

Novel Pentafluorosulfanyl-containing Triclocarban Analogs selectively kill Gram-positive bacteria.

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Abstract

The antibacterial and antibiofilm efficacy of our novel pentafluorosulfanyl-containing triclocarban analogs was explored against seven different Gram-positive and Gram-negative indicator strains. After initial screening, they had bactericidal and bacteriostatic activity against Gram-positive bacteria, especially *Staphylococcus aureus* and *MRSA* (methicillin-resistant *Staphylococcus aureus*) in a very low concentration. Our results were compared with the most common antibiotic being used (Ciprofloxacin and Gentamycin); the novel components had significantly better antibacterial and antibiofilm activity in lower concentrations in comparison to the antibiotics. For instance, EBP-59 minimum inhibitory concentration was < 0.0003 mM, while ciprofloxacin 0.08 mM. Further antibacterial activity of novel components was surveyed against 10 clinical antibiotic resistance *MRSA* isolates. Again, novel components had significantly better antibacterial and antibiofilm activity in comparison with antibiotics. Mechanistic studies have revealed that none of these novel compounds exhibit any effect on the reduced thiol, disrupting iron sulfur clusters, or hydrogen peroxide pathways. Instead, their impact is attributed to the disruption of the Gram-positive bacterial cell membrane. Toxicity and safety testing on tissue cell culture showed promising results for the safety of components to the host.

Keywords: Novel antimicrobials; Gram-positives; *Staphylococcus aureus*; *MRSA*; pentafluorosulfanyl; diarylurea scaffold; antibiofilm; antibacterial.

Introduction

Bacterial resistance to antimicrobials has become a pressing concern in healthcare, demanding the development of effective formulations targeting multi-resistant pathogens (1). To address this challenge, researchers have turned to drug repurposing, a strategy that explores novel uses for authorized or experimental pharmaceuticals beyond their initial applications, offering advantages such as reduced costs and shorter development timelines (2). In this context, the exploration of multitarget molecules has emerged as a promising approach to counteract pathogens through diverse mechanisms. Diarylureas, such as regorafenib, sorafenib, linifanib, ripretinib, and tivozanib, have long been recognized as anticancer agents (3). One intriguing avenue involves the repurposing of diarylureas with anticancer properties for novel indications, such as antimicrobial, anti-inflammatory, and antiviral applications (4).

Trifluoromethyl groups, serving as bio-isosteric replacements for chlorine atoms, have found widespread use in medicinal chemistry. Hence, it is unsurprising that specific N,N'-diarylureas that comprise a trifluoromethyl moiety have demonstrated encouraging antibacterial properties. (5). An example is cloflucarban (TFC, 3-trifluoromethyl-4,4'-dichlorocarbanilide), a trifluoromethyl-substituted diarylurea which shares a similar spectrum of activity and pharmacokinetic profile with triclocarban (TCC). In recent times, a number of diarylureas analogs of TCC have been identified as having antibacterial and antifungal properties. The presence of pentafluorosulfanyl and trifluoromethylcoumarine groups characterizes these analogs. Given the risks associated with TCC, the need for alternative antimicrobial agents has become crucial (3).

In the recent years, a novel bio-isosteric trifluoromethyl unit, the pentafluorosulfanyl group (SF_5), has emerged in medicinal chemistry, finding applications in agriculture and material chemistry (6-8). Regarded as a "super-trifluoromethyl group," the SF_5 -group possesses several advantageous properties over its isostere trifluoromethyl group, the compound exhibits a tetragonal bipyramidal morphology and possesses a greater electronegativity value of 3.65, in comparison to trifluoromethyl's value of 3.36. Additionally, it displays higher lipophilicity and notable steric volume, which is a bit smaller compared to *tert*-butyl but more than trifluoromethyl. The compound's hydrolytic and chemical stability has also been approved (9). These unique characteristics have led to a significant rise in the utilization of SF_5 in medicinal

chemistry over the past decade, making it an exceptionally appealing substituent for medicinal applications. SF₅-containing building blocks have garnered attention among medicinal chemists owing to their stated ability to decelerate metabolic rates and their eco-friendliness vis-à-vis lack of chlorine atoms, despite their elevated cost relative to analogous CF₃ compounds. Building upon the growing utilization of SF₅ in medicinal chemistry and its favorable environmental profile, the objective of this study is to incorporate this innovative group onto the N,N'-diarylurea scaffold to explore for novel antimicrobial agents (10).

This present study aims to explore the synthesis and antimicrobial and antibiofilm activity of novel diphenylurea agents, particularly inspired by TCC but bearing different aryl moieties. Mechanistic studies, such as cell wall or hydrogen peroxide assays, are conducted to elucidate the compounds' modes of action. Cytotoxicity assays using cell lines will ensure their safety as a unique class of antimicrobials.

Result and Discussion

During the recent years, overuse, and misuse of antibiotics against bacterial infections has brought up an issue regarding prevalence of multi-resistant bacteria to conventional antibiotics. This has led that development of novel antimicrobial agents and their delivery mechanism has turned into a global concern (11). Given that diarylureas analogs of TCC have been recognized for their antibacterial and antifungal properties (3), our study investigates the antibacterial efficacy of our novel pentafluorosulfanyl-containing triclocarban analogs.

Chemistry

The eighteen N,N'-diarylureas evaluated in this work were synthesized following a simple and straightforward procedure consisting of the coupling of phenyl isocyanates with the corresponding anilines, as previously described by some of us reported (5, 12, 13) (**Fig 1**). In turn, the intermediate phenyl isocyanates were either commercially available or synthesized *in situ* from the reaction of the precursor anilines with triphosgene. The analytical data of the compounds fully agreed with the data previously reported (5, 12, 13).

Antibacterial activity

We explored the antibacterial activity of compounds against several bacterial pathogens, obtaining measurements of minimum inhibitory concentration (MIC), and minimum bactericidal concentration (MBC), (**Table 1**). None of the diphenylurea compounds were effective against any of the Gram-negative bacteria evaluated in this study. The only exception was the effect of compound **59** on *P. mirabilis* with MIC and MBC of 0.002 mM. However, regarding the main outcome measure of this study, three compounds **59**, **61**, and **62** were effective against *S. aureus*, *S. epidermidis* and *MRSA* spp. Exact dosages in respect to each compound and bacteria is illustrated in **Table1**. Compound **59** showed the best efficacy against *S. aureus*, *MRSA*, and *S. epidermidis*. Compound **61** also was effective against these bacteria with relatively higher doses. Compound **62** was also effective against these Gram-positive bacteria, however, it had the highest doses comparatively (**Fig 2**).

Given the efficacy against the reference strain of *MRSA*, we tested further ten clinical isolates of *MRSA* evaluated with compounds **59**, **61** and **62**. The results were extremely promising showing similarly effective doses of MIC and MBC against the clinical isolates (**Table 2**).

Antibiofilm activity

The antibiofilm activity of antimicrobials can vary significantly compared to their effectiveness against planktonic forms of bacteria due to differences in physiology and structure of the biofilm (14). In fact, certain reports have demonstrated that biofilms require concentrations of antibiotics up to 100 times higher than those needed to eliminate planktonic bacteria (15-17). To assess the antibiofilm potential of the novel compounds, their Minimum Biofilm Inhibitory Concentration (MBIC) were determined **Table 1**. Similar to planktonic antibacterial activity, compounds **59**, **61** and **62** were effective against biofilm form of Gram-positive bacteria. Compounds **59** and **61** had the best antibiofilm activity in comparison to other compounds against *S. epidermidis*, *MRSA* and *S. aureus*. Compound **62** also showed relatively better results against these bacteria. When evaluating these compounds against ten clinical isolates of *MRSA*, all these isolates showed had their biofilms inhibited by these compounds (**Table 2**).

Antimicrobial mechanism of action

A wide variety of mechanisms of antimicrobial action are reported for antimicrobial agents. Here in this study the most common mechanisms such as oxidation of the redox buffer reflected in reduced thiol content (19), related oxidative stress as seen by increase in reactive oxygen species (20) and breakdown of redox enzyme [Fe-S] clusters (21), and cell membrane dysfunction (11) were explored to have a general view of the various diphenylurea compounds mechanism of action.

A well-established method to evaluate bacteria viability, propidium iodide (PI) staining was performed to assess the membrane permeability. Compounds **59** and **61** showed no effect on the cell membrane of Gram-negative bacteria, *P. aeruginosa*. However, both compounds were effective at cell membrane disruption of *MRSA* strains. Proving the effective bactericidal of such compounds on the Gram-positive bacteria's cell membrane (**Fig 3**). Further evaluations including reduced thiol, iron detection Ferine-S, and hydrogen peroxide levels were performed to explore other possible mechanisms of action of these compounds. These assays demonstrated that none of these novel compounds have any effect on these processes of toxicity.

Due to the considerable structural differences between the Gram-positive and Gram-negative cell membranes of bacteria(18), as well as eukaryotic cells (19), certain antibiotics, such as polycations and chelators, selectively target the distinctive structure of bacterial cell membranes (20). This targeted approach allows antibiotics to eradicate bacteria more specifically while preserving host cells (20). In our investigation of membrane disruption potency within our chosen components, we employed the membrane leakage probe (PI), which binds to DNA, to assess membrane destabilization following a 1-hour exposure to our agents. Results indicate that all selected agents disrupted the Gram-positive cell membrane compared to the untreated group ($p < 0.001$), but not sig effect of Gram-negative cell membrane.

Toxicity in Eukaryotic Cell lines

We selected compounds EPB-59, EPB-61, and EPB-62 due to their elevated antibacterial activity. Subsequently, we evaluated their cytotoxicity in eukaryotic cell lines derived from human, canine, or simian origins. In HEL, HeLa, MT4 and VERO cells, the MCC or CC₅₀ values were similar for the three compounds and in the order of 4-20 μ M. EPB-59 and EPB-61 showed lower cytotoxicity in canine MDCK cells, giving a CC₅₀ value ≥ 100 μ M (**Table 3**). The CC50 values of EPB-59 and EPB-61 on MDCK cells reveal their exceptional cytotoxic profiles in comparison to other compounds, with concentrations varying from ≤ 0.8 μ M to a > 100 μ M. Notably, EPB-59 and EPB-61 possess cytotoxicity levels 2 logs lower than that of EPB-34.

***In vitro* results**

The new diphenylurea compounds studied here were not effective against Gram-negative pathogens such as *E. coli*, *P. aeruginosa*, *K. pneumonia*, and *P. mirabilis* except compound **59**, which had acceptable potency regarding MIC, MBC, and MBIC to *P. mirabilis* spp. This could indicate that variations in diphenyl-urea antibiotics could make them efficient to use against Gram-negative bacteria. Perhaps this compound and how it differs from others could be a future topic for further study. Furthermore, promising results regarding the application of compounds **59**, **61**, and **62** were achieved with Gram-positive bacteria such as *S. aureus*, *S. epidermidis*, and methicillin-resistant strains *S. aureus* (*MRSA*) specifically with compound **59**, having relatively superior results while factoring cell cytotoxicity in all cell lines evaluated. Even though results regarding the use of such compounds on *MRSA* species did not accompany a safe dose regarding human cell lines, MIC, MBC, and MBIC measurements of 10 *MRSA* clinical isolates showed these compounds could be used in a rather safe measure when evaluated on clinical isolates. All *MRSA* clinical isolates showed susceptibility to compounds **59**, **61** and **62** regarding MBIC and MIC (**Table 2**). With compound **59** yet again proving to be the best option across all measurements regarding safety and efficacy, all these compounds showed a significantly higher potency when compared to conventional antibiotics evaluated.

Other compounds were also evaluated in this study. However, many of them did not prove as effective as the three discussed above. Compound **34** showed promising results regarding MIC in Gram-positive bacteria assessed in this study with a similar cytotoxicity profile as compound **59** regarding HeLa, HEL, and Vero cell lines. Unfortunately, the same does not apply in the context of MBIC and MBC. Compound **37** showed slightly worse results and its cytotoxicity was not

assessed. Compound **40** did not show any promising effectiveness against Gram-positive bacteria even in comparison to conventional antibiotics.

In vivo efficacy

The most effective compounds-**59**, **61**, and **62**, were selected to evaluate their *in vivo* efficacy. Mice were infected systemically with the community-associated MRSA strain MW2 at an infectious dose of 5×10^7 CFU as described (21). Treatment groups were injected intraperitoneally with the compound of interest one hour before or after infection, with a saline-injected control group. After 24 hours of infection, mice were sacrificed and blood, peritoneal fluid, spleen, liver, kidney (left), heart, and lungs were harvested for CFU enumeration to determine the bacterial burden in these organs. These experiments revealed a consistently high bacterial burden in all organs of the animals 24 hours after treatment, irrespective of the specific treatment administered (**Fig 4**). The single-dose treatment failed to reduce bacterial counts in any of the animal organs, even after experimenting with different compound concentrations. While disappointing, the lack of promising results in the single-dose treatments may have a variety of reasons including, rapid metabolism and excretion of the compounds by the liver or to the acknowledged intracellular aspect of *S. aureus* in its infectious cycle (22). Consequently, further research is imperative to determine the *in vivo* efficacy of compounds **59**, **61**, and **62**. Specifically, if a pulsatile and continuous treatment over 2-4 days may prove to be more effective. Additionally, investigating the possibility of topical application for these components presents another viable option.

Conclusion:

Overall, for the first evaluation of such compounds, the results demonstrated Gram class selectivity, a cell wall/membrane associated mechanism of antimicrobial activity. Cell toxicity and animal results indicate challenges but still suggest there is promise to explore applications of these novel compounds as useful antimicrobials moving forward.

Materials and Methods

Pentafluorosulfanyl-containing triclocarban analogs preparation and characterization.

All the pentafluorosulfanyl-containing diarylureas were synthesized as previously reported (5, 12, 13). In all cases the analytical and spectroscopic data matched with those previously reported in the above mentioned three references.

Bacterial Preparation

All bacterial strains subjected to testing had been stored at -70°C prior to the experiment. The experiment commenced by streaking small amounts of the frozen bacterial samples onto Muller-Hinton agar (MHA) culture medium (BD Bacto, Oxoid, Basingstoke, UK Cat# X296B) and allowing for overnight incubation. Bacteria were re-cultured on MHA plates as required to confirm colony morphology and lack of contamination. Strains tested were *Pseudomonas aeruginosa* ATCC 27853, *Escherichia coli* ATCC 25922, *Staphylococcus aureus* ATCC 25923, Methicillin resistant *S. aureus* (MRSA) ATCC 33591, *Staphylococcus epidermidis* ATCC 12228, *Klebsiella pneumoniae* ATCC, *Proteus mirabilis* ATCC 35659 and 10 clinical isolates of MRSA from Foothills hospital, Calgary, Canada. To prepare for susceptibility testing, a protocol modified from Lemire *et al* was used (23). MHB was swab inoculated from MHA colonies and shake incubated until turbid. Dilution of the broth with 0.9% saline was performed as necessary to match a 1.0 McFarland standard.

Planktonic Susceptibility

Minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) assessments were conducted on all bacterial strains using 96-well microtiter plates. To prepare the wells for bacteria, serial dilutions of the antibacterial agents were carried out along the rows of two microtiter plates. The antibacterials were 2-fold diluted with Mueller-Hinton broth (MHB) in each column, reaching a volume of 75 µL. The 1.0 McFarland standard was diluted 15-fold in MHB, and 75 µL of inoculum was added to the wells treated with antibacterials, resulting in a total volume of 150 µL. The microtiter plates were covered, then shake-incubated overnight at 150 rpm and 37°C. MIC was determined based on visible bacterial growth. In cases of antibacterial opacity or ambiguous results, streaking of well cultures on Mueller-Hinton agar (MHA), overnight incubation, and subsequent comparison to controls were performed. MBC was

determined by transferring 3 μ L of culture from all MIC microtiter plate wells to 147 μ L of Mueller-Hinton broth in fresh plates. These plates were shake-incubated overnight at 150 rpm and 37°C, and bacterial growth was visually inspected.

Inhibition of Biofilm formation

Minimum biofilm inhibitory concentration (MBIC) determination coincided with MIC and MBC assessments. The Calgary Biofilm Device (CBD) was employed to cover the 150 μ L wells of the MIC microtiter plate. Following overnight incubation, the MIC plates and pegs underwent three washes with 250 μ L of distilled water (dH₂O) to eliminate planktonic bacteria. Biofilms were then stained with 200 μ L of a 0.1% crystal violet solution for 30 minutes. Post-staining, microplates and pegs were washed three times with 200 μ L of ddH₂O to remove excess dye. Biofilm quantification involved sonication using a 250HT ultrasonic cleaner (VWR International), set at 60 Hz for 10 minutes, into 200 μ L of 70% ethanol. Absorbance readings at 600 nm were taken, with 70% ethanol serving as the blank (1).

Membrane Permeability Measurements

For assessing membrane permeability, propidium iodide (PI) (Invitrogen, Eugene, Oregon, USA) served as the fluorescent reporter dye. Increased PI fluorescence readings are indicative of heightened membrane disruption and permeability, as PI can penetrate cells, bind to DNA, and remain within the cells (24, 25). Bacteria were cultured in 3 mL of Mueller-Hinton broth (MHB) and incubated at 37 °C for approximately 3 hours in a shaker incubator (150 rpm) until reaching an OD₆₀₀ of 0.08. Each group received treatment with MIC concentrations of agents, the untreated groups with phosphate-buffered saline (PBS) as a negative control, and a positive control involving bacteria subjected to freezing and boiling three times (frozen for 5 minutes at -80 °C and incubated at 90 °C for 10 minutes) to disrupt the bacterial cell membrane. All treated and control groups were incubated at 37 °C for 1 hour in a shaker incubator (150 rpm). Ciprofloxacin and gentamycin were employed as internal controls, as they are known not to target the bacterial cell membrane in the initial steps. After incubation, the samples were centrifuged, washed with PBS (10,000 rpm for 2 minutes), and bacteria were stained with 0.08

mM PI for 5 minutes at 21 °C in the dark. Subsequently, 10 µL of samples were transferred onto slides and examined under a fluorescence microscope (Zeiss Axio Imager Z1) using the same exposure time (640 ms red, 1s green). Densitometry analysis was conducted using Fiji software (ImageJ).

Hydrogen Peroxide Assay

The hydrogen peroxide concentration after exposure with our agents was detected with Pierce Quantitative Peroxide Assay Kit, in the aqueous-compatible formulation according to the manufacturer's instructions (26). For preparing the standard, 1mM solution of H₂O₂ was initially made by diluting a 30% H₂O₂ stock 1:9000 (11µL of 30% H₂O₂ into 100mL of double-distilled (DD) water. This sample was then serially diluted with DD water 1:2 (100µL of DD water + 100µL of the previous dilution) for a total of 11 samples as a standard. 200µL of the working reagent (WR) from the kit was added to 20µL of the diluted H₂O₂ standards. Samples were mixed and incubated for 15 minutes at 21 °C in the dark. Absorbances were measured at 595nm using a Thermomax microtiter plate reader with Softmax Pro data analysis software (Molecular Devices, Sunnyvale, CA).

For measuring the treated and untreated samples. The bacteria were cultured in 2mL of MHB and were incubated at 37 °C for ~ 3 h in a shaker incubator (150 rpm) to reach the OD₆₀₀ of 0.08. Then treated with MIC concentrations of agents and untreated groups with PBS as a negative control, the positive control was treated with 250µM H₂O₂, and incubated at 37 °C for 1 h in a shaker incubator (150 rpm). Ciprofloxacin and gentamycin were used as antibiotic comparators. The bacterial cells were washed with PBS by centrifuging (10,000 rpm for 5 min) and discarding the supernatant. 2mL PBS was added to each sample and vortexed. 200µL of the WR was added to 20µL of each sample. Samples were mixed and incubated for 15 minutes at room temperature. Absorbances were measured at 595nm using a Thermomax microtiter plate reader with Softmax Pro data analysis software (Molecular Devices, Sunnyvale, CA). The blank value was subtracted from all sample measurements. The samples' H₂O₂ concentrations were calculated based on standard curve $R^2 = 0.93$ value.

Iron Detection Ferene-S Assay

The release of Fe^{2+} from the iron-sulfur clusters in *P. aeruginosa* and *MRSA* was detected using a Ferene-S assay with the probe, 3-(2pyridyl)-5,6-bis(2-(5-furysulfonic acid))-1,2,4-triazin (Sigma-Aldrich, St Louis, MO, USA) (27). The 10 mL of bacteria (OD600 of 0.08) were prepared in Tris-HCl buffer (20 mM, pH 7). The bacterial cells were washed with the same buffer by centrifuging (10,000 rpm for 5 min) and discarding the supernatant. The platelet (bacterial cells) was then lysed by sonication using a 250HT ultrasonic cleaner (VWR International), set at 60 Hz for 20 min in the same buffer. The samples were centrifuged (10,000 rpm for 5 min) and the supernatant was collected. The solution was treated with MIC concentration of agents, untreated control (PBS) and positive control (incubated in 90 °C for 10 min to break down the Fe-sulfur cluster). Then, a 10 mM Ferene-S probe was added to each sample in a 96 well plate, and samples were incubated at 21 °C in dark for 1 h. Absorbance was measured at 600 nm, using a Thermomax microtiter plate reader with Softmax Pro data analysis software (Molecular Devices, Sunnyvale, CA) (24).

Reduced thiol (RSH) assay

The accuracy of the assay was assessed with a standard dilution of reduced glutathione ($\geq 98\%$, Alfa Aesar, Germany) and oxidized glutathione (Sigma, USA). For preparing the standard, 1mM solution of each glutathione was serially diluted with 50 mM Tris/HCl pH 8.1:2 (150 μL of 50 mM Tris/HCl pH 8 + 150 μL of glutathione) for a total of 11 samples as a standard. Then 0.1 mM of Ellman's reagent 5,5'-dithiobis(2-nitrobenzoic acid) (DTNB) was added to each well. Samples were mixed and incubated for 30 minutes at 37 °C in dark. Absorbances were measured at 412nm using a Thermomax microtiter plate reader with Softmax Pro data analysis software (Molecular Devices, Sunnyvale, CA).

For measuring the treated and untreated samples with agents. The bacteria were cultured in 3mL of MHB and were incubated at 37 °C for ~ 3 h in a shaker incubator (150 rpm) to reach the OD600 of 0.08. Then treated with MIC concentrations of agents and untreated groups with PBS (as a negative control) and incubated at 37 °C for 1 h in a shaker incubator (150 rpm). The bacterial cells were washed with PBS by centrifuging (10,000 rpm for 5 min) and discarding the

supernatant. 1mL of 50 mM Tris/HCl pH 8.0, 5 mM EDTA, 0.1% SDS and 0.1 mM DTNB was added to each sample and vortexed. These cell suspensions were incubated at 37°C for 30 min, and then centrifuged in a microfuge for 1 min at 15000 g. Absorbances were measured at 412nm using a Thermomax microtiter plate reader with Softmax Pro data analysis software (Molecular Devices, Sunnyvale, CA). The blank value was subtracted from all sample measurements. The absorption coefficient of oxidized DTNB ($1.36 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$) at this wavelength was then used to calculate the RSH concentration of the cell.

Eukaryotic cell toxicity of pentafluorosulfanyl-containing triclocarban compounds

Five Eukaryotic cell lines were used to determine the cytotoxicity of the compounds: human embryonic lung (HEL) fibroblast cells; human cervixcarinoma-derived HeLa cells; human T-cell leukemia-derived MT4 cells; Madin-Darby canine kidney (MDCK) cells; and African Green monkey kidney-derived VERO cells. Semi-confluent cell cultures in 96-well plates were exposed to serial dilutions of the compounds or to medium (= no compound control), then incubated at 37 °C. Four days later, the cells were inspected by microscopy to determine the Minimum Cytotoxic Concentration (MCC), i.e. compound concentration that causes a microscopically detectable alteration of normal cell morphology. Next, the MTS cell viability reagent (CellTiter 96 AQueous MTS Reagent from Promega) was added. After 4 h incubation at 37 °C, optical density (OD) at 490 nm was recorded in a microplate reader. The percentage cytotoxicity was calculated as: $[1 - (\text{OD}_{\text{Cpd}})/(\text{OD}_{\text{Contr}})] \times 100$, after which the 50% cytotoxicity value (CC_{50}) was derived by extrapolation, assuming semi-log dose response.

Bacterial infection and mouse treatment

Animal experiments were performed with 6-10 week/old adult male mice. Mice were kept in a pathogen-free facility under standardized conditions: temperature of 21-22 °C, illumination of 12h light-12h dark, and access to tap water and food. Experimental animal protocols were approved by the University of Calgary Animal Care Committee and followed the Canadian Council for Animal Care Guidelines.

MW2 *Staphylococcus aureus* were grown on brain heart infusion (BHI) agar plates (BD biosciences). A single colony was grown overnight in BHI medium (BD biosciences) at 37 °C while shaking. Subcultures of 100 ml of the ON culture were grown in 3 ml of the same culture medium for 2h at 37 °C while shaking to reach the exponential growth phase. Bacteria were brought to a concentration of OD₆₆₀ 1.0 in saline. A 1:4 dilution of bacteria was prepared, and animals were infected through an intravenous catheter with 200 µl of the bacterial suspension, so that bacterial dose was controlled to a dose of 5x10⁷ bacteria. Mice were treated with and intraperitoneal injection of 100 µl of compound of interest (5 mM) 1 hour before infection or 1 hour after infection. Untreated group was injected 1-hour post-infection with 100 µl saline.

For assessing bacterial by CFU enumeration, the mice were anesthetized with isoflurane (Fresenius Kabi) and washed with 70% ethanol to collect blood, and sacrificed to further collect liver, spleen, kidney (left), heart, and lungs. Blood was collected by cardiac puncture and pooled with 40 ml heparin. 25 ml of 1:10 diluted blood samples were plated. Organ samples were weighed and homogenized in 1 ml PBS and serially diluted (spleen, liver, and lung samples 1:10, 1:100; 1:1000; 1:10 000; kidney and heart samples 1:100; 1:1000; 1:10 000; 1:100 000). 30 ml of each dilution was plated on a 30° angle to form stripes. All plates were incubated ON at 37 °C. Cultures were counted, and colony-forming units (CFUs) were calculated taking the weight of the organ into account.

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Conceived and designed the study: AP, MVC, RJT
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Analyzed the data: AP, MM, JWH, MVC, RJT

407 Wrote the paper: **AP, MM, MVC, RJT**
 408 Participated in data analysis and manuscript editing: **AP, MM, JWH, EP, LN, SV, BGS, MZ, MVC, RJT**
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 410 version of the manuscript.
 411
 412 **Conflicts of Interest:** The authors declare no competing financial interests.

413 **Table1.** Minimum inhibitory concentration (MIC), Minimum bactericidal concentration (MBC) and
 414 Minimum biofilm inhibitory concentration (MBIC) of diphenyl-urea antibiotics for *P. aeruginosa*, *S.*
 415 *aureus* and *E. coli*.

		Agents	<i>E. coli</i>	<i>P. aeruginosa</i>	<i>K. pneumonia</i>	<i>P. mirabilis</i>	<i>S. aureus</i>	<i>MRSA</i>	<i>S. epidermidis</i>
MIC (mM)	diphenyl-urea	EPB-21	160	310	>5	>5	1250	2500	U/A
		EPB-22	310	310	>5	>5	0.61	0.61	U/A
		EPB-23	310	310	>5	>5	2.4	4.9	U/A
		EBP-24	310	160	>5	>5	0.076	0.15	U/A
		EBP-25	310	U/A	>5	>5	9.8	U/A	U/A
		EBP-33	310	620	>5	>5	0.019	0.0048	U/A
		EBP-34	310	620	>5	0.625	0.0038	0.0019	0.0002
		EBP-36	310	310	>5	>5	0.076	0.61	U/A
		EBP-37	310	310	>5	0.625	0.0012	0.008	0.008
		EBP-40	310	310	>5	0.625	0.48	0.15	1.2
		EBP-41	310	310	>5	>5	0.038	0.0095	U/A
		EBP-53	310	310	>5	>5	1.2	1.2	U/A
		EBP-55	310	310	>5	>5	0.019	0.0048	U/A
		EBP-56	310	310	>5	>5	4.9	0.30	U/A
		EBP-58	310	310	>5	>5	620	20	U/A
		EBP-59	310	310	>5	0.002	< 0.0003	0.0024	0.004
		EPB-61	310	310	>5	>5	0.0095	0.16	0.001
		EBP-62	310	310	>5	>5	0.0003	0.038	0.02
		A	310	310	>5	>5	2500	310	U/A
		B	310	310	>5	>5	9.8	4.9	U/A
	Antibiotics	Cip	0.002	0.061	0.01	0.02	0.08	0.160	1.2
		Gen	0.01	0.012	0.001	0.001	0.04	0.98-0.39	0.3
MBC (mM)	diphenyl-urea	EPB-21	>2500	1250	>5	>5	> 2500	>2500	U/A
		EPB-22	>2500	2500	>5	>5	20	4.9	U/A
		EPB-23	>2500	>2500	>5	>5	20	39	U/A
		EBP-24	>2500	1250	>5	>5	78	2500	U/A
		EBP-25	>2500	U/A	>5	>5	78	U/A	U/A
		EBP-33	>2500	1250	>5	>5	20	2.4	U/A
		EBP-34	>2500	1250	>5	0.625	0.01	0.061	0.01
		EBP-36	2500	310	>5	>5	78	9.8	U/A
		EBP-37	2500	310	>5	0.625	4.9	0.04	0.04
		EBP-40	2500	310	>5	0.625	1.2	0.61	0.9
		EBP-41	2500	310	>5	>5	1.2	1.2	U/A
		EBP-53	2500	310	>5	>5	1250	2500	U/A
		EBP-55	2500	310	>5	>5	78	0.15	U/A
		EBP-56	2500	310	>5	>5	78	0.61	U/A
		EBP-58	310	>2500	>5	>5	2500	620	U/A
		EBP-59	310	1250	>5	0.002	0.0061	0.02	0.01
		EPB-61	310	>2500	>5	>5	0.095	0.30	0.04

	Antibiotics	EBP-62	310	>2500	>5	>5	0.30	0.076	0.1
		A	310	1250	>5	>5	>2500	620	U/A
		B	310	>2500	>5	>5	620	160	U/A
		Cip	0.002	2.4	0.5	0.8	0.08	0.160	1.2
		Gen	0.01	1.2	0.05	0.05	0.04	0.98-0.39	0.3
MBIC (mM)	diphenyl- urea	EPB-21	310	1250	>5	>5	620	>2500	U/A
		EPB-22	310	1250	>5	>5	4.9	20	U/A
		EPB-23	310	1250	>5	>5	4.9	39	U/A
		EBP-24	310	1250	>5	>5	4.9	2500	U/A
		EBP-25	310	U/A	>5	>5	160	u/A	U/A
		EBP-33	310	1250	>5	>5	0.15	0.15	U/A
		EBP-34	310	1250	>5	0.625	0.30	0.076	0.001
		EBP-36	160	1250	>5	<5	78	1.2	U/A
		EBP-37	>2500	1250	>5	0.625	0.0095	<0.0003	0.001
		EBP-40	>2500	1250	>5	0.625	0.15	0.61	0.9
		EBP-41	>2500	1250	>5	>5	0.30	0.30	U/A
		EBP-53	>2500	1250	>5	>5	310	1250	U/A
		EBP-55	>2500	1250	>5	>5	0.30	0.15	U/A
		EBP-56	>2500	1250	>5	>5	9.8	1.2	U/A
		EBP-58	310	1250	>5	>5	2500	39	U/A
		EBP-59	310	1250	>5	0.002	0.019	0.30	0.002
		EPB-61	310	1250	>5	>5	0.0095	0.30	0.001
		EBP-62	310	1250	>5	>5	0.38	0.038	0.01
		A	310	1250	>5	>5	>2500	620	U/A
		B	310	1250	>5	>5	20	9.8	U/A
	Antibiotics	Cip	0.002	2.4	0.01	0.02	0.08	0.160	1.2
		Gen	0.01	1.2	0.001	0.001	0.04	0.98-0.39	0.3

416

417 3-(1,2,3-benzothiadiazol-6-yl)-1-(4-pentafluoro- λ^6 -sulfanyl)urea = EPB-21, 3-(1,2,3-benzothiadiazol-6-yl)-1-(3-pentafluoro- λ^6 -
418 sulfanyl)urea = EPB-22, 3-(1,2,3-benzothiadiazol-6-yl)-1-(2-chloro-5-pentafluoro- λ^6 -sulfanyl)urea = EPB-23, 3-(1,2,3-
419 benzothiadiazol-6-yl)-1-(2-chloro-3-pentafluoro- λ^6 -sulfanyl)urea = EPB-24, 3-(1,2,3-benzothiadiazol-6-yl)-1-(4-chloro-3-
420 pentafluoro- λ^6 -sulfanyl)urea = EBP-25, 1,3-bis(4-pentafluoro- λ^6 -sulfanyl)urea = EBP-33, 1,3-di(4-chloro-3-
421 (trifluoromethyl)phenyl)urea = EBP-34, 1,3-bis(2-chloro-5-pentafluoro- λ^6 -sulfanyl)urea = EBP-36, 3-(4-chloro-3-
422 (trifluoromethyl)phenyl)-1-(2-chloro-5-pentafluoro- λ^6 -sulfanyl)urea = EPB-37, 1-(2-Chloro-5-(pentafluoro- λ^6 -sulfanyl)phenyl)-3-
423 (4-chlorophenyl)urea = EPB-40, 1-(4-chloro-3-(pentafluoro- λ^6 -sulfanyl)phenyl)-3-(4-chlorophenyl)urea = EPB-41, 1,3-bis(3-
424 (Pentafluoro- λ^6 -sulfanyl)phenyl) urea= EPB-53, 1-(3-(pentafluoro- λ^6 -sulfanyl)phenyl)-3-(4-(pentafluoro- λ^6 -sulfanyl)phenyl)urea
425 = EPB-55, 1,3-bis(4-Chloro-3-(pentafluoro- λ^6 -sulfanyl)phenyl) urea = EPB-56, 1-(2-Chloro-3-(pentafluoro- λ^6 -sulfanyl)phenyl)-3-
426 (4-chlorophenyl)urea = EPB-58, 1-(3,4-dichlorophenyl)-3-(4-pentafluoro- λ^6 -sulfanyl)phenyl) urea= EPB-59, 1-(4-Chlorophenyl)-3-
427 (4-(pentafluoro- λ^6 -sulfanyl)phenyl)urea = EPB-61, 1-(4-chlorophenyl)-3-(3-(pentafluoro- λ^6 -sulfanyl)phenyl)urea = EPB-62,
428 Ciprofloxacin = Cip, Gentamicin = Gen, Ampicillin = Amp MIC=minimum inhibitory concentration, MBC=Minimum bactericidal
429 concentration, MHB= Mueller Hinton Broth, MBIC = Minimum biofilm inhibitory concentration.

430 The **bold values** show the components are antibacterial/anti-biofilm in lawyer concentration in
431 comparison with antibiotics.

432

433

434 **Table2.** Minimum inhibitory concentration (MIC), Minimum bactericidal concentration (MBC) and
 435 Minimum biofilm inhibitory concentration (MBIC) of diphenyl-urea antibiotics against *MRSA* clinical
 436 isolates.

MRSA clinical isolates			<i>MRSA</i> #1	<i>MRSA</i> #2	<i>MRSA</i> #3	<i>MRSA</i> #4	<i>MRSA</i> #5	<i>MRSA</i> #6	<i>MRSA</i> #7	<i>MRSA</i> #8	<i>MRSA</i> #9	<i>MRSA</i> #10
MIC (mM)	diphenyl-urea	EBP-34	0.6	< 0.002	0.08	< 0.002	0.01	0.01	< 0.002	0.005	< 0.002	< 0.002
		EBP-37	0.01	< 0.002	0.08	< 0.002	0.01	< 0.002	0.01	< 0.002	0.005	< 0.002
		EBP-40	0.02	0.02	0.08	0.04	0.02	0.04	0.04	0.04	0.02	0.04
		EBP-59	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002
		EPB-61	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002
		EBP-62	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	0.005	0.005	< 0.002	0.005
	Antibiotics	Cip	0.4	0.05	0.4	0.05	0.2	0.2	0.2	1.6	0.4	0.05
		Gen	0.1	0.05	0.05	0.05	0.8	3	0.05	0.05	0.05	0.05
MBC (mM)	diphenyl-urea	EBP-34	< 0.002	< 0.002	0.08	< 0.002	0.01	0.01	0.01	0.005	< 0.002	0.01
		EBP-37	1.2	0.01	0.6	0.01	0.01	< 0.002	0.04	0.01	0.01	0.01
		EBP-40	1.2	0.04	0.16	0.16	0.04	0.16	0.08	0.08	0.08	0.01
		EBP-59	0.005	< 0.002	< 0.002	0.005	0.01	< 0.002	< 0.002	0.005	0.02	0.04
		EPB-61	0.04	0.01	0.01	0.02	0.04	0.02	0.02	0.04	0.04	0.02
		EBP-62	0.08	0.005	< 0.002	0.02	0.02	0.005	0.01	0.08	0.04	0.08
	Antibiotics	Cip	1.6	0.4	0.05	0.4	0.4	0.4	0.05	12.5	1.6	0.8
		Gen	1.6	0.2	0.05	0.05	6.25	25	1.6	0.05	0.05	0.05
MBIC (mM)	diphenyl-urea	EBP-34	1.2	< 0.002	0.08	< 0.002	0.01	0.01	0.02	0.005	< 0.002	< 0.002
		EBP-37	0.02	< 0.002	0.3	0.01	0.01	< 0.002	0.01	< 0.002	< 0.002	< 0.002
		EBP-40	< 0.002	< 0.002	0.08	< 0.002	0.02	0.04	< 0.002	< 0.002	< 0.002	< 0.002

		EBP-59	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002
		EPB-61	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002
		EBP-62	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002	< 0.002
	Antibiotics	Cip	0.05	0.05	0.2	0.05	0.05	0.2	0.05	0.05	0.05	0.05
		Gen	0.05	0.05	0.05	0.05	0.1	12.5	0.05	0.05	0.05	0.05

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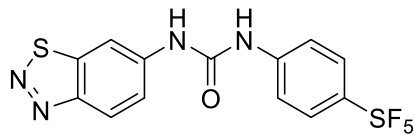
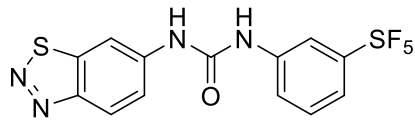
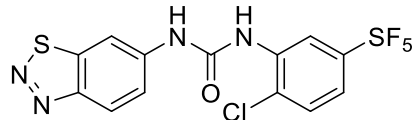
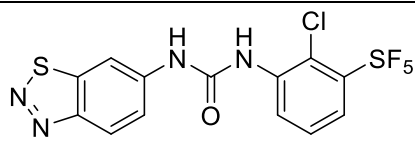
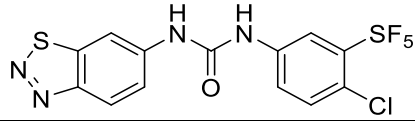
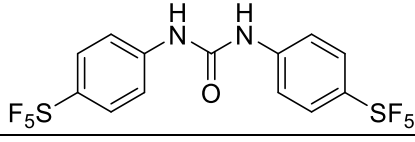
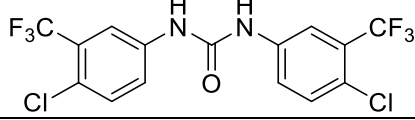
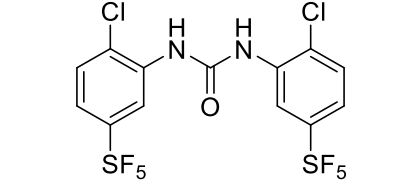
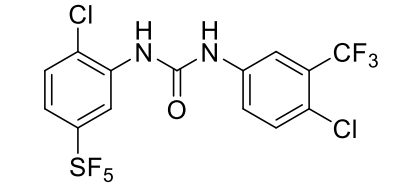
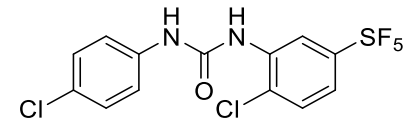
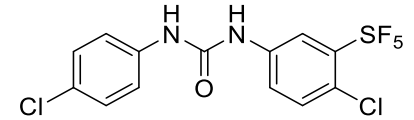
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Table 3. Eukaryotic cell Toxicity of different diphenyl-urea analogs.

	Cell line (in vitro cytotoxicity, μ M)				
	Human			Mouse	Monkey
	HEL ^a (MCC) ^d	HeLa ^b (MCC) ^d	MT4 ^c (MCC) ^d	MDCK ^e (CC ₅₀) ^g	Vero ^f (CC ₅₀) ^g
EBP-59	>4	4	3.9	>100	>4
EBP-61	>4	4	2.8	>100	>4
EBP-62	20	>4	4.7	100	>4

^aHEL human leukemia cells. ^bHeLa human cervical cancer cells. ^cMT4 human T leukemia cells. ^dMCC: minimum compound concentration that causes a microscopically detectable alteration of normal cell morphology. ^eMDCK: Madin-Darby canine kidney cells. ^fVero: Green monkey kidney cells. ^g50% Cytotoxic concentration, as determined by measuring the cell viability with the colorimetric formazan-based MTS assay. Values shown are the mean of 2 or 3 determinations.

449 **Figures:**

Chemical structure	Compound's reference (EPB)
	EPB-21
	EPB-22
	EPB-23
	EPB-24
	EPB-25
	EPB-33
	EPB-34
	EPB-36
	EPB-37
	EPB-40
	EPB-41

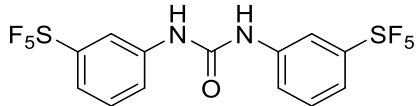
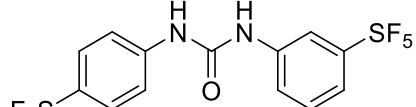
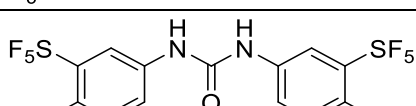
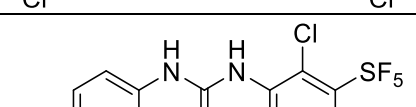
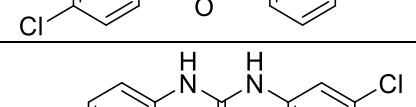
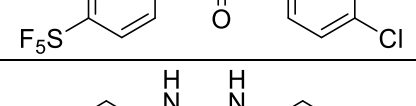
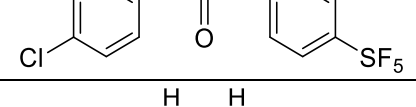
	EPB-53
	EPB-55
	EPB-56
	EPB-58
	EPB-59
	EPB-61
	EPB-62

Fig1. Compounds EPB-21, EPB-22, EPB-23, EPB-24, EPB-25, EPB-36 (12) , compounds EPB-33, PB-34, EPB-37, EPB-40, EPB-41, EPB-55, EPB-56, EPB-58, EPB-59, EPB-61, EPB-62 (5, 12) and EPB-53 have been published in our previous works (5, 28).

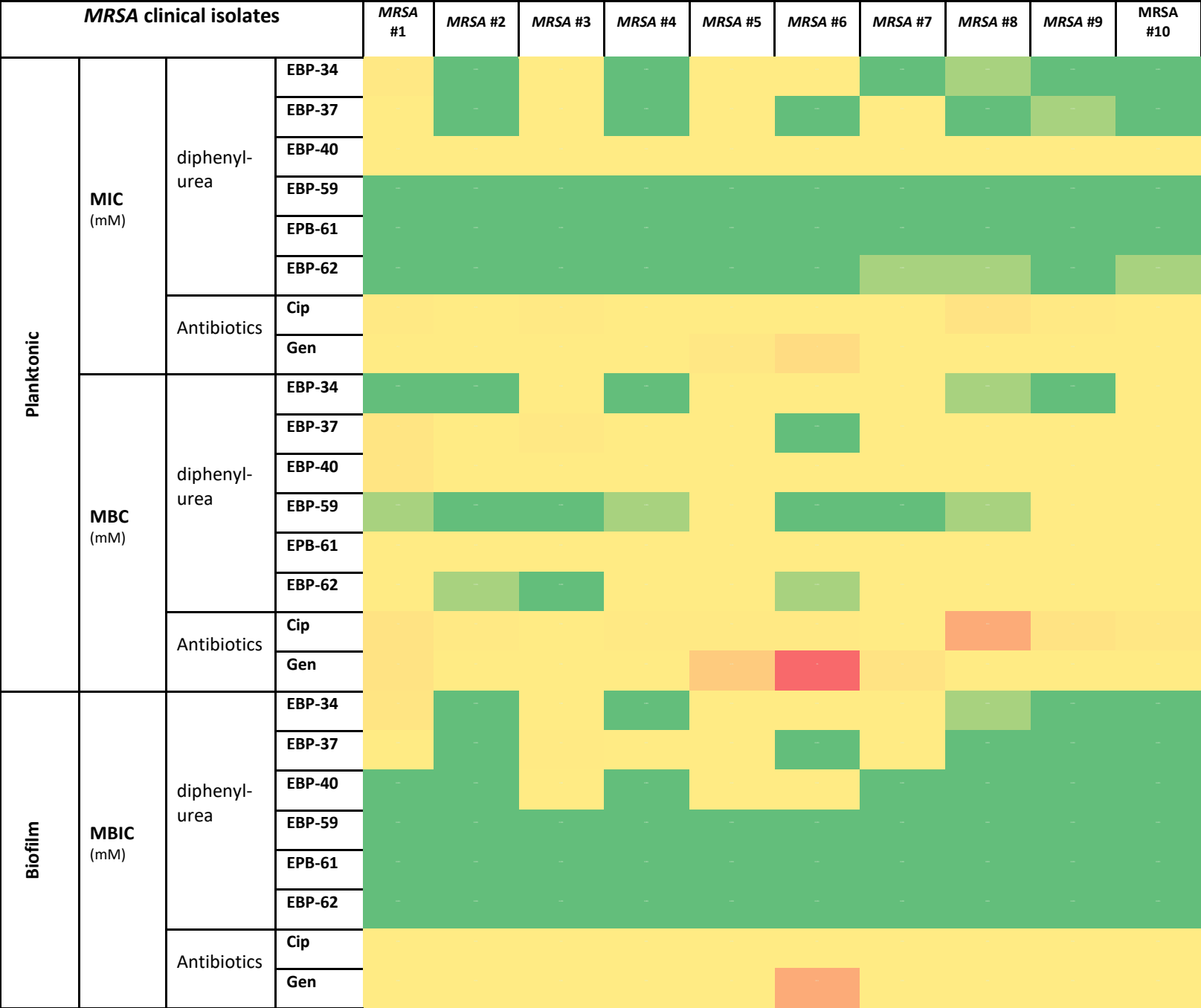


Fig 2. Minimum inhibitory concentration (MIC), Minimum bactericidal concentration (MBC), and Minimum biofilm inhibition concentration (MBIC) of diphenyl-urea antibiotics against MRSA clinical isolates.

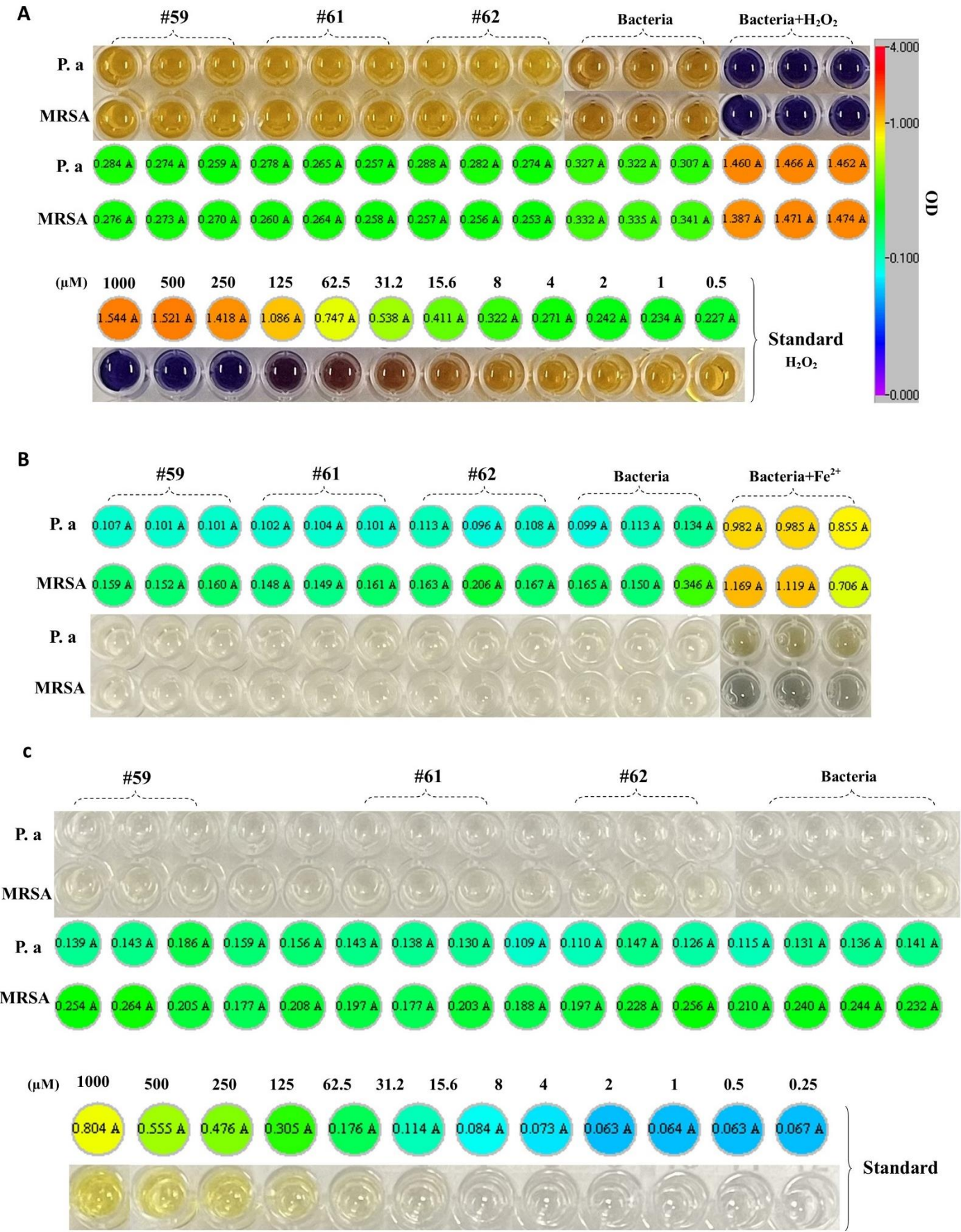


Fig 3. A. Hydrogen peroxide (H₂O₂) concentration. Shows the H₂O₂ concentrations in the samples with the naked eye (down) and plate reader (upper). The standard, 1mM solution of hydrogen peroxide serially diluted with double distilled (DD) water 1:2 for a total of 11 samples. DD water was used as the blank and working reagent (WR). **B. Free iron [(ferrous) Fe⁺²] concentration.** After 1 h treatment with selected components relative amount of ferrous measured by OD. The naked eye (upper) Heatmap (down) is illustrated in the panels. **C. Reduced thiol (RSH) level.** Shows the RSH absorbance values in the samples with the naked eye (upper) and plate reader (down). 1mM solution of glutathione reduced oxidized serially diluted with Tris/HCl 1:2 for a total of 11 samples as a standard. The 50 mM Tris/HCl pH 8 and Ellman's reagent were used as the blank.

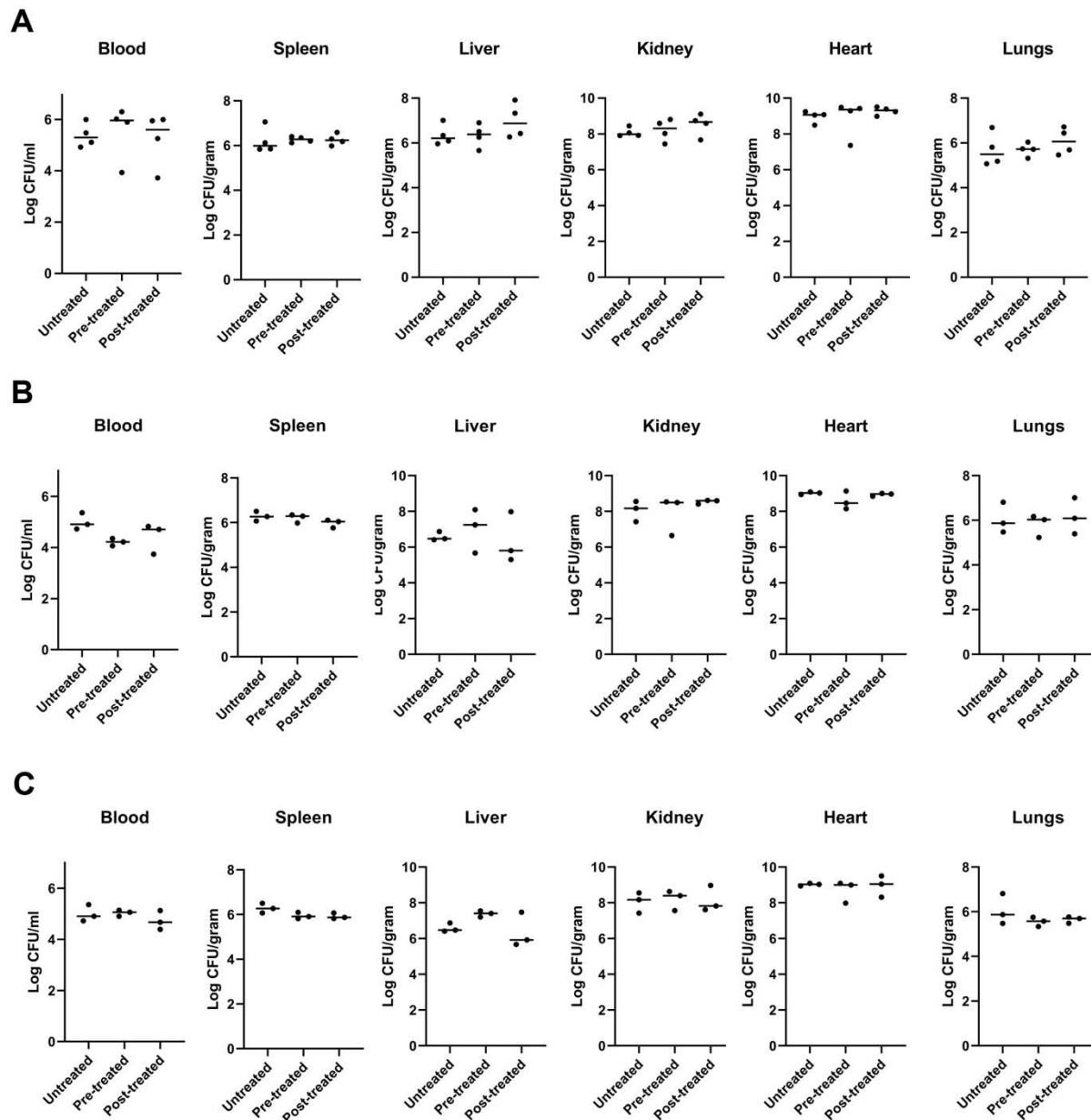


Fig 4. Intravenous MRSA Infection model to screen for *In vivo* effectivity of compound-59, 61, and 62. CFU enumerations for blood, spleen, liver, kidney (left), heart, and lungs after 24 hours of infection with *S. aureus* MW2. Mice were left untreated or were treated 1 hour before infection (pre-treated) or 1 hour after infection (post-treated) with compound **59** (A), **61** (B), or **62** (C). One-way ANOVA with Bonferroni's posttest, data not significant.

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