

# Frequency and susceptibility profile of pathogens causing urinary tract infections at a tertiary care hospital in western Nepal

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## ABSTRACT

**Introduction:** Urinary tract infection (UTI) is the most common nosocomial infection among hospitalised patients. Area-specific monitoring studies aimed to gain knowledge about the type of pathogens responsible for UTIs and their resistance patterns may help the clinician to choose the correct empirical treatment. Recent reports have shown increasing resistance to commonly-used antibiotics. We aimed to study the antibiotic resistance pattern of the urinary pathogens isolated from hospitalised patients.

**Methods:** Three urine samples were collected by the mid-stream "clean catch" method from 1,680 clinically-suspected cases of urinary tract infections from inpatients of various clinical departments during one year. The samples were tested microbiologically by standard procedures. Antibiotic susceptibility of the isolated pathogens was tested for commonly-used antibiotics by Kirby-Bauer technique according to NCCLS guidelines.

**Results:** Significant bacteriuria was present in 71.7 percent of the samples, 17 percent were sterile, 4.8 percent showed insignificant bacteriuria, and 6.5 percent non-pathogenic bacteriuria. The most common pathogens isolated were *Escherichia coli* (59.4 percent), *Klebsiella* spp (15.7 percent) and *Enterococcus faecalis* (8.1 percent). The mean susceptibility was high for amikacin (87.2 percent), ciprofloxacin (74.8 percent), ceftazidime (71.5 percent) and gentamicin (70.4 percent) but low for nitrofurantoin (35 percent), cephalexin (49.7 percent) and ampicillin (50.5 percent). *Escherichia coli* was found to be most susceptible to amikacin (98 percent) followed by gentamicin (87.9 percent), ceftazidime (80.8 percent), norfloxacin (78.4 percent) and cotrimoxazole (77.9 percent).

**Conclusion:** A high isolation rate of pathogens from urine samples of clinically-suspected UTI shows a good correlation between clinical findings and microbiological methods. The antibiotics commonly used in UTIs are less effective. Since the present study was a cross-sectional study, regular monitoring is required to establish reliable information about resistance pattern of urinary pathogens for optimal empirical therapy of patients with nosocomial UTIs.

**Keywords:** antibiotics, bacteriuria, nosocomial infection, urinary tract infection

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## INTRODUCTION

Urinary tract infection (UTI) is one of the most important causes of morbidity in the general population, and is the second most common cause of hospital visits<sup>(1)</sup>. With advancing age, the incidence of UTI increases in men due to prostate enlargement and neurogenic bladder<sup>(2)</sup>. Recurrent infections are common and can lead to irreversible damage of the kidneys, resulting in renal hypertension and renal failure in severe cases<sup>(3)</sup>. In the community, women are more prone to develop UTI. About 20% of women experience a single episode of UTI during their lifetime, and 3% of women have more than one episode of UTI per year<sup>(4)</sup>. Pregnancy also makes them more susceptible to infection<sup>(5)</sup>. Catheter-associated UTI is a trenchant problem, with about 5% of catheterised patients developing bacteriuria<sup>(6)</sup>. It is universally accepted that UTI can only be ascertained on the basis of microscopy and microbial culture. The dipstick/dip-slide method used in many centres serves only as a screening method but culture is needed for the final diagnosis<sup>(7)</sup>.

UTI is the most common cause of nosocomial infection among hospitalised patients<sup>(1)</sup>. In almost all cases, there is a need to start treatment before the final microbiological results are available.

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Area-specific monitoring studies aimed to gain knowledge about the type of pathogens responsible for UTIs and their resistance patterns may help the clinician to choose the correct empirical treatment. Studies from the eastern part of Nepal, India and Bangladesh have reported an increased resistance of the urinary pathogens to commonly-used antibiotics<sup>(8-10)</sup>. Such information was not available for a tertiary care hospital in western Nepal. Hence this study was undertaken to find out the frequency and antibiotic susceptibility pattern of the urinary pathogens isolated from the urine samples of suspected nosocomial urinary tract infections at a tertiary care hospital in western Nepal.

## METHODS

The present study was conducted at the Manipal Teaching Hospital of Manipal College of Medical Sciences, a 750-bed tertiary care hospital located at Pokhara city in western Nepal. All the urine samples received from the inpatients of various clinical departments of Manipal Teaching Hospital from August 1, 2003 to July 30, 2004 were selected for the study. If there were more than two episodes of UTI for the same patient, either due to prolonged hospitalisation or repeated hospitalisations, each episode was considered as a separate case of UTI. The patients who had symptoms and/or signs suggestive of UTI were suspected as having UTI.

Three urine samples were collected by standard mid-stream "clean catch" method from all the patients with suspected urinary tract infection. Three urine samples were also collected from catheterised patients, if the catheter was not inserted for not longer than 72 hours. The samples collected were examined microscopically for pus cells and casts. The samples were also processed using standard microbiological procedures. All the urine samples were processed on the blood agar and MacConkey's medium by standard loop method and incubated at 37°C overnight. The plates were observed for bacterial growth.

Culture results were interpreted as being significant and insignificant, according to the standard criteria. A growth of  $\geq 10^5$  colony forming units/mL was considered as significant bacteriuria<sup>(11)</sup>. Patients with significant bacteriuria and symptomatic patients with lower colony counts were also considered as having UTI. The organism was identified by routine methods from the samples showing significant bacteriuria<sup>(12)</sup>. Cultures with more than three colonies were discarded, as contaminants and their antibiotic susceptibility were not tested.

Antibiotic susceptibility tests and interpretations were carried out for bacterial isolates by the Kirby-Bauer technique<sup>(13)</sup>. Antibiotic susceptibility was tested by disc diffusion method for all the first and the second line antibiotics, according to the hospital antibiotic policy, irrespective of the isolated organism showing susceptibility/resistance to the first line antibiotics.

The first line antibiotics tested were: ampicillin, oxacillin, cephalexin, nitrofurantoin, cotrimoxazole, gentamicin, amikacin, nalidixic acid and norfloxacin, and the second line were: ciprofloxacin, ceftazidime and piperacillin (Hi-Media Pvt. Laboratories, Bombay, India). Appropriate quality control strains were used to validate the results of the antimicrobial discs. The following were the quality control strains used: *Pseudomonas* spp-NCTC 10662, *Staphylococcus aureus*-NCTC 6571, *Escherichia coli* (*E. coli*)-NCTC 10418, *Enterococcus*-NCTC 12697<sup>(14)</sup>. Guidelines of the National Committee for Clinical Laboratory Standards (NCCLS) were followed for the testing and interpreting the antimicrobial disks. Data was entered into Microsoft Excel and analysed. The results were expressed as proportions and mean susceptibility was calculated for first line and second line antibiotics for each organism isolated by culture.

## RESULTS

The age-sex distribution of the samples, according to the culture positive cases, is shown in Table I. Out of the 1,291 culture positive urine samples, 109 (8.4%) were from children aged ten years or less, and 277 (21.4%) were from those aged 60 years and above. In all age groups, except those aged 0-10 years, females were more frequently affected than males. 389 (23.2%) out of 1,680 urine samples examined did not yield any organism. There were 64 urine samples obtained from catheterised patients and 1,141 urine samples collected from non-catheterised samples. Out of the 1,680 samples processed, 1,205 (71.7%) showed significant bacteriuria, 286 (17%) were sterile, 80 (4.8%) showed insignificant bacteriuria, and 109 (6.5%) non-pathogenic bacteriuria. *E. coli* and *Klebsiella* spp were commonest organisms isolated from both catheterised and non-catheterised samples (Table II).

The organisms grown on the culture of all the 1,205 samples with significant bacteriuria were as follows: *E. coli* 716 (59.4%), *Klebsiella* spp 189 (15.7%), *Enterococcus faecalis* 98 (8.1%), *Proteus mirabilis* 89 (7.4%), *Staphylococcus aureus* 41 (3.4%), *Pseudomonas aeruginosa* 35 (2.9%), *Staphylococcus*

**Table I. Age and sex distribution of the culture-positive urine samples.**

| Age (in years) | Gender      |             | Total        |
|----------------|-------------|-------------|--------------|
|                | Male        | Female      |              |
| 0-10           | 68          | 41          | 109 (8.4%)   |
| 11-20          | 108         | 122         | 230 (17.8%)  |
| 21-40          | 81          | 189         | 270 (20.9%)  |
| 41-60          | 95          | 310         | 405 (31.4%)  |
| Above 60       | 135         | 142         | 277 (21.4%)  |
| Total          | 487 (37.7%) | 804 (62.3%) | 1,291 (100%) |

**Table II. Bacteria isolated from urine samples of catheterised and non-catheterised patients.**

| Organism                            | Catheterised | Non-catheterised | Total       |
|-------------------------------------|--------------|------------------|-------------|
| <i>E. coli</i>                      | 31 (48.4%)   | 685 (60%)        | 716 (59.4%) |
| <i>Klebsiella spp</i>               | 20 (31.2%)   | 169 (14.8%)      | 189 (15.7%) |
| <i>Enterococcus faecalis</i>        | 1 (1.5%)     | 34 (2.9%)        | 98 (8.1%)   |
| <i>Proteus mirabilis</i>            | 9 (14%)      | 80 (7%)          | 89 (7.4%)   |
| <i>Staphylococcus aureus</i>        | 1 (1.5%)     | 97 (8.5%)        | 41 (3.4%)   |
| <i>Pseudomonas aeruginosa</i>       | 2 (3.1%)     | 39 (3.4%)        | 35 (2.9%)   |
| <i>Staphylococcus saprophyticus</i> | –            | 26 (2.2%)        | 26 (2.1%)   |
| <i>Citrobacter freundii</i>         | –            | 11 (0.9%)        | 11 (0.9%)   |
| Total                               | 64           | 1,141            | 1,205       |

*saprophyticus* 26 (2.1%) and *Citrobacter freundii* 11 (0.9%) (Table II).

The antibiogram of isolated pathogens is shown in Table III. The highest susceptibility was for amikacin (87.2%), ciprofloxacin (74.8), ceftazidime (71.5%), and gentamicin (70.4%) for all the pathogens isolated; and the least was for nitrofurantoin (35%), cotrimoxazole (45.8), cephalixin (49.7%) and ampicillin (50.5%). *E. coli* was most sensitive to amikacin (98%), followed by gentamicin (87.9%), ceftazidime (80.8%), norfloxacin (78.4%) and cotrimoxazole (77.9%).

*Klebsiella spp* was most susceptible to ceftazidime (96.8%), norfloxacin (76.2%) and ciprofloxacin (95.7%). *Pseudomonas aeruginosa* was most sensitive to piperacillin (100%) amikacin (100%) ciprofloxacin (98.2%) and ceftazidime (97.3%). *Staphylococcus aureus* was most sensitive to amikacin (100%) and ciprofloxacin (85.7%) but least susceptibility was observed for cotrimoxazole (12.5%) and ampicillin (13.8%). Since *Staphylococcus saprophyticus* was not a common cause of UTI, antibiotic susceptibility was not tested.

## DISCUSSION

The age and sex distribution of the patients diagnosed with UTI among the hospitalised patients followed the natural epidemiological pattern of UTI. There was a higher number of young male children. Similarly, there was a greater predominance of young and middle-aged females, whereas in the older age group ( $\geq 60$  years), equal proportions of

**Table III. Bacteria isolated from all the urine samples and their susceptibility pattern to common antibiotics.**

| Organism isolated       | No (%)<br>n=1205 | Antibiotic susceptibility (%) |      |      |      |      |      |      |      |      |      |      |     |
|-------------------------|------------------|-------------------------------|------|------|------|------|------|------|------|------|------|------|-----|
|                         |                  | Amp                           | Cpx  | Nfx  | Gm   | Amk  | Oxc  | Nfn  | NA   | Cmz  | Cfz  | Cfx  | Ppc |
| <i>E. coli</i>          | 716 (59.4)       | 72.6                          | 65.9 | 78.4 | 87.9 | 98.0 | –    | 68.4 | 69.9 | 77.9 | 80.8 | 55.0 | –   |
| <i>Klebsiella spp.</i>  | 189 (15.6)       | 33.8                          | 11.1 | 76.2 | 60.8 | 61.3 | –    | 2.1  | 51.8 | 28.5 | 96.8 | 95.7 | –   |
| <i>P. mirabilis</i>     | 89 (7.3)         | 57.1                          | 50.0 | 62.9 | 83.3 | 87.6 | –    | 13.4 | 68.5 | 56.1 | –    | 85.7 | –   |
| <i>P. aeruginosa</i>    | 41 (3.4)         | 50.0                          | 75.0 | –    | 88.8 | 100  | –    | –    | –    | –    | 97.3 | 98.2 | 100 |
| <i>S. aureus</i>        | 98 (8.1)         | 13.8                          | 73.6 | –    | 64.5 | 100  | 96.2 | 66.3 | –    | 12.5 | –    | 85.7 | –   |
| <i>S. faecalis</i>      | 35 (2.9)         | 58.2                          | 11.2 | 8.2  | 14.2 | 68.6 | –    | –    | 9.1  | 60.0 | 11.2 | 10.5 | –   |
| <i>S. saprophyticus</i> | 26 (2.1)         | –                             | –    | –    | –    | –    | –    | –    | –    | –    | –    | –    | –   |
| <i>Citrobacter</i>      | 11 (0.9)         | 68.1                          | 61.3 | –    | 93.5 | 94.8 | –    | 24.8 | 80.3 | 39.9 | –    | 92.8 | –   |
| Mean susceptibility     |                  | 50.5                          | 49.7 | 56.4 | 70.4 | 87.2 | 96.2 | 35.0 | 55.9 | 45.8 | 71.5 | 74.8 | 100 |

Antibiotic susceptibility test was not done for *Staphylococcus saprophyticus*.

First line antibiotics: Amp: Ampicillin; Cpx: Cephalixin; Gm: Gentamicin; Amk: Amikacin; Oxc: Oxacillin; Nfx: Norfloxacin; NA: Nalidixic Acid; Nfn: Nitrofurantoin; Cmz: Cotrimoxazole

Second line antibiotics: Cfx: Ciprofloxacin; Ppc: Piperacillin; Cfz: Ceftazidime

Mean susceptibility = Sum of susceptibility of individual organisms/total no of organisms.

males and females had UTI. Diagnosis of UTI is a good example of the need for close cooperation between the clinician and the microbiologist. The presence of significant bacteriuria in 71.6% of the samples shows a good correlation between clinical and bacteriological diagnosis, and also indicates the significance of microbiological culture to clinch the diagnosis of UTI.

In the present study, insignificant bacteriuria and sterile cultures were observed in 4.8% and 17%, respectively. Prior antibiotic treatment before submitting the urine sample, and clinical conditions such as non-gonococcal urethritis or other clinical conditions that mimic UTI could be the factors responsible for such results. Furthermore, 6.5% of the urine samples showed presence of non-pathogenic bacteria. This indicates the need for educating the patients about the method collection of "clean catch" urine specimens.

*Enterobacteriaceae* have several factors responsible for their attachment to the uroepithelium. These gram-negative aerobic bacteria colonise the urogenital mucosa with adhesin, pili, fimbriae and P1-blood group phenotype receptor<sup>(15)</sup>. In the present study, the *Enterobacteriaceae* group, namely, *E. coli* (59.4%), *Klebsiella* spp. (15.7%), *Enterococcus faecalis* (8.1%), and *Proteus mirabilis* (7.4%), were the most common pathogens isolated, followed by gram-positive cocci, namely, *Staphylococcus aureus* (3.4%) and *Staphylococcus saprophyticus* (1.4%). The isolation rates of urinary pathogens are consistent with reports of the recently-published studies.<sup>(9, 16, 17)</sup>

Isolation rate for *Staphylococcus aureus*, which is a rare cause of UTI, was 3.4%. However, it is interesting to note that only one case occurred out of 64 catheterised patients, whereas the rate was 8.1% for non-catheterised patients. In our study sample, there were 321 patients who were diagnosed to have either renal calculus, hydronephrosis who could have had prior catheterisation, or may even be suffering from bacteraemic kidney infection. Hence, there could have been high isolation of *Staphylococcus aureus* from the cases of nosocomial UTI in our study.

*E. coli*, which was the principal pathogen isolated, showed high susceptibility to common antibiotics ampicillin (72.6%), cotrimoxazole (77.9%) and norfloxacin (78.4%). This is contrary to the reports of studies from other countries which have reported an increasing resistance to ampicillin<sup>(9,17-19)</sup>. A study from eastern Nepal reported a high prevalence of resistance to nalidixic acid, ampicillin and norfloxacin<sup>(8)</sup>. Another study from Bangladesh reported an

increased resistance of uropathogens to ciprofloxacin<sup>(10)</sup>. Ciprofloxacin and norfloxacin, which are used equally for UTI caused by *Klebsiella* spp has demonstrated increased resistance to both antibiotics and also sometimes cross-resistance exists. But in the present study, *Klebsiella* spp showed high susceptibility to both ciprofloxacin and norfloxacin. This is contrary to the results of the above-mentioned studies which reported an increased resistance of the uropathogens to both ciprofloxacin and norfloxacin.

*Pseudomonas aeruginosa*, which is a common cause of hospital-acquired UTI, was less sensitive to the common antibiotics but highly sensitive to amikacin, piperacillin, ciprofloxacin and gentamicin. Similar results are reported from other countries<sup>(17,20)</sup>. *Pseudomonas* was susceptible to the second line drugs and most of these are associated with high resistance to the first line antibiotics used, namely, ampicillin and ciprofloxacin. This may be due to widespread clinical use of common antibiotics in the hospital and cross-resistance existing among various classes of antibiotics. Even though *Pseudomonas aeruginosa* is known to be resistant to ampicillin, and first and second generation cephalosporins, we tested the susceptibility of *Pseudomonas* to these antibiotics to determine its susceptibility pattern in our setting.

The highest resistance was shown by *Klebsiella* spp which was the second most common organism isolated from the UTIs. *Klebsiella* spp was resistant to commonly-used antibiotics, except ceftazidime (96.8%) which was the second line antibiotic according to hospital antibiotic policy. A similar pattern was observed for *Proteus mirabilis*, which was susceptible to amikacin, ciprofloxacin and gentamicin (87.6%, 85.7 % and 83.3%, respectively). The gram-positive organisms isolated from urinary samples also showed an increased resistance against common antibiotics except for the *Staphylococcus aureus* which showed high susceptibility to oxacillin (96.2%).

In the present study, among the cephalosporins, cephalexin showed low mean susceptibility (49.7%) but ceftazidime showed high mean susceptibility (71.5%). *Enterococcus* showed low susceptibility to cephalexin (11.2%) and ceftazidime (11.2%). A report from the SENTRY Antimicrobial Surveillance Program, USA has shown similar results<sup>(17)</sup>. Various *Citrobacter* spp isolated might have had inducible beta-lactamases which was not tested in the present study. The results of the present study showed that the mean susceptibility of the uropathogens was low for cotrimoxazole

which often is one of the first line drugs in many countries. This low susceptibility could be due to widespread use of this antibiotic in the community. It is possible that low susceptibility is prevalent in the nosocomial UTI as well as community-acquired UTI. Some of the hospitalised patients may be having community-acquired UTI. In our study, we could not distinguish between community-acquired UTI and nosocomial UTI. This was the main limitation of the study.

In conclusion, a high isolation rate of pathogens from urine samples of clinically-suspected UTI shows a good correlation between clinical findings and microbiological methods. Gram-negative organisms were the commonest organisms isolated, among which *E. coli* was the principal urinary pathogen. Data presented in this study indicate that antibiotics commonly used for the treatment of nosocomial UTIs are less effective. Since this was a cross-sectional study, further regular monitoring is required to establish reliable information about resistance pattern of urinary pathogens for optimal empirical therapy of patients with nosocomial UTI.

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