Antibiotic Susceptibility of Bacterial Strains Isolated From Urinary Tract Infections in Karaj, Iran

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ARTICLE INFO

Article type: Original Article
Article history:
Received: 13 Mar 2012
Revised: 07 May 2012
Accepted: 21 May 2012

Keywords:
Urinary Tract Infections
Antibiotic Susceptibility

ABSTRACT

Backgrounds: Urinary Tract Infection (UTI) is one of the most common bacterial diseases worldwide that can be presented asymptomatic or symptomatic, characterized by a wide range of symptoms from mild irritative voiding to bacteremia, sepsis, or even death. In latter decades, widespread utilization of antibiotics has resulted in increased incidence of antibiotic resistance among urinary tract pathogens all over the world.

Objectives: The aim of this study was to determine patterns of antibiotic resistance bacteria isolated from urinary tract infections in the city of Karaj, Iran.

Materials and Methods: This descriptive study was conducted in the city of Karaj from November 2009 to August 2010. Bacterial isolates were identified by standard biochemical tests. Antibacterial susceptibility test was performed by Kirby-Bauer method against some common antibiotics for Gram negative and Gram positive bacteria and the data were analyzed by SPSS software, version 12.0.1.

Results: In this study, 230 (8.06%) out of 2852 patients showed positive urine cultures of which there were 204 (88.69%) females and 26 (11.3%) males. 180 (83.17%) cases of isolated bacteria were Gram negative bacilli while 50 (21.73%) cases were Gram positive cocci. E. coli isolates, as the most frequent Gram negative bacteria, showed high resistance to cephalothin (88.16%) and high susceptibility to nitrofurantoin (90.78%). Also Staphylococcus saprophyticus isolates, as the most frequent Gram positive bacteria, exhibited high resistance to ampicillin, tetracycline, and erythromycin (92.31%) and high susceptibility to nitrofurantoin and vancomycin (92.3%).

Conclusions: considering the results of this study, E.coli isolates were the predominant pathogens in UTI. Our data demonstrated that antibiotics with the highest activity included ciprofloxacin and nitrofurantoin and those with the least activity included ampicillin and sulfamethoxazole-trimethoprim. Our findings reinforce the need for ongoing investigations to show the trends of antibiotic resistance which can help antibiotic prescription in the clinics.

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Implication for health policy/practice/research/medical education:
The article will help Clinicians in predicting and planning correct strategies for UTI therapy.

Please cite this paper:

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DOI: 10.5812/jjm.4830
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1. Background

Urinary tract infections (UTIs) are among the most common bacterial diseases worldwide (1), and the commonest genitourinary disease in children (2). UTI can be asymptomatic or symptomatic, characterized by a wide spectrum of symptoms ranging from mild irritative voiding to bacteremia, sepsis, or even death (3). UTIs account for approximately 11.3 million office visits with an overall cost of 1.6 billion dollars annually in the US health budget (4). Infection may occur at any part of the genitourinary tract, including urethra, bladder, ureter, renal pelvis, or renal parenchyma (5, 6). Most infections are caused by retrograde ascent of bacteria from fecal flora to bladder and kidney via urethra, especially in females whom the urethra is shorter and wider (7).

Anatomical structure of female urethra and vagina make them susceptible to trauma during sexual intercourse and bacterial passage through urethra up to the bladder during pregnancy and delivery (8, 9). The infection may involve only lower or both upper and lower parts of urinary tract. The term “cystitis” has been used to describe a syndrome including dysuria, frequency, and occasionally suprapubic tenderness. “Acute pyelonephritis” describes a clinical syndrome characterized by flank pain or tenderness, or both, and fever, often accompanied by dysuria, urgency, and frequency (10). Majority of UTIs are not life threatening and do not cause any irreversible damage. However, when the kidneys are involved, there is a risk of irreparable tissue damage with an increased risk of bacteremia (11).

Bacteria are the major causative organism and are responsible for more than 95% of UTI cases (12). Escherichia coli is the most frequent infecting organism in acute infections (13, 14). Area-specific monitoring studies aimed to understanding the types of pathogens responsible for UTIs and their resistance patterns may help clinicians to select proper empirical treatment (15).

2. Objectives

The aims of this study were to investigate microbial species isolated from patients with UTI and evaluate their in vitro susceptibility patterns to commonly used antimicrobial agents in Karaj, Iran.

3. Materials and Methods

3.1. Specimen Collection and Bacterial Isolates

This study was conducted from November 2009 to August 2010 on 2852 untreated patients with clinical symptoms of UTI referred to a central diagnostic lab in Karaj, Iran. Clean catch midstream urine samples (MSU) were collected in sterile disposable containers (4-5 ml) and transported immediately to the laboratory. Urine specimens were subjected to general urine examinations using direct microscopy for white blood cell (WBC) counting. Urine samples were cultured on 5% blood agar, MacConkey, and Eosin-Methylene blue (EMB) agar (Oxoid Ltd, Basingstore, Hampshire, UK) using calibrated loops for semi-quantitative method (16) and incubated in both aerobic and anaerobic conditions for 24 hours at 37°C. Cultures without any colony at the end of 24h incubation were further incubated for 48h. Samples with colony count equal or more than 10^5 CFU/ml were considered positive. The isolates were identified and confirmed using standard microbiological methods including Gram staining, colonial morphology on media, growth on selective media, lactose and mannitol fermentation, H2S production, catalase, oxidase, coagulase, indole, and citrate utilization, and urease test (16-18).

3.2. Antimicrobial Susceptibility Testing

Antimicrobial susceptibility testing was performed on Mueller-Hinton agar (Merck, Germany) using disk diffusion (Kirby Bauer’s) technique according to Clinical and Laboratory Standards Institute (CLSI) guidelines. The antibiotic discs and their concentrations consisted of Cefoxime (CTX, 30μg, Polfa, Tarchomin, Poland), Cephalothin (CF, 30μg, Polfa, Tarchomin, Poland), Imipenem (IPM, 10μg, Zeneca, Macclesfield, UK), Ciprofloxacin (CP, 5μg), Ceftazidime (CAZ, 30μg), Ceftizoxime (CT, 30μg), Gentamicin (GM, 10μg), Ampicillin (AM, 10μg), Trimethoprim-sulfamethoxazole (SXT, 30μg), Nitrofurantoin (FM, 50μg) and, Nalidixic acid (NA, 30μg) for Gram negative isolates, and Chloramphenicol (C, 30μg, Bayer, Wuppertal, Germany), Ampicillin (AM, 10μg), Erythromycin (E, 15μg), Norfloxacin (NOR, 10μg), Gentamicin (GM, 10μg), Vancomycin (V, 30μg), Nitrofurantoin (FM, 50μg), Tetracycline (TE, 30μg), and Trimethoprim-sulfamethoxazole (SXT, 30μg) for Gram positive isolates all obtained from Hi media, India. E. coli (ATCC 25922), Staphylococcus aureus (ATCC25923), and Pseudomonas aeruginosa (ATCC 27853) were used as quality control strains.

Data were analyzed using SPSS version 12.0.1. Discrete variables were expressed as percentages and proportions were compared using the Chi-square test. Statistical significant difference was considered at value of P ≤ 0.05.

4. Results

In this study, 230 (8.06%) out of 2852 patients showed positive urine cultures of which there were 204 (88.69%) females and 26 (11.3%) males. 180 (83.17%) cases of the isolated bacteria were Gram negative bacilli while 50 (21.73%) cases were Gram positive cocci. The frequency of isolated microorganisms and their relations to sex is given in Table 1.

Patients’ age ranged from 10 days to 90 years with an average age of 26.17 years. The age distribution was 21.73% cases were Gram positive cocci. The frequency of isolated microorganisms and their relations to sex is given in Table 1.
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22.6% for patients aged ≤ 2 year, 9.56% for 2 to 14 years, 57.39% for 15 to 60 years, and 9.56% for ≥ 60 years. *E. coli* was the most common microorganism isolated in all age-groups and both sexes. No significant difference was found between isolated bacteria and age of the patients. The isolation percentages of bacterial species were found as *E. coli* (66.08%); *S. saprophyticus* (11.3%); *Klebsiella* spp. (9.6%); *S. group B* (4.34%); *Enterococcus* spp. (4.34%); *S. aureus* (1.7%); *Proteus* spp. (1.7%); and *Enterobacter* spp. (0.86%). *E. coli* isolates, the predominant cause of UTIs, showed high resistance to gentamicin, ampicillin, SXT, and cephalothin, and higher sensitivity to imipenem, ciprofloxacin, nitrofurantoin, and ceftizoxime. *Klebsiella* strains displayed a similar resistance pattern as for *E. coli* and showed high susceptibility to ciprofloxacin, ceftizoxime, nalidixic acid, and imipenem and high resistance to gentamicin, cephaptoxin, and ampicillin. On *Proteus* strains, ampicillin, nitrofurantoin, and SXT showed the least (0%, 0%, and 25%, respectively) and ciprofloxacin, ceftizoxime, and imipenem showed the highest susceptibility (100%, 100% and 75%, respectively) that were almost similar to the results with other Gram positive bacteria. The results were slightly different in *Enterobacter* as it can be seen the utmost resistance of *Enterobacter* was recorded to ampicillin and nalidixic acid while 100% susceptibility to other antibiotics was recorded.

In this study, staphylococci were responsible for about 13% of UTIs cases; among these, *S. saprophyticus* isolates as most frequent Gram positive bacteria showed high resistance to ampicillin, tetracycline, and erythromycin, and high sensitivity to nitrofurantoin, gentamicin, norfloxacin, and vancomycin. The rates of resistance to 11 selected antimicrobial agents against Gram negative bacilli and to 9 selected antibiotics against Gram positive cocci are summarized in Table 2 and Table 3, respectively.

### Table 1: Frequency of Bacterial Agents Isolated From Urine Specimens and Their Relation to Sex in This Study

<table>
<thead>
<tr>
<th>Isolated Bacteria</th>
<th>Female, No. (%)</th>
<th>Male, No. (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Gram-Negative bacilli</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td><em>Escherichia coli</em></td>
<td>136 (89.5%)</td>
<td>16 (10.5%)</td>
</tr>
<tr>
<td><em>Klebsiella</em> spp.</td>
<td>18 (81.8%)</td>
<td>4 (18.2%)</td>
</tr>
<tr>
<td><em>Proteus</em> spp.</td>
<td>4 (100%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td><em>Enterobacter</em> spp.</td>
<td>2 (100%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td><strong>Gram-Positive cocci</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td><em>Staphylococcus saprophyticus</em></td>
<td>22 (84.6%)</td>
<td>4 (15.4%)</td>
</tr>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>4 (100%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td><em>Streptococcus group B</em></td>
<td>10 (100%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td><em>Enterococcus</em> spp.</td>
<td>8 (80%)</td>
<td>2 (20%)</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td>204 (88.7%)</td>
<td>26 (11.3%)</td>
</tr>
</tbody>
</table>

### Table 2: Prevalence and Antimicrobial Resistance of Gram Negative Uropathogens Isolated in this Study

<table>
<thead>
<tr>
<th>No.</th>
<th>AM</th>
<th>CP</th>
<th>CT</th>
<th>IPM</th>
<th>GM</th>
<th>SXT</th>
<th>NA</th>
<th>CTX</th>
<th>FM</th>
<th>CAZ</th>
<th>CF</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>E. coli</em></td>
<td>52</td>
<td>73.69</td>
<td>26.32</td>
<td>15.79</td>
<td>15.79</td>
<td>73.69</td>
<td>69.74</td>
<td>60.53</td>
<td>38.16</td>
<td>9.22</td>
<td>55.27</td>
</tr>
<tr>
<td><em>Klebsiella</em> spp.</td>
<td>22</td>
<td>63.64</td>
<td>27.28</td>
<td>27.28</td>
<td>18.9</td>
<td>63.64</td>
<td>63.64</td>
<td>27.28</td>
<td>45.46</td>
<td>54.55</td>
<td>54.55</td>
</tr>
<tr>
<td><em>Proteus</em> spp.</td>
<td>4</td>
<td>100</td>
<td>0</td>
<td>0</td>
<td>25</td>
<td>50</td>
<td>75</td>
<td>50</td>
<td>50</td>
<td>100</td>
<td>50</td>
</tr>
<tr>
<td><em>Enterobacter</em> spp.</td>
<td>2</td>
<td>100</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>50</td>
<td>0</td>
<td>100</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

Abbreviations: AM, Ampicillin; CP, Ciprofloxacin; CT, Ceftizoxime; IPM, Imipenem; GM, Gentamicin; SXT, Sulfamethoxazole-trimethoprim; NA, Nalidixic acid; CTX, Cefatoxime; FM, Nitrofurantoin; CAZ, Cefazidime; CF, Cephalothin

### Table 3: Prevalence and Antimicrobial Resistance of Gram Positive Uropathogens Isolated in this Study

<table>
<thead>
<tr>
<th>No.</th>
<th>AM</th>
<th>C</th>
<th>E</th>
<th>NOR</th>
<th>GM</th>
<th>V</th>
<th>FM</th>
<th>TE</th>
<th>SXT</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Staphylococcus saprophyticus</em></td>
<td>26</td>
<td>92.31</td>
<td>61.54</td>
<td>92.31</td>
<td>30.77</td>
<td>30.77</td>
<td>7.7</td>
<td>7.7</td>
<td>92.31</td>
</tr>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>4</td>
<td>100</td>
<td>50</td>
<td>50</td>
<td>0</td>
<td>50</td>
<td>50</td>
<td>0</td>
<td>50</td>
</tr>
<tr>
<td><em>Streptococcus group B</em></td>
<td>10</td>
<td>30</td>
<td>20</td>
<td>30</td>
<td>40</td>
<td>70</td>
<td>80</td>
<td>0</td>
<td>70</td>
</tr>
<tr>
<td><em>Enterococcus</em> spp.</td>
<td>10</td>
<td>40</td>
<td>40</td>
<td>70</td>
<td>100</td>
<td>0</td>
<td>100</td>
<td>0</td>
<td>100</td>
</tr>
</tbody>
</table>

Abbreviations: AM, Ampicillin; C, Chloramphenicol; E, Erythromycin; NOR, Norfloxacin; GM, Gentamicin; V, Vancomycin; FM, Nitrofurantoin; TE, Tetracycline; SXT, Sulfamethoxazole-trimethoprim
5. Discussion

This study shows the distribution of microbial species isolated from patients with UTI at a diagnostic laboratory center in Karaj, Iran and their susceptibility pattern to antimicrobial agents. Furthermore, we described the relationship between sex and isolated bacterial agents of UTI. Microbial infection of the UTI is one of the most common infectious diseases worldwide (19). Approximately 1 in 3 women will require antimicrobial treatment for a UTI before age 24, and 40% to 50% of women will suffer from UTI during their lifetime (20). The sex distribution of patients in our study is consistent with those in other studies (2, 16, 21) showing a statistically predominance of females (88.69% of the positive cultures). It is assumed that the short urethra in girls predisposes them to ascending infection. Furthermore, management of mic- culation in women is important. Management mistakes made by women include cleaning perineum forward from the anus to the vulva (2) that can cause urinary tract infection.

Sexual activity has been reported to influence higher prevalence of UTI in females (22). Considering the fact that most of infecting organisms are commensals of per- anal and vaginal regions, emphasis on personal hygiene especially in females may be important in reducing the incidence of UTI (18). Males are less prone to UTIs possibly because of their longer urethra and the presence of anti- microbial substances in prostatic fluid (23). Our study indicates that E. coli is still the most common cause of UTI in Iran. This corresponds with the data obtained by other investigators (3, 15, 17). In addition, coagulase negative Staphylococcus spp. was the most common cause of UTI among Gram positive bacteria. Recent studies have revealed the importance of coagulase negative Staphylococcus spp. in urinary tract infections (24).

Our findings demonstrated that these pathogens play an important role in UTI as 11.3% of cases in our study were affected by these bacteria. Klebsiella spp. with the rate of 9.6% was the third species that caused UTI. In our study, as with previous studies, E. coli demonstrated a very high microbial resistance to antibiotics. The analyzed results of antibiotic susceptibility test showed that E. coli was highly resistant to gentamicin (73.69%), ampicillin (73.69%), SXT (69.74%), and cephaptothin (88.16%). This is similar to previous studies in USA (25) and Iran (17). On the other hand, very low resistance was detected to anti- biotics such as ciprofloxacin (26.32%), imipenem (15.79%), nitrofurantoin (9.22%), and ceftizoxime (15.79%).

In the present study, the coagulase negative Staphylococcus showed 92.31% resistance to ampicillin, tetracycline, and erythromycin, respectively. Results showed that the best activity against these bacteria was achieved by nitrofurantoin, gentamicin, norfloxacin, and vancomycin that corresponds with other studies in our region and other parts of the world (26, 27). Klebsiella spp. exhibited vary-
to those, demonstrate appropriate respond to other antibiotics such as ciprofloxacin and nitrofurantoin.

Acknowledgements

The authors would like to express their appreciations to Dr. Hi and Dr. M. Basaki for review of the manuscript and suggestions. We also thank Mr. A. Shahed for his technical support.

Financial Disclosure

All authors declare that they have no conflict of interest.

Funding/Support

None declared.

Authors’ Contribution

None declared.

References