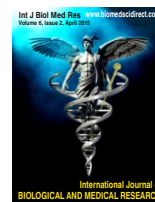




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Original article

Study on Non fermenting gram negative bacilli from various clinical samples in a tertiary care hospital.

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ABSTRACT

Non fermenting gram negative bacilli are a group of heterogenous, aerobic, non sporing bacteria. they are saprophyte in nature and are also found as commensals in man and animals. Hence this study aims at isolation, identification, and antibiotic susceptibility for non fermenting gram negative bacilli from clinical samples and to find out the clinical significance among the inpatient in meenakshi medical college and hospital kanchipuram. A total of 103 non fermenters isolated from clinical sample such as pus, sputum, urine, blood, and ear swab were included in the study. All these were identified by a battery of test as per standard laboratory technique. Aintimicrobial sensitivity testing was performed for all isolates by kirby bauer disc diffusion method. In these non fermenters, *Pseudomonas* were 83.5% followed by *Acinetobacter* 15.5% species was the second commonest non-fermenter followed by *shewanella sp.*,(0.9%). The NF GNBs are known cause of variety of nosocomial infection. Resistance pattern among the nosocomial bacterial pathogen may vary widely from place to place even within the same country over time. Thses nonfermenters are isolated from clinical specimens frequently and are resistant to most of the routinely used antibiotics showing that organisms need to be taken seriously and identified and not just regarded as contaminants.

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1. Introduction

Non Fermenting gram negative bacteria constitutes about one-fifth of all GNB's. They are ubiquitous in nature, widely distributed although frequently considered as the commensal or contaminants but are frequently isolated from clinical specimens and are associated with disease[1]. Recently there has been a tremendous interest in these organisms as they are being isolated from clinical specimens with increasing frequency. Infections caused by other species are relatively infrequent[2]. In recent years due to the indiscriminate use of antimicrobials, NFGNB have emerged as important health care associated pathogens. They have been incriminated in infections such as bacteremia, meningitis, pneumonia, urinary tract infections, surgical site infections, wound infections, osteomyelitis, etc[3].

They are common in hospital settings and may be found on the surface of humidifiers, ventilator dialysis machines and other equipments as well as from the skin of hospital person. Recently they have been reported as the responsible for the serious infections such as septicemia and pneumonia. NF-GNB's are oxidase positive known for the ability to cause infections in debilitated and immunocompromised individuals[4]. NFGNB are known to account for 15% of all bacterial isolates from clinical microbiological laboratory. The organisms belonging to the genera

Pseudomonas, *Acinetobacter*, *Burkholdaria*, *Aeromonas*, *Achromobacter* and *Stenotrophomonas* are normally present in air and on moist or dry surfaces[5] & therefore they have greater chances to infect patients in nosocomial setting.

They rarely cause serious infections in healthy persons and are infrequently identified as normal microbial floras in healthy individuals, but they are of greatest concern in hospitalized patients, particularly those in ICUs, where these opportunistic pathogens are capable of causing severe infections in critically ill and immunocompromised patients such as bacteremia, pneumonia, urinary tract infection, meningitis, endocarditis, burns, wound infection, eye infection, surgical site infection and osteomyelitis[1,5]. *Pseudomonas spp.*,having resistance to Carbapenem which is currently the most effective treatment option and the number of reported incidence are gradually increasing. Multidrug-resistant (MDR) strains of *P. aeruginosa* are often isolated among patients suffering from nosocomial infections, particularly those in the intensive care unit (ICU)[4,5].

Thus,infections caused by *P. aeruginosa* are particularly problematic because the organism is inherently resistant to many drug classes and is able to acquire resistance to all effective antimicrobial drugs. As an opportunistic infectious pathogen, *P. aeruginosa* can often lead to life-threatening diseases.The emergence in causing infection is of great concern to clinicians, spreading of resistance to commonly used antibiotics in both human populations has posed adverse impact on morbidity and mortality due to diseases caused by resistant bacteria[6]. The isolation rate of NF-GNB's were increasing in our lab, hence this

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study was undertaken to identify, speciate and to study the sensitivity pattern of NF-GNB's and also to know the clinical significance of this infected organisms.

2. Materials and methods:

The present study was undertaken at the department of microbiology, Meenakshi Medical College, Kanchipuram during the periods from Feb2013- Nov 2013 and a total of 103 Non-fermenters were isolated from clinical samples such as pus, sputum, urine and blood were included in the study. All collected specimens were processed by standard techniques. Organisms that failed to acidify the butts of triple sugar iron media were considered nonfermenters and subjected to a battery of tests like morphology, motility, oxidase, catalase, indole, urease, nitrate, citrate tests and oxidation-fermentation reactions of glucose, lactose, xylose, maltose, mannitol (Hugh-Leifson method). Antimicrobial susceptibility testing of all the isolates using Kirby-Bauer disc diffusion method and the results were interpreted as per clinical and laboratory standards institute (CLSI) recommendation.

3. Results:

Out of 103 NF isolated from pus, blood, sputum, urine were processed for a period of Feb2013-Nov2013. In these *Pseudomonas* spp., were maximum isolated followed by *Acinetobacter* spp., institute. All the cases with nonfermentative bacteraemia were immunosuppressed due to presence of one or the other underlying condition or disease.

Table- 1: Distribution of NFGNB from various clinical specimens

Isolate	Total no:	Percentage(%)
<i>P.aeruginosa</i>	86	83.5%
<i>A.baumannii</i>	13	12.7%
<i>A.lwoffii</i>	03	2.9%
<i>Sh.putrefaciens</i>	01	0.9%

Among the non-fermenters, *Pseudomonas* spp., were the maximum isolate followed by *Acinetobacter* spp.,

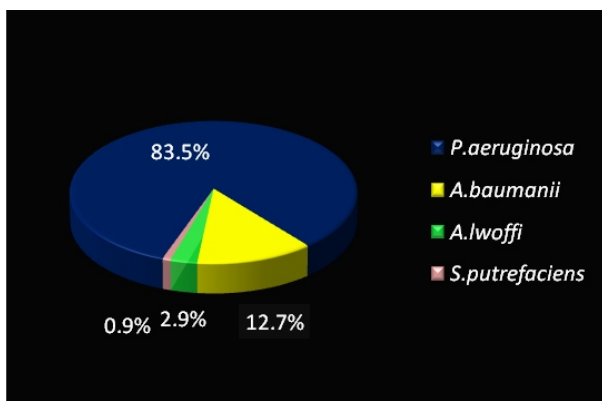


Table – 2: Distribution of different NFGNB various clinical specimens

Specimen	No. of cases	Percentage(%)
Pus	50	48.6%
Sputum	34	33%
urine	13	12.6%
Blood	06	5.8%

Among the non-fermenters, *Pseudomonas* spp., were the maximum isolate followed by *Acinetobacter* spp.,

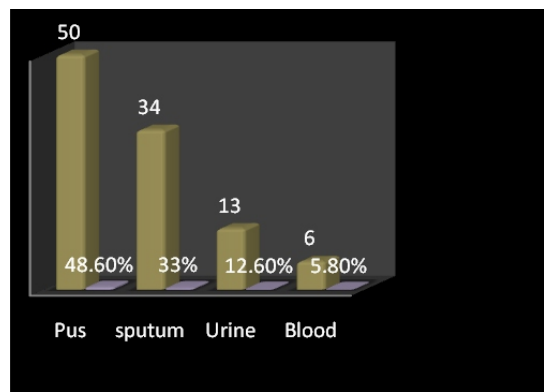


Table-3. Antibiotic resistance pattern of NFGNB against penicillin group

Antibiotics	Number of <i>Pseudomonas</i> spp.,	Percentage(%)	Number of <i>Acinetobacter</i> spp.,	Percentage(%)
Penicillin	61	75%	7	43.7%
Amoxyclav	40	46.5%	13	75%
Netilmicin	35	40.6%	8	50%
Piperacillin+tazo	28	32.5%	4	25%

Maximum resistance is seen to penicillin by *Pseudomonas* followed by Amoxyclav. In *Acinetobacter* spp., more resistance to Amoxyclav followed by Netilmicin.

Table-4 Antibiotic resistance pattern of NFGNB to Cephalosporin group.

Antibiotics	Number of <i>Pseudomonas</i> spp.,	Percentage(%)	Number of <i>Acinetobacter</i> spp.,	Percentage(%)
Cefuroxime	48	55.6%	6	37.5%
Cefotaxime	37	43%	5	33.7%
Ceftriaxone	34	39.5%	7	43.7%
Cefepime	31	36%	7	43.7%
Cefaperazone	32	37.2%	4	25%
Ceftazidime	36	41.8%	6	37.2%

Maximum resistance is seen to Cefuroxime by *Pseudomonas* followed by Cefotaxime. In *Acinetobacter* more resistance to Ceftriaxone followed by Cefepime.

Table-5. Antibiotic resistance pattern of NFGNB to aminoglycoside

Antibiotics	Number of <i>Pseudomonas</i> spp.,	Percentage(%)	Number of <i>Acinetobacter</i> spp.,	Percentage(%)
Amikacin	42	48.8%	4	25%
Gentamicin	38	44.5%	5	31.2%

Maximum resistance is seen to Amikacin by *Pseudomonas* & *Acinetobacter*.

Table -6. Antibiotic resistance pattern of NFGNB to quinolones

Antibiotics	Number of <i>Pseudomonas</i> spp.,	Percentage(%)	Number of <i>Acinetobacter</i> spp.,	Percentage(%)
Ciprofloxacin	42	48.2%	5	31.2%
Ofloxacin	48	55.2%	03	18.7%

Maximum resistance is seen to Ciprofloxacin by *Pseudomonas* & *Acinetobacter*.

Table -7. Antibiotic resistance pattern of NFGNB to carbapenems groups

Antibiotics	Number of <i>Pseudomonas</i> spp.,	Percentage%	Number of <i>Acinetobacter</i> spp.,	Percentage(%)
Imipenem	10	11.6%	04	25%
Meropenem	09	10.4%	06	37.5%

Maximum resistance is seen to Imipenem by *Pseudomonas*. In *Acinetobacter* shows high resistance to Meropenem.

PHOTOGRAPHS

Figure: 1 Gram's stain showing gram negative bacilli of *Paeruginosa*.

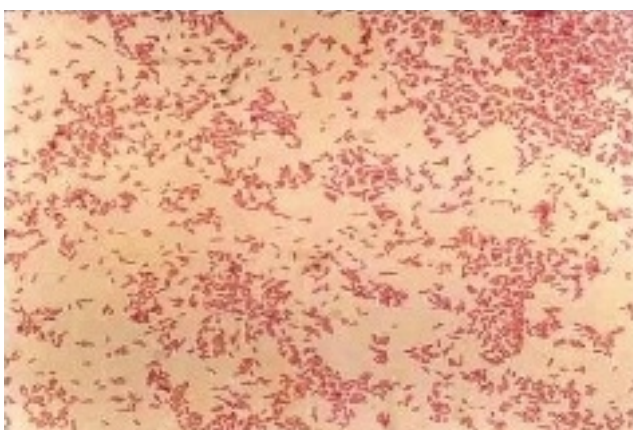


Figure: 2 Gram's stain showing gram negative Cocobacilli of *A.baumanii*

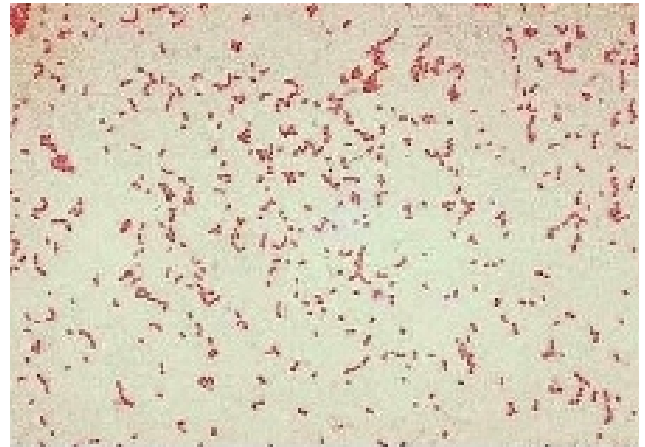


Figure: 3 Pigmented colonics of *p.aeruginosa* on Nutrient agar



Figure: 4. *A. baumannii* on Blood agar



Figure. 5

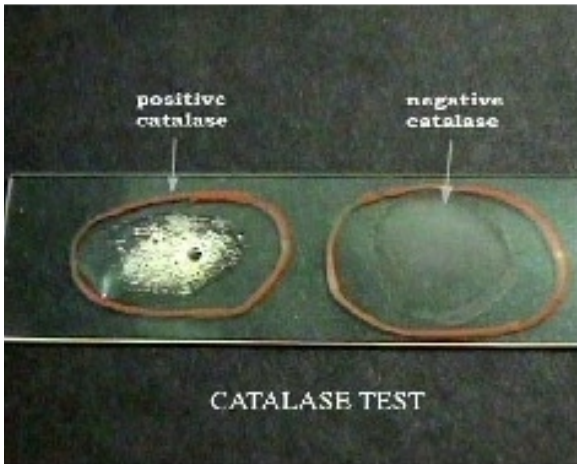


Figure. 6



Pseudomonas aeruginosa shows Catalase and Oxidase test positive

Figure. 7



Pseudomonas aeruginosa biochemical reaction from L to R: Indole, MR, VP} Negative, Citrate Positive, Urease Negative, TSI - M/NC, Mannitol - NF/M, Nitrate test Positive.

Figure. 8



Biochemical reactions of *shewanella putrefaciens* from L to R: Indole, MR, VP Negative, Citrate Positive, TSI - K/K [H₂S Production]. Mannitol - NF/NM.

Figure. 9



A.baumannii biochemical reactions from L to R Indole, MR, VP Negative, Citrate positive, Urease Positive, TSI-K/K, Mannitol - NF/NM, Nitrate positive.

Discussion:

NF GNBS were considered as a contaminant in the past but now emerged as an important health care pathogen. *Pseudomonas aeruginosa* and *Acinetobacter* sp., are known to be nosocomial pathogen. In the present study out of 103 non fermenters, *Pseudomonas* were 83.5% followed by *Acinetobacter* 15.5% species was the second commonest non-fermenter followed by *shewanella* sp.,(0.9%). This is similar to studies done by(Malini and Vijaya et al 2009)[6,26]. *Pseudomonas* isolation was predominant in the study by done (Parimal et al 2013). In our study highest number of NF GNBS were from pus samples followed by sputum. This is similar to study done by Malini et al and Parimal et al (2013)[6]. In our study NF GNBS were isolated (18.4%) from various clinical samples.

The NF GNBS are known cause of variety of nosocomial infection (Larson EL1981)[8],(Bergogne.E 2001). Resistance pattern among the nosocomial bacterial pathogen may vary widely from place to place even within the same country over time (Prashanth K, Badrinath S. 2004)[7]. In our study out of 16

isolates of *Acinetobacter* species, revealed majority isolates as *A.baumannii*(81.25%) were as 3 isolates were found to be *A.lwoffii*(18.75%). Majority of the isolates were from pus specimen followed by urine. This is similar to study done by Parandekar P et al (2012).

Resistance to 3rd generation Cephalosporin, Cephotaxime showed (43%) and Ceftazidime showed (41%) in *Pseudomonas*. But it is higher than Angadi et al (2012) who reported (24.6%) to Cefotaxime and (25.6%) to Ceftazidime in *Pseudomonas*. Similarly Jayanthi et al (2012)[9] reported Cefotaxime resistance as (37%) and Ceftazidime resistance to (34%) for *Acinetobacter* which is concordant to our resistance of *Acinetobacter* as (37.2%) resistance to Ceftazidime and (33.7%) to Cefotaxime. *Pseudomonas* showed (44.8%) resistance to Gentamicin which is concordant with Murugan et al (2010)[10] who also reported (42.8%) resistance to Gentamicin, Jeya et al (2013) reported high resistance (51.6%) to Gentamicin.

In the present study Ciprofloxacin resistance to *Pseudomonas* is (48.2%) which is very much lower than Juyal et al (2013)[11] who was reported (73.7%) resistance to Ciprofloxacin for *Pseudomonas*. In our study (11.6%) of Imipenem resistance is reported to *Pseudomonas* which is concordant with Paul et al[5] (9.8%) and Nicholas et al[5] (9.8%) which is much higher resistant with Nagaveni et al[14](32%) and (71%) by Murugan[22]. In our present study (25%) were resistant to imipenem for *Pseudomonas* which is slightly lower than Juyal et al[11] who have been reported (31.3%) resistant to imipenem.

ESBL producing *Pseudomonas aeruginosa* were (38.3%) concordant with the study of Varun Goel et al (42.3%) and Agarwal et al[15](22.2%) where as *Acinetobacter* produces low prevalence of ESBLs as (25%). This similar study was concordant with Sinha et al[16] (28%). In the present study imipenem resistant strains of *Pseudomonas aeruginosa* were MBL producers, detected by phenotypic detection method of MBL production. In our study MBL producing strains of *Pseudomonas* were (11.6%) reasonably similar rate of MBL producers done by Nagaveni et al[17](24%) and (28%) by Anuradha et al[18].

A total of 16 *Acinetobacter* spp. isolates were tested for the antibiotic susceptibility testing. Meropenem sensitivity of 90% has been reported in Shilpa .K Gokale et al[5] study, & also well correlates with study conducted by Karlwosky et al[21](90%) and Taneja et al[20] 12% of resistant which is concordant with our study(%). metallo beta lactamase was detected in 13% of the Meropenem resistant *Acinetobacter baumannii* isolates. Studies from the Indian subcontinent on the occurrence of metallo beta lactamase production by resistant *Acinetobacter* isolates are minimal.

An Indian study on the *Acinetobacter baumannii* species stated that 70.9% of these isolates produced Metallo beta lactamase (Uma KR et al 2009)[22], while another study reported from Kerala, India, states that 21% of the *Acinetobacter baumannii* isolates were found to be Metallo- β -lactamase producers (Anil VK et al 2011)[23]. A large number of clinical isolates identified tentatively as *Shewanella putrefaciens* were shown to be *S.algae* by Nozue et al[24]. Most common infections of skin and soft tissue, and are usually associated with breaches in the skin such as ulcers or following trauma (Chen SC et al 1991)[25]. Among our isolate was found to be sputum sample.

According to the literature, most *Shewanella* infections are treated easily by a combination of surgical therapy, drainage and antibiotics. Poor outcome is associated most often with underlying disease. The main reason for this is that in most of the hospitals, all gram-negative and oxidase-positive organisms are reported as *Pseudomonas* spp. And no further identification is done. A study done by Sharma et al who was also reported that *Shewanella* spp. were sensitivity to all antibiotics and which is similar to our study. Among the Non fermenters one case of *Shewanella* spp. was identified which was usually a environment organism and commensal[26].

Conclusion:

NFGNB though regarded as contaminants are important bacteria causing wide range of nosocomial infections. Variability in sensitivity pattern emphasizes the need for identification of NFGNB and to monitor their susceptibility patterns as it help in proper management of the infection caused by them. These organisms can also spread resistance to other susceptible bacteria by horizontal gene transfer. The most effective antibiotics are Amikacin, Imipenem, Ciprofloxacin, Ofloxacin, and Carbenicillin. Most of the NFs isolated were resistant to Penicillin group of drugs.

Repeated exposure of organisms to antimicrobial agents is thought to enhance the development and maintenance of resistance. Also presence of antimicrobial agent in sub lethal concentration makes an environment suitable for development of resistance. The sensitivity pattern changes from hospital to hospital and population to population. Often in clinical laboratory oxidase-positive non-fermenter gram-negative rods grown on routine laboratory culture media are considered as *Pseudomonas* spp. and further studied to exclude other oxidase-positive gram-negative rods that are thought to be rare. *Shewanella* spp. are very easy to identify and can easily grown on routine bacteriological media. So there is need to look for such rare organisms and not to dispose all oxidase-positive organisms as *Pseudomonas*. It may affect the overall outcome of the patients, but it will definitely help in better understanding of the epidemiology, pathogenesis and preventive aspects of such organisms.

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