



OPEN Antibiofilm activity of Plumbagin against *Staphylococcus aureus*

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In chronic infections caused by *Staphylococcus aureus*, biofilm is a major virulence factor. In *Staphylococcus aureus* biofilms, bacteria are embedded in a matrix of extracellular polymeric substances and are highly tolerant to antimicrobial drugs. However, the lack of effective solutions to inhibit biofilm formation remains a challenge, and the mechanism of inhibition of biofilm formation targeting extracellular polymeric substances is unclear. The aim of the present study was to investigate the inhibitory mechanisms of Plumbagin against *Staphylococcus aureus* biofilms formation by affecting secretion of extracellular polymeric substances using the high-content screening. Our results showed Plumbagin (16 µg/mL) inhibited biofilm formation, revealing a significant reduction in both biomass and bacterial metabolic activity, and disrupted the biofilm structure, leading to a significant decrease in both biological volume and average thickness ($P \leq 0.01$). High-content screening imaging indicated that the Plumbagin treatment induced alterations in the extracellular polymeric substances of *Staphylococcus aureus* biofilm, significantly reducing the quantities of extracellular polysaccharide, proteins and extracellular DNA. Interestingly, extracellular DNA within the matrix was found to be the most sensitive to Plumbagin treatment. Extracellular DNA formation was significantly inhibited at a concentration of 4 µg/mL, whereas the inhibition of extracellular polysaccharide and proteins required a higher concentration of 8 µg/mL. Overall, these results demonstrated the inhibitory effects of Plumbagin on *Staphylococcus aureus* biofilm formation and extracellular polymeric substances secretion, suggesting that extracellular DNA may be a potential target for the anti-biofilm activity of Plumbagin. These findings will provide new insights into the mode of action of Plumbagin in treating infections caused by *Staphylococcus aureus* biofilms.

Keywords *Staphylococcus aureus*, Biofilm, Plumbagin, Extracellular polymeric substance, High-content screening

Abbreviations

<i>S. aureus</i>	<i>Staphylococcus aureus</i>
EPS	Extracellular polymeric substances
PLB	Plumbagin
PIA	Polysaccharide intercellular adhesin
eDNA	Extracellular DNA
DMSO	Dissolved in dimethyl sulfoxide
FITC-WGA	Fluorescein iso-thiocyanate-conjugate wheat germ agglutinin from <i>Triticum vulgaris</i>
LB	Luria bertani
MIC	Minimum inhibitory concentrations
CLSI	The clinical and laboratory standards institute
CV	Crystal violet
PBS	Phosphate-buffered saline
OD	Optical density
PI	Propidium iodide
AUC	Area under the curve

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Chronic wounds are a growing medical problem, resulting in high morbidity and mortality. *Staphylococcus aureus* (*S. aureus*) is one of the predominant causes of persistent infections in chronic wounds, mainly due to its remarkable ability to adapt to different environments and to acquire genes associated with antimicrobial resistance^{1,2}. Bacteria of the *Staphylococcus* genus are highly efficient in establishing biofilms resulting in persistent infections that are difficult to treat³.

Biofilms are complex microbial architectures that attach to surfaces and encase microorganisms in a matrix composed of self-produced hydrated extracellular polymeric substances (EPS). In *S. aureus* biofilms, poly-N-acetyl- β -(1–6)-glucosamine (PNAG; also known as polysaccharide intercellular adhesin PIA), proteins and extracellular DNA (eDNA) are broadly viewed as the main components of the EPS matrix⁴. Previous studies suggested that removing the residual EPS matrix could be at least as crucial as killing bacteria in the management of biofilm infections⁵. Targeting the EPS may be an effective strategy to remove biofilm, disaggregate bacteria and disrupt the pathogenic environment⁶. Additionally, due to the variability in the composition of *S. aureus* EPS matrix and the interaction between their multiple components, the strategies to disrupt the matrix should ideally target several constituents of the EPS matrix simultaneously⁷. Currently, it is believed over 80% of chronic infectious diseases are mediated by biofilms, and it is known that conventional antibiotic medications are inadequate at eradicating these biofilm-mediated infections⁸. Therefore, there is an urgent need for alternative compounds capable to treat *S. aureus* biofilms.

Natural product therapeutics are currently receiving significant attention due to their antimicrobial efficacy and their ability to avoid inducing drug resistance. Examples of such natural products include polyphenols, flavonoids, quinones, terpenoids, alkaloids, and tannins, all of which have been demonstrated to possess antimicrobial properties^{9–13}. Plumbagin (5-hydroxy-2-methyl-1,4-naphthoquinone, PLB) is a naturally occurring naphthoquinone isolated from the roots of the medicinal plant *Plumbago zeylanica* L. PLB has been documented to exhibit a wide range of biological activities including anti-inflammatory, anticancer, antidiabetic, antioxidant, antibacterial, antifungal, and anti-atherosclerotic properties¹⁴. It is considered a natural agent with the potential to control chronic diseases and has shown in vitro antibacterial activity against *S. aureus* and *Candida albicans*^{15,16}.

In this study, *S. aureus* biofilm was used as a research model for chronic wound infection in vitro to examine the antibiofilm activity of PLB against *S. aureus* and to explore the mechanism of inhibiting the formation of *S. aureus* biofilm by influencing EPS secretion.

Results

MIC determination and growth curve construction

To evaluated the sensitivity of PLB to *S. aureus*, the MIC of PLB was determined. Figure 1A illustrated that the MIC of PLB against planktonic *S. aureus* was 4 $\mu\text{g/mL}$. At a PLB concentration of 4 $\mu\text{g/mL}$, the growth of *S. aureus* planktonic cells was markedly inhibited (Fig. 1B). Given that biofilm formation significantly enhanced antimicrobial resistance, we constructed a straightforward biofilm growth curve to delineate the growth characteristics of the biofilm and pinpoint the optimal time for antibiofilm research. According to Fig. 1C, the biofilm formation of *S. aureus* reached its maximum at 24 h, after which a decline in growth was noted. Consequently, we selected the 24-hour-old biofilm as an in vitro model to evaluate the inhibitory effect of PLB on biofilm formation.

Assessment of the inhibitory effect of PLB on biofilm formation

Effect of PLB on the biofilm biomass

Based on the susceptibility of planktonic *S. aureus* cells to PLB, we evaluated the impact of PLB on biofilm formation by measuring the binding of CV to the biofilms. The antibiofilm activity of PLB at concentrations ranging from 0.5 to 64 $\mu\text{g/mL}$ was shown in (Fig. 2A). In our biofilm biomass quantification assay, we observed a concentration-dependent increase in the antibiofilm activity of PLB. Specifically, biofilm formation was reduced by 2, 43, and 92% at concentrations of 4, 8, and 16 $\mu\text{g/mL}$, respectively, as shown in (Fig. 2A). However, no significant increase in antibiofilm activity was observed when the PLB concentration surpassed 16 $\mu\text{g/mL}$. Consequently, this concentration was identified as the minimum biofilm inhibitory concentration (MBIC) for PLB.

Effect of PLB on the metabolic activity

To assess the effect of PLB on bacterial metabolic activity within biofilms, we performed an XTT reduction assay on the biofilms. The XTT assay showed that the metabolic activity of bacteria within the biofilms decreased in a concentration-dependent manner with PLB treatment. The findings from the XTT reduction assay provided clear evidence that PLB, at its MBIC, specifically targeted biofilm formation without compromising the viability of *S. aureus* (Fig. 2B). Specifically, the metabolic activity of the *S. aureus* biofilm was found to be reduced by only 64% at the MBIC. Furthermore, when the concentration of PLB was elevated to 32 $\mu\text{g/mL}$, the inhibition rate of PLB on the metabolic activity within *S. aureus* biofilms increased to 90%.

Light microscope and HCS analysis

The impact on biofilm architecture is a crucial criterion for evaluating the efficacy of any antibiofilm agent. The *S. aureus* biofilm was visualized using a light microscope after staining with CV (Fig. 2C). The untreated wells exhibited a clumped structure. In contrast, samples treated with PLB displayed a significant reduction in biofilm formation, suggesting that PLB effectively inhibited both biofilm formation and adhesion.

To substantiate our findings, the HCS system was employed to assess changes in biofilm architecture between PLB-treated and untreated biofilms. Analysis of the z-stack images, captured using this microscopy technique and processed with COMSTAT software, clearly demonstrated that PLB treatment disrupted biofilm

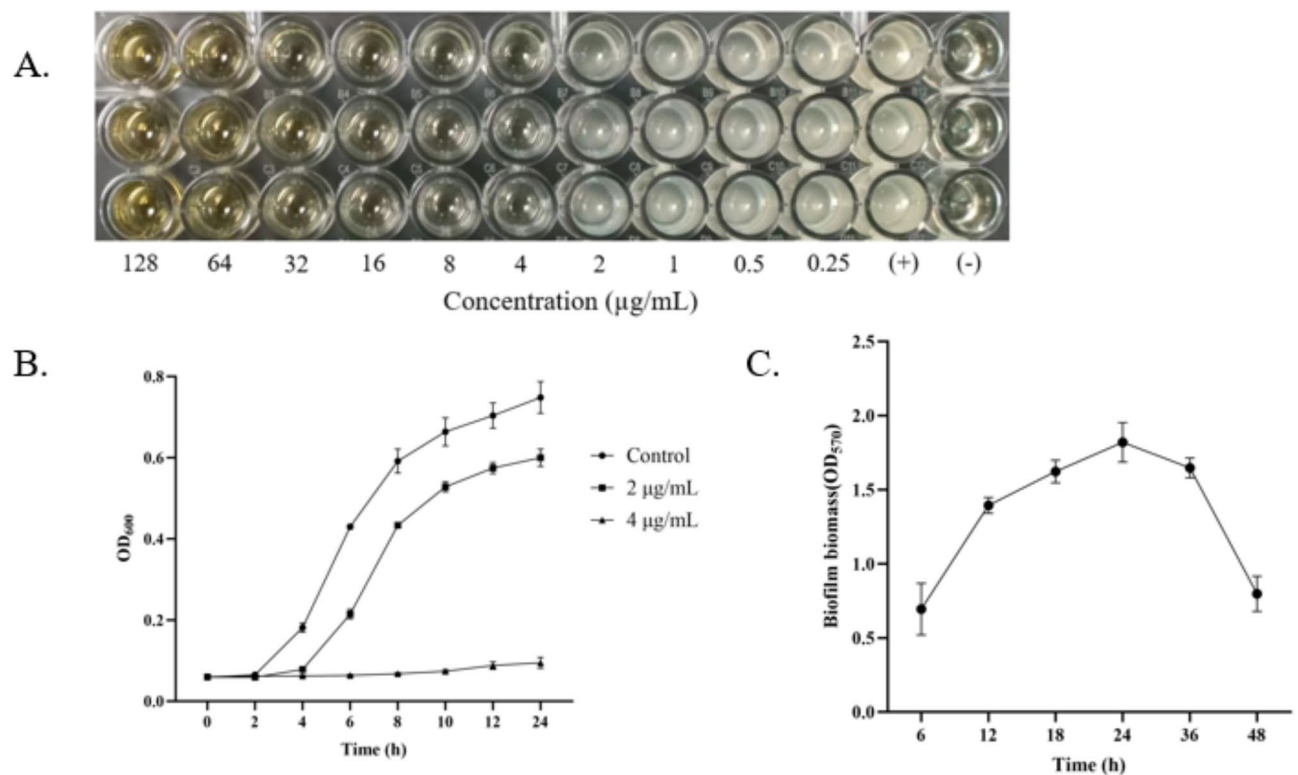


Fig. 1. MIC determination and growth curve construction. (A) The MIC of PLB against planktonic *S. aureus*. (B) The effect of PLB on growth of *S. aureus*. (C) The growth curve of the biofilm from 0 h to 48 h. Data were presented as the mean \pm standard deviation.

development. This was evidenced by a decrease in the average biofilm biovolume and thickness, as well as an increase in the roughness coefficient post-treatment, as illustrated in (Fig. 2D, E).

Effect of PLB on biofilm matrix components

EPS is crucial for maintaining biofilm architecture. To elucidate the potential mechanisms underlying the inhibition of biofilm formation, we investigated the effect of PLB on EPS secretion. Treatment of *S. aureus* biofilms with PLB led to a significant reduction in EPS production (Fig. 3). FITC-WGA staining revealed a marked decrease in PIA production in *S. aureus* biofilms treated with 8 and 16 $\mu\text{g/mL}$ of PLB when compared to the untreated control group (Fig. 3B). Additionally, the protein content within the biofilm was significantly diminished at a concentration of 8 $\mu\text{g/mL}$ of PLB (Fig. 3C). The presence of eDNA in the biofilms was assessed using PicoGreen, a novel method that involves fluorescence intensity measurement. When the stain was added, PicoGreen did not penetrate the cells but bound exclusively to eDNA. The inhibitory effect of PLB on eDNA was particularly pronounced, with a concentration of 4 $\mu\text{g/mL}$ significantly suppressing eDNA formation (Fig. 3D, E). The fluorescence intensity quantified by HCS images also observed the same trend which showed a significant change in fluorescence intensity during biofilm formation after PLB treatment (Fig. 3A).

Discussion

The presence of chronic wound infection significantly hampers the process of normal wound healing and typically necessitates antibiotic treatment. However, the excessive use of antibiotics contributes to the development of bacterial resistance against these drugs¹⁷. Biofilm formation is a bacterial resistance mechanism that acts as a physical barrier to antibiotics and leads to metabolic differences, limiting antibiotic effectiveness^{18,19}.

A variety of plant-derived compounds have been demonstrated to exhibit potential antimicrobial and antibiofilm actions, including cell membrane rupture; blockade of QS activity; inhibition of bacterial motility, cell adhesion, EPS formation, and cell proliferation; efflux pump downregulation; and synergism with other antimicrobials²⁰. PLB is a naturally occurring naphthoquinone that has been reported to exhibit antibiofilm effects. Previous studies have demonstrated that Plumbagin exhibits anti-biofilm activity against *S. aureus*, enhances the antibacterial efficacy of antibiotics, and inhibits biofilm formation by *S. aureus*^{21–23}. Our study confirms these findings and further elucidates the underlying mechanism by which PLB interferes with biofilm formation, specifically focusing on its effects on the biofilm extracellular matrix. Guo et al.²⁴ demonstrated that PLB inhibited the expression of PIA and the release of eDNA in *S. aureus* biofilms using extraction and isolation techniques. To further investigate the effects of PLB on the biofilm extracellular matrix, we employed molecular probe labeling to selectively label PIA, proteins, and eDNA. These components were then quantitatively analyzed to identify potential targets through which PLB may inhibit biofilm formation. However, its effects on biofilm

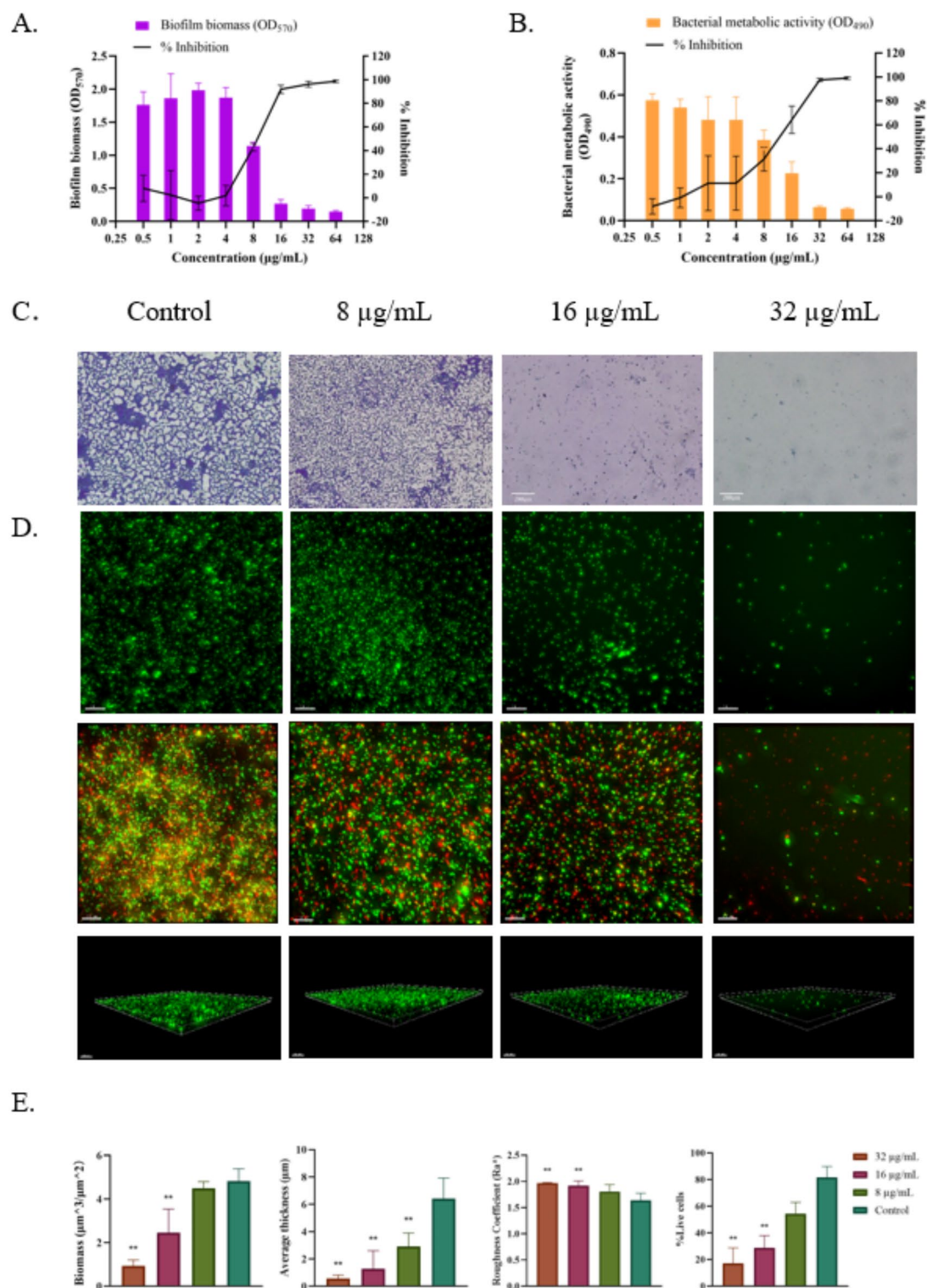


Fig. 2. Effect of PLB on *S. aureus* biofilm formation. (A) Effect of PLB on the biofilm biomass of *S. aureus* as quantified by CV staining. (B) The viability of biofilm cells of *S. aureus* was determined by XTT assay. (C) Light microscope images at 40 times magnification. (D) The HCS Z-stack maximum projections, 3D reconstructions and merge images of *S. aureus* biofilm stained with the Film Tracer™ LIVE/DEAD biofilm kit. (E) COMSTAT image analysis. Data were presented as the mean \pm standard deviation. **Indicated $P \leq 0.01$ compared to control.

formation, as well as the related molecular mechanisms, particularly against *S. aureus*, remain largely unknown. In this study, the antibacterial and antibiofilm activities of PLB were evaluated. The results demonstrated that PLB exhibited strong antibacterial activity against *S. aureus*, with a MIC of 4 µg/mL. This antibacterial activity may be related to the naphthoquinone structure of PLB. It has been reported that shikonin and luteolin, which also have the naphthoquinone structure, also have antibacterial activity against *S. aureus*^{25,26}. As demonstrated by CV staining, PLB significantly inhibited biofilm formation at a concentration of 16 µg/mL. However, the results from the XTT reduction experiment revealed that PLB had limited inhibition effects on the metabolic activity of bacteria within biofilms at this concentration. Hence, it is reasonable to state that the anti-biofilm potential of PLB in MBIC does not arise from its ability to inhibit bacterial growth. Biofilms are more resistant to antimicrobials compared to planktonic cells. The most prominent characteristic of a biofilm is the formation of a three-dimensional structure through colony aggregation and stacking²⁷. *S. aureus* biofilm is often encased in a self-produced extracellular matrix which is pivotal for the three-dimensional architecture of the biofilm, facilitating its adherence to various surfaces and ensuring the cohesion among the bacterial communities within the biofilm²⁸. The complex nature of the matrix represents a diffusion barrier for antimicrobial agents, since it limits their penetration into deeper layers of the biofilm²⁹. This indicates that the matrix components are a crucial focus in the investigation of the anti-biofilm activity exerted by plant-derived bioactive compounds (PLB) against *S. aureus*.

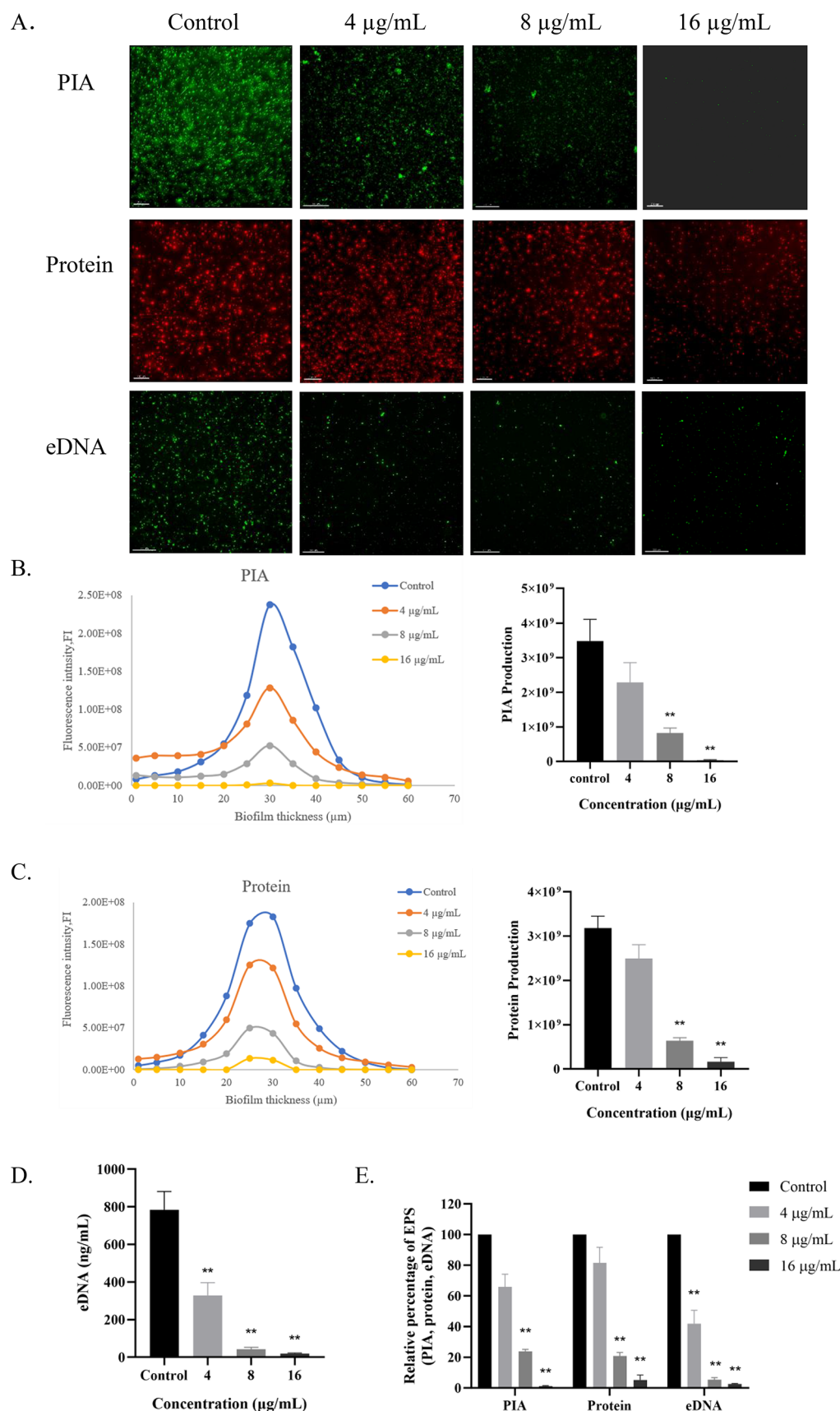
The antibiofilm effect of PLB on *S. aureus* was assessed using CV staining, a widely used method for high-throughput quantification of biofilm biomass^{30,31}. However, the use of CV has limitations including toxicity, unspecific binding to negatively charged molecules, and low reproducibility due to uneven dye extraction or differential removal of biofilm biomass throughout the washing steps^{32–34}. Traditional microscopy approaches suffer from low throughput and subjectivity in image acquisition³⁵. In this study, we utilized a novel high-throughput screening platform based on high-content screening that provided rapid, reliable, quantitative assessment of biofilm formation. SYTO9 and PI fluorescent dyes in combination with COMSTAT software were utilized to further analyze the impact of PLB on biofilm formation. The findings demonstrated that PLB disrupted the biofilm structure, leading to a significant decrease in both biological volume and average thickness. MatthewV et al.³⁵ also employed high-content screening system to assess the influence of inhibitors on *C. jejuni* biofilms viability and structural integrity. They conducted initial screening for the metabolically active living bacterial population as well as the eDNA of the biofilm using TMAMR and SytoX fluorescent dyes. Powell et al.³⁶ utilized COMSTAT image-analysis software of CLSM z-stack images to quantify the structural changes in *Pseudomonas aeruginosa* biofilms treated with oligosaccharides, indicating a disruption of the biofilm through reduction in polysaccharides and extracellular DNA components of the matrix. Although the composition of *Pseudomonas aeruginosa* biofilms is different compared to *S. aureus* biofilms, the paradigm of targeting EPS and disrupting biofilms in *S. aureus* needs to be explored. Thus, to clarify possible mechanism of inhibiting biofilm formation, the influence of PLB on the secretion of EPS was investigated.

The EPS plays a crucial role in protecting bacteria within biofilms by forming multiple layers around the cells and facilitating attachment to surfaces^{37,38}. Biochanin A³⁹ and phenyllactic acid⁴⁰ effectively prevent *S. aureus* biofilm formation through the reduction of EPS production. These polymeric substrates have been widely recognized as prominent molecular targets for the development of anti-biofilm agents.

Exopolysaccharides serve as the primary constituents of the EPS matrix, facilitating adhesion and maintaining structural integrity to form bacterial aggregates³⁹. Numerous phytochemicals have been discovered to effectively reduce biofilm formation by directly targeting exopolysaccharides within the EPS^{41–43}. Aloe-emodin⁴¹ and Quercetin⁴⁴ have demonstrated their efficacy in reducing biofilm formation by inhibiting protein synthesis. As expected, the HCS images confirmed that PLB effectively inhibited the formation of biofilm EPS.

FITC-WGA and SYPRO Ruby staining, along with fluorescence intensity measurements, revealed that PLB caused changes in the distribution of PIA and proteins during biofilm formation. At a concentration of 8 µg/mL, there was a significant reduction in PIA and protein content. For eDNA content determination, we utilized a rapid and direct method involving staining with PicoGreen followed by measuring fluorescence intensity⁴⁵. Remarkably, at a concentration of 4 µg/mL, the release of eDNA was significantly suppressed. Therefore, it can be concluded that eDNA is more susceptible to PLB. It has been demonstrated that eDNA plays an important role in the *S. aureus* biofilm matrix by acting as an electrostatic net binding cells together through the proteinaceous layer of the biofilm matrix. Given its prevalence in *S. aureus* biofilms, eDNA holds potential as a primary target for anti-biofilm therapeutics^{46,47}. This paper proposes that targeting this vital structural matrix component can be achieved through the use of a plant-derived active ingredient PLB. Since eDNA plays a crucial role in initial bacterial attachment to a surface and subsequent biofilm growth, including cell aggregation, the absence of eDNA or its inability to form a structural framework hinders the biofilm from establishing a strong foundation and maturing, ultimately inhibiting its growth^{48,49}.

Therefore, the anti-biofilm effect of PLB targeting eDNA opens up new avenues for understanding and treating *S. aureus* biofilm infections. Although our study highlights eDNA as the most sensitive EPS component to PLB, the precise mechanism by which PLB inhibits eDNA release remains to be elucidated. Based on PLB's chemical structure and prior studies on analogous compounds, we propose two non-exclusive pathways: First, PLB may directly bind to eDNA via electrostatic or intercalative interactions. The compound's fused aromatic rings resemble classical DNA intercalators such as Doxorubicin⁵⁰, while its hydroxyl and ketone groups could form hydrogen bonds with DNA bases⁵¹. Furthermore, the partial positive charge of PLB at neutral pH (predicted pKa ~ 8.2) may enhance its affinity for the negatively charged DNA backbone, enabling it to sequester eDNA and impede its integration into the biofilm matrix. Second, PLB might indirectly suppress eDNA release by targeting bacterial autolysis. Biofilm-associated eDNA is often derived from cell lysis mediated by autolysins⁵². PLB could inhibit autolysin activity or downregulate their expression, thereby reducing eDNA liberation. This hypothesis aligns with studies showing that natural compounds (e.g., Geraniol) inhibit autolysis



by downregulating autolysin genes⁵³. While the current study did not resolve whether PLB acts directly on DNA or through bacterial pathways, its structural features (e.g., planarity, charge distribution) and phenotypic effects provide a foundation for future mechanistic work.

◀ **Fig. 3.** Effect of PLB on the EPS components of *S. aureus* biofilms. **(A)** HCS images of PIA, protein and eDNA in the biofilm formation assay. **(B)** The quantitative analysis of PIA production in each sample was determined by measuring the intensity of green fluorescence using Image J software. The data was presented as the PIA production in each layer of the biofilm (1 μm sections). Total PIA production in biofilms was presented as the area under the curve. **(C)** The quantitative analysis of protein production in each sample was determined by measuring the intensity of red fluorescence using Image J software. The data are presented as the protein production in each layer of the biofilm (1 μm sections). Total protein production in biofilms was presented as the area under the curve. **(D)** The concentration of eDNA was assessed using the Quant-iT[™] PicoGreen[®] dsDNA Assay Kit. **(E)** The percentage of EPS components (PIA, protein, eDNA) in biofilms treated with various concentrations of PLB compared to the untreated control was calculated. Data were expressed as mean \pm standard deviation. **Indicates a significant difference $P \leq 0.01$ compared to control.

Our previous investigations explored the interactions between PLB and various antibiotics, including nitrofurantoin, ciprofloxacin, mecillinam, and chloramphenicol. PLB demonstrated synergistic effects with these antibiotics, significantly enhancing their growth-inhibitory activity against *S. aureus*²¹. These findings suggest that PLB could serve as a valuable adjunctive therapy in combinatorial strategies targeting biofilm-associated infections. Research indicates that PLB enhances the anti-biofilm activity of chlorhexidine (CHX) against *Klebsiella pneumoniae*. This combination reduces the MIC of CHX and significantly diminishes bacterial load in biofilms. The synergistic effect is attributed to PLB's ability to increase membrane permeability and inhibit efflux pump expression, facilitating better antibiotic penetration and efficacy⁵⁴. Integrating PLB into conventional antibiotic regimens presents a promising strategy to combat biofilm-related infections, particularly those caused by multidrug-resistant pathogens. By improving antibiotic penetration and potency, PLB may reduce required antibiotic dosages, potentially mitigating side effects and delaying the emergence of resistance.

As a natural product, PLB exhibits broad-spectrum antimicrobial and antibiofilm activities against diverse bacterial species, highlighting its potential as a novel anti-infective agent. However, its achievable concentrations in the body and potential toxicity have not been thoroughly studied. Future research should focus on the pharmacokinetic properties of PLB, assessing its absorption, distribution, metabolism, and excretion in the human body to determine its feasibility for clinical application. Additionally, preclinical and clinical trials are necessary to evaluate the safety and efficacy of PLB, providing scientific evidence for its development as an anti-infection drug.

Conclusion

Overall, this study demonstrated that PLB is an effective anti-biofilm agent. Moreover, its anti-biofilm does not arise from its ability to inhibit bacterial growth, so we investigated the mechanism by which PLB inhibits biofilm formation, focusing on its influence on EPS secretion. We used a high-content system to confirm that PLB inhibits EPS formation during biofilm development, and that PLB is more sensitive to eDNA of biofilm, suggesting that eDNA is a target for the observed PLB anti-biofilm activity. Given its enhanced antibacterial efficacy, PLB represents a promising alternative for biofilm inhibition and offers a novel strategy for treating *S. aureus* biofilm infections. However, further research is necessary to elucidate the precise mechanisms of action of PLB in biofilm inhibition and to evaluate its clinical potential as an anti-biofilm agent.

Materials and methods

Chemicals

PLB was purchased from the Aladdin Industrial Corporation (Shanghai, China). PLB was dissolved in dimethyl sulfoxide (DMSO) at a final stock concentration of 50 mg/mL. Film Tracer[™] LIVE/DEAD biofilm kit, Film Tracer[™] SYPRO Ruby, and the Quant-iT[™] PicoGreen[®] dsDNA Assay Kit were purchased from Invitrogen (Thermo Fisher Scientific, USA). Fluorescein iso-thiocyanate-conjugate wheat germ agglutinin from *Triticum vulgaris* (FITC-WGA) was purchased from Sigma-Aldrich. All other chemicals were of analytical grade.

Bacterial strains and growth conditions

S. aureus CMCC(B) 26,003 was purchased from China Institute of Food and Drug Control and stored at -80°C with 25% glycerol (Aladdin Reagent Ltd., Shanghai, China). The bacteria were inoculated in Luria-Bertani (LB) medium and incubated at 37°C for 18–22 h. The culture was then adjusted to an optical density (OD) of 0.1 at 625 nm, corresponding to approximately 10^8 CFU/mL, which represents the mid-logarithmic growth phase of *S. aureus*⁵⁵. This cell suspension was used for subsequent experiments. The bacteria were inoculated in LB supplemented with at 37°C for 16 h to obtain logarithmic phase cells for biofilm assays. Subsequently, the bacterial suspensions were incubated at 37°C for 24 h in LB medium containing 3% glucose to promote biofilm formation, as previously described with slight modifications^{56,57}.

Minimum inhibitory concentrations (MIC) of PLB against *S. aureus* in planktonic state

The MIC of PLB against *S. aureus* in planktonic state was determined using the broth microdilution technique, following the guidelines of the Clinical and Laboratory Standards Institute (CLSI), with slight modifications. Briefly, various concentration of PLB ranging from 0 to 128 $\mu\text{g/mL}$ were added into a 96-well microtiter plate with the bacterial suspensions and incubated 37°C for 24 h. The bacterial growth was measured as optical density by a full-wavelength microplate reader at 625 nm. The MIC of PLB was determined as the concentration that resulted in a 90% inhibition.

Growth curve analysis

The effect of PLB on *S. aureus* growth was tested as previously described⁵⁸. To obtain growth curve, 1% inoculum from standard cell suspension (around 1×10^8 CFU/mL) of *S. aureus* were added to 50 mL sterile LB broth with PLB at 2 and 4 µg/mL and incubated at 37 °C with 120 r/min for 24 h. The bacterial cell density (OD at 600 nm) was measured at selected time intervals (0, 2, 4, 6, 8, 10, 12 and 24 h) using a microplate reader (Tecan Spark, Männedorf, Switzerland).

To characterize the temporal progression of *S. aureus* biofilm formation, we quantified biofilm development at 6, 12, 18, 24, 36, and 48 h using crystal violet staining. The biofilm formation assay was performed in 96-well microtiter plates, as previously described with slight modifications^{30,31}. Biofilms were initiated by inoculating *S. aureus* into LB medium supplemented with 3% glucose and incubated at 37 °C. Following incubation at each time point, the biofilms were stained with 0.1% crystal violet and the optical density (OD) of the resulting solution was measured at 570 nm using a microplate reader. All assays were performed in triplicate.

Crystal Violet (CV) staining assay

The biofilm formation assay was performed in 96-well microtiter plates, as previously described with slight modifications^{30,31}. Briefly, *S. aureus* was inoculated into wells containing PLB (0–64 µg/mL) and incubated for 24 h at 37 °C. After incubation, the planktonic cells were removed and the wells rinsed twice with sterile water. Then, biofilms in the wells were stained with 0.1% crystal violet for 20 min. After removal of unbound dye, the stained biofilms were washed gently with sterile water, and crystal violet dye bound to biofilms was dissolved using 200 µL of ethanol. The optical density (OD) of the ethanol solution described above was measured at 570 nm using a microplate reader. All assays were performed in triplicate. The percentage of inhibition was calculated according to the following formula⁵⁸ (Formula(1)). Additionally, the stained biofilms were observed by using a light microscope (Olympus IX53, Tokyo, Japan) at 40 times magnification.

$$\text{inhibition}\% = \frac{\text{Control OD}_{570\text{nm}} - \text{Treated OD}_{570\text{nm}}}{\text{Control OD}_{570\text{nm}}} \times 100\% \quad (1)$$

Metabolic activity detection by XTT assay

The effect of PLB treatment on *S. aureus* metabolic activity and proliferation was determined using an XTT assay. The culture method of biofilm and the concentration of Plumbagin were same as above. After 24 h of biofilm culture, the culture medium was removed, and the wells were washed three times with PBS to remove planktonic cells. Samples were then incubated with the solution of 0.5 mg/mL XTT (Sigma-Aldrich, Shanghai, China) and 10mM menadione in the dark for 3 h. The conversion of tetrazolium salt XTT to a colored formazan derivative was measured at 490 nm in a 96-well plate. All assays were performed in triplicate. The percentage of inhibition was calculated according to the following formula⁵⁸ (Formula (2)).

$$\text{inhibition}\% = \frac{\text{Control OD}_{490\text{nm}} - \text{Treated OD}_{490\text{nm}}}{\text{Control OD}_{490\text{nm}}} \times 100\% \quad (2)$$

Representation and quantification of biofilm three-dimensional structure

For the representation and quantification of biofilm, the Film Tracer™ LIVE/DEAD biofilm kit, which includes two stains, SYTO 9 dye and propidium iodide (PI) dye, was used. Firstly, the biofilms were cultured on the black 96-well plates as described the above method. After removal of planktonic cells, the biofilm was stained with SYTO9 and PI according to the kit instructions. All samples were stained for 30 min in the dark at room temperature with an appropriate volume of a mixture of the two dyes SYTO9 and PI (3 µL of each component in 1 mL of filter-sterilized water). Furthermore, the images of the stained biofilm were obtained with an ImageXpress^{micro} high-content screening system (Molecular Devices, Sunnyvale, CA)⁵⁹. And then, the Imaris software (version 9.0.1, Oxford Instruments, Abingdon, UK) was utilized to represent a three-dimensional structure of the biofilms. For the quantification of biofilm, at least four random optical fields were chosen and observed. Additionally, a series of z-stack images were analyzed using Image J (version 2.14, Wayne Rasband, NIH, Bethesda, MD, USA) and COMSTAT⁶⁰ image analysis software, and the three-dimensional biofilms were quantified by the measurement of biofilm volume, surface roughness and biofilm depth. All assays were performed in triplicate.

Quantification of biofilm matrix components

To assess the impact of PLB on the extracellular matrix content of biofilms, the biofilms were allowed to form in black 96-well microtiter plates in the absence or presence of PLB for 24 h at 37 °C.

Quantification of PIA

FITC-WGA was used as a fluorescent probe to evaluate the formation of PIA in biofilms under the influence of PLB⁶¹. After removal of planktonic cells, biofilms were stained with a volume of 100 µL FITC-WGA (10 µg/mL) per well and plates were incubated at room temperature in the dark for 2 h. The plates were washed three times with PBS and then 100 µL of PBS were added to each well. Stained biofilms were observed with HCS system. Three-dimensional images of PIA were constructed using Imaris software. At least four random optical fields were captured and analyzed. The production of the PIA was calculated according to the fluorescence intensity using Image J. The data was presented as the PIA production in each layer of biofilm (1 µm). The percentage of PIA in biofilms treated with PLB was presented as area under the curve (AUC) and compared to untreated control. Three separate experiments were conducted, and a single set of representative data was presented.

Quantification of extracellular proteins

FilmTracer™ SYPRO Ruby (Invitrogen, Molecular Probes) was used as a fluorescent probe to evaluate the formation of biofilms proteins under the influence of PLB⁶². SYPRO Ruby stain labels most classes of proteins, including glycoproteins, phosphoproteins, lipoproteins, calcium binding proteins, fibrillar proteins, and other proteins that are difficult to stain. The biofilms were stained with a volume of 100 µL SYPRO Ruby per well and plates were incubated at room temperature in the dark for 1 h. Stained biofilms were observed with HCS system. A similar approach to the one described above for PIA was used.

Quantification of eDNA

The eDNA was quantified using the Quant-iT™ PicoGreen® dsDNA Assay Kit to assess the formation of biofilm eDNA under the influence of PLB. PicoGreen® is a fluorescent dye that selectively binds to double-stranded DNA (dsDNA) and exhibits enhanced fluorescence upon binding. Biofilms were cultivated as described in the previous section. After removing planktonic cells, the biofilms were washed twice with PBS. The quantification of eDNA involved adding 100 µL TE buffer followed by 100 µL freshly prepared PicoGreen solution (1 µL PicoGreen dye in 199 µL TE buffer). The wells were mixed and incubated for 2–5 min at room temperature in darkness before measuring fluorescence intensity (excitation 485 nm/emission 535 nm) using a fluorescence microplate reader. Lambda DNA was used to generate a standard curve for each run. Stained biofilms were observed with HCS system. All assays were performed in triplicate.

Statistical analysis

SPSS version 27.0 (SPSS Inc., Chicago, IL, USA) software was used for data quantification and analysis. GraphPad Prism 8.0.2 software was used for mapping. All results were presented as mean ± standard deviation. Analysis of variance (ANOVA) was performed to evaluate significant differences. Statistical significance compared to the untreated control was indicated using double asterisks (**) for $P \leq 0.01$.

Data availability

All raw data are available from the corresponding author upon reasonable request.

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Author contributions

SB and HY conceived the study and designed the experiments. HY and CS performed the experiments and handled data curation. HY and SB acquired the funding. CL conducted the investigation. ML, ZY, RL and DL provided critical feedback and helped shape the research, analysis tools, and manuscript. ZC, ML and TM provided the software for visualization. SB and HY wrote the first draft of the manuscript and all authors revised and commented it. All authors read and approved the final manuscript.

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Declarations

Competing interests

The authors declare no competing interests.

Additional information

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