The Droplet-Size Effect Of Squalene@cetylpyridinium Chloride Nanoemulsions On Antimicrobial Potency Against Planktonic And Biofilm MRSA

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Background: It is important to explore the interaction between antibacterial nanoparticles and microbes for understanding bactericidal activity and developing novel applications. It is possible that the nanoparticulate size can govern the antibacterial potency.

Purpose: The purpose of this study was to evaluate the antimicrobial and antibiofilm properties of cetylpyridinium chloride (CPC)–decorated nanoemulsions against methicillin-resistant Staphylococcus aureus (MRSA).

Methods: The droplet size could be adjusted by varying the percentage of squalene, the main ingredient of the oily core.

Results: We fabricated cationic nanoemulsions of three different sizes, 55, 165, and 245 nm. The nanoemulsions showed greater storage stability than the self-assembled CPC micelles. The tested nanoemulsions exhibited more antimicrobial activity against Gram-positive bacteria than Gram-negative bacteria and fungi. The killing of MRSA was mainly induced by direct cell-membrane damage. This rupture led to the leakage of cytoplasmic DNA and proteins. The nanoemulsions might also degrade the DNA helix and disturb protein synthesis. The proteomic analysis indicated the significant downregulation of DNA-directed RNA polymerase (RNAP) subunits β and β′. The antibacterial effect of nanoemulsions increased with decreasing droplet size in the biofilm MRSA but not planktonic MRSA. The small-sized nanoemulsions had potent antibiofilm activity that showed a colony-forming unit (CFU) reduction of 10-fold compared with the control. The loss of total DNA concentration also negatively correlated with the nanoemulsion size.

Conclusion: The present report established a foundation for the development of squalene@CPC nanosystems against drug-resistant S. aureus.

Keywords: cetylpyridinium chloride, nanoemulsion, MRSA, biofilm, size effect

Introduction
Pathogenic infection is the second-highest cause of death worldwide, with 16 million victims each year.1 The growing resistance to antibiotics exacerbates the health problem of microbial infection. The emergence of methicillin-resistant Staphylococcus aureus (MRSA) is a huge threat to health with everything from superficial skin wounds to toxic shock syndrome.2 The development of novel approaches to solve this problem is urgent. Nanotechnology demonstrates a promising strategy for combating the antibacterial resistance. The large surface area to mass ratio, high reactivity, and surface functionalization of the nanostructures have led to the unique properties for efficient eradication of pathogens.3 Nanoemulsions
are isotropic and thermodynamically stable nanosystems consisting of oil, water, and emulsifiers. They provide significant potential as functional additives in cosmetics, foods, and pharmaceutical products. Antimicrobial nanoemulsions show innate antibacterial activity against a broad species of pathogens.4,5 The surfactants in nanoemulsions are responsible for the strong bacteria-killing ability. These features make antimicrobial nanoemulsions feasible for application in wound treatment and surface decontamination.

The quaternary ammonium surfactants are often incorporated into nanoemulsions to form cationic nanosystems for antibacterial use. The cationic and amphiphilic characteristics of quaternary ammonium compounds can destabilize the bacterial membrane to exhibit the antimicrobial activity. Among these, cetlypyridinium chloride (CPC) as a cationic surfactant is approved by the USFDA as GRAS for extensive application in the food industry and hospitals as a disinfectant.6 It is found in many commercial products, including mouth rinses, wound dressings, nasal sprays, and cosmetics.7,8 CPC can form micelles in water via self-assembly in the amount of a higher than critical micellar concentration (CMC). However, the micelles usually reveal an unstable condition, resulting in the difficulty of quality control and application. Nanoemulsions can be ideal candidates as the vehicles of CPC. The size and composition of the nanoparticles can greatly influence the biological effects. The impact of the particulate size on the antimicrobial activity of metallic nanoparticles such as Ag, Pd, and ZnO has been widely studied.9–11 Nevertheless, no literature has been found pertaining to investigating the size effect of lipid nanosystems on the antibacterial effect. We intended to explore how the droplet size of cationic nanoemulsions affects the bacterial killing. Squalene was employed as the oil phase to produce CPC-coated nanoemulsions in the presence of blended emulsifiers (squalene@CPC nanoemulsions).

We fabricated three sizes of nanoemulsions: 55, 165, and 245 nm. MRSA was chosen as the main model microorganism in this study due to its prevalence and virulence in clinical infection. MRSA can exist in both planktonic and biofilm phenotypes, with the latter being predominant in the medical environment.12 A biofilm-forming capacity contributes to the unfavorable eradication of MRSA infection. Both phenotypes were used in this work to examine the anti-MRSA effect of the nanoemulsions. The proteomics of MRSA were established to understand the possible antibacterial mechanisms of squalene@CPC nanoemulsions. The size change of nanoemulsions under an accelerated stability test was evaluated to compare with self-assembled CPC micelles. It is possible to adjust the formulation and size of nanoemulsions in order to achieve optimized nanosystems based on the antibacterial profiles in the present study.

**Materials And Methods**

**Preparation Of squalene@CPC Nanoemulsions**

The oil and aqueous phases of nanoemulsions were fabricated separately. The oil phase consisted of soy phosphatidylcholine (SPC, 150 mg) and squalene. The squalene amount was 200, 500, and 700 mg for preparation of cationic nanoemulsions of small (CN-S), medium (CN-M), and large (CN-L) size, respectively. The aqueous phase consisted of CPC (80 mg), Poloxamer 188 (150 mg), and water for all formulations. Both phases were heated in a water bath for 20 min. The aqueous phase was added into the oil phase in the presence of high-shear homogenization at 12,000 rpm for 20 min, followed by the agitation using a probe-type sonicator (VCX600, Sonics and Materials) for 20 min at 35 W. The total volume of the final product was 10 mL.

**Size And Surface Charge Of Nanoemulsions**

The mean diameter and zeta potential of the nanoemulsion droplets were recorded by a laser-scattering technique (Zetasizer ZS90, Malvern). The recordation was done on the nanodispersion concentration after a 100-fold dilution with water. The storage stability was carried out by size measurement under the condition at 50°C and 75% relative humidity. The duration for the accelerated stability test was 14 d.

**Molecular Environment**

The molecular environment of the nanoformulations was assessed by fluorescence spectrophotometry based on the solvatochromism of Nile red.13 The CPC formulations were loaded with Nile red (1 ppm). The emission spectra of dye-incorporated systems were scanned from 550 to 700 nm. The excitation wavelength was 546 nm.

**Bacterial Strains And Culture Condition**

MRSA (ATCC 33591), *S. aureus* (ATCC 6538), *Escherichia coli* (ATCC 8739), and *Candida albicans* (ATCC 10231) were purchased from American Type Culture Collection. The drug-resistant clinical isolates of MRSA (KM-1) and vancomycin-intermediate *S. aureus* (VISA, KV-2) were gifts provided by Dr. Po-Liang Lu (Kaohsiung Medical University
Hospital). The strains were grown in tryptic soy broth (TSB) medium at 37°C and 150 rpm.

**Minimum Inhibitory Concentration (MIC) And Minimum Bactericidal Concentration (MBC)**

A broth twofold serial dilution method was utilized to measure MIC. Overnight culture of bacteria was diluted in TSB to achieve OD$_{600}$ of 0.01 (about 2x10$^6$ CFU/mL). The bacterial population was exposed to several dilutions of CPC ranging from 0.03 to 62.5 μg/mL with TSB and incubated at 37°C for 20 h. The ELISA reader at 595 nm was used to detect the absorbance. The reading value of <0.09 was the level of MIC. For the MBC assay, the bacterial suspension was diluted in PBS and plated on TSB plates. The TSB plates were incubated at 37°C for 20 h. Subsequently the CFU was counted. MBC was defined as the lowest concentration that killed ≥99.9% of the bacteria.

**Agar Diffusion Assay**

This study was performed according to the previous investigation. This assay was performed by inoculating the bacteria (OD$_{600}$=0.7) in the 0.75% TSB agar. The bacteria-agar mixture (5 mL) was poured into the dish for 15 min; then CPC (8 mg/mL) in micelles or nanoemulsions with a volume of 10 μL were pipetted onto the agar. The diameter of the inhibition zone was estimated after a 16-h incubation.

**MRSA Morphology Visualized By Scanning Electron Microscopy (SEM)**

MRSA at OD$_{600}$=0.1 was treated with CPC (50 μg/mL) in micelles or nanoemulsions at 37°C for 24 h. The microbes were fixed with 3% glutaldehyde and 2% paraformaldehyde in cacodylate buffer several times. After dehydration in an ascending series of ethanol, the samples were coated with gold and observed under Hitachi SU8220 SEM.

**Live/Dead MRSA Visualized By Fluorescence Microscopy**

The viability and death of MRSA after treatment of CPC (10 and 100 μg/mL) for 4 h were observed using a Live/Dead BacLight® kit (Molecular Probes). The pellet of bacteria was obtained by centrifugation at 12,000 rpm before CPC treatment. The treated bacteria were stained with the kit for 15 min. The samples were monitored by Leica DMI8 fluorescence microscopy. The detailed procedure of this experiment is described by Manteca et al.

**MRSA Biofilm Determination**

The biofilm was grown in a Cellview® dish by incubating MRSA (OD$_{600}$=0.1) in TSB containing 1% glucose at 37°C for 24 h. The biofilm was treated by CPC (50 μg/mL) for 24 h. The culture medium was transferred to the microtubes. The biofilm was rinsed twice with PBS to remove the loosely adherent planktonic MRSA and suspended in PBS (200 μL). The recovered bacteria outside the biofilm and the bacteria inside the biofilm were serially diluted and plated in an agar plate for 24 h to count CFU. The biofilm was also stained by SYTO9 for 15 min. The biofilm was then gently rinsed with PBS. The three-dimensional structure and thickness of the biofilm were detected by Leica TSC SP2 confocal microscopy as described previously.

**Genomic DNA Analysis Of MRSA**

MRSA at OD$_{600}$=3 was treated by CPC (100 μg/mL) for 3 h. The genomic DNA was extracted using a Presto® Mini Bacteria kit based on the manufacturer’s instructions. The aliquot of purified genomic DNA (100 ng) was analyzed by electrophoresis on an 0.8% agarose gel.

**Total Protein Analysis Of MRSA**

MRSA was grown in TSB to OD$_{600}$=6 and then treated with CPC (100 μg/mL) at 37°C for 3 h. The centrifuged pellet was resuspended with water. Following sonication for 20 min, the bacteria were centrifuged at 4°C and 10,000 rpm for 15 min. The total protein was quantified by a Bio-Rad protein assay kit in ELISA (595 nm).

**Proteomic Profiles Of MRSA**

The treatment protocol of CPC was the same as the total protein analysis. The SDS-PAGE was carried out with a 5% stacking gel and a 10% separating gel followed by Coomassie blue staining. The bands were withdrawn and digested with trypsin overnight. The digested proteins were acidified with 0.5% trichloroacetic acid. The MALDI-TOF/TOF was detected by Bruker Ultraflex spectrometer. The detailed procedure was shown in the previous work. The fold change of the protein expression after CPC treatment was estimated by band quantification based on Prodigy Samespots analysis software. The protein masses were assigned and used for a database search with MASCOT search engine (www.matrixscience.com).
Statistical Analysis
The data presented as mean and standard deviation (S.D.). The difference in the data of the different experimental groups was analyzed using the Kruskal-Wallis test. The post hoc test for checking individual differences was Dunn’s test. The significance was demonstrated as * for p<0.05, ** for p<0.01, and *** for p<0.001 in the figures.

Results
Characterization Of Micelles And Nanoemulsions
The three nanoemulsions prepared by homogenization and sonication were depicted based on droplet size and zeta potential. CPC alone could self-assemble into micelles without the fabrication process. Table 1 compares the size, polydispersity index (PDI), and surface charge between micelles and squalene@CPC nanoemulsions. Self-assembly of CPC formed 1435 nm aggregates. The directed assembly of nanoemulsions showed much smaller droplets as compared to micelles. The droplet size increased from 55 nm (CN-S) to 165 (CN-M) and 245 nm (CN-L) with the increase of squalene from 2% to 7%. The micelles revealed a broad size distribution (PDI=1). The PDI of CN-S, CN-M, and CN-L was 0.40, 0.19, and 0.26, respectively. The surface of all formulations had cationic charges with zeta potential ranging from 9.3 to 51.1 mV, mainly due to the cationic feature of CPC. The positive charge of nanoemulsions was greater than that of micelles. CN-L showed a zeta potential slightly higher than that of the nanoemulsions with smaller droplets.

Figure 1A demonstrates the change in size as a function of storage time at 50°C. The micellar size increased very quickly during the first day of storage. The average diameter of micelles at Day-1 was >5,000 nm, which was the maximum detection limit of Zetasizer. All nanoemulsions retained their droplet size after 14 d. Figure 1B shows the emission spectrum of Nile red in micelles and nanoemulsions. Nile red fluorescence is quenched in the molecular environment with high hydrophilicity. The micelles exhibited the weakest fluorescence, implying the least lipophilicity. The smaller-sized nanoemulsions revealed a higher lipophilicity compared to the larger ones.

Antibacterial Effect Of Nanoemulsions Against Planktonic MRSA
To evaluate whether the response to micelles and nanoemulsions varies among different species of pathogens, we determined MIC and MBC against Gram-positive bacteria (MRSA, S. aureus, KM-1, and KV-2), Gram-negative bacteria (E. coli), and fungi (C. albicans). Table 2 presents the MIC of the CPC formulations. MIC is defined as the lowest CPC concentration that contributes to complete inhibition of visible growth. The results demonstrated that all formulations studied showed inhibitory activity for all pathogens investigated. Although a significant difference had been achieved in the droplet size by varying squalene content, the inhibitory effect was approximate. No inhibition was detected with the treatment of bulk materials (squalene, poloxamer 188, and SPC). This suggested that the inhibitory activity was mainly due to CPC in the formulations. The organisms of Gram-positive bacteria were found to be more susceptible to CPC than Gram-negative bacteria and fungi. The MIC assay indicated a superior inhibition toward Gram-positive bacteria compared to Gram-negative bacteria by about 8-fold. The CPC inhibition against MRSA was weaker than that against drug-sensitive S. aureus. The susceptibility of clinically isolated MRSA and VISA to CPC was analyzed. VISA is a strain highly prevalent in the epidemic MRSA species. CPC could significantly suppress the growth of both clinical isolates. MBC expressed the minimum bactericidal amount to kill >99.9% bacteria. As shown in Table 2, MBC values are slightly higher or comparable to the degrees of MIC. The same as MIC, Gram-positive bacteria were more sensitive to CPC, followed by E. coli and C. albicans. CPC at 1.95 μg/mL completely killed MRSA.

The MRSA inhibition by CPC formulations was recognized by the agar diffusion study as shown in Figure 2A. CPC displayed a clear inhibition zone on the agar media against planktonic MRSA. We observed no difference in the effect of any of the droplet sizes on MRSA growth.
Application of CPC provided the inhibition zone of about 9 mm for all formulations tested. SEM allowed us to directly visualize MRSA integrity and morphology after CPC treatment. As shown in Figure 2B, the untreated MRSA reveals a smooth and bright surface without debris. The CPC treatment disrupted the intact morphology, leading to the shrinkage of some of the MRSA membrane. The enlarged image showed surface blebbing and roughening after CPC treatment (Figure 2C). The formation of vacuoles and debris was detected. The bacterial membrane damage resulted in the cytoplasmic material leakage. The different droplet sizes did not significantly affect the level of interference.

The live/dead staining was conducted to further assess the planktonic MRSA killing by micelles and nanoemulsions. Live MRSA with intact membrane would be stained green by SYTO9-DNA interaction. PI is a molecule that diffuses to the bacteria with compromised membrane. The PI uptake is an implication of cellular injury and death. Figure 3A and B represent the live/dead images of MRSA at a CPC dose of 10 and 100 μg/mL, respectively. The PI signal was increased by nanoemulsion treatment as compared to the untreated control. SYTO9 covered the same area with PI in the nanoemulsion-treated groups. This was because SYTO9 stained all cells while PI stained only dead cells. PI would cover the entire visual field of

Table 2 The Minimum Inhibitory Concentration (MIC, μg/mL) And Minimum Bactericidal Concentration (MBC, μg/mL) Of Cetylpyridinium Chloride (CPC) In Micelles And Cationic Nanoemulsions Against Different Strains Of Bacteria

| Strain | MIC | MBC |
|--------|-----|-----|
| M. aureus | 0.49–0.98 | 0.49–1.95 |
| KM-1 | 0.49–0.98 | 0.49–1.95 |
| KV-2 | 0.49–0.98 | 0.49–1.95 |
| E. coli | 3.91 | 7.81 |
| C. albicans | 1.95–3.91 | 7.81–15.63 |

**Table 2** The Minimum Inhibitory Concentration (MIC, μg/mL) And Minimum Bactericidal Concentration (MBC, μg/mL) Of Cetylpyridinium Chloride (CPC) In Micelles And Cationic Nanoemulsions Against Different Strains Of Bacteria

Note: KM-1 and KV-2 are the clinical isolates of MRSA and VISA, respectively. Each value represents the mean±S.D. (n=3).
SYTO9. An accumulation of red signal was observed in the nanoemulsion-treated MRSA. The accumulation was more obvious following the increase of droplet size. This could be due to the massive loss of cell integrity causing DNA release from the bacteria. The red signal aggregation was not visualized in MRSA treated with micelles. The SYTO9 staining was also absent in the group of micelles. This could be because micelles prompted complete MRSA damage, leaving the entire DNA disruption. The detailed explanation is explored in the Discussion section.

Antibacterial Effect Of Nanoemulsions Against Biofilm MRSA

Figure 4 presents the antibiofilm impact of micelles and nanoemulsions against one-day-old MRSA biofilm cells. Figure 4A shows the corresponding percentage of MRSA inhibition inside biofilm measured as CFU. Similar to the strong killing on planktonic cells, CPC could eliminate MRSA in biofilm. The biofilm formation was inhibited up to 89%, 82%, and 78% after application of CN-S, CN-M, and CN-L, respectively. This suggested that the size reduction led to an improvement in the nanoemulsion penetration into the biofilm matrix. The antimicrobial activity towards planktonic MRSA outside the biofilm was also examined as depicted in Figure 4B. A complete planktonic cell killing was observed by micelle treatment. Contrary to the result of MRSA inside the biofilm, treatment by nanoemulsions with smaller droplets showed less CFU reduction compared to the larger ones. CN-M and CN-L consistently produced a lower CFU percentage against MRSA outside the biofilm than that against bacteria inside the biofilm. The antibiofilm effect was seen by SYTO9 staining as illustrated in Figure 4C. The viable MRSA in the intact biofilm (control) displayed a dense and thick architecture. CPC was vital to diminishing the viability and thickness of biofilm MRSA. The SYTO9
intensity and biomass thickness was calculated as shown in Figure 4D and E, respectively. There was a 3.5-fold decrease of green fluorescence by micelle treatment. CN-S and CN-L further reduced the green signal as compared to micelles. The mid-sized nanoemulsions are less toxic to biofilm MRSA based on green intensity. The treatment of CPC reduced the average thickness of the biofilm. The biofilm treated with micelles, CN-S, CN-M, and CN-L was 27.2, 23.1, 24.4, and 21.7 μm, respectively. The trend of thickness inhibition was similar to that of SYTO9 intensity.

The Genomic And Proteomic Profiles Of MRSA Treated By Nanoemulsions
The genomic DNA detection was conducted to evaluate the anti-MRSA activity of CPC as shown in Figure 5A. The micelles exerted a significant 2.5-fold reduction of DNA concentration compared to the untreated control. The DNA elimination by nanoemulsions increased with lowering the droplet size. The application of CN-S caused a 5-fold decrease in the DNA concentration compared to the control. Figure 5B expresses that micelles and nanoemulsions decrease the cellular protein content. This reduction could be due to the bacterial membrane injury and/or the inhibition of protein synthesis after DNA denaturalization. The protein reduction by the nanoemulsions was not size-dependent. The changes in the MRSA proteins by CPC were further examined by SDS-PAGE. The control MRSA exhibited apparently clear and intense protein bands as shown in Figure 5C. The protein bands of >75 kDa had faded after treatment with CPC formulations. No significant change of protein bands of <75 kDa was observed before and after CPC treatment. The qualitative and quantitative analysis of the proteins is conducted by mass as summarized in Table 3. The protein change of >±2-fold can be regarded as significant. DNA-directed RNA polymerase (RNAP) subunits β and β’ (band No. 1 and 2) were significantly downregulated by CPC. The other proteins (band No. 3~16) generally showed negligible change after CPC application.

Discussion
Most of the investigations involved in the antimicrobial nanoemulsions have used susceptible bacteria as the model pathogens. Further study of the capability of nanoemulsions to eradicate drug-resistant microorganisms.
is required. In the present study, we employed CPC to form cationic nanoemulsions, and we attempted to optimize the formulations for achieving anti-MRSA activity. Our ability to control the droplet size gave the benefit of showing how specific sizes influence bacteria killing. We demonstrate here that squalene@CPC nanosystems could significantly inhibit the growth of MRSA and VISA. Most of the evaluation platforms showed no significant effect of size on anti-MRSA activity. In the experiment of biofilm suppression, the smaller-sized nanostructures more facilely restrained MRSA viability than did the bigger ones. A similar trend (CN-S>CN-M>CN-L) was observed when detecting the total genomic DNA loss in MRSA. The nanoemulsions potentially represented the capacity to treat MRSA and those that developed an impermeable biofilm.

The surface charge of cationic nanomaterials depends upon the number of cationic surfactants on the particulate surface and the associated degree of the counterions. Poloxamer 188 and SPC as the emulsifiers in lipid nanoparticles offer negative charge to the surface. The high positive charge in squalene@CPC nanoemulsions reported here verified the CPC association with the droplet shell. Besides the aggregates of CMC to create micelles, some monomers existed in the aqueous dispersion. This is the reason for the low zeta potential obtained from the micelles. The size of the micelles was much larger than that of the nanoemulsions. The micelles with CPC alone might produce electrostatic repulsion, resulting in the loose structure with large size. The blend of CPC with anionic emulsifiers caused electrostatic attraction, leading to an enhancement of particulate cohesion and condensation. Previous study also suggests the formation of a close surface by the introduction of phospholipids into the cationic nanostructures. The nanoemulsion size was directly proportional to the

Figure 4 Determination of the antibacterial activity of micelles and squalene@CPC nanoemulsions against biofilm MRSA: (A) MRSA CFU inside the biofilm; (B) MRSA CFU outside the biofilm; (C) the three-dimensional images of biofilm visualized by confocal microscopy; (D) the corresponding biofilm green intensity; and (E) the corresponding biofilm thickness. All data are presented as the mean of three experiments±S.D. ***p < 0.001; **p < 0.01; *p < 0.05.
squalene percentage. Compared to micelles, squalene@CPC nanoemulsions were more stable. Zeta potential serves as an important fact governing the stability as it provides electrostatic repulsion among the charged droplets. The nanoemulsions processed greater zeta potential, signifying a high-energy barrier to maintain the storage stability. The metastable solid form of CPC in the micellar dispersion suggested that the stability of micelles is easily disturbed by the environmental change.

Our results suggested a stronger growth inhibition of CPC against Gram-positive than Gram-negative microbes. Gram-negative bacteria possess an outer membrane in the cell wall, forming an extra barrier against molecular penetration. On the other hand, there are plenty of pores in the cell wall of *S. aureus* comprising teichoic acid and peptidoglycan. The greater barrier function of Gram-negative bacteria than Gram-positive bacteria had led to the poor delivery of the nanoemulsions into Gram-negative microbes. Our findings highlighted the potential of nanoemulsions for treatment of *S. aureus* that are resistant to methicillin and vancomycin. VISA is *S. aureus* that can transform the cell wall and become less susceptible to vancomycin in the presence of some pressures. This is becoming a public health problem. The antibacterial mechanism of cationic surfactants is predominantly based on the membrane damage. Cationic surfactants facilely attach to the MRSA surface with negative charge via electrostatic affinity. In addition, the long alkyl chain of CPC can partition into the lipophilic region of the membrane. The increased total surface area of nanoemulsions improves the interaction with the site of action. The evidence of MRSA membrane impairment was supported by the genomic and proteomic profiles of MRSA treated by micelles and squalene@CPC nanoemulsions.

Figure 5 The genomic and proteomic profiles of MRSA treated by micelles and squalene@CPC nanoemulsions: (A) analysis of the quality of MRSA genomic DNA by agarose gel electrophoresis; (B) total protein concentration in MRSA; and (C) the protein change of MRSA analyzed by SDS-PAGE and MALDI-TOF/TOF mass. All data are presented as the mean of three experiments±S.D. ***p < 0.001; **p < 0.01; *p < 0.05.
Table 3 Differentially Expressed Proteins In MRSA Following The Treatments Of Micelles And Cationic Nanoemulsions

| Band No. | Protein                                      | Accession No. | Mw (Da) | Matched Peptides | Coverage % (SCORE) | Ratios To MRSA<sup>a,b</sup> | Biological Function                                                                                                                                                                                                 |
|---------|----------------------------------------------|---------------|---------|------------------|--------------------|-----------------------|----------------------------------------------------------------------------------|
| 1       | DNA-directed RNA polymerase subunit β        | P60279        | 133,409 | 19               | 16% (100)          | -2.33 −3.23 −8.34 −3.45 | DNA-dependent RNA polymerase catalyzes the transcription of DNA into RNA using the four ribonucleoside triphosphates as substrates.                                                                                       |
| 2       | DNA-directed RNA polymerase subunit β'       | P60286        | 134,327 | 24               | 24% (115)          | -2.78 −0.71 −3.57 −0.64 | DNA-dependent RNA polymerase catalyzes the transcription of DNA into RNA using the four ribonucleoside triphosphates as substrates.                                                                                       |
| 3       | Transketolase                                | Q6GH64        | 72,235  | 15               | 28% (136)          | 0.64 2.07 1.16 2.44    | Catalyzes the transfer of a two-carbon ketol group from a ketose donor to an aldose acceptor, via a covalent intermediate with the cofactor thiamine pyrophosphate.                                                      |
| 4       | Phosphoenolpyruvate carboxykinase (ATP)      | Q8NVZ8        | 59,599  | 26               | 56% (168)          | -0.59 -0.21 -0.96 -0.54 | Involved in the gluconeogenesis. Catalyzes the conversion of oxaloacetate (OAA) to phosphoenolpyruvate (PEP) through direct phosphoryl transfer between the nucleoside triphosphate and OAA.                        |
| 5       | Catalase                                     | Q8NWV5        | 51,221  | 14               | 47% (141)          | -0.28 0.95 0.97 2.20   | Decomposes hydrogen peroxide into water and oxygen; serves to protect cells from the toxic effects of hydrogen peroxide.                                                                                                   |
| 6       | Enolase                                      | P64079        | 46,036  | 14               | 47% (90)           | 0.91 -1.59 1.82 1.18   | Catalyzes the reversible conversion of 2-phosphoglycerate into phosphoenolpyruvate. It is essential for the degradation of carbohydrates via glycolysis.                                                                   |
| 7       | Pyruvate dehydrogenase E1 component subunit alpha | P60090   | 36,459  | 7                | 34% (84)           | -3.03 -2.50 -3.57 -1.94 | The pyruvate dehydrogenase complex catalyzes the overall conversion of pyruvate to acetyl-CoA and CO₂.                                                                                                              |
| 8       | L-lactate dehydrogenase                      | P65257        | 34,650  | 15               | 58% (96)           | -2.33 -1.87 -2.22 -2.21 | Catalyzes the conversion of lactate to pyruvate. Appears to be the primary factor that allows <i>S.aureus</i> growth during nitrosative stress in both aerobically and anaerobically cultured cells.                           |
| 9       | Cysteine synthase                            | P63872        | 33,013  | 9                | 45% (84)           | -2.86 -0.27 -0.31 -1.85 | O-acetyl-L-serine + hydrogen sulfide = L-cysteine + acetate.                                                                              |

(Continued)
Table 3 (Continued).

| Band No. | Protein | Accession No. | Mw (Da) | Matched Peptides | Coverage % (SCORE) | Ratios To MRSA<sup>a,b</sup> | Biological Function |
|----------|---------|---------------|---------|------------------|-------------------|-----------------------------|-------------------|
|          |         |               |         |                  |                   | Micelles | CN-S | CN-M | CM-L |
| 10       | Pyridoxal 5’-phosphate synthase subunit PdxS | P60799 | 32,114 | 8                | 42% (95)  | -0.07 | 1.13 | 0.99 | 2.34 |
| 11       | 2,3-bisphosphoglycerate-dependent phosphoglycerate mutase | P65709 | 26,697 | 24               | 86% (204) | 1.08  | 1.19 | -0.12 | -0.21 |
| 12       | DUF642 domain-containing protein | PF07852.6 | 22,691 | 9                | 40% (82)  | -1.79 | 1.22 | -0.95 | 1.25 |
| 13       | Superoxide dismutase [Mn/Fe] 2 | P66832 | 23,098 | 10               | 72% (92)  | 1.06  | 1.09 | 1.99 | 1.82 |
| 14       | Ferritin, partial | Q7A0HS | 16,627 | 7                | 52% (112) | 1.04  | 1.78 | 1.55 | 2.17 |
| 15       | DNA protection during starvation protein 2 | Q8RPQ2 | 16,749 | 9                | 76% (88)  | 1.23  | 1.16 | 0.93 | 1.58 |
| 16       | DNA-binding protein HU | Q7A0U9 | 9,606  | 7                | 72% (96)  | 1.14  | 1.25 | 1.28 | 1.61 |

Notes: <sup>a</sup>Ratios to MRSA indicated the fold changes in protein volume among micelles, CN-S, CN-M, and CN-L versus MRSA samples, respectively. The ratios > 1.0 mean the proteins whose expression levels were increased upon treatments of compounds, while ratios < -1.0 indicate the proteins were downregulated under the exposure to compounds. <sup>b</sup>Analyzing the gel images using GeneTools software.
by SEM and live/dead imaging. The distortion involved the pore formation and the subsequent loss of cytoplasmic materials. Besides the leakage of molecules and ions, it is possible that DNA, proteins, and lipids in MRSA would be denatured by nanoemulsion treatment.

The aggregates of DNA were found after treatment of squalene@CPC nanoemulsions in the live/dead visualization, especially the bigger droplets. In addition to the DNA leakage from the damaged MRSA, cationic nanoemulsions acted as a crosslinker of DNA to develop the aggregates. CPC provided positive surface charge and long alkyl chains, which were capable of winding with negatively charged DNA. It has been demonstrated that cationic lipid nanoparticles can efficiently adsorb DNA in the biological medium.21 The higher positive charge of CN-L might increase this effect to form the larger aggregates. The limited zeta potential and loose structure of micelles contributed to the absence of DNA aggregates. The bactericidal behavior of nanoparticles is suggested with respect to particulate size, surface charge, and lipophilicity. As the size decreases, the surface of bacteria covered by the nanoparticles increases. The stronger nanoparticle-bacteria interaction is observed for the smaller size. This can be the pivotal factor in promoting antibacterial activity. This phenomenon can be found in the nanoparticles made of polymers and metals.29–31 This was not found in our case since no significant size effect was detected in the assays of MIC/MBC, agar diffusion, SEM, and live/dead imaging. This may suggest that the droplet diameter of cationic nanoemulsions was not the major cause of regulating the battle against planktonic MRSA.

Biofilm is a sessile structure with clusters of bacteria enclosed in extracellular polymeric substance (EPS). Biofilm formation is an essential factor contributing to the infection of S. aureus.32 Because of the rigid architecture, most of the antibiotics cannot freely diffuse into the biofilm. Biofilm also prefers gene transfer between bacteria, spreading the antibiotic resistance for virulence.33 Although CPC has been well documented to reveal antibacterial activity against planktonic culture, the investigation of biofilm is limited.34 Our results demonstrated an efficient prevention of biofilm development by squalene@CPC nanoemulsions. The nanoparticles are highly beneficial in disassembling biofilm due to the profuse penetration into the porous matrix and close contact with the biofilm surface offering a high concentration of antibacterial agents.35

The lipophilic nature of nanoemulsions can produce interaction with EPS, leading to the disruption and disengagement of the lipid layer.36 Biofilm is composed of polysaccharides, proteins, and DNA. The cationic CPC can be a chelator to interact with extracellular DNA. The CFU determination inside the biofilm showed that the matrix was not a major barrier for the entry of nanoemulsions, especially the smaller droplets. The ability of nanoparticles to penetrate into the biofilm can be affected by their size. The smaller droplets easily penetrated the biofilm mesh. The anti-MRSA effect was enhanced because of the larger contact surface of the smaller droplets. The cationic ammonium surfactants can make lipophilic interaction with S. aureus.37 The incrementation of lipophilicity may promote the nanoparticle diffusion in the biofilm. The lipophilicity of the smaller-sized nanoemulsions was greater than that of the larger-sized nanoemulsions based on the molecular environment assay. This allowed the higher biofilm deterioration of the smaller droplets.

Some of the bacteria can be dispersed from the biofilm to the perimeter in the final stage of EPS development.38 The tendency of bacterial killing outside the biofilm was quite different from that inside the biofilm. Since it was more difficult for the bigger droplets to diffuse into the EPS, these large-sized droplets remained outside the biofilm to eradicate the planktonic cells derived from the matrix. Different from the CFU determination in biofilm, the trend of SYTO9 intensity was CN-M>CN-S>CN-L. CFU evaluation is simply an indicator of bacterial viability. The SYTO9 staining manifests both the cell viability and the ability to interact with DNA. The mid-sized nanoemulsions might exhibit a strong binding with extracellular DNA in biofilm. Further study is needed to show the evidence. Overall, the smaller-sized nanoemulsions can be an ideal antibiofilm agent to deliver into the matrix and exhibit a bactericidal effect. The micelles also showed a strong activity towards biofilm MRSA. There was almost no viable MRSA inside and outside the biofilm after micelle treatment. Both monomers and micelles coexisted in the aqueous CPC dispersion. The monomers of cationic surfactants quickly bind to the bacterial surface, disturbing the membrane structure.32,39 The tiny volume of monomers is also advantageous for diffusing into the biofilm mesh.

It was apparent that the DNA concentration reduction was magnified when the droplet size decreased. However, the size effect on the total protein concentration was not apparent. Both DNA and protein could be lost by the capability of CPC to damage the cellular membrane. According to SEM, some bacteria maintained surface integrity after CPC treatment. The mechanisms other
than DNA and protein leakage should be discussed. The intercalation of antibacterial agents or nanoparticles into the DNA helix can block DNA replication to exert anti-MRSA activity. CPC is known to bind to and precipitate double-stranded DNA. This interaction results in the bacterial DNA degradation and denaturing. This effect may be more significant for the small-sized nanoemulsions. The DNA degradation can interfere with the DNA-protein binding in cytoplasm, inhibiting the enzymatic and metabolic activity. It is possible that CPC produced both direct and indirect membrane collapse. In order to understand the possible mechanism, proteomic profiles were established in CPC-treated MRSA. CPC significantly reduced the expression of RNAP subunits β and β′. Transcription is an important process for tight control of gene expression for S. aureus survival. All transcriptional activity is maintained by RNAP. Bacteria consist of four subunits of RNAP: α, β, β′, and σ. The downregulation of this pathway suggested that squalene@CPC nanoemulsions had a profound effect on the transcription of DNA to RNA. A similar example is rifamycin binding to the β subunit of RNAP as the antibacterial mechanism of action. Compared to RNAP, we did not find any significant concordance with the proteins involving oxidative stress, Fenton reaction, and the Krebs cycle. In-depth work on the anti-MRSA mechanisms of squalene@CPC nanoemulsions is required for further elucidation.

Conclusions
The antimicrobial nanoparticles have attracted great interest as one of the suitable candidates to substitute for conventional antibiotics. In this study, we examined how the size difference in CPC-decorated nanoemulsions influences bacterial eradication. The nanoemulsions with small, medium, and large droplet sizes had high positive zeta potential, presenting great colloidal stability. The squalene@CPC nanosystems demonstrated substantial activity against bacteria and fungi, with the greatest potential to eliminate Gram-positive bacteria. Although the self-assembled micelles also showed broad and strong antibacterial activity, the poor stability led to the limited application. The killing of MRSA by the nanoemulsions was caused by membrane disruption initiated by electrostatic force. The degradation of DNA and interference in protein synthesis were also the possible mechanisms. The enhancement of anti-MRSA impact of the nanoemulsions with size reduction was observed in biofilm form but not in planktonic form. Compared to the larger ones, the small-sized nanoemulsions could facilely enter into the biofilm matrix to exert the stronger MRSA inhibition. The total DNA concentration in MRSA was also decreased following the reduction of droplet size. A slight difference in the bactericidal activity of different nanoformulations was detected in the evaluation platforms of MIC/MBC, agar diffusion assay, and SEM. The experimental results of this study can lead to the rational design of antibacterial nanoemulsions based on the desired application in antimicrobial coatings, foods, cosmetics, and mouth rinses. The nanoemulsions also have therapeutic significance to treat MRSA infection and act as the carriers of antibiotics.

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Disclosure
The authors report no conflicts of interest in this work.

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