

# Detection and Control of Biofilm Formation by *Staphylococcus aureus* from Febrile Neutropenic Patient

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**Introduction:** Febrile neutropenia (FN) is the evolution of fever in a patient with neutropenia over 38.0°C. Neutropenia is diagnosed when absolute neutrophil count (ANC) <1500 cells/ $\mu$ L. FN represents a common complication of cancer treatment. Hence, it is featured to be a major cause of morbidity and mortality in cancer patients. *Staphylococcus aureus* is one of the most important microorganisms isolated from the blood of febrile neutropenic patients. Infections caused by *S. aureus* range from mild to life-threatening diseases. Biofilm production by *S. aureus* is one of the most significant virulence factors of the bacterium as it prevents the penetration of antibiotics. Recently, it has been shown that *S. aureus* carries the *ica* operon responsible for biofilm production. The aim of the work is to determine a genotypic characterization that includes not only the detection of *icaA* and *icaD* genes in *S. aureus* but also the determination of their relation to clinical and microbiological features. Empiric antibacterial treatment was recommended for cancer patients receiving chemotherapy.

**Materials and Methods:** The relation between the presence of *icaA* and *icaD* and biofilm production was determined in a collection of 66 *S. aureus* samples from febrile neutropenic patients. Biofilm-forming ability was tested on Congo Red agar plates. Also, the effect of the most sensitive antibiotics on the bacterial cells was determined by an electron microscope.

**Results:** Of the bacterial samples, 48 were biofilm-productive and 18 were non-biofilm productive. For the biofilm productive bacteria, 37.5% were positive for *icaA*, 22.9% were positive for *icaD* and 10.4% were positive for both. Linezolid was the most effective antibiotic and it is highly recommended for the treatment of febrile neutropenia caused by biofilm-productive *S. aureus*. Severe changes were found on the bacterial cell after being treated with Linezolid. The *icaA* and *icaD* genes were present in only 50% of biofilm-productive bacteria.

**Conclusion:** The *ica* operon is present in only 50% of biofilm-productive *S. aureus* and Linezolid is the best antibiotic against these bacteria.

**Keywords:** febrile neutropenia, *Staphylococcus aureus*, *icaA*, *icaD*, biofilm production

## Introduction

FN is defined as the rising of body temperature with other symptoms of infection in a neutropenic patient. It is also defined as an oral fever measurement higher than 38.3°C in a neutropenic patient without any environmental factors, where the fever persists for at least an hour.<sup>1</sup> Neutropenia is characterized by abnormally low numbers of neutrophil granulocytes in patient's blood; less than 1500 cells/mm<sup>3</sup> and less than 500 cells/mm<sup>3</sup> in severe cases.<sup>1</sup> The most common cause can be

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detected with a chemotherapy receiving complication especially in case of being myelosuppressive.<sup>2</sup>

The Multinational Association for Supportive Care in Cancer (MASCC) has divided the neutropenic patients into low-risk and high-risk according to the risk index. This is very important to determine the most suitable therapeutic strategies for treatment.<sup>3</sup> Treatment of febrile neutropenic patients starts with empirical antibiotics until the neutrophil count becomes normal (more than 500 cells/mm<sup>3</sup>). The body temperature also becomes normal. Antifungal agent must be added in cases of recurrence or continuous fever.<sup>4</sup>

Staphylococci are the most common organism associated with febrile neutropenia. *S. aureus* is considered to be the most dangerous of the numerous types of staphylococci. *S. aureus* is transmitted directly from infected persons by using contaminated objects or by droplet infections when sneezing or coughing.<sup>5</sup>

Infections caused by *S. aureus* range from mild to life-threatening diseases. A skin infection is the most common staphylococcal infection and it often causes abscesses. Bacteremia can also occur if the bacteria enter the bloodstream, and this can affect almost all the body parts.<sup>6</sup>

Staphylococcal infections occur by two mechanisms including direct invasion to the tissue, e.g. endocarditis and septic arthritis, and through the production of exotoxins. Some of these exotoxins have local effects and others have severe systemic effects as they induce release of cytokine from T cells, causing disease to all parts of the host body.<sup>7</sup>

The laboratory diagnostics of infections caused by *S. aureus* depend on the detection of the pathogen while species diagnostics depend on the delimitation of coagulase-negative *Staphylococcus* spp. either by the phenotypic method or by the genotypic method.<sup>8</sup>

*S. aureus* has the ability to form a biofilm of settled microbes. Those microbes are characterized by their ability to attach to a substrate or surfaces and to any available moist solid surface, including host tissue, forming a matrix using extracellular polymer substances. These microbes exhibit a special phenotype depending on the growth phase and gene expression and play a serious role in providing resistance to both host immune response and antibiotics.<sup>9</sup>

On one level, the most common genes known to play an important role in biofilm production are *icaA* and *icaD*. Extracellular matrix excreted by biofilm-producing bacteria consists mainly of polysaccharides, especially intercellular adhesion polysaccharide, i.e. poly- $\beta$  (1-6)-N-acetylglucosamine which is coded by the *ica* operon.<sup>10</sup>

On the other level, the biofilm matrix is built passively or actively. Passive matrix rearrangement occurs due to the sorption of substances that strengthen it or as a result of damaging factors,<sup>11</sup> while active modification is carried out by biofilm microbes. Micro-dependent matrix reconstruction is realized due to anti-directional processes include increasing production of matrix biopolymers, synthesizing matrix enzymes, enhancing cell reproduction, preventing autolysis, and rejecting of biofilm constituents into the environment. Moreover, these ways are the evolution of the skeleton of the biofilm takes place.<sup>12</sup> The matrix not only mechanically links the biofilm into a single structure, but also fills the intercellular spaces, forming a three-dimensional filtering system.<sup>13</sup>

Identification of the biofilm matrix depends on a number of methods based on its color. These methods are traditionally used and include the utilization of definite stains or sets of fluorescent dyes marketed by manufacturers as biofilm matrix detection agents or by the detection of specific antigens.<sup>14</sup>

Controlling the formation of biofilm occurs by direct or indirect pathways. Direct pathways include using substances that directly react with biofilm matrix molecules. These substances may be pharmaceutical preparations like Dispersin B or polysaccharides oxidative substances like sodium periodate.<sup>15</sup> Indirect pathways involve controlling the reactions of microbial metabolism through suppression of the synthesis of biofilm matrix polymers and the production of enzymes that destabilize it.<sup>16</sup>

## Materials and Methods

The study was carried out at the Microbiology Diagnostics and Infection Control Unit at the Medical Microbiology and Immunology Department, Oncology Center, Mansoura University. The study includes 414 blood samples collected from inpatients with fever (>38°C) receiving chemotherapy attending Mansoura University Hospitals. The patients were 165 males and 249 females. A written approval was taken before sample withdrawal. The study was performed in accordance with the ethical standards laid down in Mansoura Oncology Center.

## Collection of Blood Samples and Selection of Target Organism

Blood samples were collected from these patients – 1 mL in medical disposable blood collection EDTA tube under aseptic conditions – and a Complete Blood Picture (CBC)

test was done to determine the ANC for each sample so that we would be able to select febrile neutropenic patients. The CBC test was performed automatically using Sysmex KX 21 which is best known for its high accuracy in determining white blood cells count with differential. A blood culture test was performed using two systems. The first system was BacT/Alert 3D 60 as an automated microbial detection system based on the colorimetric detection of CO<sub>2</sub> produced by growing microorganisms. The second one was Vitek 2 compact that has the capacity to improve therapeutic success and patient outcomes through reliable microbial identification.

## Confirmatory Tests to Ensure the Isolated Bacteria

Isolated bacteria were subjected to several tests including microscopic examination after Gram-stain, catalase test, coagulase test, Api staph test (A strip utilizes a battery of miniaturized biochemical tests), and mannitol salt agar. Each of these tests has a standard result for *S. aureus* so that we were able to confirm our samples.

## Congo Red Test

Congo Red test was performed to detect biofilm production by *S. aureus*. Although the Congo Red test is not the most sensitive test to determine biofilm production, it can be reliably used because it is acceptably sensitive and specific.<sup>17</sup> Tested bacterial isolates were inoculated then incubated at 35°C for 24 h. Black and brown colonies with a dry crystalline consistency indicated biofilm production while red and dark red colonies are non-biofilm producers.

## Antibiotic Sensitivity Testing

Bacterial isolates were cultivated on Blood Agar as an enriched medium to encourage their growth. The standard disc diffusion microbial sensitivity test based on the Kirby–Bauer method on Mueller–Hinton agar was done for all *S. aureus* isolates to assess the antibiotic resistance as per Clinical and Laboratory Standard Institute guidelines.<sup>18</sup> Cells of 3 to 5 colonies were inoculated into 5 mL nutrient broth and incubated 2 to 8 hours until turbidity of the suspension matched 0.5 McFarland turbidity standards. This resulted in a suspension containing approximately 1–2 × 10<sup>8</sup> CFU/mL. Mueller–Hinton agar plates were inoculated by the inoculum using a sterile cotton swab. The antibiotic-impregnated discs were placed on the agar plates within 15 minutes of inoculation using

sterilized forceps. The plates were then inverted and incubated for 16–18 hours at 37°C. The diameters of inhibition zones were measured in millimeters using a ruler on the undersurface of the plate. Interpretation was done by referring to the zone diameter interpretive standards for *S. aureus* and the isolates were reported as sensitive, intermediate, or resistant. The antibiotics used were Tetracycline 30 mg, Ciprofloxacin 5 mg, Vancomycin 30 mg, Erythromycin 15 mg, Ceftriaxone 30 mg, Levofloxacin 5 mg, Ofloxacin 5 mg, Ampicillin 10 mg, Cefoxitin 30 mg, and Linezolid 30 mg.

## Specimen Preparation for Scanning Electron Microscope (SEM) and Transmission Electron Microscope (TEM)

Two samples were prepared to be examined with an electron microscope (EM). Each sample was examined two times using TEM and SEM. The first sample was normal bacterial isolates and the second one was treated with linezolid 30 mg. Both samples were examined to determine the recent changes of the bacterial cell with the effect of the antibiotic and to ensure the presence of biofilm so as to determine its thickness. Ultrathin sections were examined at 80 kV using a JEOL 2100 EM in the Faculty of Agriculture, Mansoura University, Egypt.

## Specimen Preparation for Electron Microscope

All samples were fixed in 4F1G fixative for electron microscopy studies, samples processed and examined as follows:<sup>19</sup>

### Fixation

- Fix 1 mm tissue blocks in 4% formaldehyde and 1% glutaraldehyde in 0.1 M PB (pH 7.4) for at least 2 hours to overnight.
- Immerse in 8% (0.2 M) sucrose in 0.1 M PB 3×15 min or overnight.
- Post-fix in 1% osmium tetroxide in 0.1 M PB for 1 hour.

### Dehydration

The dehydration was in 50% ethanol for 15 min, 70% ethanol for 15 min, 95% ethanol for 15 min, 100% ethanol for 2×15 min, 100% acetone for 1 hour, 1:1 Embed 812: propylene oxide for 1–2 hour, and 2:1 Embed 812: propylene oxide overnight in a desiccator with the top off.

## Embedding

Samples were embedded in beam capsules then were baked in a 60°C oven for 48 h.

## Sectioning

Firstly, thick sections (0.5–1.0 µm) were cut on a slide, and sections were dried on a slide warmer, and stained with Toluidine blue for 2–5 min. Secondly, sections were observed under a microscope for the precise location to cut ultrathin sections at 60–90 nm thick, and sections were collected onto grids. Finally, sections were examined under a transmission electron microscope (JEOL JEM-2100).

## Coating

This step was used for SEM only; samples after dehydration were coated with gold by a Sputter Coating Evaporator instrument. Examination was done by scanning electron microscope (JEOL JSM 6510 Iv).

## Molecular Characterization

Molecular characterization was performed to determine if *icaA* and *icaD* genes are present in *S. aureus* isolates that was defined to be able to produce biofilm around them. The relation between the presence of these genes and biofilm production was determined. The process consists of three steps: bacterial DNA extraction from a colony on blood agar with a genomic BYF DNA extraction mini kit according to manufacturer instructions. The second step was centered on polymerase chain reaction (PCR) for detection of *icaA* 5-TCTCTTGCAGGAGCAATCAA which was used as the forward primer (corresponding to nucleotides 4796–4815) and 5-TCAGGCACTAAC ATCCAGCA which was also used as the reverse primer (corresponding to nucleotides 4964–4983) and detection of *icaD* ATGGTCAAGCCCAGACAGAG was used as the forward primer (corresponding to nucleotides 5422–5441), and 5-CGTGTTTTCAACATTTAATGCAA was used as the reverse primer (corresponding to nucleotides 5616–5597). PCR was performed using the method described previously<sup>20</sup> using a thermal cycler. PCR cycling conditions used were 30 cycles of 1 min of denaturation at 94°C, annealing for 1 min at 58°C followed by 1 min extension at 72°C with final extension for 5 min at 72°C. PCR was performed with Taq polymerase (Helini). The third step was centered on analyzing the reaction mixtures by 1% agarose gel electrophoresis to resolve PCR products. The amplified product sizes were estimated

by comparison with the 50 bp DNA ladder. Amplicons for *icaA* and *icaD* produced fragments of 189 and 195 bp, respectively.

## Results

### Bacterial Isolation

This study was carried out on 414 febrile patients. CBC tests allowed us to select neutropenic patients. The distribution of these patients was 180 non-neutropenic and 234 neutropenic. The distribution of the neutropenic patients according to blood culture was 28.2% *S. aureus*, 14.5% Gram-positive other than *S. aureus*, and 57.3% were Gram-negative isolates as shown in Table 1.

### Confirmatory Test for Isolated Bacteria

Isolated bacteria were purple in color and cocci in shape by microscopic examination, catalase and coagulase positive (showed colt appearance in the test tube), the Api-Staph test strip after inoculation with bacterial isolates and occurrence of chemical reactions matched standard results for *S. aureus*, and mannitol salt agar changed to yellow as an indication of fermentation occurrence.

### Isolation of Biofilm-Productive *S. aureus*

Congo Red test results were black and brown colony color for 72.7% of the isolates, indicating biofilm-productive bacteria (positive result) and red or dark red colonies were 24.4% of the isolates, representing non-biofilm-productive bacteria (negative result), as shown in Table 2.

### Antibiotic Sensitivity Test

Table 3 summarizes the number and percentage of interpretation for each antibiotic. The table shows that Linezolid and Vancomycin are the most suitable antibiotics recommended to be used against *S. aureus* in febrile neutropenia.

### Electron Microscope Examination

Untreated *S. aureus* examination displayed condensed growth with rod-shaped undamaged cells. Also, it showed

**Table 1** Distribution of 234 Samples Collected from Febrile Neutropenic Patients According to Causative Organism

	Febrile Neutropenic Patients	Number	%
Causative organism	<i>S. aureus</i>	66	28.2
	Other Gram-positive	34	14.5
	Gram-negative	134	57.3

**Table 2** Type of Isolates According to Congo Red Test

Colony Color	Result	Conclusion	Number	%
Red and dark red	Negative	Non-slime productive bacteria	18	24.4
Black and brown	Positive	Slime productive bacteria	48	72.7

a smooth and intact surface without significant variation in length that seems to be several microns in thickness as shown in Figure 1. Moreover, *S. aureus* showed normal cell morphology and undamaged ultra-structure intact inner membrane, slightly waved outer membrane and each bacterial cell seems to be divided into two parts surrounded with cell membrane as shown in Figure 2.

After anti-microbial treatment with antibiotic, treated *S. aureus* was prepared for examination in the border line between growth and clear zone that showed enlarged malformed and uncharacterized cell shape. Some bacterial cells had burst with deep craters in their cell wall. Also, many lysed cells were observed. The cells condensed in the shape of a cluster and no growth was observed after the border line. Furthermore, non-uniform electron density cytoplasm and malformed enlarged cells with slightly ruptured as shown in Figures 3 and 4.

### Molecular Detection for *icaA* and *icaD* in Biofilm-Productive *S. aureus*

Table 4 shows that 37.5% of the isolates carry the *icaA* gene, 22.9% of the isolates carry *icaD* genes, and 10.4% of the isolates carry both genes at the same time. The percentage of the samples carrying at least one gene was 50%. Table 5 shows the mean age of febrile neutropenic patients who carry at least one gene, and those who do not carry any genes, and determines the significance of age in these patients. The mean age for patients who carry genes was  $51.58 \pm 17.90$  and the *P*-value was 0.019, so it is significant.

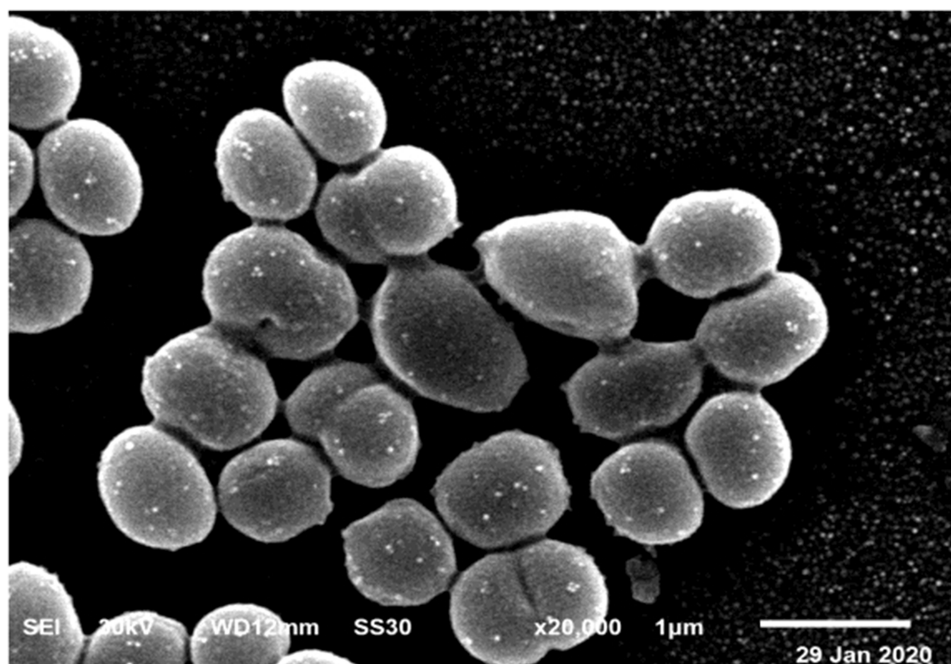
### Discussion

FN is the development of fever in a patient with neutropenia. Fever is an oral temperature measure of  $\geq 38.3^\circ\text{C}$  or a temperature of  $\geq 38.0^\circ\text{C}$  for a period of an hour. Neutropenia is characterized by abnormally low numbers of neutrophil granulocytes in the patient's blood.<sup>21</sup>

**Table 3** Antibiotic Susceptibility of the Isolated Bacteria Against Ten Antibiotics (Interpretation by Inhibition Zone)

Antibiotic	Inhibition Zone (mm)	No. of Isolates	%	Interpretation
Tetracycline (TE) (30 mg)	<11	18	37.5	Resistant
	12–14	15	31.25	Intermediate sensitive
	>15	15	31.25	Sensitive
Ciprofloxacin (CIP) (5 mg)	<15	42	87.5	Resistant
	16–20	3	6.25	Intermediate sensitive
	>21	3	6.25	Sensitive
Vancomycin (VA) (30 mg)	<9	3	6.25	Resistant
	10–11	24	50	Intermediate sensitive
	>12	21	43.75	Sensitive
Erythromycin (E) (15 mg)	<13	39	81.25	Resistant
	14–22	–	0	Intermediate sensitive
	>23	9	18.75	Sensitive
Ceftriaxone (CRO) (30 mg)	<13	24	50	Resistant
	14–20	15	31.25	Intermediate sensitive
	>21	9	18.75	Sensitive
Levofloxacin (LEV) (5 mg)	<13	9	18.75	Resistant
	14–16	30	62.5	Intermediate sensitive
	>17	9	18.75	Sensitive
Ofloxacin (OFX) (5 mg)	<12	39	81.25	Resistant
	13–16	3	6.25	Intermediate sensitive
	>17	6	12.5	Sensitive
Ampicillin (AM) (10 mg)	<13	36	75	Resistant
	14–16	9	18.75	Intermediate sensitive
	>17	3	6.25	Sensitive
Cefoxitin (FOX) (30 mg)	<21	42	87.5	Resistant
	21–22	3	6.25	Intermediate sensitive
	>22	3	6.25	Sensitive
Linezolid (LZD) (30 mg)	<20	–	0	Resistant
	21–22	12	25	Intermediate sensitive
	>23	36	75	Sensitive

FN is caused by different microorganisms, including Gram-positive and Gram-negative bacteria. From blood



**Figure 1** Examination of normal bacterial cells under scanning electron microscope.

cultures of FN cancer patients, staphylococci were some of the most common isolated organisms. *S. aureus* is considered to be the most dangerous of the numerous types of staphylococci. It causes skin infections, bone infections, pneumonia, and heart valve infections.<sup>22</sup>

*S. aureus* is the most common causative organism of FN in our results, representing 28.2%. This result confirms the results of Boyce<sup>23</sup> and Hakim et al<sup>24</sup> that *S. aureus* is the most common isolated organism from blood cultures of FN cancer patients.

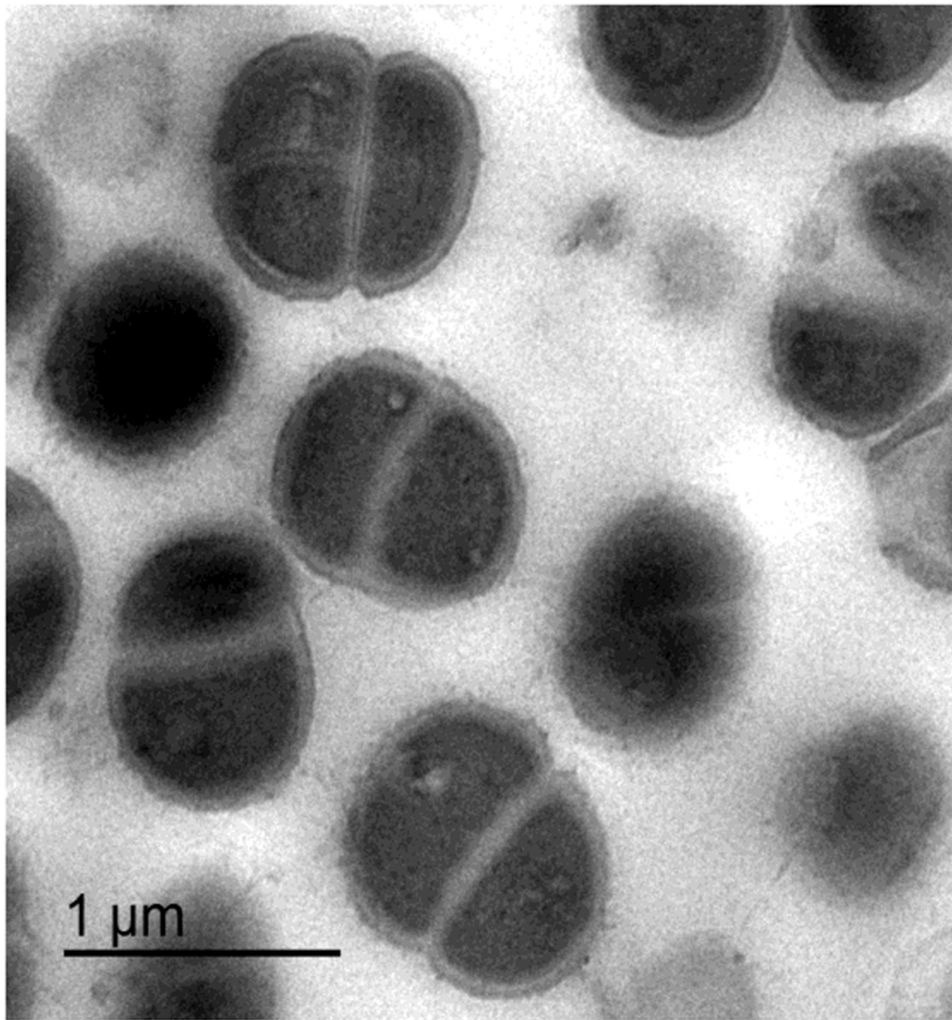
The distribution of microorganisms isolated from samples collected from febrile neutropenic patients in our research was 57.3% Gram-negative bacteria and 42.7% were Gram-positive. This result matches the previous results that confirmed the predominance of Gram-negative bacteria as the causative organisms of febrile neutropenia reported by Bodey et al<sup>25</sup> and Glauser et al,<sup>26</sup> and is in contrast to the result of Durnaa et al<sup>27</sup> that Gram-positive bacteria are the most common causative organism.

Congo Red test result showed that 72.7% of *S. aureus* were positive (biofilm producer) and 24.4% were negative (non-biofilm producer). This result is very similar to the result of Darwish et al<sup>28</sup> that 70.4% of staphylococcal isolates were positive to Congo Red test (biofilm producers) and completely in contrast to the result of El-Hendi et al,<sup>29</sup> which includes 23.6% biofilm productive staphylococcal isolates only.

These results explained the fact that biofilm formed by *S. aureus* is connected into a single system using the extracellular matrix. This helps the bacteria to mechanically link the biofilm into a single structure and fill the intercellular spaces, forming a three-dimensional filtering system. This is considered as virulence factors of *S. aureus* that enable bacteria to resist the effect of antibiotics.<sup>30</sup>

Antibiotic culture and sensitivity tests were performed to determine the most effective antibiotic recommended for the treatment of FN caused by *S. aureus*. The results showed that Linezolid and Vancomycin were the most effective antibiotics against isolated bacteria, while Ciprofloxacin and Cefoxitin were the antibiotics that have least effect on isolated bacteria.

As a comparison between the recent results and previous results, antimicrobial susceptibility patterns for *S. aureus* isolates exhibited 92.8% susceptibility to ciprofloxacin, Levofloxacin and Ofloxacin, 85.7% to tetracycline and a high resistance pattern of 71.4% was recorded in ceftriaxone and 42.8% in Vancomycin<sup>31</sup> Meanwhile, in our research, percentages of ciprofloxacin, Levofloxacin, Ofloxacin, and tetracycline sensitivity were 12%, 81.25%, 18.75%, and 62.5%, respectively, and the percentages of Ceftriaxone and Vancomycin resistance were 50% and 6.25%, respectively. So, our results match these results for Levofloxacin and approach results for Tetracycline and Ceftriaxone.



**Figure 2** Examination of normal bacterial cells under transmission electron microscope.

EM has long been used in the discovery and description of microorganisms like bacteria. Use of EM has been known to exist since the late 19th century. One of the main advantages of using EM for diagnosis is that it does not require organism-specific reagents for recognizing the pathogenic agent.

Shalamanou<sup>32</sup> reported that there was formation and/or degradation of the cell wall as well as pleomorphism, giant cell formation, and clustering of the damaged bacterial cell wall of three Gram-negative bacteria, i.e. *Enterobacter cloacae*, *Pseudomonas aeruginosa*, and *Serratia marcescens* after treatment with a minimal inhibitory concentration of chlorhexidine gluconate and examination by SEM.

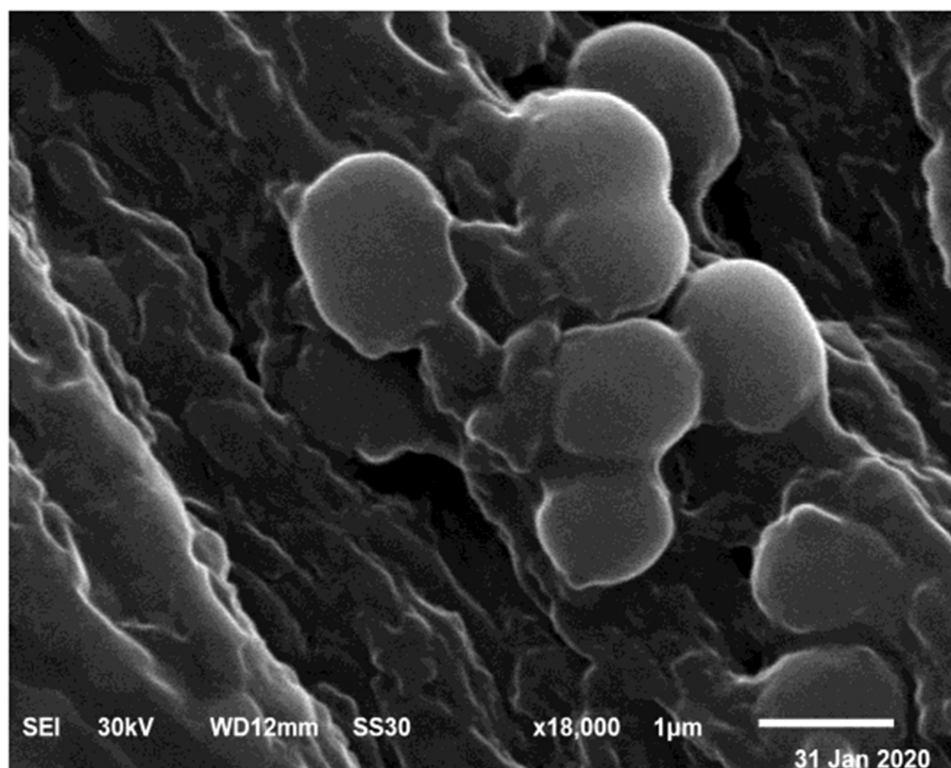
Bhaskar et al<sup>33</sup> treated *Klebsiella pneumoniae* with Cefotaxime (30 μg). He found wrinkled with rough-surfaced bacterial cells when examining with SEM. Previous researchers noticed loose attachment causing loss of permeability of bacterial cell membrane in addition

to crinkling and malformation of bacterial cell wall after being examined with TEM.

In this research, non-treated control bacterial isolates were examined to show normal cell morphology with undamaged ultra-structure intact inner membrane and slightly wavy outer membrane. Each bacterial cell seems to be divided into two parts surrounded by cell membrane.

The effect of Linezolid (30 mg) was tested by SEM and TEM in the border line between growth and clear zone. Differences between the two samples were reported. Observations showed enlarged malformation in the cell shape so that the shape became uncharacterized.

On one level, some bacterial cells had burst with deep craters in their cell wall. Also, many lysed cells were observed. The cells were condensed in the shape of a cluster.



**Figure 3** Examination of treated bacterial cells under scanning electron microscope.

On the other level, some other bacteria have an apparent hole in the cell surface, others are abounded, varying markedly in size and shape. Many of them were smaller than usual while others were abnormally large. No growth was observed after the border line. These changes in the bacterial cell are similar to those reported by Hartmann et al<sup>34</sup> and similar to the result of usage silver nanoparticles reported by Mirzajani et al.<sup>35</sup>

A PCR detection test was performed to detect the prevalence of *icaA* and *icaD* genes for *S. aureus* isolates isolated from febrile neutropenic patients. The results were 37.5% of these patients carry *icaA* gene, 22.9% carry *icaD* gene, and 10.4% of the isolates carried both genes. These results indicate that biofilm production by *S. aureus* was associated with the presence of *icaA* or *icaD* genes with a percentage of 50%. This percentage is less than the percentage found by El-Mahallawy et al,<sup>36</sup> who reported that 83.3% of biofilm productive *S. aureus* carry *ica* genes.

On the other hand, the results were proportionally equal to the result of Arciola et al<sup>21</sup> which determined a relation between the presence of the gene and slime production with a percentage of 60%.

Febrile neutropenic patients carrying *ica* genes showed a significant indication to the age of those patients. The death rate in older patients receiving chemotherapy was

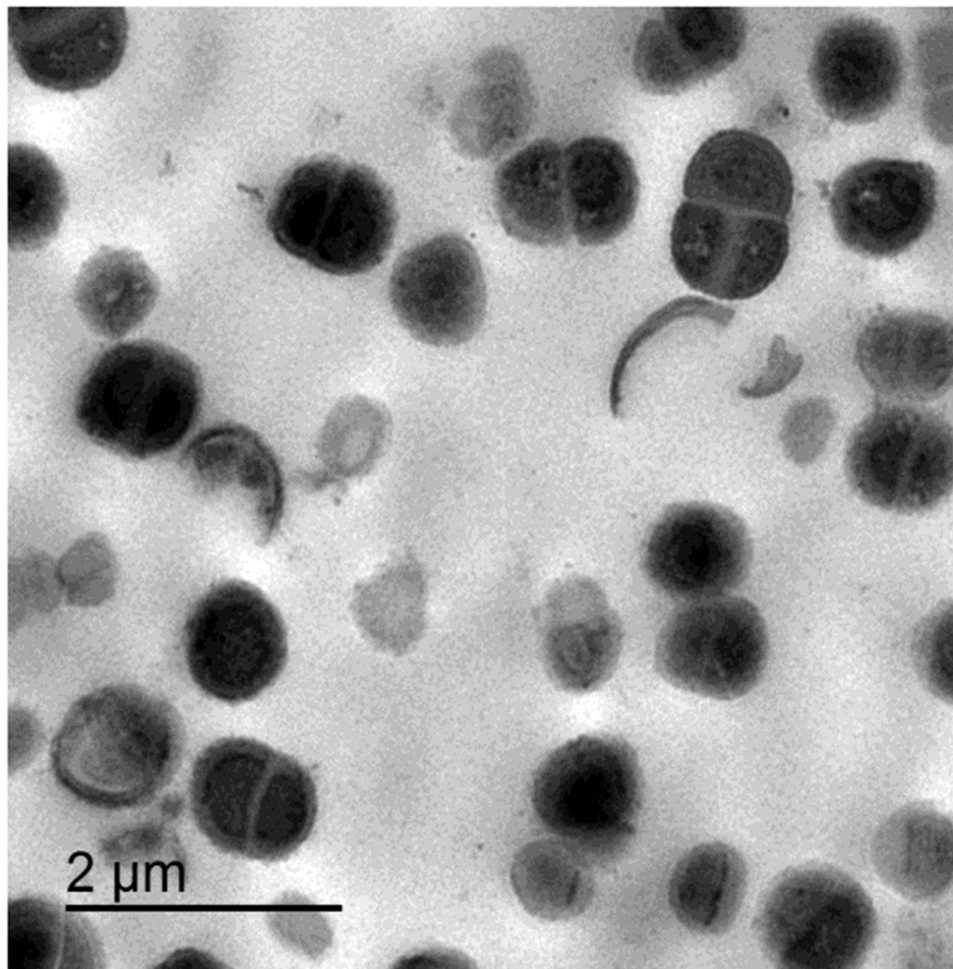
very high due to immune deficiency, cytokine storm, inappropriate diet, mutations, etc. This explains that the numbers in the last two groups were lower than expected. These results match the result of Aapro et al<sup>37</sup> in the sense that older age is considered an important patient-related risk factor that should be considered in the overall assessment of FN risk before administering chemotherapy.

## Conclusion

*S. aureus* is a Gram-positive bacterium, facultative anaerobic mostly aerobic, catalase, coagulase and hemolysis positive, mannitol, sucrose. It is the most common organism associated with febrile neutropenia. *S. aureus* is considered to be the most dangerous of the numerous types of staphylococci. Infections caused by *S. aureus* range from mild to life-threatening diseases. One of the most dangerous infections caused by *S. aureus* is bacteremia. Bacteremia occurs when the bacteria enter the bloodstream of the patient and affect almost all the body parts.

Biofilm production is one of the most important virulence factors of *S. aureus* as it enables them to resist or inhibit the effect of different antibiotics. The *icaA* and *icaD* genes are considered to be responsible for the





**Figure 4** Examination of treated bacterial cells under transmission electron microscope.

production of biofilm. In our research, we found that they were present in 50% of biofilm-productive *S. aureus*.

The most suitable antibiotic recommended to be used against *S. aureus* in febrile neutropenic patients is Linezolid. It has a destructive effect on the biofilm. In addition to its effect, it has the ability to make enlarged malformations in the cell shape and cytometry as it was clear in the electron microscopic examination.

**Table 4** Distribution of *icaA* and *icaD* Genes in Staphylococcal Isolates

Genes	No of Isolates	%
<i>IcaA</i>	18	37.5
<i>IcaD</i>	11	22.9
<i>IcaA</i> and <i>icaD</i> at the same time	5	10.4
Samples carrying at least one <i>ica</i> genes	24	50
Total biofilm-productive <i>S. aureus</i>	48	100

**Table 5** Significance of Ages in Cases Carrying at Least One Gene

Age	Both Genes				$\chi^2$	P
	Negative (n = 24)		Positive (n = 24)			
	No	%	No	%		
1–15y	3	12.5%	0	0%	7.822	0.166
16–30y	5	20.8%	4	16.7%		
31–45y	9	37.5%	6	25%		
46–60y	4	16.7%	5	20.8%		
61–75y	3	12.5%	6	25%		
>75y	0	0%	3	12.5%		
Mean $\pm$ SD	38.92 $\pm$ 18.34		51.58 $\pm$ 17.90			

**Note:** \*Significant  $P < 0.05$ .

**Abbreviations:** FN, febrile neutropenia; ANC, absolute neutrophil count; MASCC, Multinational Association for Supportive Care in Cancer; CBC, complete blood picture; SEM, scanning electron microscope; TEM, transmission electron microscope; PCR, polymerase chain reaction.

## Ethics Approval and Consent to Participate

This study was approved by the official ethics committee of the Oncology center – Mansoura University Institutional Review Board (IRB), reference number; R/17.1.39. Written approval was signed by a parent or a legal guardian of pediatric patients less than 18 years old. The study procedures were done in accordance with the Declaration of Helsinki. Patients above 18 years old have provided written approval.

## Acknowledgments

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## Disclosure

The authors report no conflicts of interest for this work.

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