

Determination Pattern of Antimicrobial Resistance of Pseudomonas Isolated from Patients in a University Tertiary Hospital

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Abstract

Introduction: Pseudomonas are ubiquitous bacteria widely present in nature. However, they have emerged as an opportunistic pathogen for humans. This bacterium is accountable for many localized and disseminated diseases, especially wounds in burn patients, respiratory infections, septicemia and bacteremia. Among all Pseudomonas species, *P. aeruginosa* is one of the important and most virulent species in hospital settings, while other pathogenic species include *stutzeri*, *putida* and *fluorescence*. The aim of this study was to assess pattern of antibiotic resistance in these bacteria isolated from a University teaching and treatment center. This cross-sectional study was conducted on 99 Pseudomonas strains (68 strains *P. aeruginosa* and 31 other Pseudomonas species) isolated from various clinical specimens. Antimicrobial drug susceptibility test was performed using the disc agar diffusion method according to CLSI recommendations. In this study, among various clinical specimens sent to microbiology laboratory, wound (59.59%) was found as a major source of Pseudomonas spp. Among various wards, Pseudomonas spp. was isolated more from patients admitted to burns ward (48.48%). Antibiotic susceptibility assay results revealed non susceptibility pattern towards most of the antibiotics; however, among all antibiotics tested, most common resistance was observed towards ceftazidime (76.76%). The results of this study shows the presence of Pseudomonas infection in the hospital setting and their developed resistance towards many conventional antibiotics, which is a concern at this treatment center. Thus, there exist a need for evaluation of careful and accurate measurement of resistance and assessment of exact drug administration policies. Therefore, to control the infection and prevent from increased prevalence of resistant strains appropriate resolution should be followed.

Keywords: Pseudomonas; Antibiotic Resistance; Nosocomial infection;

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Introduction

Pseudomonas is an opportunistic pathogen involved in many localized and invasive infections envisioning a serious concern in patient care environment (1). Presence of many cellular components as well as virulent genes are some of the factors responsible for fever, motility and adhesion of bacteria to the biological cell membrane (2). In addition, *Pseudomonas aeruginosa* (*P.aeruginosa*) produces a variety of exo-enzymes which are one of the factors responsible for augmenting pathogenicity. The attachment of bacteria to medical supplies is accelerated through pilus and fimbriae and secretion of exopolysaccharide which binds to water and forms a gel assisting bacteria to remain suspended onto the artificial devices (3). *Pseudomonas* spp. are widely distributed in nature too. Various species of *Pseudomonas* can survive in different environmental conditions, but they are often found in moist environments of Hospital and considered as opportunistic pathogens for humans (4). The bacterium is also involved in broad spectrum of diseases, including infections especially in burned patients, respiratory infections, septicemia and bacteremia (5). These bacteria are also a primary cause of ventilator-associated pneumonia in the special care unit (ICU) (6). Amongst the various species of *Pseudomonas*, *P.aeruginosa* is the most important bacteria in many nosocomial infections (7).

One of the most serious concerns associated with *Pseudomonas* infections is the emergence of antibiotic resistance towards many traditional therapeutic agents which has doubled their importance particularly in nosocomial infections (8). *P.aeruginosa*, especially among patients with immunosuppression or long-term hospitalization, leads to irreparable consequences (7).

The treatment of patients with *P.aeruginosa* infection is problematic because these organisms are innately resistant to several antibiotic classes and, on the other hand, are able to acquire resistance to many other antimicrobial drugs (9, 10). The development of antibiotic resistance, particularly resistance to aminoglycosides which is plasmid mediated, is of interest since such resistance can cause the transfer of resistance genes in sensitive *Pseudomonas* spp. and other Gram-negative bacteria and turn them non-susceptible to most therapeutic regimens (11). *P.aeruginosa* is also notorious for being intrinsically resistant to many structurally unrelated antimicrobial agents by exhibiting low permeability of its outer membrane, the constitutive expression of various efflux pumps and the naturally occurring chromosomal AmpC β lactamase, and it can acquire additional resistant gene from other organisms via plasmids, transposons, and bacteriophages. It also shows a very high innate resistance to antimicrobial drugs (jcdr-8-DC30) through biofilm production (12). Horizontal gene transfers via mobile genetic elements coding for Class B Carbapenemase which hydrolyses all beta-lactams except aztreonam is one of the carbapenem resistance mechanisms (13).

Identifying common microorganisms and their antibiotic resistance pattern assists physicians with selecting the appropriate antibiotic and preventing the unnecessary prescription of antibiotics and spread of antibiotic resistance. This study aimed to study prevalence and antibiotic resistance pattern in *Pseudomonas* species isolated from different clinical specimens taken from in-patient hospitalized patients admitted to various wards of Sina Hospital, one of University Teaching Hospitals of Tabriz University of Medical Sciences, Iran. Sina Hospital is a tertiary



hospital which receives patients from North West Iran.

Materials and Methods

In this study, 99 clinical isolates including *P.aeruginosa* and other *Pseudomonas* species were collected from various wards of the hospital including, burn, infectious, surgery, internal and emergency wards and all ICUs from July 2015 until June 2016. These bacteria specimens were isolated from blood, urine, sputum, ulcers, abscesses and tracheal tubes. The samples were processed for the isolation and identification of *Pseudomonas* spp. along with other pathogenic bacteria according to phenotypic methods. Wound and urine specimens yielding >10⁵ colony forming units (cfu/ml) were included in this study. Duplicate isolates from the same patient were not enrolled. Antibiotic susceptibility testing was performed on all isolates using the Kirby-Bauer method. The tests were performed according to the guidelines of clinical and laboratory standards institute (CLSI) (14), with a panel of following antibiotics: Ofloxacin (10µg), Ciprofloxacin (5µg), Gentamicin (10µg), Amikacin (30µg) Imipenem (10µg), Ceftriaxone (30µg), Ceftazidime (30µg), Levofloxacin (5µg), Tobramycin (10 µg), and Carbenicillin (25 µg). Statistical analysis: Categorical variables were compared by means of either χ^2 analysis or Fisher's exact test when needed. A 2-tailed P value of <0.5 was considered significant. All statistical calculations were done using standard programs in SPSS 16 software.

Results

A total of 99 *Pseudomonas* species were isolated from diverse clinical specimens. Of these, 68 (68.7%) were *Pseudomonas aeruginosa* and 31 (31.3%) other species of *Pseudomonas*. *P.aeruginosa* was isolated from 18 (26.5%) ICU patients (burn and infectious), 35 (51.45%) burn unit, 6 (8.8%) internal ward, and 9 (13.2%) emergency unit patients. Other

species of *Pseudomonas* were isolated from 11(35.5%) ICU (burn and infectious), 13 (41.9%) burn, 4 (12.9%) internal ward and 3 (9.7%) emergency unit cases. Source of majority of *Pseudomonas* isolates was wound (59.9%). Of these, 46 (63.23%) isolates were *P.aeruginosa* and 16 (51.61%) other *Pseudomonas* species (Table 1).

Regarding antibiotic susceptibility of isolates, our results revealed *P.aeruginosa* to be non-susceptible to cephalosporins, while majority of other *Pseudomonas* species were found resistant to ceftazidime (Table 2).

Discussion

It is noticeable that *Pseudomonas* play significant role in causing severe infections, especially in immunocompromised patients. Unfortunately, despite the progress in the field of antibiotics against *Pseudomonas*, mortality is still a concern (15). *Pseudomonas* are opportunistic bacteria which have become one of the most important pathogens in the second half of the last century, considering their resistance towards traditional therapeutic agents (16). In our study, majority of *Pseudomonas* spp. were isolated from wound specimen, which is in concordance with other published reports. A study conducted in Saudi Arabia disclosed majority of *Pseudomonas* to be isolated from wounds, followed by respiratory tract specimens and wound from out-patients (17). Another study performed in Turkey reported *Pseudomonas* from respiratory specimens of patients admitted in ICU to be common (18). Having substantially increased in recent years, acquired resistance to ceftazidime has been estimated to be 10 to 40% (19). In our study, among the 99 isolates of *Pseudomonas*, 76 isolates were resistant to ceftazidime. Majority of *P.aeruginosa* isolates were non-susceptible to ceftriaxone (80.88%); among other *Pseudomonas* species, non-susceptibility was related to ceftazidime (83.87%). In the present study, all of the resistance among



Samples	<i>P.aeruginosa</i>	<i>Pseudomonas.spp</i>	Total
	N (%)	N (%)	
Wound	43(63.23)	16(51.61)	59(59.59)
Urine	15(22.05)	11(35.48)	26(26.26)
Blood	8(11.76)	3(9.67)	11(11.11)
Tracheal tube	1(1.47)	1(3.22)	2(2.02)
Abscess	1(1.47)	0	1(1.01)
Total	68(68.68)	31(31.31)	99(100)

Table1: *Pseudomonas* isolates from different specimen

isolates of *P.aeruginosa* and other *Pseudomonas* species were related to ceftazidime (76.76%). The sensitivity of the bacteria to at least 3 of the antibiotics namely, amikacin, gentamicin, ciprofloxacin, ceftazidime and imipenem was increased from 13% in 1997 to 21% in 2000 in the US. (20). In a study conducted by Mihani and colleagues, it was reported that 71% of *P. aeruginosa* isolates obtained from the patients in Ahvaz Taleghani Hospital were resistant to ceftazidime, 41% to imipenem and 67% towards ciprofloxacin (21) which is similar to our findings. However, non-susceptibility varies from ward to ward, even among diverse hospitals. The published literature from Tehran, Kashan, Iran and Turkey reported *Pseudomonas* resistance to ceftazidime of 72.9% (16), 86% (22) and 84%, respectively (18); these are consistent with our study.

Most sensitivity of *P.aeruginosa* isolates was related to carbenicillin (33.8%), and among the other *Pseudomonas* isolates were related to Imipenem (32.2%). Among all isolates of *P.aeruginosa*, the sensitivity to imipenem was much more than other antibiotics in this study. In a study conducted by Shojaipour et al., 48% of *P.aeruginosa* isolates were resistant to imipenem (23). Rate of resistance to imipenem was reported (26%) in Latvia, 20.5% in South Korea (24) and 14%-18% in Spain (25). In Turkey also, the sensitivity of the isolates decreased and the resistance of *Pseudomonas*

to imipenem increased in recent years (26). In another study conducted by Munir and colleagues, cases of sepsis in newborns were investigated; later, the resistance of *Pseudomonas* isolates to imipenem reported to be 83.3% (23). In this study, the amount of imipenem resistance among isolates of *Pseudomonas* (*aeruginosa* and other species) was 61.61%, which indicates the high resistance to these antibiotics.

About 20.5% of *P.aeruginosa* isolates had intermediate resistant to ciprofloxacin and other *Pseudomonas* isolates had the most intermediate resistant to ceftriaxone. Generally, among all isolates of *Pseudomonas* the highest intermediate resistance was related to ciprofloxacin (16.1%). Based on the results presented in this article, antibiotics such as ceftazidime, ceftriaxone, ofloxacin, gentamicin and tobramycin, have lost their effectiveness against *Pseudomonas* to some extent, and probably this class of drugs will be useless in the foreseeable future. Although the use of antibiotics such as ceftazidime and ceftriaxone for treatment of infections caused by gram-negative bacteria are not recommended since long time ago, resistance to them is impressive. Imipenem, carbenicillin and ciprofloxacin can be more effective than other antibiotics. Nevertheless, according to the recent studies, it can be predicted that resistant bacteria will be replaced by the partly sensitive strains soon and treatment will be more difficult.

Antibiotics	<i>P.aeruginosa</i> isolated N (%)			<i>Pseudomonas.spp</i> isolated N (%)			Total <i>Pseudomonas</i> antibiotics Resistance N (%)		
	Resistant	Interme diate	Sensitive	Resistant	Interm ediate	Sensitive	Resistant	Intermediate	Sensitive
Ofloxacin	47(69.11)	3(4.41)	18(26.47)	25(80.64)	0	6(19.35)	72(73.72)	3(3.03)	24(24.24)
Ciprofloxacin	37(54.41)	14(20.58)	17(25)	25(80.64)	2(6.45)	4(12.90)	62(62.62)	16(16.16)	21(21.21)
Gentamicin	50(73.52)	2(2.94)	16(23.52)	23(74.19)	1(3.22)	7(22.58)	73(73.73)	3(3.03)	23(23.23)
Amikacin	43(63.23)	5(7.35)	20(29.41)	22(70.96)	2(6.45)	7(22.58)	65(65.65)	7(7.07)	27(27.27)
Imipenem	41(60.29)	5(7.35)	22(32.35)	20(64.5)	1(3.22)	10(32.25)	61(61.61)	6(6.06)	32(32.32)
Ceftriaxone	55(80.88)	1(1.47)	12(17.64)	18(58.06)	5(16.12)	8(25.80)	73(73.73)	6(6.06)	20(20.20)
Ceftazidime	50(73.52)	2(2.94)	16(32.52)	26(83.87)	2(6.45)	3(9.67)	76(76.76)	4(4.04)	19(19.19)
Levofloxacin	46(67.64)	4(5.88)	18(26.47)	21(67.74)	2(6.45)	8(25.80)	67(67.67)	6(6.06)	26(26.26)
Tobramycin	48(70.58)	0	20(29.41)	24(77.41)	0	7(22.58)	72(72.72)	0	27(27.27)
Carbenicillin	42(61.76)	3(4.41)	23(33.82)	21(67.74)	2(6.45)	8(25.80)	63(63.63)	5(5.05)	31(31.31)

Table2: Antibiotic resistance profile among *Pseudomonas* isolated from patients

Antibiotic resistance of *Pseudomonas* to different antibiotics is distinct around the world and even this resistance is also different between various species of *Pseudomonas*. As a result, the correct treatment against this type of infection requires careful consideration of drug resistance of infectious agent and then prescription of the useful medicine. On the other hand, emergence of resistant strains can

be partly avoided by completing the treatment duration and not using antibiotics inappropriately as much as possible.

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