

Characterization and Antibiotic Susceptibility Pattern of Nonfermenting Gram Negative Bacilli from Blood Stream Infections in Children in a Tertiary Care Hospital, Punjab

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Abstract

Purpose: Bacteraemia due to nonfermentative gram negative bacilli appears to be increasing in frequency particularly in hospitalized patients with severe underlying illness.

Objectives: Therefore purpose of this study was to characterize various nonfermenters and to know their antibiotic susceptibility pattern causing blood stream infections in children.

Materials and Methods: This retrospective study was conducted in the Microbiology department of a tertiary care hospital, in Punjab. A total of 2267 paediatrics blood cultures were received from suspected cases of septicaemia in the laboratory. All the blood samples were processed in the Microbiology Laboratory by RAPID BD BACTEC PLUS system and organisms were identified as per standard microbiological techniques. Antimicrobial susceptibility testing was done using Kirby Bauer disc diffusion method as per CLSI guidelines.

Results: Out of the 2267 samples from paediatrics patient's 187 (8.24%) blood cultures were positive. Gram negative bacilli were the predominant organism 96 (51.3%). Of these 32 (33.3%) were nonfermenters which were further characterized. Among the 32 NFGNB, *Pseudomonas* spp was the commonest isolate among the nonfermenters accounting for 71.8 % (23/32), followed by *Acinetobacter* spp 15.6% (5/32) and *Shewanella putrefaciens* 12.50% (4/32). These were 100% sensitive to Imipenam, Meropenam, Polymixin B and Colistin. It showed (87.50%) sensitivity to Chloramphenicol followed by Ciprofloxacin (84.38%), Piperacillin-Tazobactam (84.38%) and Amikacin (81.25%). The isolates showed least sensitivity to Ampicillin (12.50%).

Conclusion: *P.aeruginosa* and *A.baumannii* were the common NFGNB isolated in our study. Therefore identification of NFGNB and monitoring their susceptibility pattern are important for proper management of these infections caused by them.

Key words: Bacteraemia, Nonfermentative Gram Negative Bacilli

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I. Introduction

Nonfermenting gram negative bacilli (NFGNB) are a taxonomically diverse group of aerobic, nonsporing bacilli that either do not utilize glucose as a source of energy or utilize it oxidatively^[1]. They occur as saprophytes in the environment and some are also found as commensals in the human gut^[2,3]. Nonfermentative gram negative bacilli (NFGNB), although frequently considered to be commensals or contaminants, but the pathogenic potential of these organisms has been established beyond doubt because of their frequent isolation from clinical specimens and their association with the disease^[4]. Gram negative nonfermenting aerobic bacteria are being increasingly implicated in human disease. The complex physicochemical properties of these organisms necessitate a battery of tests for their precise identification.^[5] In addition there is still much confusion regarding the taxonomic status of many of these nonfermenters. Identification of these nonfermenters has often been neglected^[6]. In the recent years due to liberal and empirical use of antibiotics, NFGNB have emerged as important health care associated pathogens. NFGNB are innately resistant to many antibiotics and are known to produce extended spectrum β -Lactamases and metallo β -Lactamases^[3,5,7,8]

II. Objectives

Therefore purpose of this study was to characterize various nonfermenters and to know their antibiotic susceptibility pattern causing blood stream infections in children.

III. Material & Methods

This retrospective study was conducted in the Microbiology Department of a tertiary care hospital in Punjab. A retrospective analysis was conducted from September 2012 to December 2013. During this period 2267 paediatrics blood cultures were received from suspected cases of septicaemia in the laboratory. All the blood samples were processed in the Microbiology Laboratory by RAPID BD BACTEC PLUS system and organisms were identified as per standard microbiological techniques^[9]. Every positive vial flagged by the Bactec 9120 (Becton Dickinson, USA) was removed for sub culturing in the safety cabinet. The lid of the positive blood culture vial was disinfected with an alcohol swab and 1 ml of blood was aseptically removed from the vial and plated on blood agar, and MacConkey's agar. The Blood agar and MacConkey agar, incubated aerobically in incubator at 37°C for 24 to 48 hours and then observed for growth of bacteria. All the plates showing growth were processed according to standard Microbiological techniques including gram staining^[10]. All the gram negative bacilli/coccobacilli, oxidase positive or negative were inoculated on to triple sugar iron (TSI) medium. Organisms that failed to show any change, both in slant and butt of medium were considered nonfermenters. The isolates were further identified up to the species level based on motility, pigment production, enzyme production (catalase, urease, nitrate reductase)^[11] and various biochemical tests included oxidative-fermentative test, growth at 42°C, nitrate or nitrite reduction, gelatin liquefaction, arginine dihydrolase, lysine decarboxylase, citrate utilization, indole test, production of hydrogen sulphide and esculin hydrolysis. Antimicrobial susceptibility testing of all the isolates using Kirby-Bauer disc diffusion method^[12] was carried out with disc containing, ampicillin(10µg), cotrimoxazole(25µg), amikacin(10µg), cefotaxime(30µg), ceftazidime(30µg), ciprofloxacin(5µg), piperacillin-tazobactam (100/10 µg), and imipenem (30 µg) and the results were interpreted as per clinical and laboratory standards institute (CLSI) recommendation^[13].

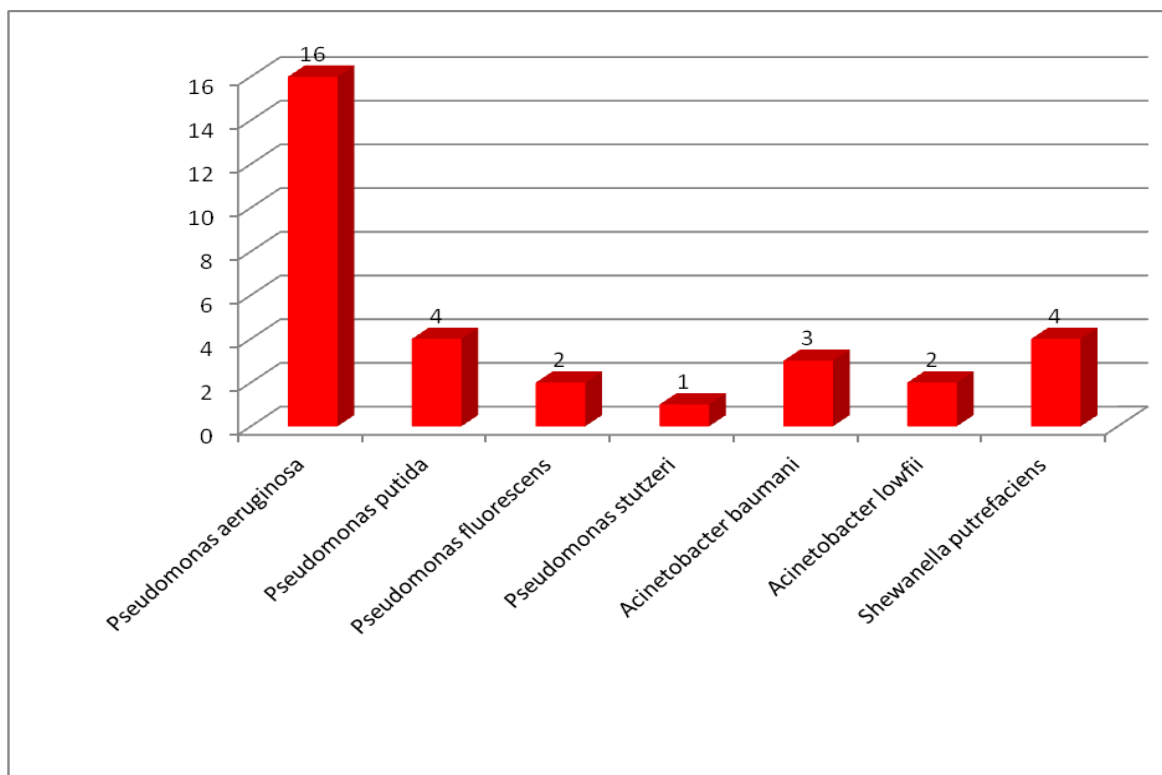
IV. Results:

Out of the 2267 samples from paediatrics patient's 187 (8.24%) blood cultures were positive. Gram negative bacilli were the predominant organism 96 (51.3%). Of these 32 (33.3%) were nonfermenters which were further characterized. Among the 32 NFGNB, *Pseudomonas aeruginosa* was the commonest isolate among the nonfermenters accounting for 50.0% (16/32), followed by *Pseudomonas putida* 12.50% (4/32), *Shewanella putrefaciens* 12.50% (4/32), *Acinetobacter baumannii* 9.38% (3/32), *Acinetobacter Iwofii* 6.25% (2/32), *Pseudomonas fluorescens* 6.25% (2/32) and *Pseudomonas stutzeri* 3.12% (1/32). These were 100% sensitive to Imipenam, Meropenam, Polymixin B and Colistin. It showed (87.50%) sensitivity to Chloramphenicol followed by Ciprofloxacin (84.38%), Piperacillin-Tazobactam (84.38%) and Amikacin (81.25%). The isolates showed least sensitivity to Ampicillin (12.50%).

Table 1: Nonfermentative gram negative bacilli in blood culture isolates

Si.no	Isolate	Number
1.	<i>Pseudomonas aeruginosa</i>	16(50.0)
2.	<i>Pseudomonas putida</i>	4(12.50)
3.	<i>Pseudomonas fluorescens</i>	2(6.25)
4.	<i>Pseudomonas stutzeri</i>	1(3.12)
5.	<i>Acinetobacter baumani</i>	3(9.38)
6.	<i>Acinetobacter lowfii</i>	2(6.25)
7.	<i>Shewanella putrefaciens</i>	4(12.50)
	TOTAL	32(100)

Percentage in parentheses represent out of total isolates (n=32)



Graph 1: Nonfermentative gram negative bacilli in blood culture isolates

Table 2 : Antibiotic sensitivity pattern of nonfermenters

Antibiotic	Sensitivity
Ampiciline (amp)	4 (12.5)
Cotrimoxazole (cot)	19 (59.3)
Amikacin (amk)	26 (81.25)
Cefotaxime (ctx)	23 (71.8)
Cefperazone (cpz)	20 (62.5)
Pipracillin /tazo (pit)	27 (84.3)
Ciprofloxin (cip)	27 (84.3)
Imipenem (imp)	32 (100)
Meropenem (mrp)	30 (93.7)
Polymixin b(p)	32 (100)
Colistin (c)	32 (100)

Percentage in parentheses represent out of total isolates (n=32)

Table 3: Antibiotic sensitivity pattern of pseudomonas spp, acinetobacter spp and schwanella spp

Antibiotic	Pseudomonas spp n (23)	Acinetobacter spp n(5)	Schwanella spp n(4)
Ampiciline (amp)	2(8.69)	1(20)	1(25)
Cotrimoxazole (cot)	14(60.8)	1(20)	4(100)
Amikacin (amk)	23(100)	1(20)	2(50)
Cefotaxime (ctx)	20(86.9)	1(20)	2(50)
Cefperazone (cpz)	17(73.9)	1(20)	2(50)
Pipracillin /tazo (pit)	22(95.6)	2(40)	3(75)
Ciprofloxin (cip)	21(91.3)	2(40)	4(100)
Imipenem (imp)	23(100)	5(100)	4(100)
Meropenem (mrp)	23(100)	3(60)	4(100)
Polymixin b(p)	23(100)	5(100)	4(100)
Colistin (c)	23(100)	5(100)	4(100)

Percentage in parentheses represent out of isolates

V. Discussion:

In the present study, 32 (33.3%) strains of nonfermenters were isolated which is in accordance with the results of study done by Chang & Huang^[14] isolated 31.62% of nonfermenters. Various workers have reported variable results in their studies. Rao & Shivananda^[15] reported higher positivity rate of 66.88% while Seetha et

al^[16] isolated 45% of nonfermenters. These variations might be because of the hospital infection control practices of that institute. *Pseudomonas aeruginosa* is an opportunistic pathogen which causes severe infections in patients with immune deficiency and newborns. The present study identified 11.50% (23/200) *Pseudomonas* spp and 2.50 % (5/200) *Acinetobacter* spp. Similar results were seen in a retrospective study conducted by (Asghar AH).^[17] Nonfermenters are resistant to a large number of commonly used antibiotics. In our study they are 100% sensitive to Imipenam, Meropenam, Polymixin B and Colistin. Most of the strains showed sensitivity to Chloramphenicol (87%) followed by Ciprofloxacin (84.3%), Piperacillin-Tazobactam (84.3%) and Amikacin (81.2%). The isolates showed least sensitivity to Ampicillin (12.5%). This contrasts with the antibiotic sensitivity pattern of isolates obtained from other studies^[18-20], where 70-80% of the isolates were sensitive to Ciprofloxacin & Amikacin.

In the present study, *Pseudomonas* spp showed (90%) resistance to Ampicillin and (56%) to Aztreonam. They are (60%) sensitivity to Cotrimoxazole, Cefotaxime (86%), Ceferazone(73%), , Piperacillin/Tazobactam (95%), Ciprofloxacin (91%). (100%) isolates were sensitive to Amikacin, Gentamicin, Imipenam, Meropenam, Polymixin B and Colistin. (Karlowsky JA et al)^[21] have shown that *P. aeruginosa* was highly susceptible to Amikacin (92.3%) and Piperacillin-Tazobactam (91%) similar to the present study. This contrasts with the antibiotic sensitivity pattern of isolates from other studies^[18-20] where (70-80%) of the isolates were sensitive to Ciprofloxacin & Amikacin.

In our study the frequency of *Acinetobacter* species was 2.5% (5/200) among the culture positive blood samples. The total number of *Acinetobacter* species isolated during the study period was 5, out of which 3 were *A.baumannii* and 2 were *A.Iwofii*. *Acinetobacter* species were less frequent in blood samples as compared to other bacteria causing blood stream infections. A study conducted in paediatric intensive care unit (PICU) of a tertiary care teaching hospital reported (8.6%) frequency of *Acinetobacter* species isolated from blood samples.^[22] Another study done in National Cheng Kung University, Tainan reported the similar result of 9.9% for *Acinetobacter* species.^[23] The frequency of *Acinetobacter* in our study was lower than the other studies. This could be due to clean hospital environment and hygienic conditions and good infection control practices. In the present study, *Acinetobacter* spp showed (80%) resistance to Ampicillin, Cotrimoxazole, Amikacin, Cefotaxime and Ceferazone of the isolates were resistant to Piperacillin/Tazobactam and Ciprofloxacin. It showed (60%) sensitivity to Meropenam. 100% isolates were sensitive to Imipenam, Polymixin B and Colistin. This finding is in conformity with the observation of other workers in 2009 a study from Rohtak showed that the resistance of *Acinetobacter* to Meropenem had increased to 25.6%(Goel et al).^[24] In the same study the resistance to Amikacin was (87.2%). Our study showed (100%) sensitivity to Polymyxin. However there have been reports of Polymyxin resistant *Acinetobacter* from Greece, Slovakia and other parts of Europe (Gales et al)^[25], (Souli et al)^[26]. *Shewanella* spp showed 75.0% resistance to Ampicillin. It showed 50.0% sensitivity to Amikacin, Cefotaxime and Ceferazone and 75% sensitive to Piperacillin/Tazobactam.. All the isolates were 100% sensitive to Cotrimoxazole, Ciprofloxacin, Imipenam, Meropenam, Polymixin B and Colistin. This contrasts with the antibiotic sensitivity pattern of isolates from other studies.

VI. Conclusions:

P.aeruginosa and *A.baumannii* were the commonest NFGNB isolated in our study. The different species of NFGNB have shown a varied sensitivity pattern in our study. Therefore identification of NFGNB and monitoring their susceptibility pattern are important for proper management of these infections caused by them. Our study highlights that it is essential to establish the clinical relevance of the NFGNB isolated, before they are considered as pathogens. This would avoid unnecessary usage of antibiotics and emergence of drug resistant strains.

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