

Asian Journal of Research in Medical and Pharmaceutical Sciences

3(1): 1-8, 2018; Article no.AJRIMPS.39732

ISSN: 2457-0745

Antimicrobial Susceptibility Profiles of Pseudomonas aeruginosa Isolates from Patients Attending Health Care Facilities, Ebonyi State, Nigeria

V. U. Olugbue^{1*}, V. O. Nwaugo², M. O. Okata¹, M. C. Korie³, I. Oko⁴ and N. U. Okoro⁵

¹Microbiology Unit, Department of Science Laboratory Technology, Akanu Ibiam Federal Polytechnic, Unwana, P.M.B. 1007, Afikpo, Ebonyi State, Nigeria.

²Department of Microbiology, Faculty of Biological and Physical Sciences, Abia State University, Uturu, P.M.B 2000 Uturu, Abia State, Nigeria.

³Department of Science Laboratory Technology, Imo State Polytechnic, Umuagwo-Ohaji, Imo State, Nigeria.

⁴Medical Clinic, Akanu Ibiam Federal Polytechnic, Unwana, P.M.B 1007, Afikpo, Ebonyi State, Nigeria.

⁵Department of Food Technology, Akanu Ibiam Federal Polytechnic, Unwana, P.M.B 1007, Afikpo, Ebonyi State, Nigeria.

Authors' contributions

This work was carried out in collaboration between all authors. Authors VUO and VON designed and wrote the protocol and first draft of the manuscript. Author VUO managed the literature search and analysis of the study. Authors MOO, MCK, NUO and IO revised the manuscript. All authors read and approved the final manuscript.

Article Information

DOI: 10.9734/AJRIMPS/2018/39732

Editor(s):

(1) Aurora Martinez Romero, Professor, Department of Clinical Biochemistry, Juarez University, Durango, Mexico.

(1) Ronald Bartzatt, University of Nebraska, USA.

(2) Eddie, Chi Man Leung, Prince of Wales Hospital, Hong Kong.

(3) Philippe Huber, Grenoble-Alpes University, France.

Complete Peer review History: http://www.sciencedomain.org/review-history/23446

Received 19th December 2017 Accepted 21st February 2018 Published 5th March 2018

Original Research Article

ABSTRACT

Aim: To investigate the antibiotic susceptibility profile of *Pseudomonas aeruginosa* isolates from wounds of patients and to determine the age group commonly infected with the bacteria.

Study Design: This is a cross-sectional study conducted among patients suspected of having wound infection.

Place and Duration of Study: The study was conducted between May, 2015 and June, 2016 at the Microbiology Laboratory of Akanu Ibiam Federal Polytechnic, Unwana, Ebonyi State, Nigeria. **Methodology:** A total of 165 wound swabs were analysed for the presence of *P. aeruginosa*. Standard microbiology laboratory tests were used to isolate and identify the isolates. Antibiotic susceptibility testing of the isolates was carried out using the disc diffusion method.

Results: A total of 56 (33.94%) P. aeruginosa isolates were identified. The age group, 31 - 40 years had the highest number 28 (50.00%) of P. aeruginosa infection. Within this age group, females 15 (51.72%) were slightly more infected than males 13 (48.15%). In the tertiary hospital (MMH), the highest sensitivity was observed for ofloxacin (32 strains, 78.05% of P. aeruginosa isolates) followed by ciprofloxacin (29 strains, 70.73 % of P. aeruginosa isolates) and ceftazidime (26 strains, 63.41% of P. aeruginosa isolates). The number of isolates resistant to amoxicillin/clavulanate, cefixime and cefuroxime were 34 strains (82.93% of P. aeruginosa isolates), 26 strains (63.41% of P. aeruginosa isolates) and 23 strains (56.10% of P. aeruginosa isolates) respectively. The number of P. aeruginosa isolates from the teaching hospital (FETHA) sensitive to ofloxacin and ciprofloxacin were 9 strains (60.00% of P. aeruginosa isolates) and 8 strains (53.33% of P. aeruginosa isolates) respectively. In FETHA, the isolates showed their highest resistance to amoxicillin/clavulanate (14 strains, 93.33% of P. aeruginosa isolates) followed by cefixime (12 strains, 80.00% of P. aeruginosa isolates). A total of 32 strains (57.14% of P. aeruginosa isolates) were found to be multidrug-resistant (MDR). Regular monitoring of antimicrobial susceptibility profile is essential to guide the physicians in drug prescription against P. aeruginosa strains.

Keywords: Susceptibility; infection; P. aeruginosa; multidrug resistance.

1. INTRODUCTION

Wound infection is universal and the bacterial type varies with geographical location, resident flora of the skin, clothing at the site of wound, time between wound and examination [1]. The development of wound infection depends on the integrity and protective function of the skin [1]. Many bacterial pathogens are involved in wound infection [2].

P. aeruginosa can be found in most environments including soil, water and various types of vegetation. The organism is a clinically important pathogen responsible for a variety of systemic infections such as urinary tract infections. respiratory system infections. gastrointestinal infections, dermatitis, bacteremia, soft tissue infections, bone and joint infections [3]. P. aeruginosa causes infection in immunocompromised patients such as those suffering from burn wounds or receiving cancer chemotherapy [4]. It is a major nosocomial pathogen, particularly dangerous to cystic fibrosis patients [5].

This organism is resistant to many antibacterial drugs [6]. Multidrug-resistant *P. aeruginosa*

phenotype is defined as resistant to one antimicrobial agent in three or more antipseudomonal anti-microbial classes (carbapenems, penicillins/ fluoroquinolones, cephalosporins and aminoglycosides) [7,8]. Multidrug-resistant P. aeruginosa is an important public health issue because the organism is inherently resistant to many drug classes and is able to acquire resistance to all effective antimicrobial drugs [9]. Multidrug-resistant P. aeruginosa develops resistance by various mechanisms like multidrug resistance efflux pumps, biofilm formation, production of βlactamases and aminoglycoside modifying enzymes [8]. Broad-spectrum anti-pseudomonal drugs such as imipenem, ceftazidime, amikacin have been recommended for treatments of infections caused by multiple drug resistant P. aeruginosa [10]. However, resistance to one or more of these anti-pseudomonal drugs during therapy has been widely observed.

The alarming rate of resistance in bacteria pathogens raises concern for the effectiveness of antibiotic therapy [11]. The spread of multidrugresistant pathogens is a real threat to public health and a major concern for infection control

practitioners globally [12]. This spread has paved way for the re-emergence of previously controlled diseases and a high frequency of opportunistic and chronic infection cases in developing countries like Nigeria [12].

This study aims at investigating the antibiotic susceptibility profile of *P. aeruginosa* isolates from wounds of patients and to determine the age group commonly infected with the organism.

2. MATERIALS AND METHODS

2.1 Study Area

One tertiary hospital (MMH) and one teaching hospital (FETHA) in Ebonyi State, Nigeria was used for this study. These hospitals were selected because they receive a large number of patients seeking medical attention and also serve as a referral centre for the state and neighbouring states.

2.2 Sample Collection

A total of 165 wound swabs were collected from patients aged 11 years and above. No specimen was collected from the age group 0 -10 years. All wound swabs collected at the hospitals were inoculated into 10 ml cooked meat broth and incubated at 37°C for 48 hours [13].

2.3 Isolation of Bacteria

A loopful of organisms growing in the cooked meat broth above was inoculated onto Cetrimide agar plates using the streaking technique. The inoculated Petri dishes were incubated at 37°C for 24 hours [14]. Suspected discrete colonies of *P. aeruginosa* that appeared green on cetrimide agar was inoculated into the nutrient broth and nutrient agar slants and incubated at 37°C for 24 hours. The organisms were confirmed by adopting a standard microbiological procedure which includes: colony morphology, Gram stain reaction and biochemical reaction such as

catalase test, indole test, motility test, citrate utilization test, oxidase test, triple sugar iron agar test and urease test.

2.4 Antibiotic Susceptibility Testing

In vitro, antibiotic susceptibility testing of the isolates was carried out using the disc diffusion method as recommended by the Clinical and Laboratory Standard Institute [15]. The antibiotic disc used include: Ceftazidime (30 Cefuroxime (30 µg), Gentamycin (10 μq), Cefixime Ofloxacin (5 (5 μg), μg), (30 Amoxycillin/clavulanate μg), and Ciprofloxacin (5 µg) (Abtek Biologicals Ltd, Liverpool, UK). A sterile Pasteur pipette was used to drop 0.2 ml of the standardized inoculum equivalent to 0.5 McFarland turbidity standards (1.0 x10⁸ cfu/ml) on the surface of dry Mueller-Hinton agar. The inoculum was evenly spread using Hockey stick shaped glass rod. The agar was left for about 10 minutes for the inoculum to dry. Thereafter, antibiotic discs were aseptically placed on the surface of the inoculated Mueller-Hinton agar plate using heat-sterilized forceps. They were incubated at 37°C for 18-24 hours. The diameter of zones of inhibition each antibiotic was for measured millimetre and compared with values provided by the Clinical and Laboratory Standard Institute [15].

2.5 Statistical Analysis

Statistical analysis was carried using the statistical package for social sciences (SPSS) version 15. P < 0.05 was considered significant.

3. RESULTS AND DISCUSSION

Out of 165 wound swabs analysed, 56 (33.94%) *P. aeruginosa* isolates were obtained (Table 1). The most frequent isolates of *P. aeruginosa* was from trauma 39 (44.32%) followed by burn wound 3 (25.00%).

Table 1. Distribution of P. aeruginosa from wound swabs

Type of wound	Number	Number of bacterial isolates obtained	Percentage (%) bacterial isolates
Burn wound	12	3	25.00
Abscess	65	14	21.54
Trauma	88	39	44.32
Total	165	56	33.94

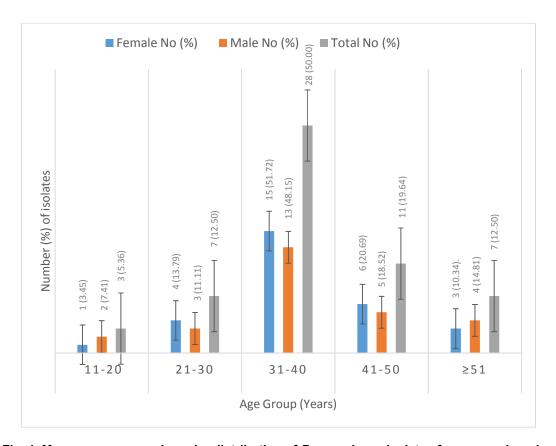


Fig. 1. Mean age group and gender distribution of *P. aeruginosa* isolates from wound swabs.

Key: % = Percentage in bracket

The age group 31 - 40 years had the highest number 28 (50.00%) of *P. aeruginosa* infection. (Fig. 1). Within this age group, females 15 (51.72%) were slightly more infected than males 13 (48.15%).

Antibiotic susceptibility profile of *P. aeruginosa* from MMH is as shown in Table 2. The highest sensitivity was observed for ofloxacin (32 strains, 78.05% of *P. aeruginosa* isolates) followed by ciprofloxacin (29 strains, 70.73% of *P. aeruginosa* isolates) and ceftazidime (26 strains, 63.41% of *P. aeruginosa* isolates). The number of isolates resistant to amoxicillin/clavulanate, cefixime and cefuroxime were 34 strains (82.93% of *P. aeruginosa* isolates), 26 strains (63.41% of *P. aeruginosa* isolates) and 23 strains (56.10% of *P. aeruginosa* isolates) respectively.

The number of *P. aeruginosa* isolates from FETHA sensitive to ofloxacin and ciprofloxacin were 9 strains (60.00% of *P. aeruginosa* isolates) and 8 strains (53.33% of *P. aeruginosa* isolates) respectively (Table 3). The isolates showed their highest resistance to amoxicillin/clavulanate (14 strains, 93.33% of *P. aeruginosa* isolates)

followed by cefixime (12 strains, 80. 00% of *P. aeruginosa* isolates).

A total of 57.14% of *P. aeruginosa* isolates from the two hospitals were found to be multidrug resistant. (Table 4). The prevalence was slightly higher in MMH (58.54%) than FETHA (53.33%).

P. aeruginosa is known to be associated with wound infections. The result of this study was in contrast with a similar study carried out by Garba et al. [2] who isolated 11% of P. aeruginosa from wounds of patients attending Ahmadu Bello University Teaching Hospital, Zaria, Nigeria. Also, Hyder et al. [16] and Mama et al. [17] reported a lower prevalence of 6 (3.95%) and 11 (8%) respectively. Similarly, Ezebialu et al. [18] isolated 27.3% of P. aeruginosa from wounds in Enugu, which is slightly lower than the 33.94% observed in this study. The findings of this study is similar to the reports of Nwachukwu et al. [19] and Anupurba et al. [1] who isolated 32.90% and 32% P. aeruginosa respectively from wound swabs.

Table 2. Antibiotic Susceptibility Profile of P. aeruginosa isolates from Wound swabs, MMH.

Antibiotic	Disc content	Number of sensitive strains (%)	Number of intermediate strains (%)	Number of resistant strains (%)
Ceftazidime	30 µg	26 (63.41)	4 (9.76)	11 (26.83)
Cefuroxime	30 µg	13 (31.71)	5 (12.20)	23(56.10)
Gentamycin	10 µg	15 (36.59)	5 (12.20)	21 (51.22)
Cefixime	5 µg	6 (14.63)	9 (21.95)	26 (63.41)
Ofloxacin	5 µg	32 (78.05)	0 (0.00)	9 (21.95)
Amoxicillin/	30 µg	3 (7.32)	4 (9.76)	34 (82.93)
Clavulanate				,
Ciprofloxacin	5 µg	29 (70.73)	3 (7.32)	9 (21.95)

Key: % = Percentage in bracket; MMH = Code of tertiary hospital used

Table 3. Antibiotic susceptibility profile of *P. aeruginosa* isolates from wound swabs, FETHA.

Antibiotic	Disc content	Number of sensitive strains (%)	Number of intermediate strains (%)	Number of resistant strains (%)
Ceftazidime	30 µg	6 (40.00)	1 (6.67)	8 (53.33)
Cefuroxime	30 µg	5 (33.33)	1 (6.67)	9(60.00)
Gentamycin	10 μg	4 (26.67)	2 (13.33)	9(60.00)
Cefixime	5 µg	2 (13.33)	1 (6.67)	12 (80.00)
Ofloxacin	5 µg	9 (60.00)	1 (6.67)	5 (33.33)
Amoxicillin/	30 µg	1 (6.67)	0 (0.00)	14 (93.33)
Clavulanate	. •	-	•	
Ciprofloxacin	5 µg	8 (53.33)	0 (0.00)	7 (46.67)

Key: % = Percentage in bracket; FETHA = Code of tertiary hospital used

Table 4. Prevalence of MDR isolates of P. aeruginosa from wound swabs

Hospital	Total no. of <i>P. aeruginosa</i> isolates	Number of MDR <i>P. aeruginosa</i> strains	Percentage of MDR P. aeruginosa strains
MMH	41	24	58.54
FETHA	15	8	53.33
Total	56	32	57.14

Key: MDR = Multidrug-Resistant

In a similar study carried out by Kirecci and Kareem [20], the age group mostly infected with *P. aeruginosa* was 41 – 60 years representing 45.33%. Their finding differ from the present study where the age group of 31 – 40 years recorded the highest number (50.00%) of *P. aeruginosa* infection. Rajat et al. [21] reported an isolation rate of 29% in the age group of 31–45 years. This result is similar to our study and that carried out by Chander and Raza [22] that found 20% *P. aeruginosa* infection in age group of 21–40 years.

P. aeruginosa has been reported to have an innate resistance to several antibiotics due to the presence of lipopolysaccharide in the outer membrane [16]. In Nigeria, the trend in resistance phenotype of P. aeruginosa to commonly prescribed antibiotics in various hospitals is increasing since the last decade [10]. In line with the present study, Jombo et al. [23]

and Garba et al. [2] reported a high resistance of respectively, and 81.8% amoxicillin/clavulanate among P. aeruginosa isolates from urine and wound swabs. Rajeevan et al. [24] found 40% and 20% resistance to ceftazidime and gentamycin respectively. This is low compared to the finding of this study in FETHA with ceftazidime and gentamycin resistance of 53.33% and 60.00% respectively. In MMH, the resistance to gentamycin was high (51.22%) compared to 20% resistance reported by Rajeevan et al. [24]. Ahmed et al. [8] reported a high (91%) resistance to ceftazidime. In comparison to the present study in the two hospitals, Hyder et al. [16] observed a higher (70.83%) resistance to gentamycin and a lower (31.25%) resistance to ciprofloxacin in Hilla Teaching hospital, Babylon. The resistance observed against ceftazidime is worrisome, being among the broad-spectrum anti-pseudomonal drugs recommended for treatment of infections caused by MDR *P. aeruginosa* [10]. The high resistance of the isolates to antibiotics may be due to the practice of self-medication, lack of diagnostic laboratory services or unavailability of guidance regarding the selection of drugs, thereby leading to inappropriate use of antibiotics [17].

Various researchers have reported lower rates of MDR P. aeruginosa in their study. Compared to this study, Ahmed et al. [8] detected a slightly low prevalence rate of 52% in patients with nosocomial infections at a University hospital, Egypt. In Iran, Zahra and Moniri [25] reported 30 % prevalence of MDR P. aeruginosa. Siveraj et al. [26] reported 12% MDR among P. aeruginosa isolates from India. In Abeokuta, Nigeria, Okonko et al. [27] reported multidrug-resistant to antibiotics (ampicillin, chloramphenicol, cotrimoxazole, nitrofuratoin and tetracycline) by P. aeruginosa. The level of MDR recorded in this study shows that P. aeruginosa can develop resistance to many antibacterial. P. aeruginosa can develop resistance to many antibacterial both through the resistance genes on extrachromosomal genetic elements and through mutational processes [28].

4. CONCLUSION

In this study, 56 (33.94%) of *P. aeruginosa* isolates were obtained from the wound swabs analyzed. The age group mostly infected with *P. aeruginosa* was 31–40 years. The *P. aeruginosa* strains were more sensitive to ofloxacin while they showed their highest resistance to amoxicillin/clavulanate. Exactly 57.14% of *P. aeruginosa* isolates were found to be multidrug resistant. A regular monitoring of antimicrobial susceptibility profile is essential to guide the physicians in prescribing the right drugs and prevent the emergence of multidrug-resistant strains of *P. aeruginosa*.

CONSENT

All authors declare that written informed consent was obtained from the patients for publication of this case report.

ETHICAL APPROVAL

It is not applicable.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

- Anupurba S, Bhattacharjee A, Garg A, Sen MR. Antimicrobial susceptibility of Pseudomonas aeruginosa isolated from wound infections. Indian Journal of Dermatology, 2010;51:286-288.
- 2. Garba I, Lusa YH, Bawa E, Tijjani MB, Aliyu MS, Zango UU, Raji MIO. Antibiotics Susceptibility Pattern of Pseudomonas aeruginosa isolated from Wounds in Patients Attending Ahmadu Bello University Teaching Hospital, Zaria. Nigeria. Nigerian Journal of Basic and Applied Science. 2012;20(1):32-34. Available: http://www.ajol.info/index.php/njb as/index
- 3. Nasreen M, Sarker A, Malek MA, Ansaruzzaman MD, Rahman M. Prevalence and Resistance Pattern of *Pseudomonas aeruginosa* Isolated from Surface Water. Advances in Microbiology, 2015;**5**:74-1. DOI:http://dx.doi.org/10.4236/aim.2015.51
- Abouelfetouch AY, Moussa NK. Enhancement of antimicrobial activity of four classes of antimicrobials combined with garlic. Asian Journal of Plant Science, 2012.

DOI: 10.3920/ajps.2012.

800

- Jayaraman P, Meena KS, Chu SL, Thean HT, Kishore RS. Activity and interactions of antibiotic and phytochemical combinations against *Pseudomonas aeruginosa* in vitro. International Journal of Biological Sciences, 2010;6(6):556-568.
- Mesaros N, Nordmann P, Plésiat P, Roussel-Delvallez M, Van Eldere J, Glupczynski Y. Pseudomonas aeruginosa: resistance and therapeutic options at the turn of the new millennium. Clinical Microbiology of Infection. 2007;13:560-578.
- 7. Magiorakos AP. Multidrug-Resistant (MDR), Extensively Drug Resistant (XDR) and Pandrug-1 Resistant (PDR) Bacteria in Healthcare Settings. Expert Proposal for a Standardized International Terminology, 2011.

Available: www.escmid.org.

 Ahmed BM, Wafaa AZ, Ghada RH, Aza ZL, Rasha G. Prevalence of multidrugresistant *Pseudomonas aeruginosa* in patients with nosocomial infections at a University Hospital in Egypt, with special reference to typing methods. Journal of Virology and Microbiology. 2013;13. Article ID 290047.

DOI: 10.5171/2013.290047

- Gad GF, El- Domany RA, Zaki S, Ashour HM. Characterization of *Pseudomonasa* eruginosa Isolated from clinical and environmental samples in Minia, Egypt: Prevalence, antibiogram and resistance mechanisms. Journal of Antimicrobial Chemotherapy. 2007;60:1010–1017.
- Odumosu BT, Bolanle AA, Dada-Adegbola H, Ram C. Multidrug resistant Pseudomonas aeruginosa from Southwest Nigeria Hospitals. International Journal of Pharmaceutical Science and Review Research. 2012;15(2):11-15.
- Nmema EE. Peculiar pattern of antibiotic resistance in bacteria isolated from various sources in South-East Nigeria and the implications in health and economy. Journal of Applied Science and Environment. 2013;17(4):529-534.
- Vishal G, Trivedi NA. In vitro evaluation of antimicrobial effect of fresh garlic extract and its interaction with conventional antimicrobials against Escherichia coli isolates. International Journal of Current Research and Reviews. 2012;05(1):106-114.
- 13. Saana SBBM, Adu F, Agyare C, Gbedema CY, Boamah VE, George DF. Antibiotic resistance patterns of strains of Staphylococcus aureus isolated from patients in three hospitals in Kumasi, Ghana. Journal of Bacteriology Research. 2013;5(3):35-40.

DOI: 10.5897/JBR2012.0081

- Betty F, Daniel S, Alice W. Bailey & Scott's Diagnostic Microbiology. 12th edition. Elsevier Inc; New York. 2007;322-325.
- CLSI. Performance standards for antimicrobial susceptibility testing; twentieth informational supplement CLSI document M100–S20 Wayne, PA. 2012.
- Hyder H, Tarrad JK, Banyan HA. Isolation of *Psudomonas aeruginosa* from clinical and environmental samples, and analysis of its antibiotic resistant spectrum at Hilla Teaching Hospital. Medical Journal of Babylon. 2011;8(4):618-624.
- 17. Mama M, Abdissa A, Sewunet T. Antimicrobial susceptibility pattern of bacterial isolates from wound infection and their sensitivity to alternative topical agents at Jimma University Specialized Hospital, South-West Ethiopia. Annals of Clinical

Microbiology and Antimicrobials, 2014; 13:14.

Available: http://www.ann-clinmicrob.com/content/13/1/14

18. Ezebialu CU, Chukwurah EI, Ezebialu IU. Bacterial pathogens associated with wound infections at National Orthopaedic Hospital, Enugu. Nigerian Journal of

Microbiology. 2010;24(1):1987-1992.

- Nwachukwu NC, Orji FA, Okike UM. Antibiotic susceptibility patterns of bacterial isolates from surgical wounds in Abia State University Teaching Hospital (ABSUTH), Aba – Nigeria. Research Journal of Medicine and Medical Sciences. 2009;4(2): 575-579.
- Kireçci E, Kareem RD. Antibiotic susceptibility patterns of *P. aeruginosa* strains isolated from various clinical specimens. Sky Journal of Microbiology Research. 2014;2(2):013–017.
 Available:
 http://www.skyjournals.org/SJMR ISSN 2315-876X.
- 21. Rajat RM, Ninama GL, Mistry K, Parmar R, Patel K, Vegad MM. Antibiotic resistance pattern in *Pseudomonas aeruginosa* species isolated at a tertiary care hospital, Ahmadabad. National Journal of Medical Research. 2012;2(2):156–159.
- 22. Chander A, Raza MS. antimicrobial susceptibility patterns of *Pseudomonas aeruginosa* clinical isolates at a tertiary care hospital in Kathmandu, Nepal. Asian Journal of Pharmaceutical and Clinical Research. 2013;6(3):235–38.
- 23. Jombo GTA, Jonah P, Ayeni JA. Multidrug resistant *Pseudomonas aeruginosa* in contemporary medical practice: findings from urinary isolates at a Nigerian University Teaching Hospital. Nigerian Journal of Physiological Sciences. 2008;23:105-109.
- 24. Rajeevan S, Ahmad SM, Jasmin PT. Study of prevalence and antimicrobial susceptibility pattern in blood isolates from a tertiary care hospital in North Kerala, India. International Journal of Current Microbiology and Applied Sciences. 2014;3(4):655-662.
- Zahra T, Moniri R. Detection of ESBLs and MDR in *Pseudomonas aeruginosa* in a Tertiary-Care Teaching Hospital. Iranian Journal of Clinical Infectious Diseases. 2011;6(1):18-23.
- 26. Sivaraj S, Murugesan P, Muthuvelu S, Purusothaman S, Silambarasan A. Comparative study of *Pseudomonas*

- aeruginosa isolate recovered from clinical and environmental samples against antibiotics. International Journal of Pharmacy and Pharmaceutical Sciences. 2012;4(3):103-107.
- 27. Okonko IO, Soleye FA, Amusan TA, Ogun AA, Ogunnusi TA, Ejembi J. Incidence of Multi-Drug Resistance (MDR) Organisms in Abeokuta, Southwestern Nigeria.
- Global Journal of Pharmacology. 2009;3(2):69-80.
- 28. Lister PD, Wolter DJ, Hanson ND. Antibacterial-resistant Pseudomonas aeruginosa: Clinical impact and complex regulation of chromosomally encoded resistance mechanisms. Clinical Microbiology Reviews. 2009;22(4):582–610.

© 2018 Olugbue et al.; This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Peer-review history:
The peer review history for this paper can be accessed here:
http://www.sciencedomain.org/review-history/23446