# Bacteriological Profile of Orthopedic Patients in a Tertiary Care Hospital, Bengaluru

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## Abstract

**Background:** In developing countries, the problem of changes in pathogenic microbial flora and the emergence of bacterial resistance have created major problems in the management of orthopedic diseases. The aim of this study was to determine the type of bacterial pathogens isolated from surgical site infection (SSI), open fractures and non-operative wound infections in our hospital and their antibiotic sensitivity profiles. **Materials And Methods:** This study was conducted in the department of microbiology, RajaRajeswari Medical College & Hospital, Bengaluru, over a period of 1 year from February 2014 to January 2015. During this period around 195 samples received as surgical site infection, from open fracture and wound infected cases. Standard microbiological techniques were used to isolate and identify the organisms and to determine the antibiotic resistance pattern. **Results:** 45.13% (88/195) specimens showed culture positivity. Surgical site infection rate was 46.59% (41/88). The age of patients ranged from 1 year to 75 years with a mean age of 39 years. The most commonly isolated organisms were Staphylococcus aureus (40.90%), Escherichia coli (15.9%), Pseudomonas aeruginosa (13.6%). 12.5% of Staphylococcus aureus were methicillin resistant. All Staphylococci were susceptible to vancomycin, linezolid and teicoplanin. All gram negative bacilli were sensitive to colistin and tigecycline. **Conclusion:** High rates of antibiotic resistance observed in our study, due to widespread usage of broad spectrum antibiotics. While deciding antibiotic therapy many factors must be considered like previous antibiotic history, knowledge of most common causative organism in these infections, and their antibiotic profile.

## Keywords

Antimicrobial sensitivity, Implant infections, Orthopedic infections, Surgical site infection, Staphylococcus aureus.

## 1. Introduction

Orthopedic infections are one of the commonest which can occur in approximately one percent of all orthopedic operations.¹ The commonest orthopedic infections are surgical site infections and implant infections in open or closed wounds.²³ Incidence of Surgical site infection (SSI) is reported to vary from 3.6% to 22.5%⁴⁻⁵ and the implant infection varies from 0.5-2% in fixation of closed fractures to 30% after fixation of open fractures.⁶⁻⁷⁻⁹⁻¹⁰

Surgical site infection (SSI) as defined by US Centers for Disease Control (CDC) in 1992 is an infection occurring within 30 or 90 days after a surgical operation (or within 1 year if an implant is left in place after procedure) and affecting either incision or deep tissues at the operation site. These infections may be superficial or deep incisional infection or infections involving organ or body space.¹¹ Open or compound fractures are fractures that communicate with the outside environment through skin wounds.¹² The main causes of open fracture include road traffic accidents, fall from height, assaults, machine injury and others. Anglen JO et al reported 3-4% of all fractures are open fractures, and the development of infection is favored by devitalization of bone and soft-tissue. Use of implants and prosthesis during the orthopedic surgeries can pose greater risk of microbial contamination and infection.¹³

During the course of surgery the exogenous or endogenous microorganisms that enter the operative wound are responsible for these infections.¹² The factors that influence the nature and frequency of infection are type of wound, nature of surgery, the dose and virulence of infecting organism, host resistance and drug resistance of organisms.¹³ The bacteriological profile of the orthopedic cases are changing day by day all over the world, so the need to know the pathogen profile involved in the infections of orthopedic cases remains challenging.

Despite advances in diagnostic technologies patients with orthopedic infections have been given empirical therapy which can lead to drug resistance. So this study is aimed at determining the type of bacterial pathogens in SSI, open fractures and non-operative wound infections in our hospital and their antibiotic sensitivity profiles. And also to formulate and develop an antibiotic policy for the chemotherapeutic management of orthopedic infections.

## 2. Materials and Methods

The study was conducted in the department of microbiology, RajaRajeswari Medical College & Hospital from February 2014 to January 2015. Swabs from surgical site infection, open fractures, bedsore and infected wounds were collected with aseptic precautions and immediately transported to the laboratory for culture and antibiotic susceptibility testing.

Swabs were inoculated onto 5% sheep blood agar, MacConkeys agar and Brain Heart Infusion broth. The plates were incubated at 37 °C for 24-48 hours and...
examined for the growth of bacteria. All positive cultures were identified by their characteristic appearance on their respective media, Gram staining reaction and confirmed by the pattern of biochemical reactions. If no growth was observed on the plates, subcultures were made from the Brain Heart infusion broth onto 5% sheep blood agar and MacConkey agar, which were observed after 24 hours of incubation. Antimicrobial susceptibility testing done on Mueller Hinton Agar by Kirby Bauer disk diffusion method (means and percentages) were used wherever necessary. 

All the confirmed Staphylococcus aureus and coagulase-negative Staphylococcus spp (CONS) strains were subsequently screened for Methicillin resistance based on Kirby-Bauer disk diffusion method as per CLSI guidelines.

3. Interpretation

The isolates were considered Methicillin Resistant Staphylococcus aureus (MRSA) if the zone of inhibition was less than 21 mm and Methicillin Sensitive Staphylococcus aureus (MSSA) if it was ≥ 22 mm. For coagulase-negative Staphylococcus (CoNS), if the zone of inhibition was less than 24 mm considered as Methicillin resistant coagulase-negative Staphylococcus (MRCONS) and if it was ≥ 25 mm Methicillin sensitive coagulase-negative Staphylococcus (MSCONS).

The antibiotics tested against Staphylococcus spp were penicillin-G, cephalaxin, cefazolin, erythromycin, clindamycin, gentamicin, amikacin, vancomycin, teicoplanin, linezolid, rifampicin and chloramphenicol. The following antibiotics were used for Gram Negative bacilli: ampicillin, cephalaxin; ceftriaxone; cefotaxime; amoxicillin-clavulanate; ciprofloxacin; gentamicin; amikacin; imipenem; meropenem; piperacillin-tazobactam and the antibiotics tested against Pseudomonas spp were gentamicin, amikacin, ciprofloxacin, aztreonam, ceftazidime, piperacillin-tazobactam, imipenem, meropenem, netilmicin and tobramycin. The tests were interpreted as Sensitive, Intermediate susceptible or Resistant in accordance with standard recommendation.

4. Statistical Analysis

Data was entered into a computerized Excel (Microsoft Excel 2009) spread sheet, and subsequently it was analyzed using SPSS (trial version 20) software. Descriptive statistics (means and percentages) were used wherever necessary.

5. Results

During the one year study period, a total of 195 specimens were received from orthopedic department which included specimens from open fractures (18/88), surgical site infection (41/88), deep bed sores involving bones and miscellaneous ones (31/88). 45.13% (88/195) specimens showed culture positivity and 54.88% (107/195) specimens did not show any growth. The age of patients ranged from 1 year to 75 years with a mean age of 39 years.

In our study, both Gram positive and Gram negative organisms were isolated in equal numbers (44/88). The common isolates found in our study are Staphylococcus aureus, Escherichia coli, Pseudomonas aeruginosa, Klebsiella Spp, Non fermenting gram negative bacilli and Proteus spp. Majority of the organisms isolated were from surgical site infections (46.59%) followed by wounds (35.22%) and open fracture (18.18%). The incidence of various microbes in relation to orthopedic illness and procedures are detailed in the table 1. The antibiotic susceptibility pattern of gram positive organisms and gram negative organisms are detailed in the table 2 and 3 respectively.

6. Discussion

Infections are known to occur in spite of aseptic precautions by the orthopedicians. Eighty eight (45.13%) samples showed positive culture in our study which coincides with Abraham Y et al study which showed 41% positivity, whereas Gomez et al and Zimmeli et al reported positive cultures in 60% and 89% respectively.

In 1950’s and 60’s Staphylococcus aureus used to be the most common strain. In late 70’s there was a shift from gram positive infections to gram negative infections among orthopedic patients.

Staphylococcus aureus (40.90% of the total number of isolates) is the most common organism isolated from our study. Anterior nares, palm acts as important reservoirs for Staphylococcus aureus, about 10-20% of the healthy individuals will harbour this organism. Bed sheets, instruments and dressings have been found to act as reservoirs. Bergqvist et al and Dan et al found that 29.8% of hospitalized patients and 26.6% of hospital staff respectively are carriers. 12.5% (11/88) of our isolates are Methicillin Resistant Staphylococcus aureus. Other studies have observed MRSA ranging from 5.6% to 37.9 %, thus indicating lower range of prevalence of MRSA during this study period.

Escherichia coli is the second most common pathogen (15.9% cases), especially in SSI and patients with wound infection. Escherichia coli is a commensal of gut and as many patients are admitted for prolonged periods, contamination of wounds, dressings, linen, clothes and hands during perineal hygiene plays a major role in increasing chances of transmission of infection.

Pseudomonas aeruginosa (13.6% cases) is the third most common cause, which commonly isolated from SSI and bed sores. Pseudomonas can multiply on common objects in a hospital environment such as dressings materials, buckets used for soaking Plaster of Paris bandages and foreceps, has also been isolated in a significant number (10.2% cases) in our study. Other gram negative organisms like NFGB (5.7% cases), Proteus spp. (4.6% cases) isolated from SSI.

In our study, all the MRSA (12.5%) isolates were sensitive to vancomycin, linezolid and teicoplanin. Many other studies
have reported all the staphylococcal isolates being sensitive to vancomycin and linezolid.\(^{(27)}\) Currently vancomycin resistance \textit{Staphylococcus aureus} (VRSA) is not widespread. Vancomycin remains the first choice of treatment for MRSA. There was good sensitivity of MRSA for doxycycline (90.90%), and clindamycin (63.63%), so these drugs are also useful for SSI by MRSA. Among coagulase negative \textit{staphylococcus} (5.9%) isolates only 2.3% of isolates were methicillin resistant, which were sensitive to vancomycin, teicoplanin and linezolid.

In this study, \textit{E.coli} showed more resistance to ampicillin (93%), piperacillin (93%), cephalexin (93%), cefuroxime (86%), amoxicillin/clavulanic acid (79%), ciprofloxacin (71%), cefotaxime (72%), and less resistance to ceftazidime (57%), piperacillin/tazobactum (30%), gentamicin (29%), meropenem (14%) and imipenem (7%), while amikacin, colistin and tigecycline were 100% sensitive (Table No.3). Similar finding were observed by Aratikalakutakar, Vishwanath LYemul.\(^{(28)}\) 36% ESBL \textit{E.coli} were isolated.

\textit{Pseudomonas aeruginosa} showed 50% resistant to piperacillin, piperacillin and cefotaximoz, 33% to ceftazidime and cefepime, 25% to gentamicin and netilmicin, 17% to piperacillin/tazobactum and meropenem, while tobramycin, imipenem, colistin were 100% sensitive (Table No.3). Similar observation was reported by Aratikalakutakar, Vishwanath LYemul.\(^{(28)}\)

From our results, we observed that amoxicillin/clavulanic acid, ceftriaxone and ceftazidime cannot be recommended for use as an empirical therapy in SSI and open fracture infections because these drugs were inactive against most strains. Based on the antimicrobial susceptibility data, we suggest that piperacillin/tazobactum and imipenem are the most effective agents against most of gram negative bacteria and doxycline, vancomycin, linezolid are the most effective agents against gram positive organisms. Colistin and tigecyclin showed 100% sensitivity by all gram negative bacteria, but these drugs are kept as reserve, should be used judiciously.

### 7. Conclusion

High rates of antibiotic resistance were observed in our study, due to widespread usage of broad spectrum antibiotics. While deciding antibiotic therapy many factors must be considered like previous antibiotic history, knowledge of most common causative organism in these infections, and their antibiotic profile. By multidisciplinary collaboration involving: the orthopedic surgeons, infectious disease specialist and clinical microbiologist we can further reduce the incidence of infection in our hospital.

### 8. Recommendations

There is a need for formulation of antibiotic policy in tandem with clinicians/orthopedicians and antibiotic sensitivity pattern. A strict adherence to the antibiotic policy and formulary restriction is a must.

### References


Table 1: Frequency of Gram positive and gram negative organisms isolated from different sites.

<table>
<thead>
<tr>
<th>Organism</th>
<th>Open Fracture</th>
<th>SSI</th>
<th>Bedsore</th>
<th>Total</th>
</tr>
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<tbody>
<tr>
<td>Staphylococcus aureus (MSSA)</td>
<td>7</td>
<td>5</td>
<td>13</td>
<td>25 (28.4)</td>
</tr>
<tr>
<td>MRSA</td>
<td>5</td>
<td>3</td>
<td>3</td>
<td>11 (12.5)</td>
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<td>2 (2.3)</td>
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<td>1</td>
<td>0</td>
<td>1 (1.2)</td>
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<tr>
<td>Coagulase negative Staphylococcus (MSCONS)</td>
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<td>3 (3.4)</td>
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<td>MRCONS</td>
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<td>2</td>
<td>0</td>
<td>2 (2.3)</td>
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Table 2: Antibiotic sensitivity pattern of Gram positive organisms

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<th>AMC</th>
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<th>LEX</th>
<th>CXM</th>
<th>ERY</th>
<th>CLI</th>
<th>CIP</th>
<th>DOX</th>
<th>SXT</th>
<th>CHL</th>
<th>GEN</th>
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Table 3: Antibiotic sensitivity pattern of Gram negative organisms

<table>
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<th>Organism</th>
<th>AMK</th>
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<th>TOB</th>
<th>CIP</th>
<th>SXT</th>
<th>COL</th>
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<td>21</td>
<td>7</td>
<td>14</td>
<td>28</td>
<td>43</td>
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<td>P.aeruginosa (12)</td>
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<td>-</td>
<td>-</td>
<td>-</td>
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<td>-</td>
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<tr>
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<td>29</td>
<td>29</td>
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<td>57</td>
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<td>-</td>
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