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Bacteriological Profile of UTI and their Antimicrobial Resistance Pattern in Pediatric Age group patients at RMCH

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ABSTRACT: Background: Urinary tract infections (UTIs) are common pediatric infections and contribute to high morbidity. Gram negative bacilli predominating the infections. At present, the antimicrobial resistance has quadrupled worldwide and poses a serious threat to the treatment of patients. Objectives: The present study was designed to find out the bacteriological profile of urinary tract infection and their antibiotic resistance pattern in children. Material and Methods: A cross-sectional study was done in the department of Microbiology, Rajshahi Medical College from July to December 2024. Midstream urine samples were collected from clinically suspected cases of UTI in the age group of 0 to 18 years from various indoor and outdoor patients attending at Rajshahi Medical College hospital. Results: Out of 984 cases, 195 (19.8%) were culture positives, where females were 110(56.4%) and male were 85(43.6%). Most common isolate identified was Escherichia coli 113(57.9%), followed by Klebsiella spp.31(15.9%), Pseudomonas aeruginosa 18(9.2%) and Enterococci spp. 19(9.8%). Gram negative organisms show highest resistance to cefixime, ceftriaxone, cefuroxime, azithromycin, amoxiclav and ciprofloxacin and highest sensitivity to Colistin, nitrofurantoin and imipenem. Gram positive bacteria were highly resistant against cefixime, Cotrimoxazole, ceftriaxone, cefuroxime, azithromycin, amoxiclav and ciprofloxacin. Conclusion: This study indicates that gram-negative bacteria, particularly E. coli are the commonest isolated organism, and all isolates show resistance to commonly used antimicrobial agents. Urinary tract infections as well as the significant growth and spread of resistant bacterial pathogens in children should be monitored regularly.

Keywords: Paediatric Age Group, Urinary Tract Infection, Uropathogens, Antimicrobial Resistance.

Article at a glance:

Study Purpose: To contribute to existing knowledge or propose new ideas.

Key findings: Among gram-negative bacteria, predominant gram-positive bacteria were isolated in children Escherichia Coli 113 (57.9%).

Newer findings: In this study, Gram negative bacteria showed higher resistance to commonly used antibiotics from previous study. **Abbreviations:** UTI: Urinary Tract Infection.

INTRODUCTION

Urinary Tract Infection (UTI) is a common disease in pediatric practice. Early diagnosis and prompt treatment can reduce the risk of renal scarring and its long-term sequelae such as hypertension and end stage renal failure. The estimated incidence of UTI is 3% in girls and 1% in boys during the first ten years of life. After the initial UTI, the prevalence of UTI during the first 6-12 months is upto 30% in infants and children. The risk of having a UTI before the age of 14 years is approximately 3-10% in girls and 1-3%

in boys. The prevalence of UTI among paediatric patients worldwide was 2-20%.² The clinical and laboratory diagnosis of UTI in children poses some difficulties as presentation varies among different age groups. Diagnosis is often missed in infants and young children because urinary symptoms are minimal and often nonspecific. Collection of appropriate samples of urine for investigations is also often difficult in this age group. The presence of significant bacteriuria in urine culture is essential to diagnose UTI and patients then must be evaluated for

anomalies of the urinary tract to prevent recurrence and complications.3 Most common cause of UTI in children is Escherichia coli followed by other organisms like Klebsiella species, Enterococcus species, Pseudomonas Staphylococcus species, aureus, Enterobacter species, Proteus species and Coagulase negative staphylococcus.4 In suspected cases of UTI, empirical treatment with antibiotics is started after collecting urine samples for culture and antimicrobial sensitivity. The selection of antibiotics should be based on the commonly prevalent urinary pathogens and their antibiotic sensitivity in the region. This is important because injudicious use of antibiotics has led to bacterial resistance to antibiotics. Depending upon the extent of use of antibiotics, the antibiogram pattern is different in different countries and in different regions of the same country.5 This study is therefore undertaken to study the current scenario of UTI in children in Rajshahi region. The aim of the study is clinical profile, bacteriology and antibiotic sensitivity pattern of bacteria causing UTI in children attending Rajshahi medical college hospital.

MATERIAL AND METHODS

This cross-sectional study was carried out in the Microbiology department of Rajshahi medical college over a period from July to December 2024. Patients from the age of 0 upto 18 years presenting with urinary symptoms (dysuria, urgency, frequency, incontinence, hematuria and suprapubic pain) and those with fever without a focus were enrolled in the study. Wet mount microscopy of urine was done to detect pyuria, hematuria and presence of any other abnormal cells. Clean catch midstream urine samples were collected in older children while transurethral bladder catheterization was done to obtain urine samples in infants and younger children. Urine culture was done on Mac conkey agar, Nutrient agar and Hichrome UTI agar with a calibrated loop. A growth of greater than 10 colony forming units/ml of a single organism for midstream urine samples and greater than 5x10 colony forming units/ml for samples obtained by catheterization considered was significant bacteriuria and UTI. The antibiotic sensitivity test was done on Mueller-Hinton agar by Kirby-Bauer disc diffusion test.6

RESULTS

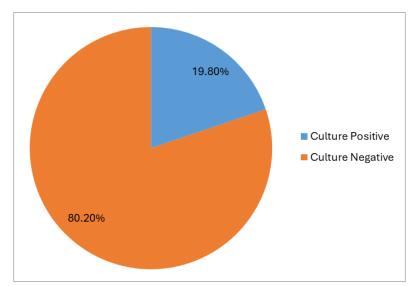


Figure 1: Frequency of Culture Positive and Negative Cases (N=984)

Figure 1 shows culture positivity of isolated organisms. Out of 984 samples, 195(19.80%) samples

were culture positive while 789(80.20%) samples were culture negative.

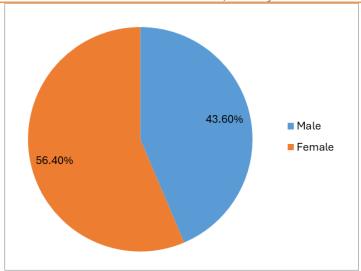


Figure 2: Sex Distribution of Culture Positive Cases (N=195).

Figure 2 shows sex distribution of culture positive cases. Males were 43.6% and females were 56.4% giving a male and female ratio 1:1.3.

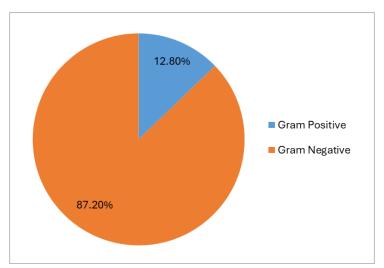


Figure 3: Frequency of Gram Positive and Gram-Negative Bacteria (N=195)

Figure 3 shows the distribution of grampositive and gram-negative isolate among culture positive cases. Among the total 195 isolates, Gram negative bacteria were 170(87.20%) and gram-positive bacteria were 25(12.80%).

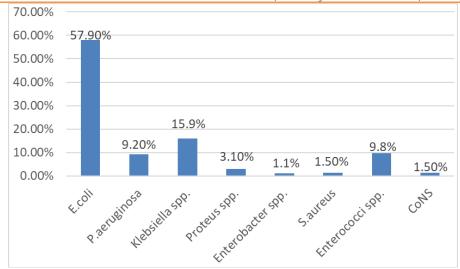


Figure 4: Bacteria Isolated from Culture Positive Cases (N=195).

Figure 4 shows the identified species of bacteria isolated from culture positive cases. Out of 984 samples, a total of 195 bacteria were identified. *E.*

coli was 113(57.9%) followed by Klebsiella spp. 31 (15.9%), Enterococci spp. 19(9.8%), P. aeruginosa 18(9.2%) and proteus spp. 6(3.1%).

Table 1: Antimicrobial Resistance Pattern of Gram-Negative Bacteria

Antimicrobial agents	E. coli	Klebsiella	Pseudomonas	Proteus spp.	Enterobacter
	(N=113)	<i>spp.</i> (N=31)	aeruginosa (N=18)	(N=06)	<i>spp.</i> (N=2)
Imipenem	19(16.8%)	06(19.4%)	05(27.8%)	01(16.7%)	00
Azithromycin	93(82.3%)	27(87.1%)	17(94.4%)	05(83.3%)	02(100%)
Ciprofloxacin	55(48.7%)	16(51.6%)	09(50%)	03(50%)	01(50%)
Ceftriaxone	92(81.4%)	29(93.5%)	15(83.3%)	05(83.3%)	02(100%)
Cefepime	52(46.1%)	19(61.3%)	16(88.8%)	03(50%)	01(50%)
Piperacillin/tazobactam	7(32.7%)	09(29.1%)	06(33.3%)	02(33.3%)	01(50%)
Cefixime	95(84.1%)	30(96.8%)	18(100%)	05(83.3%)	02(100%)
Aztreonam	47(41.6%)	14(45.2%)	10(55.6%)	04(66.7%)	01(50%)
Amikacin	57(50.4%)	15(48.4%)	08(44.4%)	03(50%)	01(50%)
Cefuroxime	98(86.7%)	30(96.8%)	18(100%)	06(100%)	18(100%)
Colistin	07(6.2%)	02(6.5%)	02(11.1%)	00	00
Levofloxacin	33(29.2%)	07(22.6%)	06(33.3%)	02(33.3%)	01(50%)
Amoxiclav	67(59.3%)	17(54.8%)	11(61.1%)	04(66.7%)	01(50%)
Nitrofurantoin	13(11.5%)	04(12.9%)	03(16.6%)	01(16.7%)	00

Table 1 shows the antimicrobial resistance pattern among gram negative bacteria. All the gramnegative bacteria were highly resistant against cefixime, ceftriaxone, cefuroxime, azithromycin,

amoxiclav and ciprofloxacin. Colistin, nitrofurantoin, imipenem and levofloxacin were showed lower resistance against gram negative bacteria.

Table 2: Antimicrobial Resistance Pattern of Gram-Positive Bacteria

Antimicrobial agents	Enterococci spp. (N=19)	CoNS (N=03)	S. aureus (N=03)
Imipenem	04(21.1%)	01(33.3%)	01(33.3%)
Azithromycin	16(84.2%)	03(100%)	03(100%)
Ciprofloxacin	09(47.4%)	01(33.3%)	16(66.7%)
Ceftriaxone	15(78.9%)	02(66.7%)	02(66.7%)
Vancomycin	00	00	00

Linezolid	02(10.5%)	00	00
Nitrofurantoin	03(15.7%)	00	00
Amoxiclav	12(63.2%)	01(33.3%)	02(66.7%)
Amikacin	10(52.6%)	01(33.3%)	02(66.7%)
Cefuroxime	18(94.7%)	02(66.7%)	03(100%)
Cefixime	17(89.5%)	02(66.7%)	03(100%)
Levofloxacin	05(26.3%)	01(33.3%)	01(33.3%)
Cotrimoxazole	15(78.9%)	02(66.7%)	03(100%)

Table 2 shows the antimicrobial resistance pattern among gram positive bacteria. Gram positive bacteria were highly resistant against cefixime, cotrimoxazole, ceftriaxone, cefuroxime, azithromycin, amoxiclav and ciprofloxacin. Vancomycin, linezolid, nitrofurantoin and imipenem were showed lower resistance against gram positive bacteria.

DISCUSSION

Out of 984 urine samples obtained in the Microbiology laboratory from RMCH, Rajshahi for aerobic culture and sensitivity, 19.8% yielded positive culture whereas 80.2% yielded no growth. This study was nearly similar with the study of Rafi et al. Nag et al. and Venugopal et al., but dissimilar with the study of Sharmin et al. and Koli et al.7-11 Figure 2 shows sex distribution of culture positive cases. Among them, 85 (46.6%) were male and 110 (56.4%) were female. Females are at increased risk due to a shorter urethra and its proximity to the anus which encourages contamination and ascent of fecal flora into the urinary tract. This study was nearly similar with the study of Sharmin et al.10 and Patel et al.3 but dissimilar with the study of Nazme et al. and Koli et al. 11, 12 Out of a total 984 samples, Gram negative bacteria were accounted for higher isolation rate (Gram-positive 12.8% and Gram-negative 87.2%) than gram positive bacteria. This study was nearly similar with the study of Nahar et al. and Arora et al. but nearly dissimilar with the study of Thaddanee et al. and Khandelwal et al. 13-16 The reason for this high occurrence of culture positivity may be due to the fact that most of the study population belonged to lower middle and lower socioeconomic group with poor knowledge about hygiene, poor sanitation personal overcrowding of patients in hospital, inadequate measures for prevention of the spread of resistant pathogens in hospital environment. Among gram negative bacteria, E. coli were the most frequent isolates 113(57.9%) .Study were similar with the study of Nazme et al. and Patel et al. but findings were dissimilar with Thaddanee et al. and Khandelwal et

al.^{3, 12, 15, 16} *E. coli* is a common cause of UTIs in children because it resides in the digestive tract and can easily migrate to the urethra and urinary tract, where it can multiply and cause infection and also due to their developing immune systems, frequent hand-tomouth contact, and exposure to contaminated environments. Among gram positive bacteria, Enterococci spp. was the most common bacterial isolates 19(9.8%). This study was similar with the study of Nazme et al. and Patel et al. Study was dissimilar with the study of Islam et al. Arora et al.3, 12, 14, 17 The isolated gram-negative bacteria were highly resistant to cefixime, ceftriaxone, cefuroxime, azithromycin, amoxiclav and ciprofloxacin. Colistin, nitrofurantoin, imipenem and levofloxacin are effective against gram negative bacteria. This study was nearly similar with Nazme et al. and Islam et al. 12, ¹⁷ The isolated gram-positive bacteria were highly resistant to cefixime, Cotrimoxazole, ceftriaxone, cefuroxime, azithromycin, amoxiclav ciprofloxacin. But relatively lower resistance was against vancomycin, observed nitrofurantoin, linezolid and imipenem. This study was nearly similar with Nazme et al. and Patel et al.3, 12 These variations may be due to differences in local conditions, prevention protocols, antibiotic policy as well as duration of study, variation in host and immune status of the host.

Ethical approval: Ethical clearance for the study was taken from the Institutional Review Board and concerned authorities, Rajshahi Medical College & Hospital.

Conflict of interest: None declared.

Consent: Informed written consent was taken from each patient or patient's attendant.

REFERENCES

1. Hellersteim S. Urinary tract infection, old and new concepts. Pediatr Clin North Am. 1995;42(6):1433-57.

- Shams F, Afroz S, Akter R, Tangim SF, Sarkar D, Paul S. Bacteriological Profile and Antibiotic Resistance Pattern in Children with Urinary Tract Infection in a Tertiary Care Hospital. Sir Salimullah Med Coll J 2021; 29: 112-116.
- 3. Patel AH, Bhavsar RH, Trivedi P, Mehta SR. Urinary Tract Infections in Children: Clinical Profile, Bacteriology and Antibiotic Sensitivity Pattern. GCSMC J Med Sci. 2015; 4(2):75-81
- 4. Patel P, Garala RN. Bacteriological profile and antibiotic susceptibility pattern (antibiogram) of urinary tract infections in paediatric patients. Journal of Research in Medical and Dental Science.2014;2(1):20-23.
- Kalal BS, Patel RB.Microbiological and Antimicrobial Profile of Urinary Tract Infection in Children from A Teaching Hospital in South India. J Pediatr Inf. 2017; 11: 19-22.
- 6. Patel JB, editor. Performance standards for antimicrobial susceptibility testing. Clinical and laboratory standards institute; 2017.
- Rafi MA, Rahman N. High Prevalence Of Multidrug Resistant Bacteria Causing Urinary Tract Infection Among Children In Northern Bangladesh.icddrb.
- 8. Nag BC, Rahman MM, Pervez M, Halder AK. Clinical and bacteriological profile of urinary tract infection in children at a tertiary care hospital. Int J Contemp Pediatr. 2022;9(1):1-5.
- 9. Venugopal P, Cherain CS, Raghunath P. Clinicoetiological profile of urinary tract infection in pediatric population in a teaching hospital in south India. Int J Contemp Pediatr. 2021;8(12):1958-1964.
- Sharmin LS, Haque MA, Begum F, Parvez KS, Md. Uddin MB. Bacteriological Profile and Antimicrobial Sensitivity Pattern of Urinary Tract Infection in a Tertiary Care Hospital of Bangladesh. TAJ. 2020; 33(2): 27-31.

- Koli G, Ninama G, Vaja K, Duttaroy B. Bacteriological Profile of Urinary Tract Infection in Paediatric Age group along with their Antimicrobial Susceptibility Pattern. International Journal of Pharmaceutical and Clinical Research. 2023; 15(7): 849-852.
- Nazme NI, Amin AA, JALILF, Sultana J, Fatema NN. Bacteriological Profile of Urinary Tract Infection in Children of a Tertiary Care Hospital. BANGLADESH J CHILD HEALTH. 2017; 41 (2): 77-83.
- 13. Nahar K, Hasan AR & Jahan NK. Bacteriological Profile and Antibiotic Sensitivity Patterns in Children with Urinary Tract Infection: A Cross-Sectional Study in the Northern Part of Bangladesh. Global Journal of Medical Research.2023;23(4):5-12.
- 14. Arora M, Gugnani P, Singh P, Neki NS. Bacteriological profile and Antibiogram of Urinary Tract Infections in children in a Tertiary Care Hospital in North India. Int. J. Curr. Res. Biol. Med. 2018; 3(3): 30-36.
- 15. Thaddanee R, Khilnani G, Shah N, Khilnani AK. Antibiotic sensitivity pattern of pathogens in children with urinary tract infection in a tertiary care hospital in Kachchh, Gujarat, India. Int J Contemp Pediatr. 2017; 4(6):2103-2108.
- 16. Khandelwal N, Sutariya D. Retrospective study of bacteriological profile and their antibiotic sensitivity pattern in urine of pediatric patients (0-12 years) in a tertiary care hospital of South Gujarat. Indian Journal of Microbiology Research. 2020;7(4):373–376.
- 17. Islam MA, Begum S, Parul SS, Bhuyian T, Islam MT, Islam MK. Antibiotic Resistance Pattern in Children with UTI: A Study in a Tertiary Care Hospital, Dhaka, Bangladesh. American Journal of Pediatrics.2019; 5(4): 191-195.