Staphylococcus epidermidis Prevails *Staphylococcus aureus* in Multispecies Biofilm under Gentamicin Stress

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Abstract: *Out of 60 staphylococcal isolates isolated form different sources, the results revealed that 20 and 10 were identified as S. aureus and S. epidermidis, respectively. These results were confirmed by detecting 16sRNA gene. Remarkably, 80% of S. aureus and 80% of S. epidrmidis isolates developed Methicillinresistance. Findings of the current work demonstrated that most of methicillin resistant S. aureus (MRSA) and methicillin resistant S. epidermidis (MRSE) formed weak biofilm. The competition between S. aureus and S. epidermidis in multispecies biofilm were tested at several conditions encompassed temperature, pH, and starvation. Results showed that the numbers of biofilm cells have intensely decrease to undetectable limits in the presence of Gentamicin, approximately, at all tested conditions. With few exceptions, S. epidermidis showed a noticeable dominance in all species combinations.*

1. Introduction

Staphylococci defined as a diverse group of bacteria that cause different types of diseases which could be minor skin infections up to life-threatening bacteraemia. Staphylococci still the major cause of both hospital and community acquired infections worldwide even the high continued efforts to control their spread. *Staphylococcus aureus* and *Staphylococcus epidermidis* considered as the two major opportunistic pathogens of this genus [1].

Fifty years ago, Methicillin and different semisynthetic penicillins were commonly used which led to the appearance of Methicillin Resistance *Staphylococcus aureus* (MRSA) and *S. epidermidis* (MRSE), and that continue to persevere in both the healthcare and community environments. Currently, the results revealed that 80% (16/20) of *S. aureus*, 80 % (8/10) *S. epidrmidis*, and 86.6% (26/30) other CONS isolates developed Methicillinresistance.

Biofilm formation may be determined to detect the capability of staphylococci to colonize the biomedical devices. The microtiter plate (MtP) test devised by Christensen *et al.* [2].Were most commonly used as the phenotypically methods for slime and/or biofilmproduction.

The organization of the biofilm into complex structures is regulated by the exchange of chemical signals between cells in a process known as quorum sensing. Quorum sensing is a strategy of cell-cell communication benefiting the biofilm community by controlling unnecessary overpopulation and competition for nutrients with important implications for the infectious process [3].

This study was designed to achieve the following aims; 1) studying the competence between isolates (resistant vs sensitive to environmental conditions (pH, temperature, and starvation). 2) detecting the role of Gentamicin on competition at the same environmental conditions and 3) investigating the competence capacity between the two species in vivo using murine model.

2. Materials and Methods

Specimen'scollection
Sixty clinical specimens referring to Flow Catheter, Burn, Indwelling Devices, Midstream Urine, Wounds Swabs, and Sputum, were collected from patients attending several hospitals in Baghdad (AL Karama Hospital, Teaching Laboratories of Medical City) for the period from October 2015 to January 2016.

Isolation and identification ofstaphylococci
All specimens were streaked on mannitol salt agar and Blood Agar. Thereafter, all plates were incubated aerobically for 24 h at 37°C. The grown coloines were identified according to Bergey's Manual [4] depending on the morphological features on culture media and biochemical tests[5, 6] as follows:

Detection of MRSA by Cefoxitin Disk Diffusionmethod
All the isolates were subjected to cefoxitin disk diffusion test using a 30 µg disk. A 0.5 McFarland standard suspension of the isolate was made and lawn culture was done on Mueller-Hinton agar plate. Plates were incubated at 37ºC for 18 hr. and zone diameters were measured. An inhibition zone diameterx of ≤ 19 mm was reported as Methicillin-resistance and ≥20 mm was considered as methicillin-sensitive *S. aureues*, and an inhibition zone diameter of ≤ 24 mm was reported as Methicillin-resistance and ≥25 mm was considered as methicillin-sensitive *S. epidermidis* [7].

Extracting and purifying of genomicDNA
Genomic DNA from all isolates (n= 60) was extracted and purified by using Genomic DNA Extraction (Geneaid, USA), using manufacturer protocol, which can be summarized asfollows:

Multiplex PCR
The presence of 16sRNA for *S. aureus*. 16sRNA for *S. epidermidis* genes and *mecA*in both species were detected by amplifying them as described by Arciola *et al.* [8].

*mecA*and 16sRNA for *S. aureus* and 16sRNA for *S. epidermidis* primers (Table 1) were provided in lyophilized form (Bioneer, Korea), dissolved in DNase free distilled water to give a final concentration of 100 pmol/ µl as recommended by provider and stored in a freezer until use.

Table 1: DNA primers used in the present study [8]

Id	Primers	Target gene	Amplicon size (pb)
1	SA1 (5'-AATCTTTGTCGGTACACGATATTCTTC ACG-3') SA2 (5'-CGT AATGAGATTTCAGTAGAT AATACAACA-3')	16srRNA of *S. aureus*	108
2	SE1 (5'-ATCAAAAAGTTGGCGAACCTTTTC A-3') SE2 (5'-CAAAAAGAGCGTGGAGAAAAGT ATCA-3')	16srRNA of *S. epidermidis*	124
3	MRS1 (5'-TAG AAA TGACTGAACGTCCG-3') MRS2 (5'-TTG CGA TCAATGTTACCGTAG-3')	*mecA*	154

PCR mixture was set up in a total volume of 20 µl included 2 µl of each primer (10 micromole/ µl) and 6 µl of template DNA (50 ng/µl) have been used. The rest volume was completed with free nuclease water. Negative control contained all material except DNA, in which D.W. was added instead of template DNA. Program of DNA amplification was as follows: initial denaturation at 92°C for 3 min., 30 cycle at 92°C for 1 min., 56°C for 1 min., 72°C for 1 min. and a final extension at 72°C for 3 min.

Determination of Gentamicin minimal inhibitory concentration (MIC).
Depending on microdilution method described by CLSI[9], different concentrations of Gentamicin (2 – 1024 µg/ml) were dissolved in Mueller-Hinton broth. The microtiter plate was incubated overnight at 37°C. Thereafter, the lowest concentration that inhibits bacterial growth was considered as the MIC.

Biofilm formationassay
Quantification of biofilm formation by *S. aureus* and *S. epidermides* on abiotic surfaces was assessed as previously described by Nakao *et al.*[10]. In brief; The wells of sterile 96-well U shaped-bottomed polystyrene microplates were filled with 200 µl of an overnight nutrient broth (bacterial concentration was adjusted to be equivalent to McFarland standard no. 0.5) before the plates were covered and incubated aerobically at 37°C for 24 h. Each bacterial isolate was tested in triplicate. Control wells were performed by adding bacteria-free nutrient broth. To visualize biofilms, the contents of the wells were aspirated and the wells washed three times with distilledwater to remove loosely adherent cells. The remaining attached bacteria were fixed with 200 µl methanol for 30 min. After drying in air, the wells were stained with 200 µl 0.1% crystal violet solution for 20 min at room temperature. Excess stain was rinsed off by placing the plate under running tap water. Thereafter, the plates were dried in a 37°C incubator for approximately 30 min to ensure they were completely dry. Subsequently, the adherent cells were resolubilized with 200 µl absolute ethanol for 10 minute. Finally, the optical density of each well was obtained at 630 nm using microplate reader.

Classification summarized in Table 2 based on OD_{630} values obtained for individual isolate of *Staphylococcus* spp. were used for the purpose of data simplification andcalculation [11].

Table 2: Classification of bacterial adherence by tissue culture plate method [11]

Mean OD_{630}	Adherence Biofilm Formation
OD ≤ ODc	Non –adherent
ODc< OD≤2*ODc	Weakly adherent
2*ODc < OD ≤4*ODc	Moderately adherent
4* ODc <OD	Strongly adherent

ODc = average OD of negative control + (3 *SD of negative control)

After calculating the biofilm formation capacity for all tested isolates and negative controls, the cut- off value (ODc) was established. It is defined as a three standard deviations (SD) above the mean OD of the negative control: ODc value was calculated for each microtiter plate separately. When a negative value was obtained, it presented as zero, while any positive value was an indicator for biofilmproduction.

Competition experiments

Bacterial preparation
MRSA, MRSE, methicillin sensitive *S. aureus* (MSSA), and methicillin sensitive *S. epidermidis* (MSSE) cells were diluted to the desired concentration (1.5×10^8 CFU/ml) and validated by dilution plating in mannitol salt agar. Thereafter, equal volumes of each bacterial isolate in accordance to the following combinations were prepared: A) MRSA and MRSE. B) MRSA and MSSE. C) MSSA and MRSE. and D) MSSA and MSSE.

The aforementioned protocol for biofilm formation assay was adopted to form biofilm for each previously mentioned combination. However, Different pH values (6, 7, and 8), incubation temperature (10, 37, and 45°C) and starvation (culture medium was diluted 1:100 and 1:1000 with D. W.). Viable count was accomplished for each assay prior and post Gentamicin addition.

Statistical analysis
Means were compared to each other via ANOVA and T test. Differences were considered significant when P ≤ 0.05.

3. Results and Discussion

Bacterial isolation andidentification
In accordance to biochemical, morphological, and cultural as well as 16sRNA,20 coagulase producers and mannitol fermenters were identified as *S. aureus*; whereas the other 40

non-mannitol fermenters, non-hemolytic (gamma haemolysis), and coagulase negative were comprised as *S. epidermidis* (n= 10) and other*Staphylococcus* species (n=30).

Numerous studies have investigated the isolation of staphylococci from diverse clinical specimens. Onanuga and Temedie [12]stated that nearly 33.3%of *S. aureus* isolates were isolated from 120 nares swabs. Babakir-Mina *et al.* [13] reported that *S. aureus* was isolated from 22% of patients attending Sulaimania burn hospital, and formed about 36% from burn specimens. In another workcarried out byVaez *et al.* [14]recorded nearly 30.8%, 26.5%, 22.7%, 10.8 and 9.2% of *S. aureus* isolates were isolated from urine, wound, blood, sputum, and abscess, respectively.

Concerning *S. epidermidis*,Eftekhar and Mirmohamadi[15] reported that 40% of isolates were isolated form blood; while 14% of isolates were recovered from urine specimens and another 14% from wound specimens. Nevertheless, similar percentages (8%) were recorded from intravascular catherter and exudate, and other unknown sources (16%). On the contrary,a study conducted byDiemond-Hernandez *et al.* [16], showed that 79.5% of *S. aureus* and 73.77% of CONS were detected in different sources.

LikewiseGad *et al.* [17] found that out of 292 bacterial isolates collected from urine and catheterspecimens, *S. aureus* and *S. epidermidis* represented 6.2%, 12%, respectively.

Whilst,Adesida *et al.* [18] reported that 14.0% *S. aureus* nasal colonization in medical students in Lagos,Nigeria. Chopra *et al.* [19] found that *S. aureus* comprised 31.11%; while CONS covered 13.33%. In spite of that, Rabelo *et al.* [20] collected 404 samples of 202 health professionals, in which 14.6% (59/404) were *S. aureus* isolates and 6.2% (25/404) were classified as *S. epidermidis*;while Onanuga and Onalapo[21]andAl-Geobory[22] declared that about 36% of 150 urine sample was identified as *S. aureus*. Pant and Rai's [23] findings revealed higher *S. aureus* nasal colonization rate (43.8%) in staffs of teaching hospital in Nepal.

These variations may be attributed to the characteristics of the population under study. A population with medication with antibiotics as at the time of sampling may yield a much lower prevalence of *S. aureus* while a population deal with hospital settings may yield a much higher prevalence because of the high number of infectious patients in that environment.

Detection of methicillin resistance
Cefoxitin (30 μg/disk) disk diffusion method was adopted to investigate the methicillin resistance to all staphylococcal isolates (n = 60). Markedly, variable methicillin resistance was noticed among *Staphylococcus* sp. The results revealed that 80% (16/20) of *S. aureus*, 80 % (8/10) *S. epidrmidis*, and 86.6% (26/30) other CONS isolates developed Methicillinresistance.

Infections because of MRSA are globallyexpanding problematic issue either inside or outside clinical settings.

Therefore, it is very important for laboratories at epidemiological and clinical levels to distinguish MRSA from MSSA. The reason lying behind such distinguishing in not limited to picking suitable and efficient antibiotic treatment policy for the individual patient, but correspondingly to keepthe transmission of MRSA under control[24].

It has been reported in literature that the results of cefoxitin disk diffusion methodis highly compatible with PCR technique that detect the presence of *mecA* more than do the results of oxacillin disk diffusion method. Given that, the specificity and sensitivity for detecting methicillin resistance for both methods (Ca. cefoxitin disk and PCR) are 100% [7, 25, 26].

Resistance to methicillin in *Staphylococcus* sp. is primarily mediated by the presence of penicillin-binding protein 2a, encoded by the *mecA* gene. In certain MRSA strains, the *mecA* gene is heterogeneously expressed *in vitro*[27].

A lot of local studies dealt with MRSA isolation. Among them a study done byAl-Hasani[28]demonstrated that 41/49 (83.7%) were MRSA and 30/37 (81%) were identified as Methicillin resistance coagulase negative staphylococci (MRCONS). Another local study reported that MRSE covered84%[29]. Likewise,Babakir-Mina *et al.*[13]stated that among *S. aureus* positive cases, 88% were MRSA. Upon the results ofAl-Dahbi and Al-Mathkhury [30] MRSA constitutes about 94.3% among *S. aureus*isolates.Muhammad[31] revealed that 68% (49/72), 93% (26/28), and 92% (34/37) of *S. aureus*, *S. epidrmidis*, and other CONS isolates developed Methicillinresistance.

Studies in Iran also paid attention to MRSA isolation, it has been noticed that the incidence MRSAin burn patients was ranging between 60% - 80% [32, 33,34]. Similarly, in Isfahan, Iran, Khorvash [35] demonstrated that the prevalence of MRSA in nosocomial infections reached67.2%. Diverse epidemiological factors like geographical location, health system proficiency in administrating infection control programshave crucial role in such noticed variability of MRSA prevalence. MRSA isolates were mostly isolated form wound infections (43%). For instant, [36]) reported that MRSA covered around 33% in a burn center in USA[36].

The results of Al-Hasani [37] (using Methicillin disks (5μg/disk) revealed that from all 49 isolates, 41 isolates (83.7%) showed resistant to the Methicillin while 5 isolates (10.2%) were sensitive to the Methicillin and only 3 isolates (6.1%) were intermediate resistance. A study carried out byAl-Maliki[38] who showed that the rate of MRSA was 80.3% and the Methicillin sensitive *S. aureus* was 16.4% while the intermediate resistance covered 3.3%. Additionally the same author demonstrated that the rate of MRSE was 86.6% and the Methicillin sensitive *S. epidermidis* was 4.4% and *S. epidermidis* that intermediate resistance to the Methicillin was8.8%.

When this study compared with the present study two important findings could be observed; the first one is the emergence increased in the Methicillin resistance

staphylococci (both COPS and CONS) strains and decreased of sensitive strains in the very short period of time in our country (Iraq). Additionally, this finding is agreed with many other studies and reports in the world that suggested over the last few decades, there has been an enormous increase and emergence in the prevalence of Methicillin resistance both COPS and CONS strains all over the world and such strains are prone to cause serious outbreaks [39, 40,41]. For instance; CA-MRSA infections have been reported in North America, Europe, Australia, New Zealand, United States and South Korea [42, 43].

The second important finding concluded from the present results, is that the rate of coagulase-negative staphylococci isolates that resistant to Methicillin were higher than the rate of Methicillin resistance coagulase-positive and this finding is agreed with many other studies that mention that *S. epidermidis* is more often resistant to antimicrobial drugs than *S. aureus* and resistance to Methicillin occurs in about 35% of *S. aureus* while occurs in approximately 75% of *S. epidermidis* isolates [44, 45]. Some studies mention that 90% of coagulase-negative staphylococci, mostly *S. epidermidis*, being Methicillin resistant [46, 47].

The variations are due to differences of local conditions, such as climate or microbial prevalence, but others are likely to be caused by differing prevention protocols, topical and systemic treatment of burn wounds, sampling regimens as well as study lengths [48].

Methicillin resistance of *S. aureus* remains to be a significant problem. Rapid and accurate determination of Methicillin resistance is vital for the establishment of successful treatment policy. Misdiagnosing this resistance leads to treatment failures and spread of infections with these resistant strains. This means that disk diffusion testing by using cefoxitin disk is far superior to most of the currently recommended phenotypic methods and is now an accepted method for the determination of MRSA by many authors and related centers including CLSI [49].

Estimation of Minimal Inhibitory Concentration(MIC)
This estimation was performed for four isolates, two S. *aureus* (MRSA and MSSA) isolates and two *S. epidermidis* (MRSE and MSSE) isolates. Different concentrations of Gentamicin (2 – 1024 µg/ml) were used, the break point of Gentamicin is ≥ 4µg/ml depending on CLSI[7].

Findings of this work revealed that all isolates were Gentamicin resistance (≥ 64µg/ml), hence sub-MIC (32 µg/ml) was used for further experiments.

Biofilmassay
The ability of *S. aureus, S.epidermidis* and other *staphylococcus* spp.biofilm producing isolates were evaluated using pre-sterilized 96-well polystyrene microtiter plates. In order to estimate biofilm degree, absorbance was determined at 630 nm by an aid of microtiter plate reader. Given that, absorbance values represented the degree of the biofilm thickness that formed by the studied isolates on the surface of the microtiter well. Obtained results were categorized into four groups based on limits summarized in Table 3.

Table 3: Biofilm degree based on estimated cut off value* of this study

Id	Biofilm degree	OD_{630} limits
1	Non adherent	< 0.043
2	Weak adherent	0.043 – 0.086
3	Moderate adherent	0.086 – 0.17
4	Strong adherent	≥ 0.17

*cut off value= Mean of controlOD_{630} + 3*Standard deviation= 0.036 + 3*0.0025= 0.043

Regarding limits stated in Table 3, results of the present study revealed that out of 16 MRSA, 12 formed weak biofilm while 4 isolates formed mild biofilm. However, 2 MSSA produced weak biofilm and another two formed mild biofilm. Concerning *S. epidermidis*, two MRSE isolates formed weak biofilm; while six and two MSSE formed weak and mild biofilms, respectively. In regard to MRCONS, seven isolates formed weak biofilm and another seven also made mild biofilm. Additionally, one MSCONS formed mild biofilm. Noticeably, 11 MRCONS isolates and one MSCONS were unable to form biofilm. Nevertheless, no isolate has the ability to produce strong biofilm Table 4.

Table 4: Biofilm forming degree of *S. aureus*, *S. epidermidis* and other CONS*

Id	Isolate sp.	Methicillin susceptibility	Biofilm degree			
			No Biofilm	Weak Biofilm	Mild Biofilm	Strong Biofilm
1.	*S. aureus*	MRSA	0	12	4	0
		MSSA	0	2	2	0
2.	*S. epidermidis*	MRSE	0	2	0	0
		MSSE	0	6	2	0
3.	Other CONS	MRCONS	11	7	7	0
		MSCONS	1	0	4	0
	Total		12	29	19	ZERO

*CONS; coagulase negative staphylococci

A study performed by Muhammad [31] showed that 100% MRSA isolates were able to form weak biofilm; whereas, MRSE isolates couldn't develop strong biofilm. Nevertheless, 96% and 4% of MRSE isolates formed weak and moderate biofilm, respectively.

In the study of Al-Dahbi [50], the biofilm-forming ability of *S. aureus* was evaluated for all isolates (106 isolates) by microtiter plates and the result revealed that 31%, 45%, and 22% of isolates were weakmild, and strong producers, respectively. Noticeably, 2% of the tested isolates were highly-strong producers.

Gad *et al.* [17] reported that the biofilm producing in both *S. aureus* and *S. epidermidis* produce high, moderate and weak biofilm for *S. aureus* (66.7%, 16.7 %, and 16.7%, respectively) and *S. epidermidis* (51.4%, 37.1%, and 11.4%,respectively).

The findings of this study was incompatible with a study achieved byGad *et al.* [51], that they stated the biofilm production assessed by microtiter plates MtP revealed 30 (90.9%) strains of *S. aureus* were biofilm positive and 19 strains (95%) of CONS were biofilm former. Quantitative biofilm production demonstrated that 22 (76.7%) of strains

were strong biofilm producers, 8 (24.2%) strains were moderate biofilm producer, and 3 (9.1%) strains were non-biofilm producers. In CONS, 13 (65%) strains were strong biofilm producers, 6 (30%) moderate biofilm producers and one (5%) strain was non-biofilm producer.

Liduma *et al.*[52] suggested that the percentage of the biofilm producing strains was 47.6%. It is very close to the results that what was observed byArciola *et al*. [8], who revealed that 46% of isolates were biofilm producers. Furthermore, in a study of Oliveira and Cunha [53] among 100 isolates studied, 35 (35%) of them were classified as weakly adherent and 46 (46%) as strongly adherent,foratotalof81(81%)positiveisolatesand19(19%)ne gative isolates.
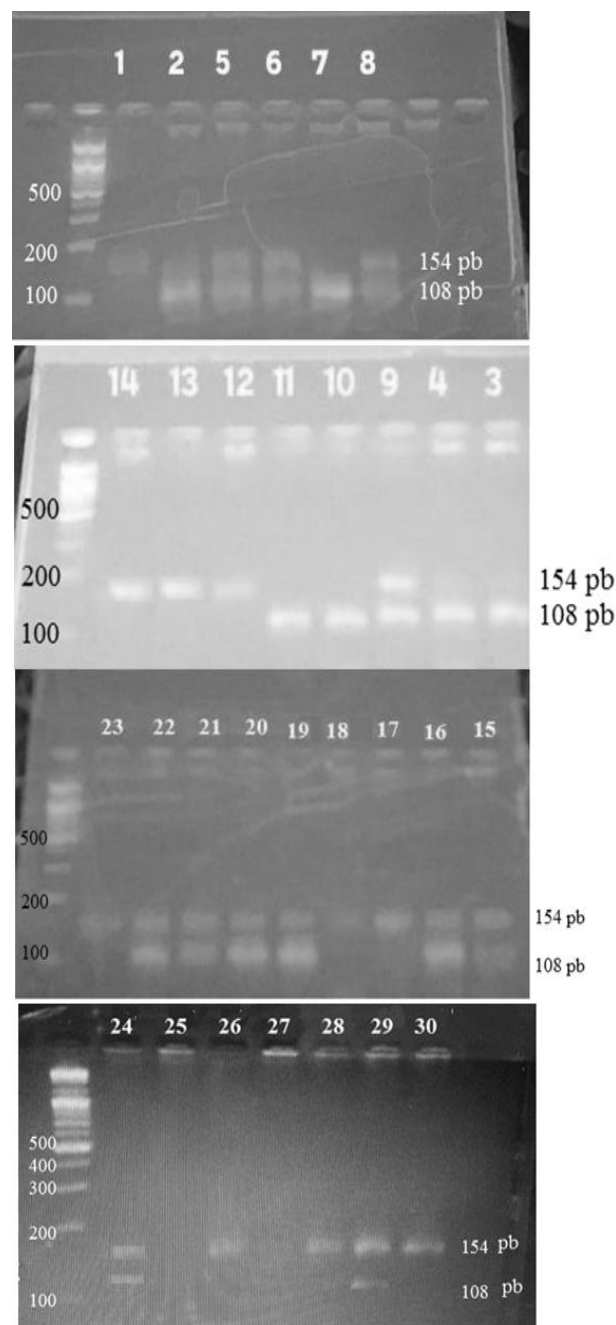
Detection of *mecA* gene and16sRNA for *S. aureus* and 16sRNA for *S.epidermidis*

Results depicted in Figure 1revealed that 20 isolates of *S. aureus* and 10 isolates of *S. epidermidis* were confirmed by 16srRNA amplification. Nearly 80% (16/20) of *S. aureus* isolates, 80 % (8/10)of *S. epidermidis*, and 86.6% (26/30) of other CONS isolates were *mecA* positive. Noticeably, the result of PCR completely matched those resulted from cefoxitin disk diffusion method.

Most of staphylococcal isolates (83.3%) of the present study appeared to be resistant to cefoxitin (alternative to methicillin). Such findings are in agreement withFey *et al.* [54] who stated that 81% of *S. aureus* were MRSA, whileAl-Geobory[22] found that 90.9% of *S. aureus* isolates were resistant to methicillin. Additionally,Jain *et al.* [55] observed about 75.3% of isolates were methicillin resistant. Yet, these observed differences might be due to the variation in the geographic area, sources of clinical specimens, genetic background and the collection site of isolates. The reason behind continuous increasing in resistant to β-lactam antibiotics is caused by the overuse or misuse of these antibiotics and by the use of poor quality antibiotics. It also results from natural genetic changes, or mutations, within the organisms that cause diseases.

Bockstael and van Aerschot [56] have been describe four major mechanisms of resistance; Target site alterations, such as changes to the penicillin-binding protein, are common, inactivation of antimicrobials, as by penicillinases or the new carbapenemases, is often seen, Prevention of antimicrobial access to their targets by changing the cell wall permeability that denies access to antimicrobials during the course of therapy and Finally, newly described efflux mechanisms pump the antimicrobial out of the cell before it can reach its target site.

Nevertheless, several studies have shown that resistance in bacteria also acquired by a mutation and acquisition. Acquisition may be passed vertically by selection to daughter cells or horizontally by transfer of resistance genes between strains and species. Exchange of genes is possible by transformation, transduction or conjugation [57].
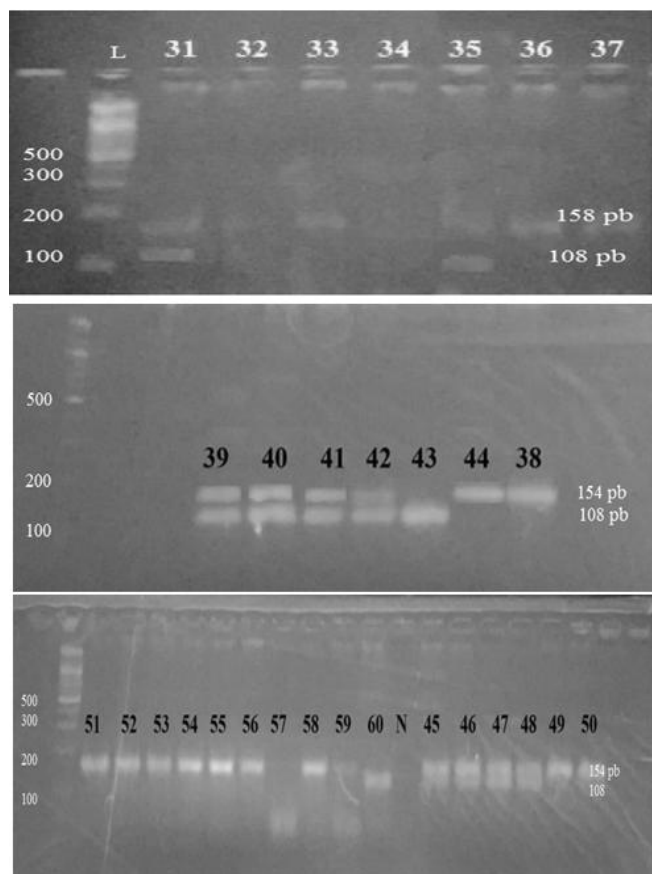
Figure 1: Agarose gel electrophoresis analysis of *mecA* gene (154bp) and 16sRNA for *S. aureus* (108bp) and 16sRNA for *S. epidermidis* (124bp). Ladder denotes to 100 pb DNA ladder. Detection was accomplished on agarose gel (2.5%) at 75 V for 1.5 hour, stained with ethidium bromide and visualized on a UV transilluminator documentation system.

Competition experiments

Given that the numbers of biofilm cells have dramatically decrease to undetectable limits nearly at all tested conditions (Tables 5 to 11), it can be concluded that Gentamicin has a significant ($P < 0.05$) impact on viability of cells inside the biofilm. In general, *S. epidermidis* showed a noticeable dominance ($P < 0.05$) in all species combinations at all tested conditions.

Table 5: Effect of Gentamicin* on Biofilm cell viability at pH 7 and 10°C

Species combinations	Without Gentamicin ($\times 10$ CFU/ml)		With Gentamicin ($\times 10$ CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	984	12406	Undetectable	Undetectable
MRSA and MSSE	30	181	Undetectable	Undetectable
MSSA and MRSE	63	123	Undetectable	Undetectable
MSSA and MSSE	42	1294	Undetectable	Undetectable

***Gentamicin was added 32µg/ml (subMIC)**

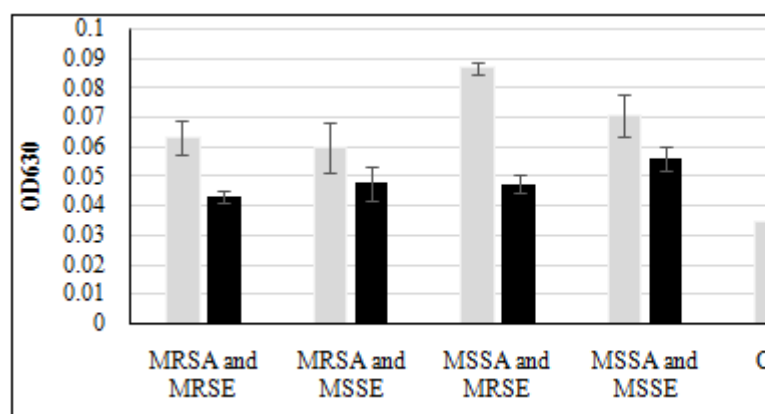Markedly, biofilm degree was highlyaffected ($P \leq 0.05$) by Gentamicin (Figure 2).

Figure 2: Effect of Gentamicin (32µg/ml) on Biofilm at pH 7 and 10°C

Nevertheless, at 37°C, MRSA and MSSA outnumbered MRSE and MSSE, respectively (Tables 6). However, Gentamicin has decreased the biofilm thickness ($P \leq 0.05$) in exception for MRSA and MSSE coexistence, in which biofilm thickness increased insignificantly ($P \geq 0.05$) as it is depicted in Figure 3.

Table 6: Effect of Gentamicin* on Biofilm cell viability at pH 7 and 37°C

Species combinations	Without Gentamicin ($\times 10$ CFU/ml)		With Gentamicin ($\times 10$ CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	2015	885	Undetectable	Undetectable
MRSA and MSSE	565	1018	Undetectable	Undetectable
MSSA and MRSE	25724	2868	Undetectable	Undetectable
MSSA and MSSE	571	805	Undetectable	Undetectable

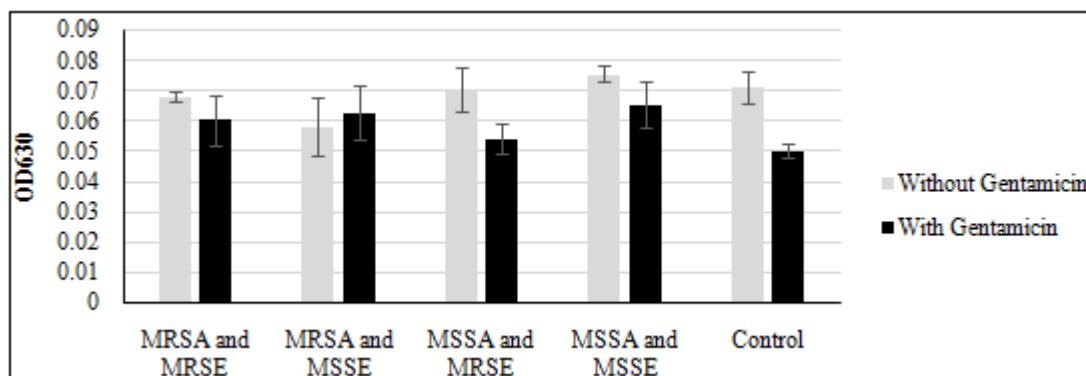***Gentamicin was added 32µg/ml (subMIC)**

Figure 3: Effect of Gentamicin (32μg/ml)on Biofilm at pH 7 and 37°C

At 45°C, a marked dominance of *S. epidermidis* over *S. aureus* was noticed (Table 7). Moreover, biofilm thickness was affected ($P \leq 0.05$) by the presence of Gentamicin as it depicted in Figure 4.

Table 7: Effect of Gentamicin* on Biofilm cell viability at pH 7 and 45°C

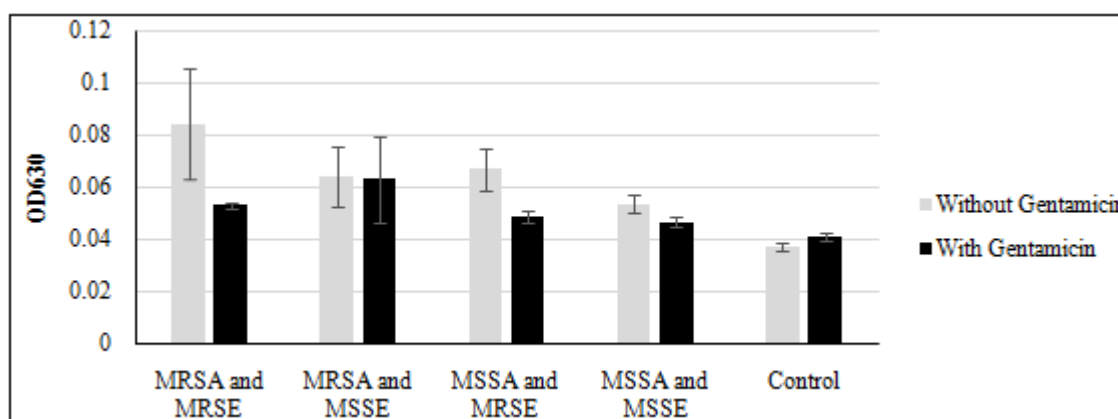| Species combinations	Without Gentamicin (× 10 CFU/ml)		With Gentamicin (× 10 CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	20	31	Undetectable	Undetectable
MRSA and MSSE	24	34	Undetectable	Undetectable
MSSA and MRSE	31	39	Undetectable	Undetectable
MSSA and MSSE	19	28	Undetectable	Undetectable

*Gentamicin was added 32μg/ml (subMIC)

Figure 4: Effect of Gentamicin (32μg/ml) on Biofilm at pH 7 and 45°C

In Gentamicin free medium, MRSA and MSSA overnumbered ($P < 0.05$) MSSE when they incubated at pH. Whereas, when Gentamicin was added, MRSA overnumbered MRSE in addition to numbers of cells is highly reduced post Gentamicin addition (Table 8). Evidently, biofilms formed by all combinations were highly influenced ($P < 0.05$) by the presence of Genamicin as it is illustrated in Figure 5.

Table 8: Effect of Gentamicin* on Biofilm cell viability at pH 6 and 37°C

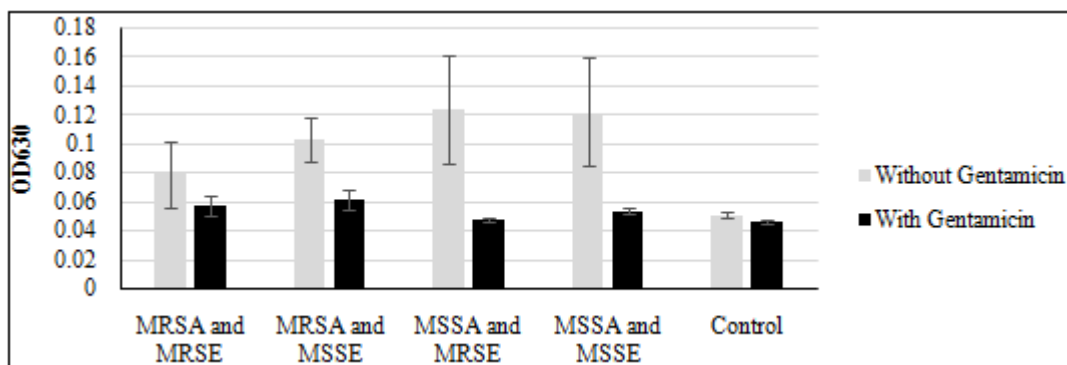| Species combinations	Without Gentamicin (× 10 CFU/ml)		With Gentamicin (× 10 CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	682	800	136	59
MRSA and MSSE	112	84	Undetectable	Undetectable
MSSA and MRSE	469	812	Undetectable	Undetectable
MSSA and MSSE	1056	397	Undetectable	Undetectable

*Gentamicin was added 32μg/ml (subMIC)

Figure 5: Effect of Gentamicin (32μg/ml)on Biofilm at pH 6 and 37°C

Once again, at pH 8, *S. epidermidis* had a superiority (P ≤ 0.05) in biofilm over *S. aureus* with and without Gentamicin addition (Table 9). However, Gentamicin highly affected (P ≤ 0.05) biofilm thickness as shown in Figure 6 and cell numbers (Table 9).

Table 9: Effect of Gentamicin* on Biofilm cell viability at pH 8 and 37°C

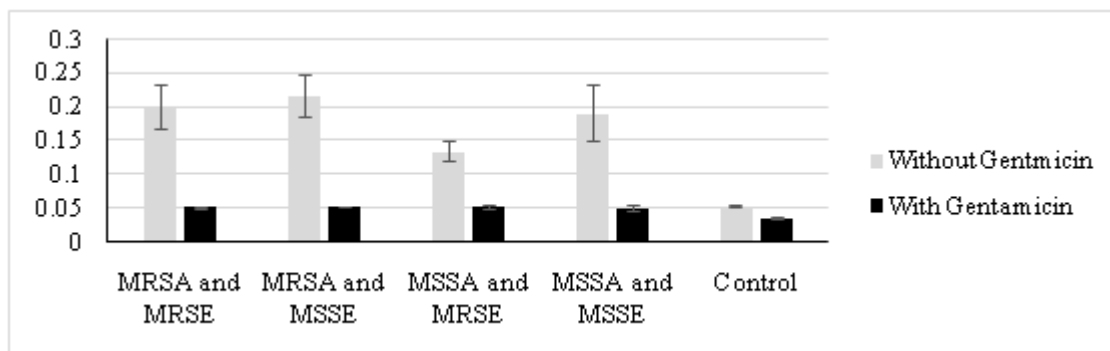| Species combinations	Without Gentamicin (× 10 CFU/ml)		With Gentamicin (× 10 CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	620	694	82	92
MRSA and MSSE	363	964	25	39
MSSA and MRSE	607	1055	77	154
MSSA and MSSE	80	152	80	169

*Gentamicin was added 32μg/ml (subMIC)

Figure 6: Effect of Gentamicin (32μg/ml)on Biofilm at pH 8 and 37°C

When biofilm was starved (1:100), MRSA was in greater number (P < 0.05) than MRSE in Gentamicin free medium; however, in the presence of Gentamicin, *S. epidermidis* numbers were more than those of *S. aureus* (Table 10). Gentamicin reduced the biofilm thickness in all species combination (Figure 7).

Table 10: Effect of Gentamicin* on starved** Biofilm cell viability at pH 7 and 37°C

Species combinations	Without Gentamicin (× 10 CFU/ml)		With Gentamicin (× 10 CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	602	53	197	987
MRSA and MSSE	16	32	12	27
MSSA and MRSE	488	675	85	189
MSSA and MSSE	66	150	570	589

*Gentamicin was added 32μg/ml (subMIC);**Brain Heart Infusion at 1:100 dilution.
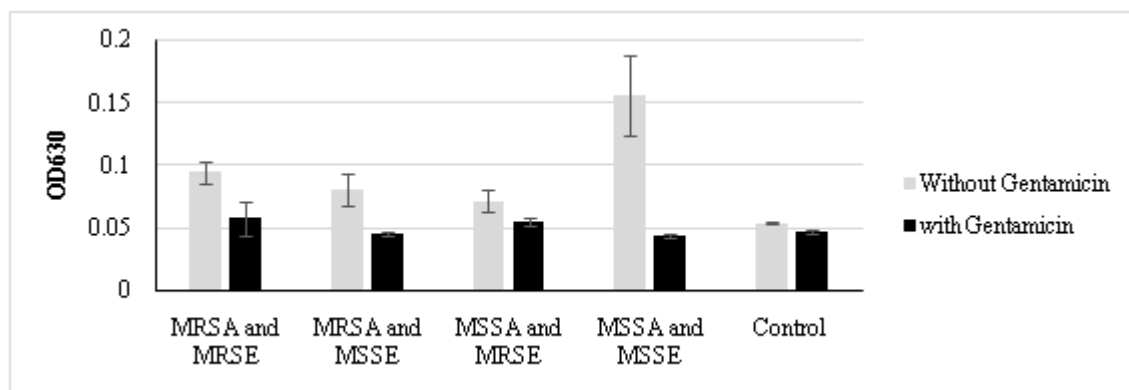
Figure 7: Effect of Gentamicin (32µg/ml)on starved (1:100 dilution) Biofilm at pH 7 and 37°C

But when biofilm was starved at 1:1000,*S. epidermidis* exceeded *S. aureus*in all species combinations except for MSSA; which hadtopped ($P < 0.05$) MRSE in the presence of Gentamicin (Table 11). In regard to biofilm thickness, Gentamicin reduced thickness of all biofilms in exception for MRSA and MRSE (Figure 8)

Table 11: Effect of Gentamicin* on starved** Biofilm cell viability at pH 7 and 37°C

Species combinations	Without Gentamicin (× 10 CFU/ml)		With Gentamicin (× 10 CFU/ml)	
	S. aureus	*S. epidermidis*	*S. aureus*	*S. epidermidis*
MRSA and MRSE	45	142	674	867
MRSA and MSSE	81	183	163	1336
MSSA and MRSE	117	208	135	90
MSSA and MSSE	173	1062	453	847

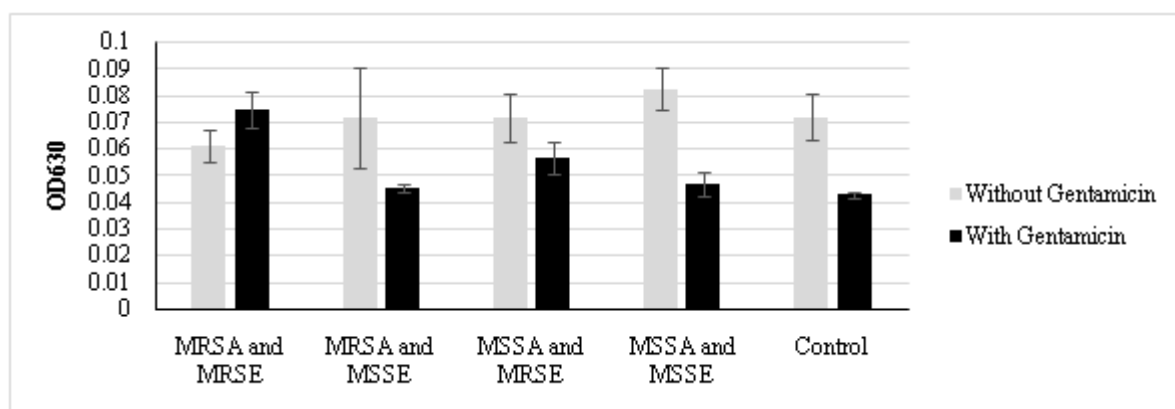***Gentamicin was added 32µg/ml (subMIC);**Brain Heart Infusion at 1:1000 dilution.**

Figure 8: Effect of Gentamicin (32µg/ml)on starved(1:1000 dilution)Biofilm cell viability at pH 7 and 37°C

The dominance of *S. epidermidis* over *S.aureus* reflects the ability of *S. epidermidis* to adapt and overcome the adverse conditions on the expense of *S. aureus*. Perhaps, it possesses colonizing factors or other determinants enable them to compete successfully other microorganisms which might found in vicinity.

Sato *et al.*, [58]stated that MRSA displayed a lowered surface electronegativity alongside with a shortage in teichoic acids amount. Even though the physicochemical features of such surfaces could hypothetically affect adhesion.

Iwase *et al.* [59] reported that the serine protease Esp secreted by *S.epidermidis* have inhibited the biofilm formation of *S.aureus* as well as nasal colonization. Moreover, purified Esp has the ability to destroy the pre-existing *S. aureus* biofilms.Sugimoto *et al.* [60] demonstrated that Esp degraded certain proteins found in the

biofilm matrix and cell wall of *S. aureus*. Additionally Esp developed an ability to degrade colonization factors, such as fibronectin-binding protein A and protein A. Furthermore, Esp selectively degraded many host receptors of *S. aureus* (e.g., vitronectin, fibrinogen, and fibronectin).

Burmolle *et al.*[61]demonstrated that synergistic increases in biofilm mass in mixed- versus single-species biofilms. However, Macleod and Stickler [62] specifically investigated a mixed culture of *Proteus mirabilis* and*P. aeruginosa* in catheter biofilms, and they found minimal antagonism between the two species.

In a study achieved byVarposhti *et al.* [63], *Pseudomonas aeruginosa, Acinetobacter baumannii*, and *Stenotrophomonas maltophilia* were investigated togetherand inter-species communication among them, comprising some secreted factors (e.g. proteins, quorum-sensing molecules, secondarymetabolites, and

carbohydrates), which affect geneexpression, physicalcontact, the production of antimicrobials, metabolic cooperation and competition, maylead to improved biofilm formation. The precise mechanism ofsynergism among these species is unknown. However, one or more of the interactions stated above may haveoccurred.

Lehman and Donlan [64] stated that the numbers of biofilm-associated *P. aeruginosa* and *P. mirabilis* were not affected by the presence of the other species at the 2-h, 24-h, and 48-h time points. The elimination of *P. aeruginosa* by 72 h in the two-species catheter reactors, regardless of phage treatment, was likely driven by the high pH that developed between 48 and 72 h due to *P. mirabilis* urease activity, as evidenced by *P. aeruginosa* growth inhibition in medium with pH 10. However, the high pH did not appear to inhibit lytic activities for either the *P. mirabilis* or *P. aeruginosa* phages in our catheter model, suggesting that phage application in indwelling urinary catheters with high pH conditions might be feasible.

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