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Original Research Article

Antiboitic susceptibility pattern of common uropathogens isolated at a tertiary care hospital of Bihar

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Abstract

Introduction: Urinary tract infections (UTIs) are amongst the most common infections encountered in clinical practice. The present study was undertaken to assess the current antibiotic resistance pattern in the common uropathogens isolated in a tertiary care hospital in south India. Material & Methods: A secondary data analysis was carried out in the department of Microbiology, Jan Nayak Karpuri Thakur Medical College, Madhephura, Bihar, India, for the samples collected during January 2019 to December 2019. Samples received included mid-stream clean catch urine, suprapubic aspirate, urine collected from Foley's catheter and from the nephrostomy tube under sterile precautions, in patients who had undergone percutaneous nephrostomy. Samples were processed and isolates were identified as per standard methods. The chi-square test or Fisher's exact test was used to compare different groups. Statistical software SPSS 16.0 (SPSS Inc, Chicago, Illinois, USA) was used to analyse the data. Results: Data from a total of 1000 consecutive urine samples received in the dept during the period of data collection were included in the study. Of these, 680 (68%) were sterile, 180 (18%) showed significant growth, 20 (2%) showed insignificant growth and 120 (12%) were found contaminated. Of the 180 culture positives, E. coli was the most common isolate. The percentage of Klebsiella pneumoniae, Acinetobacter spp, Pseudomonas spp, Staphylococcus spp, Enterococcus faecalis was higher in patient females and patients with history of prior treatment with antibiotics, compared to the rest. Almost three-fourth of all E.coli isolates were found to be resistant to ciprofloxacin. Ciprofloxacin resistance was comparatively less among the other Gram-negative uropathogens like Pseudomonas spp, Proteus spp and K. pneumoniae as mentioned. Resistance to the aminoglycosides, amikacin and gentamicin was also considerable especially among isolates of Acinetobacter. Conclusion: The present results in increasing antibiotic resistance trends in UTI patients indicate that it is imperative to rationalize the use of antimicrobials and to use these conservatively.

Key Words: Antiboitic Susceptibility, Uropathogens

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Urinary tract infections (UTIs) are amongst the most common infections encountered in clinical practice[1]. The commonest bacterial agent involved in causation of UTIs is Escherichia coli, being the principal pathogen both in the community as well as in the hospital[2, 3].

The treatment of UTIs varies according to the age of the patient, sex, underlying disease, infecting agent and whether there is lower or upper urinary tract involvement. Trimethoprim/sulphamethoxazole is the recommended drug for the treatment of UTIs in settings where the prevalence of resistance is <10-20 per cent and ciprofloxacin is recommended where this resistance is >20 per cent, according to the Infectious Diseases Society of America (IDSA) guidelines[4, 5]. The other agents used in the treatment of UTI include fluoroquinolones, cephalosporins and other β -lactams with or without β -lactamase inhibitors, nitrofurantoin[4, 5]. Recently, several studies have revealed increasing trends of resistance to many antimicrobials including the fluoroquinolones[6-8]. The increase in bacterial resistance to fluoroquinolone is multifactorial[9-13]. With the increasing trend of antibiotic-resistance in E. coli, the management of urinary tract infections is likely to become complicated with limited therapeutic options.

The present study was undertaken to assess the current antibiotic

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resistance pattern in the common uropathogens isolated in a tertiary care hospital in south India.

Material & Methods

A secondary data analysis was carried out in the department of Microbiology, Jan Nayak Karpuri Thakur Medical College, Madhephura, Bihar, India, for the samples collected during January 2019 to December 2019. This was an analysis of data generated from the records of consecutive urine samples received in the laboratory during the study period. Only the initial sample of an individual received was included to avoid duplication. Analysis of the data was carried out focusing on the age, gender, whether admitted or not, whether received prior antibiotic therapy, any surgical or gynecological intervention performed in the recent past, and any history of urinary tract infection in the past. The antibiotic susceptibility data of all isolates were also reviewed and analyzed. Samples received included mid-stream clean catch urine, suprapubic aspirate, urine collected from Foley's catheter and from the nephrostomy tube under sterile precautions, in patients who had undergone percutaneous nephrostomy. Samples were processed and isolates were identified as per standard methods[14]. All urine samples were inoculated onto cysteine lactose electrolyte deficient (CLED) medium (Himedia, Mumbai, India) using a calibrated loop (volume-0.005 ml) and were incubated for 18-24 h at 37°C.Wet mount preparations were also made from all urine samples to look for pus cells and epithelial cells. Depending upon the number of the colonies grown on the CLED medium, the interpretations of urine culture were made as insignificant (<50 colonies), doubtful significance (>50 - <500 colonies) and significant (≥500 colonies) with due clinical correlation as per recommendations[14, 15]. The

Kumari V et al International Journal of Health and Clinical Research, 2022; 5(3):836-839 antibiotic susceptibility testing of the isolated bacteria was carried out by the Kirby Bauer method[15, 16].

Statistical analysis

The chi-square test or Fisher's exact test was used to compare different groups. Statistical software SPSS 16.0 (SPSS Inc, Chicago, Illinois, USA) was used to analyse the data.

Results

Data from a total of 1000 consecutive urine samples received in the dept during the period of data collection were included in the study. Of these, 680 (68%) were sterile, 180 (18%) showed significant growth, 20 (2%) showed insignificant growth and 120 (12%) were found contaminated.

Of the 180 culture positives, E. coli was the most common isolate. The percentage of Klebsiella pneumoniae, Acinetobacter spp,

Pseudomonas spp, Staphylococcus spp, Enterococcus faecalis was higher in patient females and patients with history of prior treatment with antibiotics, compared to the rest. In addition to these isolates, the percentage of Proteus spp was more in males, especially in cases with prior history of antibiotics administration. [Figure 1]

Almost three-fourth of all E.coli isolates were found to be resistant to ciprofloxacin. Ciprofloxacin resistance was comparatively less among the other Gram-negative uropathogens like Pseudomonas spp, Proteus spp and K. pneumoniae as mentioned. Resistance to the aminoglycosides, amikacin and gentamicin was also considerable especially among isolates of Acinetobacter. [Table 1] Substantial number of Staphylococcus isolates were resistant to ciprofloxacin. Resistance to nitrofurantoin was comparatively more amongst the Enterococcus spp. [Table 1]

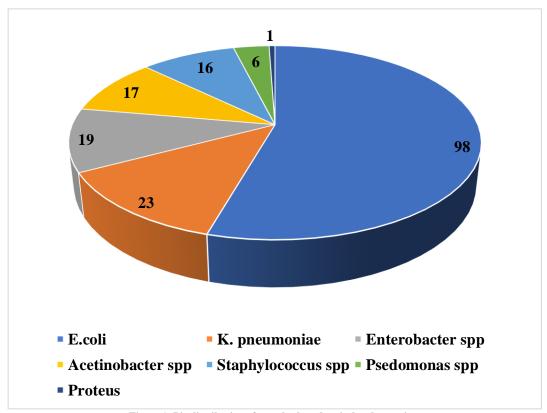


Figure 1: Pie distribution of samples based on isolated organism

Table 1: Antiboitic susceptibility pattern of isolated organism

	E.coli	K.pneumoniae	Pseudomonas	Enterobacter	Acitenobacter	Staphylococcus
Ceftriaxone	57/98	15/23	-	16/19	-	-
Ceftazidime	52/98	14/23	2/6	8/19	11/17	-
Gentamycin	56/98	17/23	3/6	11/19	13/17	-
Nitrofurantoin	25/98	16/23	-	10/19	-	7/16
Meropenam	8/98	5/23	3/6	6/19	2/17	-
Ciprofloxacin	76/98	13/23	4/6	9/19	11/17	12/16
Amikacin	19/98	7/23	2/6	5/19	10/17	-
Ampicillin	82/98	-	-	17/19	-	8/16
Vancomycin	-	-	-	-	-	1/16
Penicillin	-	-	-	-	-	14/16

Discussion

This study showed that E. coli was the commonest pathogen causing complicated and uncomplicated UTI as described previously[1-3].

There are several organisms known to cause UTIs, including P. aeruginosa, S. saprophyticus, S.epidermidis, Enterococcus spp, P. mirabilis, Klebsiella spp., Citrobacter spp, etc. as reported by earlier

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workers[17,18]. This study also demonstrates that Pseudomonas spp as major uropathogens especially in the patients admitted in the hospitals, more so in the intensive care units. Such findings have been documented elsewhere[5-16, 19-22]. The percentage of isolates of E.coli resistant to ampicillin was found to be as much as 80 per cent in our set up. Such high levels of resistance to ampicillin have been quoted by many other studies from different parts of India. Gupta et al[23] in a study from the northern part of the country reported 76 per cent resistance to E.coli isolates for ampicillin. A more recent study from Karnataka reported a resistance rate of 80.6 per cent for ampicillin[24].

The emergence of resistance for fluoroquinolones multifactorial[7,17,19,20]. Resistance to ciprofloxacin has emerged in a variety of genera of the family Enterobacteriaceae[25, 26]. Apart from the notable resistance of E. coli to ciprofloxacin, other organisms were also found to be resistant to ciprofloxacin especially K. pneumoniae, Citrobacter spp, Pseudomonas spp, Acinetobacter spp, Proteus spp and Enterobacter spp, Staphylococcus spp, and E. faecalis. Also, fluoroquinolone resistance in E. coli has emerged particularly in patients with urinary tract infections who have received fluoroquinolone prophylaxis[5-12]. An association between the increase in quinolone prescriptions and an increase in bacterial resistance has been reported from several countries[5-8, 12]. Usually, the prevalence of fluoroquinolone resistance is related to the intensity of antibiotic use[5], Resistance rates for ciprofloxacin against uncomplicated UTI pathogens were reported as 0-14.7 per cent in the ECO-SENS Project, 2.5 per cent in the USA and 1.2 per cent in outpatients in Canada[27, 28].

Conclusion

The present results in increasing antibiotic resistance trends in UTI patients indicate that it is imperative to rationalize the use of antimicrobials and to use these conservatively. [29]

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