

Study of the virulence factors and antimicrobial susceptibility profile of staphylococcus aureus isolated from clinical samples at tertiary care centre

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Abstract

Introduction: Staphylococcus aureus is an important cause of healthcare-associated infections. The increasing resistance of this pathogen to various antibiotics complicates treatment. Present study was conducted to isolate staphylococcus aureus from various clinical samples, to study the few virulence factors, antibiogram and to detect methicillin resistance among them. **Materials and Methods:** Present study was done over a period of six months. The study was conducted on 100 staphylococcus aureus isolated from various clinical samples. The organisms were identified as per standard conventional methods. The antibiotic sensitivity testing of the isolates was done by Kirby Bauer's Disk Diffusion method according to CLSI guidelines. These strains were screened for DNase production, beta hemolytic property and slime production. **Results:** Among 100 clinical isolates of staphylococcus aureus 82% of them showed beta hemolysis, 79% produces DNase, 72% produces slime. The proportion of MRSA amongst S. aureus isolates was found to be 53%. MSSA were 100% sensitive to vancomycin and linezolid 85.10% to Doxycycline, 82.98% to clindamycin, 72.34% to chloramphenicol, 68.09% Co- trimoxazole, 65.95% to erythromycin, 61.70% gentamicin, 51.06% to Ciprofloxacin, 12.77% topenicillin. MRSA were 100% sensitive to vancomycin and linezolid, 71.69% Doxycycline, 64.15% to Chloramphenicol, 60.37% to Clindamycin, 43.39% to Ciprofloxacin, 39.62% Gentamicin, 37.74% Co- trimoxazole, 35.84% to Erythromycin and all the isolates were resistant to penicillin. **Conclusion:** Understanding of virulence mechanisms and antibiotic susceptibility pattern of Staphylococcus aureus is important for effective management of infections.

Key words: Antibiotic sensitivity, D Nase test, Staphylococcus aureus, Slime production, Virulence factors.

Introduction

Staphylococcus aureus is one of the most prevalent and clinically significant pathogen, causing variety of infections ranging from mild skin and soft-tissue infections to serious life-threatening systemic infections and is a leading cause for hospital associated (HA) and community associated (CA) infections worldwide [1,2]. It causes skin, bone, urinary tract infections, soft tissue infections, pneumonia, bacteremia and other invasive infections in community and hospital settings [3].

Emergence of methicillin resistance among Staphylococcus aureus has reduced therapeutic alternatives available to treat staphylococcal infections [4]. Methicillin-resistant S. aureus (MRSA) infections

account for 40-60% of all health care associated S. aureus infections in many centres across the world [1]. Report from the National Nosocomial Infections Surveillance System of the Centers for Disease Control and Prevention, showed that MRSA in India and USA accounts for > 60% of S. aureus isolates causing health care associated infection in ICUs. Drug-resistant strains limit the therapeutic options, creating an economic and social burden to the healthcare system [3].

Virulence factors of S. aureus include slime formation, secreted enzymes, toxins like lipases, DNase, proteases, hemolysins and some super antigens like toxic shock syndrome toxin and enterotoxin [5]. The pathogenicity of S. aureus depends on various virulence factors associated with adherence, evasion of the immune system and damage of the host [6,7]. The presence of

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two or more virulence factors could increase the pathogenic ability of organism in relation to those that express only one virulence factor [8].

It is important for treating physicians to know the virulence factors and antibiotic susceptibility patterns of *S. aureus* isolates in their region as it shows differences among regions and it will help the clinician to choose appropriate antibiotic and to control the emergence of drug resistant strains.

Materials & Methods

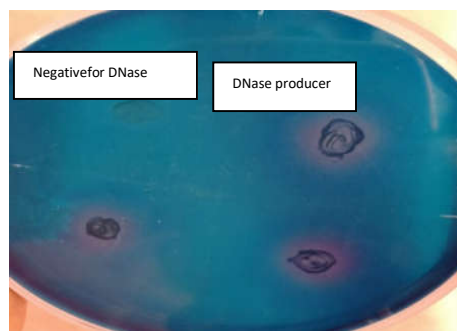
Place of study: Present study was done at Shimoga Institute of Medical Sciences, Shivamogga, for six month duration from January 2018 to June 2018.

Ethical consideration & permission: Study was done after obtaining institutional ethics committee clearance.

Statistical Methods: All Data were analysed and expressed in terms of percentage.

Collection of Samples and processing: The study was conducted on 100 staphylococcus aureus isolates from various clinical samples like blood, exudates, urine and respiratory samples received at Microbiology laboratory from Mc Gann teaching hospital, attached to Shimoga institute of medical sciences. These samples were processed on blood agar and Mac Conkey agar media and incubated at 37°C under aerobic conditions. The organisms were identified as per standard conventional methods [9].

Detection of DNase: This test was carried out by using DNase agar. The organisms were spot inoculated in DNase agar and incubated at 37°C for 24 hours. Then flood the plate with 1N HCl. After standing a few minutes, examine the plate against a dark background. Clearing zone around the spotted colony is considered as positive for DNase [5,9,10].



Detection of beta hemolysin: Staphylococcus aureus were spot inoculated onto 5% sheep blood agar and incubated at 37°C overnight. After overnight incubation the plates were kept at 4°C to observe hot-cold type of hemolysis produced by beta hemolysin [5].



Detection of Slime formation: Slime formation was detected using congo red agar (CRA) containing brain heart infusion broth, 5% sucrose, agar and 0.08% congo red. Strains were inoculated and plates were incubated at 37°C

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overnight. Dry black colonies of *Staphylococcus aureus* strains with crystalline consistency were considered as slime producers and those that produced pink colonies were considered as negative for slime formation [5,11].



Antibiotic susceptibility testing: Antimicrobial susceptibility testing of isolates was done on Mueller-Hinton agar by Kirby-Bauer disc diffusion method, according to the Clinical and Laboratory Standards Institute (CLSI) guidelines. In the present study the susceptibility testing was carried out using the following antibiotics: Penicilline (P) Co-trimoxazole (COT), Clindamycin (CD), Erythromycin (E), Gentamicin (G), Chloramphenicol (c), Ciprofloxacin (CIP), Doxycycline (DO), Vancomycin (VA), Linezolid (LZ), Cefoxitin (cx)[12].

Detection of MRSA: Methicillin resistance was detected by disc diffusion method using 30µg cefoxitin disc which remains as a surrogate marker to identify *mecA* mediated resistance. The strains were considered as methicillin susceptible (MSSA-Methicillin susceptible staphylococcus aureus) if the zone of inhibition is ≥ 22 mm and considered as MRSA (Methicillin resistant staphylococcus aureus) if the zone of inhibition was ≤ 21 mm [12].

Results

Present study was carried out in the department of microbiology, Shimoga Institute of Medical Sciences, Shivamogga. The observations made from the study are shown in following tables.

Table-1: Staphylococcus aureus isolates obtained from various clinical specimens.

Different clinical specimens	Staphylococcus aureus isolates (%)
Exudate	69(69%)
Blood	21(21%)
Urine	6(6%)
Respiratory samples	4(4%)
Total	100(100%)

Among 100 clinical isolates of staphylococcus aureus 69% were from exudates samples, 21% from blood, 6% from urine, 4 from respiratory samples.

Table 2: Distribution of various virulence factors among staphylococcus aureus

Virulence factors	No of isolates N- 100	%
Beta hemolysin	82	82%
DNase production	79	79%
Slime production	72	72%

Among 100 clinical isolates of staphylococcus aureus 82% of staphylococcus aureus showed Beta hemolysis, 79% produced DNase, 72% were slime producers.

Table-3: Antibiotic sensitivity pattern of Staphylococcus aureus.

Antibiotics tested	Organisms isolated	
	MSSA N=47 (%)	MRSA N=53 (%)
Penicillin	6 (12.77)	0
Co-trimoxazole	32 (68.09)	20 (37.74)
Clindamycin	39 (82.98)	32(60.37)
Ciprofloxacin	24(51.06)	23(43.39)
Cefoxitin	47(100)	0
Chloramphenicol	34 (72.34)	34(64.15)
Doxycycline	40(85.10)	38(71.69)
Erythromycin	31 (65.95)	19 (35.84)
Gentamicin	29 (61.70)	21(39.62)
Linezolid	47 (100)	53(100)
Vancomycin	47(100)	53 (100)

The proportion of MRSA among Staphylococcus aureus isolates was found to be 53%. MSSA were 100% sensitive to vancomycin and linezolid, 85.10% to Doxycycline, 82.98% to clindamycin, 72.34% to chloramphenicol, 68.09% Co-trimoxazole, 65.95% to erythromycin, 61.70% gentamicin, 51.06% to Ciprofloxacin, 12.77% to penicillin. MRSA were 100% sensitive to vancomycin and linezolid, 71.69% Doxycycline, 64.15% to Chloramphenicol, 60.37% to Clindamycin, 43.39% to Ciprofloxacin, 39.62% Gentamicin, 37.74% Co-trimoxazole, 35.84% to Erythromycin and all the isolates were resistant to penicillin. The antibiotic sensitivity results showed that all MRSA isolates were significantly more resistant to antibiotics as compared to MSSA isolates.

Discussion

MRSA is an important clinically significant pathogen, incidence of which is increasing every year. The organism has an ability to spread and cause outbreaks. Knowledge about virulence factors and prevalence of staphylococcus aureus and their current antimicrobial profile is necessary in the selection of appropriate empirical treatment for these infections, formulation of antibiotic policy, infection control, and patient management [13,4,14].

In our study, 79% of the staphylococcus aureus isolates were positive for DNase test. Another study by Kateete David P et al., reported 75% positivity for DNase test [15]. Study by Alice P. Selvabai et al. reported 86% positivity for the DNase test among MRSA isolates [5]. In the present study 82% of the Staphylococcus aureus isolates showed beta hemolysis. Study by Alice P. Selvabai et al. reported 67% of the MRSA isolates showed beta hemolysis [5]. In our study 72% of the staphylococcus aureus strains were found to be slime producers. Study by Adrianna Podbielska et al. reported 69% of staphylococcus aureus were slime producer [11]. Study by Alice P. Selvabai et al. reported 61% were slime producers among MRSA isolates [5]. Slime producing strains of Staphylococcus aureus has the

ability to form intact biofilm and also have higher rate of colonization in host tissues. Slime production is considered as a significant virulence factor for some strains of staphylococci [16]. The pathogenicity of S. aureus depends on various virulence factors associated with adherence, evasion of the immune system and damage of the host [6,7].

In our study the proportion of MRSA among Staphylococcus aureus isolates was 53%. MSSA were 100% sensitive to vancomycin and linezolid, 85.10% to Doxycycline, 82.98% to clindamycin, 72.34% to chloramphenicol, 68.09% Co-trimoxazole, 65.95% to erythromycin, 61.70% gentamicin, 51.06% to Ciprofloxacin, 12.77% to penicillin.

MRSA were 100% sensitive to vancomycin and linezolid, 71.69% Doxycycline, 64.15% to Chloramphenicol, 60.37% to Clindamycin, 43.39% to Ciprofloxacin, 39.62% Gentamicin, 37.74% Co-trimoxazole, 35.84% to Erythromycin and all the isolates are resistant to penicillin. Study done by Rajadurai pandiet al reported almost all clinical MRSA strains (99.6%) were resistant to penicillin, 93.6% to ampicillin and 63.2% towards gentamicin, co-

trimoxazole, cephalixin, erythromycin, and cepho-taxime [13]. In a Study done by Gitau et al. A total of 944 *S. aureus* isolates were analyzed. High sensitivity of *S. aureus* was observed for quinupristin/ dalfopristin (100%), tigecycline (98.2), imipenem (98%), nitrofurantoin (97.6%), linezolid (97.3%), teicoplanin (97.1%) and vancomycin (95.1%). High resistance was recorded against penicillin G (91.9%), trimethoprim/ sulfamethoxazole (56.9%) and tetra-cycline (33.2%). MRSA prevalence was 27.8%. Both MRSA and MSSA were highly susceptible to quinupristin/ dalfopristin, tigecycline, linezolid, nitrofurantoin, ampicillin/ sulbactam and vancomycin and showed high resistance to commonly used antibiotics such as gentamycin, erythromycin, levofloxacin and tetracycline [3].

In a study done by Kitara LD et al. reported antibiotic susceptibility to Ampicillin (75.0%), Chloramphenicol (34.4%), Ciprofloxacin (1.6%), Erythromycin (7.8%), Gentamycin (0%), Methicillin (1.6%), Tetracycline (45.3%) and Co-trimoxazole (50.0%)[17]. Study by Alain C. Juayang et al evaluated the antimicrobial resistance of *S. aureus* isolated from clinical specimens and to put emphasis on the prevalence of MRSA and Inducible Clindamycin Resistance.

A total of 94 cases from 2010 to 2012 were diagnosed to have *S. aureus* infection using conventional bacteriologic methods. From these cases, 38 (40.6%) were identified as MRSA.

Wounds and abscesses were considered to be the most common specimens with MRSA infections having 71.05% while blood was the least with 5.3%. For drug susceptibility, out of the 94 *S. aureus* cases, including MRSA, 100% were susceptible to linezolid. It was then followed by tetracycline having a mean susceptibility of 95%; while penicillin G was ineffective with 94 cases having 0% susceptibility [18].

Study by Edet E. Udo et al showed All MRSA isolates were susceptible to linezolid, vancomycin, and teicoplanin. However, some isolates were resistant to kanamycin (2,979; 43%), ciprofloxacin (2,955; 42.7%), erythromycin and clindamycin (2,935; 42.4%), fusidic acid (2,858; 41.2%), gentamicin (2,665; 38.5%), tetracycline (2,652; 38.3%), and trimethoprim (2,324; 33.5%).

Whereas the prevalence of resistance to most antibiotics showed annual variations, those resistant to chloramphenicol and rifampicin increased from 2.6 and 0.1% to 9.6 and 1.6%, respectively, and high-level mupirocin resistance declined from 9.3% in 2011 to 3.6% in 2015 [19].

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For the effective management of *Staphylococcus aureus* infection, knowledge about virulence factors and antibiotic sensitivity patterns play a vital role. Conducting regular studies to know the changing trends of antibiotic sensitivity will help the treating physician to start empirical therapy.

Conclusion

MRSA is an important cause of healthcare-associated and community associated infections. It is important to do culture and sensitivity of specimens when *Staphylococcus aureus* infection is suspected. Continuing surveillance to assess the prevalence, geographic distribution, antibiogram, effective infection control measures are necessary to reduce the incidence of infections due to methicillin resistant *Staphylococcus aureus* and it will also help the clinician to choose appropriate treatment options and to control the emergence of drug resistant strains.

What this study adds to existing knowledge? The publication of such surveillance study, data describing regional antimicrobial susceptibility/ resistance rates in clinical isolates of *Staphylococcus aureus* essential to stimulate antimicrobial stewardship efforts as well as to identify emerging resistance trends and geographic diversity over time.

Contribution by authors

- Study concept, Data collection, Manuscript writing, Data compiling, literature review: Dr. Vedavati B. I.
- Study concept, Manuscript writing: Mallikarjun koppad.
- Manuscript writing: Dr. D. E Premalatha.
- Data collection, Manuscript writing, Data compiling: Akshatha. Y. J.
- Manuscript editing, final review and approval: Dr Halesh L.H.

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