

Prevalence of Drug Resistant Uropathogenic Escherichia coli from Immunocompromised **Diabetic Patients Attending Selected Health Facilities in Benue State**

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Abstract

Escherichia coli is the commonest bacterial uropathogen of UTIs, the commonest infections in immunocompromised diabetic patients. Better understanding of their main resistance mechanisms to commonly used antibacterial agents will help to reduce the burden of this infection. The prevalence of drug resistant uropathogenic Escherichia coli isolates from immunocompromised diabetic patients attending selected health facilities in Benue State was investigated. Two hundred and ninety-six midstream urine samples were collected for both study and control diabetic patients. Bacterial isolation was done using semi-quantitative method. Drug resistant *Escherichia coli* were identified as multidrug resistant (MDR), extensive drug resistant (XDR) and pan-drug resistant organisms (PDR). Statistical significance was considered at p < 0.05. Results showed sixty-two and twenty-five *E. coli* isolates from the study and control subjects with overall prevalence of 20.9% and 8.4% respectively. The isolates were highly resistant to penicillin (ampicillin), monobactam (aztreonam), older quinolone (nalidixic acid) whereas the majority of them showed high susceptibility to aminoglycoside (streptomycin), cephalosporin (cefotaxime) and carbapenem (imipenem). None showed complete susceptibility to all the tested antibiotics. Twenty-five E. coli were identified in this MDR, eight, XDR while 5 were PDR. High numbers of drug resistant E. coli isolates were identified in the study group of which 25 were MDR, 8 XDR while 5 were PDR isolates. High prevalence of UTI and drug resistant isolates occur in diabetic patients with hyperglycemic condition.

Keywords

Urinary Tract Infection, Immunocompromised, Diabetic Mellitus, Antibiotic

Resistance, Escherichia coli

1. Introduction

A healthy immune system can defeat invading disease-causing pathogens as well as cancer cells. However, when the number of pathogens is overwhelming or there is any defect in the immune system such as defects in antibody production, complement and neutrophils, infections can occur [1]. Individuals who are immunocompromised have reduced ability to fight infections and other diseases [2]. This inability to fight infection may be caused by diseases or conditions such as HIV/AIDS, cancer, diabetes mellitus and so on [3].

Diabetic patients' immune response is disrupted as a result of the prevailing hyperglycemic condition. In addition to the risk of natural barrier damage due to neuropathy, diabetes especially type 2 can also affect cellular immunity [4]. Diabetes mellitus results from the inability of β -cells in the pancreas to produce sufficient insulin or the insulin produced cannot be properly utilized [5]. According to the American Diabetes Association (ADA), infections are an important issue for individuals with diabetes due to the failure of the immune system to fight off invading pathogens [6].

People living with diabetes are more susceptible to infections than people without diabetes [7]. One of the commonest infections in these immunocompromised diabetic patients is UTI [8]. This results from decreased cellular responses and poor pathogen clearance that are very common in these patients [9]. Urinary tract infection is a major cause of death in these groups of immuno-compromised patients [9].

Studies have reported *E. coli* to be the commonest bacterial uropathogens of UTIs [10]. This is due to the presence of a number of virulence [11]. and because *E. coli* is a common flora in the gastrointestinal tract, rectal and vagina area from where it could ascend to the urinary tract.

Bacteria that are resistant to multiple antimicrobial agents are called multi-drug resistant (MDR), those with extensive drug resistant (XDR) or totally drug resistant (TDR) are sometimes called "superbugs" [12]. The study was undertaken to determine the prevalence of drug resistance uropathogenic *E. coli* from diabetic patients with hyperglycemic condition attending three selected health facilities in Benue State.

2. Materials and Methods

2.1. Study Design, Area and Population

This study used a cross-sectional survey design carried out from November, 2021 to April, 2022. It includes patients seeking medical services at Federal Medical Centers, Makurdi, General Hospital at Vandeikya and Otukpo, Benue State. It includes both in-patients and out-patients showing symptoms of urinary tract infections. Patients selected for the study and control group were diabetic patients with blood glucose levels $\geq 200 \text{ mg/100ml}$ and between 126 mg/100ml and 200 mg/100ml respectively, who were not on any antibiotic therapy for the past two weeks before the study period. Patients with extreme age below 18 and above 60 years were excluded.

A total of 296 diabetes mellitus patients with blood glucose levels \geq 11.1 mmol/L (200 mg/dl) were the test and other 296 diabetes patients with blood glucose levels between 7.0 - 11.1 mmol/L (126 mg/dl and 200 mg/dl) were the control. Sample size was determined using the simple population proportion formula described by Charan and Biswas [13].

$$n = \frac{Z^2 p \left(1 - p\right)}{d^2}$$

where *n* = the study sample size,

Z = Z score for 95% confidence interval (degree of confidence) = 1.96

P = prevalence of UTI in the study area (26.03%) [14].

d = margin of error (of setting a significant level of 0.05) = 5% (0.05)

The participants for both studies were selected using systematic random sampling [15].

2.2. Ethical Clearance

Ethical approval was obtained from the Health Research Ethical Committee at Federal Medical Centre (FMH/FMC/MED/108/VL.1) and Benue State Hospitals Management Board, Makurdi (HMB/OFF/215/VOL.II/455). Permission was also obtained from the Principal Medical Officer of the respective health care facilities. Informed written consent was obtained from all the study participants. Questionnaire was also given to each patient to obtain demographic and clinical information.

2.3. Bacterial Isolation

Midstream urine samples from patients were cultured on Cysteine Lactose Electrolyte Deficient (CLED) (Oxoid, CM 0398), MacConkey and Blood Agar (Oxoid) to determine the bacterial uropathogens present in each urine sample. Each urine sample was aseptically inoculated onto the culture media and incubated aerobically at 37°C for 24 hours.

The media which show no bacteria growth after 24-hours incubation were further incubated up to 48 hours before reporting no bacterial growth. The numbers of isolated bacterial colonies were. Only urine samples that gave single pure with colony $\geq 10^5$ CFU/ml were taken as significant growth [16] [17].

2.4. Identification of Bacterial Uropathogens

E. coli isolates were identified following standard bacteriological procedures as described by Cheesbrough [16], Kolawale *et al.* [18] and through molecular approaches as reported by Adeoti [19].

2.5. Antibiotic Susceptibility Testing (AST)

The isolates were subjected to in-vitro antibiotic susceptibility testing using the modified Kirby-Bauer disc diffusion method according to the guidelines recommended by the CLSI [20]. Antibiotics used in this study were Mast discs product. They included ampicillin (AMP) 10 μ g, cefoxitin (FOX) 30 μ g, cefotaxime, (CTX) 30 μ g, ceftriaxone (CRO) 30 μ g, imipenem (IMI) 10 μ g, aztreonam (ATM) 30 μ g, augmentin (AUG) (amoxicillin/clavulinic acid 20/10 μ g), genticn (GM) 10 μ g, streptomycin (S) 10 μ g, chloramphenicol (C) 30 μ g, cotrimoxazole (trimethoprim-sulfamethoxazole 1.25/23.75) (COT) 25 μ g, nalidixic acid (NA) 30 μ g, ciprofloxacin (CIP) 5 μ g, ofloxacin (OFX) 5 μ g, levofloxacin (LEV) 5 μ g, nitrofurantoin (NI) 300 μ g and tetracycline (T) 30 μ g.

Antibiotic susceptibility testing was done on a sterile Muller Hinton Agar (Oxoid, CM00339) plate using an equivalent of 0.5% McFarland standard (10⁸ CFU/mL), incubated and interpreted as recommended by the CLSI [20].

2.6. Identification of Drug Resistant Isolates

All drug resistant uropathogenic *E. coli* isolates were identified as multi-drug resistant isolates (MDR), extensive drug resistant isolates (XDR) and pan-drug resistant isolates following the combined guidelines of European Centre for Disease Prevention and Control (ECDC) and the Centers for Disease Control and Prevention (CDC). The *E. coli* were identified as MDR when resistant to 1 agent in 3 antibiotic classes. They were extensively drug resistant organisms (XDR) when non-susceptibility to at least one agent in all but two or fewer antibacterial categories. Pan-drug resistant organisms (PDR) were defined as non-susceptibility to all agents in all antibiotic classes [21].

2.7. Statistical Analysis

Data generated were analyzed using SPSS software version 20.0 for Windows. Chi-square test was used to assess and compare relationships between variables. In all the comparisons, *p-value* less than 0.05 were taken as statistically significant levels. Sample size was determined using the formula [12]

3. Results

3.1. Prevalence of Urinary Tract Infections

The overall prevalence of UTIs in this study was 20.9%. Female diabetic patients had the highest occurrence of UTI (54.8%) (Table 1). Age group 46 and 50 years had the highest cases 21 (33.9%), followed by age group 51 and 55, 16 (25.8%), age group 41 and 45 years 9 (14.5%), age groups 36 - 40 and 56 - 60 years had 8 (12.9%) each, and all participants in the age groups 18 - 25, 26 - 30 and 31 - 35 had no positive UTI case. The female participants in the age group 46 and 50 years had the highest occurrence of UTIs (12), followed by the males in the age groups 46 - 50 and 51 - 55 years each having 9 while age group 41 and 45 of the male participants had the lowest with only a patient (Figure 1).

 Table 1. Socio-demographic characteristics and prevalence of UTI in the studied DM patients.

Demographic characteristics	Total (%)	Negative UTI (%)	Positive UTI (%)	X²	P-value	
Age group (years)						
18 - 25	0	0	0			
26 - 30	1 (0.3)	1 (0.4)	0			
31 - 35	5 (1.7)	5 (2.1)	0			
36 - 40	55 (18.6)	47 (20.1)	8 (12.9)	7.438	0.282	
41 - 45	57 (19.3)	48 (20.5)	9 (14.5)			
46 - 50	78 (26.4)	57 (24.4)	21 (33.9)			
51 - 55	57 (19.3)	41 (17.5)	16 (25.8)			
56 - 60	43 (14.5)	35 (15.0)	8 (12.9)			
Gender						
Male	110 (37.2)	82 (35.0)	28 (45.2)	2.149	0.143	
Female	186 (62.8)	152 (65.0)	34 (54.8)			
Aarital status						
Single	6 (2.0)	5 (2.1)	1 (1.6)			
Married	272 (91.9)	214 (91.5)	58 (93.5)	0.466	0.926	
Divorce	1 (0.3)	1 (0.4)	0			
Widowed	17 (5.7)	14 (6.0)	3 (4.8)			
esidence						
Urban	169 (57.1)	132 (56.4)	37 (59.7)	0.214	0.644	
Rural	127 (42.9)	102 (43.6)	25 (40.3)			
ducational status						
Illiterate	26 (8.8)	19 (8.1)	7 (11.3)			
Literate Primary	35 (11.8)	25 (10.7)	10 (16.1)	2.336	0.506	
Secondary	79 (26.7)	63 (26.9)	16 (25.8)			
Tertiary	156 (52.7)	127 (54.3)	29 (26.8)			
Occupational status						
Civil servant	111 (37.5)	96 (41.0)	15 (24.2)			
Business	53 (17.9)	36 (17.5)	12 (19.4)			
Student	1 (0.3)	0	1 (1.6)	9.706	0.046	
Farmer	69 (23.3)	52 (22.2)	17 (27.4)			
House wife	62 (20.9)	45 (19.2)	17 (27.4)			
Others	0	0	0			
Fotal	296 (100.0)	234 (79.1)	62 (20.9)			

Key: χ^2 : Chi-square %: percent, UTI: Urinary tract infection and DM: Diabetic mellitus.

The married participants had the highest occurrence of UTI 58 (93.5%), followed by the widowed 3 (0.5%) and single. Participants who are more educated gave the highest occurrence of UTI 29 (26.8%), while the illiterates showed the least occurrence of UTI 7 (11.3%). Urban dwellers patients had UTI of 37 (59.7%) while 25 (40.3%) lived in rural areas. Farmers and house wives had the highest occurrence of UTIs 17 (27.4%), followed by civil servants 15 (24.2%), business owners 12 (19.4%) and students 1 (1.6%). There was significant association between occurrence of UTIs and occupation (p = 0.046) of the enrolled diabetic patients (**Table 1**).

3.2. Bacterial Profile of UTI in Immunocompromised Diabetic Patients

Out of the total number of urine samples analyzed, 62 (20.9%) and 25 (8.4%) of them were positive for bacterial culture for the study population and control subject respectively (**Table 2**). From the study, 14, 20 and 28 *E. coli* isolates were identified from Vandeikya, Otukpo and Makurdi location respectively.

3.3. Antibiotic Susceptibility Pattern of Uropathogenic *E. coli* Isolates

Antibiotic susceptibility test was performed for all the *E. coli* isolates using 17 different antibiotics. From Vandeikya location, the isolates showed high sensitivity to cefotaxime 12 (85.7%), followed by imipenem 10 (71.4%), chloramphenicol 9 (64.3%), nitrofurantoin 9 (64.3%), ceftriaxone 9 (64.3%), moderate susceptibility to genticin, streptomycin, tetracycline and ciprofloxacin. There was no sensitivity to ampicillin, aztreonam and nalidixic acid.

Escherichia coli isolates from the health facilities at Otukpo location demonstrated

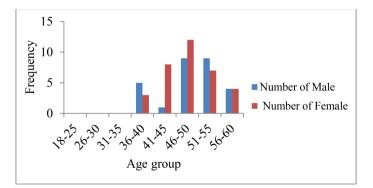


Figure 1. Age/sex specific prevalence of UTIs from diabetic patients in the study area.

Bacterial profiles		Fre	quency	Percentage (%)		
		Test	Control	Test	Control	
Bacterial isolates	Escherichia coli	62	25	20.9	8.4	
	Negative	234	271	79.1	91.6	

Table 2. Bacterial profiles in the diabetes mellitus patients.

high susceptibility to streptomycin 12 (60%), cefotaxime 11 (55%) and imipenem 10 (50%). High resistance was exhibited to ampicillin 16 (80%), aztreonam 16 (80%), levofloxacin 16 (80%), nalidixic acid 14 (70%), ofloxacin 14 (70%), ciprofloxacin 13 (65%), ceftriaxone 12 (60%) and tetracycline 11 (55%). *Escherichia coli* from Makurdi were most susceptible to streptomycin (75%) followed by cefotaxime (64.3%), and no susceptibility to ampicillin and aztreonam. Similarly, the rate of resistance ranged from ampicillin 89.3%, aztreonam 78.6%, nalidixic acid 71.4%, cefoxitin 57.1%, ofloxacin 53.6%, levofloxacin 53.6%, and tetracycline 50% while other antibiotics yielded less than 50% resistance pattern (**Table 3**).

3.4. Percentage of Isolated *E. coli* Showing Resistance to the Different Antibiotics Used in This Study

The percentile distribution (Figure 2) showed the prevalence of resistance of

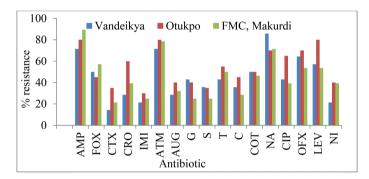
Mode of action	Antibiotic class	Antibiotics	<i>Escherichia coli</i> , n = 14 (Vandeikya)		<i>Escherichia coli</i> , n = 20 (Otukpo)			<i>Escherichia coli</i> , n = 28 (Makurdi)			
			S	I	R	S	Ι	R	S	Ι	R
Cell wall synthesis inhibitors	Penicillins	AMP	0 (0)	4 (28.6)	10 (71.4)	0 (0)	4 (20)	16 (80)	0 (0)	3 (10.7)	25 (89.3)
	Cephalosporins	FOX	4 (28.6)	3 (21.4)	7 (50)	7 (35)	4 (20)	9 (45)	10 (35.7)	2 (7.1)	16 (57.1)
		CTX	12 (85.7)	0 (0)	2 (14.3)	11 (55)	2 (10)	7 (35)	18 (64.3)	4 (14.3)	6 (21.4)
		CRO	9 (64.3)	1 (7.1)	4 (28.6)	5 (25)	3 (15)	12 (60)	10 (35.7)	7 (25)	11 (39.3)
	Carbapenem	IMI	10 (71.4)	1 (7.1)	3 (21.4)	10 (50)	4 (20)	6 (30)	17 (60.7)	4 (14.3)	7 (25)
	Monobactam	ATM	0 (0)	4 (28.6)	10 (71.4)	0 (0)	4 (20)	16 (80)	0 (0)	6 (21.4)	22 (78.6)
	eta-lactams inhibitors	AUG	8 (57.1)	2 (14.3)	4 (28.6)	9 (45)	3 (15)	8 (40)	15 (53.6)	4 (14.3)	9 (32.1)
Protein synthesis inhibitors	Aminogylcoside	G	7 (50)	1 (7.1)	6 (42.9)	9 (45)	3 (15)	8 (40)	17 (60.7)	4 (14.3)	7 (25)
		S	7 (50)	2 (14.3)	5 (35.7)	12 (60)	1 (5)	7 (35)	21 (75)	0 (0)	7 (25)
	Tetracycline	Т	7 (50)	1 (7.1)	6 (42.9)	7 (35)	2 (10)	11 (55)	13 (46.4)	1 (3.6)	14 (50)
	Phenicol	С	9 (64.3)	0 (0)	5 (35.7)	8 (40)	3 (15)	9 (45)	12 (42.9)	8 (28.6)	8 (28.6)
Folate pathway inhibitors	Sulfonamides	СОТ	5 (35.7)	2 (14.3)	7 (50)	8 (40)	2 (10)	10 (50)	13 (46.4)	2 (7.1)	13 (46.4)
Nucleic acid inhibitors		NA	0 (0)	2 (14.3)	12 (85.7)	4 (20)	2 (10)	14 (70)	4 (14.3)	4 (14.3)	20 (71.4)
		CIP	7 (50)	1 (7.1)	6 (42.9)	5 (25)	2 (10)	13 (65)	13 (46.4)	4 (14.3)	11 (39.3)
	Quinolones	OFX	3 (21.4)	2 (14.3)	9 (64.3)	5 (25)	1 (5)	14 (70)	12 (42.9)	1 (3.6)	15 (53.6)
		LEV	3 (21.4)	3 (21.4)	8 (57.1)	2 (10)	2 (10)	16 (80)	9 (32.1)	4 (14.3)	15 (53.6)
	Nitrofurans	NI	9 (64.3)	2 (14.3)	3 (21.4)	8 (40)	4 (20)	8 (40)	14 (50)	3 (10.7)	11 (39.3)

Table 3. Antibiotic susceptibility pattern of bacterial uropathogens from diabetes mellitus patients in the study area.

Key: AMP: Ampicillin, FOX: Cefoxitin, CTX: Cefotaxime, CRO: Ceftriaxone, IMI: Imipenem, ATM: Aztreonam, AUG: Augmentin, GM: Genticin, S: Streptomycin, T: Tetracycline, C: Chloramphenicol, ATH: Azithromycin, E: Erythromycin, COT: Cotrimoxazole, NA: Nalidixic acid, CIP: Ciprofloxacin, OFX: Ofloxacin, LEV: Levofloxacin, NI: Nitrofurantoin, S: Sensitive, I: Intermediate, R: Resistant. *E. coli* isolates to the different classes of antibiotics used. The isolated *E. coli* gave the highest percentage resistance to ampicillin (89.3%) in Makurdi, followed by nalidixic acid (85.7%) in Vandeikya, ampicillin, aztreonam and levofloxacin from Otukpo 80%. *Escherichia coli* demonstrated the lowest percentage resistance to cefotaxime (14.3%) in Vandeikya, followed by imipenem (21.4%) in Vandeikya and cefotaxime (21.4%) in FMC, Makurdi (**Figure 2**).

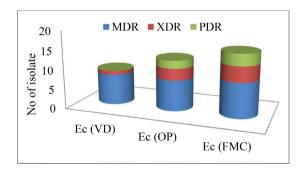
3.5. Occurrence of MDR, XDR and PDR Isolates among Uropathogenic *E. coli* Isolates

Out of the thirty-eight (38) drug resistant *E. coli* isolates detected from study group (diabetic patients with hyperglycemic condition), 25 were MDR strains, 8 XDR strains and 5 PDR strains. The 25 MDR isolates were distributed as follows: 8 (21.0%) isolates from Vandeikya, 8 (21.0%) isolates from Otukpo and 9 (23.7%) from Makurdi. One (2.6%) XDR isolates were from Vandeikya, 3 (7.9%) from Otukpo and 4 (10.6%) from Makurdi while 2 (5.3%) and 3 (7.9%) were positive PDR strains from Otukpo and Makurdi respectively (**Figure 3**).



Key: AMP: Ampicillin, FOX: Cefoxitin, CTX: Cefotaxime, CRO: Ceftriaxone, IMI: Imipenem, ATM: Aztreonam, AUG: Augmentin, GM: Genticin, S: Streptomycin, T: Tetracycline, C: Chloramphenicol, ATH: Azithromycin, E: Erythromycin, COT: Cotrimoxazole, NA: Nalidixic acid, CIP: Ciprofloxacin, OFX: Ofloxacin, LEV: Levofloxacin and NI: Nitrofurantoin.

Figure 2. Percentage of uropathogenic *E. coli* showing resistance to the different antibiotics used in the three different locations.



Key: VD: Vandeikya, OP: Otukpo, FMC: Federal Medical Centre, Makurdi, Ec: *E. coli*, MDR: multi-drug resistance, XDR: Extensive drug resistance, PDR: Pan-drug resistance.

Figure 3. Occurrence of MDR, XDR and PDR isolates in the isolated uropathogenic *E. coli* from diabetic patients in the study area.

4. Discussion

The present study was undertaken to ascertain the prevalence of UTI and associated antibiotic resistance mechanisms in these patients. The goal is to determine the most effective approach towards management of UTI to reduce the burden of this infection in this group of immunocompromised patients. The study showed the prevalence of UTI among immunocompromised diabetic patients (blood glucose levels \geq 11.1 mmol/L) to be higher (20.9%) than the diabetic patients with blood glucose levels between 7.0 mmol/L and 11.1 mmol/L (8.4%).

The prevalence of UTI recorded for diabetes mellitus study population is comparable to the reported prevalence in other studies such as 22.6% recorded in a study in Addis Ababa [22]. and 19.5% among diabetic in Khartoum, Sudan [23]. It was consistent with studies of Forson *et al.* [24] who reported 9.2%, 10.1% recorded at Derbe Tabor General Hospital, Northwest Ethiopia [25] and 10.5% in Southern Ethiopia [26]. In contrast, much higher prevalence of 35.3% was reported in Iraq [27] 40.2% in Malaysia [28] and 54.7% in Nepal [29].

The prevalence of UTI based on age group showed that age group 46 and 50 had the highest prevalence (33.9%) while age groups 18 and 25, 26 and 30, 31 and 35 had no recorded positive UTI cases. No significant association was observed in relation to age group of the participants. Study by Al-Rubeaan *et al.* [30] carried out in Saudi reported similar findings while another study in Sweden by Hamdan *et al.* [23] reported significant association between age groups and positive UTI cases among diabetic patients. Some studies have reported that as age increases, UTI cases in diabetic patients also increased probably due to decreasing state of immunity of the patients as a result of the hyperglycemic condition [31].

In this study, UTI occurs only in the age groups 36 - 40, 41 - 45, 46 - 50, 51 - 55 and 56 - 60 with age groups 46 - 50 and 51 - 55 highest. This observation was similar to the studies of Njunda *et al.* [32] in Cameroon who also recorded high prevalence among age group 41 - 60 years. This was attributed to increased sexual activity in this age group [33]. Contrary to this, Walelgn *et al.* [34] in South Wollo, Northeast Ethiopia, Chita *et al.* [35] in the UK and Forson *et al.* [24] in Ghana obtained higher occurrence of UTIs among diabetic patients in the age group 61 - 70 years.

Farmers and house wives each had prevalence of 27.4% UTI being the highest. There was significant association between occurrence of urinary tract infection and occupation (p = 0.046) of the diabetic patients.

All the *E. coli* isolates in this study showed resistance to at least one antibiotic and none showed complete susceptibility to all the antibiotics tested. *Escherichia coli* from the location at Vandeikya were sensitive to β -lactams (cefotaxime and imipenem), chloramphenicol and nitrofurantoin. Slightly higher sensitivity has been reported in a study among diabetic patients in Ethiopia to nitrofurantoin 95%, cefotaxime 85% and ceftriaxone 85%. [22]. Another study in Ethiopia also reported high susceptibility (100%) to nitrofurantoin, genticin and meropenem [25]. This study also reported high resistance to ampicillin 71.4%, ofloxacin 64.3%, cefoxitin 50% and cotrimoxazole 50%. Worku *et al.* [25] reported similar high resistance pattern to ampicillin 100%, cefoxitin 100%, augmentin 92.9% and cotrimoxazole 61.1% in Ethiopia.

The results of antibiotic susceptibility pattern from Otukpo revealed *E. coli* to be moderately susceptible to streptomycin 60%, cefotaxime 55% and imipenem 50% while at the same time showing moderate to high resistance to cotrimox-azole 50%, tetracycline 55%, ceftriaxone 60%, ciprofloxacin 65%, ofloxacin 70%, nalidixic acid 70%, levofloxacin 80%, aztreonam 80%, ampicillin 80%. This was contrary to a study in India [36] who reported complete sensitivity (100%) to imipenem but comparable resistant pattern has been observed in Ethiopia to na-lidixic acid 70% and cotrimoxazole 64.7% [26].

The isolated *E. coli* from Makurdi exhibited moderate to slightly high susceptibility pattern to nitrofurantoin, augmentin, genticin, imipenem and streptomycin which partly agreed with a study in Ethiopia where high sensitivity was demonstrated to nitrofurantoin 95.5% and genticin 73.3% [37]. Another study in Ethiopia also reported 100% sensitivity to imipenem and nitrofurantoin [25]. *Escherichia coli* also showed moderate to high resistance to tetracycline, levofloxacin, ofloxacin, cefoxitin, nalidixic acid, aztreonam and ampicillin in this study. A similar high resistance pattern was reported in a study conducted among diabetic patients in Jos, Nigeria where resistance to ofloxacin was 70% and ampicillin 90% [38].

The antibiotic susceptibility test results showed streptomycin and cefotaxime as the most effective antibacterial against *E. coli* isolates, hence could be used as therapeutic agents in the empirical treatment of UTIs in diabetic patients in the study area. Also, most resistance were expressed to antibacterial agents like ampicillin, cotrimoxazole, tetracycline and the quinolones which might be due to their ready availability, being agents commonly used for self-medication. These antibacterial agents are also commonly used agents for treatment of other bacterial infections.

This study showed high resistance to older quinolone, nalidixic acid than to newer quinolones like ciprofloxacin, ofloxacin and levofloxacin. Findings by Woldemariam *et al.* [22] reported similar observations in their study. Quinolone antibiotics like ciprofloxacin, ofloxacin and levofloxacin are commonly prescribed antibiotics in this country for the treatment of UTIs in most diabetic patients. This could be one of the major reasons the bacteria isolate from diabetic patients in this study are resistant to these antibiotics.

Relatively low rates of resistance were expressed to the 2 aminoglycosides, genticin and streptomycin used in this study. Studies in Nigeria [39], Ethiopia [40] [41] have all reported comparable antibiogram pattern to these antibiotics. The study also demonstrated the low resistance to cefotaxime which disagrees with a study in Iran [42] where the *E. coli* isolates exhibited the highest resistance to cefotaxime. This low resistance exhibited to the third-generation ce-

phalosporin could be because they are not among the commonly used antibiotics for the treatment of UTIs in most communities in the study area. It could also be as a result of their non-availability and the high cost of purchasing them.

This study showed 38 drug resistant *Escherichia coli* isolates among the study diabetic patients consisting of 25 MDR, 8 XDR and 5 PDR isolates. This implies danger with respect to the global fight to reduce antibiotic resistance by organisms. The findings are comparable to that of Aly and Balkhy [43] who reported similar isolates of *E. coli* as the most common MDR isolates. It also correlates well with another study conducted in a tertiary care hospital, Central India [44] where *E. coli* was reported to be most predominant MDR pathogens followed by *K. pneumoniae*. Studies by Abbas *et al.* [45] from Pakistan and Effah *et al.* [46] reported contrary findings of *K. pneumoniae* isolates as most common XDR strains.

The high number of MDR, XDR and PDR isolates in this study could be a possible reflection of the increasing antibiotic resistance among the uropathogens. The health facilities used for this study are all referral centers that attend to patients who may have been treated with one antibiotic or the other from smaller health facilities scattered all around the different study health centres.

One major limitation of the study is that it was conducted in Benue State, north-central Nigeria instead of all the states of the federation. The selected immunocompromised patients were diabetes mellitus patients with blood glucose levels $\geq 200 \text{ mg/dl}$ (11.1 mmol/L). The study was restricted to this group of immunocompromised patients seeking medical attention and those on hospital admission in Federal Medical Centre (FMC), Makurdi, General Hospitals at Otukpo and Vandeikya all within the State. The study focused only on one bacterial pathogen (*E. coli*) excluding other bacterial uropathogens, and other pathogens such as fungi and protozoan which might have added more value to the research. Other useful clinical and laboratory tests like HbA1c estimations, cytokine profile and possibly white blood cell counts were not included. The study did not provide information about extended spectrum β -lactamase (ESBL) production of the isolated *E. coli*. Besides, the study did not conduct detailed study on the isolates from the control group just as it was done on those from the study diabetic group.

5. Conclusion

High levels of resistance were expressed to ampicillin, aztreonam, nalidixic acid. Multidrug resistant, XDR and PDR urinary *E. coli* isolates were also high. However, the isolates were susceptible to streptomycin and cefotaxime thereby making these drugs useful for UTI treatment in immunocompromised diabetic patients.

Conflicts of Interest

The authors declare no conflicts of interest regarding the publication of this paper.

References

- [1] Perdue, S.S. and Humphrey, J.H. (2016) Immune System. Encyclopedia Britanica. https://www.britanica.com/science/immunesystem
- Singh, D. and Bonomo, R.A. (2016) Infections in Cancer Patients. In: Hoag, J.B., Ed., Oncologic Critical Care, IntechOpen, London, 117-143. https://doi.org/10.5772/64372
- [3] Zhang, S., Yang, H., Rehman, M.U., Yang, K., Dong, M., Yang, J., Wang, M., Jia, R., Chen, S., Liu, M., Zhu, D., Zhao, X., Yang, Q., Wu, Y., Zhang, L., Liu, Y., Yu, Y., Tian, B., Pan, L., Chen, X. and Cheng, A. (2019) Class 1 Integrons as Predominant Carriers in *Escherichia coli* Isolates from Waterfowls in Hainan, China. *Ecotoxicology and Environmental Safety*, **183**, Article ID: 109514. https://doi.org/10.1016/j.ecoenv.2019.109514
- [4] Wong, S.L. and Wagner, D.D. (2018) Peptidylarginine Deiminase 4: A Nuclear Button Triggering Neutrophil Extracellular Traps in Inflammatory Diseases and Aging. *Federation of American Societies Experimental Biology Journal*, **32**, 6258-6370. <u>https://doi.org/10.1096/fj.201800691R</u>
- [5] Dall, T.M., Yang, W, Gillespie, K., Mocarski, M., Bryne, E., Cintina, I., Beronja, K., Semilla, A.P., Iacobucci, W. and Hogan, P.F. (2019) The Economic Burden of Elevated Glucose Levels in 2017: Diagnosed and Undiagnosed Diabetes, Gestational Diabetes Mellitus, and Prediabetes. *Diabetes Care*, **42**, 1661-1668. https://doi.org/10.2337/dc18-1226
- [6] Lachmandas, E., Thiem, K., van den Heuvel, C., Hijmans, A., de Galan, B.E., Tack, C.J., Neetea, M.G., van Crevel, R. and van Diepen, J.A. (2018) Patients with Type 1 Diabetes Mellitus Have Impaired IL-1beta Production in Response to *Mycobacterium tuberculosis. European Journal of Clinical Microbiology and Infectious Disease*, **37**, 371-380. <u>https://doi.org/10.1007/s10096-017-3145-y</u>
- [7] Wallet, S.M., Puri, V. and Gibson, F.C. (2018) Linkage of Infection to Adverse Systemic Complications: Periodontal Disease, Toll-Like Receptors, and Other Pattern Recognition Systems. *Vaccines*, 6, 21. <u>https://doi.org/10.3390/vaccines6020021</u>
- Yamamoto, S. (206) Prevention and Treatment of Complicated Urinary Tract Infection. *European Urology Open Science*, 27, 186-189. <u>https://doi.org/10.1016/j.urols.2016.07.001</u>
- [9] Gołebiewska, J.E., Debska-Slizien, A. and Rutkowski, B. (2014) Urinary Tract Infections during the First Year after Renal Transplantation: One Center's Experience and a Review of the Literature. *Clinical Transplantation*, 28, 1263-1270. https://doi.org/10.1111/ctr.12465
- [10] Awean, G.Z., Salameh, K., Elmohamed, H., Alshmayt, H. and Omer, M.R.B. (2019) Prevalence of ESBL Urinary Tract Infection in Children. *Journal of Advances in Pediatrics and Child Health*, No. 1, 4-7. https://doi.org/10.29328/journal.japch.1001004
- [11] Nongrum, S., Thaledi, S., Singh, V.A., Narang, V.K., Mehta, S., Garg, R. and Sharma, T. (2016) Association of Uropathogens with Asymptomatic Urinary Tract Infection in Diabetes Mellitus Patients. *International Journal of Current Microbiological and Applied Sciences*, 5, 355-361. <u>https://doi.org/10.20546/ijcmas.2016.510.040</u>
- [12] Fentie, A., Wondimeneh, Y., Balcha, A., Amsalu, A. and Adanke, B.T. (2018) Bacterial Profile, Antibiotic Resistance Pattern and Associated Factors among Cancer Patients at University of Gondar Hospital, Northwest, Ethiopia. *Infection and Drug Resistance*, **11**, 2169-2178. <u>https://doi.org/10.2147/IDR.S183283</u>

- [13] Charan, J. and Biswas, T. (2013) How to Calculate Sample Size for Different Study Designs in Medical Research? *Indian Journal of Psychological Medicine*, 35, 121-126. <u>https://doi.org/10.4103/0253-7176.116232</u>
- [14] Mgbakogu, R.A. and Eledo, B.O. (2015) Studies on Urinary Tract Infection among Diabetes in Some Eastern States of Nigeria. *Advances in Life Sciences and Technol*ogy, 34, 42-47.
- [15] Yates, F. (1948) Systematic Sampling, Rothamsted Experimental Station, Communicated by Thornton, HG. 346-377.
- [16] Cheesbrough, M. (2007) Biochemical Testes to Identify Bacteria. In: Cheesbrough, M., Ed., *District Laboratory Practice in Tropical Countries*, 2nd Edition, Cambridge University Press, Cambridge, 50-55. <u>https://doi.org/10.1017/CBO9780511543470</u>
- [17] Public Health England, PHE (2014) UK Standards for Microbiology Investigations. *Investigation of Urine*, **B41**, 2.
- [18] Kolawole, A., Kolawole, O., Kandaki, O., Babatunde, S., Durowadw, K. and Kolalowole, C. (2009) Prevalence of Urinary Tract Infections (UTIs) among Patients Attending Dalhatu Araf Specialist Hospital, Lafia, Nasarawa State, Nigeria. *International Journal of Medicine and Medical Sciences*, 1, 163-167.
- [19] Adeoti, I. (2023) Determination of Resistance Genes in Gram-Negative Bacterial Uropathogens from Immunocompromised Patients Attending Selected Health Care Facilities in Benue State. PhD Thesis, Joseph Sarwuan Tarka University, Makurdi.
- [20] Clinical and Laboratory Standards Institute (CLSI) (2020) Performance Standards for Antimicrobial Susceptibility Testing. 30th Edition, CLSI Supplement M100, Clinical and Laboratory Standards Institute, Wayne, 32-131.
- [21] Magiorakos, A.P., Srinivasan, A., Carey, R.B., Carmeli, Y., Falagas, M.E. and Giske, C.G. (2012) Multidrug-Resistant, Extensively Drug-Resistant and Pandrug-Resistant Bacteria: An International Expert Proposal for Interim Standard Definitions for Acquired Resistance. *Clinical Microbiology and Infection*, 18, 268-281. https://doi.org/10.1111/j.1469-0691.2011.03570.x
- [22] Woldemariam, H.K., Geleta, D.A., Tulu, K.D., Aber, N.A., Legese, M.H., Fenta, G.M. and Ali, I. (2019) Common Uropathogens and Their Antibiotic Susceptibility Pattern among Diabetic Patients. *BioMed Central Infectious Diseases*, **19**, Article No. 43. <u>https://doi.org/10.1186/s12879-018-3669-5</u>
- [23] Hamdan, H.Z., Kubarr, E., Adam, A.M., Hassan, O.S., Suliman, O.S. and Adam, I. (2015) Urinary Tract Infections and Antibiotic Sensitivity among Diabetic Patients at Khartoum, Sudan. *Annals of Clinical Microbiology and Antimicrobials*, 4, 26-32. https://doi.org/10.1186/s12941-015-0082-4
- [24] Forson, A.O., Menkah, D.A., Quarchie, M.N., Dhikrullahi, S.B., Olu-Taiwo, M. and Codjoe, F.S. (2021) Bacterial Drug Resistance Patterns and Genetic Diversity of Bacteria-Associated Bacteriuria in Diabetic Patients in Ghana. *International Journal* of Infectious Disease Regions, 1, 142-149. https://doi.org/10.1016/j.ijregi.2021.10.007
- [25] Worku, G.Y., Alamneh, Y.B. and Abegaz, W.E. (2021) Prevalence of Bacterial Urinary Tract Infection and Antimicrobial Susceptibility Patterns among Diabetes Mellitus Patients Attending Zewditu Memorial Hospital, Addis Ababa, Ethiopia. *Infection and Drug Resistance*, 14, 1441-1454. https://doi.org/10.2147/IDR.S298176
- [26] Mohammed, A., Getnet, B., Teshager, L. and Daka, D. (2020) Urinary Pathogenic Bacterial Profile, Antibiogram of Isolates and Associated Risk Factors among Diabetic Patients in Hawassa Town, Southern Ethiopia: A Cross-Sectional Study. Urology and Nephrology Open Access Journal, 8, 84-91.

https://doi.org/10.15406/unoaj.2020.08.00282

- [27] Ahmed, A. (2013) Prevalence of Urinary Tract Infection in Diabetic Patients and Identification of the Causal Microorganisms. *Zanco Journal of Medical Sciences*, 17, 363-369. <u>https://doi.org/10.15218/zjms.2013.0016</u>
- Shah, M.A., Kassab, Y.W., Anwar, M.F., Al-Dahoul, K.H., Menon, S., Kaur, H.J., If-zaal, M., Muthanna, F.M. and Searan, W.M. (2019) Prevalence and Associated Factors of Urinary Tract Infections among Diabetic Patients. *Health Sciences Journal*, 13, Article No. 646. <u>https://doi.org/10.22207/JPAM.13.2.20</u>
- [29] Kumar, J.P., Baral, R. and Khanal, B. (2014) Prevalence of Uropathogens in Diabetic Patients and Their Susceptibility Pattern at a Tertiary Care Center in Nepal: A Retrospective Study. *International Journal of Biomedical Laboratory Science*, 3, 29-34.
- [30] Al-Rubeaan, K.A., Moharram, O., Al-Neqeb, D., Hassan, A. and Rafiullah, M.R.
 (2013) Prevalence of UTI and Risk Factors among Saudi Patients with Diabetics. World Journal of Urology, 31, 573-578. <u>https://doi.org/10.1007/s00345-012-0934-x</u>
- [31] Mageto, V.M., Mbuthia, O.W., Ngetsa, C.J., Moraa, D.O., Okoyo, E.O., Mathenge, S.G. and Njoroge, W.G. (2020) Evaluation of Socio-Demographic Factors among Diabetic Patients with UTIs in Kisii Referral Hospital, Kenya. *International Journal* of Microbiology, 2020, Article ID: 5053867. <u>https://doi.org/10.1155/2020/5053867</u>
- [32] Njunda, A.L., Assob, N.J.C., Nsagha, S.D., Nde, F.P., Kamga, F.H.L., Nkume, A.F. and Kwenti, T.E. (2012) Uropathogens from Diabetic Patients with Asymptomatic Bacteriuria and UTIs. *Scientific Journal of Microbiology*, 1, 141-146.
- [33] Clare, C.E., Juliet, A.O., Maureen, O.C., Romeo, O.J. and Ada, K.E. (2021) Urinary Tract Infections and Antimicrobial Sensitivity Patterns of Uropathogens Isolated from Diabetic and Non-Diabetic Patients Attending Some Hospitals in Awka. *American Journal of Microbiological Research*, 9, 83-91. https://doi.org/10.12691/ajmr-9-3-3
- [34] Walelgn, B., Abdu, M. and Kumar, P. (2021) The Occurrence of UTI and Determinant Factors among DM Patients at Dessie Referral, South Wollo, Northeast Ethiopia. SAGE Open Medicine, 9, 1-7. https://doi.org/10.1177/20503121211060614
- [35] Chita, T., Timar, B., Muntean, D., Badijoiu, L., Horhat, F., Hogea, E., Moldovan, R., Timar, R. and Licker, M. (2017) UTIs in Romanian Patients with Diabetes: Prevalence, Etiology and Risk Factors *Therapeutics and Clinical Risk Management*, 13, 1-7. <u>https://doi.org/10.2147/TCRM.S123226</u>
- [36] Nath, T., Das, S.K. and Hazra, S. (2021) Patterns of Uropathogens and Antibiotics Sensitivity in Diabetic Patients Attending to Out-Patients Department and Diabetes Clinic of a Teaching Hospital: A Cross Sectional Study. *Journal of Family Medicine* and Primary Care, 10, 3638-3840. <u>https://doi.org/10.4103/jfmpc.jfmpc_71_21</u>
- [37] Kibret, M. and Abera, B. (2014) Prevalence and Antibiogram of Bacterial Isolates from Urinary Tract Infections at Dessie Health Research Laboratory, Ethiopia. *Asian Pacific Journal of Tropical Biomedicine*, 4, 164-168. <u>https://doi.org/10.1016/S2221-1691(14)60226-4</u>
- [38] Anejo-Okopi, J.A., Okojokwu, O.J., Ramyil, S.M., Bakwet, P.B., Okechalu, J., Agada, G., Bassi, P.A. and Adeniyi, S.D. (2017) Bacterial and Antibiotic Susceptibility Pattern of UTI Isolated from Asymptomatic and Symptomatic Diabetic Patients Attending Tertiary Hospital in Jos, Nigeria. *Trends in Medicine*, **17**, 1-5. https://doi.org/10.15761/TiM.1000108
- [39] Ifeanyichukwu, I., Emmanuel, N., Chika, E., Anthonia, O., Esther, U.I., Ngozi, A. and Justina, N. (2013) Frequency and Antibiogram of Uropathogens Isolated from Urine Samples of HIV Infected Patients on ART. *American Journal of Bioscience*, 1, 1000 (2010) (20

50-53. https://doi.org/10.11648/j.ajbio.20130103.11

- [40] Getu, Y., Ali, I., Lema, T., Belay, H. and Yeshetela, B. (2017) Bactreiuria and Antimicrobial Susceptibility Pattern among HIV Patients Attending ALERT Center, Addis Ababa, Ethiopia. *American Journal of Health Research*, 5, 76-82. https://doi.org/10.11648/j.ajhr.20170503.14
- [41] Hantalo, A.H., Taassaw, K.H., Bisetegen, F.S. and Mulate, Y.W. (2020) Isolation and Antibiotic Susceptibility Pattern of Bacterial Uropathogens and Associated Factors among Adult People Living with HIV/AIDS Attending the HIV Center at Wolaita Sodo University Teaching Referral Hospital, South Ethiopia. *HIV/AID-Research* and Palliative Care, **12**, 799-808. <u>https://doi.org/10.2147/HIV.S244619</u>
- [42] Shams, S., Hashemi, A., Esmkhani, M., Kermani, S., Shams, E. and Piccirillo, A.
 (2018) Imipenem Resistance in Clinical *Escherichia coli* from Qom, Iran. *BioMed Central Research Note*, 11, 314-319. https://doi.org/10.1186/s13104-018-3406-6
- [43] Aly, M. and Balkhy, H.H. (2012) The Prevalence of Antimicrobial Resistance in Clinical Isolates from Gulf Corporation Council Countries. *Antimicrobial Resistance and Infection Control*, 1, Article No. 26. https://doi.org/10.1186/2047-2994-1-26
- [44] Basak, S., Singh, P. and Rajurkar, M. (2016) Multidrug Resistant and Extensively Drug Resistant Bacteria: A Study. *Journal of Pathogens*, 2016, Article ID: 406560. https://doi.org/10.1155/2016/4065603
- [45] Abbas, A.K. and Lichtman, A.H. (2003) Cellular and Molecular Immunology. 6th Edition, Saunders, Philadelphia.
- [46] Effah, C.Y., Sun, T., Liu, S. and Wu, Y. (2020) *Klebsiella pneumoniae*: An Increasing Threat to Public Health. *Annals of Clinical Microbiology and Antimicrobials*, 19, Article No. 1. <u>https://doi.org/10.1186/s12941-019-0343-8</u>