Prevalence and Antibiotic Sensitivity Pattern of Methicillin-Resistant Staphylococcus aureus in Kathmandu Medical College –Teaching Hospital

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Abstract

Introduction: Staphylococcus aureus is found to be a major source of community as well as hospital acquired infection. Staphylococcal isolates from tertiary care hospital are found to be resistant to commonly used antimicrobial agents. Methicillin resistant S. aureus (MRSA) with intrinsically developed antimicrobial resistance has been associated with an increase in morbidity and mortality of the patients in the hospital. This study was undertaken to know the antibiotic sensitivity pattern of staphylococcal isolates with special reference to Methicillin resistant S. aureus.

Methods: Clinical specimens received from July 2009 to July 2010 in Kathmandu Medical college-Teaching Hospital were processed and all S. aureus isolates were included in the study. The isolates were identified by standard laboratory procedure. The antibiotic susceptibility pattern of all staphylococcal strain was determined by modified Kirby Bauer antibiotic sensitivity method.

Results: Of 111 S. aureus isolates 29(26.12%) were identified to be MRSA. The rate of multi drug resistance was 75.86% for MRSA and 6.09% for MSSA. All the staphylococcal isolates were resistant to penicillin. However, all strains were sensitive to vancomycin.

Conclusions: This study showed a high prevalence of MRSA in tertiary care hospital of Kathmandu valley. Regular surveillance of hospital-associated infection and monitoring of antibiotic sensitivity pattern is mandatory to reduce MRSA prevalence in hospital and its spread to community as well. Present study conclusively shows that vancomycin remains the first choice of treatment for MRSA infection. To preserve its value, use of vancomycin should be limited to those cases where there are clearly needed.

Key words: MRSA, MSSA, Antibiotic Sensitivity, Vancomycin, Staphylococcus aureus

Introduction

S. aureus remains a potent human pathogen, since it is one of the most common cause of nosocomial as well as community acquired infection.1 It is also the most significant pathogen known for causing sporadic infections and epidemics.2 Most of the S. aureus infections are caused by Methicillin-sensitive S. aureus strains (MSSA) that are usually susceptible to major classes of anti-staphylococcal antibiotics. But Resistance to multiple antibiotics among the staphylococci isolates in hospitals has been recognized as one of the major challenges in hospital infection control.3

In the recent years, the widespread use of antibiotics has
undoubtedly accelerated the evolution of MRSA and led to the emergence of strains that have systematically acquired multiple resistance genes. With the current emergence of multi-drug resistant MRSA in hospitals on the one hand and the dramatically increased incidence of hyper-virulent community-associated MRSA on the other hand, MRSA has been able to evolve rapidly and create new clinical problems. These strains are frequently implicated in serious infections and nosocomial outbreaks, which appear to be disseminated globally in adult, pediatric, and neonatal intensive care units (ICUs).

The prevalence of MRSA infection varies depending on the characteristics and size of the hospital. In Australia, 31.9% of S. aureus samples taken from 32 laboratories from all states and territories of the country were resistant to methicillin. The number of infections with MRSA in United States hospitals alone rose to nearly 369,000 in 2005.9 While study done in India also shows prevalence rate as high as 31.1% in clinical samples.10 In an earliest study from Nepal, Rai et al.6 found that 29% of S. aureus isolates were resistant to methicillin.11 MRSA are usually found to be resistant to most commonly used antibiotics against staphylococcal isolate. In fact, many strains of MRSA exhibit resistance to both β-lactams and aminoglycosides.12 These strains are seen possessing elevated resistance to a wide range of antibiotics, limiting the treatment options to very few agents such as vancomycin and teicoplanin.13 Hence, knowledge of prevalence of MRSA and MSSA and their antimicrobial profile becomes necessary in the selection of appropriate empirical treatment of these infections and controlling nosocomial infection. This study was done to know the antibiotic sensitivity pattern of staphylococcal isolates with special reference to Methicillin resistant S. aureus at KMC Teaching Hospital.

Methods

Clinical specimens from July 2009 to July 2010 received in clinical microbiology lab of KMC Teaching Hospital were processed and all S. aureus isolates were included in the study. One hundred and eleven (111) isolates of S. aureus were collected from culture samples received from different departments of the hospital. The isolates were consecutive and non repetitive (One per patient). One sample from one patient was inclusion criteria of study data, second sample from other site of same patient was not considered for study.

In Microbiology Lab, samples were cultured on Blood, Mac-Conkey, and Chocolate agars for 24-48 hours. Samples for blood culture were inoculated in Brain Heart Infusion broth and subcultured on 24 and 72 hours on Blood, Mac-Conkey agar. Identification of organisms was carried out by standard laboratory operating procedures.14 (Gram staining, Catalase test, Mannitol fermentation, Slide coagulase and Tube coagulase test).

The antibiotic susceptibility pattern of all the strains was determined by modified Kirby Bauer disc diffusion method against the following antibiotics: penicillin (10 units), gentamicin (10mcg), erythromycin (15mcg), tetracycline (30mcg), co-trimoxazole (25mcg), amikacin (30mcg), cephalexin (30mcg), ciprofloxacin (5mg) and vancomycin (30mg). For identification of MRSA, oxacillin discs (1unit) obtained from Hi-Media Laboratories Pvt. Ltd. was used. A zone of inhibition less than 10 mm. or any discernible growth within zone of inhibition was indicative of methicillin resistance. S. aureus ATCC 25923 was used as a standard control strain. Methicillin resistance was confirmed for all the MRSA isolates by the agar screening method using Mueller Hinton agar supplemented with 4% NaCl and 6 μg/L of oxacillin.15 Multidrug resistance (MDR) is defined for this report as resistance to three or more antimicrobial classes. Statistical analysis tool (SPSS17) was used to calculate P-value (≤0.05 significant) was calculated using Pearson Chi Square test.

Results

Out of 111 S. aureus isolated from various clinical specimen. Highest number of isolates was from pus and wound swab and least number from urine. (Table 1) 29 (26.12%) of isolates were MRSA (Table 1). Out of 29 MRSA isolated 22(75.86%) were MDR strain. Only 5(6.09%) among 82(73.87%) MSSA were MDR strain (Fig. 1).

Table 1: Distribution of S aureus and MRSA in various clinical specimens

<table>
<thead>
<tr>
<th>Clinical specimen</th>
<th>No of S aureus isolate (%)</th>
<th>MRSA (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Pus and wound swab</td>
<td>87 (78.37)</td>
<td>25 (28.73)</td>
</tr>
<tr>
<td>Blood</td>
<td>19 (17.11)</td>
<td>3 (15.78)</td>
</tr>
<tr>
<td>Urine</td>
<td>05 (04.50)</td>
<td>1 (20.00)</td>
</tr>
<tr>
<td>Total</td>
<td>111 (100)</td>
<td>29 (26.12)</td>
</tr>
</tbody>
</table>
Prevalence and Antibiotic Sensitivity Pattern of Methicillin-Resistant

Fig 1: Detection of multidrug resistance in MSSA and MRSA

The antimicrobial susceptibility pattern of MRSA and MSSA isolates against antimicrobial agents are summarized in Table 2. More than 25% of MRSA isolates were resistant to penicillin, oxacillin, cephalaxin, co-trimoxazole, erythromycin. Least amount of resistance was observed in vancomycin (0%) ciprofloxacin (17.03%) tetracycline and gentamicin (20.68%). Last but not the least amikacin (24.13%). β-lactam antibiotics like penicillin (100% resistance) and cephalaxin (30.48% resistance) were found to be ineffective against MSSA too. Rest of the antibiotic showed less than 20% resistance towards the isolated MSSA.

Table 2: Antibiotic resistance pattern of MRSA and MSSA

<table>
<thead>
<tr>
<th>Antibiotic</th>
<th>MRSA</th>
<th>%</th>
<th>MSSA</th>
<th>%</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Penicillin</td>
<td>29</td>
<td>100</td>
<td>82</td>
<td>100</td>
<td>N/A*</td>
</tr>
<tr>
<td>Oxacillin</td>
<td>29</td>
<td>100</td>
<td>00</td>
<td>00</td>
<td>0.0001</td>
</tr>
<tr>
<td>Vancomycin</td>
<td>00</td>
<td>00</td>
<td>00</td>
<td>00</td>
<td>N/A*</td>
</tr>
<tr>
<td>Gentamicin</td>
<td>06</td>
<td>20.68</td>
<td>05</td>
<td>0.6</td>
<td>0.024</td>
</tr>
<tr>
<td>Amikacin</td>
<td>07</td>
<td>24.13</td>
<td>02</td>
<td>2.43</td>
<td>0.001</td>
</tr>
<tr>
<td>Co-trimoxazole</td>
<td>13</td>
<td>44.82</td>
<td>17</td>
<td>20.07</td>
<td>0.012</td>
</tr>
<tr>
<td>Tetracycline</td>
<td>06</td>
<td>20.68</td>
<td>09</td>
<td>10.97</td>
<td>0.188</td>
</tr>
<tr>
<td>Cephalaxin</td>
<td>22</td>
<td>75.86</td>
<td>25</td>
<td>30.48</td>
<td>0.0001</td>
</tr>
<tr>
<td>Erythromycin</td>
<td>13</td>
<td>44.82</td>
<td>07</td>
<td>8.53</td>
<td>0.0001</td>
</tr>
<tr>
<td>Ciprofloxacin</td>
<td>05</td>
<td>17.03</td>
<td>14</td>
<td>17</td>
<td>0.984</td>
</tr>
</tbody>
</table>

Discussion

MRSA has emerged as a serious threat to public health worldwide. It has added to the burden of patient by prolonging hospital stay and increasing morbidity and mortality rate. Present study showed prevalence rate of MRSA to be 26.12%. Another study done in Kathmandu valley by Shrestha et. al. reported 44.9% as MRSA from nosocomial S. aureus.16 Rajbhandari et. al. also reported 54.9% MRSA isolates at Bir Hospital.1 Study done in Eastern Nepal showed comparable result of 26.14% MRSA.17 While recent study done in Bharatpur, Nepal had made known worrisome isolation rate of 39.6%.18 MRSA was isolated at the rate 75.5% from clinical samples in a study conducted by Rijal et. al. in Pokhara Valley.19 Similar study done in western parts of Nepal by Tiwari et. al. also had shown alarming high rate of MRSA isolate (69.1%) which the authors has attributed to indiscriminate use of antibiotics and its accessibility in these.20 Above studies show considerable variations between institutions, often in the same geographical areas, exist, demonstrating that MRSA prevalence, in some settings, significantly exceeds previous estimate. There could be many explanations for these differences: infection control measures, antibiotic prophylaxis and treatments used in each ward/hospital and, not less important, the clonal and often epidemic nature of these microorganisms.21,22
Present study also shows maximum number of S.aureus and MRSA isolation from pus and wound swabs (25/29) ascertaining the role of the organism as cause of pyogenic infection. This is similar to the study done in Nepal, Bharatpur, India and Pakistan. 16, 18, 20, 23, 24

Analysis from previous studies revealed a relationship between methicillin resistance and resistance to other antibiotics. 16, 18, 22, 24. This study showed that all MRSA isolates were significantly less sensitive to antibiotics as compared with MSSA isolates. Significant difference (P-value < 0.05) was observed in case of oxacillin, gentamicin, amikacin, co-trimoxazole, cephalexin, erythromycin. However, the difference observed in case of tetracycline and ciprofloxacin was statistically insignificant (P-value > 0.05). Homogeneous insusceptibility to beta-lactams like penicillin and cephalexin, characteristic MRSA was also observed in our study. This may be due to presence of intrinsically developed β-lactamase in MRSA strain. It also showed the high resistance to crotimoxazole and erythromycin as these antibiotics are usually used at random to cure generalized and pyogenic infection.

Antimicrobials such as amikacin, gentamicin and tetracycline with resistance less than 25% could be used against MRSA infection. But due to their mode of action, have limited use for empirical therapy of MRSA related infection.

Resistance to ciprofloxacin was observed to be 17.93% in this study. Limiting its indiscriminate use and doing antibiotic susceptibility testing it could be considered as a drug of choice for MRSA infection and can be recommended for empirical therapy in this setting.

The multi-drug resistant phenotype is a particular characteristic of the methicillin-resistant S.aureus strains. It has added to the burden of hospital personnel to control infection associated with MDR-MRSA. Present study shows alarmingly high rate of MDR strain among MRSA isolates (75.86%). Studies conducted in eastern and western part of Nepal also have reported MDR-MRSA to be as high as 65-78%. 16, 19 Indian literature also shows the isolation of MDR-MRSA as high as 77% 23

Though these MDR strains are not found with additional virulence properties, their characteristic multidrug resistance restricts the options available to treat infections caused by this organism. 24

Vancomycin a glycopeptides seems to be the only antimicrobial agent which showed 100% sensitivity through all parts of Nepal and may be used as the drug of choice for treating multidrug resistant MRSA infections. But its toxic side effects like renal impairment and prohibitive cost has limited its use. When vancomycin is considered for treatment, choice inevitably requires the need for in vitro susceptibility testing of every isolate of MRSA in the clinical laboratories owing to emergence of Vancomycin resistant Staphylococcus aureus (VRSA) in various parts of world.

**Conclusion**

In view of the high rates of isolation of MRSA and its resistance to penicillin, erythromycin, cephalexin, and trimethoprim, treatment of MRSA infections in this province with these antibacterial agents would be unreliable. Vancomycin remains the first choice of treatment for MRSA infection world wide and to preserve its value, use of vancomycin should be limited to those cases where there are clearly needed. Further, the regular surveillance of hospital associated infections including monitoring antibiotic sensitivity pattern of MRSA and MSSA and formulation of definite antibiotic policy may be helpful in reducing the incidence of MRSA infection. Our study is an opening to facilitate epidemiologists to understand the nature of MRSA isolates in this part of Nepal.

**References**


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