Extended-Spectrum β-Lactamase-Producing 
*Escherichia coli* and *Klebsiella* Species in Pediatric Patients Visiting International Friendship 
Children’s Hospital, Kathmandu, Nepal

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**ABSTRACT**

**INTRODUCTION:** Emergence and spread of antimicrobial resistance (AMR) is a global threat and significantly affects the treatment options for common infectious diseases. Inappropriate use of antibiotics, particularly third-generation cephalosporins, has contributed to the development of AMR. This study aims to determine the prevalence of extended-spectrum β-lactamase (ESBL) production in *Escherichia coli* and *Klebsiella* species isolated from various clinical samples.

**METHODS:** This cross-sectional study was conducted at International Friendship Children’s Hospital, Kathmandu, Nepal, from August 2017 to January 2018. A total of 1443 samples that included urine, pus, wound swab, endotracheal tip, catheter tip, and blood were collected from pediatric patients below 15 years and processed by standard microbiological methods. Following sufficient incubation, isolates were identified by colony morphology, gram staining, and necessary biochemical tests. Identified bacterial isolates were then tested for antibiotic susceptibility test by modified Kirby-Bauer disk diffusion method and were subjected to ESBL screening by using 30µg cefotaxime and ceftazidime. The ESBL production was confirmed by combination disk method.

**RESULTS:** From a total of 103 nonduplicated clinical isolates, *E. coli* (n = 79), *Klebsiella pneumoniae* (n = 18), and *Klebsiella oxytoca* (n = 6) were isolated from different clinical specimens. Of which, 64 (62.1%) exhibited multidrug resistance, and 29 (28.2%) were ESBL producers. All ESBL-producing isolates were resistant toward ampicillin, cefotaxime, ceftriaxone, and ceftazidime. Most ESBL producers were susceptible toward imipenem (89.7%; 26/29), nitrofurantoin (82.8%; 24/29), piperacillin/tazobactam (79.3%; 23/29), and amikacin (72.4%; 21/29).

**CONCLUSIONS:** A high prevalence of multidrug-resistant ESBL organisms was found in this study among pediatric patients. Treatment based on their routine identification and susceptibility to specific antibiotics is critical to halt the spread of AMR and ESBL.

**KEYWORDS:** Antibiotic resistance, ESBL, *Escherichia coli*, *Klebsiella oxytoca*, *Klebsiella pneumoniae*, MDR

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**Introduction**

*Escherichia coli* and *Klebsiella* species are the most common causative pathogens for most of the infections, especially in countries with poor health care system.¹ *E. coli* is a normal flora of human and animal gut but can also be found in water, soil, and vegetation.² *Klebsiella* species are considered as major opportunistic pathogens that can cause infections mostly in children. *Klebsiella pneumoniae* is an important cause of human infections among all *Klebsiella* species, followed by *Klebsiella oxytoca, Klebsiella ozaenae,* and *Klebsiella rhinoscleromatis*. Several common bacterial infections such as gastroenteritis, urinary tract infection (UTI), septicemia, and neonatal meningitis are mainly caused by *E. coli* and *Klebsiella spp* in children.³,⁴

Commonly used antimicrobial agents against these pathogens are tetracycline, β-lactams, fluoroquinolones, aminoglycosides, and cotrimoxazole. However, antimicrobial resistance (AMR) among Enterobacteriaceae has increased dramatically in recent years, limiting the therapeutic options. Isolates that are not susceptible to at least 3 or more groups of antimicrobials are known as multidrug resistant (MDR) organisms.⁵

Extended-spectrum β-lactamases (ESBLs) are class A β-lactamases, a rapidly evolving group of β-lactamases with the ability to hydrolyze and cause resistance to the oxy-imino cephalosporins (cefotaxime, ceftazidime, ceftriaxone, and cepalexime) and monobactams (aztreonam).⁶ ESBL-producing *E. coli* and *K. pneumoniae* are the predominant cause of childhood infections and present significant challenges⁷ such as development of adverse outcomes,⁸ treatment failure due to multidrug resistance, and high morbidity and mortality.⁹ Empirical and symptomatic (without a diagnosis) use of antibiotics in resource poor settings is responsible for higher incidence of antibiotic resistance among bacteria.¹⁰
Several studies in the past have investigated the prevalence of ESBL organisms among inpatients, mostly focused in adult patients. Studies have shown the varying prevalence of ESBL organisms, for instance, the prevalence was 27.7% in Pokhara, 18% in Kathmandu, 14% in pediatric hospital in Kathmandu. Another study reported 35.9% ESBL in E. coli isolates among outpatients at tertiary care hospital in Kathmandu. However, similar study from Lalitpur district reported 6.8% ESBL-producing isolates. Studies have shown the wide range in the prevalence of ESBLs (10%-43%) in different hospitals/settings from various samples.

Although it is deemed to be essential to have a routine diagnosis and monitoring of ESBL-producing clinical isolates in clinical laboratories, ESBL screening as a routine test has not yet been practiced in Nepal. In addition, very few studies have reported on ESBL-producing clinical isolates from pediatric patients in Nepal. Only 1 study in the past has reported ESBLs (prevalence: 38.9%) from urine samples in pediatric patients from a tertiary teaching hospital in Kathmandu. Expanding and building on the previous research, this study focused to isolate both E. coli and Klebsiella spp from wider and larger number of clinical specimens from pediatric patients. The main objectives of this study were to explore the prevalence of ESBL-producing organisms, including the resistance types among pediatric patients attending a tertiary care pediatric hospital at Kathmandu.

Methods
Study design, area, and sample population
This was a cross-sectional study conducted at International Friendship Children’s Hospital, Maharajgunj, Kathmandu, Nepal, during August 2017 to January 2018. The study population comprised children below 15 years attending the hospital for treatment.

The specimens were collected adhering to a standard protocol from pediatric patients below 15 years of age. Children who attended the hospital for treatment and provided assent (from parents) or consent for the study were included in the study. Children who had chronic diseases such as leukemia, malnutrition, and neuropsychiatric disorders based on the diagnosis made by clinicians, including if they were initiated with the antibiotic treatment after the admission, were excluded.

A total of 1443 different samples that included urine (n = 1050), pus (n = 50), wound swabs (n = 40), endotracheal tip (n = 83), catheter tip (n = 40), and blood (n = 140) were collected and processed by standard microbiological methods.

Sample collection and transport
Special measures were taken to collect the urine samples from children who were not able to use toilet on their own. An adhesive, sealed, sterile collection bag was placed underneath the genitalia to collect urine sample. Toilet-trained children were requested to collect mid-stream urine assisted by their parents in a sterile, dry, wide-necked, and leak-proof container. In either condition, genitalia were cleaned with alcohol swab to reduce contamination.

In the case of infected wounds, in addition to wound swab, pus was aspirated in syringe by trained medical personnel. In case pus was not discharging, cotton swab was gently rolled over the surface of the wound approximately 5 times, focusing on areas where there was evidence of pus or inflamed tissue. Two swabs were taken from each patient, one for culture and another for direct gram staining.

About 2 mL of blood from children was withdrawn and dispensed into sterile screw capped culture bottles containing BHI (brain heart infusion) broth. Specimens were collected from other sources such as endotracheal and catheter tips by trained medical personnel. The collected samples were labeled properly and were immediately delivered to a laboratory for further processing. When immediate delivery was not possible, the specimens were refrigerated at 4°C to 6°C.

Laboratory examinations of samples
Culture. For processing of each sample, microbiological protocols were followed according to standard microbiological guidelines.

Urine sample: Using a sterile calibrated loop, urine sample was inoculated on MacConkey agar (MA) and blood agar (BA), and then incubated aerobically at 37°C for 24 hours. Colony count was made, and positive result was considered for plates showing more than or equal to 10⁵ colony forming units (CFU)/mL of urine based on Kass, Marple, and Sanford criteria.

Blood sample: Blood sample was inoculated on BHI broth for 7 days at 37°C. Bottles showing turbidity during the period were subcultured aerobically in MA and BA at 37°C for 24 to 48 hours.

Pus, wound swab specimens were inoculated into MA and BA plate sand incubated at 37°C overnight.

Other specimens: Endotracheal and catheter tips were first incubated on BHI broth at 37°C for 24 hours and subcultured on MA and BA plates and incubated at 37°C overnight.

Identification of E. coli and Klebsiella spp. Presumptive identification of E. coli and Klebsiella spp was done on the basis of colony color and Gram staining morphology. Then, obtained pure cultures of isolates were assessed for various biochemical tests (indole, methyl red, Voges-Proskauer, citrate, triple sugar iron agar, oxidative/fermentative, urease test for confirmation).

Antibiotic susceptibility testing
All identified isolates of E. coli and Klebsiella spp were treated for susceptibility testing against ampicillin (10 μg), gentamicin (10 μg), amikacin (30 μg), aztreonam (30 μg), cefoxitin (30 μg), aztreonam (30 μg), and amikacin (30 μg) disks. Susceptibilities were interpreted according to the Sanford criteria.
ceftazidime (30 µg), cefotaxime (30 µg), ceftazidime (30 µg), ciprofloxacin (5 µg), imipenem (10 µg), piperacillin/tazobactam (100/10 µg), nitrofurantoin (300 µg), and cefepime (30 µg) (HiMedia India Pvt. Ltd, Bengaluru, India) following Kirby-Bauer method on Mueller-Hinton Agar (HiMedia India Pvt. Ltd). Results were interpreted based on the Clinical and Laboratory Standards Institute (CLSI) 2016 guidelines. Those isolates which were not susceptible (either resistant or intermediate) to 3 or more antibiotic classes were considered as MDR.

Screening and confirmation of ESBL producers. Bacterial isolates exhibiting reduced susceptibility to ceftazidime (30 µg), cefotaxime (30 µg), ceftazidime (30 µg), and aztreonam (30 µg) were considered as potential ESBL producers. The ESBL production was further phenotypically confirmed by combination disk method as described by CLSI 2014. The disk used was cefotaxime and ceftazidime alone and cefotaxime and ceftazidime in combination with clavulanic acid. A ≥5 mm increase in growth inhibition zone for any antimicrobial associated with clavulanic acid in comparison with the inhibition zone of antimicrobial tested alone confirmed ESBL production.

Quality control
Each batch of media and reagents was subjected to sterility and performance testing. During antibiotic susceptibility test, quality control was done using the control strains of *Escherichia coli* ATCC 25922.

Data management and statistical analysis
Data were entered and analyzed by using IBM SPSS Statistics for Windows, Version 21.0 (IBM Corp, Armonk, NY, USA). Descriptive and inferential statistics were analyzed.

Results
Prevalence of bacterial isolates. A total of 1443 different clinical specimens were processed during the study, of which 299 (20.7%) samples showed bacterial growth. Of the 299 isolates, 79 (26.4%), 18 (6.0%), and 6 (2%) were identified as *Escherichia coli*, *K. pneumoniae*, and *K. oxytoca*, respectively. *E. coli* was predominant bacteria isolated from urine samples (86.0%; 68/79), followed by pus/wound pus samples (8.8%; 7/79). *K. pneumoniae* (77.8%; 14/18) and *K. oxytoca* (83.3%; 5/6) were mostly isolated from urine samples (Table 1).

### Antibiotic susceptibility pattern of bacterial isolates
Among 103 bacterial isolates, 90.3% (*n* = 93) were found to be susceptible to imipenem, followed by piperacillin/tazobactam (88.3%; *n* = 91), nitrofurantoin (85.5%; *n* = 88), and amikacin (82.5%; *n* = 85). Most *E. coli* isolates (92.4%; 73/79) were found to be susceptible to imipenem, followed by nitrofurantoin (91.2%; 72/79). Similarly, 88.9% (16/18) of *K. pneumoniae* were found to be susceptible to amikacin. *K. oxytoca* were found to be 100% (6/6) susceptible to gentamicin, piperacillin/tazobactam, and imipenem (Table 2).

### MDR profile in bacterial isolate
Among the total of 103 bacterial isolates, 62.1% (*n* = 64/103) were found to be MDR; the highest MDR strains were detected in *K. pneumoniae* (88.9%; 16/18), followed by *E. coli* (57%; 44/79) and *K. oxytoca* (50%; 3/6) (Figure 1).

ESBL production among *E. coli* and *Klebsiella* spp
Among 103 *E. coli* and *Klebsiella* isolates, 28.2% (*n* = 29/103) were confirmed as ESBL producers by combination disk diffusion method. The highest percentage of ESBL production was found among *K. pneumoniae* (33.3%; 6/18), followed by *E. coli* (27.9%; 22/79) and *K. oxytoca* (16.7%; 1/6) (Table 3).

Distribution of ESBL producers according to different age of patient. Among the 103 isolates, 77.7% (*n* = 80) were isolated from children ≤5 years age, followed by 6 to 10 years age group children (15.5%; *n* = 16). Of 103 bacterial isolates, 28.1% (*n* = 29) were ESBL producers and the most (82.8%; *n* = 24) were isolated from children ≤5 years of age. There was no association between ESBL producers and age of patients (*P* < .05) (Table 4).

### Distribution of ESBL producers in different wards of hospitals.
Of 29 isolates of ESBL producers, 51.7% (*n* = 15) were from...
inpatients, whereas 48.3% (n = 14) were from outpatient department. There was no significant association between ESBL production and type of the patients (P > .05) (Table 5).

**Antibiotic susceptible pattern of ESBL producers.** All of ESBL producers’ isolates were found to be resistant toward cefotaxime, ceftriaxone, ceftazidime, and ampicillin. Most ESBL producers were found to be susceptible toward imipenem (89.7%; 26/29), nitrofurantoin (82.8%; 24/29), piperacillin/tazobactam (79.3%; 23/29), and amikacin (72.4%; 21/29) (Figure 2).

**Discussion**

**Overall findings**

This study found the high prevalence (>60%) of MDR bacteria in clinical specimens isolated from the tertiary care hospital of children in Kathmandu valley. Among MDR isolates, half of the isolates were ESBL producers. Most ESBL-producing isolates were found to be resistant toward cefotaxime, ceftriaxone, ceftazidime, and ampicillin. Similar findings were reported in previous studies in different clinical settings of Nepal.12,17,22,23

Most isolates (>80%) in this study were found susceptible to imipenem, piperacillin/tazobactam, and amikacin. The high efficacy of amikacin and imipenem against *E. coli* and *Klebsiella* was also reported from studies conducted in Chitwan24 and Lumbini,25 Nepal. The findings were also in line with a study from Manipal Teaching Hospital, Pokhara.11

Most isolates (77.5%) were resistant to ampicillin and more than half of the cephalosporin group of antibiotics. Similar findings were observed in a tertiary hospital in Pokhara, Nepal.11 This type of resistance could be due to the production of several β-lactamase enzymes. As ampicillin is the first-line

| ORGANISMS         | NO. OF ISOLATES | ESBL PRODUCER CONFIRMED (%) |
|-------------------|-----------------|-----------------------------|
| *E. coli*         | 79              | 22 (27.8)                   |
| *Klebsiella* spp. | 18              | 6 (33.3)                    |
| *Klebsiella* oxytoca | 6              | 1 (16.7)                    |
| Total             | 103             | 29 (28.2)                   |

Abbreviation: ESBL, extended spectrum β-lactamase.
β-lactam drug for Enterobacteriaceae, it can be easily hydrolyzed by β-lactamase enzymes. Resistance to fluoroquinolones is due to mutation at the target site, ie, gyrA (gyrase subunit gene) and parC (topoisomerase subunit gene) and efflux.26

The AMR, including MDR, is a global problem, and its burden varies between the regions; however, low- and middle-income countries share a disproportionate burden due to multitude of factors embedded in the characteristics of the health system, policy, and the practice.27 Moreover, MDR pathogens are more common in hospital settings and are mostly of nosocomial origin which is often difficult to treat.28 MDR pose a major threat in the management of uropathogens.29-31 More than two-thirds of the isolates in this study were MDR, mostly being E. coli, K. pneumoniae, and K. oxytoca. A high prevalence of MDR strains have been reported consistently from past studies from within Kathmandu10,12,17,28 and outside.8,14,16

Table 4. Distribution of ESBL according to sex and age of children.

| AGE GROUPS, Y | NO. OF ISOLATES (%) | ESBL PRODUCTION | P-VALUE |
|--------------|---------------------|-----------------|---------|
| ⩽5           | 80 (77.7)           | 24 (82.8)       |         |
| 6-10         | 16 (15.5)           | 4 (13.8)        | .837*   |
| 11-15        | 7 (6.8)             | 1 (3.4)         |         |
| Total        | 103 (100.0)         | 29 (28.1)       |         |

Abbreviation: ESBL, extended spectrum β-lactamase.

*χ² test.

Table 5. Distribution of ESBL producers in different wards of hospitals.

| WARDS         | NO. OF ISOLATES (%) | ESBL PRODUCER BACTERIAL ISOLATES (%) | P-VALUE |
|---------------|---------------------|-------------------------------------|---------|
| Outpatients   |                     |                                     |         |
| OPD           | 52 (50.5)           | 14 (48.3)                           |         |
| Inpatients    |                     |                                     | .737*   |
| ICU           | 13 (12.6)           | 5 (17.2)                            |         |
| Other than ICU| 38 (36.9)           | 10 (34.5)                           |         |
| Total         | 103 (100.0)         | 29 (28.1)                           |         |

Abbreviations: ESBL, extended spectrum β-lactamase; ICU, intensive care unit; OPD, outpatient department.

*χ² test.

Figure 2. Antibiotic susceptibility pattern of ESBL producers. ESBL indicates extended-spectrum β-lactamase.
E. coli and K. pneumoniae are major ESBL producers posing serious threat to the treatment regimen. ESBL enzymes are becoming increasingly expressed by many strains of pathogenic bacteria presenting diagnostic challenges to the clinical microbiology laboratories. The highest bacterial isolates were found in children less than 5 years age, including the prevalence of ESBL organisms which was above 80%. The reason for this may be due to the immunological status of the children below 5 years of age who are more vulnerable to infections. The higher prevalence of bacterial growth in inpatients may have been added by nosocomial infections. Nosocomial infections are associated with prolonged hospital stay, intensive care unit admission, extensive use of invasive medical devices, and over-consumption of antibiotic among inpatients.

Most ESBL organisms were susceptible to imipenem, piperacillin/tazobactam, amikacin, and nitrofurantoin. However, ESBL producers were resistant to ampicillin and cephalosporin group of antibiotics. These findings are consistent with similar studies reported from Nepal. The high proportion of resistance to third-generation cephalosporins reported for E. coli and K. pneumoniae means that treatment of severe infections likely to be caused by these bacteria in many settings must rely on carbapenems, the last resort to treat severe community- and hospital-acquired infections.

Increasing spectrum of ESBL drug-resistant bacterial isolates can cause major problems for physicians in choosing from the available therapeutic options, if these organisms are not routinely isolated. Reporting of ESBL-producing isolates from clinical samples is thus critical for the clinicians to select appropriate antibiotics for the treatment, including to take proper precaution to prevent the spread of these resistant organisms to other patients.

Strengths and limitations

This study will be a useful reference for future studies to explore and expand on the wider prevalence of ESBL organisms in clinical and nonclinical settings. As our study was based on phenotypic detection of AMR and ESBL production that excluded identification and characterization of wide sorts of lactamases and pathogenic strains, genotypic characterization is recommended in future studies.

Implications for AMR and its control

This study has identified one of the major determinants of burgeoning AMR in Nepal. All antibiotics are available over the counter (OTC) in Nepal without medical prescriptions, and this is a major challenge as it contributes to antibiotic pressure and development of resistance. The availability of OTC antibiotics and its consumption before arriving to hospitals may also confound the clinical presentation, including general culture and sensitivity tests. Thus, cautious evaluation of preceding treatment history, combined with strong suspicion for ESBL and MDR and its diagnosis, may inform the appropriate treatment. The findings in this study warrant a relevant stakeholder’s engagement to strengthen the health policy to rationalize the use of antibiotics, including promoting diagnostic-based antibiotic prescriptions. Specifically, in pediatric patients with UTIs, it is critical to establish the diagnosis of ESBL organisms before initiating the antibiotic treatment.

Conclusion

A high prevalence of MDR ESBL organisms was found among pediatric patients in this study. Identification of ESBL producers in routine treatment of infectious diseases in pediatric patients can reduce unnecessary and inappropriate antimicrobial use. Hospitals treating infectious diseases can benefit by integrating antimicrobial stewardship programs to combat the emergence of AMR and ESBLs.

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Author Contributions

All the authors made substantial contribution to the study. K.K. and B.D. conceived and designed the study. K.K. collected samples, investigated, and recorded the laboratory findings. K.R.R., S.K., M.R.B., and P.G. advised and formulated the methodology for the study. K.R.R. and B.A. are responsible for reviewing several versions of the article. Others helped to review and amend this article. All authors read and approved the final article.

Availability of Data and Materials

All data pertaining to this study are within the article.

Ethics Approval and Consent to Participate

Institutional Review Committee of Nepal Health Research Council (NHRC) approved this research. Written consent was applicable to literate people while verbal consent was approached for the rest subjects. Parents/Guardians were interviewed in case of children. Strict adherence to the ethical guidelines was taken, and we declare that this research is free from selection bias.

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