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PREVALENCE OF MULTI DRUG RESISTANT GRAM-NEGATIVE BACTERIA AND ITS SUSCEPTIBILITY PATTERN ISOLATED FROM ADMITTED PATIENTS IN A HOSPITAL OF CENTRAL INDIA; AN EMERGING GLOBAL PROBLEM

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Abstract

Introduction: The multi-drug resistant gram-negative bacterial infections are the principle threats to the critically ill patients. Multidrug-resistant Gram-negative rods (MDR-GNR) are emerging as a major challenge to human health. The prevalence of drug-resistant cases is increasing globally. However, the impact of these infections on the patient's clinical outcome has not yet been clearly evaluated.

Aim: To determine the incidence and associated clinical outcome of multi-drug resistant gramnegative bacterial infections.

Materials and Methods: This record-based retrospective cross-sectional study was designed to analyse MDR-GNB positive cases at a tertiary care hospital of central India. MLB Medical college with attached. Hospital is a 700-bed referral tertiary care centre. An increase of MDR-GNB was seen from September 2022 to June 2023 in the hospital. A retrospective analysis of blood culture GNB-positive samples was performed to evaluate MDR-GNB-positive cases at admission.

Results: The total number of positive blood cultures in September 2022 to June 2023, November to January 2023 and February to April 2023 were 236, 186 and 206, respectively, with 76.83%,80.% and 71.83% GNB-positive. Total MDR-GNB-positive cases were 27.08%, 34.98% and 32.65%,

respectively, and amongst these MDR-GNB, 24%, 32% and 5% where positive at time of admission to the hospital. The MDR-GNB were *Escherichia coli*, *Klebsiella*, *Acinetobacter*, *Pseudomonas* and *Enterobacter*, *Klebsiella pneumoniae*, *Acinetobacter* spp., *Pseudomonas aeruginosa* and *Escherichia coli* have alarming degrees of antimicrobial resistance and are associated with high mortality and morbidity.

Conclusion: The incidence of multi-drug resistant gram-negative bacterial infections was remarkably high in our hospitals and showed a significant association with healthcare-associated infections and in-hospital-mortality.

Key words: Multi drug resistant, Pulmonary infiltrate, Excessive use of antibiotics.

Introduction

Multidrug-resistant Gram-negative bacterial infections are emerging as a major challenge to clinicians, especially intensive care units for neonates, infants, children, young adults as well as for adults. Bacteraemia may be associated with high level of morbidity and mortality. Emperical antibiotic therapy without confirming susceptibility pattern may be life threatening, increases the hospital stay, increases morbidity and in many cases it may leads to death.

Its an emerging health issue globally and commonly associated organism are *Klebsiella pneumoniae*, *Acinetobacter* baumannii, *Pseudomonas aeruginosa* and *Escherichia coli*. The global scenario shows that gram-positive infections are common in the developed countries ICUs [1]. However, multidrug-resistant gram-negative bacteria (MDR-GNB) infections dominate in the Asia-Pacific region [2, 3]. Among MDR-GNB, extended-spectrum beta-lactamases (ESBL) organisms, carbapenemase producing enterobacteriaceae, carbapenem-resistant *Acinetobacter species*, multidrug- resistant

Pseudomonas aeruginosa are the major culprits. Unfortunately, new antibacterial agents have not been developed in pace with the growth of multidrug-resistant (MDR) organisms [4].

As per the World Health Organization (WHO) fact sheet, infections by antimicrobial-resistant organisms can result in failure of treatment, increased cost of medical treatment, increased hospitalization stay and increased socioeconomic burden [5]. Infections caused by multidrug-resistant Gram-negative bacteria (MDR-GNB) have been reported to have increased significantly worldwide in recent years. The increase in MDR-GNB cases has become a serious challenge for health care professionals. Excessive use of antibiotics including the use without treatment indication is believed to be one of the major factors accelerating the spread of antibiotic resistance. A survey by Van Boeckel et al. on total antibiotic sales from 2000 to 2010 in 71 countries indicated that India is the country with the highest consumption of antibiotics (approx. 13 billion standard units, i.e. pill/capsule/or ampoule), followed by China (approx. 10 billion standard units) and the United States (more than 6 billion standard units). Consumption of antibiotics increased by 36% in this time period. Brazil, Russia, India, China and South Africa accounted for 76% of this increase [6,7].

Material and methods

This study was conducted at Government Prakash Chandra Sethi Hospital, Indore it is multispecialty, referral secondary care centre. It was observed that the MDR-GNB cases were rising from September 2022 to June 2023 in the hospital. However, there was no outbreak. The study was designed to analyze the prevalence and epidemiology of increase in MDR-GNB cases. Cases resistant to carbapenem, third and fourth generation cephalosporins were considered as MDR-GNB cases.

Study design

A retrospective, cross-sectional study design was used for the research. A retrospective analysis of data of all hospitalized patients from September 2022 to June 2023 was done for presence of MDR-GNB-positive blood culture samples seen at the time of admission to hospital. As a part of routine

practice, blood culture was drawn within 24 h for all the patients admitted with sepsis. Routine conventional Method of blood culture was used for antibiotic susceptibility and bacterial identification.

Data review procedure

Records of hospitalized patients from September 2022 to June 2023 were reviewed stepwise with the focus on results of blood culture analysis. Blood culture data showing positive MDR-GNB was separated and analyzed further for the type of GNB.

Statistical analysis

Data were entered in the MS Excel 2007 and analyzed with STATA version 14.

MICROBIOLOGICAL PROCEDURES

Pathogenic bacteria isolated from the clinical specimens from the various wards and ICU,NICU,SNCU,PICU etc. were further characterized by conventional biochemical tests to identify the specific GNB by using standard microbiologic methods [08]. Antibiotic susceptibility test of GNB strains was done by the Kirby Bauer disc diffusion method on Mueller Hinton agar (MHA) as per the Clinical Laboratory Standard Institute (CLSI) guidelines [09]. Antibiotics of following concentrations were used: ampicillin (10µg), amikacin (30µg), gentamycin (10µg), tobramycin(10µg), ciprofloxacin (5µg), levofloxacin (5µg), chloramphenicol (30µg), co-trimoxazole (25µg), ceftazidime (30µg), cefotaxime (30µg), cefepime (30µg), piperacillin (100µg), carbenicillin (100µg.), piperacillintazobactam (100/10µg), imipenem (10µg), tigecycline (30µg), polymyxin B (300unit), and colistin sulphate (10µg) from Hi Media Laboratories, India. Disk zone diameters were interpreted according to the CLSI 2017 recommendations. Quality control for culture plates and antibiotic susceptibility was performed using Escherichia coli ATCC 25922 and Pseudomonas aeruginosa ATCC 27853. All the strains were subjected to various phenotypic methods for the screening and confirmation of the beta lactamases. Strains showing decreased sensitivity to ceftazidime/ cefotaxime were considered as screen positive for ESBL production and were subjected to the following confirmatory phenotypic tests as per the CLSI guidelines [10].

- ESBL- A difference in the zone size of 5 mm between ceftazidime and ceftazidime+ clavulanic acid and cefotaxime and cefotaxime+clavulanic acid discs was considered as confirmed ESBL producer [10].
- Carbapenemase- The screen positive for carbapenemase production was considered for strains showing resistance to carbapenems. A positive modified hodge test (MHT) with appearance of clover leaf at the streaking line was considered as carbapenemase producer as per the CLSI guidelines .A difference in the zone size of 7 mm between Imipenem and Imienem+ EDTA disc in the EDTA disk synergy test was considered as MBL producer [10].

RESULT

A total 710 patient's samples were analyzed which included blood 174 ,body fluids 64 ,urine 232 pus 108 ,sputum 35,and Other samples(Vaginal Swab ,Catheter tip, Peritoneal Fluid, ET Suction Tip) 97 (Table 1). Total 362 samples were positive for growth of the MDR organisms. Mostly MDR organism isolated from urine samples around 89,followed by blood samples, around 58 isolates from pus and around 45 from different body fluids i.e. Ascitic fluid, synovial fluid, pleural fluid from patients admitted in intensive care units.

On culturing of 97 various catheters like ET Tube, Umbilical catheter tip, Foley's catheter & samples of Vaginal swab around 72 isolates were obtained.

Table.1 Sample prome from which MDK organishis isolated				
S. No.	Sample	Isolates	s MDR Isolates	
1.	Blood	174	72	

Table.1 Sample profile from which MDR organisms isolated

2.	Urine	232	89
3.	Pus	108	58
4.	Body fluids	64	45
5.	Sputum	35	26
6.	Other samples(Vaginal Swab ,Catheter tip, Peritoneal Fluid, ET Suction Tip)	97	72

From above given 710 isolates around 362 (52%) are Multi drug resistance organisms were isolated, out of which Escherichia coli was most pravelent organism followed by Citrobacter *spp*. and Klebsiella *spp*.

Among non-fermenters Pseudomonas *aeruginosa* is most prevalent followed by Acinetobacter *spp*. Around 38.8% of Pseudomonas are multidrug resistant shown in Table-2.

Among antibiotics ampicillin shows highest number of resistance pattern i.e. in E.coli. followed by Klebsiella.

S. No.	Organism	Isolates	MDR Isolates	Percentage
1.	Escherichia coli	256	159	62.10 %
2.	Klebsiella spp	124	52	41 %
3.	Citrobacter spp	180	98	54.4 %
4.	Enterobacter spp	09	02	2.2 %
5.	Proteus spp.	13	03	2.3 %
6.	Pseudomonas spp.	108	42	38.8 %
7.	Other Non-fermenting GNB	20	06	3 %

Table 2: Pattern of isolates Organisms Isolates

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Table 3: Number of	gram negative	bacteria resistant to	given anfimicrobial	agent
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Antibiotic	E.coli	Klebsiella	Citrobacter	Proteus	Pseudomonas
Amikacin	115(72%)	39(75%)	77(78%)	2(66%)	33(78%)
Ampicillin	146(91%)	44(84%)	67(68%)	3(100%)	17(40%)
Ciprofloxacin	76(47%)	24(46%)	52(53%)	1(33%)	27(64%)
Chloramphenicol	55(34%)	34(65%)	82(83%)	2(66%)	18(42%)
Carbenicillin	58(36%)	22(42%)	24(25%)	2(66%)	08(19%)
Cefepime	82(51%)	42(80%)	46(47%)	2(66%)	32(76%)
Cefotaxime	75(47%)	32(61%)	72(73%)	1(33%)	26(61%)
Co-Trimoxazole	37(23%)	42(80%)	52(53%)	1(33%)	22(52%)
Ceftazidime	67(42%)	25(46%)	84(85%)	1(33%)	29(69%)
Colistin Sulphate	39(24%)	22(42%)	39(40%)	1(33%)	07(16%)
Gentamicin	42(26%)	34(65%)	46(47%)	1(33%)	22(52%)
Imipenem	36(22)	12(23%)	26(27%)	1(33%)	08(19%)
Levofloxacin	58(36)	43(81%)	51(52%)	1(33%)	27(64%)
Tobramycin	23(14%)	18(34%)	26(27%)	1(33%)	10(27%)
Tigecycline	12(7.5%)	22(42%)	23(24%)	00	08(19%)
Piperacillin-	82(51%)	42(80%)	72(73%)	2(66%)	27(64%)
Tazobactam					
Polymyxin B	42(26%)	28(53%)	32(33%)	1(33%)	32(76%)

Discussion

Antimicrobial resistance is increasing worldwide, threatening global public health and the effective prevention and treatment of infections [8]. The U.S. National Healthcare Safety Network reported

increasing occurrence of MDR-GNB (*E. coli, K. pneumoniae, Enterobacter* spp., etc.) amongst which more than 60% were *Acinetobacter* spp. Similarly, in Europe the European Antimicrobial Resistance Surveillance Network reported recognizable resistance trends for Gram-negative bacteria, with highest levels of resistance reported for *Acinetobacter* spp., followed by *E. coli* and *K. pneumoniae* [2]. Reports from various studies from hospitals in India suggest that the prevalence of ESBL-producing GNB range between 19% and 60%, and that of carbapenem-resistant GNB between 5.3% and 59% [6]. In Mumbai, west India, the prevalence of drug-resistant Enterobacteriaceae was about 18.5% [9]. The prevalence of ESBL- and carbapenemase-producers in Kolkata was estimated to be 70% and 39%, respectively [10]. In south India, the occurrence of drug-resistant GNB was 53% of isolates from patients with community-acquired bacteraemia caused by *E. coli* and *Klebsiella* spp. [11]. This indicates the increased burden of MDR-GNB globally including India. In the present study, it was revealed that the numbers of blood culture-positive cases in hospital were increasing consecutively in the years 2012–2014, amongst which the GNB-positive cases were more than 70%, showing a similar trend of increase in GNB-positive cases as reported globally. *Acinetobacter*,

Klebsiella, E. coli and *Enterobacter* are the most common MDR-GNB isolated from neonatal septicaemia at tertiary hospitals [12]. Similarly, *E. coli* and *Klebsiella* are the most predominant organisms causing urinary tract infection (UTI) in children. There are several studies showing *E. coli* as the significant pathogen causing UTI [13], [14]. Fifty-three percent of Gram-negative organisms isolated from children were found to be multidrug-resistant [15]. Mandell et al. reported that Gram-negative bacteria were the predominant cause of UTI when compared with Gram-positive bacteria [16]. Our present study also supported the findings by various studies that *E. coli*, *Klebsiella*, *Acinetobacter* and *Pseudomonas/Enterobacter* were the GNB which were the most prominent cause of infections in hospitalized patients; however, there was a greater incidence of *Acinetobacter* and *Klebsiella* MDR-GNB amongst all other GNB. *Pseudomonas* spp. and

Acinetobacter spp. are the most common organisms isolated from ICU in south India and Delhi [17]. Javeri et al. also reported Acinetobacter as the second most common isolate in ICU of tertiary care centres in Ahmadabad [18] indicating the wide spread of MDR-GNB throughout India, supporting our findings. Overall, MDR-GNB cases increased during the period of 2021–2022. The number of MDR-GNB *Klebsiella* and Acinetobacter increased in consecutive years over the study period, indicating the need for analysis of MDR-GNB cases at primary and secondary care units to prevent the further spread of resistance.

Conclusion

MDR-GNB blood cultures positive at admission increased from July 2021 to June 2023 and hence there is an urgent need for possible contact isolation of all patients coming from primary and secondary to tertiary health care centres to be made compulsory until screening to rule out MDR-GNB has been performed, to prevent spread of MDR organisms in the hospital. The study findings will be part of a strict Antibiotic Stewardship (AMS) programme and also indicate that AMS should begin at primary and secondary health care centres to prevent antimicrobial resistance.

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Conflict of interest

The authors declare no conflict of interest.

Ethical approval Not required

Limitations

Patients who were in Incubation Period of nosocomial infections on discharge, who manifests it after discharge, were not covered in current study. Contribution of their load to current study prevalence is unknown.

References

- 1. Chaudhry D, Prajapat B. Intensive care unit bugs in India: How do they differ from the Western world? J Assoc Chest Physicians. 2017;5:10–7. https://doi.org/10.4103/2320-8775.196645.
- Mendes RE, Mendoza M, Banga Singh KK, et al. Regional resistance surveillance program results for 12 Asia Pacific nations (2011). Antimicrob Agents Chemother. 2013; 5 7(11):5721– 5726. doi: https://doi.org/10.1128/AAC.01121-13.
- 3. Parajuli NP, Acharya SP, Mishra SK, et al. High burden of antimicrobial resistance among gramnegative bacteria causing healthcare associated infections in a critical care unit of Nepal. Antimicrob Resist Infect Control. 2017;6:67. https://doi.org/10.1186/s13756-017-0222-z.
- 4. Khanal S, Joshi DR, Bhatta DR, et al. 훽-lactamase-producing multidrug-resistant bacterial pathogens from tracheal aspirates of intensive care unit patients at National Institute of neurological and allied sciences. Nepal ISRN Microbiology. 2013; https://doi.org/10.1155/2013/847569.
- 5. Boucher HW, Talbot GH, Bradley JS, et al. Bad bugs, no drugs: no ESKAPE! An update from the Infectious Diseases Society of America. Clin Infect Dis. 2009;48:1–12.
- 6. Martin-Loeches, E. Diaz, J. Valles Risks for multidrug-resistant pathogens in the ICU Curr Opin Crit Care, 20 (2014), pp. 516-524
- 7. T.P. VanBoeckel, S. Gandra, A. Ashok, Q. Caudron, B.T. Grenfell, S.A. Levin, *et al.* Global antibiotic consumption 2000 to 2010: an analysis of national pharmaceutical sales data Lancet Infect Dis, 14 (2014), pp. 742-750 Google Scholar
- 8. Washington CW Jr, Stephen DA, William MJ, et al. Koneman's color atlas and text book of diagnostic microbiology. 6th ed. Philadelphia: Lippincott Williams and Wilkins; 2006. Google Scholar
- 9. Clinical and Laboratory Standards Institute. Performance standards for antimicrobial susceptibility testing; 27th ed. CLSI supplement. *CLSI Document M100-S27*. Wayne, PA: Clinical and Laboratory Standards Institute; 2017.
- Yong D, Lee K, Yum JH, et al. Imipenem-EDTA disk method for differentiation of metallo-betalactamase-producing clinical isolates of pseudomonas spp. and Acinetobacter spp. J Clin Microbiol. 2002;40(10):3798–801 PMID: 12354884 Article CAS PubMed Pub Med Central Google Scholar
- 11. WHO Fact Sheet Antimicrobial resistance (2018) http://www.who.int/en/news-room/fact-sheets/detail/antimicrobial-resistance
- 12. M. Kazi, L. Drego, C. Nikam, K. Ajbani, R. Soman, A. Shetty, *et al.* Molecular characterization of carbapenem-resistant Enterobacteriaceae at a tertiary care laboratory in Mumbai Eur J Clin Microbiol Infect Dis, 34 (2015), pp. 467-472 Cross Ref View in Scopus Google Scholar
- G. Goel, L. Hmar, M. Sarkar De, S. Bhattacharya, M. Chandy Colistin-resistant *Klebsiella pneumoniae*: report of a cluster of 24 cases from a new oncology center in eastern India Infect Control Hosp Epidemiol, 35 (2014), pp. 1076-1077 View article Cross Ref View in Scopus Google Scholar
- K.P. Abhilash, B. Veeraraghavan, O.C. Abraham Epidemiology and outcome of bacteremia caused by extended spectrum beta-lactamase (ESBL)-producing *Escherichia coli* and *Klebsiella* spp. in a tertiary care teaching hospital in south India J Assoc Physicians India, 58 (2010), pp. 13-17
- 15. R. Srivastava, J. Agarwal, S. Srivastava, M. Kumar, M. Singh Multidrug resistant Gramnegative bacilli from neonatal septicaemia at a tertiary care centre in North India: a phenotypic and genotypic study Indian J Med Microbiol, 32 (1) (2014), pp. 97-98
- 16. P. Patel, R.N. Garala Bacteriological profile and antibiotic susceptibility pattern (antibiogram) of urinary tract infections in paediatric patients J Res Med Dent Sci, 2 (2014), pp. 20-25

- 17. R. Rajbhandari, J. Shrestha Bacteriological study of urinary tract infection and its antibiotic sensitivity test: a hospital based study J Nepal Asso Med Lab Sci, 4 (2002), pp. 26-32
- 18. S. Srinivasan, N.S. Madhusudhan Prevalence of multidrug resistant pathogens in children with urinary tract infection: a retrospective analysis Int J Med Res Health Sci, 3 (2014), pp. 954-958
- 19. G.L. Mandell, J.E. Bennett, R. Dolin Principles & practices of infectious diseases (7th ed.), Elsevier, St Louis (2010) Google Scholar
- 20. K. Moolchandani, A.S. Sastry, R. Deepashree, S. Sistla, B.N. Harish, J. Mandal Antimicrobial resistance surveillance among intensive care units of a tertiary care hospital in Southern India J Clin Diagn Res, 11 (2017) DC01-DC07 Google Scholar
- 21. J.R. Javeri, S.M. Patel, S.N. Nayak, K. Desai, P. Patel A study on bacteriological profile and drug sensitivity & resistance pattern of isolates of the patients admitted in intensive care units of a tertiary care hospital in Ahmadabad Nat J Med Res, 2 (2012), pp. 330-334 Google Scholar