



GLOBAL JOURNAL OF MEDICAL RESEARCH: K
INTERDISCIPLINARY

Volume 15 Issue 3 Version 1.0 Year 2015

Type: Double Blind Peer Reviewed International Research Journal

Publisher: Global Journals Inc. (USA)

Online ISSN: 2249-4618 & Print ISSN: 0975-5888

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GJMR-K Classification: NLMC Code: QY 185



COMMUNITY ACQUIRED URINARY TRACT INFECTION PREVALENCE IN A TERTIARY INSTITUTION BASED IN EVBUOBANOSA EDO STATE NIGERIA

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Otajewwo, F. D^α & Amedu, S. S^σ

Abstract- The prevalence rate of asymptomatic UTI among students of a community tertiary institution based in Evbuobanosa near Benin City, Nigeria was the focus of this work. A total of 390 freshly voided midstream urine samples collected into sterile plastic screw capped universal containers containing boric acid as preservative from students aged between 15 – 30 years were used for the study. Samples were screened for significant bacteriuria and puscells (neutrophil) counts. All positive samples with pyuria were aseptically cultured by standard methods on sterile Cystine Lactose Electrolyte Deficient (CLED) agar, MacConkey agar and Sabouraud Dextrose agar plates and incubated appropriately. Antibiotic susceptibility testing was done on isolated and identified uropathogens by Kirby – Bauer agar diffusion disc method. While 231(59.2%) samples yielded bacterial growth, 159(40.8%) yielded no growth. The uropathogens isolated included *Staphylococcus aureus* (33.1%), *Escherichia coli* (20.8%), coagulase negative Staphylococci (15.6%), *Klebsiella aerogenes* (7.9%), *Coliform* organisms (7.9%), *Candida albicans* (7.9%) and *Proteus* spp (6.8%) of which gram negative bacilli and gram positive bacteria accounted for 43.4% and 48.7% respectively. *Candida albicans*, a fungal uropathogen accounted for 7.9%. Occurrence of UTI in male and female students was 57.1% and 42.9% respectively of which UTI occurred highest in the 21 – 25 age group. More than 50% of isolated bacterial strains were sensitive to gentamicin only with more than 90% resistant to augmentin and nalidixic acid. *Staph aureus*, *Kleb aerogenes*, *Coliform* organisms and *Proteus* spp were multidrug resistant as they resisted 4, 7, 5 and 7 drugs respectively. Findings emphasize the need for regular and routine isolation of uropathogens in health institutions and subjection of same to invitro antibiotic sensitivity testing in order to obtain updated antibiograms of UTI pathogens. This will address the emergence of resistance genes and reduce healthcare cost and long stay in hospitals.

I. INTRODUCTION

Urinary tract infection(UTI) is the most common infection experienced by humans after respiratory and gastro-intestinal infections and also the most common cause of both community-acquired and hospital acquired (nosocomial) infections for patients admitted to the hospitals (Najaret *et al.*,2009). UTI can be asymptomatic or symptomatic characterized by a wide range of symptoms from mild voiding irritation to

bacteraemia, sepsis or even death (Ranjbar *et al.*, 2009).

Infection of the urinary tract could manifest differently depending on the site of the infection and length of time involved (Takhar, 2011). Those that affect the lower urinary tract are called cystitis(i.e. involving the bladder alone with symptoms including painful urination, burning sensation, frequent urination or urge to urinate or both while those that affect upper urinary tract are referred to as pyelonephritis(i.e. involve the kidneys and other organs (Sarah,2010). The symptoms of the upper urinary tract infection include fever and flank pain during urination in addition to those of the lower urinary tract (Sarah 2010).

Urinary tract infection occurs more frequently in females than males due to the shortness and width of the female urethra to the vagina which makes it liable to trauma during sexual intercourse as well as bacteria being passed from the urethra into the bladder during pregnancy (Ebie *et al.*, 2001). The moist environment of the female's perineum favours microbial growth and predisposes the female bladder to bacterial contamination (Ebie *et al.*, 2001). In addition, urine of females was found to have more suitable pH and osmotic pressure for the growth of *Escherichia coli* than urine from males (Obiogbolu *et al.*, 2004).

Most UTIs are caused by gram negative bacteria like *Escherichia coli* and *Klebsiellaspp* (Omonighoet *et al.*, 2001; Ebie *et al.*,2001). Other bacterial pathogens frequently isolated include gram positive bacteria such as *Staph. aureus*, *Staph. epidermidis* and *Enterococcus*spp formerly called *Strept. faecalis* (Kashefet *et al.*, 2010; Theodros, 2010;Mulugeta and Bayeh,2014) as well as *Proteus mirabilis*, *Pseudomonas aeruginosa*, coagulase negative staphylococci, *Acinetobacterspp* and *Serratiaspp* (Shankel,2007).

Drug resistance among bacteria causing UTI has increased since the introduction of UTI chemotherapy (Nerurkar *et al.*,2012; Sood and Gupta,2012; Bahadin *et al.*,2011; Haideret *et al.*,2010). The aetiological agents and their susceptibility patterns vary in various regions and geographical locations (Mulugeta and Bayeh, 2014). Knowledge of the local bacterial aetiology and susceptibility pattern is required to trace any change that might have occurred with time so that

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updated recommendation for optimal empirical therapy of UTI can be made (Leegardet *et al.*, 2000).

A number of studies have been done on the prevalence and antimicrobial resistance patterns of UTIs (Onhet *et al.*, 2006; Mbata, 2007; Okonko *et al.*, 2009). Be that as it may, there is no documented UTI study that has been carried out in Ewobanosa community. Ewobanosa community is a bini speaking community situated about 20km from Benin City, Edo State of Nigeria. The community has a privately owned polytechnic situated tangentially on the express road that connects Benin City to Asaba (in Delta State of Nigeria). It was therefore to extend the frontiers of knowledge of UTI that this work aimed at studying community acquired urinary tract infection prevalence among students in a tertiary educational institution in Evbuobanosa, Nigeria was carried out with the following objectives:

- Determine the frequency distribution of microbial uropathogens in urine samples obtained from the students.
- Determine the sex distribution of uropathogens isolated.
- Determine the age distribution of the uropathogens isolated.
- Determine the antibiotic susceptibility profiles of isolated uropathogens.

II. MATERIALS AND METHODS

a) Sampling

A total number of the three hundred and ninety (390) midstream urine samples were collected from students (both residential and non-residential) of Lighthouse Polytechnic situated in Evbuobanosa, a community in Orhionwon local government area of Edo State of Nigeria. Samples (which were freshly voided), were collected into sterile screw capped plastic universal containers containing a few crystals of boric acid as preservative. Recruited students were instructed on how to collect the samples. All samples were appropriately labeled and processed immediately in the Microbiology laboratory of Western Delta University, Oghara, Delta State. Students recruited were grouped into 15-20, 21-25 and 26-30 age brackets. The study designed was a descriptive cross sectional study. Samples were collected in March, 2014.

b) Ethnical Clearance

Ethnical clearance was sought and granted by the ethnical committee of the Polytechnic. In addition, the recruited students gave their verbal informed consent after thorough explanation of the rationale for the study.

c) Processing of Samples

Samples were tested for significant bacteriuria by use of a modified semi quantitative technique

described by Mbata (2007). A standard bacteriological loopful of each urine sample (0.01ml) was spread over the surface of sterile Cystine Lactose Electrolyte Deficient (CLEDI) agar plates (LabM, UK). After inoculation, the plates were inverted and incubated at 37°C for 18-24 hrs. The number of bacterial colonies was counted and multiplied by 100 to give an estimate of the number of bacterial organisms per millilitre of urine. A significant bacterial count was taken as any count equal to or in excess of 10⁵ per millilitre.

d) Confirmation of Significant Bacterial Count by Microscopy

All samples that recorded significant bacterial counts were subjected to urine microscopy test to detect presence of five pus cells per high power focus (5PC/HPF) or 10 white blood cells (pus cells) /mm³ in urine sediments or deposits (Smith, 2004) using x40 objective microscopically. All samples that were positive for significant bacterial count and also recorded 5PC/HPF or 10PC/mm³ or more were cultured on suitable laboratory media.

e) Cultural Studies

Urine sample that recorded 5PC/HPF or 10PC/mm³ and were positive for significant bacteriuria test were cultured aseptically on sterile CLED agar (Lab M, UK), MacConkey agar (Lab M, UK) and Saboraud Dextrose agar (Lab M, UK) plates according to standard methods. All inoculated plates were incubated at 37°C for 24hrs. Pure isolates were then obtained and identified according to schemes provided by Cowan and Steel (1993) and Alexopoulos (1996). All identified isolates were subjected to antibiotic sensitivity testing. Fungal isolates were excluded from antibiotic sensitivity testing. All urine samples that yielded no bacterial growth were noted.

f) Antibiotic Sensitivity Testing

Antibiotic susceptibility testing was carried out on confirmed uropathogens according to the agar diffusion disc technique described by Bauer *et al.* (1966). A colony of each pure (axenic) isolate was streaked on sterile Mueller Hinton agar plates aseptically using sterile inoculating wire loop. The relevant multidiscs containing their respective minimum inhibitory concentrations (MICs) were used and included erythromycin (5ug), cefuroxime (30ug), gentamicin (10ug), nalidixic acid (30ug), cefixime (5ug), ceftazidime (30ug), ofloxacin (10ug), augmentin (30ug), nitrofurantoin (200ug), ciprofloxacin (10ug), streptomycin (10ug) chloramphenicol (30ug), tetracycline (10ug) and cotrimoxazole (25ug). Discs were aseptically placed (impregnated) firmly onto the surface of the dried plates using sterile forceps. The plates were left at room temperature for one hour to allow diffusion of the different antibiotics from the disc into the medium (Mbata, 2007).

The plates were then incubated at 37°C for 18hrs. Interpretation of results was done using the length of diameter of zone of inhibition (NCCLS, 2000). Zones of inhibition greater than 10mm were considered sensitive, 5-10mm moderate sensitive and no zone of inhibition, resistant.

g) *Statistical Analysis*

Simple percentages were used throughout.

III. RESULTS

Table 1 shows the isolated identified uropathogens from the samples that yielded growth after 24hrs incubation period. A total of 390 midstream urine samples were processed of which 159 (40.8%) yielded no growth at the end of 48hrs incubation at 37°C. A total of 231 (59.2%) samples yielded growth of seven microbial uropathogens were identified as follows: Bacterial colonies that were in clusters, gram positive, catalase positive, coagulase positive, DNAase negative, phosphatase positive, mannitol fermenting, raised, round and smooth colonies were identified as *Staphylococcus aureus*. All gram negative raised entire, circular, motile, lactose, glucose fermenting, indole positive, methylred positive, voges praskauer negative, citrate negative and urease negative bacilli strains were identified as *Escherichia coli*.

Bacterial colonies that were positive for all the characteristics of *Staph aureus* colonies but were negative for slide and tube coagulase tests were identified as coagulase negative staphylococci. Gram negative mucoid non-motile, lactose, adonitol, inositol, glucose fermenting, voges praskauer positive, urease positive, citrate positive and indole negative bacilli strains were identified as *Klebsiella aerogenes*. Coliform organisms were mixed cultures made of *Escherichia coli*, *Enterococcus faecalis* and *Clostridium perfringens*. Gram positive yeast cells, pseudohyphae positive, chlamydo spores positive, germtube positive, glucose, maltose galactose and sucrose assimilation positive strains were identified as *Candida albicans*. Gram negative, swarming, fish odour colonies on sodium chloride containing media, indole negative and urease positive strains were identified as *Proteus* spp. The frequency occurrence of the isolates and their strains therefore were as follows: *Staphylococcus aureus* (33.1%) *Escherichia coli* (20.8%), coagulase negative Staphylococci (15.6%), *klebsiella aerogenes* (7.9%), Coliform organism (7.9%), *Candida albicans* (7.9%) and *Proteus* spp (6.8%). On the whole, total gram negative and gram positive bacilli isolated represented 43.4% and 48.7% respectively of which 43.4% and 56.6% belonged to enterobacteraceae and non enterobacteraceae respectively.

Table 1 : Frequency of distribution of microbial uropathogens isolated from Samples

Isolated Uropathogens	No of Strains	Frequency of Strains
<i>Staphylococcus aureus</i>	75	33.1%
<i>Escherichia coli</i>	48	20.8%
Coagulase negative Staph	36	15.6%
<i>Klebsiella aerogenes</i>	18	7.9%
Coliform organisms	18	7.9%
<i>Candida albicans</i>	18	7.9%
<i>Proteus</i> spp	15	6.8%
Total n=7	231	100.0%

Gram negative bacteria: 43.4%
 Gram positive bacteria: 48.7%
 Enterobacteriaceae: 43.4%
 Non-Enterobacteriaceae: 56.6%

Table 2 shows the sex distribution of the microbial strains isolated. Out of the total 231 strains isolated, 132 (57.1%) and 99 (42.9%) were obtained from male and female students respectively. In a

decreasing order, the highest occurring uropathogens in male students were *Staph aureus* (34.1%), *Escherichia coli* (22.7%), coagulase negative staphylococci (18.2%), Coliform organisms (9.1%), *Klebsiella aerogenes* (6.8%),

Candida albicans (4.6%) and *Proteus* spp (4.6%). Similarly *Staph aureus* (30.3%), *Escherichia coli* (18.2%), coagulase negative staphylococci (12.1%), *Candida albicans* (12.1%), *klebsiella aerogenes* (9.1%), *Proteus* spp (9.1%) and Coliform organisms (6.1%) uropathogens were isolated from female students.

Table 2 : Sex Distribution of uropathogens Isolated from Processed Samples

Isolated Uropathogens	No of Strains males (%)	No of Strains females (%)	Total %
<i>Staphylococcus aureus</i>	45 (34.1)	30 (30.3)	75(32.5)
<i>Escherichia coli</i>	30 (22.7)	18 (18.2)	48 (20.8)
Coag. Neg. Staph	24 (18.2)	12 (12.1)	36 (15.6)
<i>Klebsiella aerogenes</i>	09(6.8)	09 (9.1)	18 (7.8)
Coliform organisms	12 (9.1)	06 (6.1)	18 (7.8)
<i>Candida albicans</i>	06 (4.6)	12 (12.1)	18 (7.8)
<i>Proteus</i> spp	06 (4.6)	09 (9.1)	15 (6.5)
Total n=7	M 132 (57.1%)	F 99(42.9%)	231(100.0%)

The age distribution of the entire 231 uropathogen strains isolated is shown in Table 3. Recruited students were grouped into 15-20, 21-25 and 26-30 age brackets. Hence, 102 (44.2%), 87 (37.7%) and 42 (18.2%) of the sampled students belonged to 21-25, 15-20 and 26-30 age groups respectively. Seventy two (70.6%), 45(51.7%) and 15(35.7%) students of 21-25, 15-20 and 26-30 age group respectively were males while 30 (29.4%), 42(48.3%) and 27(64.3%) students of the same age groups were females. Consequently, the highest number of uropathogen strains of 72(70.6%) was isolated from male students belonging to 21-25 age group. Fifteen (35.7%) uropathogen strains were the least isolated from male students who occurred in the 26-30 age bracket. The highest and lowest uropathogen strains of 42 (48.3%) and 27 (64.3%) respectively were isolated from female students of 15-20 and 26-30 age group respectively.

Table 3 : Age Distribution of uropathogens Isolated

Uropathogens Isolated	15 -20		21 -25		26 – 30	
	M	F	M	F	M	F
<i>Staph aureus</i>	18 (40.0)	12 (28.6)	21 (29.2)	09 (30.0)	06 (40.0)	09 (33.3)
<i>Escherichia coli</i>	09 (20.0)	12 (28.6)	18 (25.0)	06 (20.0)	03 (20.0)	0 (0.0)
<i>Coag. neg Staph</i>	09 (20.0)	03 (7.2)	12 (16.7)	03 (10.0)	03 (20.0)	06 (22.2)
<i>Klebsiella aerogenes</i>	06 (13.3)	03 (7.2)	03 (4.2)	03 (10.0)	0 (0.0)	03 (11.1)
<i>Coliform organisms</i>	0 (0.0)	06 (14.3)	09 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)
<i>Candida albicans</i>	0 (0.0)	03 (7.2)	03 (4.2)	06 (20.0)	03 (20.0)	06 (22.2)
<i>Proteus spp</i>	0 (0.0)	03 (7.2)	06 (8.3)	03 (10.0)	0 (0.0)	03 (11.1)

In Table 4, the antibiotic susceptibility patterns of all isolated uropathogens (with exception of *Candida albicans* and coagulase negative staphylococci) to some selected antibiotics are shown. The antibiotic sensitivity profiles of the 231 uropathogen strains to ciprofloxacin, cefuroxime, gentamicin, cefixime, ceftazidime, ofloxacin, streptomycin, erythromycin, tetracycline, cotrimoxazole, augmentin, nalidixic acid, nitrofurantoin, chloramphenicol are shown. Twenty four (32.0%), 11 (22.9%), 06 (33.4%) 09(50.0%) and 06(40.0%) strains of *Staphylococcus aureus*, *Escherichia coli*, *Klebsiella aerogenes*, Coliform organisms, and *Proteus spp* respectively were sensitive to all the 14 (fourteen) antibiotics used.

With respect to *Staph aureus*, 12 antibiotics used were twelve because nalidixic acid and nitrofurantoin were not tested on it and whereas erythromycin and streptomycin were used for *Staph aureus*, they were not used for the other uropathogens. On the average, a higher percentage of pathogens resisted all fourteen antibiotics compared to those that were sensitive.

In terms of effectiveness of each of the antibiotics used, 108(62.1%), 60(34.5%), 48(27.6%), 45(25.9%), 39 (22.4%), 24(13.8%), 24 (13.8%), 21 (12.1%), 15(8.6%), 15(8.6%), 12(6.9%), 12(6.9%), 06(3.5%) and 03(1.7%) strains of all isolated pathogens were sensitive to gentamicin, ofloxacin, streptomycin, nitrofurantoin, ciprofloxacin, tetracycline, cotrimoxazole, chloramphenicol, ceftazidime, cefixime, erythromycin, cefuroxime, augmentin and nalidixic acid in that decreasing order. The two most sensitive drugs in this study were therefore gentamicin and ofloxacin whereas augmentin and nalidixic acid were the least sensitive. Hence, augmentin and nalidixic acid were the most

resisted antibiotics with ofloxacin and gentamicin as the least resisted.

Table 4 : Susceptibility Profiles of Isolated Uropathogens to Selected Antibiotics after 24hrs Incubation at 37°C

Isolated uropathogens	SELECTED ANTIBIOTICS USED (%)														MEAN (%)
	ERY	CRX	GEN	NAL	CXM	CAZ	OFL	AUG	NIT	CPR	STR	CHL	TET	COT	
<i>Staph aureus</i> n = 75	12 (16.0)	0 (0.0)	66 (88.0)	NA	0 (0.0)	0 (0.0)	12 (16.0)	0 (0.0)	NA	06 (8.0)	48 (64.0)	18 (24.0)	12 (16.0)	18 (24.0)	24 (32.0)
<i>Escherichia coli</i> n = 48	NA	06 (12.5)	24 (50.0)	03 (6.3)	09 (18.8)	03 (6.3)	30 (62.5)	06 (12.5)	27 (56.3)	12 (25.0)	NA	03 (6.3)	06 (12.5)	03 (3.3)	11 (22.9)
<i>Klebsiella aerogenes</i> n = 18	NA	0 (0.0)	06 (33.4)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	06 (33.4)	06 (33.4)	NA	0 (0.0)	06 (33.4)	03 (16.7)	06 (33.4)
<i>Coliform organisms</i> n = 18	NA	06 (33.4)	12 (66.7)	0 (0.0)	06 (33.4)	12 (66.7)	12 (66.7)	0 (0.0)	06 (33.3)	09 (50.0)	NA	0 (0.0)	0 (0.0)	0 (0.0)	09 (50.0)
<i>Proteus spp</i> n = 15	NA	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	06 (40.0)	0 (0.0)	06 (40)	06 (40)	NA	0 (0.0)	0 (0.0)	0 (0.0)	06 (40.0)
Total n = 174	12 (6.9)	12 (6.9)	108 (62.1)	03 (1.7)	15 (8.6)	15 (8.6)	60 (34.5)	06 (3.5)	45 (25.9)	39 (22.4)	48 (27.6)	21 (12.1)	24 (13.8)	24 (13.8)	24 (13.8)

Table 5 shows the frequency occurrence of multidrug resistance among the isolated bacterial uropathogens. All the bacterial uropathogens apart from *Escherichia coli* were resistant to more than three antibiotics at a time. For clarity and for the purpose of this study, a pathogen is described as multidrug resistant to any of the selected antibiotics if more than 50% of its strains are resistant to it and hence, a

pathogen is multidrug resistant if it resists up to three drugs at a time. Therefore, only four out of the five bacterial uropathogens isolated were multidrug resistant. None was resistant to 3 drugs. *Staph aureus* strains were resistant to 4 drugs while *Coliform* organisms were resistant to 5 drugs. *Klebsiella aerogenes* and *Proteus* spp strains were resistant to 6 drugs each.

Table 5 : Occurrence of Multidrug Resistance among bacterial Uropathogens Isolated

Bacterial Uropathogens	Resistance to				
	3 drugs	4 drugs	5 drugs	6 drugs	>6 drugs
<i>Staph aureus</i> n = 75	-	+	-	-	-
<i>Escherichia coli</i> n = 48	-	-	-	-	-
<i>Coliform orgs</i> n = 18	-	-	+	-	-
<i>Proteusspp</i>	-	-	-	-	+

IV. DISCUSSION

In this study, out of 390 urine sample processed, 159(40.8%) samples yielded no growth and no significant microbial growth combined. Hence, 231(59.2%) samples yielded growth and significant growth all together. The absence of bacterial growth in 40.8% of processed samples was established as such when microscopically, a non-significant pus cell count (less than 5) was observed (Otajewwo, 2014). That 231 (59.2%) processed samples yielded significant microbial growth suggested a UTI prevalence rate of 59.2% in this study. This finding is consistent with 55%, 54,58%,60% and 66% prevalence rates reported by earlier authors (Onuoha and Fatokun, 2014;Tolkoff and Rubin, 1986; Naylor, 1984; Warren, 1987; Kunin 1985) and inconsistent with higher prevalence rates of 77.3% 71.3%, 71.6%, 77.9%, and 86.0% as reported by some authors (Otajewwo, 2014; Jellheden and Norrby, 1996; Mbata 2007; Alabi *et al.*, 2014). It is on record that lower rates of 30.0%, 35.5%, 38.5%, 38.6%, 39.0%, 39.0% and 39.7% have been reported by some authors (Anochie *et al.*, 2001; Ebie *et al.*, 2001; Anigilaje and Bitto 2013; Akinyemi *et al.*, 2000; Ogomaka *et al.*, 2014; Otajewwo 2013 and Oladeinde *et al.*, 2011). Other previous authors recorded much lower UTI prevalence rates of 11.9%

26.7, 28.2% and 28.9% (Aiyegero*et al.*, 2007; Sobazak 2010;Harvna*et al.*, 2014; Okon *et al.*, 2014). The high or low incidence rates may be attributed to poor environmental conditions where the subjects reside in terms of lack of proper personal and environmental hygiene, low socio – economic status, sexual intercourse or sexual promiscuity, pregnancy e.t.c among Nigerian men and women (Akinyemi *et al.*, 2000; Kolawole *et al.*, 2009; Andriole, 1985).

Seven uropathogens were isolated in this study (Table 1) of which gram negative bacilli constituted 43.4% while gram positive bacteria accounted for 48.7%. Isolates that belonged to enterobacteriaceae and non-enterobacteriacease families were 43.4% and 56.6% respectively (Table1). Whereas *Staphylococcus aureus* and coagulase negative staphylococci were the only gram positive bacteria, *Candida albicans* represented the only member of the non-enterobacteriacease. Finding is consistent with the report of a previous author who isolated 86.1% gram negative bacilli and 13.9% gram positive bacteria of which enterobactereaceae accounted for 49.9% (Otajewwo, 2013). The report of this study does not agree with a report of Oluremi *et al.* (2011) which stated 85.0% gram negative bacilli and 15.0% gram negative of which enterobactereaceae organisms constituted

66.7%. The report of Otajewwo (2014) stating 65.2% and 34.8% gram negative and gram positive bacteria respectively as well as 55.7% and 44.3% enterobactereaceae and non – enterobactereaceae respectively as isolates is also not consistent with the findings in this study.

In this study, the most and second most occurring uropathogens were *Staphylococcus aureus* (33.1%) and *Escherichia coli* (20.8%). Other uropathogens isolated were coagulase negative staphylococcus (15.6%), *Klebsiella aerogenes* (7.9%), *Coliform* organisms (7.9%), *Candida albicans* (7.9%) and *Proteus* spp. This result suggests that the two least isolated uropathogens in the study area are *Candia albicans* and *Proteus* spp. The occurrence of *Staphaureus* and *E.coli* as the most commonly occurring uropathogens is consistent with the report of a previous study (Otajewwo, 2014) as well as with reports (though in reverse order) of some earlier authors which stated *E.coli* and *Staphaureus* as the most and second most implicated organisms in UTI (Onuoha and Fatokun, 2014; Okon *et al.*, 2014; Akobi *et al.*, 2014; Ogomaka *et al.*, 2014, Okonko *et al.*, 2010; Nwanze *et al.*, 2007).

The present report however, does not agree with results of other studies (Omgie *et al.*, 2009; Abubakar, 2009; Otajewwo, 2013; Oluremi *et al.*, 2011; Daza *et al.*, 2001; Dimitrov *et al.*, 2004; Inabo and Obanibi, 2006; Dash *et al.*, 2008; Nwadioha *et al.*, 2010). Report in this study does not also agree with some other previous reports which stated *Klebsiella* spp and *Proteus* spp (Alabi *et al.*, 2014), *E.coli* and *Streptococcus faecalis* (Anigilaje and Bitto, 2014), *E.coli* and *Proteus* spp (Haruna *et al.*, 2014), *E.coli* and *Pseudomonas aeruginosa* (Habibu, 2014) as the most and second most occurring uropathogens in UTI. In this study, *Proteus* spp, *Klebsiella* spp and *Staph aureus* were isolated and these organisms have been incriminated in hospital acquired infections often following catheterization or gynaecological surgery (Cheesbrough 2000; Tapsak *et al.*, 1995). *Proteus* infection is also associated with renal stones (Cheesbrough, 2000). In this study and previous similar studies, it is yet again confirmed the prominent involvement of *Staphaureus* and *E.coli* in UTI cases (Logoria and Gonzalez, 1987; Obiogbolu *et al.*, 2009; Otajewwo and Eriagbor, 2014; Okon *et al.*, 2014; Onuoha and Fatokun, 2014; Ogomaka *et al.*, 2014; Akobi *et al.*, 2014).

Overall, UTI prevalence rate was higher in males (57.1%) than in the female students (42.9%) of which *Staphaureus*, *E.coli* and *Coliform* organisms occurred more in the male students than in the female students (Table 2). The earlier report of Otajewwo (2014) which recorded higher UTI prevalence rate does not agree with the finding of this report. The reason for a higher UTI prevalence rate in males than females in this study though not clear, may be due to lack of

circumcision, receptive anal intercourse (as in homosexuals) and HIV infection (Orret and Davis, 2006). Also in this study, *Candida albicans* occurrence was higher in female students compared to the males. According to Ochei and Kolhatkar (2008), yeast cells appear in urine as a result of contamination from women with vaginal candidiasis (occasionally seen in the urine of men) or may be seen in the urine of diabetic patients due to presence of sugar in the urine. Yeasts may also cause recurrent infections in debilitated and immune-compromised patients (Cheesbrough, 2000; Ochei and Kolhatkar, 2008). The occurrence of *Coliform* organisms is crucial in pregnant women although its occurrence in females is lower compared to males. The isolation of *Candida albicans* and *Coliform* organisms in this study is consistent with the report of a previous work (Otajewwo and Eriagbor, 2014). It has been documented that *Coliforms* and *Enterococcus* spp can cause UTI when present in high numbers on the perineum (Behzardi and Behzardi, 2010; Moore *et al.*, 2002).

This study, strangely, did not confirm reports of other studies which stated that UTI occurs more in females than in males except at the extremes of life (Ebie *et al.*, 2001; Kolawole *et al.*, 2009). In terms of age bracket, UTI occurred highest (44.2%) in the 21–25 age group followed by 37.7% occurrence rate of students in the 15 – 20 age bracket. This finding does not agree with the reports of some previous authors which stated that UTI is more frequent in females than in males (Mbata, 2007; Ibeawuchi and Mbata, 2002; Asinobi, 2002; Olaita, 2006). Findings in this work however, agree with the reports of these same authors which stated that UTI is most prevalent during youth and adulthood as indicated by the 21 – 25 age group occurring as the most implicated in UTI in this study. This age group as well as the 15 – 20 group consist of teenagers, adolescents and young people who are characteristically vulnerable to increased sexual activities that predispose them to UTI (Oladeinde *et al.*, 201, Oluremi *et al.*, 2011).

There is considerable evidence of practice variation in the use of diagnostic tests, interpretation of signs or symptoms and initiation of antibiotic treatment such as drug selection, dose, duration and route of administration (Jamieson, 2006). For patients with symptoms of UTI and bacteriuria, the main aim of treatment is to get rid of bacteria causing the symptoms. Besides, there is need to obtain sensitivity reports before the start of antibiotic treatment to checkmate emergence of resistance strains and thus help in proper patient management. The decision to use a particular antibiotic however depends on its toxicity, cost and attainable level (Ibeawuchi and Mbata, 2002).

Therefore, antibiotic sensitivity testing was done on all seven isolates (minus *Candida albicans* – a fungal uropathogen in this study) and the resulting sensitivity

profile (antibiogram) showed susceptibility reactions of 108 (62.1%), 60(34.5%), 48(27.6%), 45(25.9%), 39(22.4%), 24(13.8%), 24(13.8%), 21(12.1%), 15(8.6%), 12(6.9%), 06(3.5%) and 03(1.7%) for gentamicin, ofloxacin, streptomycin, nitrofurantoin, ciprofloxacin, tetracycline, cotrimoxazole, chloramphenicol, cefixime, ceftazidime, erythromycin, cefuroxime, augmentin and nalidixic acid respectively. This suggests that more than 50% of the bacterial uropathogens implicated in UTI in this study were sensitive to gentamicin only. This finding with respect to gentamicin is consistent with the reports of some previous studies (Okon *et al.*, 2014; Akobi *et al.*, 2014; Onuoha and Fatokun, 2014; Otajewwo and Eriagbor, 2014). Less than 50% of the isolated uropathogens were sensitive to ofloxacin and streptomycin

It is cheering and surprising that gentamicin recorded more than 50% sensitivity. Cheering because, it is cheap and readily available and hence, patients who go down with UTI can afford it if prescribed. The use and administration of gentamycin however, should be closely monitored in order to avoid any possible abuse owing to its accessibility, availability and cheapness. It is surprising because, gentamicin ought to be prone to abuse (due to its cheapness) with the attendant development of resistance genes by pathogens to it. It can be reasoned however that gentamycin though cheap, is not easily accessible for abuse because it is available in injection form only. The fluoroquinolone–ofloxacin recorded below moderate sensitivity (i.e. 27.6%). Although this finding agrees with that of a previous study which recorded 30.2% (Otajewwo and Eriagbor, 2014), it is quite low when compared with 100% ofloxacin sensitivity recorded by Anigilaje and Bitto (2014) and more than 50% ofloxacin sensitivity reported by Okon *et al.* (2014). Sensitivity of streptomycin which was less than 30% was also very low when compared to 100% streptomycin sensitivity reported recently by an author (Habibu, 2014). Nitrofurantoin (22.4%) and ciprofloxacin (13.8%) sensitivities were quite low. The low sensitivities recorded for the fluoroquinolones (ofloxacin and ciprofloxacin) in this study was worrisome because these drugs are expensive and therefore are not readily accessible for abuse. In a similar study, Nakhjavani *et al.* (2007) reported that the widespread use of fluoroquinolones in medical centres is a possible cause of high level resistance to fluoroquinolones in UTI patients. Nwadioha *et al.* (2010) surprisingly, recorded a high sensitivity (80.0%) and above of UTI bacterial agents with ciprofloxacin. This is however open to re–evaluation through further studies by other workers.

The low sensitivity recorded for nitrofurantoin in this study is at variance with 100%, 97.6% and more than 50% sensitivities recorded by previous authors (Oluremi *et al.*, 2011; Haruna *et al.*, 2014; Alabi *et al.*,

2014). The least sensitive antibiotics were augmentin and nalidixic acid. The low sensitivity of nalidixic acid is surprising because it is a drug that is not commonly and routinely used in medical practice. The less than 4% sensitivity of augmentin is very low and disturbing in view of its usefulness in the treatment of UTI's and other diseases. The almost total resistance of augmentin is alarming as it may have lost its value in the treatment of UTI (Oluremi *et al.*, 2011).

Four out of the seven isolated uropathogens were resistant to more than three drugs and hence they were all multi–drug resistant (MDR). In this study, *Staph aureus*, *kleb.aerogenes*, *Coliform* organisms and *Proteus* spp were resistant to four drugs, seven drugs, five drugs and seven drugs respectively (Table 5). Multi drug resistance of *Staph aureus* may be due to beta–lactamase (penicillinase) encoding genes it carries on its plasmids as well as other extracellular and intracellular factors the organism elaborates. Besides, multi–drug resistant *Staph aureus* strains have been widely reported in some studies (Abubakar, 2009; Aiyegoro *et al.*, 2007; Gales *et al.*, 2000). Multi–drug resistance of *Kleb. aerogenes* and *Proteus* spp are documented. The prevalence of multiple antibiotic resistant strains in this study is a possible indication that very large population of bacterial isolates has been exposed to several antibiotics (Oluremi *et al.*, 2011).

V. CONCLUSION

A prevalence UTI rate of 59.2% in this study (though not very high), indicates that UTI may be a health problem among students (who are mainly residential) of Lighthouse Polytechnic, Evbuobanosa, near Benin City. The findings suggest that most (if not all) the uropathogens isolated in this study are excretable through urine to the environment of the study area, Ewobanosa and its environs. Besides, the students should be advised to step up their personal hygiene in their hostels and immediate environment. Where signs and symptoms of UTI are noticed notwithstanding the above, healthcare providers of the Polytechnic may administer doses of gentamicin only or ofloxacin or perhaps streptomycin for therapy. A prompt therapeutic intervention in this regard, will prevent asymptomatic UTI cases becoming symptomatic with the accompanying renal damage.

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