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Isolation Of Bacterial Pathogen And Antibiotic Sensitivity Analysis Of Breast Abscess In Lactating And Non-Lactating Women

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Abstract

This study aimed to isolate and identify bacterial pathogens from breast abscess samples of women attending different tertiary care hospital in Rawalpindi, Pakistan, and to evaluate the antibiotic susceptibility of these isolates. Pus samples were collected from both lactational and non-lactational women and processed using standard microbiological techniques, including culture on selective media, Gram staining, and biochemical tests. Antibiotic susceptibility testing was performed using the Kirby-Bauer disc diffusion method following CLSI guidelines. The predominant bacterial isolates were Staphylococcus aureus, Escherichia coli, Streptococci, and Pseudomonas aeruginosa. S. aureus exhibited high susceptibility to ciprofloxacin, levofloxacin, and amikacin, but resistance to streptomycin and amoxicillin. P. aeruginosa was most susceptible to ciprofloxacin and clarithromycin, while showing resistance to streptomycin, amoxicillin, and amikacin. E. coli was susceptible to ciprofloxacin and clarithromycin, but resistant to amikacin, amoxicillin, and streptomycin. Streptococci showed susceptibility to ciprofloxacin, levofloxacin, and clarithromycin, but resistance to moxifloxacin and amikacin. The findings of this study highlight the importance of routine microbiological culture and antibiotic susceptibility testing for effective management of breast abscess infections.



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Introduction

Breast abscess is a localized collection of purulent material within the breast tissue, commonly encountered in women, and represents a significant clinical concern in both lactating and non-lactating populations [1]. This condition can be classified into two primary categories: lactational and non-lactational abscesses. Lactational abscesses predominantly affect women of reproductive age, particularly those who are breastfeeding, while non-lactational abscesses are more frequently observed in perimenopausal women and those with underlying health conditions [2].

The clinical presentation of breast abscesses typically includes painful, swollen, and reddish masses within the breast, often accompanied by systemic symptoms such as fever and malaise [3]. These abscesses can significantly impact a woman's quality of life, leading to physical discomfort and emotional distress. [4] The pathogenesis of breast abscess often involves a bacterial infection, with bacteria entering the breast tissue through cracks or fissures in the nipple, which can occur due to trauma, improper latch during breastfeeding, or other irritative factors. [5] In the case of lactational abscesses, milk stasis plays a critical role by creating a nutrient-rich environment that facilitates bacterial growth and proliferation [6].

The most common causative organism identified in breast abscesses is Staphylococcus aureus, particularly methicillin-resistant strains (MRSA), though other bacteria, including Streptococcus species and Escherichia coli, can also be implicated [7,8]. The presence of these pathogens necessitates prompt and accurate identification through cultures and sensitivity testing. [9] Effective treatment hinges on the timely initiation of appropriate antibiotic therapy, which is crucial not only for the resolution of the abscess but also for the prevention of potential complications, such as systemic infection or recurrence of the abscess.[10] Understanding the underlying mechanisms and risk factors associated with breast abscesses is essential for healthcare providers to implement effective prevention and management strategies.

Materials and Methods

Study Area and Population

The study was conducted from November 2024 to February 2025 at the Amjad naeem Laboratory, Clinic, Rawalpindi, Pakistan. The study population comprised women visiting the clinic in Rawalpindi, Khyber Punjab, with breast abscess. Sample Collection

Pus samples were collected from patients with breast abscess at different Hospitals in, Rawalpindi. Eighty-Eight pus samples were collected in sterile universal containers and transported to the laboratory following standard protocols.

Bacterial Isolation and Identification

The pus samples were subjected to Gram staining and cultured on Nutrient Agar and Mannitol Salt Agar (MSA) [1]. Inoculated plates were incubated at 37°C for 18-24 hours, and if no growth was observed, incubation was extended for another 24 hours. Colonies were examined for color, shape, size, margins, elevation, and hemolytic pattern. Isolated organisms were identified by spot tests, including catalase and coagulase tests [1].



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Antimicrobial Susceptibility Testing

Antimicrobial susceptibility testing of the isolated organisms was performed on Mueller-Hinton agar using the Kirby-Bauer disc diffusion technique [1]. The following panel of antimicrobials was used: ampicillin, ciprofloxacin, levofloxacin, amikacin, clarithromycin, sulfamethoxazole, streptomycin, and moxifloxacin. The tests were conducted, and results were interpreted according to the Clinical and Laboratory Standards Institute (CLSI) 2019 guidelines [1].

Results

The mean age of the patients was 30.3 years (range 15-50 years). The majority of patients were in the 24-35 age group, and two patients were over 50 years of age. Out of 84 samples, 68 showed bacterial growth, and polymicrobial growth was observed in 3 out of 71 samples. The predominant isolates were Gram-positive bacteria, including *Staphylococcus aureus* and Streptococci. The Gram-negative bacteria isolated were *Escherichia coli* and *Pseudomonas aeruginosa*. *S. aureus* isolates showed the highest susceptibility to ciprofloxacin (80%), levofloxacin (80%), and amikacin (93.33%), while exhibiting high resistance to streptomycin (73.33%) and amoxicillin (93.33%).

P. aeruginosa isolates were most susceptible to ciprofloxacin (90%) and clarithromycin (85%) and most resistant to streptomycin (90%), amoxicillin (95%), and amikacin (90%). Streptococci isolates showed susceptibility to ciprofloxacin (60%), levofloxacin (90%), and clarithromycin (75%) and resistance to moxifloxacin (60%) and amikacin (60%).

E. coli isolates were susceptible to ciprofloxacin (90%) and clarithromycin (90%) and resistant to amikacin (90%), amoxicillin (95%), sulfamethoxazole (50%), and streptomycin (90%).







Frequency distribution of resistance and susceptible *P. Aeruginosa* towards A panel of antibiotics



Percent frequency distribution of resistant and susceptible of *streptococci* towards a panel of antibiotics









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Conclusion

S. aureus was the predominant pathogen isolated from breast abscesses in both lactational and non-lactational women. The bacterial isolates associated with breast abscess were *S. aureus*, *E. coli*, Streptococci, and *P. aeruginosa*. Age and gender had little or no effect on susceptibility to breast abscess.

Recommendations

Continuous screening and surveillance of breast abscess patients are recommended to prevent serious infections. Similar studies should be conducted in other healthcare facilities to provide a comprehensive understanding of the pathogens causing breast abscess in the region.

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