



A Study on VRSA Prevalence in Hospital Settings- Bangalore India

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Abstract

Objective: *Staphylococcus aureus* (SA) is one of the most common pathogenic Gram-positive bacteria. In this study, SA isolated from three different hospitals in Bangalore, India were tested against various antibiotics including vancomycin, ceftazidime, co-trimoxazole and ciprofloxacin for their sensitivity, especially against vancomycin to show the presence of vancomycin resistance *S. aureus* (VRSA) in clinical samples.

Methods: A total number of 25 samples from gauze bandage of skin lesions and wounds dressings were collected from various hospitals in Bangalore. The samples included pus swabs from skin lesions, swabs or open wounds from treated patients. These samples were inoculated into selective media for *S. aureus* (MSA-Mannitol salt agar) and 37 morphologically SA isolates were recovered. These putative SA isolates were then further characterized using traditional biochemical tests including Catalase, Oxidase, Coagulase and Mannitol fermentation tests. Subsequently, each *Staphylococcus aureus* isolates was tested against antibiotics for their susceptibility.

Results: A total of 37 isolates were characterized using the traditional biochemical test methods and 16 were identified as *S. aureus*. Antimicrobial resistance profiling using minimum inhibitory concentration (MIC) measurements showed that eight isolates of SA were resistant to vancomycin. and five isolates were resistant to ciprofloxacin.

Conclusion: Vancomycin resistant SA (VRSA) is becoming more prominent in hospitals in Bangalore, India.

Keywords: *Staphylococcus aureus*; VRSA; Vancomycin; Ciprofloxacin

Introduction

Staphylococcus aureus (SA) is among the most common pathogens isolated from skin, eyes, cerebrospinal fluid, blood, respiratory and gastrointestinal tract, bone, and connective tissues. *S. aureus* is the leading cause of community acquired skin and soft tissue infections such as abscesses (boils), furuncles and cellulitis. Skin infections are common, but bacteria can spread through the bloodstream infect distant organs [1]. Surgical site infections are a serious nosocomial infection and significantly contribute to patient morbidity and healthcare costs. When disseminated, SA is

responsible for diverse conditions such as toxic shock syndrome, osteomyelitis, meningitis, bacterial endocarditis. SA septicemia may enhance the mortality rate up to 40% [2]. Spread is by direct contact with an infected person, by a contaminated object, or by inhaling infected droplets dispersed by sneezing or coughing.

Many strains have developed resistance to antibiotics. Genetically bacteria use two main strategies to adopt for the antibiotic activity; i. *De novo* mutation in the genome where it interferes with the mechanism of activity of the compound, ii. Exchange of the for-

eign DNA containing resistant determinants via horizontal gene transfer (HGT) [3]. *De novo* mutations can alter target sites causing them to be insensitive to the action of antibiotics or causes activation of latent resistant mechanisms. One frequent *de novo* mutation to be observed is mutations in penicillin binding proteins which renders them to resistant to beta lactams. Moreover, some antibiotic efflux pumps are repressed at the transcription level. Mutations in repressor proteins activate efflux pump expression resulting in broad spectrum antimicrobial resistance. SA uses all three methods of HGT: Conjugation, Transformation and Transduction.

Beta lactams are drug of choice for treating bacterial infections due to their low toxicity and high therapeutic indices. SA uses few mechanisms to combat the action of beta lactams. Beta lactamases are enzymes that open the lactam ring in the active site of the drug and render them inactive. Plasmid bourn or chromosomal located beta lactamase BlaZ confer resistance to broad spectrum of beta-lactamses such as first and second generation cephalosporins and commonly used penicillins such as benzyl penicillins and ampicillin derivatives and second or third generation penicillins. The advent of Beta lactamase resistant penicillins such as methicillin and its derivatives against SA was a key event in therapy but due to their heavy use, methicillin resistant SA (MRSA) became a problem in both nosocomial and community acquired SA infections. Methicillin resistance is due to the presence of a heterologous horizontally acquired homolog of penicillin binding protein 2 (PBP2) which cannot be inhibited by beta lactams.

Centers for Disease Control and Prevention (CDC) reported a prevalence of SA in hospital infections of about 18% [4]. Asia has been reported as a continent which has higher number of MRSA in the world [5]. Vancomycin-intermediate *S. aureus* (VISA) strains and vancomycin-resistant *S. aureus* (VRSA) strains are also being increasingly identified in certain countries in this region [6,7]. A fast-growing number of VRSA mostly evolved from MRSA has also emerged and thus both Methicillin and vancomycin [8,9]. It was identified that excess accumulation of peptidoglycan which thicken the cell wall due to unidentified mutations paves the way for the resistance against vancomycin by *S. aureus* making it's the most common resistance mechanism for VRSA strains [10].

MRSA infection that is acquired in a hospital is treated with antibiotics that are effective against MRSA. MRSA infections can be treated with less toxic antibiotics such as trimethoprim-sulfamethoxazole, 4-amino quinolones or cephalosporins which are

relatively cheaper and can be orally administered. Vancomycin and daptomycin are the first-line therapies for MRSA bacteremia if drug resistance profiles indicates other simpler and cheaper antimicrobials are useless against MRSA [11]. MRSA isolates that are also multidrug resistant and also resistant to vancomycin are treated with antimicrobials which are recently developed or expensive. Some of these are linezolid, tedizolid, quinupristin plus dalfopristin, ceftaroline, telavancin, or daptomycin. Daptomycin, which is a lipopeptide with a poorly characterized antimicrobial mechanism has become the main stay therapy against MRSA which are also resistant to vancomycin.

Two shifts in epidemiology of SA can be witnessed in last past two decades: first, a growing number of health care-associated infections, particularly seen in infective endocarditis and prosthetic device infections, and second, an epidemic of community-associated skin and soft tissue infections driven by strains with certain virulence factors and resistance to β -lactam antibiotics [12]. On the other hand, Ciprofloxacin has been used invitro experiments in treating against VRSA [13] but further research is ongoing on whether it could be a better option on treating VISA and VRSA strains. Considering the recent studies showing the emergence of VRSA, we determined the prevalence of VISA and VRSA strains in three different hospitals in Bangalore, India and additionally compared the resistance profile between vancomycin and Ciprofloxacin.

Materials and Methods

Sample collection and isolation of *Staphylococcus aureus*

A total number of 25 samples from different gauze bandages with skin lesions and wounds were collected from various hospitals in Bangalore which includes Cratis hospital, Avietha and Chris hospital. Swabs of pus from wounds and surgical sites were collected after dressings. Aseptic plating on nutrient agar was done for all samples in the laboratory. About 37 different isolates was derived from the initial plates using colony morphology and Gram staining morphological characteristics.

Mannitol fermentation test and other biochemical tests

All 37 Gram-positive isolates were plated in Mannitol Salt agar (a selective and differential media for SA) and incubated in 37 degrees Celsius for 24hours. The observation of growth and color change from red to yellow indicates fermentation of Mannitol by the bacterial culture. The pH indicator used is phenol red, therefore the positive color change from red to yellow indicates acid produc-

tion and pH change in the media by bacteria. Subsequently, further biochemical tests were carried out which included catalase, oxidase, Indole, Methyl red, Voges-Proskauer, Citrate utilization test (IMViC) and Coagulase test.

Test	Medium	Reagent	Result
Indole	Tryptophan broth	Ko Vac's reagent	Positive
Methyl-red	methyl red broth	Methyl red reagent	Positive
Voges-Proskauer	Voges-Proskauer broth	Barritt's A and Barritt's B	Positive
Citrate	Simmons citrate agar		Positive
Catalase	Saline solution	Hydrogen Peroxide	Positive
Oxidase	Oxidase disc		Negative
Mannitol Fermentation	Mannitol Salt agar	phenol red	Positive
Coagulase	Blood Agar media		Positive

Table 1: Summary of biochemical tests that were carried out to differentiate SA from other Gram-positive cocci.

Bacterial antimicrobial sensitivity tests.

All samples were plated on Muller Hinton agar media and grown for antimicrobial resistance profiling. To determine vancomycin resistance, disc diffusion method using vancomycin discs (Va30 Hi-media) was used. Moreover, using disc diffusion method, four other laboratory antibiotics including cefoxitin (Cx30), ketoconazole (Kt30), co-Trimoxazole (Cot25) and ciprofloxacin (Cip5) were tested against SA isolates for the MIC determination. Measurement of each zone of growth inhibition was measured and the diameter was recorded as millimeters for MIC determination.

Results and Discussion

In this study, a total of 25 samples were taken from three different hospitals in Bangalore, India. 37 isolates were derived from these samples and sixteen (48.6%) were identified as SA using variety of biochemical tests (Table 2). MIC determination indicated that eight (50%) were resistant to vancomycin while the other half (51.4%) were susceptible to vancomycin (Table 3). Further testing with 4 other antimicrobials showed 100% resistance to cefoxitin (Cx30), ketoconazole (Kt30) and Co-Trimoxazole (Cot25), but surprisingly, five of the SA isolates that are VRSA showed sensitive to Ciprofloxacin (Cip5).

Bacterial isolates	Media	Citrate test	Indole test	Vogues Pr test	Methyl red test	Oxidase test	Catalase test	Coagulase	Mannitol fermentation
CH1	N.A, MSA	-	-	+	-	+	-	-	-
CH 10-3	N.A, MSA	-	+	-	+	+	+	-	-
CH 10-4	N.A, MSA	-	+	-	-	+	-	-	-
CH3	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
A	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
AV1 10-3N	N.A, MSA	+	-	+	+	+	+	-	-
AV1 10-3M	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
AV1 10-4M	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
AV2 10-3	N.A, MSA	-	+	-	-	+	-	-	-
AV3 10-3	N.A, MSA	-	-	-	+	+	-	-	-
AV2 10-4	N.A, MSA	-	+	-	-	+	-	-	-
CR1 10-3 M	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
CR1 10-4 M	N.A, MSA	-	+	+	+	+	+	-	+
CR2 10-3	N.A, MSA	-	-	-	-	+	-	-	+
CR2 10-4	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+

CR3 10-3	N.A, MSA	+	-	-	-	+	-	-	-
CR4	N.A, MSA	+	+	+	-	-	+	-	-
CR5	N.A, MSA	+	+	+	-	+	-	-	-
CR6	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
CR7	N.A, MSA	+	+	+	-	+	-	-	-
CR8	N.A, MSA	-	+	+	-	+	-	-	-
B1CR4	N.A, MSA	-	+	-	-	+	-	-	-
B2CR5	N.A, MSA	-	+	-	-	+	+	-	-
B3CR6	N.A, MSA	+	-	+	+	+	-	-	-
B4CR7	N.A, MSA, Blood Agar	+	-	-	-	-	+	+	+
B5CR8	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
B6CR9	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
B7CR1	N.A, MSA	-	+	-	-	+	+	+	-
CR11	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
CR12	N.A, MSA	-	+	-	-	+	-	+	-
CR13	N.A, MSA, Blood Agar	+	-	-	+	-	+	+	+
CR14	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
CR15	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
CR16	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
UK	N.A, MSA, Blood Agar	+	-	+	+	-	+	+	+
10-3M	N.A, MSA	+	+	+	+	+	-	-	-
10-4M	N.A, MSA	-	+	-	-	+	-	-	-

Table 2: Biochemical test results for identification of SA strains in patient samples.

<i>S. aureus</i> culture	Vancomycin (zone of inhibition) in cm	Cefoxitin Cx30 (zone of inhibition) in cm	Ketoconazole Kt30 (zone of inhibition) in cm	Co-Trimoxazole Cot25 (zone of inhibition) in cm	Ciprofloxacin Cip5 (zone of inhibition) in cm
B5	0	0	0	0	1.3
b6	2	0	0	0	2.5
A	0	0	0	0	0
Ch3	0	0	0	0	0
AV10-3m	0	0	0	0	1.1
Cr10-3M	1.6	0	0	0	1.6
Cr2.10-4	0	0	0	0	0
cr15	0	0	0	0	0
UK	2.1	0	0	0	2
Cr16	1	0	0	0	1.6
Cr13	0	0	0	0	0.6
cr11	1.8	0	0	0	2
Av10-4m	1.3	0	0	0	2.3
Cr6	1.1	0	0	0	1.7
Cr14	1.1	0	0	0	2
B4	0	0	0	0	0

Table 3: MIC results derived from disc diffusion assays. The bacterial clearing diameter is recorded in centimeters measured from the edge of the antimicrobial disc.

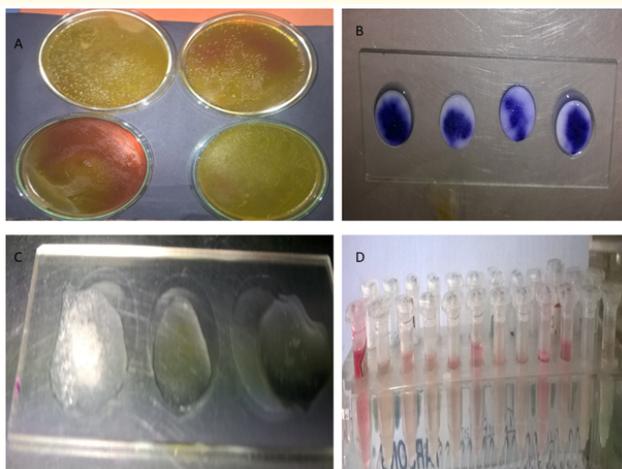


Figure 1: (A) Showing Mannitol fermentation test. (B) Shows oxidase test. (C) Shows Catalase test. (D) Shows oxidase test.

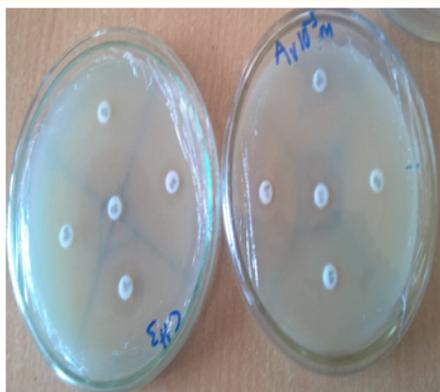


Figure 2: Plates showing clear zones of inhibition and no zones of inhibition.

Conclusion

This study shows prevalence of vancomycin resistance strain of *Staphylococcus aureus* from clinical isolates derived from hospitals in Bangalore, India. The high percentage (50%) of VRSA strain shows that vancomycin use should be always preceded by MIC testing for its efficacy. There wasn't a definite correlation between the hospital locality or that of the patients from the sampling en-

viron. In comparison to previous years, the resistance of *S. aureus* strain to vancomycin has been increasing, indicating that thorough MIC determination against antimicrobials is paramount important in hospital practices. Moreover, we demonstrated that Ciprofloxacin could be an alternative antimicrobial against VRSA. Ciprofloxacin is relatively cheaper and orally administered and has less serious side effects than both vancomycin and daptomycin. This study should be followed up by a more extensive cross-sectional studies characterizing the antimicrobial resistance profiling using vancomycin, methicillin, daptomycin and ciprofloxacin to determine the MIC profiling of MRSA and VRSA strains. This study which demonstrated (i) vancomycin resistance has high prevalence and, (ii) some of vancomycin resistant strains are sensitive to quinolones could be the basis for a larger prevalence study in nation-wide study for recording VISA and VRSA strains in India.

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