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PREVALENCE AND ANTIBIOTIC RESISTANT PATTERN OF STAPHYLOCOCCUS AUREUS IN CLINICAL ISOLATES FROM PESHAWAR: A SINGLE CENTER STUDY

KHAN WA¹, ULLAH J^{*2}, ULLAH MJ², ULLAH K³, RABNAWAZ M², ULLAH I², AHMAD N⁴, SHUMAILA⁵, ULLAH A²

¹Pakistan Red Crescent Society, National Headquarter Islamabad, Pakistan
²Department of Allied Health Sciences, Iqra National University Peshawar Pakistan
³Department of Pathology, Aga Khan Hospital Karachi, Pakistan
⁴Department of Surgery, Lady Reading Hospital Peshawar, Pakistan
⁵Department of Health and Biological Sciences, Abasyn University Peshawar, Pakistan
*Correspondence Author Email Address: jamshidawar0071@gmail.com

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Abstract The health authorities recognize S. aureus as a severe public health problem because of its varied pathogenicity while antibiotics lose their effectiveness against this pathogen. The main goal of this research was to analyze S. aureus resistance patterns among hospital patients from Peshawar's tertiary care facility in Pakistan. The research work was carried out in the Pathology section of Rehman Medical Institute from July 2022 to December 2022. The study processed 400 samples were obtained. The research team performed morphological as well as biochemical identification of the isolates followed by antibiotic susceptibility testing through Kirby-Bauer disk diffusion testing methods. The positive results for bacteria reached 78.25% from 400 collected samples and S. aureus emerged as one of the 10% types of isolates found. The study showed resistance towards Cefoxitin, Levofloxacin, and Erythromycin 74.35%, 69.23%, and 64.10% respectively. The bactericidal activities of Vancomycin, Linezolid, as well as Rifampicin, demonstrated high efficiency in combating bacterial pathogens. The research demonstrates antibiotic-resistant S. aureus strains have appeared in this region demonstrating the necessity of enhancing antibiotic-use strategies and steady monitoring for S. aureus resistance patterns to stop global S. aureus infection threats.

Keywords: Staphylococcus aureus; Antibiotic resistance; Methicillin-resistant Staphylococcus aureus (MRSA); Vancomycin; Antimicrobial stewardship

Introduction

Public health teams identify the gram-positive bacterium Staphylococcus aureus as an important threat because of its wide variety of infections that progress from skin surface infections to severe systemic diseases. Research indicates that 30% of people carry these bacteria without symptoms, yet they enable the bacterium to spread throughout healthcare facilities and community areas (Ahmad-Mansour et al., 2021). The discovery of methicillinresistant S. aureus (MRSA) in 1961 accelerated the problem of the bacteria developing into a significant pathogen that spread within hospitals and communities (Lawal et al., 2021; Wu et al., 2019). The resistance mechanism of MRSA arises from the mecA gene that produces the PBP2a protein, which makes the bacterium immune to β -lactam antibiotics. Medical treatment is challenging because resistance from patients surpasses 50% of treatments that use macrolides, fluoroquinolones, lincosamides, and aminoglycosides (Khatoon et al., 2018; Zaky et al., 2020).

The emergence of both VISA and VRSA strains of *S. aureus* creates advanced therapeutic hurdles for physicians (Lalitha *et al.*, 2020). The pathogenic capabilities of S. aureus intensify through biofilm formation because this mechanism helps the bacteria avoid immune response while blocking the entry of medication into the infection site (Lalitha *et al.*, 2020). Treatment of community-associated MRSA strains is challenging because these strains possess distinct virulence factors that make them difficult to combat (Hwang, 2023). Healthcare strategies require immediate approval of antimicrobial stewardship alongside constant monitoring of resistance patterns to reduce the global effects of *S. aureus* infections (Setiabudy, 2023).

Materials and Methods

The research followed a descriptive cross-sectional design conducted in the Pathology Department of Rehman Medical Institute (RMI) Peshawar from July 2024 to December 2024. All participants who had their blood samples tested in this department constituted the examined population. A total of 400

random blood cultures were collected and processed comprising patients of different ages, and both female and male patients. Individuals who received antibiotics before 72 hours of blood culture analysis were excluded from the experiment.

Specialized growth media within Bactec blood culture vials served as the collection medium for blood culture specimens. The laboratory followed standard protocols to create several bacterial culture media for identification purposes. Morphological identification was done by ZN staining while biochemical identification was performed by catalase, followed by tube and slide coagulase and DNase testing Figure 1. The Kirby-Bauer disk diffusion method was used to analyze antibiotic susceptibility according to the CLSI (2017) protocol Figure 2. A sterile saline suspension of bacterial colonies was used to prepare the Inocula until the solution reached a McFarland standard of 0.5. Antibiotic disks were applied to inoculated Mueller Hinton agar plates using an oxide antimicrobial susceptibility disk dispenser. Temperature-controlled conditions of 35-37°C maintained the plates for a timeframe extending from 18 h to 24 h. Bacterial sensitivity was determined by measuring inhibition zones using millimeter values.

Results

A total of n=400 single selected non-repetitive samples were collected to determine antimicrobial resistance in which 313 (78.25%) were positive and 87 (21.75%) showed no growth Figure 3. Male patients were 255 (63.75%), (p = 0.016) higher than (36.25%). The female patients 145 total staphylococcal positive samples in males and females were 27/255(10.58%) and 12/145(8.27%) shown in Figure 4. Both male and female patients were distributed in different age groups shown in Table 1 in which the highest number of patients that showed infections was under ten years. The pathogen distribution examined in this study displayed a substantial statistical difference (Chi-square = 73.1, p < 0.001) because Salmonella Typhi (14%) followed by skin contaminants (13%) and Staphylococcus aureus was (10%) Table 2. The susceptibility tests demonstrated Vancomycin's total effectiveness while Linezolid and Rifampicin showed excellent effects on the bacteria (100%, p < 0.001) Figure 5. The hospital needs to enhance antibiotic care management through strategic planning and ongoing antimicrobial resistance pattern monitoring due to emerging resistance against three important antibiotics: Cefoxitin (74.35%), Levofloxacin (69.23%), and Erythromycin (64.10%) respectively.

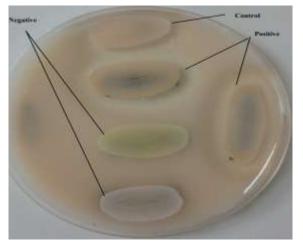
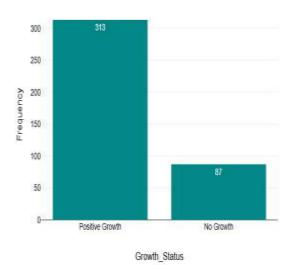


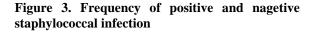
Figure 1. Positive and negative Staphylococcus aureus on DNase media



Figure 2. Antibiotic resistance behavior on Kirby-Bauer disk diffusion method

Frequency of Growth Status





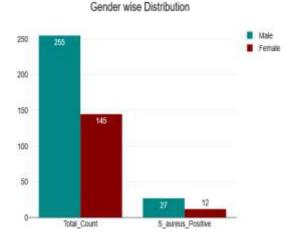
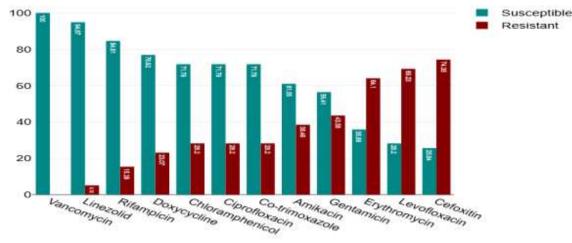
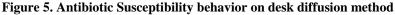


Table 1: Age wise distribution	
Age Group	Count (%)
<10 Years	120 (30%)
11-20 Years	85 (21.25%)
21-30 Years	75 (18.75%)
31-40 Years	70 (17.50%)
>41 Years	50 (12.50%)
Total	400 (100%)
Table 2: Pathogen Distribution	
Pathogen	Percentage
Skin contamination	13%
Salmonella Typhi	14%
Staphylococcus aureus	10%
Escherichia coli	9%
Others	54%
Total	100%

Table 1. Age wice distribution

Figure 4. Gender wise distribution of culture positive and negative patients





Discussion

A cross-sectional study performed at a Peshawar tertiary care hospital generated important data about bacterial pathogen occurrences as well as their antimicrobial resistance patterns for *Staphylococcus aureus*. A total of 78.3% of blood culture samples showed growth in the hospital along with S. aureus which emerged as one of the leading bacterial isolates with a frequency of 10% of the total samples. *S. aureus* is a major cause of bloodstream infections (septicemia) worldwide (Coombs and Others, 2020; Singh and Others, 2018). The studies show that pediatric patients under ten years old along with male patients make up the majority of reported cases therefore pediatric populations need specific intervention strategies.

Resistance testing demonstrated Vancomycin and Linezolid and Rifampicin maintained perfect effectiveness yet Cefoxitin presented moderate resistance (74.35%), while Levofloxacin showed 69.23% resistance and Erythromycin demonstrated 64.10% resistance to the bacteria. The research confirms previous studies showing increasing *S. aureus* isolate resistance in developing regions with active antibiotic misuse issues (Ansari and Others, 2014; Mnyambwa and Others, 2021). The study conducted by Naik *et al.*, (2020). showed comparable findings to this research by reporting high resistance levels against standard antibiotic treatments thus demonstrating a critical need for proper antibiotic management practices (Naik and Others, 2020).

A critical healthcare concern about methicillinresistant *Staphylococcus aureus* (MRSA) can be seen directly through the study results. Cefoxitin resistance serves as a public health warning since MRSA infections create higher mortality rates and worsen illness severity (Singh, 2018; Tarai *et al.*, 2013). Various investigations in different areas show that *S. aureus* clinical isolates commonly demonstrate MRSA resistance at rates surpassing 50% according to scientific findings (<u>Castro and Others, 2016</u>; <u>Mnyambwa and Others, 2021</u>). The widespread occurrence of MRSA requires continuous monitoring to support best infection control practices in healthcare facilities.

The research demonstrates positive findings about the antibiotic efficacy of Vancomycin and Linezolid since these drugs function as primary options to address severe S. aureus infections that include MRSA. The increasing detection of vancomycinintermediate S. aureus (VISA) strains in medical facilities represents an important threat to antibacterial treatment strategies according to research findings by Bhattacharya (Bhattacharya and Others, 2013). The antibiotic susceptibility patterns detected at Peshawar strongly emphasize ongoing antibiotic monitoring programs for developing treatment protocols that stop resistance strain spread. The study results about male patient infections match findings in other studies which demonstrate genderbased differences in S. aureus infection rates. The research conducted by Kengne et al. (2020) supports this finding by demonstrating male patients experience increased numbers of S. aureus skin and soft tissue infections (Kengne and Others, 2020). The findings support several explanations about male behavioral patterns as well as health conditions that create susceptibility to infections.

The identification of *Salmonella Typhi* at 14% among the leading isolates demands improved diagnostic programs for handling different bacterial infections effectively. The detection of *Salmonella* in blood cultures requires urgent implementation of suitable antibiotic treatments because it shows evidence of serious systemic infections (Bessa and Others, 2013). Different pathogens identified in the same patient group demonstrate how difficult it is to control infections inside hospitals so medical practices must use personalized antibiotic choices based on what each microorganism can survive.

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Declaration

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Author's contribution

WA K, and JU, conducted research and wrote the initial draft of manuscript. MJ, UA, S, MR, and IU, NA and AU collected the literature and wrote the manuscript, and edited the manuscript in original. All authors have read and approved the final manuscript. The author have read and approved the final manuscript.

Conflict of Interest

The authors state that there is no conflict of interests with regard to this study. There is no conflict of interest in any financial or personal manner concerning the development of this project, the gathering of data, the interpretation of the data, or the writing and publishing of this paper.

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Data Availability statement

All authenticated data have been included in the manuscript.

Ethics approval and consent to participate

These aspects are not applicable in this paper. **Consent for publication**

Consent for publicat





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