Roles of Mannosylerythritol Lipid-B Components In Antimicrobial Activity Against Bovine Mastitis-Causing Staphylococcus Aureus

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Roles of mannosylerythritol lipid-B components in antimicrobial activity against bovine mastitis-causing *Staphylococcus aureus*

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Abstract

Mannosylerthritol lipid-B (MEL-B), which comprises ester-bonded hydrophilic ME and hydrophobic fatty acids, is a bio-surfactant with various unique properties, including antimicrobial activity against most gram-positive bacteria. The gram-positive Staphylococcus aureus is a causative pathogen of dairy cattle mastitis, which results in considerable economic loss in dairy industry. Here, we demonstrate the efficacy of MEL-B as a disinfectant against bovine-derived S. aureus and elucidate a mechanism of action of MEL-B in inhibition of bacterial growth. The growth of bovine mastitis causative S. aureus BM1006 was inhibited when cultured with MEL-B above 10 ppm (equivalent to 0.015 mM). The activity of MEL-B required fatty acids (i.e., caprylic and myristoleic acids) as ME, the component of MEL-B lacking fatty acids, did not inhibit the growth of S. aureus even at high concentrations. Importantly, ME-bound fatty acids effectively inhibited the growth of S. aureus when compared with free fatty acids. Specifically, the concentrations of ME-bound fatty acids and free caprylic and myristoleic acids required to inhibit the growth of S. aureus were 0.015, 10, and 1 mM, respectively. The involvement of ME in the antimicrobial activity of MEL-B was confirmed by digestion of MEL-B with lipase, which dissociated ME and fatty acids. These results indicated that a mechanism of action of MEL-B in inhibiting the growth of S. aureus could be explained by the effective transporting of antimicrobial fatty acids to the bacterial surface via hydrophilic ME.

Key words Mannosylerthritol lipid-B (MEL-B), bio-surfactant, S. aureus, mastitis, dairy cattle
Introduction

In the livestock industry, large amounts of antibiotics have been extensively used to cure domestic animals suffering from infectious diseases such as mastitis, diarrhea, and pneumonia (Berendsen et al. 2015; Kromker and Leimbach 2017). Livestock have been also administered various antibiotics as feed additives to gain body weight rapidly; however, most countries recently proposed to discontinue using antibiotics as feed additives because of multiple safety concerns (Maron et al. 2013). The most serious issue caused by excess use of antibiotics is the development of antimicrobial resistant bacteria (AMR) that may then infect human beings (Graham et al. 2009; Lee 2003). The number of human deaths predicted in 2050 due to AMR infection has been estimated to be over 10 million if no actions are taken at the current time (de Kraker et al. 2016). Considering that contagious bacteria can spread globally, the idea of “One Health” that pursues the health of animals, humans, and the environment without the use of excess antibiotics has been widely accepted (Collignon 2015; McEwen and Collignon 2018).

Mastitis is caused primarily by infection with pathogens in the udder of mammals such as dairy cattle (Schrick et al. 2001). Bovine mastitis can be divided into two types based on the classification of causative bacteria. One type is caused environmentally (so-called environmental mastitis) by infection of bacteria that are commonly found in farms, such as environmental streptococci and Escherichia coli. The other type is infectious mastitis, which is mostly caused by Staphylococcus aureus (Hospido and Sonesson 2005). Of note, methicillin-resistant S. aureus (MRSA), a well-known AMR, has been frequently isolated from milk of dairy cattle with infectious mastitis (Vanderhaeghen et al. 2010). S. aureus forms a biofilm to facilitate infection and may survive latently in target cells, such as macrophages, by escaping from the host immune machinery (Scherr et al. 2015; Thurlow et al. 2011). Intracellular S. aureus is believed to account for why dairy cattle with subclinical mastitis cause large economical loss due to the reduction of milk production.
Therefore, the development of novel strategies that can prevent infection by *S. aureus* in dairy cattle without using antibiotics not only contributes to “One health,” especially in suppressing the outbreak of AMRs such as MRSA, but also facilitates reducing the economic loss caused by both clinical and subclinical mastitis (Sinha et al. 2014).

Mannosylerythritol lipid B (MEL-B) is a glycolipid bio-surfactant produced by the cultivation of *Pseudozyma tsukubaensis* (NBRC1940) in the presence of olive oil (Fukuoka et al. 2008). MEL-B is composed of hydrophilic 4-O-β-D-mannopyranosyl-D-erythritol (ME) and two types of hydrophobic fatty acids (e.g., caprylic acid and myristoleic acid in most cases) (Yamamoto et al. 2012). MEL-B possesses several biological characteristics because of the amphiphilic properties and is therefore attracting attention in various fields (Coelho et al. 2020). For instance, MEL-B has been commercialized in the cosmetic industry as a natural ingredient with a moisturizing effect, clearly indicating that there are no safety concerns (Yamamoto et al. 2012). Importantly, the properties of MEL-B include antibacterial activity against most gram-positive bacteria (Kitamoto et al. 1993); however, the mechanism of action is yet to be identified. Furthermore, the inhibitory effect of MEL-B on the growth of *S. aureus* strains derived from dairy cattle has not yet been investigated.

Here, we show that MEL-B inhibits the growth of a bovine mastitis causative *S. aureus* BM1006 and demonstrate that the hydrophilic ME acts as a delivery vehicle to transfer the antimicrobial fatty acids that affect bacterial growth. These results indicate that, as a bio-surfactant, MEL-B could be useful as a possible disinfectant capable of suppressing outbreaks of *S. aureus* in dairy farms.
Materials and Methods

MEL-B, ME, and fatty acids

*Pseudozyma tsukubaensis* (NBRC1940) was cultured in production medium containing 10% (v/v) olive oil, 0.1% (w/v) yeast extract, 4% (w/v) glucose, 0.3% (w/v) NaNO₃, 0.03% (w/v) MgSO₄•7H₂O, and 0.03% (w/v) KH₂PO₄ at 25℃ on a rotary shaker (180 ppm) for 7 days to generate MEL-B, which was composed of ME and fatty acids, mainly caprylic (C8:0) and myristoleic (C14:1) acids. Purification of MEL-B from the culture medium of *P. tsukubaensis* was performed using high-performance liquid chromatography (HPLC) on a silica gel column. A portion of purified MEL-B was reacted with 4-N-chloroformylmethy-N-methyl(amino)-7-nitro2,1,3-benzoxadiazole, a fluorescent reagent with λex = 470 nm and λem = 540 nm, in dry acetone for 60 min to synthesize NBD-labeled MEL-B. ME was synthesized according to the previous study (Fukuoka et al. 2008).

Bacterial strains and culture

*S. aureus* BM1006 (MAFF913131) and *E. coli* JM109 (ATCC53323) were used in this study. Both bacteria were cultured overnight in Trypto-Soya (TS) broth (Nissui) at 37℃ on a rotary shaker (120 ppm). Pre-cultured bacteria (30 μL) were added to fresh TS broth (3 mL) containing MEL-B at various concentrations (0, 0.1, 1, 10, 100, and 1000 ppm) to address the effect of MEL-B on bacterial growth for 24 h at 37℃. Additionally, *S. aureus* was cultured in TS broth containing ME at various concentrations (0.1, 1, 10, 100, and 1000 ppm). *S. aureus* was also cultured for 5 h in the presence of either caprylic acid (1 and 10 mM) or myristoleic acid (0.1 and 1 mM). Survival of bacteria in the presence of either MEL-B, ME, caprylic, or myristoleic acid was determined by obtaining the numbers of colony-forming units (CFUs). Aliquots of bacterial broth were collected and seeded on TS agar plates (three replicates) after dilution with saline to obtain CFUs.
Binding analysis

*S. aureus* BM1006 and *E. coli* JM109 were treated with 10 ppm of NBD-labeled MEL-B (NBD-MEL-B) for 30 min at room temperature (RT). After washing three times with phosphate-buffered saline, bacteria were fixed in 4% (w/v) of paraformaldehyde (Nacalai Tesque) for 30 min at RT and affixed on glass slides for fluorescence microscopy (BZ-9000, Keyence). Flow cytometry analysis (Accuri C6, DB) was also used to assess the mean fluorescence intensity of bacteria.

Scanning electron microscopy analysis

The effect of MEL-B on the surface structure of *S. aureus* BM1006 was investigated by scanning electron microscopy (SEM) analysis. Specifically, bacteria were cultured for 0.5, 2, 5, and 24 h with or without 100 ppm of MEL-B or were treated with 100 U/mL of lysostaphin (WAKO) for 30 min at RT. Bacteria were fixed in 2.5% (v/v) of glutaraldehyde (Nacalai Tesque) for 1 h at 4℃. After washing, bacteria were affixed to glass slides, coated with platinum and palladium, and analyzed using an SEM SU8000 (Hitachi) at 3.0 kV.

HPLC analysis

To quantify MEL-B contained in TS broth during the culture of *S. aureus*, HPLC was performed. Culture medium (250 μL) collected after 0.5, 2, 5, and 24 h of culturing was mixed with ethyl acetate (500 μL) for 3 min, and the supernatant was collected by centrifugation (13,000 rpm, 4℃, 3 min). The process was repeated three times, and MEL-B was extracted by evaporating from the supernatant. Specifically, the quantification of the extracellular glycolipids was performed using the HPLC of the MEL-B extracts loaded onto a silica gel column (Inertsil SIL 100 A 5 μm, 4.6 × 250 mm; GL Science, Japan) using chloroform/methanol as the solvent system with a gradient flow (1 mL/min) controlled from 100:0 to 0:100. A low-temperature evaporate light-scattering
Detector (Shimadzu, Kyoto, Japan) was used for detecting MEL-B.

**Structural analysis of MEL-B on the antimicrobial activity**

MEL-B (1000 ppm) was treated with porcine pancreas-originated lipase (Sigma) at various concentrations (0.2, 2, and 20 mg/mL) for 1 h at 37°C at 1:1 ratio to cleave the ester bonds between ME and fatty acids. *S. aureus* was cultured with the reactant including MEL-B equivalent to 10 ppm (=0.015 mM) and also with a mixture of ME, caprylic acid, and myristoleic acid (0.015 mM each) to compare the efficacy between the mixture and intact MEL-B (0.015 mM). After 5 h, the survival of bacteria was determined by obtaining the CFU count.

**Statistics**

Statistical analyses were conducted using one-way analysis of variance (ANOVA) with the Kruskal–Wallis test and two-way ANOVA with Tukey’s multiple comparison test using Prism 7 (GraphPad).
Results

MEL-B inhibits the growth of a bovine mastitis causative *S. aureus* strain BM1006

We first sought to demonstrate the efficacy of MEL-B as a possible disinfectant for use with dairy cattle in their breeding environment. Bovine milk-derived *S. aureus* BM1006, a bovine mastitis causative bacterium (Kiku et al. 2016), was cultured *in vitro* in the presence of MEL-B at 0, 0.1, 1, 10, 100, and 1000 ppm, and the influence of MEL-B on the growth of *S. aureus* (via CFU counts) was monitored at 0.5, 2, 5, and 24 h. The growth of *S. aureus* was significantly delayed when cultured with MEL-B, and the minimum concentration of MEL-B required for complete inhibition of bacterial growth was 10 ppm (Fig. 1). Consistent with a previous study that demonstrated the antimicrobial activity of MEL-B against most gram-positive (but not gram-negative) bacteria (Kitamoto et al. 1993), the growth of *E. coli* JM109, a representative gram-negative bacterium, was not significantly affected in the presence of MEL-B even at high concentrations (Fig. 2). These results support previous findings demonstrating the efficacy of MEL-B in inhibiting the growth of *S. aureus* (not *E. coli*) using bovine-derived *S. aureus*.

MEL-B is stable during *in vitro* culture

Although MEL-B possessed antimicrobial activity against *S. aureus*, the growth of *S. aureus* gradually continued during culture even in the presence of MEL-B at high concentrations. Therefore, we next addressed the condition of MEL-B during culture. Normal phase HPLC analysis showed that two peaks were present, one of which corresponded to intact MEL-B, whereas the other eluted at an earlier retention time corresponded unknown molecule(s) excluded by HPLC separation because of the low polarity (Fig. 3a). Importantly, the retention time of MEL-B did not change during culture with *S. aureus*, indicating that MEL-B is intact in our culture condition. In contrast, the first peak with a shorter retention time was slightly but obviously seen even in water (Fig. 3a).
Furthermore, the amount of first peak increased during culture with *S. aureus*, resulting in a decrease in the proportion of MEL-B (Fig. 3b). These results suggest that hydrophobic molecule(s) produced by *S. aureus* may affect the antimicrobial effect of MEL-B *in vitro*.

**MEL-B associates with the cell surface of both *S. aureus* and *E. coli***

To understand the molecular mechanism whereby MEL-B inhibits the growth of *S. aureus*, we labeled MEL-B with NBD and used this in a binding assay to confirm the association between MEL-B and *S. aureus*. The incubation of *S. aureus* with 10 ppm of NBD-labeled MEL-B (NBD-MEL-B) made the bacterial surface clearly fluorescent under a microscope (Fig. 4a). Interestingly, *E. coli* JM109 treated with the same concentration of NBD-MEL-B displayed weak (but apparent) fluorescence despite the lack of inhibitory effect of MEL-B on the growth of *E. coli* (Fig. 4a). No fluorescence was seen in either untreated *S. aureus* or *E. coli* (Fig. 4a). We confirmed this result using flow cytometry analysis with *S. aureus* and *E. coli*, both of which were treated with NBD-MEL-B at various concentrations (0, 0.1, 1, 10, and 100 ppm). The analysis indicated that NBD-MEL-B was associated with both *S. aureus* and *E. coli* in a dose-dependent manner (Fig. 4b). Furthermore, the fluorescence intensities observed in *S. aureus* were significantly higher than those in *E. coli* at NBD-MEL-B concentrations of 1 ppm or higher (Fig. 4c). These results suggest that the actions mediated by the binding of MEL-B to *S. aureus* (rather than simply the act of binding) may be necessary to inhibit bacterial growth.

**MEL-B does not damage the structure of *S. aureus***

We then used SEM to chronologically examine the influence of MEL-B treatment on the structure of *S. aureus* during culture with MEL-B at relatively high concentration (100 ppm). Although MEL-B was anticipated to affect the surface structure of *S. aureus* by binding, no apparent
differences could be seen in *S. aureus* during culture for 24 h with MEL-B when compared with *S. aureus* cultured without MEL-B (Fig. 5a). The typical shape of a grape-like cluster of *S. aureus* was observed regardless of MEL-B treatment (Fig. 5a). Moreover, these structures were completely different from those of damaged *S. aureus* treated with lysostaphin, which specifically disrupts the cell wall of *S. aureus* (Fig. 5b) (Watanakunakorn et al. 1971). These results suggest that the effect of MEL-B on the inhibition of the growth of *S. aureus* may not be due to the disruption of the overall bacterial structure.

**Fatty acids, components of MEL-B, are involved in inhibiting the growth of *S. aureus***

Since MEL-B is composed of mannopyranose-erythritol (ME), which has a hydrophilic property, and two different fatty acids [i.e., caprylic acid (C8:0) and myristoleic acid (C14:1)], which have hydrophobic properties (Yamamoto et al. 2012), we then identified the critical components of MEL-B that affect the growth of *S. aureus*. Supplementation of ME, obtained as an intermediate product in MEL-B synthesis, to *S. aureus* culture at various concentrations (0, 0.1, 1, 10, 100, and 1000 ppm) did not have any inhibitory effects, even at high concentrations (Fig. 6). Unfortunately, the binding capability of ME to the cell surface of *S. aureus* could not be investigated because of the technical difficulty of synthesizing fluorescence-conjugated ME; however, this result suggested that the hydrophobic fatty acid components of MEL-B may directly or indirectly inhibit the growth of *S. aureus*.

**ME acts as a delivery vehicle for caprylic acid and myristoleic acid to *S. aureus***

To elucidate the mechanism of action of MEL-B on the growth inhibition of *S. aureus*, three related experiments were conducted to clarify the functions of ME and fatty acids. Both caprylic and myristoleic acids are known to possess antimicrobial activity against *S. aureus* (Kabara,
1984). Therefore, we determined the minimum concentration of caprylic and myristoleic acids required to inhibit the growth of \textit{S. aureus}. Since the efficacy of MEL-B in inhibiting the growth of \textit{S. aureus} was clearly observed 5 h after culture (as shown in Fig. 1), \textit{S. aureus} was cultured for 5 h in the presence of either caprylic acid (1 or 10 mM) or myristoleic acid (0.1 or 1 mM). There was a significant decrease in CFU when 10 mM of caprylic acid or 1 mM of myristoleic acid was added to the culture (Fig. 7a). We then compared the efficacy of MEL-B and a mixture of ME, caprylic acid, and myristoleic acid at the same concentration. The molecular weight of MEL-B composed of ME, caprylic acid, and myristoleic acid is 634 Da, and therefore, we used 10 ppm (the minimum dose of MEL-B required to inhibit the growth of \textit{S. aureus}), which is equivalent to 0.015 mM. However, a decrease in \textit{S. aureus} CFU occurred when MEL-B at 0.015 mM (but not the mixture of ME, caprylic acid, and myristoleic acid at 0.015 mM each) (Fig. 7b). We then demonstrated the reduction of MEL-B efficacy by digestion of the ester bond that connects the ME with the two fatty acids. Specifically, 1000 ppm of MEL that had been pretreated with lipase, which digests the ester bond, at various concentrations (0.2, 2, and 20 mg/mL) and 10 ppm (=0.015 mM) equivalent of MEL-B was added in the culture medium of \textit{S. aureus}. Importantly, MEL-B did not inhibit the growth of \textit{S. aureus} when pretreated with the high concentration of lipase (Fig. 7c); the lipase itself did not affect the growth of \textit{S. aureus} (Fig. 7c). These results indicated that the efficacy of MEL-B could be explained by the effective association of the antimicrobial caprylic and myristoleic acid components with the surface of \textit{S. aureus} via the hydrophilic ME moiety.
Discussion

Our study aimed to evaluate MEL-B as a possible disinfectant for use in the livestock industry to control the spread of *S. aureus* strains, including MRSA. We demonstrated that the growth of *S. aureus* BM1006, which belongs to the sequence type 352 that often causes mastitis in dairy cattle (Hata et al. 2010), is significantly inhibited when cultured in the presence of MEL-B in vitro. Notably, the pathogenicity of *S. aureus* BM1006 has been confirmed by in vivo infectious studies where mastitis was induced in dairy cattle when *S. aureus* BM1006 was inoculated into the udder (Nagasawa et al. 2018). A previous study has also shown that inflammatory chemokines (e.g., IL-8, CXCL6, and CCL2), which are involved in the recruitment of polymorphonuclear leukocytes into the udder, are abundantly secreted by mammary epithelial cells when stimulated with *S. aureus* BM1006 (Kiku et al. 2016). Given the virulence of *S. aureus* BM1006 as a causative bacteria of bovine mastitis, our results can provide insight into the development of a novel disinfectant to prevent the onset of mastitis in dairy cattle.

MEL-B had no effect on the growth of a representative gram-negative bacteria, *E. coli* JM109, in this study, which is consistent with previous results (Kitamoto et al. 1993). Our results also demonstrated that MEL-B bound to the cell surface of *E. coli* despite a lack of antimicrobial activity against the bacteria. Therefore, we hypothesized that further actions following the binding of MEL-B to the bacterial cell surface are required to inhibit the growth of gram-positive *S. aureus* (but not gram-negative *E. coli*). Recently, atomic force microscopy was used to demonstrate that the diameter of peptidoglycan pores can be up to 60 nm (Pasquina-Lemonche et al. 2020). Hence, MEL-B may diffuse freely into the cross-linked cell wall structure of gram-positive bacteria by penetrating the peptidoglycan interspaces and then interfering the cell membrane, where functional molecules (e.g., transporters and receptors) are involved in nutrient uptake and signal transduction are expressed. In contrast, lipopolysaccharide found in the outer membrane of gram-negative bacteria...
may interfere with the free diffusion of MEL-B, leaving the cell membrane intact in these bacteria even after binding.

One of the significant advances obtained in this study was to elucidate how MEL-B inhibits the growth of \textit{S. aureus}. Specifically, the ME moiety plays an important role in effectively exerting antimicrobial activity of the caprylic and myristoleic acids against \textit{S. aureus}. Nevertheless, \textit{S. aureus} still grew for 24 h when cultured in the presence of MEL-B at high concentration. A key structural feature of MEL-B is the ester bonding that connects ME and the two fatty acids, which is important for exhibiting antimicrobial activity. Lipase produced by \textit{S. aureus} during the culture was initially suspected as a factor that could reduce the antimicrobial activity of MEL-B. In this regard, pre-treatment of MEL-B with commercially available lipase from porcine pancreases reduced the antibacterial activity of MEL-B against \textit{S. aureus}. However, the concentration of commercial lipase required to inhibit MEL-B activity was extremely high compared with that produced normally from \textit{S. aureus} (Flanagan et al. 2016; Hu et al. 2012). Another possible hypothesis is that hydrophobic molecule(s) may affect the antimicrobial activity of MEL-B. HPLC analysis showed two major peaks corresponding to the MEL-B and unknown molecules(s) excluded by HPLC separation due to the low polarity (= high hydrophobicity). The peak with a shorter retention time was found even in water, and the amount increased during culture with \textit{S. aureus}. These results suggest that the reduced antimicrobial activity of MEL-B against \textit{S. aureus} during culture may be due to the effect of \textit{S. aureus}-derived hydrophobic molecule(s) (but not impurities) in our culture condition. Therefore, further studies are needed to identify appropriate solvent conditions that maintain or increase the efficacy of MEL-B for practical application as a disinfectant for dairy cattle.

A recent study demonstrated that the apoptosis of \textit{S. aureus} was induced by treatment with MEL-A (Shu et al. 2020). By contrast, in our study using MEL-B, there was no sign to demonstrate apparent cell death, such as membrane disruption, although the growth of \textit{S. aureus} was successfully
inhibited by the treatment of MEL-B. This discrepancy may be due to the structural difference between MEL-A and MEL-B. MELs are divided into four types (i.e., MEL-A, MEL-B, MEL-C, and MEL-D), based on the number and position of the acetyl group (Saika et al. 2018), which depends on the microorganism responsible for synthesis (Coelho et al. 2020). One of the notable characteristics of MEL-A, which differentiates this from MEL-B, MEL-C, or MEL-D, is that MEL-A possesses an additional acetyl group, resulting in a higher hydrophobicity compared with that of the others (Imura et al. 2007; Saika et al. 2018). Since we did not compare the antimicrobial activity between MEL-A and MEL-B, further studies should be conducted to determine which form of MEL is most appropriate in controlling the spread of *S. aureus*. Nevertheless, it should be emphasized that MEL-B has been extensively used in the cosmetic industry, where it is used to suppress skin perspiration and increase water content (Yamamoto et al. 2012). Although verification is required to determine the use of MEL-B as a disinfectant for controlling mastitis caused by *S. aureus*, the safe use of MEL-B as a moisturizer for skin care should be positively appreciated for potential application in the dairy industry.

In conclusion, we elucidated the mechanism of action of MEL-B, composed of ME and antimicrobial fatty acids, in inhibiting the growth of bovine-derived mastitis causative *S. aureus*. ME acts as a delivery vehicle for antimicrobial fatty acids. Therefore, the binding between ME and fatty acids via ester bond is essential for the effective transport of antimicrobial fatty acids that then associate with the surface of *S. aureus*. These results provide a novel insight into developing a disinfectant that could be used in farms to prevent dairy cattle from contracting mastitis.

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**Author contribution** ShiY and TN conceived and designed the research and wrote the manuscript. ShiY and MF performed experiments and analyzed the data. EH provided *S. aureus* BM1006 and supports experiments. ShuY, MK, AK, TS, and AS synthesized MEL-B. KW, HY, and HA support experiments.

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**Data availability** All data generated or analyzed during this study are included in this manuscript.

**Declarations**

**Conflict of interest** The authors declare no competing interests.

**Ethical approval** This study does not contain any experiments with human participants or animals.
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Figure legends

**Fig. 1** MEL-B inhibits the growth of *Staphylococcus aureus* in vitro. Bovine-derived *S. aureus* BM1006 was cultured in the presence of MEL-B at 0, 0.1, 1, 10, 100, or 1000 ppm for 0.5, 2, 5, and 24 h. The effect of MEL-B on the growth of *S. aureus* was assessed by counting colony-forming units (CFUs). Growth was significantly delayed when cultured with MEL-B above 10 ppm. Experiments were performed in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using two-way ANOVA with Tukey’s multiple comparison test. ****p < 0.0001

**Fig. 2** MEL-B does not inhibit the growth of *Escherichia coli* in vitro. *E. coli* JM109, a representative of gram-negative bacteria, was cultured in the presence of MEL-B at 0, 0.1, 1, 10, 100, or 1000 ppm for 0.5, 2, 5, and 24 h. The effect of MEL-B on the growth of *E. coli* was assessed by counting colony-forming units (CFUs). MEL-B did not affect the growth of *E. coli* even at high concentrations. Experiments were performed in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using two-way ANOVA with Tukey’s multiple comparison test.

**Fig. 3** MEL-B is stable during culture in TS broth. a *S. aureus* BM1006 was cultured in the presence of MEL-B at 1000 ppm for 0.5, 2, 5, and 24 h. HPLC retention times of MEL-B in culture media were compared with those of fresh MEL-B undiluted or diluted with water or TS broth. In addition to the typical peak corresponding to MEL-B, another peak with a shorter retention time was detected when MEL-B was dissolved in water or TS broth, and the ratio of peak with a shorter retention time increased during culturing. b The proportion of the original peak corresponding to intact MEL-B was calculated. Statistical analyses were conducted using one-way ANOVA with the Kruskal–Wallis test.
Fig. 4 MEL-B binds to the surface of both *S. aureus* BM1006 and *E. coli* JM109 with different binding affinities. **a** Fluorescence signals were detected from *S. aureus* and *E. coli* when bacteria were incubated for 30 min *in vitro* with MEL-B that was pre-conjugated with a fluorescence molecule NBD (NBD-MEL-B). No fluorescence signals were seen from untreated *S. aureus* or *E. coli*. **b** Flow cytometric analysis at 0, 0.1, 1, 10, or 100 ppm of MEL-B showed that NBD-MEL-B associated with *S. aureus* and *E. coli* in a dose-dependent manner. **c** Mean fluorescence intensity (MFI) obtained from *S. aureus* was higher than that obtained from *E. coli* when bacteria were individually incubated with NBD-MEL-B above 1 ppm. Experiments were performed in triplicate. The representative data were shown in histograms and all MFIs obtained from three experiments were summarized in a dot graph. Statistical analyses were conducted using *t*-test. ***p* < 0.001, ****p* < 0.0001

Fig. 5 MEL-B does not disrupt the surface structure of *S. aureus* BM1006. **a** Scanning electron microscopic (SEM) analysis showed that the morphology of *S. aureus* did not change through *in vitro* culture in the presence of a relatively high concentration (100 ppm) of MEL-B. **b** *S. aureus* was treated with 100 U/mL of lysostaphin to confirm the disruption of the bacterial structure as a control in the SEM analysis. Scale bar = 1 μm

Fig. 6 ME, a component of MEL-B, does not inhibit the growth of *S. aureus* BM1006 *in vitro*. *S. aureus* was cultured in the presence of ME at 0, 0.1, 1, 10, 100, or 1000 ppm for 0.5, 2, 5, and 24 h. The effect of ME on the growth of *S. aureus* was assessed by counting colony-forming units (CFUs). ME did not affect the growth of *S. aureus* even at high concentrations. Experiments were performed
in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using two-way ANOVA with Tukey’s multiple comparison test.

**Fig. 7** Modification of ME with two fatty acids is essential for effectively inhibiting the growth of *S. aureus* BM1006. **a** Caprylic acid (C8:0) and myristoleic acid (C14:1), both of which associate with ME, showed clear antimicrobial activity against *S. aureus* when cultured for 5 h at 10 and 1 mM, respectively. **b** A mixture of ME with caprylic and myristoleic acid, which were adjusted to the same concentration as MEL-B (10 ppm = 0.015 mM), did not inhibit the growth of *S. aureus*. **c** Pretreatment of MEL-B with a high concentration of lipase to digest the ester bond between the two fatty acids and ME reduced the antibacterial activity against *S. aureus*. Experiments were performed in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using one-way ANOVA with the Kruskal–Wallis test. **p < 0.01, ***p < 0.001, ****p < 0.0001
MEL-B inhibits the growth of Staphylococcus aureus in vitro. Bovine-derived S. aureus BM1006 was cultured in the presence of MEL-B at 0, 0.1, 1, 10, 100, or 1000 ppm for 0.5, 2, 5, and 24 h. The effect of MEL-B on the growth of S. aureus was assessed by counting colony-forming units (CFUs). Growth was significantly delayed when cultured with MEL-B above 10 ppm. Experiments were performed in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using two-way ANOVA with Tukey’s multiple comparison test. ****p < 0.0001
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culturing. The proportion of the original peak corresponding to intact MEL-B was calculated. Statistical analyses were conducted using one-way ANOVA with the Kruskal–Wallis test. *p < 0.05, **p < 0.01

Figure 4

MEL-B binds to the surface of both S. aureus BM1006 and E. coli JM109 with different binding affinities. a Fluorescence signals were detected from S. aureus and E. coli when bacteria were incubated for 30 min in vitro with MEL-B that was pre-conjugated with a fluorescence molecule NBD (NBD-MEL-B). No fluorescence signals were seen from untreated S. aureus or E. coli. b Flow cytometric analysis at 0, 0.1, 1, 10, or 100 ppm of MEL-B showed that NBD-MEL-B associated with S. aureus and E. coli in a dose-dependent manner. c Mean fluorescence intensity (MFI) obtained from S. aureus was higher than that obtained from E. coli when bacteria were individually incubated with NBD-MEL-B above 1 ppm. Experiments were performed in triplicate. The representative data were shown in histograms and all MFIs obtained from three experiments were summarized in a dot graph. Statistical analyses were conducted using t-test. ***p < 0.001, ****p < 0.0001
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ME, a component of MEL-B, does not inhibit the growth of S. aureus BM1006 in vitro. S. aureus was cultured in the presence of ME at 0, 0.1, 1, 10, 100, or 1000 ppm for 0.5, 2, 5, and 24 h. The effect of ME on the growth of S. aureus was assessed by counting colony-forming units (CFUs). ME did not affect the growth of S. aureus even at high concentrations. Experiments were performed in triplicate, and the mean ± standard error of the mean obtained from representative data is shown. Statistical analyses were conducted using two-way ANOVA with Tukey's multiple comparison test.
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