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Natural Berberine-Hybridized Benzimidazoles as Novel Unique Bactericides against Staphylococcus aureus

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ABSTRACT: Natural berberine-hybridized benzimidazoles as potential antibacterial agents were constructed to treat Staphylococcus aureus infection in the livestock industry. Bioassay showed that some new berberine-benzimidazole hybrids exhibited potent antibacterial efficacies, especially, the 2,4-dichlorobenzyl derivative 7d not only showed strong activity against S. aureus ATCC 29213 with the MIC value of 0.006 mM but also effectively eradicated bacterial biofilm and exhibited low toxicity toward mammalian cells. The drug combination experiments showed that compound 7d together with norfloxacin could enhance the antibacterial efficacy. Moreover, the 2,4-dichlorobenzyl derivative 7d did not show obvious propensity to develop bacterial resistance. Preliminary mechanism studies revealed that the active molecule 7d could damage the membrane integrity, stimulate ROS generation, and bind with DNA as well as S. aureus sortase A, thus exerting powerful antibacterial ability. In light of these facts, berberine-benzimidazole hybrid 7d showed a large potentiality as a new bactericide for treating S. aureus in the livestock industry.

KEYWORDS: berberine, benzimidazole, antibacterial, resistance, Staphylococcus aureus

■ INTRODUCTION

Antibiotics play a crucial role in ensuring the safe supply of food-producing animal in sustainable animal production systems.^{1,2} However, extensive and uncontrolled exposure of various antibiotics has caused an unprecedented risk from bacterial resistance.^{3,4} In particular, Staphylococcus aureus has shown resistance to most currently approved antibiotics, such as penicillin and tetracycline.^{5,6} Therefore, it is imperative to exploit new effective and innovative antibacterial agents to defend the invasion caused by drug-resistant Staphylococcus aureus strains

Natural products were significant resources of alternative classes of antibiotics with new antibacterial scaffolds.⁷ Berberine, a natural isoquinoline alkaloid isolated from several medicinal herbs such as Berberis vulgaris and Rhizoma coptidis, has been used as a nonprescription medicine to treat infectious diseases (such as bacillary dysentery, acute gastroenteritis, and cholera) in China for 2000 years.⁸⁻¹⁰ Currently, various clinical applications of berberine have been discovered, especially in antibacterial usage, which revealed that berberine with a quaternary nitrogen, polycyclic, and planar system could helpfully increase membrane permeability and strengthen the bind affinities with amino acids in biomolecules.¹¹⁻¹⁵ Therefore, the unique and favorable structural features endowed natural berberine superiorities to be used as a novel promising scaffold for further antibacterial drug developments.

The successful applications of antibacterial azoles greatly hindered the invasion of bacteria in humans, terrestrial and aquatic animals, as well as plants.¹⁶⁻¹⁹ In particular, benzimidazoles have attracted considerable attention because of the purine-like structure in benzimidazole ring that enabled benzimidazole derivatives to block the biosynthesis of proteins and nucleic acids.^{20,21} A variety of bactericides containing benzimidazole fragment have been launched into the market, such as carbendazim and thiabendazole,²² demonstrating the enormous potentiality in the antibacterial aspect.

SUPPORTING Information

Herein, by incorporating benzimidazoles into the C-12 position of berberine backbone, a series of berberinehybridized benzimidazoles were constructed in this work, which were expected to overcome drug resistance (Figure 1). Moreover, a variety of substituents including chlorine and bromine atoms, methyl and methoxy groups, aliphatic chains, halogen-containing phenyl moieties, as well as amide and ester groups were introduced to adjust the physicochemical properties and to investigate the effect on bioactivities. Antibacterial evaluation, bactericidal kinetic assay, drug combination study, resistance development evaluation, and cytotoxicity assay were conducted to verify the antibacterial potentiality of new berberine-hybridized benzimidazoles. Finally, the preliminary explorations of antibacterial mechanism were performed by experiments including membrane depolarization and permeabilization assays, reactive oxygen species (ROS) generation, glutathione (GSH) activity determination, docking study, and interaction with DNA.

MATERIALS AND METHODS

Instruments and Chemicals. NMR spectra were recorded on Bruker AVANCE III 10 600 spectrometers (600 and 400 MHz). High-resolution mass spectra (HRMS) were recorded on a Bruker

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Figure 1. Design concept of novel berberine-hybridized benzimidazoles as new potential bactericides.

impact II 10200 spectrometer. Melting points of new compounds were measured by an X-6 melting point apparatus (Beijing Focus Instrument Co., Ltd., China). A TU-2450 spectrophotometer (Puxi Analytic Instrument Ltd., Beijing, China) and an F-7000 spectrofluorometer (5J1-0004 model, Hitachi, Tokyo, Japan) were utilized to record UV spectra and fluorescence spectra, respectively. The tested bacterial strains were provided by Sichuan Provincial People's Hospital, Chengdu, China. All reagents and solvents were commercially available.

Chemistry. Synthesis of Berberrubine 1. Compound 1 was synthesized according to the reported procedure.²³

Synthesis of Compound 2. The mixture of berberrubine 1 (1.10 mmol) and hexamethylenetetramine (5.50 mmol) in trifluoroacetic acid (10 mL) was refluxed for 8 h. The reaction system was cooled to room temperature, and then the diluted hydrochloric acid (10%, 5 mL) was added with stirring for 20 min. A saturated solution of sodium hydrogen carbonate was added to neutralize the mixture, and the solid was precipitated. The resulting precipitation was purified by column chromatography on silica gel using dichloromethane/ methanol (V/V, 15/1, Rf = 0.5) as an eluent to produce target compound 2 with a yield of 55.4%. The corresponding data are provided in the Supporting Information.

Synthesis of Compounds 3a-e. The mixture of intermediate 2 (0.65 mmol), corresponding *o*-phenylenediamines (1.30 mmol), and sodium pyrosulfite (1.30 mmol) in *N*,*N*-dimethyl formamide (10 mL) was heated at 110 °C for 6 h. After the reaction was completed, water (20 mL) was added and the solid was precipitated. The residue was dissolved in chloroform (30 mL) and washed with saturated sodium chloride solution (3 × 30 mL). The organic phase was dried over anhydrous sodium sulfate and then evaporated. Crude product was purified by silica gel column chromatography using dichloromethane/ methanol (V/V, 15/1, Rf = 0.2–0.4) as an eluent to give pure target compounds 3a-e with yields of 37.6–80.2%. The corresponding data are provided in the Supporting Information.

Synthesis of Compounds 4a-f, 5a-d, 6a-c, 7a-e, 8a-b, 9a-b, and 10. Berberine-hybridized benzimidazoles 4-10 were prepared according to the procedure described for 3a using intermediate 2 (0.65 mmol) and the corresponding mono-substituted *o*-phenylenediamines 11-17 (1.30 mmol) as starting materials in the presence of sodium pyrosulfite (1.30 mmol). Pure target compounds 4-10 were obtained with yields from 20.2 to 87.5%. The corresponding data are provided in the Supporting Information.

Synthesis of Intermediates 11-17. Mono-substituted *o*-phenylenediamine derivatives 11-17 were synthesized according to the reported procedure.²⁴

Antibacterial Activity Assay. The prepared intermediate 2, berberine-hybridized benzimidazoles 3-10, berberine, and norfloxacin determined the in vitro antibacterial activities against Grampositive bacteria (methicillin-resistant *S. aureus* N315, *S. aureus*, *S. aureus* ATCC 29213) and Gram-negative bacteria (*Pseudomonas aeruginosa, Klebsiella pneumonia, Acinetobacter baumannii, Escherichia coli*), and detailed procedure is given in the Supporting Information.

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Biofilm Disruption Evaluation. The biofilm disruption activity of active compound 7d against *S. aureus* ATCC 29213 was measured using crystal violet staining assay,²⁵ and the detailed procedure is provided in the Supporting Information.

Metabolic Activity Evaluation. The preformed *S. aureus* ATCC 29213 biofilm was incubated with increasing concentration of compound 7d for 24 h at 37 °C. Resazurin solution (5 μ g/mL, 25 μ L) was added into control and treated cells to stain the biofilm for 40 min at 37 °C, and the absorbance was measured at 571 nm.

Bactericidal Activity Evaluation. An overnight culture of *S. aureus* ATCC 29213 was diluted in a Mueller-Hinton broth medium and incubated at 37 °C with aeration at 225 rpm for 2 h (early exponential) or 5 h (late exponential). The bacteria were then treated with compound 7d at $6 \times$ MIC and $8 \times$ MIC in culture tubes at 37 °C and 225 rpm. At different intervals, the bacteria solution (100 μ L) was moved to a 96-well plate, centrifuged at 4000 rpm for 5 min, and resuspended in sterile phosphate-buffered saline (100 μ L). Tenfold serially diluted suspensions were plated on Mueller-Hinton agar plates and incubated at 37 °C overnight. Colonies were counted, and CFU per mL was calculated.

Cytotoxicity Evaluation. MTT assay was conducted to evaluate the cytotoxicity of active compound 7d and berberine against human embryonic kidney (Hek) 293T and human fibroblast (HFC) normal cells, ¹⁴ and the hemolysis was also evaluated. The detailed procedure is depicted in the Supporting Information.

Drug Combination. The drug combination study of compound 7d and norfloxacin was determined by the checkerboard titration method,²⁶ and the detailed procedure is provided in the Supporting Information.

Propensity to Develop Bacterial Resistance. The propensity of compound 7d, norfloxacin, and drug combination to induce resistance in *S. aureus* ATCC 29213 was evaluated, and the detailed procedure is depicted in the Supporting Information.

Membrane Depolarization Assay. The membrane depolarization ability of compound 7d against *S. aureus* ATCC 29213 was tested using 3,3'-dipropylthiadicarbocyanine iodide (diSC35) as fluorescence probe, and the detailed procedure is provided in the Supporting Information.

Membrane Permeabilization Assay. The membrane permeabilization ability of compound 7d against *S. aureus* ATCC 29213 was tested using propidium iodide (PI) as fluorescence probe, and the detailed procedure is provided in the Supporting Information.

Scheme 1. Synthetic Route for Target Berberine-Hybridized Benzimidazoles 3-6



Scheme 2. Synthetic Route for Target Berberine-Hybridized Benzimidazoles 7-10



Molecular Simulation. The docking investigation between compound 7d and *S. aureus* sortase A (PDB code: 1T2W) was conducted by Sybyl-x 2.0.

Supramolecular Interaction of Active Molecule 7d and Calf Thymus DNA. The interaction mode of active compound 7d with DNA was explored through UV-vis and fluorescence spectra using acridine orange (AO) as spectral probe, and the detailed procedure is provided in the Supporting Information.

Determination of the Intracellular Reactive Oxygen Species (**ROS**). The intracellular ROS of *S. aureus* ATCC 29213 was measured using the standard 2,7-dichlorofluorosceindiacetate (DCFH-DA) assay,²⁷ and the detailed procedure is provided in the Supporting Information.

Determination of Intracellular Glutathione (GSH) Activity. The activity of intracellular GSH in *S. aureus* ATCC 29213 was determined by the standard Ellman's assay,²⁷ and the detailed procedure is provided in the Supporting Information.

RESULTS AND DISCUSSION

Chemistry. Synthetic routes to prepare berberine-hybridized benzimidazoles are outlined in Schemes 1, 2, and S1. Commercially available berberine underwent demethylation at 190 °C under vacuum to generate berberrubine 1, which was further reacted with hexamethylenetetramine in trifluoroacetic acid to afford berberrubine-12-carbaldehyde 2 in 55.4% yield. The cyclization reaction of formyl derivative 2 and the corresponding *o*-phenylenediamines produced target berberine-hybridized benzimidazoles 3a-e in the yields of 37.6-80.2%.

In a similar method to that described for N-1-alkylated benzimidazoles, the synthesis of compound 4a was initially attempted by direct nucleophilic substitution of compound 3a with bromoethane using potassium carbonate as base and tetrabutylammonium bromide as a phase transfer catalyst. However, no desired compound was obtained. Variations of

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Table 1	. Antibacterial	Activities	of Intermediate	2, Ber	berine-Hybridized	Benzimidazoles	3-10, and	l Reference	Drugs	(MIC,
mM)										

		Gram-positiv	e bacteria	Gram-negative bacteria					
compds	MRSA N315	S. aureus	S. aureus ATCC 29213	Klebsiella pneumoniae	E. coli	Pseudomonas aeruginosa	A. baumannii		
2	0.166	0.166	0.664	0.332	0.664	0.166	0.664		
3a	0.068	0.135	0.068	0.540	0.270	0.270	0.270		
3b	0.031	0.252	0.504	0.252	0.504	0.252	0.252		
3c	0.247	0.247	0.247	0.495	0.247	0.124	0.247		
3d	0.033	0.131	0.016	0.131	0.262	0.131	0.033		
3e	0.254	0.254	0.127	0.508	0.254	0.127	0.254		
4a	0.016	0.127	0.008	0.255	0.510	0.127	0.127		
4b	0.121	0.121	0.015	0.241	0.483	0.241	0.121		
4c	0.059	0.059	0.029	0.235	0.471	0.235	0.118		
4d	0.115	0.007	0.229	0.029	0.029	0.014	0.029		
4e	0.218	0.218	0.055	0.437	0.437	0.218	0.218		
4f	0.208	0.104	0.208	0.208	0.417	0.104	0.104		
5a	0.242	0.015	0.030	0.121	0.015	0.004	0.015		
5b	0.225	0.028	0.056	0.112	0.112	0.056	0.056		
5c	0.249	0.031	0.498	0.249	0.249	0.249	0.249		
5d	0.250	0.031	0.250	0.250	0.250	0.250	0.250		
6a	0.247	0.247	0.124	0.494	0.494	0.124	0.247		
6b	0.470	0.470	0.235	0.235	0.470	0.235	0.470		
6c	0.218	0.218	0.436	0.436	0.436	0.218	0.436		
7a	0.014	0.027	0.055	0.110	0.220	0.055	0.055		
7b	0.107	0.053	0.007	0.107	0.107	0.107	0.107		
7 c	0.214	0.003	0.214	0.107	0.428	0.107	0.027		
7 d	0.006	0.013	0.006	0.101	0.202	0.051	0.013		
7e	0.101	0.025	0.025	0.202	0.202	0.101	0.101		
8a	0.219	0.219	0.219	0.219	0.219	0.219	0.219		
8b	0.213	0.213	0.213	0.426	0.426	0.213	0.213		
9c	0.218	0.218	0.218	0.436	0.055	0.218	0.218		
9b	0.208	0.208	0.208	0.208	0.416	0.208	0.208		
10	0.218	0.218	0.218	0.218	0.218	0.218	0.218		
berberine	0.689	0.086	0.172	0.172	0.086	1.377	0.172		
norfloxacin	0.025	0.025	0.006	0.013	0.050	0.006	0.025		

reaction conditions (temperature, bases, solvents, and catalysts) were tried, but the reaction failed. This situation resulted in an alternative strategy from the reaction of mono-substituted *o*-phenylenediamines with compound **2** to prepare target molecules. Commercially available *o*-phenylenediamine was reacted with corresponding chlorinated or brominated compounds together with potassium carbonate to smoothly afford intermediates **11–17** (Scheme S1), which were further reacted with compound **2** in the presence of sodium pyrosulfite in *N*,*N*-dimethyl formamide at 110 °C to successfully access target products **4–10** (Schemes 1 and 2). The structures of all new berberine-hybridized benzimidazoles were confirmed through NMR and HRMS spectra, and the detailed synthetic processes and spectral data are provided in the Supporting Information.

Antibacterial Activity Assay. Table 1 shows that some target berberine-hybridized benzimidazoles exhibited more potent antibacterial effects in comparison to berberine and norfloxacin against the tested bacterial strains. This indicated that the introduction of benzimidazoles to berberine skeleton was greatly helpful for the antibacterial activities. Berberrubine-12-formaldehyde 2 displayed high inhibitory concentration toward the tested bacterial strains, while MIC values decreased when the formyl group was cyclized to afford benzimidazole derivative 3a, suggesting the great contribution of benzimidazole in exerting antibacterial activity. Compound 3a could

obviously inhibit the growth of MRSA and *S. aureus* ATCC 29213 in a low concentration of 0.068 mM, which was much lower than berberine. The inhibition concentration of **3b** against MRSA N315 strain was shown to be comparable to norfloxacin and exhibited a profound effect of 0.031 mM. Significantly, compound **3d** bearing electron-donating methyl group exhibited profound antibacterial effect with low MIC values (0.016–0.033 mM) toward MRSA, *S. aureus* ATCC 29213, and *Acinetobacter baumannii*. However, compound **3e** with methoxy group abolished the antibacterial activity partially or completely.

Alkyl chains were known to exert advantageous effects on biological activity via regulating lipid-water distribution coefficient and binding affinity to targets.^{28,29} Therefore, compounds 4a-f were synthesized to investigate the spacer lengths of alkyl chains on the antibacterial activities. Ethyl derivative 4a could effectively inhibit the growth of *S. aureus* 29213 and MRSA with low MIC values of 0.008 and 0.016 mM, respectively. Pentyl derivative 4c exhibited good potencies with MIC values ranging from 0.029 to 0.059 mM against MRSA, *S. aureus* and *S. aureus* 29213. Particularly, compound 4d exhibited excellent inhibitory effects (MIC = 0.029 mM) against *K. pneumoniae, E. coli*, and *A. baumannii*. Meanwhile, hexyl derivative 4d was active toward *S. aureus* with a low MIC value of 0.007 mM that was 12 and 4 times higher than those for berberine and norfloxacin, respectively.



Figure 2. (A) Eradication of the preformed S. aureus ATCC 29213 biofilms by compound 7d. (B) Images of S. aureus ATCC 29213 treated with 7d after staining by crystal violet.

When the number of carbon atoms increased, octyl and decyl compounds 4e and 4f were essentially inactive toward the tested bacteria, which might result from large lipophilicity. The data strongly indicated that the lengths of the alkyl chains significantly affected antibacterial activity and moderate lengths might be helpful for improving antibacterial activities. Inspired by the improved biological potentiality by incorporating cyclanes,²⁵ compounds 5a and 5b were synthesized and tested for their antibacterial potencies, where cyclopropyl derivative 5a was active toward the tested bacteria except for MRSA and K. pneumoniae. It was highly sensitive to S. aureus, A. baumannii, E. coli, and S. aureus ATCC 29213 at low concentrations (0.015-0.030 mM). Moreover, this compound exerted comparable activity (MIC = 0.004 mM) to norfloxacin (MIC = 0.006 mM) for *P. aeruginosa*. When cyclopropane was expanded to cyclohexane, the antibacterial activity of compound 5b significantly reduced.

Compounds **5c** and **5d** bearing allyl and propargyl groups were prepared to investigate the effects of unsaturated bonds on the antibacterial effect, respectively. Molecules **5c** and **5d** both showed a 3-fold decrease in anti-*S. aureus* with a low MIC value (0.031 mM) in comparison to berberine (0.081 mM). To explore the effect of hydrophilic groups on the antibacterial activity of berberine derivatives, compounds **6a**–**c** were prepared. Both of them were inactive toward the tested strains, suggesting that the increase of water solubility was deleterious to the antibacterial activities.

The structure and activity relationship (SAR) investigations were further moved onto the effect of halogen-containing benzyl groups, which were helpful for increasing lipid solubility and membrane penetrability,^{30,31} and compounds 7a-e were synthesized. Fluorobenzyl derivative 7a gave relatively low MIC values (0.014-0.055 mM) against MRSA, S. aureus, S. aureus ATCC 29213, P. aeruginosa, and A. baumannii, which were comparable to or even higher than norfloxacin. When the number of fluorine atom was increased, the antibacterial potency of compound 7b dropped obviously except that the inhibitory concentration toward S. aureus ATCC 29213 was decreased by 8-fold up to 0.007 mM compared to the fluorobenzyl molecule 7a (MIC = 0.055 mM). It seemed that the displacement of the fluorine atom (7a) by a chlorine atom (7c) reduced the activity of the molecule, but surprisingly enhanced its effect against S aureus (8 times in comparison to norfloxacin) and A. baumannii (the same as for norfloxacin). The introduction of more chlorine atoms resulted in compound 7d, which could effectively suppress S. aureus

ATCC 29213 growth with a low concentration of 0.006 mM. Furthermore, this molecule possessed 114- and 4-fold antibacterial potency than berberine and norfloxacin in inhibiting MRSA growth, respectively. Additionally, compound 7d gave a low MIC value (0.013 mM) toward *S. aureus* and *E. coil*, which was more than 2-fold higher than norfloxacin. Moreover, it showed relatively good inhibitory effect on *P. aeruginosa* with a MIC value of 0.051 mM, which was 27 times more potent than berberine. However, the 3,4-dichlorobenzyl derivative 7e only exhibited good inhibitory effect (MIC = 0.025 mM) against *S. aureus* and *S. aureus* ATCC 29213, which suggested that the number and location of halogen atoms on benzyl group would result in different effects on bioactivities.

Encouraged by the enhanced antimicrobial activity through the incorporation of amino derivatives,³² compounds 8 and 9 were synthesized to study the impact of amide groups on biological activities. Unfortunately, all of them abolished the activity completely or partially, indicating that the introduction of amides at the N-1 position of benzimidazole nucleus was not favorable for antibacterial potency. Stimulated by the lipophilic ester group, tert-butyl ester derivative 10 was synthesized and tested the antibacterial activity; however, it was not active toward the tested strains.

Biofilm Disruption Evaluation. The biofilm formation of *S. aureus* made conventional antibiotics hard to adequately attack and destroy infectious biofilm populations.^{33–35} Thus, new antibacterial agents inhibiting biofilm formation were imperative to confront *Staphylococcus* infection. The ability of compound 7d to disrupt the preformed biofilm of *S. aureus* ATCC 29213 was evaluated. As shown in Figure 2, the 2,4-dichlorobenzyl derivative 7d could eradicate more than 40% biofilm of *S. aureus* ATCC 29213 at concentrations of 4 × and 8 × MIC. Especially, when its concentration was increased to 16 × and 32 × MIC, the disruption of *S. aureus* ATCC 29213 biofilm exceeded 90%. These results demonstrated that compound 7d could effectively disrupt bacterial biofilm to exert antibacterial activity.

Metabolic Activity Evaluation. The alamar blue assay was conducted to determine the viability of *S. aureus* ATCC 29213 within the matured biofilm. As shown in Figure S1, the cell viability of *S. aureus* ATCC 29213 within biofilm reduced about 47% in the presence of compound 7d at the concentration of $8 \times \text{MIC}$, which indicated that active molecule 7d possessed certain ability to disrupt matured biofilm and then inhibit the viability of *S. aureus* ATCC 29213 cells.



Figure 3. Time-dependent killing of 7d ($6 \times$ MIC and $8 \times$ MIC) against early (A) and late (B) phase exponential S. aureus ATCC 29213.

Bactericidal Activity Evaluation. Efficient bactericidal activity is one of the most important features of an effective antibacterial agent.³⁶ The bactericidal performance of active molecule 7d ($6 \times$ MIC and 8 \times MIC) against S. aureus ATCC 29213 (early and late exponential phases) was evaluated, and the results are recorded in Figure 3. Compound 7d could rapidly kill early exponential phase S. aureus ATCC 29213 (2 h, $6 \times MIC$; 1 h, $8 \times MIC$), and the late exponential phase bacteria were also killed in the presence of molecule 7d (5 h, 6 \times MIC; 3 h, 8 \times MIC). In addition, it could be clearly observed in Figure 3 that the bactericidal rate of compound 7d was more rapid than norfloxacin against S. aureus ATCC 29213. These results demonstrated that compound 7d possessed strong bactericidal potency against S. aureus ATCC 29213, which would effectively reduce the probability of bacterial resistance development.

Cytotoxicity Evaluation. Excellent safety of antibacterial agents would reinforce their potential applications in the agricultural industry.³⁷ Therefore, the cytotoxicity of highly active compound 7d against Hek 293T, HFC, and red blood cells was evaluated with berberine as the positive control. As shown in Figure S2, the viabilities of Hek 293T and HFC cells still maintained 77.3 \pm 7.0 and 52.3 \pm 1.7% in the presence of compound 7d with the concentration of 50 μ M (8 × MIC), suggesting that compound 7d exhibited relatively low cytotoxicity toward the mammalian cells. In addition, no hemolysis was observed with compound 7d at concentrations \leq 512 µg/mL, indicating that compound 7d possessed good selectivity for S. aureus ATCC 29213 over mammalian cells. Moreover, a higher cell viability was observed for compound 7d than berberine, which indicated the advantage of this modification of chemical structure.

Drug Combination. Combination therapy has been considered as an alternative method to enhance the antibacterial efficacy, reduce the usage doses, and combat serious drug resistance.³⁸ Therefore, drug interaction assays were conducted to determine the synergistic effects of active compound 7d with norfloxacin. It could be clearly observed that the treated bacteria with drug combination were more susceptible in contrast to their individual use (Table S1), which suggested that the tolerance of norfloxacin might be reduced by compound 7d. The most efficient potentiation was obtained on the co-administration of norfloxacin with compound 7d against E. coli with outstanding synergism (FIC = 0.0157). Moreover, the MIC values of compound 7d toward MRSA, S. aureus, and S. aureus ATCC 29213 were reduced to 0.0004 mM. The synergy of target berberinehybridized benzimidazole 7d provided a promising strategy for searching how the limited therapeutic range of antibacterial

drugs could be broadened to more effectively address bacterial infections. Furthermore, the availability of this type of synergistic use needs further studies to optimize the antibacterial potencies, and deep mechanism investigation is indispensable to explain the drug combination.

Propensity to Develop Bacterial Resistance. The slow propensity to develop bacterial resistance is a crucial feature of antibacterial agents.^{39,40} Therefore, the probability of compound 7d or combination use to induce resistance in *S. aureus* ATCC 29213 was conducted with norfloxacin as positive control. The results in Figure 4 showed that the susceptibility



Figure 4. Propensity of *S. aureus* ATCC 29213 resistance development treated by compound 7d, norfloxacin, and drug combination.

of *S. aureus* ATCC 29213 to the 2,4-dichlorobenzyl derivative 7d was nearly unchanged after 15 passages, while the MIC values for norfloxacin were obviously increased by 32-fold after 14 passages, suggesting that compound 7d was hard to develop resistance in *S. aureus* ATCC 29213. Moreover, the emergence of resistance in norfloxacin was effectively reversed upon combination with compound 7d, which was consistent with the drug interaction that the 2,4-dichlorobenzyl derivative 7d could potentially elevate bacterial inhibitory efficacy.

Membrane Depolarization Assay. The similar chemical properties and action mechanisms of berberine to quaternary ammonium antiseptics make it target cytoplasmic membrane and DNA.⁴¹ To explore whether benzimidazole-hybridized berberines target bacterial membrane or not, a membrane depolarization study was performed using the DiSC35 fluorescence dye, which can be accumulated within the bacterial membrane, causing fluorescence quenching.⁴² Fluorescence intensity could be sharply increased when the membrane was depolarized. As shown in Figure S3A, when *S. aureus* ATCC 29213 was treated with the 2,4-dichlorobenzyl derivative 7d at increasing concentrations, a growing trend in fluorescence intensity was observed in a concentration-dependent manner, which demonstrated that compound 7d

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Figure 5. Preliminary summary of structure and activity relationships of berberine-hybridized benzimidazoles.

could interact with bacterial membrane and cause membrane depolarization.

Membrane Permeabilization Assay. Prodium iodide (PI) is a membrane-permeable dye that can only bind to nucleic acids by penetrating incomplete bacterial membranes, resulting in significant fluorescence changes.^{43,44} When *S. aureus* ATCC 29213 was treated with active molecule 7d, a remarkable concentration-dependent increase of the PI fluorescence was found (Figure S3B), which indicated that compound 7d could promote the inner membrane permeability of *S. aureus* ATCC 29213, thereby breaking the bacterial membrane integrity and resulting in the bacterial cell death. Moreover, the concentration-dependent effect of this compound on membrane permeabilization was correlated with its dose-dependent bactericidal behavior, suggesting that the rapid bactericidal activity of compound 7d might be attributed to its membrane-disrupting activity.

Docking Simulation. S. aureus sortase A (PDB ID: 1T2W) as an attractive target in anti-S. aureus drug exploitation was utilized in a flexible ligand-receptor docking investigation to further rationalize the observed antibacterial activity of active molecule 7d. As shown in Figure S4, active molecule 7d could well match the active pocket of sortase A through various weak interactions including the hydrophobic effect, $\pi - \pi$ stacking, and van der Waals. Noticeably, the hydroxyl group at the C-9 position of berberine skeleton could pick up a hydrogen bond with the oxygen atom of the carbonyl group in ASP-82 residue with a distance of 2.1 Å, and a binding energy was as low as -6.79 kJ/mol (total score, 5.25), which suggested the great contribution of the hydroxyl group in exerting antibacterial activity. In addition, the alkyl interactions between the chlorine atoms of benzyl group and various residues (ASN-127, PRO-126, PRO-79, and ARG-125) were observed. These interactions might promote the stabilization of compound-enzyme complex, which might be responsible for the excellent antibacterial activity of compound 7d.

Supramolecular Interactions between Active Molecule 7d and Calf Thymus DNA. DNA has been considered as a potent drug target and is widely utilized for the design and development of new efficient antibacterial drugs.^{45–47} Therefore, the interaction mode of compound 7d and DNA was explored using AO spectral probe. The absorption spectra of DNA–AO complex varying with compound 7d concentrations are recorded in Figure S5A. It could be clearly observed that the absorption spectra of the DNA–AO complex (499.5 nm) did not show redshift and hypochromic effect with the addition of compound 7d, which indicated that the interaction of active molecule 7d with DNA was not the typical intercalative mode.

Fluorescence quenching experiment was further performed to explore the target mode of compound 7d toward DNA. As the concentration of compound 7d increased, the fluorescence intensity of DNA–AO complex was significantly quenched (Figure S5B); however, no blueshift or redshift was observed. Meanwhile, the calculated fluorescence quenching data in the Supporting Information indicated that compound 7d could quench the DNA–AO complex via a static quenching mechanism. The decrease in the fluorescence intensity of the DNA–AO complex might be attributed to the result of the substitution of AO by compound 7d or the formation of a 7d– DNA–AO complex. The results in absorption spectrum showed that the binding of compound 7d to DNA–AO was not a typical intercalative action, so compound 7d was likely to interact with DNA via groove binding action.

Chloride quenching experiments were also conducted to further confirm the binding of compound 7d with DNA. As shown in Figure S6C, the fluorescence intensity of compound 7d–DNA complex remained basically unchanged with the addition of sodium chloride, which suggested that the electrostatic effect was not the interaction mode of compound 7d and DNA. These results suggested that compound 7d was sufficient to interact with DNA to form a supramolecular complex via groove binding action, thus exerting good antibacterial activity.

Intracellular ROS Determination Assay. ROS may cause disruption of membrane integrity and increase membrane permeability when the dose produced exceeds the antioxidant capacity of protective enzymes.⁴⁸ To test whether active compound 7d possessed potential in perturbing the bacterial metabolism process through facilitating ROS generation, the intracellular ROS level of *S. aureus* ATCC 29213 was detected when treated by active compound 7d using DCFH-DA dye. The results showed that the treatment of *S. aureus* ATCC 29213 cells by 2,4-dichlorobenzyl derivative 7d led to significant intracellular ROS production (Figure S6A). The increasing concentration of compound 7d was found to be highly correlated with the extent of ROS generation. There

Determination of GSH Activity. To further investigate the role of oxidative stress in interfering with the normal function of bacterial cells, the activity of GSH was determined. As shown in Figure S6B, when bacteria were treated with compound 7d, a gradual reduction of GSH activity in *S. aureus* ATCC 29213 was observed, which might be attributed to the ROS-dependent oxidation. These results showed that the increase of oxidative stress in *S. aureus* ATCC 29213 might also be responsible for the good antibacterial activity of 2,4dichlorobenzyl derivative 7d.

In summary, a unique type of novel berberine-hybridized benzimidazoles was developed that exhibited potent antibacterial effects. Particularly, the 2,4-dichlorobenzyl derivative 7d showed excellent inhibitory effects with a low MIC value of 0.006 mM against S. aureus ATCC 29213. The SARs indicated that the substituents on benzimidazole nucleus were important factors on antibacterial activities and the introduction of hydrophobic groups was favorable for antibacterial activity (Figure 5). The combination use of active compound 7d with norfloxacin was found to be synergistic against the tested bacteria, and resistance study revealed that compound 7d exhibited lower propensity to develop S. aureus ATCC 29213 resistance than norfloxacin. Preliminary mechanism studies demonstrated that compound 7d with low toxicity might damage membrane integrity, increase oxidative stress, and bind with DNA as well as S. aureus sortase A, empowering favorable biofilm disruption ability and rapid bactericidal activity. These results provided powerful information for further development of compound 7d as an alternative agricultural available agent to confront the growing challenges in S. aureus infections.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jafc.1c02545.

Results of metabolic activity, drug combination, cytotoxicity, membrane depolarization and permeabilization, docking simulation, supramolecular interaction between active compound 7d and calf thymus DNA, intracellular ROS generation and intracellular glutathione oxidation; synthetic procedures and characterization data of all new compounds; specific operation steps of in vitro antibacterial activity, biofilm disruption, bactericidal activity, membrane depolarization, membrane permeabilization, supramolecular interaction between active compound 7d and calf thymus DNA, intracellular ROS generation, intracellular glutathione oxidation and drug interaction, as well as resistance development and cytotoxicity assay; and NMR and HRMS spectra of some representative compounds (PDF)

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Notes

The authors declare no competing financial interest.

■ ABBREVIATIONS USED

MRSA N315, methicillin-resistant *Staphylococcus aureus* N315; S. aureus, *Staphylococcus aureus*; S. aureus ATCC 29213, *Staphylococcus aureus* ATCC 29213; P. aeruginosa, *Pseudomonas aeruginosa*; K. pneumoniae, *Klebsiella pneumoniae*; A. baumannii, *Acinetobacter baumannii*; E. coli, *Escherichia coli*; MIC, minimum inhibitory concentration; SARs, structure– activity relationships; CLSI, Clinical and Laboratory Standards Institute; diSC35, 3,3'-dipropylthiadicarbocyanine iodide; PI, propidium iodide; ROS, reactive oxygen species; GSH, glutathione; DCFH-DA, 2',7'-dichlorodihydrofluorescein diacetate; AO, acridine orange; DNA, deoxyribonucleic acid; FIC, fractional inhibitory concentration

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