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RESEARCH ARTICLE



Drug Resistance Mechanism among Acinetobacter Species

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Abstract

Acinetobacter species cause infections that are difficult to control due to multi-drug resistance and are noted for their intrinsic resistance to antibiotics and for their ability to acquire genes encoding resistance for the production of beta-lactamases and Aminoglycoside-modifying enzymes. MBLs are molecular class B and functional group 3 beta-lactamases which have the capability of hydrolyzing all β-lactams except the Monobactam, Aztreonam. Of several MBLs, only IMP, VIM and SIM types have been detected in these species. To analyze the antibiotic resistance patterns among Acinetobacter isolates and to detect Carbapenemase and MBL among MDR Acinetobacter isolates. The descriptive study of all phenotypically identified strains and multi-drug resistant strains of Acinetobacter species was conducted. A total of 303 isolates were isolated from various samples. They were processed and identified by standard Microbiological procedures. The antibiotics susceptibility testing was performed by Kirby- Bauer disc diffusion method using CLSI guidelines. Carbapenemase production was detected by employing 3 phenotypic test methods (MHT, CDM and DDST). Of 6355 samples processed, 303 were found to be Acinetobacter species, among those 50 were multi-drug resistant strains. The highest isolation of MDR Acinetobacter was from endotracheal tube tip (42%) and pus sample (32%). Majority of MDR Acinetobacter infection was found in male patients 36 (72%) compared to female patients 14 (28%). Majority of the strains were isolated from patients >/ 60 years of age group (%). A number of these isolates were more from ICU wards (30%) followed by Surgery wards (24%). Higher resistance for the Piperacillin/tazobactam ((82%), followed by Ceftazidime (80%), Imipenem (76%) etc. and the most susceptible drug was found to be the Tigecycline (82%) followed by Colistin (80%). Carbapenemase production was detected by MHT and 24 (48%) isolates were MHT positive. MBL production was detected by CDM and 34 (68%) isolates were CDM positive and by DDST 30 (60%) isolates were positive. Acinetobacter species are increasingly important nosocomial pathogens and are capable of rapid adaptation to the hospital environment. The variety of potential source of contamination or infection with these species in the hospital environment makes control of outbreaks caused by these difficult.

Keywords: Clinical samples, MBL, CDM, DDST, Antibiotic susceptibility

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INTRODUCTION

The genus Acinetobacter contains strictly aerobic, non-motile, Gram-negative, coccobacillary rods that are oxidase negative, nitrate negative and do not ferment sugars¹. This species is the second most frequent nonfermenter encountered in clinical laboratories but with only about 1/10th the frequency of Pseudomonas aeruginosa². In humans, it has been isolated from all culturable sites. It can be part of the bacterial flora of the skin, particularly in moist regions such as the axillae, groin and toe webs, and up to 43% of healthy adults can have colonization of skin and mucous membranes with higher rates among hospital personnel and patients. Predisposing factors include the presence of a prosthesis, endotracheal intubation, intravenous catheter and prior antibiotic therapy in seriously ill patients in hospital³. The colonization of bacteria on human mucosal surface tissues is a prerequisite for an infection. The attachment of bacteria to human, animal or plant host tissues requires adhesion factors. The property of adhesion to human epithelial cells in the presence of fimbriae and / or capsular polysaccharide^{4,5}. These species cause infections that are difficult to control due to multi-drug resistance. They are noted for their intrinsic resistance to antibiotics and for their ability to acquire genes encoding resistance for the production of beta-lactamases and Aminoglycoside-modifying enzymes⁶. In addition, diminished expression of outer membrane proteins, mutations in topoisomerases, and upregulation of efflux pumps also play an important part in antibiotic resistance. Common to all strains of A. baumannii is chromosomally encoded cephalosporinase (AmpC type). In addition to the class C cephalosporinase, other β-lactamases like TEM-1 type, SHV type, CTX-M type, PER-1, and VEB-1 have been reported.7 The carbapenem resistance was associated with the overexpression of OXA-51 and OXA-23 class D carbapenemases as well as of metallo- β -lactamases (MBLs).8 Aminoglycoside resistance is due to presence of no less than nine distinct modifying enzymes, which can be found in different combinations in some strains. Resistance to fluoroguinolones is correlated with point mutations of genes encoding DNA gyrase and topoisomerase IV. Tetracyclines resistance has been linked with genes that encode tetracycline-specific efflux pumps.9 Outer membrane proteins (OMPs) alters bacterial membrane permeability which is associated with loss or decreased expression of porins. This group is represented by OmpA, OprD, and CarO proteins.¹⁰.

In clinical practice, Acinetobacter infections are associated closely with surgery or the use of artificial devices. The local circumstances of clinical units and their environment determine the type of infection, and the consequent risk of dissemination and an outbreak occurring. It is considered to be a low-grade pathogen, and can remain on or in the human body without causing illness¹¹. This species can cause suppurative infections in virtually every organ system. Although it is acknowledged to be an opportunist in hospitalized patients, communityacquired infections are reported¹². Multi-drug resistant strains of this isolates are a growing problem and have been widely reported and the rapid development of significant Quinolone resistance in France, Aminoglycoside resistance in Germany and Carbapenem resistance in selected regions worldwide raises an important therapeutic problem¹³.

MATERIALS AND METHODS

The study was undertaken at the Department of Microbiology, JSS Hospital, Mysore for a period of 6 months from July to December, 2013.

Type of study: Descriptive study Inclusion criteria

Phenotypically identified strains of *Acinetobacter* species isolated from different samples only are included for study.

Multidrug resistant *Acinetobacter* species isolated from different samples.

Exclusion criteria

Other gram negative bacilli isolated are excluded.

Various clinical samples such as urine, pus, sputum,pleural fluid, etc.., were examined for isolation and identification of *Acinetobacter* and antimicrobial susceptibility testing.lsolates were processed and identified by standard Microbiological procedures¹⁴. The antibiotic susceptibility testing was performed by Kirby-Bauer disc diffusion method using Clinical and Laboratory Standard Institute (CLSI) guidelines¹⁵.

The following antibiotics were tested: imipenem(10µg)), tigecycline(15µg), gentamycin (10µg), ampicillin (10µg), cotrimoxazole(23.75/1.25µg), Piperacillin/ Tazobactum(100/10µg), ceftriaxone (30µg), ceftazidime(10µg), norfloxacin(10µg), nitrofurantoin (300µg), amikacin (30µg), levofloxacin (5µg), minocycline (30µg), colostin(10µg). Resistance data were interpreted according to Clinical laboratory Standards Institute (CLSI, 2013).

Detection of MBL

Carbapenemase production was detected by employing 3 phenotypic test methods (Modified Hodge Test, Combined Disk Method and Double Disk Synergy Test).

 Table 1. Total number of isolates obtained from different samples.

Sample	Total Number	No of Acinetbacter	MDR Acinetobacter
ET	205	87	21
Pus	1113	67	16
Urine	2089	22	06
Sputum	1178	89	05
PF	122	06	01
Blood	1648	32	01
Total	6355	303	50

Modified Hodge Test

Escherichia coli ATCC 25922, at a turbidity of 0.5 McFarland standards, was swabbed on the surface of Mueller Hinton agar plate, a 10 μ g Imipenem disc was placed at the centre and the test strain was streaked from the edge of the disc to the periphery of the plate in four directions. After overnight incubation, the plates were observed for the presence of "Clover leaf shaped" zone of inhibition. The plates with the presence of a distorted inhibition zone were interpreted as a positive result for Carbapenem hydrolysis screening.

Imipenem-EDTA Combined Disk Method

The test organisms were inoculated on Mueller-Hinton agar as recommended by the CLSI. Two 10 μ Imipenem discs were placed on the plates and appropriate amount of 10 μ l of EDTA solution were added to one of them to obtain the desired concentration. After 16-18hrs of incubation at 35°C.if the increase in inhibition zone size was more than 7mm than the Imipenem disc alone, it was considered as MBL positive.

Double Disk Synergy Test

The test inoculums was spread onto Mueller-Hinton agar. A disc of Imipenem $10\mu g$ + sterile disc were placed on the surface of MHA, 0.5M EDTA solution were added to the sterile disc. Discs were kept at a distance between 20mm. The plate was incubated at 37°C overnight. The organisms were considered to be forming the zone Imipenem discs showed a clear-cut increase (synergy) towards the sterile disc. The increase

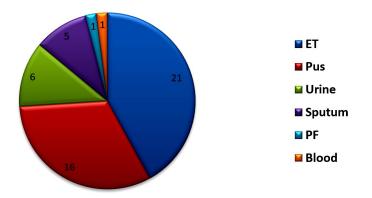




Fig. 1. Number of MDR Acinetobacter isolates isolated from various clinical Samples.

occurs because the clavulanic acid inactivates the ESBL produced by the organism resulting in the formation of extended inhibitory zone.

Statistical analysis

Data analysis was done using MS Excel. Ethical considerations

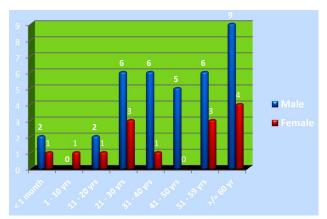
Ethical clearance was obtained from Institutional Ethical Clearance Committee of JSS Medical college, Mysore.

RESULTS

Of 6355 bacterial isolates from different clinical samples 303 (4.76%) were found to be Acinetobacterspecies. Among the 303 of these species 50(16.5%) isolates were recognized as MDR Acinetobacter species. shown in Table 1. Highest isolation of MDR Acinetobacterwas from endotracheal tube tip (42%) and pus sample (32%). Other samples included urine, sputum, pleural fluid and blood samples as shown in Fig. 1. Table 2 shows that out of 50 isolates, 36 (72%) were isolated from male patients and 14 (28%) from female patients and the Male to female ratio was 2.5:1. As shown in Fig. 2 majority of the strains were isolated from patients > 60 years of age group probably because of increased ICU admission in this age group and presence of other co-morbid conditions. Wardwise distribution of

 Table 2. MDR Acinetobacter isolates in relation to gender

Male	Female	Total	
36	14	50	



Acinetobacterstrains were showed in Table 3 and isolates were more from ICU wards (30%) followed by surgical wards (24%). Table 4 shows antibiotic sensitivity pattern. The resistance pattern of these isolates showed that there was high resistance for the Piperacillin/tazobactam (82%), followed by Ceftazidime (80%) and Imipenem (76%) etc. The most susceptible drug was found to be the Tigecycline(82%) followed by Colistin (80%). Out of 50 MDR Acinetobacter isolates, 24 (48%) isolates were MHT positive. MBL production was detected in 34 (68%) isolates by CDM and 30 (60%) isolates by DDST as shown in Table 5. and in photograhs 1 and 2. Test results obtained by different combination of the test methods for MBL production was shown in Table 6 which shows 7 (14%) isolates were positive by both MHT and CDM, 12 (24%) isolates were positive by both CDM and DDST and 6 (12%) isolates were positive by MHT and DDST. Number of Positive and Negative

 Table 3. Ward wise distribution of MDR Acinetobacter isolates

Wards	Total No. of isolates	
Private ward	02 (4%)	
Medical	11 (22%)	
Surgical	12 (24%)	
Pulmonology	02 (4%)	
ICU (SICU, RICU, etc)	15(30%)	
Orthopaedic	02 (4%)	
Pediatric	02 (4%)	
NICU	03 (6%)	
OBG	02(4%)	
Total	50	

Fig. 2. Number of MDR Acinetobacter isolates in relation to Age-gender group.

isolates by all the 3 methods is shown in table 7 which shows total number of MBL positive isolates obtained by all the 3 methods were 8 (16%) and MBL negative isolates by all the 3 methods were found to be 03 (6%).

DISCUSSION

The discovery and development of antibiotics was undoubtedly one of the greatest advances of modern Medicine. Unfortunately, the emergence of antibiotic resistance in general and also among Nonfermenters especially *Acinetobacter* is threatening the effectiveness of the antimicrobial agents in the treatment. To prevent cross transmission via the hands of the staff, strict hand washing policy and practice should be enforced.¹⁶ Emergence of newer microbes and ability of the bacteria to gain multidrug resistance by production of enzymes and other mechanisms has posed a serious threat in the management of nosocomial infections. Recently few Gram negative non fermentative bacilli with acquired metallo beta lactamase production have been increasingly reported in some countries.

In our study *Acinetobacter* species constituted 4.7% of the total bacteria isolated from different samples. Similar observations were



Plate 1. Mueller Hinton agar plate showing Modified hodge test positive.



Plate 2. Mueller Hinton agar plate showing Combined disk test positive and Double disk synergy test positive

Antibiotics	Sensitive	Intermediately sensitive	Resistance	
		Sensitive		
Imipenem		12 (24%)	38 (76%)	
Tigecycline	41 (82%)	09 (18)		
Gentamycin	09 (18%)	08 (16%)	33 (66%)	
Ampicillin	15 (30%)	06 (12%)	29 (58%)	
Cotimoxazole	11 (22%)	06 (12%)	30 (60%)	
Piperacillin/Tazobactum	05 (10%)	04 (8%)	41 (82%)	
Ceftriaxone	02 (4%)	13 (26%)	35 (70%)	
Ceftazidime	05 (10%)	05 (10%)	40 (80%)	
Norfloxacin	05 (10%)	10 (20%)	35 (70%)	
Nitrofurantoin	03 (6%)	11 (22%)	36 (72%)	
Amikacin	06 (12%)	19 (38%)	25 (50%)	
Levofloxacin	10 (20%)	19 (38%)	21 (42%)	
Minocycline	19 (38%)	11 (22%)	20 (40%)	
Colistin	40 (80%)	10(20%)		
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Table 4. Antibiotic sensitivity pattern of MDR Acinetobacter isolates to different antibiotics

 Table 5. Test Result of Carbapenemase and MBL

 detection by different Methods

Name of the Test	Total number of isolates	Number of positive isolates	Number of negative isolates
MHT	50	24	26
CDM	50	34	16
DDST	50	30	20

made by Rubina et al. and Hisham et al which showed 4.8% and 3.8% respectively^{17,18}. Most of the strains were isolated from ET samples (42%). Queenam et al isolated more MDR Acinetobacter isolates from lower respiratory tract¹⁹. 16.5% MDR Acinetobacter were isolated in the present study. Lt Col KK Lahiri et al studied antibiotic resistance pattern of Acinetobacter in their study and has shown multidrug resistance in 106 (69.7%) isolates from a total of 152²⁰. In our study out of 50 isolates, 36 (72%) were isolated from male patients, and 14(28%) were from female patients. Similar gender preponderance was observed by Purti et al²¹. Majority of the strains were isolated from patients > 60 years of age group. Similar findings were observed by Queenan AM and Pillar et al who isolated MDR Acinetobacter more among elderly patients (65%)¹⁹.

In this study, MDR *Acinetobacter* were isolated more from Critical care area (30%) followed by Surgical ward (24%). Similar observations were made by Purti et al, and Patwardhan et al.^{19,22}. These isolates showed increased resistance to Carbapenems, Aminoglycosides and Cephalosporins (Imipenem (76%), Gentamicin (66%),Ceftriaxone(70%). Concordant findings were observed by Rubina Lone et al ¹⁷and Nazmul MHM et al.⁶.

In our study it was noticed that CDM was a better method for detection of MBL among the 3 methods performed. Irfan et al. in a study on Metallo- β -lactamase – producing clinical isolates of *Acinetobacter* species and *Pseudomonas aeruginosa* from intensive care unit patients of a tertiary care hospital concluded in their study that, by using imipenem-EDTA disk method (CDM) as a screening test for metallo- β -lactamase production they found a very high percentage of metallo- β -lactamase producing isolates among **Table 6.** Test results obtained by different combination

 of the test methods for MBL production

Positive by	Positive by	Positive by
MHT and CDM	CDM and DDST	MHT and DDST
07	12	06

 Table 7. Number of Positive and Negative isolates by all the 3 methods

Positive by all the 3 Methods	Negative by all the 3 Methods	
08	03	

multidrug-resistant Acinetobacter species and *Pseudomonas aeruginosa* isolates²³. Priyam et al conducted study on comparison of genotypic and phenotypic methods of metallo- β - lactamase detection in Acinetobacter spp. PCR analysis of the 100 strains showed that 3 (3%), 5 (5%), 7 (7%), 26 (26%), and 51 (51%) strains had IMP gene, VIM gene, KPC gene, OXA gene, and NDM-1 gene, respectively and also found that MHT on MacConkey agar (MAC) had better sensitivity when compared with the gold standard PCR than MHT on Mueller–Hinton agar (MHA).²⁴

As PCR cannot be done in every laboratory, because of its high cost, the disc potentiation test for MBL detection should be introduced in any routine Microbiology laboratory for effective infection control and to prevent therapeutic failure. The Disc potentiation test is easy to perform, is cost effective and quite specific amongst other phenotypic methods which are used for MBL detection. The Disc potentiation test can also be done along with routine antibiotic sensitivity tests.

CONCLUSION

Acinetobacter are the "superbugs" of the modern hospital environment causing significant proportion of infections in specific patient populations, especially in critically-ill patients in the ICU's. As ubiquitous organisms (fortunately of low virulence), with few requirements for growth and survival, Acinetobacter species. are prone to persist indefinitely in the hospital environment and to cause infections periodically when iatrogenic factors are present, i.e., overuse of broad spectrum antibiotics and high-risk patients. Antibiotic resistance is attributed to production of extended spectrum beta-lactamase, MBL, loss of outer membrane proteins, efflux pumps and biofilm formation.

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CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

AUTHORS' CONTRIBUTION

JS collected the statistical data, drafted the manuscript, compiled the information and designed the tables. RKG gave her valuable advice, constant guidance, moral support and inspiration in the preparation of this work. MDW supervised, supported and reviewed the manuscript and tables.

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None.

DATA AVAILABILITY

All data analysed during this study are included in the manuscript.

ETHICS STATEMENT

Ethical clearance was obtained from Institutional Ethical Clearance Committee, India.

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