PHENOTYPIC ASSAYS FOR DETECTION OF ESBL AND MBL PRODUCERS AMONG THE CLINICAL ISOLATES OF MULTIDRUG RESISTANT PSEUDOMONAS AERUGINOSA FROM A TERTIARY CARE HOSPITAL

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ABSTRACT

Introduction: Pseudomonas aeruginosa has been a major nosocomial pathogen associated with nosocomial pneumonia, surgical site infections and UTI in patients admitted to intensive care units in the recent past(1). Major risk factors includes prolonged hospitalization, ventilation, underlying immunocompromised state and inadequate or irrational antimicrobial therapy(2). Despite improvements in therapy due to introduction of newer antimicrobials, P. aeruginosa is intrinsically resistant to number of antimicrobials, they aroused a major challenge to overcome the morbidity and mortality caused by multidrug and pan drug resistant P. aeruginosa(3). Drug resistance in turn leads to prolonged hospital stay and increased expenditure, which causes increased cross infections and poorer clinical outcomes. The present study investigated the prevalence of resistance mechanisms among Multi Drug Resistant Pseudomonas aeruginosa (MDRPA) clinical isolates from a tertiary care hospital.

Methods: Seventy-five MDRP. aeruginosa isolates were obtained from 226 patients admitted in various wards. Antimicrobial susceptibility testing was performed by disk diffusion method and all these isolates were found to be MDR. All the isolates were subjected to different phenotypic assays to detect the production of enzymes such as ESBL, AmpC and MBL,. MIC determination was done by agar dilution method for Meropenem and Polymyxin B. Further, quantitative evaluation of biofilm production by was carried out by microtitre plate assay, since many studies have shown positive correlation between MDR and biofilm formation.

Results: Of the 75 MDR P. aeruginosa, 36% were resistant to imipenem and 80% to meropenem. All the isolates were sensitive to polymyxin B. MBL production (38.67%) was found to be the predominant resistance mechanism followed by ESBL production (26.67%). None of them showed AmpC production. Ninety three percent (93%) of the strains produced abundant biofilms.

Conclusion: P. aeruginosa was shown to be predominant nosocomial pathogen showing resistance to most of the available antibiotics including carbapenems. MBL is shown to be predominant mechanism for development of resistance in the present study.

Keywords: MDR, Pseudomonas aeruginosa, ESBL, MBL.

INTRODUCTION

Pseudomonas aeruginosa is one of the significant Gram-negative bacteria causing hospital-associated infections(4). Among the common multidrug resistant (MDR) nosocomial pathogens emerged in medical centers, P. aeruginosa is the most frequent pathogen causing life threatening infections markedly, respiratory tract infections, surgical site and urinary tract infections in patients from intensive care units (ICUs)(5). It has
significant role in causing chronic debilitating respiratory infections in cystic fibrosis patients due to mucoid strains, which leads to increased mortality. Prolonged endotracheal intubation, associated with exposure to inappropriate antimicrobial therapies leads to colonization of the upper respiratory tract thereby complicates the eradication. 

A major challenge has aroused regarding the treatment of infections caused by opportunistic pathogens, predominantly those with pan drug resistant \textit{P. aeruginosa} and \textit{Acinetobacter baumannii} strains, which has extreme ability to acquire resistance. \textit{P. aeruginosa} possesses the ability to acquire resistance genes from the environment as well as from other bacteria. High mortality may be attributable to the inherent virulence of the organism as well as the fact that it often occurs with immunosuppression and co morbidity conditions. In addition, \textit{P. aeruginosa} is susceptible to a limited number of antimicrobial agents, which increases the likelihood of inappropriate empirical antimicrobial therapy. Reported rates of Multidrug Resistant \textit{Pseudomonas aeruginosa} (MDRPA) varied from 0.6% - 32% according to various surveillance studies held in different geographic locations. The prevalence of MDRPA has increased over the past decade and has become a major concern among hospitalized patients. Several mechanisms can contribute to resistance in \textit{P. aeruginosa}, including β-lactamase production, up regulation of efflux systems, biofilm formation and decreased outer membrane permeability.

However, production of β- lactamases such as Metallo- β- lactamases (MBL) and Extended spectrum β-lactamases (ESBL) are the most common resistant mechanisms documented in \textit{P. aeruginosa}. As carbapenems are the most potent β-lactams against \textit{P. aeruginosa}, intensive use of carbapenems facilitated the emergence of carbapenem-resistant \textit{P. aeruginosa}. Resistance to all antibiotics except polymyxins is now a reality in many medical centers. Despite of abundant literature, little is known about the prevalence of these mechanisms among the clinical isolates of \textit{P. aeruginosa} from India. Hence, this study was carried out to understand the prevailing mechanisms of resistance among the clinical isolates of MDRPA from in-patients of a tertiary care hospital. This possibly will help to prevent the associated morbidity and mortality caused by this organism by implementing proper infection control measures, which can reduce the duration of hospital stay and expenditure.

\section*{MATERIALS AND METHODS}

\subsection*{Study design}

A descriptive study was conducted during the period of November 2009 to October 2010. Seventy-five MDR \textit{P. aeruginosa} isolates were obtained from 226 patients admitted in various wards of the hospital. All the clinically significant MDRPA isolates collected in the Department of Clinical Microbiology laboratory from wound swabs, endotracheal aspirates, urine, blood, bronchoalveolar lavage, drain tip and tissue samples were investigated. Repeat isolates were excluded from the study. All the isolates were identified by standard microbiological techniques. ATCC \textit{P. aeruginosa} 27853 strain was used as quality control reference strain for all experiments with satisfactory results.

Antibiotic susceptibility testing was performed by Kirby-Bauer disc diffusion method. All these isolates were found to be MDR and were responsible for clinically significant infections. Isolates were also tested for carbapenemase production, if they were found to be intermediate or resistant to either Imipenem or meropenem in disc diffusion method. MIC values were determined by agar dilution method for clinically relevant antibiotics such as meropenem (Astra Zeneca, Bangalore) and polymyxin B (Hi-Media, Mumbai) as per CLSI guidelines.

All the MDR \textit{P. aeruginosa} were subjected to different phenotypic assays to detect the
production of enzymes such as ESBL, AmpC and MBL, which are implicated for causing multiple drug resistance. Phenotypic confirmatory test for ESBL production was performed by placing ceftazidime (30 μg) and ceftazidime + clavulanic acid disc. Detection of Metallo-β-lactamases was carried out by combined disc diffusion test. AmpC detection was done using AmpC discs method.

**Microtitre plate assay for Biofilm production**

Since many studies have shown positive correlation between biofilm and multiple drug resistance, quantitative evaluation of biofilm production by *P. aeruginosa* isolates was carried out by microtitre plate assay (16), (17). Biofilm negative *E. coli* isolate from our collection and ATCC *P. aeruginosa* 27853 were used as negative and positive controls respectively. Based on the OD values, the extent of biofilm formed by the clinical isolates were classified as follows,

- OD ≤ ODc - Non adherent
- ODc < OD ≤ 2 × ODc - Weakly adherent
- 2 × ODc < OD4 × ODc - Moderately adherent
- 4 × ODc < OD - Strongly adherent

**RESULTS**

Among the seventy-five patients with MDRPA infection, males were found to be more predominant (66.7%) than females (33.3%). MDR *P. aeruginosa* infections were found to be 34.7% between 21 to 40 years of age group. MDR *P. aeruginosa* were mainly isolated from wound infections (43%), followed by endotracheal aspirates (19%), urine (16%), and blood (11%), bronchoalveolar lavage (BAL)( 7%), drain tip (3%) and tissue (1%) respectively(Fig.1). Various risk factors for MDRPA infection were patients with prolonged hospitalization (52%), patients who are on Foley’s catheter (45%), patients with diabetes mellitus (37%), on ventilator (25%), on tracheostomy (21%), post operative patients (16%), on central line catheter (12%) and on steroids (1%). Thirty-three percentage of patients (n=25) were found to have more than two risk factors.

MDRPA isolates showed markedly high-level resistance towards ciprofloxacin (95%), tobramycin (92%), ceftiraxone and gentamicin, (83%). Forty-four isolates (59%) showed resistance to amikacin and 51% resistance was noticed for piperacillin + tazobactum. Among carbapenems, imipenem showed 36% resistance and meropenem 53% resistance. None of the isolates was resistant to polymyxin B. Among the 75 MDRPA isolates, 13 isolates, which were isolated from urine samples, showed 77% of resistance to norfloxacin and carbenicillin. MIC for meropenem ranged from 0.5μg/ml to >64μg/ml. Forty out of seventy five (53.33%) isolates of MDRPA were found to be resistant to meropenem. The isolates were categorized resistant if the MIC value was more than 8μg/ml. Thirty two (42.67%) isolates showed sensitive MIC value (≤4μg/ml) and 4% (3) isolates showed intermediate MIC value (8μg/ml). All the isolates showed lower MIC of 0.5μg/ml to 1μg/ml for polymyxin B (Break point MIC for *Pseudomonas aeruginosa* ≤2μg/ml to ≥8μg/ml) and none of these isolates showed resistant or intermediate MIC values (Fig.2).

Thirty-nine percent of isolates were found to produce MBL and only, 27% isolates showed ESBL production, none of the isolates showed AmpC production. Twelve percent (9) of isolates were found to be negative for phenotypic production of all β-lactamases (fig.3). Almost all isolates (93%) from our collection were biofilm producers, wherein 75% of the isolates were strongly adherent, 8% moderately adherent and 11% weakly adherent. The percentage of non-adherent cells or biofilm negative isolates was found to be a meager of 7%.
DISCUSSION

P. aeruginosa is a leading cause of nosocomial infections, and it exhibits intrinsic resistance to almost all commercially available antimicrobial agents. However, acquired resistance to anti-pseudomonal β-lactam antibiotics such as ticarcillin, piperacillin, ceftazidime, cefepime, aztreonam and especially carbapenems can be a major challenge in managing MDRPA infections, mostly when it is associated with co-resistance with other classes of drugs namely aminoglycosides and quinolones. Acquired ESBL and metallo-β-lactamases are the major β-lactamases produced by P. aeruginosa.

The prevalence of MDR P. aeruginosa was found to be 33.2% in our investigation, which is in accordance with a similar recent study from India that showed the predominance rate of 44% (18). However, comparatively lesser prevalence of MDRPA (26.7%) responsible for burn wound infections in Iran (19). Likewise, one more investigation from India reported 22% MDRPA and 4% Pandrug resistant P. aeruginosa (20), wherein, we recorded a higher level of MDRPA incidence. Factors such as age and sex among MDRPA infection were found to have significant association with MDRPA, wherein the incidence was more among the age groups between 21 and 40 (34.7%) and males being predominant (67%) which is the likely case in earlier reports (21). Possibly, it may be due to high incidence of road traffic accidents among males, leading to hospitalization thereby high incidence of P. aeruginosa infection through catheterization.

Earlier investigations have reported the major source of MDRPA to be sputum, tracheostomy specimen, pus, respiratory tract, surgical sites and endotracheal aspirate (22, 23 24, 25). In the present study, the major source of MDRPA was found to be wound swabs (43%), followed by endotracheal aspirate (19%) implying that wound infections and respiratory tract infections are most significant infections caused by MDRPA in most of the hospitals including our setup. The major risk factors were prolonged hospitalization followed by patients on Foley’s catheter. Foot infections and surgical site infections were found to be common source of MDRPA among the diabetic patients.

MDRPA isolates showed markedly high-level resistance towards ciprofloxacin (95%), tobramycin (92%), ceftriaxone and gentamicin, (83%). Forty-four isolates (59%) showed resistance to amikacin and 51% resistance was noticed for piperacillin/ tazobactum combination. Among carbapenems, imipenem and meropenem resistance was observed to be 36% and 53% respectively. None of the isolates were resistant to polymyxin B. Among the 75 MDRPA isolates, 13 isolates, which were isolated from urine samples, showed 77% of resistance to norfloxacin and carbenicillin.

MIC for meropenem ranged from 0.5µg/ml to >64µg/ml. Forty out of seventy five (53.33%) isolates of MDRPA were found to be resistant to meropenem. The isolates were categorized resistant if the MIC value was more than 8µg/ml. Thirty two (42.67%) isolates showed sensitive MIC value (≤4µg/ml) and 4% (3) isolates showed intermediate MIC value (8µg/ml). All the isolates showed lower MIC of 0.5µg/ml to 1µg/ml for polymyxin B (Break point MIC for Pseudomonas aeruginosa ≤2µg/ml to ≥8µg/ml) and none of these isolates showed resistant or intermediate MIC values.

Among various mechanisms of resistance, MBL and ESBL enzymes were found to be more effective and the incidence of MBL production in P. aeruginosa has been reported to be 10-30% from different clinical setups in India (26). Recent studies have shown a very high incidence of MBL (47%), AmpC (50%) and ESBL (13.3%) among the MDRPA isolates tested (29). In our study, also we have observed a high prevalence of MBL (39%), ESBL (27%) producers and 22% of MDRPA isolates were found to produce both MBL and ESBL, which appears to be significant. Around 12 % of isolates did not show any of the
mechanisms studied which might follow altogether different resistance mechanisms like formation of biofilms and/or cell wall permeability defects and efflux pump mechanisms (27).

High percentage of biofilm producers were observed in our study, which may be due to the increase number of MDRPA isolates encountered. Morten Hentzer et al. earlier reported the strong correlation between biofilm formation and multiple drug resistance in Gram-negative pathogens (29). Thus, biofilm formation appears to be one of the mechanisms among these strains to develop multi drug resistance, which is evident from the earlier reports from India (28).

CONCLUSION
In summary, MDR P. aeruginosa is a notable cause of hospital acquired infections and known to cause a wide spectrum of life threatening diseases. These organisms are resistant to almost all commonly available antibiotics with limited treatment options. Thirty-six percent of isolates showed resistance to imipenem and 53% to meropenem, which is an “alarming sign”, since carbapenems were the present drug of choice. Furthermore, 93% of isolates had the ability to form biofilm that might aid in the persistence of MDRPA thereby imparts resistance.

REFERENCES

Figure 1: Percentage of the various specimen from which MDRPA isolates were obtained

Figure 2: Percentage of Imipenem and Polymyxin B resistant and susceptible isolates based on their MIC values
Figure 3: Prevalence of various β-lactamses mediated resistance mechanisms among the clinical isolates of MDRPA based on phenotypic tests