

A Retrospective Study on Bacteria Causing Blood Stream Infection: Antibiotics Resistance and Management

S. K. SARKAR, A. BHATTACHARYYA¹, K. PARIYA², AND S. M. MANDAL^{3*}

Sanjiban Hospital, Fuleswar, Howrah-711 316, ¹TechnoIndia DAMA Hospital, Kolkata-700 091, ²Department of Zoology, Vidyasagar University, Midnapore-721 101, ³Central Research Facility, Indian Institute of Technology Kharagpur, Kharagpur-721 302, India

Sarkar *et al.*: Management of β -lactamase Positive Bacteria in Blood Stream Infection

Sepsis is one of the major cause of morbidity and mortality from infants to adults. It is most important to determine the infected bacterial species found in bloodstream and its antibiotic susceptibility pattern for appropriate treatment. Overall fifteen hundred patients' data were incorporated into our study, collected from Sanjiban Hospital, Howrah, West Bengal, India. Out of 1500 samples, 190 neonates and 250 adults were positive for bacterial sepsis. The strains expressing extended-spectrum β -lactamases are major threat as therapeutic options are limited. Among the isolates, Gram-positive bacteria were predominated (53.36 %) over Gram-negative bacteria (46.64 %) and *Staphylococcus aureus* was the most frequent isolate (37.6 %) followed by *Pseudomonas aeruginosa* (16 %), *Escherichia coli* (13.60 %), *Klebsiella* sp. (12.8 %), coagulase-negative *Staphylococci* (12.4 %), *Acinetobacter* sp. (4 %) and *Enterococci* sp. (2.8 %). Levofloxacin was revealed to be more active against all the Gram-negative isolates along with carbapenems, aminoglycosides (except *Klebsiella* sp.) and polymyxin-resistant strains. Levofloxacin gave superior coverage against both Gram-positive and Gram-negative bacteria whereas most of the penicillins and cephalosporins were found to be ineffective against both Gram-positive and Gram-negative isolates.

Key words: Bacteraemia, antibiotic resistance, blood stream infection, levofloxacin

Blood stream infection (BSI) is a significant reason of increased morbidity and mortality, one of the most frequently observed cases in the tertiary hospitals^[1]. It causes millions of deaths globally each year^[2]. In case of neonatal sepsis it contributes nearly 13-15 % of all deaths and in developing countries where it contributes between 30-50 %^[3]. The causative microorganisms of BSI have been found to be associated with increased drug resistance propensity to the conventionally used antibiotics and emergent global threats to control those multi-drug resistant pathogens^[4]. Several antibiotics have been successfully used for the treatment of sepsis, but recently most of the Gram-negative bacilli responsible for BSI became a serious therapeutic challenge due to development of multi-drug resistance^[5]. In most cases of the neonatal sepsis, extended-spectrum β -lactamase (ESBL) producing Gram-negative bacilli and methicillin-resistant *Staphylococcus aureus* (MRSA) further complicated the treatment resulting in higher mortality and morbidity rate^[6-8]. In addition, *Klebsiella pneumoniae* carbapenemases, an ESBL capable of hydrolysing a broad spectrum β -lactam antibiotics

including third-generation cephalosporins. These are plasmid-mediated enzymes, which can be transferred to other Gram-negative species, such as *Escherichia coli* and *Pseudomonas* sp.^[9,10]. Such plasmid-mediated enzymes further complicate antimicrobial therapy^[11]. Such isolates often exhibit resistance to other classes of drugs such as aminoglycosides (gentamicin, amikacin), cotrimoxazole, tetracycline and fluoroquinolones, which create a dreadful challenge with restricted therapeutic options especially in India^[12]. Some studies revealed that few bacteria (*E. coli* and *K. pneumoniae*) become highly resistant as they produce metallo- β -lactamases, which make the bacteria resistant to a broad range of β -lactams including the carbapenems^[13]. Presence of efflux pumps also explains high-level intrinsic resistance against a

This is an open access article distributed under the terms of the Creative Commons Attribution-NonCommercial-ShareAlike 3.0 License, which allows others to remix, tweak, and build upon the work non-commercially, as long as the author is credited and the new creations are licensed under the identical terms

*Address for correspondence
E-mail: mandalsm@gmail.com

Accepted 01 April 2018
Revised 17 August 2017
Received 28 December 2016
Indian J Pharm Sci 2018;80(3):547-551

broad variety of substrates. The overproduction of the efflux pump is usually accompanied by an increase in resistance to two or more structurally unrelated antibiotics and significantly contribute to the emergence and spread of multi-drug resistance pathogens^[14]. The expression of these efflux pumps are important for pathogens, which can be coupled to other mechanisms of resistance^[15,16].

Therefore, these hospital-based descriptive studies are essential for determining the pathogens prevalent in the respective hospital community and their resistance pattern to various antibiotics need to be documented. This will in turn help in formulating guidelines for management of BSI in local hospitals and the choice of antibiotic therapy in the community as well.

This study was done using the data of blood culture and antibiograms carried out at Sanjiban Hospital in Howrah district, West Bengal, India. All data were retrospectively collected and de-identified to ensure patient confidentiality. All the media, reagents and antibiotic-discs were procured from HiMedia, Mumbai, India.

A total number of 1500 blood culture samples were included in this study. This study assessed resistance in a total number of 30 antimicrobials. Two sets of blood specimens were collected from independent venepuncture sites of about 08-10 ml of blood for adult and for a child depending upon age but less than 1 % of total blood volume. The samples were sent to the laboratory as soon as possible after blood was aseptically introduced into aerobic media, brain heart infusion broth with sodium polyanethol sulfonate^[17]. The inoculated media were incubated for 7 d at 37° and blood cultures were considered negative if there was no growth after subcultures being made each day. For positive cultures, the identification of the isolates was performed through biochemical analysis and antimicrobial sensitivity test of the bacterial isolates were determined following Kirby-Bauer disc diffusion method as per CLSI guidelines^[14-18].

ESBL detection was carried out using routine susceptibility test following the method recommended by the NCCLS. Two discs, ceftazidime (30 µg) and cefotaxime (30 µg), were used. An inhibition zone of ≤22 mm for ceftazidime and ≤27 mm for cefotaxime indicated that the strain probably produced ESBL.

Confirmation test was also conducted following NCCLS recommendations, on Mueller-Hinton agar.

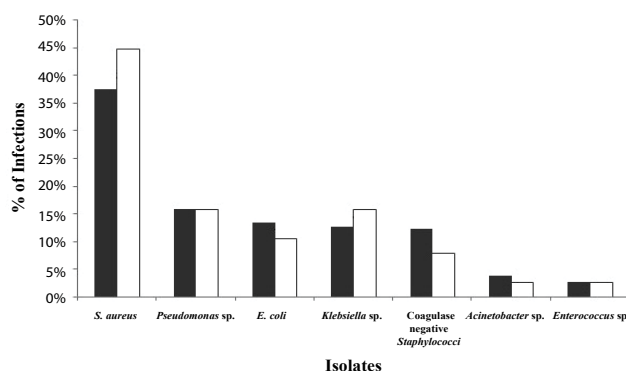


Fig. 1: Percent of different isolated organisms from blood culture

■ Total; □ neonates

Four discs, containing cefotaxime (30 µg), cefotaxime/clavulanic acid (30/10 µg), ceftazidime (30 µg) and ceftazidime/clavulanic acid (30/10 µg), were used. An inhibition zone of ≥5 mm increase in diameter for either antimicrobial tested in combination with clavulanic acid versus its zone when tested alone confirmed ESBL production^[19].

Blood culture samples were taken from patients who were admitted with clinical features suggestive of sepsis in our different wards including intensive care unit and neonatal care unit. A total of 1500 samples were critically screened and analysed. Among them, 190 samples were from neonatal patients with 15.2 % positive (38) for bacterial growth and out of the total blood cultures, 250 (16.67 %) were positive for bacterial growth, among them 123 (*E. coli*: 11, *Klebsiella sp.*: 27, *Acinetobacter sp.*: 7, *Pseudomonas sp.*: 21 and *S. aureus*: 57) isolates were ESBL positive and 2 were positive for *Candida sp.*

Amongst the 250 positive cultures, Gram-positive sp. (53.36 %) was predominant over Gram-negative sp. (46.64 %). By analysing the data, it was found that *S. aureus* was the most common bacterial pathogen causing BSI and accounted for 37.6 % of the total isolates. The other bacteria were, *Pseudomonas sp.* 16 %, *E. coli* 13.6 %, *Klebsiella sp.* 12.8 %, coagulase-negative *Staphylococci* 12.4 %, *Acinetobacter sp.* 4 % and *Enterococcus sp.* 2.8 % (fig. 1).

The antibiotic sensitivity patterns of the isolates were shown in Table 1. *S. aureus* was most sensitive to imipenem (98.76 %), meropenem (98.24 %), levofloxacin (97.56 %) but resistant to cefepime (18.75 %). The antibiotic susceptibility pattern of *Pseudomonas sp.* revealed sensitivity to levofloxacin (97.36 %), along with anti-*Pseudomonas* β-lactams, ceftazidime and cefoperazone-sulbactam. Most

TABLE 1: PERCENT ANTIMICROBIAL SUSCEPTIBILITY OF DIFFERENT ISOLATES

Bacteria	PI	AMC	CXM	CX	CTR	CPM	CTX	IPM	MRP	E	AZM	VA	CD	LZ
<i>S. aureus</i>	-	77.38	59.37	50	87.5	18.75	73.07	98.76	98.24	41.11	43.75	97.81	81.05	97.8
<i>Pseudomonas</i> sp.	42.1	37.14	61.11	-	64.87	24.13	72.73	83.33	97.14	-	-	-	-	-
<i>E. coli</i>	41.18	60.72	23.33	-	34.75	22.22	46.15	96	89.47	-	-	-	-	-
<i>Klebsiella</i> sp.	14.32	20.68	10.35		31.04	17.39	29.3	95.28	60					
<i>Staphylococcus</i> Coagulase (-ve)	-	85.18	66.67	29			71.42	100	100	42.4	45.83	100	90.33	100
<i>Acinetobacter</i> sp.	11.25	20	-	-	10	-	-	86.5	87.5	-	-	-	-	-
<i>Enterococci</i>	-	85.71	100	33.33	-	-	83.33	100	83.33	40	20	100	42.86	100
Bacteria	LE	GEN	RIF	COT	PIT	CZ	CAZ	CFS	OF	TGC	DO	CL	AK	
<i>S. aureus</i>	97.56	92.77	82.3	68.23	-	-	-	-	-	-	97.59	-	-	
<i>Pseudomonas</i> sp.	97.36	-	-	78.37	60.53	8	48.48	44.44	11.9	100	91.42	68	-	
<i>E. coli</i>	100	76.93	-	64.28	68.96	42.86	-	-	53.58	100	85.72	-	96.3	
<i>Klebsiella</i> sp.	65.25	32.14		38.46	22.24	8.1	-	-	52	100	76.92	70.25	25.2	
<i>Staphylococcus</i> , Coagulase (-ve)	96.42	88.89	87.88	69.23	-	-	-	-	-	-	96.29	-	-	
<i>Acinetobacter</i> sp.	77.75	25	-	28.5	22.5	-	-	-	44.5	100	100	100	66.75	
<i>Enterococci</i>	100	100	-	-	-	-	-	-	-	-	100	-	-	

Abbreviations are AMC: amoxicillin-clavulanate, PI: piperacillin, PIT: piperacillin-tazobactam, CZ: ceftazidime, CXM: cefuroxime, CTR: ceftriaxone, CTX: cefotaxime, CAZ: ceftazidime, CFS: cefoperazone-sulbactam, CPM: ceftazidime, CX: ceftazidime, IPM: imipenem, MRP: meropenem, OF: ofloxacin, LE: levofloxacin, GEN: gentamicin, AK: amikacin, CL: colistin, DO: doxycycline, TGC: tigecycline, E: erythromycin, AZM: azithromycin, VA: vancomycin, CD: clindamycin, LZ: linezolid, RIF: rifampicin, COT: cotrimoxazole

TABLE 2: PERCENT ANTIMICROBIAL RESISTANCE OF ESBL PRODUCERS FROM DIFFERENT ISOLATED BACTERIAL PATHOGENS

Antibacterial drugs	<i>E. coli</i> (n=11)	<i>Klebsiella</i> sp. (n=27)	<i>Acinetobacter</i> sp. (n=7)	<i>Pseudomonas</i> sp. (n=21)	<i>S. aureus</i> (n=57)
Ampicillin	81.81	100	100	90.47	82.45
Piperacillin	63.63	74.07	100	80.95	71.92
Ticarcillin	63.63	29.69	100	52.38	73.68
Amox/clavulanic acid	27.27	7.40	100	47.61	36.84
Ticarcillin/clavulanic acid	36.36	7.40	100	38.09	45.61
Piperacillin/tazobactem	27.27	40.74	100	42.85	35.08
Imipenam	0	7.40	14.28	14.28	12.28
Meropenam	0	7.40	14.28	9.52	7.01
Ceftazidime	72.72	77.77	71.28	52.38	43.85
Cephotaxime	72.72	70.37	71.28	47.61	36.84
Cefixime	36.36	25.95	71.28	61.90	57.89
Ceufuroxime	18.18	11.11	57.14	47.61	36.84
Cefpodoxime	18.18	22.22	57.14	47.61	43.85
Cotrimoxazole	45.45	40.74	85.71	38.09	49.12
Levofloxacin	0	0	28.51	19.04	5.26
Amikacin	9.09	11.11	71.48	23.80	15.78
Tobramycin	18.18	22.22	71.48	33.33	43.85

β -lactams were found ineffective against *Pseudomonas* sp. In case of *E. coli*, all the isolates were sensitive to levofloxacin, most were sensitive to amikacin and less than 40 % isolates were found sensitive to β -lactam antibiotics. Almost similar results were found with the *Pseudomonas* isolates. The sensitivity profile of *Klebsiella* isolates were quite unlike comparing with *Pseudomonas* and *E. coli* isolates. Most of the *Klebsiella* isolates were found sensitive to imipenem, meropenem and polymyxins. Nearly 60 % of the

isolates were found sensitive to levofloxacin and 25 % strains were sensitive to amikacin. Most β -lactams were found ineffective against *Klebsiella* isolates. Like *S. aureus*, coagulase-negative *Staphylococci* were found susceptible to levofloxacin along with meropenem, imipenem, vancomycin, linezolid, clindamycin amoxicillin-clavulanic acid. The antibiotic sensitivity pattern of *Acinetobacter* isolates indicated sensitivity to tigecycline, doxycycline, polymyxin B, colistin (100 %), and a few of them were also sensitive

to meropenem, imipenem, and levofloxacin. In case of *Enterococci*, it was found that all the isolates were sensitive *in vitro* to imipenem, vancomycin, linezolid, levofloxacin, gentamycin. Very few number of isolates were found sensitive to erythromycin (40 %), azithromycin (20 %).

The antimicrobial resistance patterns of ESBL-producing isolates were presented in Table 2. Among the total bacterial isolates (250), ESBL producing strains (123) were 49.20 % where individual strains were *E. coli* (39.28 %), *Klebsiella* sp. (77.14 %), *Acinetobacter* sp. (100 %), *Pseudomonas* sp. (52.5 %) and *S. aureus* (56 %), a significant number of percentage. All the ESBL positive strains are resistant to cephalosporins, ceftazidime and cefotaxime and sensitive to carbapenems, meropenem or imipenem. Few of them showed resistant to both cephalosporins and carbapenems, which could be the cause of cross-resistance or by the production of other β -lactamases.

The Surviving Sepsis Guidelines state that initial empiric antibiotic therapy is one or more drugs, which are active against all likely pathogens (bacterial and/or fungal or viral) and that penetrate in adequate concentrations into tissues presumed to be the source of sepsis (grade 1B). To ensure both the above, locally prevalent pathogens and their antibiograms should be made available to practice evidence-based medicine. The present study revealed a change in the sensitivity pattern of the common pathogens to commonly used antibiotics in these hospitals. In the last decade, ESBLs were of great medical importance as these hydrolyse β -lactam antibiotics. In recent years, the spread and incidence of ESBLs have increased due to the overuse of expanded-spectrum cephalosporins in the hospital setting. It is a major therapeutic challenge to treat infections by strains of Gram-negative bacteria that express ESBLs.

Third generation quinolone such as levofloxacin was the cheapest amongst the most potent antimicrobial agents effective against both Gram-negative and positive organisms in this unit and the least sensitive antibiotics were ampicillin and the first generation cephalosporins. The third generation cephalosporins are commonly used as second line antibiotics in many centers (and even as first line therapy in some centers) have been observed to become increasingly ineffective as shown in the present study, which is comparable to published reports^[12,20]. However, *Klebsiella* sp. showed different results, 25 % sensitivity to amikacin

and 32.14 % sensitivity to gentamicin that supported some earlier reports. Increasing resistance pattern of aminoglycosides to *Klebsiella* sp. has also been reported by other observers^[21].

Therefore, the quinolones, especially levofloxacin was found to be the most potent empirical antibiotic for sepsis patients. The major limiting factors of the third generation cephalosporins were high-price and toxicity. Amongst aminoglycosides, even in the face of gentamicin resistance, amikacin might be recommended because of its good susceptibility patterns. The carbapenems especially meropenem was found to be collectively effective in almost all types of Gram-positive and negative bacterial isolates in this investigation. However, the high cost of therapy with meropenem advocate the use of levofloxacin as an initial empirical therapy in and around this region. Furthermore, steps needed to be taken to prevent or control the emergence of resistance strains by using antibiotics rationally. Laws therefore should be enforced to discourage the indiscriminate use of antibiotics seen commonly in India as well as discourage inadequate doses through proper training of personnel in order to prevent emergence of resistant strains. In the present study resistance to three or more antibiotics was common in both ESBL producers and non-ESBL producers. About 80 % strains were resistant to penicillins and cephalosporins. Septicaemia is a leading cause of mortality when caused by ESBL-positive strains. The enormous use of cephalosporins might become one of the major factor for increasing rate of ESBL producing microorganisms^[22].

In conclusion, there is an overall shift in the antibiotic susceptibility to the commonly isolated bacterial pathogens. Antibiotic resistance becoming a major public health concern and hazard all over the world, so it is imperative to take more caution and evidence-based judgment, while starting antibiotics therapy. From this study, it could be concluded that levofloxacin is the drug that should be advocated when starting antibiotic empirically for BSI followed by amikacin as these were found to give good coverage against both Gram-positive and Gram-negative organisms. Increased awareness of the ESBL problem among hospital settings is crucial to control them as soon as possible.

Acknowledgements:

Authors acknowledge the Director of Sanjiban Hospital for granting permission and support. Special thanks are

to Mr. Amit Chakraborty and Miss Anannya Ghosh for their help throughout the work.

Conflict of interest:

Authors have no any conflict of interest and none financial involvement in this work.

Financial support and sponsorship:

Nil.

REFERENCES

1. Del Rio A, Cervera C, Moreno A, Moreillon P, Miro JM. Patients at risk of complications of *Staphylococcus aureus* bloodstream infection. *Clin Infect Dis* 2009;48:S246-S53.
2. Dellinger RP, Levy MM, Rhodes A, Annane D, Gerlach H, Opal SM, *et al.* Surviving Sepsis Campaign: international guidelines for management of severe sepsis and septic shock. *Crit Care Med* 2008;34:17-60.
3. West BA, Peterside O. Sensitivity pattern among bacterial isolates in neonatal septicaemia in Port Harcourt. *Ann Clin Microbiol Antimicrob* 2012;11:7-13.
4. Marin H, Kollef MH, Golan Y, Micek ST, Shorr AF, Restrepo MI. Appraising contemporary strategies to combat multidrug resistant gram-negative bacterial infections—proceedings and data from the Gram-Negative Resistance Summit. *Clin Infect Dis* 2011;53:S33-S5.
5. Suárez CJ, Lolans K, Villegas MV, Quinn JP. Mechanisms of resistance to beta lactams in some common gram negative bacteria causing nosocomial infection. *Expert Rev Anti Infect Ther* 2005;3:915-22.
6. Jain A, Roy I, Gupta MK, Kumar M, Agarwal SK. Prevalence of extended-spectrum beta-lactamase-producing Gram-negative bacteria in septicaemic neonates in a tertiary care hospital. *J Med Microbiol* 2003;52:421-25.
7. Kim YK, Pai H, Lee HJ, Park SE, Choi EH, Kim J, *et al.* Bloodstream infections by extended-spectrum beta-lactamase-producing *Escherichia coli* and *Klebsiella pneumoniae* in children: epidemiology and clinical outcome. *Antimicrob Agents Chemother* 2002;46:1481-91.
8. Kapil A. Risk factors associated with MRSA infection in Children. *Indian Pediatr* 2015;52:22-4.
9. Navon-Venezia S, Chmelnitsky I, Leavitt A, Schwaber MJ, Schwartz D, Carmeli YA. Plasmid-mediated imipenem-hydrolyzing enzyme KPC-2 among multiple carbapenem-resistant *Escherichia coli* clones in Israel. *Antimicrob Agents Chemother* 2006;50:3098-101.
10. Villegas MV, Lolans K, Correa A, Kattan JN, Lopez JA, Quinn JP. Nosocomial Resistance Study Group: First identification of *Pseudomonas aeruginosa* isolates producing a KPC-type carbapenem-hydrolyzing beta-lactamase. *Antimicrob Agents Chemother* 2007;51:1553-55.
11. Pitout JD. The latest threat in the war on antimicrobial resistance. *Lancet Infect Dis* 2010;9:578-9.
12. Chandel DS, Johnson JA, Chaudhry R, Sharma N, Shinkre N, Parida S, *et al.* Extended-spectrum beta-lactamase producing Gram-negative bacteria causing neonatal sepsis in India in rural and urban settings. *J Med Microbiol* 2011;60:500-07.
13. Kumarasamy KK, Toleman MA, Walsh TR, Bagaria J, Butt F, Balakrishnan R, *et al.* Emergence of a new antibiotic resistance mechanism in India, Pakistan, and the UK: a molecular, biological, and epidemiological study. *Lancet Infect Dis* 2010;10:597-602.
14. Sarkar SK, Bhattacharyya A, Mandal SM. YnfA, a SMR family efflux pump is abundant in *Escherichia coli* isolates from urinary infection. *Indian J Med Microbiol* 2015;33:139-42.
15. Wen-Sen L, Wei-Yu C, Tsong-Yih O, Fu-Lun C, Yung-Wei L. Malakoplakia in a patient with complicated urinary tract infection caused by extended-spectrum β -lactamase-producing *Escherichia coli*. *J Microbiol Immunol Infect* 2015;48:345-6.
16. Doumith M, Ellington MJ, Livermore DM, Woodford N. Molecular mechanisms disrupting porin expression in carbapenem-resistant *Klebsiella* and *Enterobacter* sp. clinical isolates from the UK. *J Antimicrob Chemother* 2009;63:659-7.
17. Wasilauskas BL, Floyd J, Richard Roberts T. Use of Sodium Polyanethol Sulfonate in the preparation of 5% Sheep Blood Agar Plates. *Appl Microbiol* 1974;28:91-4.
18. Performance Standards for antimicrobial susceptibility testing, Seventeenth informational supplement. *Clinical and Laboratory Standards Institute* 2007;M-100 S-16(26):3.
19. Deepthi R, Deepthi N. Extended-spectrum β -lactamases in Gram Negative Bacteria. *J Glob Infect Dis* 2010;2:263-74.
20. Gea-Banacloche JC, Opal SM, Jorgensen J, Carcillo JA, Sepkowitz KA, Cordonnier C. Sepsis associated with immunosuppressive medications: An evidence-based review. *Crit Care Med* 2004;32:578-90.
21. Livermore DM, Yuan M. Antibiotic resistance and production of extended-spectrum Beta lactamases amongst *Klebsiella* spp. from intensive care units in Europe. *J Antimicrob Chemother* 1996;38:409-24.
22. Paterson DL, Bonomo RA. Extended-spectrum beta-lactamases: a clinical update. *Clin Microbiol Rev* 2005;18:657-86.