

Molecular Identification of 16S rRNA gene in Staphylococcus aureus Isolated from Wounds and Burns by PCR Technique and Study Resistance of Fusidic acid

Shaymaa Khudhr Al-Alak, Darem kadhim Qassim

University of Al-Mustansiriyah \ College of Science \ Department of Biology

Abstract:

Thirty six samples were collected from patient suffering from wounds and burns in Baghdad Hospitals. The samples were analyzed for the presence of Staphylococcus aureus. Ten positive samples screened by phenotypic microbiological and Biochemical tests and genotypic by DNA extracted from all isolates and the PCR carried out using 16S rRNA gene for S. aureus. In this method for identifying and confirmed all the staphylococcal isolates as S. aureus. The present study was carried out in an attempt to detect the distribution of antibiotic-resistant and we have tried to cover the ever increasing problems facing the treatment and containment of bacterial skin infections. Therefore we are study Susceptibility of Staphylococcus aureus to antibiotics. The results showed relatively high resistance to Methicillin ME, Amoxicillin-Clavilonic acid, Azithromycin with frequencies of 100%, 100% and 70% respectively and moderate resistance to Vancomycin VA, Amikacin AK with frequencies of 60% and 62% respectively, low resistance to Fusidic acid with frequencies of 30% because that we determined the Minimum Inhibitory Concentration for Fusidic acid, the isolates incorporated in this test were those have intermediate resistant to Fusidic acid.

Keyword: wounds and burns, Staphylococcus aureus, 16S rRNA gene, PCR. Fusidic acid.

Introduction:

Staphylococcus aureus is a Gram positive coccus (Steinberg et al., 1996) with circular chromosome located on those pathogenesis and antibiotic-resistant genes (Novick, 1990). It is one of the most important bacteria in Micrococcaceae family, which is also responsible for a wide variety of community- and hospital-acquired infections (Abed EI-Jalilet al., 2008). This bacterium is one of the most versatile organisms. It is found world wide and is a leading cause of disease. Even though it is not classified as a true pathogen (an organism that is expected to always cause disease in humans), (Abed EI-Jalilet al., 2008). but an opportunistic pathogen, it has a diverse repertoire of possible infections. Normally, it is a transient colonizer of the skin and body entry portals (ears, eyes, nasal passages, etc.), and an estimated

20% of humans are carriers (asymptomatic permanent colonization). However, any break in the skin, or colonization of individuals with compromised immune systems can provide an opportunity for this bacteria to cause infection.

The disease process can be mediated via two possible mechanisms (Carboneau et al.; 2010) Staphylococcus aureus from the Staphylococcus genus of Micrococcaceae family is a leading cause of community-acquired infections in humans and a cause of mastitis and skin diseases in milk producing animals (Ebrahim Rahimiet al., 2013). S. aureus is commonly resident on skin and mucous membranes of cattle (Kluytmans, J. A 2010). Despite its generally benign nature, in changing circumstances, as response to damage and exposure of structures below the epithelial or mucosal surface, S. aureus can behave as a pathogen (Ebrahim Rahimiet al., 2013; Kluytmans, J. A 2010). Illness through S. aureus range from minor skin infection such as pimples, boils, cellulites, toxic shock syndrome, impetigo, and abscesses to life threatening disease such as pneumonia, meningitis, endocarditis and septicemia (Adwan, G ;2005). The majority of skin and

Corresponding Address:

Shaymaa Khudhr Al-Alak

University of Al-Mustansiriyah \ Collage of science \ Department of biology

Email: Shaymaakhudhr@yahoo.com

soft tissue infections are abscesses or cellulitis, but up to one quarter are superficial infections such as impetigo (Naimiet al., 2001).

Fusidic acid belongs to the fusidanes, which have molecular structures similar to corticosteroids without the steroid-like effects (Wilkinson JD., 1998). It is able to achieve a high penetration and concentration at the site of infection, and is highly effective against *S. aureus*. Many guidelines suggest fusidic acid as first line in the treatment of superficial skin infections and infected eczema, as the main bacterial culprit is *S. aureus*. (Abeck D., 1998). Topical fusidic acid and mupirocin appear to be equally effective in cases of primary cutaneous infections (Morley P., 1988). Both ointments appear to be effective against Gram-positive, Gram-negative or a combination of these organisms. The only adverse effect was that of greasiness, which was higher in the mupirocin group (Perera. G and Hay. R.; 2005).

Recently many PCR based molecular methods were developed as an alternative ways for accurate identification (Emelet al; 2010). Amplification of 16S rRNA gene sequences (228 bp) is the most commonly used method for identifying and classifying bacteria, including staphylococci (Mohammad et al; 2007).

The aim of present study was to Presence of *S. aureus* in patients wounds and burns, can be detected by simple molecular analysis of 16S rRNA gene. Based on the culture methods and PCR technique and study the resistance for fusidic acid due to the most common topical treatment for wounds and burns.

Materials and Methods:

Bacterial Isolation:

Thirty six samples were obtained from patient suffering from wounds and burns in Baghdad during November 2015 obtaining 10 from isolate as *Staphylococcus aureus*. Swabs of wounds and burns samples were collected in sterilized screw-cap containers, then the samples were inoculated on culture media (Mannitol agar and blood agar) and incubated aerobically at 37°C for 24hr. Biochemical tests were performed by using API 20 E system which provides a typical identification for all isolates.

Antibiotic Susceptibility Testing

Stock solution of the antibiotics Fusidic acid was prepared at final concentration 10 mg/ml according to Clinical Laboratory Standard Institute (CLSI) recommendations using distilled water. Sterilized by filtration and stored at -20°C

in small container. Susceptibility testing was conducted by disc diffusion according to the guidelines of the Clinical and Laboratory Standards Institute (CLSI 2013). Antibiotic discs (Becton Dickinson, USA) were placed on Mueller-Hinton agar plates, incubated at 37°C for 24 h, and the diameter of each zone was measured in millimeters. The following antibiotic discs were used: 8 Antimicrobial disks tested included Methicillin ME (10µg), Vancomycin VA (10µg), Azithromycin Azi (15 µg), (30 µg), Fusidic acid; FA (10µg), Amikacin AK (10 µg), Amoxicillin + clavulanic acid (30 µg).

Determination of the Minimum Inhibitory Concentration (MIC):

Stock solution of the antibiotics Fusidic acid was prepared at final concentration 10 mg/ml according to Clinical Laboratory Standard Institute (CLSI) recommendations using distilled water. MIC was determined for all isolates according to the CLSI (2013) criteria by a standard agar dilution method as follows: Serial two fold dilution of each antibiotic being tested ranged from 0.5 to 256 µg/ml were prepared of (Fusidic acid). Mueller-Hinton agar medium was prepared, sterilized by autoclave and cooled to 45°C, then the antibiotics were added in appropriate amount from their stock solution, mixed well and poured into the plates. Few colonies (2-4) from overnight culture were transferred to 2ml of normal saline in order to prepare the bacterial suspension and were adjusted to 0.5 McFarland turbidity equal to 1.5×10^8 CFU/ml. Five microliter of each inoculum was spotted on the agar surface by micropipette. The plates were left to dry for 5 minutes and then incubated at 37°C for 18-24 hr. Mueller-Hinton agar medium without any antibiotic was used as a positive control. MIC results were read after 18-24 hr. The lowest concentration of antibiotic inhibiting the bacterial growth was recorded as the MIC.

Genomic DNA extraction

Total DNA was extracted from *S. aureus* grown on mannitol salt agar plates by using the boiling approach as described previously (Zhang et al., 2004). In brief, for rapid DNA extraction, one to five colonies of each freshly subcultured strain were suspended in 50 µl sterile distilled water and heated at 99°C for 10 min. After centrifugation at 30,000 xg for 1 min, the supernatant was used as a DNA template and stored at -20°C until PCR was performed.

PCR amplification procedure

The sequence of oligonucleotide primers that were used in PCR to detect the presence of 16s rRNA gene were synthesized in Alpha DNA (Canada). Table (1) shows primer sequence and product.

Table (1): Sequence and molecular size of PCR products of 16s rRNA

Gene	Primer Sequence	Size of product (bp)	Cof
16srDNA	F : GTAGGTGGCAAGCGTTACC R : CGCACATCAGCGTCAG	228	Mastafa Al-musawiet al., (2014)

Attend each solution separately primer concentration of (10) Pmol/μl by dissolving 10μl of stock solution for each primer and add to 90μl distilled water unionic mix well and keeping in the ice until use .the stock solutions of the prim-

ers were store in -20 ,then mixing by vortex to homogenized before uses.

Primers obtained from Alpha DNA (USA) using the conditions in table 2 using Conventional PCR

Table 2: The conditions used for the amplification of 16s rRNA gene by Conventional PCR.

Initial denatur-ation	Denaturation in each cycle	Annealing	Primers exten-sion	Final extension
94 °C, 5 min	94°C, 1 min	51.5°C,30s	72°C, 1min	72 °C,10 min
	30 cycles			

Amplification products then separated by electrophoresis on agarose gel 1% stained with ethidium bromide (0.5 μg/ml) in Tris - borate - EDTA TBE buffer. The bands were detected using UV transillumination.

Gradient PCR amplification procedure

The reaction was performed using PCR thermal cycler apparatus. The following program was adopted:-PCR consisted of a preheating at 94°C after this initial denaturation step, the mixture was subjected to 30 amplification cycles as follows:-

Table 3: The Program of gradient PCR for 16s rRNA gene by Conventional PCR.

Initial denatur-ation	Denaturation in each cycle	Annealing gradient	Primers extension	Final extension
94 °C, 5 min	95°C, 1 min	(55°C Gradient Δ10°C)1 min	72°C, 1min	72 °C,10 min
	30 cycles			

Result and Discussion:

Isolating and Diagnosing of Staphylococcus aureus

In this study, Thirty six specimens were collected from burn and wound patients from Baghdad Hospitals. Swabs of wounds and burns samples were collected and absorbed by other report Rittenhouse et al.(2006) that wounds are common infections caused by S. aureus . Then the samples were

inoculated on culture media (Mannitol agar and blood agar) ,as shown in Figure1,they were initially diagnosed on blood agar some of the isolates produce a clear translucent area surrounding the colonies as result for β-haemolysin production.Hemolysis by this bacteria considered the most important features for diagnosis and identification the pathogenic isolates of S.aureus from other Staphylococcuspp(Atlas et al .,1995).

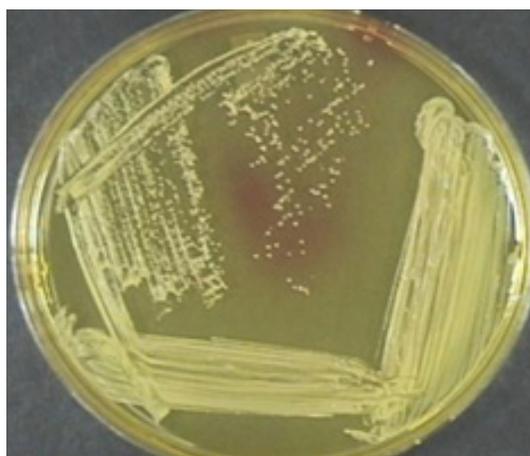


Figure 1: S.aureus on Mannitol Salt agar.

Microscopic examination was applied to the all isolates after staining by Gram stain and the cells appeared as Gram-positive cocci arranged in pairs or grape-like irregular clusters, as represent the phenotypic characteristics for this genus (Baron et al., 1994). Mannitol salt agar is selective differential growth medium. It encourages the growth of Staphylococcus isolates, while inhibiting the growth of others. It contains a high concentration 7.5-10.0 % of NaCl salt, making it selective for Gram-positive bacterium Staphylococcus and Micrococcus.

S.aureus produces yellow colonies, whereas other Staphylococci produces small pink or red colonies with no color change to the medium. If an organism can ferment mannitol, an acid is formed that will cause the phenol red in the agar to turn yellow (Anderson and Cindy, 2013).

Susceptibility of Staphylococcus aureus to antibiotics:

According to the recommendation of (CLSI, 2013), ten isolates of S.aureus isolated from burn and wound patients

were tested for antibiotic sensitivity by disc diffusion method (Kirby-Bauer test). The results showed relatively high resistance to Methicillin ME, Amoxicillin-Clavulonic acid AMC, Azithromycin AZM with frequencies of 100%, 100% and 70% respectively. Kader et al. (2011) reported that 88.24% of isolates were resistant to methicillin and Oxacillin discs, while Odonkora and Addob (2011) reported that methicillin disc diffusion test detected 54 MRSA (21.6%), while relative moderate resistance to Vancomycin VA, Amikacin AK with frequencies of 60% and 62% respectively has been observed in our study. Alebachew et al. (2012) found a resistance of S.aureus isolates towards Vancomycin and Amikacin 9.7% and 10.0%, respectively.

S.aureus showed relatively low resistance to Fusidic acid with frequencies of 30% has been detected with other study of Mason et al. (2003) who reported low value for Fusidic acid. as shown in Figure 2.

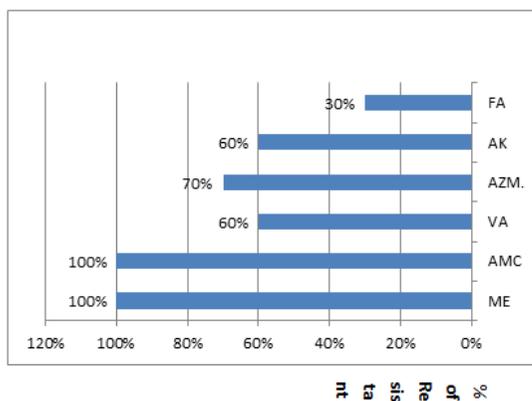


Figure 2: Percentage of Resistance antimicrobial agents for Staphylococcus aureus

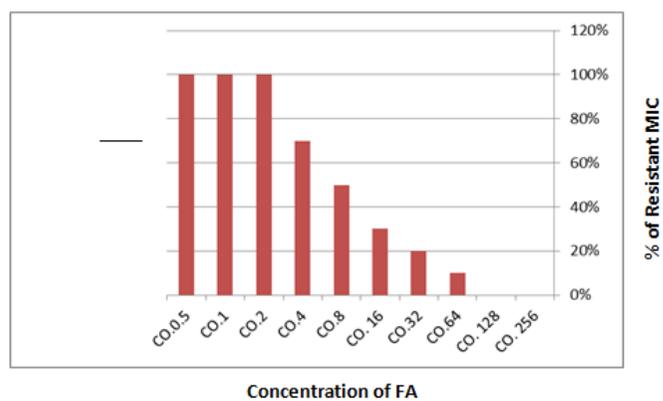


Figure 3: Percentage of Resistance Minimum inhibitory concentration agar diffusion method for Fusidic acid (co; concentration 0.5-256) for Staphylococcus aureus.

Determination of MIC for Fusidic acid antibiotic of S.aureus:

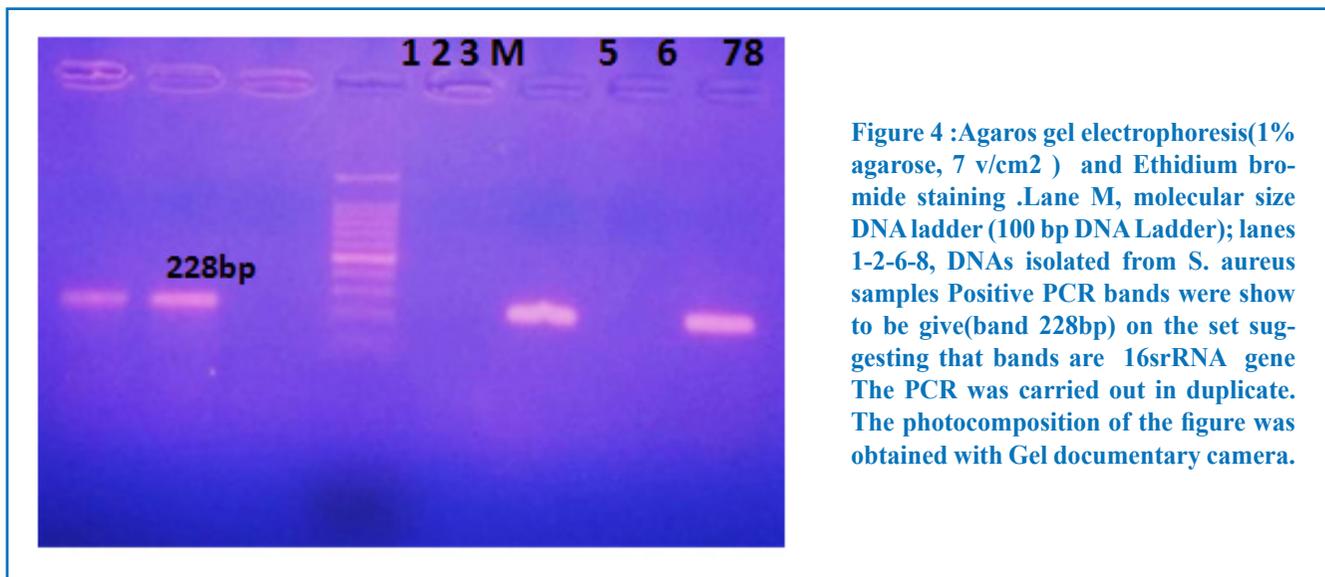
Determination of MIC for Fusidic acid resistant isolates of S. aureus shown intermediate resistant to Fusidic acid CLSI (2013). The results show MIC range from 8-256 µg/ml. The resistance was presented in Figure 3 which summarized the result of MIC experiment. Perera, G and Hay, R (2005) highlighted this concern overgrowing fucidin resistance. The West Yorkshire study found that 50% of fusidic acid-resistant

strains were from dermatology patients exposed to topical fucidin in the 6 months prior. The lowest rate of resistance to Fusidic acid was 10% in the concentration 64 µg/ml the MIC values. Mason (2003) indicated that the resistance level to fucidin is low at present, most likely due to its unique molecular structure and therefore is less likely to share resistance mechanisms with other antibiotics. Prolonged treatment with fucidin ointment should be avoided, even in the community setting.

4:Genetic Detection of S. aureus by 16s-rDNA

The total DNA (chromosomal) will be served as a template since they will be released outside the bacterial cell. The present research findings pertain to the isolation of *S. aureus*, samples were confirmed as *S. aureus* on the basis of morphological, biochemical and molecular characterization. After gradient

PCR has been performed (55°C Gradient Δ 10°C), bands have been obtained as presents in (Fig-4). Showing that the best annealing temperature to give 16s-rDNA was (51.5°C, 30s) depending on results of gradient PCR amplification procedure



Several studies have used PCR for the detection of 16S-rDNA by PCR method more rapidly and reliably. The results of current study agree with result of a same study in Iran reported by (Maliheet al;2011) who noticed that amplification of 16s-rDNA confirmed all the 126 staphylococcal isolates as *S. aureus*.

Conclusions

PCR is a rapid, sensitive, specific and inexpensive method; we suggest that it can be replaced to traditionally assays

for detecting *S. aureus*. Although the 16S rDNA procedure differentiated all tested clinical isolates identified 100%. Therefore, if an unknown organism needs to be identified, 16S rDNA sequencing is the method of choice because of the availability of universal primers. Furthermore, antibiotic susceptibility tests and MIC should be done besides detecting bacterial factors in order to enhance treatments for decreasing infections of human staphylococci and study the resistance for fusidic acid due to the most common topical treatment for wounds and burns

References:

1. Abeck, D.; Mempel, M. (1998). Staphylococcus aureus colonization in atopic dermatitis and its therapeutic implications. *Br J Dermatol*; 139: 13–16.
2. Abed El-Jalil, H.; Jallad, M.; Thwaini, A.J. (2008). Nasal carriage of methicillin resistant Staphylococcus aureus in individuals exposed and not exposed to hospital environments. *Eur. J. Sci. Res.*, 22: 570-574.
3. Adwan, G. B.; Abu-Shanab and Adwan, K. Enterotoxigenic-Staphylococcus aureus in raw milk in North of Palestine. *Turk. J. Biol.* 2005. Vol. 29. Pp: 229-232.
4. Alebachew, T.; Yismaw, G.; Derabe, A. and Sisay, Z. (2012). Staphylococcus aureus burn wound infection among patients attending Yekatit 12 Hospital Burn unit, Addis Ababa, Ethiopia. *J. Health. Sci.*, 22(3).
5. Anderson and Cindy (2013). *Great Adventures in the Microbiology laboratory* Pearson. pp:175-176.
6. Atlas, R.M.; Brown, A.E & Parks, L.C. (1995). *Laboratory manual of experimental microbiology*. 1st ed. Mosby, St. Louis U.S.A.
7. Baron, E.J.; Peterson, L.R. and Finegold, S.M. (1994). *Bailey and Scott's Diagnostic Microbiology*, 9th ed. Mosby, Library of Congress Cataloging in Publication. pp:215-332.
8. Carboneau, C.; Bengel, E.; Jaco, M.T.; Robinson, M. (2010). A lean six sigma team increases hand hygiene compliance and reduces hospital-acquired MRSA infections by 51%. *Journal for Healthcare Quality*; 32:61-70.
9. Clinical and Laboratory Standards institute. *Performance Standards for Antimicrobial susceptibility Testing: Twenty-Third Informational Supplement M100-S23*. CLSI, Wayne, PA, USA, 2013.
10. Ebrahim Rahimi and Forough Alian. (2013). Presence of enterotoxigenic Staphylococcus aureus in cow, camel, sheep, goat, and buffalo bulk tank milk. *Veterinarski Arhiv*. Vol. 83 (1). Pp: 23-30.
11. Emel, Banu B.B.; Mehmet, A.B.; Taner, I.; Erkan, Y. (2010). Application of PCR-RFLP of gap gene method as a molecular typing tool for coagulase negative Staphylococci from bovine and human origin identified with VITEK 2. *Afr J Microbiol Res* :775–82.

