spread of MDR isolates may include lack of adherence to approved infection control policies in hospitals, increasing or cumulative antimicrobial use, and changes in the public health infrastructure.¹²⁻¹⁶

In conclusion, susceptibility of antipseudomonal agents against different isolates has decreased. Significant reduction in susceptibilities of *P. aeruginosa* isolates may compromise the ability to choose efficacious empirical regimens for treatment of this formidable pathogen especially in critically ill patients. The present study provides valuable information related to emerging trends in antimicrobial resistance to monotherapy, which is vital for clinicians in the selection of reliable empirical regimen for treating *P. aeruginosa* infections.

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