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Design and biological evaluation of a novel type of potential multi-targeting antimicrobial sulfanilamide hybrids in combination of pyrimidine and azoles

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ABSTRACT

This work explored a novel type of potential multi-targeting antimicrobial three-component sulfanilamide hybrids in combination of pyrimidine and azoles. The hybridized target molecules were characterized by ¹H NMR, ¹³C NMR and HRMS spectra. Some of the developed target compounds exerted promising antimicrobial activity in comparison with the reference drugs norfloxacin and fluconazole. Noticeably, sulfanilamide hybrid **5c** with pyrimidine and indole could effectively inhibit the growth of *E. faecalis* with MIC value of 1 µg/mL. The active molecule **5c** showed low cell toxicity and did not obviously trigger the development of resistance towards the tested bacteria strains. Mechanism exploration indicated that compound **5c** could not only exert efficient membrane permeability, but also intercalate into DNA of resistant *E. faecalis* to form **5c**-DNA supramolecular complex, which might be responsible for its antimicrobial action. The further investigation showed that this molecule could be effectively transported by human serum albumins through hydrogen bonds and *van der Waals* force.

Sulfonamides are the first class of artificial synthetic antibacterial agents, whose core skeleton is *para*-aminobenzene sulfonamide that can compete with dihydrofolate synthetase, hinder the synthesis of dihydrofolate, effectively prevent the synthesis of nucleic acids and proteins, and then inhibit the growth of various microorganisms. The hybridization of core antibacterial skeleton sulfanilamide with various types of aromatic heterocycles like six-membered pyrimidine and diazine as well as five-membered thiazole and oxazole produced a variety of sulfanilamide types of antibacterial drugs, such as sulfadiazine, sulfamethoxypyridazine, sulfamoxole and sulfathiazole. Especially, the hybrids of sulfanilamide and pyrimidine with simple substituents have achieved great success, and a large number of pyrimidine derivatives as clinical antibacterial drugs are being frequently employed in clinic for the treatment of bacterial infections (Figure 1).¹ However, this kind of antibacterial drugs were found to cause some allergic reactions and show high toxicity to liver, kidney and central nervous system of human body.² Moreover, their prevalent clinical use or abuse developed the resistance with a significant public threat to health. This highlighted our keen interest to develop new hybrids of sulfanilamide and pyrimidine as antimicrobial agents with

lower toxicity, less bacterial resistance, higher safety and better activity.

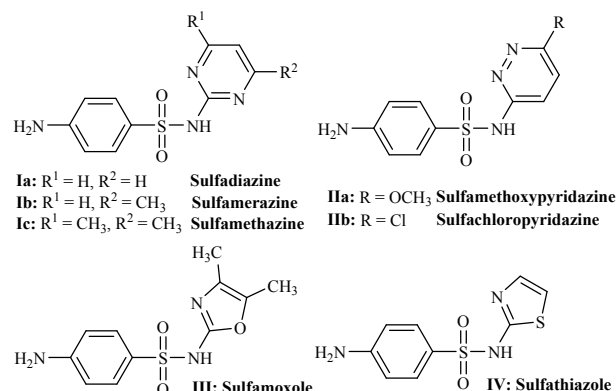


Figure 1. Sulfanilamide-aromatic heterocycle hybrids as clinical antimicrobial drugs.

Pyrimidine with a unique six-membered aromatic framework containing two nitrogen atoms is an important part in lots of biomolecules such as deoxyribonucleic acid (DNA) and ribonucleic acid (RNA),³ and as a crucial block is prevalently employed in the design of new drug molecules. It has been extensively investigated towards pyrimidine in combination with a variety of functional groups and a lot of pyrimidine derived molecules such as antibacterial trimethoprim and brodimoprim⁴

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as marketed in currently treating microbial infections. More importantly, numerous pyrimidine derivatives were reported to be potential as promising antimicrobial agents.⁷ Especially the combination of pyrimidine and heterocyclic azoles was revealed to result in strong bioactivity and high safety profile.⁸

Nitrogen-containing azole heterocycles and their analogs as antimicrobial agents⁹ like metronidazole, ornidazole and fluconazole have been widely used in clinic to benefit human health. It has been well validated that nitrogen heterocyclic

because this kind of heterocyclic fragment is able to make use of weak noncovalent interactions such as coordination bonds, hydrogen bonds, ion-dipole, cation- π , π - π stacking and hydrophobic effect as well as *van der Waals* force to bind with the enzymes and receptors in organisms by multiple sites and thus produce good biological potency.¹⁰ However, so far novel three-component hybrids of sulfanilamide with pyrimidine and azole ring or its analogs have been rarely reported.

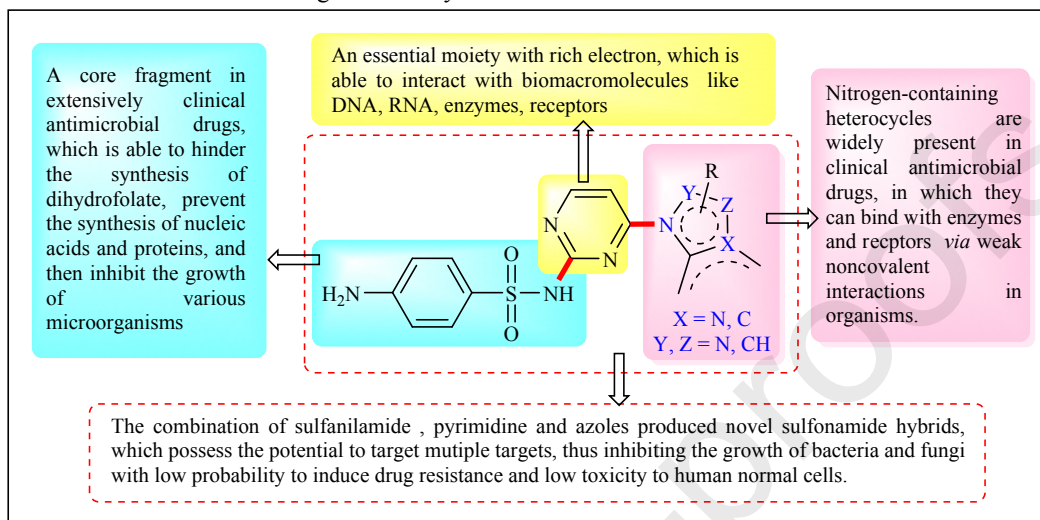
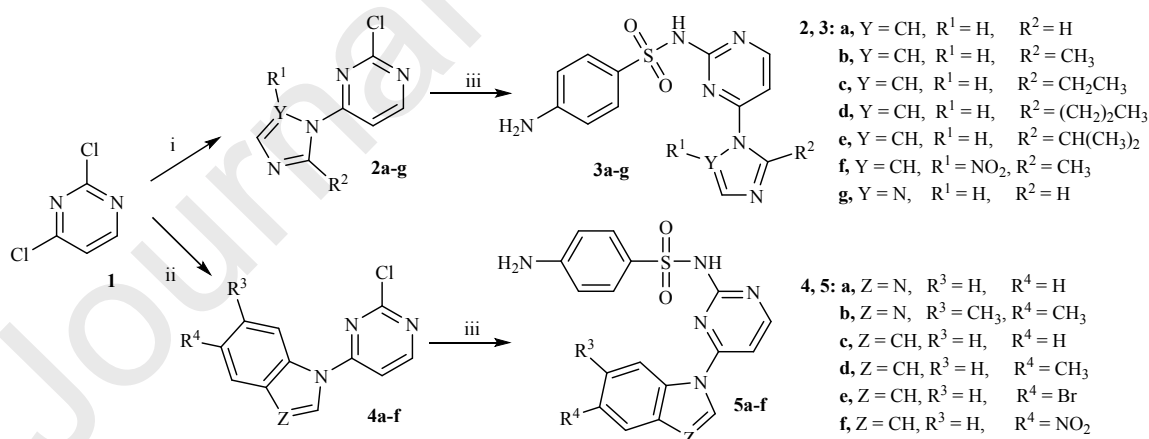


Figure 2. Design of novel three-component sulfanilamide hybrids with pyrimidine and azoles as potential multi-targeting antimicrobial agents.

In view of such stimulating overwhelming properties and as an extension of our studies towards novel antimicrobial agents, herein a new effort was explored towards a novel type of potential multi-targeting antimicrobial three-component sulfanilamide hybrids in combination of pyrimidine and azole ring or its analogs, the target azole derivatives and their analogs included imidazoles¹¹ and triazoles¹² as well as benzene-fused

azoles like benzimidazoles¹³ and their analog indoles¹⁴ (Figure 2). These new structural compounds might exert multi-site bindings with biomolecules, regulate the interaction and affinity between small molecules and biomolecules, therefore the target molecules might be conducive to overcome increasingly severe drug resistance and broaden antimicrobial spectrum.



Scheme 1. Synthetic route of sulfanilamide derivatives **3** and **5**. Reagents and conditions: (i) imidazoles and triazole, potassium carbonate, acetonitrile, 80 °C; (ii) benzimidazoles and indoles, potassium carbonate, acetonitrile, 80 °C; (iii) *para*-aminobenzenesulfonamide, cesium carbonate, acetonitrile, 80 °C.

The desired target sulfanilamide hybrids were constructed via easy synthetic procedures from commercial *para*-aminobenzenesulfonamide, azoles or indoles and 2,4-dichloropyrimidine according to the preparative route in Scheme 1. The reaction of 2,4-dichloropyrimidine with azoles or indoles in the presence of potassium carbonate efficiently produced five-membered azoles **2a-g**, fused five-membered azoles and their analog indoles **4a-f** with good yields,¹⁵ and then the coupling of intermediates **2a-g** and **4a-f** respectively with sulfanilamide in acetonitrile at 80 °C

with cesium carbonate as base afforded the target three-component sulfanilamide hybrids **3a-g** and **5a-f** in moderate to good yields ranging from 43.8% to 75.1%.

The spectral analysis from ¹H NMR, ¹³C NMR and HRMS data supported the structures of synthesized target compounds. All the protons of NH groups gave downfield singlets at 11.61–11.91 ppm. The chemical shifts of 3-H and 5-H in benzene ring appeared at 6.56–6.62 ppm, while the 2-H and 6-H protons in benzene ring showed downfield signals at 7.59–7.69

The electronegativity of nitrogen atom at 1-position in pyrimidine resulted in larger chemical values at 8.47–8.68 ppm for the 6-position proton in comparison with the 5-position proton (6.76–7.38 ppm) in pyrimidine. In ¹³C NMR spectra, the chemical shifts of carbons in benzene ring gave expected signals of 130.3–129.6 ppm and 113.2–112.8 ppm, which should be assigned to the 2,6-C and 3,5-C, respectively. The 4-C in the pyrimidine ring gave expected signals at 159.7–159.0 ppm. The other carbons ideally gave the signals at the expected regions.

All the synthesized hybrids **3a-g** and **5a-f** in combination of pyrimidine and azoles were evaluated for their antimicrobial activities *in vitro* by two folds serial dilution technique according to Clinical and Laboratory Standards Institute (CLSI) with the positive control of clinical drugs. Minimum inhibitory concentration (MIC, µg/mL) is defined as the lowest concentration of new compounds that completely inhibit the growth of bacteria and fungi. The tested strains including clinical

Gram-positive bacteria, six Gram-negative bacteria and five fungi.

As depicted in Table 1, sulfanilamide derivatives **3a-g** and **5a-f** showed inhibitory efficiency against some tested bacterial strains to some extent, while indolyl sulfanilamide **5c** displayed the most prominent efficiency. Imidazolyl compounds **3a-f** could not effectively inhibit the growth of all the tested bacteria, except for 2-propyl-imidazolyl derivative **3d**, which gave good activities with MIC value of 8 µg/mL against *S. aureus* and *S. aureus* ATCC 29213, and this might be beneficial from propyl group on the imidazole ring. The replacement of imidazolyl group in compound **3a** by triazolyl moiety yielded compound **3g**, resulting in decreased efficiency against most of the tested strains but increased activities towards *P. aeruginosa* (MIC = 4 µg/mL) and *P. aeruginosa* ATCC 27853 (MIC = 32 µg/mL), which might be attributed to the influence of nitrogen atom in triazolyl ring.

Table 1. The obtained antibacterial data as MIC (µg/mL) *in vitro* for sulfanilamide derivatives **3-5**^{a,b,c}

Comps	Gram-positive bacteria					Gram-negative bacteria					
	MRSA	<i>S. A.</i>	<i>S. A.</i> 25923	<i>S. A.</i> 29213	<i>E. F.</i>	<i>K. P.</i>	<i>E. C.</i>	<i>E. C.</i> 25922	<i>P. A.</i>	<i>P. A.</i> 27853	<i>A. B.</i>
3a	32	128	256	128	128	256	256	256	128	256	64
3b	128	256	256	256	128	512	128	256	128	256	256
3c	128	128	128	64	64	64	128	512	128	128	64
3d	16	8	64	8	64	64	16	128	128	128	32
3e	32	256	256	32	128	256	64	128	128	32	64
3f	64	128	128	128	128	64	128	256	256	128	64
3g	128	512	256	64	256	32	128	256	4	32	128
5a	8	256	512	256	256	256	128	256	512	256	512
5b	64	256	256	128	256	256	256	512	256	128	128
5c	2	32	4	8	1	32	32	32	8	1	16
5d	4	16	4	16	8	8	8	32	4	4	64
5e	4	64	2	16	4	32	4	2	4	4	128
5f	16	32	4	64	8	32	8	128	16	8	128
A	8	0.5	1	2	4	4	16	8	2	0.5	8

^a Minimal inhibitory concentrations (MIC, µg/mL) were determined by micro broth dilution method for microdilution plates.

^b MRSA, methicillin-resistant *Staphylococcus aureus*; *S. A.*, *Staphylococcus aureus*; *S. A.* 25923, *Staphylococcus aureus* ATCC 25923; *S. A.* 29213, *Staphylococcus aureus* ATCC 29213; *E. F.*, *Enterococcus faecalis*; *K. P.*, *Klebsiella pneumoniae*; *E. C.*, *Escherichia coli*; *E. C.* 25922, *Escherichia coli* ATCC 25922; *P. A.*, *Pseudomonas aeruginosa*; *P. A.* 27853, *Pseudomonas aeruginosa* ATCC 27853; *A. B.*, *Acinetobacter baumannii*.

^c **A** = Norfloxacin.

It was shown that the benzimidazolyl sulfanilamide hybrids **5a-b** gave relatively lower inhibitory potencies with MIC values ranging from 64 to 512 µg/mL, except for towards MRSA strains. Benzimidazolyl sulfanilamide hybrid **5a** exhibited obvious suppression to the growth of MRSA with MIC value of 8 µg/mL, being more active than 5,6-dimethyl-benzimidazolyl derivative **5b**, which indicated that the methyl substitution might be unfavorable for the antibacterial activity.

In comparison with benzimidazolyl sulfanilamide derivatives **5a-b**, most of benzimidazolyl analogous indolyl derivatives **5c-f** showed relatively better activities in inhibiting the growth of the test bacterial strains and gave broader antibacterial spectrum. It was noticed that indolyl derivative **5c** showed apparent inhibition with the same MIC value of 1 µg/mL towards *E. faecalis* and *P. aeruginosa* ATCC 27853, which was superior or comparable to standard drug norfloxacin. In comparison to imidazolyl and benzimidazolyl sulfanilamides **3a** and **5a**, indolyl derivative **5c**

gave more efficient potency against all the tested bacteria strains, manifesting that the indole ring played a crucial role in improving antibacterial activity.

Moreover, MRSA was more sensitive to compounds **5c-e** than norfloxacin (MIC = 8 µg/mL) with MIC values ranging from 2 to 4 µg/mL. Methylindolyl derivative **5d** gave good activities against MRSA, *S. aureus* ATCC 25923, *P. aeruginosa* and *P. aeruginosa* ATCC 27853 with MIC value of 4 µg/mL. Bromoindolyl derivative **5e** displayed relatively good inhibitory potency (MIC = 2 µg/mL) against *S. aureus* ATCC 25923 and *E. coli* ATCC 25922. The displacement of bromoindolyl moiety by nitroindolyl group would afford compound **5f** with good inhibitory efficiency (MIC = 4 µg/mL) against *S. aureus* ATCC 25923. Noticeably, Gram-negative *A. baumannii* was insensitive to sulfanilamide derivatives **3a-g** and **5a-f** with MIC values of 32–512 µg/mL, which showed that these substituent groups in

Antifungal assay revealed that the sulfanilamide derivatives **3a-g** and **5a-f** showed weak to good antifungal activities *in vitro* against the tested fungi (Table S1). It was noticed that the antifungal abilities of these compounds displayed similar potency to their antibacterial efficiencies. Towards *C. albicans* ATCC 90023, 2-propylimidazolyl derivative **3d** (MIC = 4 µg/mL) was found to be more efficient than other imidazolyl compounds (MICs = 16–128 µg/mL). In addition, the triazolyl derivative **3g** exerted good antifungal efficacy with MIC value of 8 µg/mL against *C. albicans*. It was found that benzimidazolyl derivatives **5a-b** exhibited weak inhibitory activities against the tested fungi. Among the indolyl derivatives **5c-f**, the low MIC values of 2–8 µg/mL were observed towards *C. albicans*, *C. tropicalis* and *A. fumigatus*, which suggested that these compounds should be

they might be potential to become new promising antifungal agents through deep investigation.

In order to identify the safety profile, the highly active indolyl pyrimidine hybridized sulfanilamide derivative **5c** was further evaluated for toxicities against lung cancer cell (A549) and human normal lung epithelium cells (BEAS-2B) with doxorubicin (0.5 µM) as a positive standard control using the colorimetric cell proliferation MTT assay. The results of these assays are depicted in Figure 3. The cell viability of hybrid **5c** towards these two cell lines was at least 70% after being incubated at the concentration of 64 µg/mL for 24 h, which manifested that compound **5c** displayed relatively low toxicity to A549 and BEAS-2B cell lines compared to standard drug doxorubicin.

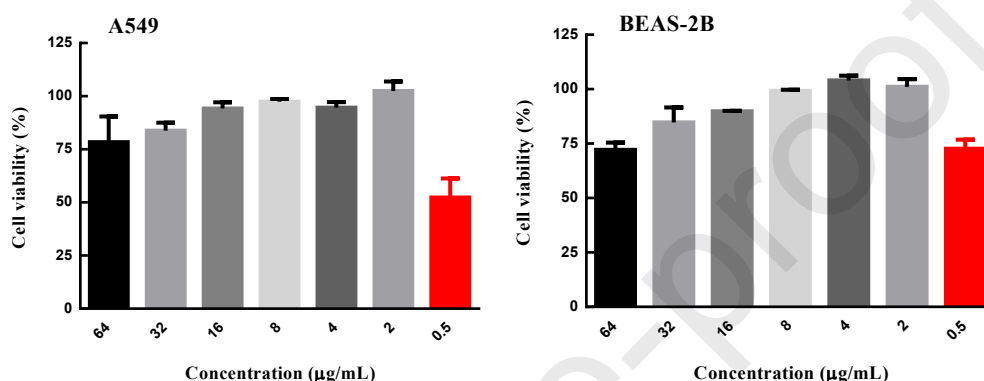


Figure 3. Cell viability of lung cancer cells (A549) and human normal lung epithelium cells (BEAS-2B) incubated with the active compound **5c** at different concentrations.

It is important for potentially antimicrobial agents that the development of resistance to their action is not readily attained.¹⁶ To evaluate the probability of these molecules to elicit resistance in bacteria, the drug resistance study of indolyl pyrimidine hybridized sulfanilamide derivative **5c** against *E. faecalis* was carried out. As shown in Figure 4, the susceptibility of *E. faecalis* bacteria to indolyl derivative **5c** remained nearly unchanged even after 16 passages. By contrast, the drug resistance developing ability of *E. faecalis* against norfloxacin was obvious, indicating that sulfanilamide hybrid **5c** with pyrimidine and indole did not significantly trigger the development of drug resistance as easily as the reference drug norfloxacin.

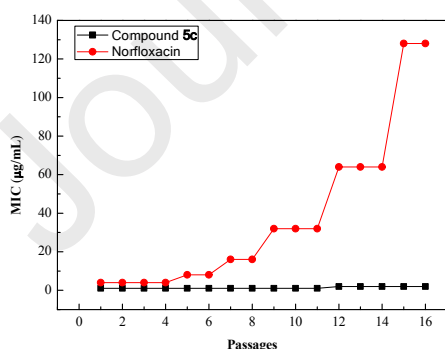


Figure 4. Development of *E. faecalis* resistance to compound **5c** and norfloxacin.

In order to study the bactericidal action of indolyl pyrimidine hybridized sulfanilamide derivative **5c**, a time-kill kinetics assay was performed against *E. faecalis* at different concentrations (MIC and 6 × MIC). As shown in Figure S1, there was more than 3 log (CFU/mL) reduction in the number of viable bacteria at concentrations of MIC and 6 × MIC against *E. faecalis*. These

results clearly revealed that the active hybrid **5c** was capable of killing Gram-positive bacteria *E. faecalis* rapidly.

DNA-gyrase (PDB code: 2XCS) as an attractive target in antibacterial drug exploration was applied to be a flexible ligand-receptor docking investigation to further rationalize the observed antibacterial activity of indolyl derivative **5c**. The molecular simulation results (Figure 5) showed that nitrogen atom at 3-position in the pyrimidine ring and the hydrogen atom of the NH₂ group at the 4-position on benzene formed two hydrogen bonds with DC11 and Asp1083B residues, respectively. The value of binding energy is -6.9 kcal/mol. These non-covalent binding might be conducive to stabilize the **5c**-enzyme complex, which might contribute to the superior inhibitory efficacy.

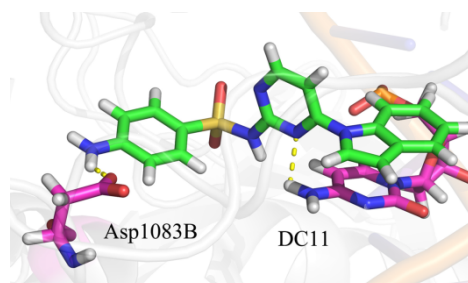


Figure 5. Supramolecular binding of compound **5c** docked in the bacterial DNA-gyrase (PDB code: 2XCS).

Bacteria with biofilms are inherently insensitive to antimicrobial agents and they are much more resistant to conventional antibiotic treatment.¹⁷ Permeabilization of the bacterial membrane of bacteria *E. faecalis* was carried out with propidium iodide (PI) as fluorescent indicator which can only pass through the membrane of compromised cells and then fluoresce red after binding with double-stranded DNA.

treatment of cells with active molecule **5c**. The results of bacterial membrane permeabilization are shown in Figure 6. Obvious fluorescence enhancements at 617 nm of the mixtures of indolyl pyrimidine hybridized sulfanilamide derivative **5c** with PI-treated *E. faecalis* appeared and became steady after 60 min, which demonstrated that sulfanilamide hybrid **5c** with pyrimidine and indole could permeate the membranes of *E. faecalis* strains efficiently.

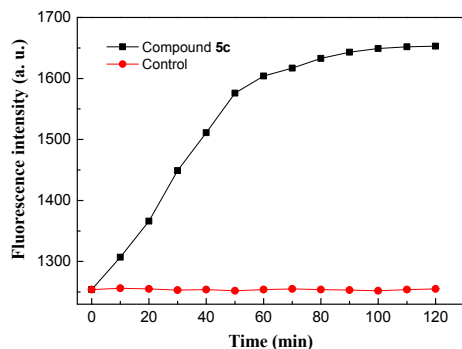


Figure 6. Bacterial membrane permeabilization of compound **5c** ($12 \times \text{MIC}$) against *E. faecalis*.

The detection of hyperchromism and hypochromism by absorption spectroscopy, which are critical spectral features to distinguish the change of DNA double helical structure, is one of the most important and useful method in DNA binding studies. The observed strong hyperchromism might be attributed to the denaturation of DNA caused by the strong interaction between the electronic states of intercalating chromophore and that of the DNA base. As shown in Figure S2, the UV-vis spectra revealed that the maximum absorption peak of DNA at 260 nm showed proportional increase and slight blue shift with the increasing concentration of hybrid **5c**. Remarkably, the absorption values of the **5c**-DNA complex were lower than that of sum of free DNA and free molecule **5c**. This hyperchromism effect might be attributed to the formed binary complexes between **5c** and DNA bases. In general, the observed spectral properties could be explained by the potent interaction of indolyl derivative **5c** with *E. faecalis* DNA (Figure S3).

To further elucidate the supramolecular interaction between hybrid **5c** and *E. faecalis* DNA, neutral red (NR, a planar phenazine dye with a confirmed intercalative binding mode with DNA) was employed as a spectral probe due to its lower toxicity and higher stability. Therefore, NR was chosen to investigate the binding mode of compound **5c** with DNA in present work, and the absorption spectra of the NR dye after the addition of DNA were shown in Figure S4.¹⁸ The absorption peak of NR at around 460 nm gradually decreased with the increasing concentration of DNA, and a new band at around 530 nm appeared, which could be attributed to the formation of the new DNA-NR complex. The isosbestic point at 500 nm provided the evidence for the formation of DNA-NR complex.

The absorption spectra of a competitive binding between NR and **5c** with DNA in Figure S5 showed that with the increasing concentration of hybrid **5c**, the maximum absorption around 530 nm of the DNA-NR complex decreased, but a slight intensity increase was observed in the developing band around 460 nm. In comparison with the absorption band of the free NR in the presence of the increasing concentrations of DNA (Figure S4), the spectra in the inset of Figure S5 exhibited the reverse process. The apparent changes of spectra suggested that indolyl pyrimidine hybridized sulfanilamide derivative **5c** could

in the DNA-NR complex.

Many promising new drugs have been rendered ineffective because of their unusually high affinity to the protein human serum albumins (HSA), which is the principal extracellular protein of the circulatory system. As a result, the investigations of interactions between drugs or bioactive small molecules and HSA are not only beneficial to provide a proper understanding of the absorption, transportation, distribution, metabolism and excretion properties of drugs, but also significant to design, modify and screen drug molecules.¹⁹

The binding mode between HSA and the highly active molecule **5c** was investigated with UV-vis spectroscopy on the molecular level. In the binding experiment (Figure S6), the aromatic rings in Tryptophan (Trp-214), Tyrosine (Tyr-411) and Phenylalanine (Phe) residues in HSA gave absorption peak at 278 nm and the peak intensity increased with the addition of hybrid **5c**, which indicated the supramolecular interaction of indolyl pyrimidine hybridized sulfanilamide derivative **5c** with HSA and the extension of HSA peptide strands (Figure S6).

The unchanged maximum absorption wavelength implied that there was a non-covalent interaction between indolyl derivative **5c** and HSA. The fluorescence intensity of Trp-214 may change when HSA interacts with other small molecules, which could be reflected in the fluorescence spectra of HSA in the UV region. The different mechanisms of quenching are usually classified as either dynamic quenching or static quenching, which can be distinguished by their different dependence on temperature (Figure S7, Table S2).

The Modified Stern-Volmer plots were showed in Figure S8 and the calculated results were depicted in Table S3. As shown in Figure S9 and Table S3, with the increase of temperature, the diffusion coefficient increased while the stability of the **5c**-HSA complex reduced. The results suggested that there was a single class of binding sites on HSA for sulfanilamide hybrid **5c**. In addition, the thermodynamic parameters were obtained in Table S4. Both hydrophobic interactions and hydrogen bonds played vital roles in the binding of molecule **5c** to HSA and seemed to contribute to the stability of the **5c**-HSA complex.²⁰

In conclusion, this work employed a facile synthetic procedure to develop a novel type of three-component sulfanilamide hybrids with pyrimidine and azoles. Some prepared new sulfanilamide derivatives could significantly inhibit the growth of the tested strains with equipotent or superior activities to the current clinical drugs norfloxacin and fluconazole. Notably, the indolyl pyrimidine hybridized sulfanilamide derivative **5c** not only showed superior inhibitory behaviors against *E. faecalis* ($\text{MIC} = 1 \mu\text{g/mL}$) to norfloxacin, but also exhibited low toxicity towards A549 and BEAS-2B cell lines and no obvious propensity to trigger development of resistance towards *E. faecalis*. Moreover, the active sulfanilamide hybrid **5c** was also membrane active and could kill bacteria in short time. The specific interaction of indolyl derivative **5c** with *E. faecalis* DNA displayed that it could intercalate into DNA to form **5c**-DNA complex which might further block DNA replication to exert their powerful antimicrobial activities. Binding investigations revealed that compound **5c** was able to be carried by HSA, which could generate fluorescent quenching by the active hybrid **5c** as a result of the formation of ground-state, and the calculated parameters indicated that the binding process was spontaneous. These results revealed that indolyl pyrimidine hybridized sulfanilamide derivative **5c** could be regarded as a promising lead compound for the further development as potential

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higher potency against *E. faecalis* than

norfloxacin.

- Multi-targeting antimicrobial sulfanilamide hybrids in combination of pyrimidine and azoles were developed.
- Indolyl sulfanilamide derivative **5c** displayed

- Hybrid **5c** exhibited low toxicity and no obvious propensity to trigger development of resistance.
- The active molecule **5c** could disturb the cell membrane, intercalate into *E. faecalis* DNA.
- Compound **5c** could be effectively transported by human serum albumins.

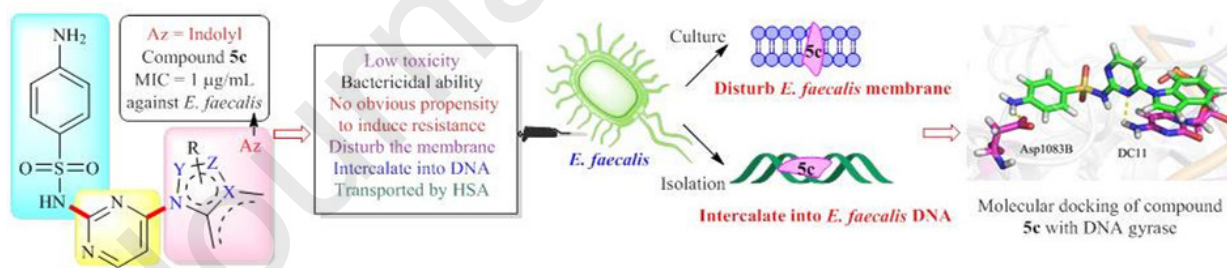
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☐ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

Graphical Abstract

Design and biological evaluation of a novel type of potential multi-targeting antimicrobial sulfanilamide hybrids in combination of pyrimidine and azoles

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