Urinary Tract Infection among Patients Visiting Ganesh Man Singh Memorial Hospital and Research Center, Lalitpur, Nepal

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ABSTRACT

Objectives: The objective of this study was to determine the prevalence of urinary tract infection (UTI) and antibiotic sensitivity pattern among the suspected UTI cases visiting at Ganeshman Singh Memorial Hospital Lalitpur, Nepal.

Methods: A total of 300 mid-stream urine, catheter and suprapubic aspirate from UTI suspected patients were included and processed for routine microscopy and culture and then identified by standard microbiological methods. Antibiotic susceptibility test was performed by Kirby-Bauer disc diffusion method.

Results: Out of 300 samples, 55(84.6%) mid-stream urine and 10(15.4%) catheter sample had significant bacterial growth. *E. coli* (32,49.2%) was the most common isolate followed by *Staphylococcus aureus* (10,15.3%), *Enterobacter* spp. (8,12.3%), *Klebsiella* spp. (7,10.7%), *Pseudomonas aeruginosa* (3,4.6%), *Proteus* spp. (3,4.6%), *Acinetobacter* spp. (1,1.5%) and *Enterococcus* spp. (1,1.5%). Most of the Gram-negative bacterial isolates were sensitive to Ceftriaxone (88.8%) followed by Gentamicin (72.2%), and Nitrofurantoin (64.8%) and resistant to Amoxicillin (68.5%) followed by Nalidixic Acid (53.7%). Gram positive isolates were sensitive to Amikacin (72.7%) followed by Imipenem (63.6%) and Gentamicin (63.6%) whereas resistant to Amoxicillin (72.7%) and Ciprofloxacin (63.6%).

Conclusion: The main cause of the UTIs was found as Gram negative bacteria. Prescription of antibiotics based on susceptibility tests would help in reduction of antibiotic resistance.

Key words: Antimicrobial susceptibility, *Escherichia coli*, *Staphylococcus aureus*, Urinary tract infection

INTRODUCTION

Urinary tract infection (UTI) is an infection caused by the presence and growth of microorganisms anywhere in the urinary tract. Urinary Tract Infection (UTI) remains the commonest bacterial infection in human population with a high rate of morbidity and financial cost as this disease encounter with both community and hospitalized patients of all age group. In contrast to men, women are more susceptible to UTI, and this is mainly due to short urethra, absence of prostatic secretion, pregnancy and easy contamination of urinary tract with faecal flora (Haider et al. 2010).

UTI is usually classified by the infection site: bladder(cystitis), kidney(pyelonephritis), and urethra(urethritis). UTIs that occur in a normal genitourinary tract with no prior instrumentation are considered as “uncomplicated,” whereas “complicated” infection is diagnosed in genitourinary tracts that have structural or functional abnormalities, including instrumentation such as indwelling urethral catheters (Haider et al. 2010; Taher et al. 2009).

The common pathogens that cause UTI are *E. coli*, *Klebsiella* spp, *Staphylococcus* spp and other pathogens. Other pathogens include *Pseudomonas*, *Streptococcus* and MRSA. More than 95% of UTI cases are caused by bacterial pathogens, among which *E. coli*, the
Most leading causative organism, is responsible for the urinary tract infection. More than 80% of urinary tract infections get caused by *Klebsiella* spp. (Ramesh et al. 2008). Other Gram-negative bacteria that cause infection include *Enterobacter* spp., *Pseudomonas aeruginosa*, *Proteus* spp., *Citrobacter* spp., *Morganella morganii*, and so on. The total account of Gram-positive bacteria to cause urinary tract infection is 5 to 15% of the total bacteria, which include *Enterococcus* spp., *Staphylococcus*, and *Streptococcus* (Akram et al. 2007).

Majorities of UTIs are not life threatening and do not cause any serious disease to the human health. Nevertheless, when the bacterial pathogens that affect kidneys are involved, there is a risk of serious disease like tissue damage with an increased risk of bacteremia (Manikandan et al. 2011). Presence of bacteria, fungi and viruses, among others, could be involved most often to cause UTI. These bacteria enter the urethra and then travel to the bladder and kidneys (Benjamin 2009).

UTI is a common disease ailment among Nepalese population as well as one of the commonest nosocomial infection (Kattel et al. 2008). Nowadays, antimicrobial resistance is a global problem that threatens individual and social well-being. The changing patterns in the etiological agents of urinary tract pathogen and their sensitivities to commonly prescribed antibiotics are reported (Manikandan et al. 2011). The problems of antimicrobial resistance may be due to the fact that antibiotics can be obtained and used without medical authorization or supervision in developing countries (Pokhrel et al. 2006) such as Nepal.

This study was conducted to assess the bacteria causing UTI among the suspected patients visiting Ganeshman Singh Memorial Hospital and Research Center, Lalitpur, Nepal.

**MATERIALS AND METHODS**

This study was carried out among the patients visiting at Ganeshman Singh Memorial Hospital and Research Center, Lalitpur, Nepal. Three hundred samples, i.e. clinically suspected UTI defined by physician, were investigated from December 2016 to March 2017. The patients with age group ten years and more were included and mid-stream urine, catheter urine and suprapubic aspirate were collected for this study. Under macroscopic examination of urine, the specimens were observed for its colour and appearance and reported accordingly (Cheesbrough 2000).

During microscopic examination of urine, 10ml of urine sample was taken in a clean test tube and the sample was centrifuged at 3000 rpm for 5 minutes. Then the supernatant was discarded and sediment was used for the wet mount preparation to detect RBC, pus cell and epithelial cell. Culture of each urine sample was done in the MacConkey agar and blood agar medium by semi-quantitative method using standard sterile inoculating loop of standard dimension (0.001ml). The plates were then incubated at 37°C for overnight. Samples showing ≥10^5 colony forming unit (CFU) per milliliter (ml) of urine were taken as significant. Low count significant bacteriuria (10^4–10^5 CFU/ml) was taken into consideration if there was any indication which can lower the concentration of bacteria in the urine.

Identification of significant isolates was done based on morphological appearance of the colonies, Gram’s staining reactions and different biochemical reactions.

**Antibiotic susceptibility testing:** Antibiotics sensitivity testing of bacterial isolates was done by modified Kirby Bauer disc diffusion method as recommended by CLSI (2014) on MHA.

**Statistical analysis:** Data were entered into SPSS version 19.0 and analyzed for descriptive statistics.

**RESULTS**

Among total samples received in laboratory for culture, 269 (89.7%) samples were mid-stream urine, 30 (10%) samples were catheter and remaining 1 (0.3%) samples were suprapubic aspirate.

Out of 300 samples, 65 (21.6%) samples had significant growth. Among 65 samples, 55 (84.6%) were mid-stream urine samples (MSU) and 10 (15.3%) were catheter samples. Out of 201 samples from outdoor patients, 45 (22.3%) samples and out of 99 samples from indoor patients, 20 (20.2%) samples had significant growth respectively. Among 138 samples from male patients, 26 (18.8%) samples had significant growth. Similarly, 39 (24.1%) out of 162 samples from female patients showed significant growth. Among the 65 significant growth cultures, high percentage (35.8%) was obtained from age group 31-40 years.

**Bacterial isolates causing UTI**

Among the bacterial isolates, *E. coli* (49.2%) was found to be the most predominant organism followed by *Staphylococcus aureus* (15.3%), *Klebsiella* spp. (10.7%) and others (24.6%).
Antibiotic susceptibility profile

More proportion of *E. coli* was found to be sensitive towards Ceftriaxone (93.5%) followed by Nitrofurantoin (87.5%), Gentamicin (78.1%), Ciprofloxacin and Norfloxacin (71.8%), Imipenem and Chloramphenicol (65.6%), Cotrimoxazole (59.3%), and Cefixime (56.2%). *E. coli* was found resistant towards Nalidixic acid (84.3%) followed by Amoxycillin (75%) and Amikacin (50%).

*S. aureus* was found sensitive towards Amikacin (80%) followed by Imipenem and Gentamicin (70%) whereas it was found resistant to Amoxycillin (70%). All *Enterococcus* isolates were found sensitive towards Cotrimoxazole and Nitrofurantoin.

*Klebsiella* spp was found sensitive towards Imipenem and Ceftriaxone (71.4%) followed by Ciprofloxacin, Gentamicin (57.1%) whereas it was found resistant towards Amoxycillin (85.7%) followed by Norfloxacin (71.4%), Cotrimoxazole (57.1%) and Amikacin (57.1%).

*Pseudomonas aeruginosa* was sensitive towards Gentamicin whereas it was resistant towards Levofloxacin (66.6%) followed by Amoxycillin, Norfloxacin and Piperacillin (33.3%).

*Acinetobacter* spp. were sensitive to Norfloxacin, Ciprofloxacin and Ceftriaxone whereas were resistant towards Cotrimoxazole, Nitrofurantoin, Amoxycillin and Nalidixic acid.

Table 1: Antibiotic resistance pattern of isolated bacteria in percentage

<table>
<thead>
<tr>
<th>Bacteria</th>
<th>Ceftriaxone</th>
<th>Nitrofurantoin</th>
<th>Gentamicin</th>
<th>Ciprofloxacin</th>
<th>Norfloxacin</th>
<th>Imipenem</th>
<th>Chloramphenicol</th>
<th>Cotrimoxazole</th>
<th>Cefixime</th>
<th>Levofloxacin</th>
<th>Amoxicilin</th>
<th>Nalidixic Acid</th>
<th>Amikacin</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>E. coli</em></td>
<td>6.2</td>
<td>12.5</td>
<td>21.8</td>
<td>28.1</td>
<td>28.1</td>
<td>4.3</td>
<td>34.3</td>
<td>40.6</td>
<td>42.7</td>
<td>46.6</td>
<td>75</td>
<td>84.3</td>
<td>50</td>
</tr>
<tr>
<td><em>S. aureus</em></td>
<td>-</td>
<td>40</td>
<td>30</td>
<td>-</td>
<td>-</td>
<td>30</td>
<td>-</td>
<td>50</td>
<td>-</td>
<td>-</td>
<td>70</td>
<td>20</td>
<td></td>
</tr>
<tr>
<td>Enterococcus spp</td>
<td>-</td>
<td>0</td>
<td>100</td>
<td>-</td>
<td>100</td>
<td>0</td>
<td>-</td>
<td>100</td>
<td>-</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td></td>
</tr>
<tr>
<td>Klebsiella spp</td>
<td>28.5</td>
<td>-</td>
<td>42.8</td>
<td>42.8</td>
<td>71.4</td>
<td>28.5</td>
<td>-</td>
<td>57.1</td>
<td>-</td>
<td>85.7</td>
<td>-</td>
<td>57.1</td>
<td></td>
</tr>
<tr>
<td><em>P. aeruginosa</em></td>
<td>-</td>
<td>33.3</td>
<td>-</td>
<td>33.3</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>66.6</td>
<td>33.3</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>Acinetobacter spp</td>
<td>0</td>
<td>100</td>
<td>0</td>
<td>0</td>
<td>100</td>
<td>0</td>
<td>-</td>
<td>100</td>
<td>-</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td></td>
</tr>
<tr>
<td>Enterobacter spp</td>
<td>0</td>
<td>37.5</td>
<td>37.5</td>
<td>-</td>
<td>37.5</td>
<td>12.5</td>
<td>-</td>
<td>12.5</td>
<td>75</td>
<td>-</td>
<td>50</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>Proteus spp</td>
<td>33.3</td>
<td>33.3</td>
<td>33.3</td>
<td>66.6</td>
<td>0</td>
<td>-</td>
<td>-</td>
<td>66.6</td>
<td>-</td>
<td>-</td>
<td>33.3</td>
<td>33.3</td>
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</tr>
</tbody>
</table>
DISCUSSION
Age group 31-40 years had got the high prevalence of UTI which is in contradictory to the study by Daniyan and Ojha (2013). More than one third patients of the total UTI positive cases were found in this age group which is similar to the study of Leigh (1990). Leigh (1990) had found that nuns and unmarried women have lower prevalence of UTI in compared to married women. In addition, use of spermicidal coated condoms dramatically alters the normal bacterial flora and has been associated with marked increase in vaginal colonization with *E. coli* and in the risk of UTI (Braunwald et al. 2001).

The majority cases were from outpatient department and 33% were from patients admitted to the hospital. In comparison to males, higher number of urine samples was collected from females which is also similar to other studies (Arjunan et al. 2010; Alzohairy and Khadri 2011). The increased incidence of the urinary tract infection in women is conditioned by favoring anatomic factors, by hormonal changes and by the urodynamic disturbance occurring with age (Bobos et al. 2010).

In this study, 21.6% urine specimens from suspected UTI patients gave significant growth. Similar results were reported by other investigators from Nepal (Chhetri et al. 2001; Kumari et al. 2005) and rest of the world (Bashir et al. 2009). Among the bacterial isolates, *E. coli* was found to be the most predominant organism followed by *S. aureus*, *Klebsiella* spp. and other which is similar to the study of Kattel et al. (2008). However, Aboderin et al. (2009) reported *Pseudomonas aeruginosa* and *Klebsiella* spp respectively as the predominant bacteria.

Majority of *E. coli* were isolated from outpatient department. *E. coli* accounts for 50% - 90% of all the uncomplicated urinary tract infections (Vgaardali et al. 2008). In our study, *Staphylococcus aureus* was found the second most common isolates. UTI due to *Enterococcus faecalis* are usually associated with the use of instrument or catheterization (Collier et al. 1998).

CONCLUSION
The main cause of the UTIs is Gram negative bacteria. Gentamicin and Ceftriaxone may be drug of choice for Gram negative bacteria whereas Amikacin and Ciprofloxacin may be drug of choice for Gram positive bacteria.

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CONFLICT OF INTEREST
The authors declare no conflict of interest.

REFERENCES


