

Emergence of *bla* TEM resistance gene in ESBL-producing *Escherichia coli* clinical isolates from Health facilities in Makurdi, Benue State Nigeria

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ABSTRACT

The emergence of extended-spectrum- β -lactamases (ESBLs)-producing *Escherichia coli* represents a serious clinical concern in healthcare. β -lactamases produced by these strains of *E. coli* render ineffective cephalosporins and other β -lactam antibiotics used to treat infections caused by Gram-negative bacteria. We determined the presence of ESBL in 400 clinical isolates of *E. coli* isolated from various clinical specimens (urine, stool, blood, sputum, throat and wound swabs) from 216 female and 184 male patients with mean age of 28.1 ± 16.8 years (age range: 2 – 71 years), who were attending 6 selected health facilities in Makurdi, Benue State Nigeria. Antibiotic susceptibility test was carried out on the isolates using Kirby Bauer diffusion method. Presence of ESBLs was determined by the double disc synergy test (DDST). Specific primers were employed to characterize the ESBL gene using PCR. The isolates showed high level of resistance to all the antibiotics tested except mipenem. Highest resistance was to penicillin 392(98.0%) followed by ceftriaxone 385(96.3%). Out of the 400 isolates, 64 (16.0%) tested positive for ESBLs by DDST method, while PCR technique confirmed 47(11.8%) to harbour *bla* TEM genes. Isolates from blood specimens harboured highest percentage of ESBL genes 5(26.3%) and also plasmid-mediated *bla* TEM genes 5(26.3%), followed by wound swabs 9(17.3). The least percentage of plasmid-mediated *bla* TEM genes was carried by isolates from sputum specimens 1(8.3). Age group 45 to 58 years harboured the highest percentage of *bla* TEM genes 15(14.6%), while female patients, 27(12.5%) carried more *bla* TEM resistance genes than the male patients 20(10.9%). A prevalence of 11.8% ($n=47$) of *bla* TEM resistance gene has been reported for the present study. In view of multidrug resistant ESBL-producing *Escherichia coli* bacteria circulating in the study location, prescription of antibiotics, especially cephalosporins should be based on laboratory results of antibiotic susceptibility tests that are carried out along with ESBL detection. Infection prevention and control strategies should be stepped up in the health facilities under study.

Key words: Antibiotic resistance, *E. coli*, ESBL, Makurdi, TEM.

1.0 INTRODUCTION

The joy that heralded the introduction of the third-generation cephalosporins into clinical practice in the early 1980s was short-lived because reports of plasmid-encoded β -lactamases capable of hydrolyzing the extended-spectrum cephalosporins began to emerge in 1983 [10]. The gene encoding the β -lactamase showed a mutation of a single nucleotide compared to the gene encoding SHV-1. Other β -lactamases were soon discovered which were closely related to TEM-1 and TEM-2, but which had the ability to confer resistance to the extended-spectrum cephalosporins [24], hence these new β -lactamases were coined extended-spectrum β -lactamases (ESBLs). Infectious diseases that became curable at the advent of antibiotics chemotherapy are again becoming killers of patients of all ages especially in developing countries [4].

Extended spectrum beta lactamases (ESBLs) which are mainly produced by *Escherichia coli* (*E. coli*) and *K. pneumoniae* render these antibiotics ineffective when used to treat infections caused by ESBL producing organisms, consequently increasing cost of therapy, morbidity and mortality [5]. Resistance to antibiotics affect all countries, but worse in developing countries because they

have access to limited antibiotic options, engage in irrational antibiotic use, have poor drug quality and patronize inadequate health care systems [3]

Bla TEM resistance genes have been reported to confer resistance to ceftazidime, a third generation cephalosporin antibiotic [6]. The first report of *bla* TEM genes was made in Liverpool, England in 1982. Well over 100 TEM-type β -lactamases have been described, and majority are ESBLs. TEM-1 is able to hydrolyze ampicillin at a greater rate than carbenicillin, oxacillin, or cephalothin, and has negligible activity against extended-spectrum cephalosporins. It is inhibited by clavulanic acid.

Reports of *bla* TEM resistance genes circulating in health facilities abound. [12] reported a prevalence of 24.6% in Ouagadougou, Burkina Faso. A study by [13] reported a prevalence of 87.1% *bla* TEM in Thailand, and a prevalence of 81.3% *bla* TEM genes was reported by [19] in South-West Nigeria. The burden of ESBL and *bla* TEM resistance organisms in clinical practice is poorly documented in Nigeria. Therefore, we determined the presence of ESBLs, and particularly *bla* TEM resistance genes, its distribution in clinical isolates of various clinical specimens and patterns of susceptibility to commonly used antimicrobial agents and β -lactam antibiotics in selected health facilities in Makurdi, North-Central Nigeria.

2.0 MATERIALS AND METHODS

This cross-sectional study was carried out in the department of Microbiology, Benue State University Teaching Hospital, Makurdi. The study was approved by the ethics committee of Benue State University Teaching Hospital Makurdi. A total of 400 *Escherichia coli* clinical isolates from blood, urine, stool, sputum, wound and throat swabs were collected from 6 selected health facilities in Makurdi, Benue State Nigeria, namely Benue State University Teaching Hospital, Federal Medical Centre, General hospital, Bishop Murray Medical Centre, Family support clinic and Primary Health Care Centre. Presumptive *E. coli* isolates from these specimens were identified by standard microbiological methods including colonial morphology, Gram reaction, biochemical-TSI, indole and motility.

Antibiotic susceptibility testing was performed by the Kirby Bauer disk diffusion method, using Mueller-Hinton agar, according to BSAC guidelines. The following agents were tested-Penicillin, ceftazidime, cefotaxime, ceftriaxone, Cefuroxime, ciprofloxacin, chloramphenicol, gentamycin and mipenem. ATCC strain 25922 *E. coli* was used as control strain and the breakpoint was determined by measurement of zone of inhibition.

The presence of ESBL was detected by placing a 30 μ g ceftazidime disk (the best indicator for TEM) on the left of each plate and a 30 μ g cefotaxime disk was placed on the right; an amoxicillin-clavulanic acid disk (AMC) (20/10 μ g) (Oxoid Ltd was placed in the centre between the other discs. The disks were placed 25 mm apart, centre-to-centre. Following overnight incubation in air at 37^o C ESBL production was inferred when the zone of inhibition around the ceftazidime and cefotaxime disks was expanded by \geq 5 mm by the presence of clavulanic acid.

The DNA used for PCR molecular detection of *bla* TEM gene was extracted by the alkaline lysis method of Birnboim and Doly (1979). Specific primers for *bla* TEM, CTX and SHV were employed for the determination of *bla* TEM resistance genes. In multiplex PCR, 2 μ L whole cell lysate DNA of *E. coli* was used in 25 μ L PCR master mix and the specific primers. The PCR conditions were: Initial step of denaturation at 95^o C for 5 minutes followed by 35 cycles of denaturation at 95^o C for 1 minute, then annealing at 56^o C for 1.5 minutes, extension at 95^o C for 1 minute, and then final extension at 95^o C for 10 minutes. The PCR products were separated electrophoretically using 1.5% agarose gel and stained with ethidium bromide. *Bla* TEM gene was identified by comparing the separated PCR products with the marker.

3.0 STATISTICAL ANALYSIS.

The data obtained were analyzed with SPSS version 20.0 applying Chi square at 95% to test degree of association.

4.0 RESULTS.

The isolates exhibited high level of resistance to all antimicrobial agents tested. Highest resistance was to penicillin (98.0%, n=392), followed by ceftriaxone (96.3%; n= 385). The isolates were almost all susceptible to mipenem, a carbapenem (Table 1). Out of 400 isolates tested, 64 (16.0%) were positive for ESBL production. Isolates from blood specimens harboured the highest percentage 5(26.3%), of ESBLs followed by wound specimen isolates 9(17.3%). Highest percentage of *bla* TEM resistance genes were also harboured by blood specimen isolates (Table 3). Demographic data showed that higher number of isolates were gotten from female population 12.5% (n = 27) compared with the male population 10.9% (n= 20). Age group of 45 to 58 (14.8%; n = 8) years were prevalent (Table 5). General hospital Makurdi, 15.1% (n = 8) which is a secondary care hospital harboured higher percentage of plasmid-mediated *bla* TEM resistance genes followed by Benue State University Teaching Hospital, 13.6% (n = 20) a tertiary care health facility. The primary Health Care centre 3.3% (n = 1) harboured the least percentage of *bla* TEM resistance genes (Table 6).

Table 1: Susceptibility Profile of Isolated *Escherichia coli* to Antibiotics (N=400)

Antibiotics	Disc content	Resistance (%)	Susceptible (%)
Penicillin	10µg	392(98.0)	8(2.0)
Ceftriaxone	30 µg	385(96.3)	15(3.8)
Cefuroxime	30 µg	376(94.0)	24(6.0)
Cefotaxime	30 µg	371(92.8)	29(7.2)
Chloramphenicol	30 µg	348(87.0)	52(13.0)
Ciprofloxacin	5 µg	336(84.0)	64(16.0)
Ceftazidime	30 µg	333(83.3)	67(16.8)
Gentamycin	10 µg	331(82.7)	49(14.8)
Mipenem	10 µg	02(0.5)	398(99.5)

Table 2: Distribution of ESBL-producing *Escherichia coli* by Clinical Specimens

Specimen	ESBL DETECTION		
	Number positive (%)	Number negative (%)	Total number examined (%)
Blood	5(26.3)	14(73.7)	19(100)
Wound swabs	9(17.3)	43(82.7)	52(100.0)
Urine	33(16.5)	167(83.5)	200(100.0)
Stool	16(14.5)	94(85.5)	110(100.0)
Sputum	1(8.3)	11(91.7)	12(100.0)
Throat swabs	0(0.0)	7(100)	7(100.0)
Total	64(16.0)	336(84.0)	400(100.0)

$\chi^2 = 3.64$; df = 5; p = 0.60

Table 3: Distribution of Plasmid-mediated *Bla* TEM Gene By Specimen Types

Specimens	Plasmid-mediated <i>bla</i> TEM		
	Number negative (%)	Number positive (%)	Total (%)
Blood	14(73.7)	5(26.3)	19(100.0)
Wound swabs	45(86.5)	7(13.5)	52(100.0)
Urine	178(89.0)	22(11.0)	200(100.0)
Stool	98(89.1)	12(10.9)	110(100.0)
Sputum	11(91.7)	1(8.3)	12(100.0)
Throat swabs	7(100.0)	0(0.0)	7(100.0)
Total	353(88.2)	47(11.8)	400(100.0)

$\chi^2 = 5.29$, df = 5, p = 0.3

Table 4: Distribution of Plasmid-mediated *Bla* TEM Gene by Age

Age (yrs)	Plasmid-mediated <i>bla</i> TEM		
	Number negative (%)	Number positive (%)	Total (%)
≤ 16	95(88.8)	12(11.2)	107(100.0)
17-30	109(90.8)	11(9.2)	120(100.0)
31-44	88(85.4)	15(14.6)	103(100.0)
45-58	46(85.2)	8(14.8)	54(100.0)
>59	15(93.8)	1(6.3)	16(100.0)
Total	353(88.2)	47(11.8)	400(100.0)

$\chi^2 = 2.54$, df = 4, p = 0.637.

Table 5: Distribution of Plasmid-mediated *bla* TEM Gene by Sex.

Sex	<i>Bla</i> TEM		
	Number negative (%)	Number positive (%)	Total (%)
Female	189(87.5)	27(12.5)	216(100.0)
Male	164(89.1)	20(10.9)	184(100.0)
Total	353(88.2)	47(11.8)	400(100.0)

$\chi^2 = 0.255$; df = 1; p = 0.614.

Table 6: Distribution of Plasmid-mediated *Bla* TEM Gene According to Health Facilities

Health Facilities	Plasmid-mediated <i>Bla</i> TEM		
	Number negative (%)	Number positive (%)	Total (%)
GH Makurdi	45(84.9)	8(15.1)	53(100.0)
BSUTH Makurdi	127(86.4)	20(13.6)	147(100.0)
FMC Makurdi	88(88.0)	12(12.0)	100(100.0)
BMMC Makurdi	27(90.0)	3(10.0)	30(100.0)
FSP Makurdi	37(92.5)	3(7.5)	40(100.0)
PHC Makurdi	29(96.7)	1(3.3)	30(100.0)
Total	353(88.2)	47(11.8)	400(100.0)

$\chi^2 = 3.9$; df = 5; p = 0.564.

Key:

GH = General Hospital; BSUTH = Benue State University Teaching Hospital

FMC = Federal Medical Centre; BMMC = Bishop Murray Medical Centre; FSP = Family Support Programme Clinic and
PHC = Primary Health Care.

5.0 DISCUSSION.

Out of 400 isolates tested, 64 (16.0%) produced extended spectrum β - lactamases (ESBLs). Isolates that produce ESBLs are known to resist antimicrobial agents more than on- ESBL producers. This carriage of ESBLs is lower than some reports seen in some parts of Nigeria and other parts of the world: [19] reported 20.9% (n = 28) in South West Nigeria, [23] reported 44.3% in Benin- City Southern Nigeria, 36.8% was reported for Kano State [18], 59.4% in Enugu State [7], 66% in Lagos [1], 41% in Pakistan [8]. It is however, higher than 3.3% reported from Gaza strip [2], 11.8% from Macedonia [9]. Our finding is similar to the report of 18.6% by [20] from a neighboring Plateau State, Jos, North-Central Nigeria and 18.5% by [17] in India.

The susceptibility profiles of ESBL-producing *Escherichia coli* showed mipenem to be the most active antimicrobial agent. This is because mipenem is stable in the presence of ESBLs. Previous use of antimicrobial agents, especially cephalosporins and quinolones have been reported to be risk factors associated with emergence of ESBLs [22]. These antibiotics are the drug of choice for the treatment of infections by Gram-negative bacteria in Nigeria [19]. Sales and use of antibiotics is unregulated in Nigeria [14]. These might be the reason for the high prevalence of ESBL infections in Nigeria, especially in the present study location. Demographic data showed that female population carried a higher burden of ESBL infections than the male counterpart and the age group most affected is 45 to 58 years. This is in agreement with the report of [20] from Jos, North-Central Nigeria.

The PCR analysis yielded 11.8% (n = 47) of plasmid-mediated *bla* TEM variants. The prevalence of *bla* TEM resistance genes from this study is lower than the reports of other researchers from different parts of Nigeria and world: [21] reported a prevalence of 42.1% (n = 48) from Ekiti State, Nigeria; [21], reported 32% from Osun State, Nigeria where the highest percentage of *bla* TEM was recovered from sputum specimen (32%) isolates and the next was from blood specimens (21.5%). Maiduguri, Borno State in North-Eastern Nigeria had a prevalence of 31.4% (n =38) as reported by [18] to ceftazidime in his study. Though the prevalence of 11.8% for *bla* TEM in this study appear to be low compared to reports by other researchers; it is still important to consider it significant being the first report from the study location. With the free availability of antibiotics and unrestricted use, the prevalence will likely rise in the near future.

6.0 CONCLUSION.

The result from this study reported a relatively low occurrence of *bla* TEM resistance gene compared with reports from other parts of Nigeria and world. The emergence of ESBL- producing *Escherichia coli* poses a serious public health challenge, which will likely result in substantial limitations in the efficacy of therapeutic interventions. This study is the first report of the emergence of *bla* TEM gene in Makurdi, Benue State. There is need to establish screening strategies for early identification of patients that may be carriers of ESBL producing bacteria in the hospitals. Prescription of mipenem should be done with caution and carefully

monitored, as it remains a viable option in a situation of overwhelming resistance of *Enterobacteriaceae* to other classes of routinely used antibiotics. This study has also shown the importance of infection, prevention and control measures to reduce the dissemination of antibiotic resistance within the hospital referral system in Makurdi, Benue State, Nigeria.

Authors' contribution.

P. O. Abba conceived, designed and executed the study; G. M. Gberikon and E. B. Agbo supervised data collection while E. U. Umeh analyzed the data and supervised manuscript writing. All authors read and approved the final manuscript.

Conflict of interest.

The authors declare none.

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