

A study of wound infections and its antibiogram in surgical intensive care unit of a tertiary care hospital

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Background: Wound infections can be caused through two major sources: exogenous and endogenous bacteria. The probability of wound infection largely depends on the patients systemic host defenses, local wound conditions and microbial burden. Despite modern surgical techniques and the use of antibiotic prophylaxis, Surgical Site Infection (SSI) is one of the most common complications encountered in surgery. SSI places a significant burden on both the patient and health system. SSI is thus a major cause of morbidity, prolonged hospital stay and increased health costs. Objective of this study was to identify and isolate various bacteria from wound infections in a surgical intensive care unit and to study their antibiogram. **Methods:** Two wound swabs were collected from the wound and from a drop of aspirate, smear was made on clean glass slide and Gram staining was done for direct microscopic examination under oil immersion 100X objective to know various morphological types of bacteria and presence or absence of inflammatory cells. Second swab/drop of aspirate was used for culture by inoculating it on routine media like Blood Agar, Nutrient Agar and Mac Conkeys agar, incubated at 37° C for 24 hours aerobically. **Result:** Out of 238 pus samples, 209 (88.6%) were culture positive for bacterial growth and no growth was observed in 29(12.4%) cases. Out of 209 bacterial culture positive cases, 201 were monobacterial and 8 were poly bacterial. Out of 201 bacterial isolates; *S. aureus* (56/26.6) was the commonest followed by *P. aeruginosa* (47/22.4%). **Conclusion:** The study concludes that variety of aerobic bacteria is responsible for wound infection with predominance of *Staphylococcus aureus* followed by *Pseudomonas aeruginosa*

Keywords: Staphylococcus aureus, Pseudomonas aeruginosa, Monobacterial

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Introduction

Infection is defined as invasion and multiplication of micro organisms in the body tissues, which may be clinically in apparent or result in local cellular injury because of competitive metabolism, toxins, intra-cellular replication or antigen antibody response [1]. This series of events lead to progressive tissue destruction and eventual host demise if left unchecked. The infection process begins with a disruption of the host mechanical barriers to micro organisms, the availability of microorganisms and colonization [2]. Wound infections can be caused through two major sources: exogenous and endogenous bacteria. The probability of wound infection largely depends on the patients systemic host defenses, local wound conditions and microbial burden [3,4]. Despite modern surgical techniques and the use of antibiotic prophylaxis, Surgical Site Infection (SSI) is one of the most common complications encountered in surgery [5]. SSI places a significant burden on both the patient and health system. SSI is thus a major cause of morbidity, prolonged hospital stay and increased health costs [6].

Skin and soft tissue infections (SSTIs) may also contribute to longer hospital stays increase the cost of medical care and play an important role in development of antimicrobial drug resistance. Common examples of SSIs includes cellulitis, abscesses, impetigo, folliculitis, furuncle, carbuncle, necrotizing fasciitis, diabetic foot infections and surgical site infections. Complicated SSI may prove fatal and require hospitalization, intravenous antibiotics or surgery. An SSI is classified as complicated if the infection has spread to the deeper soft tissue, if surgical intervention is necessary or if the patient has co-morbid conditions Hence, this study could play a significant role in the early recognition of the Problem and hence, there is need for early intervention for better management of wound infections.

Material and Methods

Duration & Setting: This study was done in a tertiary care hospital for a period of one year from Jan 2016-Dec 2016.

Ethical clearance: Ethical clearance from the Institutional Ethical committee was obtained.

Sample size: Two hundred & thirty eight cases of pussamples were taken.

Isolation and identification of aerobic bacterial pathogens was done, from various departments like Surgery, Gynecology & Orthopedics.

Data collection: Wound swabs were collected as per the procedure described below.

Procedure: Two wound swabs were collected from the wound and from a drop of aspirate, smear was made on clean glass slide and Gram staining was done for direct microscopic examination under oil immersion 100X objective to know various morphological types of bacteria and presence or absence of inflammatory cells. Second swab/drop of aspirate was used for culture by inoculating it on routine media like Blood Agar, Nutrient Agar and MacConkeys agar, incubated at 37° C for 24 hours aerobically [7-9].

The plates were examined the next day for growth. Plates not showing any growth were further incubated at 37°C aerobically for another 24 hrs. Plates not showing any growth after 48 hrs on aerobic incubation were considered to be lacking aerobic bacterial pathogens. Smears were made, stained by Gram stain and examined under oil immersion microscope 100X objective.

Antibiotic sensitivity testing was performed on Mueller Hinton Agar according to CLSI guidelines. MRSA was detected using Cefoxitin (30ug) disc and ESBL production in Gram negative bacteria was detected by using Potentiated Disc Diffusion Test (PDT)[8-10].

Inclusion criteria: All the swabs for aerobic bacterial pus culture were obtained

Exclusion criteria: Patients already on antibiotics were excluded from the study

Results

Two hundred & thirty eight cases of pus samples were taken. Aerobic bacterial pathogens were isolated and identified from various departments like Surgery, Gynecology and Orthopedics. Surgical wound swabs were 139 (58.4%) and Non Surgical wound swabs were 99(41.5%) in number. Out of 99 Non-surgical wounds, 69(69.6%) were soft tissue infections wound and burn wounds, and 30(30.3%) were traumatic wounds. Out of 139 surgical wounds, 86(61.8%) were post-operative wounds and 43(30.9%) were various other surgical site wounds.

Table-1: Distribution pattern of bacterial isolates (n= 238).

| Organisms | Total No. of cases |
|----------------------------------|--------------------|
| Staphylococcus aureus | 52 |
| Pseudomonas aeruginosa | 41 |
| Escherichia coli | 26 |
| Klebsiella pneumoniae | 25 |
| Proteus mirabilis | 17 |
| Acinetobacter spp. | 11 |
| Coagulase negative Staphylococci | 11 |
| Enterococcus faecalis | 10 |
| Citrobacter freundii | 08 |
| P.aeruginosa + S.aureus | 03 |
| P. aeruginosa +K.pneumoniae | 03 |
| S. aureus + P. mirabilis | 01 |
| K.pneumoniae + P. mirabilis | 01 |
| No growth | 29 |

Pus discharge was collected from 238 patients were identified, out of which 167 (70.1%) were males and 71(29.9%) were females.

Cases of pus discharge came mainly from rural areas (199/83.61%) as compared to urban areas (39/16.38%).

Bacterial isolates: Out of 238 pus samples, 209(88.6%) were culture positive for bacterial growth and no growth was observed in 29(11.4%) cases as shown in Table 1. Out of 209 bacterial culture positive cases, 201 were monobacterial and 8 were poly bacterial. Out of 209 bacterial isolates; *S. aureus* (56/26.6%) was the commonest followed by *P. aeruginosa* (47/22.4%). The prevalence of monomicrobial isolates from various wound infections is depicted in table 2, wherein *Staphylococcus aureus* was the predominant organism when samples were collected from post operative wounds, burns, traumatic wounds and soft tissue infections.

Table-2: Monomicrobial isolates in various wound infections.

| Types of wounds | S.aureus | P.aeruginosa | E.coli | K.pneumoniae | P.mirabilis | Acinetobacter spp. | CONS | E.fecalis | C.freundii |
|------------------------|----------|--------------|--------|--------------|-------------|--------------------|------|-----------|------------|
| Post operative wounds | 12 | 6 | 17 | 17 | 8 | 7 | 6 | 2 | 1 |
| Burns | 5 | 31 | 3 | - | 4 | - | 2 | 1 | 5 |
| Traumatic | 16 | 4 | 4 | 1 | 3 | 2 | 2 | 1 | 1 |
| Soft tissue infections | 19 | - | 2 | 7 | 2 | 2 | 1 | 6 | 1 |
| Total | 52 | 41 | 26 | 25 | 17 | 11 | 11 | 10 | 08 |
| | 25.8% | 20.3% | 12.9% | 12.4% | 8.4% | 5.4% | 5.4% | 4.9% | 3.9% |

Antibacterial susceptibility profile: Gram positive bacteria showed maximum susceptibility to Vancomycin whereas gram negative isolates showed its maximum susceptibility to Meropenem and Amikacin. *P. aeruginosa* isolates showed maximum susceptibility to Doripenem and Piperacillin tazobactam. 44.2% of *S.aureus* isolates were MRSA and 56% of Gram negative isolates were ESBL producers as shown in Table 3.

Table-3: Antibiotic sensitivity pattern of bacterial isolates.

| Antibiotics | Gram positive isolates n =73 | Gram negative isolates n =87 | Pseudomonas isolates n=41 |
|------------------|------------------------------|------------------------------|---------------------------|
| Vancomycin (VA) | 100% | - | - |
| Clindamycin (CD) | 26.5% | - | - |
| Linezolid (LZ) | 80.5% | - | - |

| | | | |
|---------------------------------|-------|-------|-------|
| Erythromycin (E) | 28.9% | - | - |
| Ampicillin (AMP) | 11.8% | - | - |
| Amoxyclav (AMC) | 32.9% | - | - |
| Ceftriaxone (CTR) | 53.5% | 55.6% | 61% |
| Cefoxitin (CX) | 55.8% | - | - |
| Cefotaxime (CTX) | - | 44.4% | - |
| Ceftazidime (CAZ) | - | 38.9% | 61.1% |
| Gentamicin (GEN) | 71.2% | 38.9% | 33.3% |
| Amikacin (AK) | 82.4% | 72.2% | 69.4% |
| Imipenem (IPM) | - | 74.4% | 86.1% |
| Meropenem | - | 87.1% | 86.7% |
| Doripenem | - | 90.6% | 91.4% |
| Piperacillin + Tazobactam (TZP) | - | 66.7% | 90.7% |
| Cefopodoxime (CPZ) | - | 50% | - |
| Ciprofloxacin (CIP) | - | 33.3% | 36.1% |
| Netlimicin (NET) | - | - | 44.4% |
| MRSA | 44.2% | - | - |
| ESBL producers | - | 56% | - |

Discussion

In the present study an attempt was made to study the bacteriological profile of wound infections and antimicrobial susceptibility pattern of the isolates. In this study along with the identification of aerobic bacterial organisms, changing pattern of antibiotic sensitivity with special reference to Methicillin Resistant *Staphylococcus aureus* (MRSA) and Extended Spectrum of Beta lactams (ESBLs) were also identified.

It was observed that the commonest age group affected is 21-40 years which is correlated with the studies done by Shute Malik et al [11] and Dr. Zarrin Afroz et al [12]. Males (70.1%) were affected more than females (29.9%). This study was correlated with Ramesh Rao et al [15] which found males (60%) more affected than females, N. Sowmya et al [14] 66.6% and Shruti Malik et al [11] 51.9% also showed the predominance of males over females probably because of their more exposure to life.

Monomicrobial etiology was more common 96.1%, than polymicrobial 3.8%. This study is correlated with N. Soumya et al [14] 91.7% and Mehta V.J. et al [15] 70.4% wherein the monomicrobial etiology was more common than polymicrobial which may be due to the prior use of antibiotics.

In the present study *Staphylococcus aureus* 26.6% was the predominant organism followed by *Pseudomonas aeruginosa* 22.4%, *Escherichia coli* 12.9%, *Klebsiella pneumoniae* 12.4%, *Proteus mirabilis* 8.4% and CONS 5.4%. *Staphylococcus aureus* (26.6%) was the most predominant isolate which correlated with the other studies done Shruti Malik (30.1%), Mehta V.J. (38.3%), Ramesh Rao (27.8%) and Dr. Pravin (48.4%) [13-16].

Second most predominant organism in the present study was *Pseudomonas aeruginosa* 22.4% which correlated with the studies of Gayathree Naik (20%), Shruti Malik (17.8%), Mehta V.J (21.3%), Ramesh Rao (18.5%) and Dr. Pravin (17.52%) [11-15]. In the present study, polymicrobial 8 cases included combination of *Pseudomonas aeruginosa* with *Staphylococcus aureus* (37.5%) which correlates with the study of Anbumani et al. Other polymicrobials included *Pseudomonas aeruginosa* with *Klebsiella pneumonia* accounting for 37.5% cases, *Klebsiella pneumonia* with *Proteus mirabilis* 12.5% and *Staphylococcus aureus* with *Proteus mirabilis* 12.5% case. Out of 52 *Staphylococcus*

Aureus isolates, 23(44.3%) were MRSA producers and remaining 29 (55.7%) were MSSA producers. The present study correlates with the study of Rajadurai et al [19] with 31%, Anupurba et al [18] with 32% and N. Soumya et al [14] with 27.5%, as MRSA producers. Among 77 Enterobacteriaceae isolates, 56% were ESBL producers and 44% were Non-ESBL producers which correlated with the studies done by Mehta V.J [15] with 44.6% as ESBL producers.

In the present study, Vancomycin (100%) was the most sensitive antibiotic among all gram positive isolates which was correlated with the studies of Amrita Shriyan et al [21], Shahnooshi Javed et al [22] and Jeena Amatya et al [23]. Amikacin was the second most sensitive antibiotic to many gram positive as well as gram negative isolates accounting for 82.4%, which is correlated with the study of Amrita Shriyan et al [21] 95% and Shruti Malik et al [11] 90%. Meropenem was the most sensitive drug among gram negative isolates accounting for 87.1% which was correlating with the study of Shruti Malik et al [11], Amrita Shriyan et al [21] and Ramesh Rao et al [13]. Limitation of this study was that only bacterial samples were isolated where in fungal isolates could also be isolated for better understanding if surgical infections.

Conclusion

Wound infections are one of the most common hospital acquired infections, and are an important cause of morbidity & account for 70-80% mortality. Development of such infections represent delayed healing, cause anxiety & discomfort for patient, longer stays at hospitals & add to cost of healthcare services significantly. This study was carried out to determine the antibacterial susceptibility of bacteria isolated from wound infections as well as update the clinicians in the various antimicrobial alternatives available in the treatment of wound infections, thus helping to reduce the burden of infection on patients and in long term, it may reduce the cost of treatment.

The study concludes that variety of aerobic bacteria is responsible for wound infection with predominance of *Staphylococcus aureus* followed by *Pseudomonas aeruginosa*. Also there were mixed infections with different bacteria. Antibiotic sensitivity pattern of the study revealed that Amikacin was the most sensitive drug among both gram positive and gram negative isolates.

Meropenem was the most sensitive drug among all gram negative isolates and Vancomycin was the most sensitive drug among gram positive isolates. This study shows that the organisms are becoming resistant to commonly used antibiotics and also developing resistance to newer antibiotics.

What the study adds to the existing knowledge?

Nowadays automated methods of identification of the pathogen and antibiotic sensitivity are available which aids in quick results, thus reducing the time of treatment and helping the patients. More comprehensive studies are required from time to time to define the magnitude of problem & produce data for policy decision on optimal intervention modalities. Vital use for formulation of antibiotic policy and implementation of antimicrobial stewardship program-mmes is need of the hour. Hospital must have an active antibiotic stewardship programme implemented. Further based on the antibiogram of hospital, antibiotic policy should be framed and followed. Such a treatment policy if followed will lead to reduction in mortality, morbidity & health care cost associated with wound infections.

Author's contributions

All the authors contributed equally in the study design, analysis and manuscript preparation

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