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Molecular design of antimicrobial conjugated oligoelectrolytes with enhanced selectivity toward bacterial cells†

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A series of cationic conjugated oligoelectrolytes (COEs) was designed to understand how variations in molecular dimensions impact the relative activity against bacteria and mammalian cells. These COEs kept a consistent distyrylbenzene framework but differed in the length of linker between the core and the cationic site and the length of substitute on the quaternary ammonium functional group. Their antimicrobial efficacy, mammalian cell cytotoxicity, hemolytic activity, and cell association were determined. We find that hydrophobicity is a factor that controls the degree of COE association to cells, but *in vitro* efficacy and cytotoxicity depend on more subtle structural features. **COE2-3C-C4butyl** was found to be the optimal structure with a minimum inhibitory concentration (MIC) of 4 $\mu\text{g mL}^{-1}$ against *E. coli* K12, low cytotoxicity against HepG2 cells and negligible hemolysis of red blood cells, even at 1024 $\mu\text{g mL}^{-1}$. A time-kill kinetics study of **COE2-3C-C4butyl** against *E. coli* K12 demonstrates bactericidal activity. These findings provide the first systematic investigation of how COEs may be modulated to achieve low mammalian cell cytotoxicity with the long-range perspective of finding candidates suitable for developing a broad-spectrum antimicrobial agent.

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Introduction

Antimicrobial resistance is expected to cause up to 10 million deaths per year by 2050, along with a massive economic burden.^{1,2} Indeed, learned opinions have placed the problem at par with possible damage by climate change.^{3–5} Despite the foreseeable crisis, there have been few new antibiotics introduced to the market, in part because of the long drug development cycle, rapid acquisition of drug resistance, and poor investment return.^{6,7} Studies of new antimicrobial compounds at a basic research level are thus warranted.^{8,9}

Different approaches are being considered for designing new antibiotics that target different machineries in bacterial cells. Examples include cell wall synthesis,^{10,11} DNA and RNA synthesis,^{12,13} protein biosynthesis,¹⁴ folic acid metabolism,¹⁵ and the cell membrane.^{16,17} Compounds that target cell membranes typically kill bacterial cells by selectively disrupting bacterial membranes over mammalian cell membranes due to difference in lipid compositions between the two cell types.¹⁸

Potential advantages of this mechanism include that bacteria are known to be less likely to develop resistance to membrane-disrupting compounds^{19,20} and its potential to eradicate persistent infections in which bacteria are in dormant states.²¹

Conjugated oligoelectrolytes (COEs) are a class of synthetic water-soluble conjugated molecules containing a conjugated core and pendant ionic groups. COEs have been reported to spontaneously interact with lipid bilayers and modify the properties of bilayers.^{22,23} The extent of membrane perturbation can be fine-tuned *via* molecular design approaches.^{24,25} A general guiding principle is that the mismatch between the bilayer thickness and length of the COE leads to membrane disruption.^{26,27}

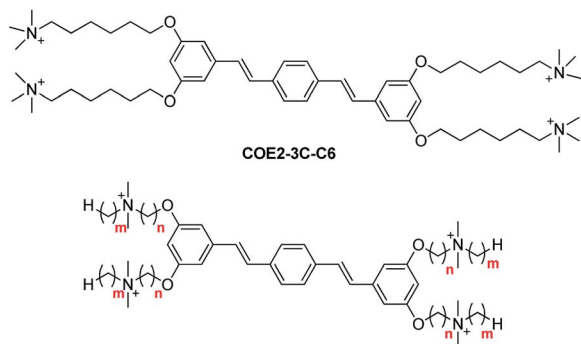
In particular, the distyrylbenzene derivative **COE2-3C-C6** (previously referred to as **COE2-3C**, see Scheme 1) was reported to have the lowest minimum inhibitory concentration (MIC) against *E. coli* K12 among a homologous series of structures that differed in the number of phenylenevinylene units.²⁵ We also recently reported a subsequent study that illustrated the relationship between antimicrobial activity and the hydrophobicity of stilbene-based COEs with an apparent correlation between increased hydrophobicity and increased antimicrobial potency.²⁸ Up to now, the cytotoxicity of COEs against mammalian cells has yet to be explored despite its importance in determining suitable chemical structures for drug development.²⁹

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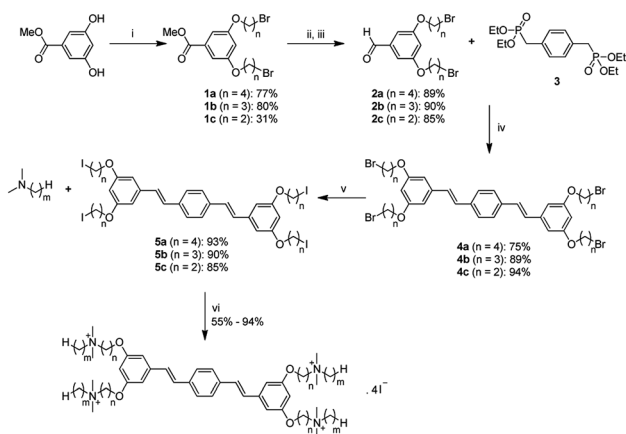
Scheme 1 Chemical structure of COE2-3C-C6 (top) and general structure of COEs in this study (bottom).

With prior information in hand, we describe herein a series of COEs based on the distyrylbenzene core that is generated by modulating two molecular features: linker length (parameter n) and terminal alkyl chains length (parameter m), see Scheme 1. We examined antimicrobial efficacy and mammalian cell cytotoxicity. Bacterial selectivity was determined and the optimal compound in the series was identified thereafter.

Results and discussion

Synthetic procedures

The general synthetic approach to generate the new series of COEs is shown in Scheme 2. Aldehyde precursors were first synthesized by alkylation of methyl 3,5-dihydroxybenzoate with α,ω -dibromoalkanes. Isolated aryl ether products **1a–1c** were subsequently reduced by using diisobutylaluminium hydride (DIBAL) and then oxidized with MnO_2 to obtain the aldehyde derivatives **2a–2c**. Next, the aldehydes were reacted with compound **3** via Horner–Wadsworth–Emmons (HWE) reaction to obtain distyrylbenzene intermediates **4a–4c**. Finkelstein halide exchange reactions were then used for bromide/iodide



Scheme 2 Preparation of COEs reported in this study: (i) α,ω -dibromoalkanes, K_2CO_3 , acetone, reflux, 2 d; (ii) DIBAL, THF, -78°C to rt, overnight; (iii) MnO_2 , DCM, rt, overnight; (iv) NaOtBu , THF, -78°C to rt, overnight; (v) NaI , acetone, reflux, 2 d; (vi) DMF, 45°C , 2 d.

exchange, thereby yielding neutral intermediates **5a–5c**. Finally, **5a–5c** underwent quaternization reactions with various tertiary amines to obtain target molecules. COE2-3C-C3propyl ($n = 3$, $m = 3$) and COE2-3C-C4propyl ($n = 4$, $m = 3$) were synthesized *via* a slightly different pathway (see ESI†). Abbreviations of the COEs are listed in Table 1.

Antimicrobial efficacy

The antimicrobial efficacies of COEs were screened by determining minimum inhibitory concentrations (MICs) against Gram-negative bacteria *E. coli* K12 (Fig. 1a). When the terminal alkyl chains, *i.e.* N substituents, are short ($m \leq 2$), one observes that the MIC gradually increased (less potent antibiotics) in response to decreasing linker length (parameter n). For example, at $m = 1$, the MIC values increase from $16 \mu\text{g mL}^{-1}$ for COE2-3C-C6 ($n = 6$, $m = 1$) to $128 \mu\text{g mL}^{-1}$ for COE2-3C-C2 ($n = 2$, $m = 1$). Similarly, for $m = 2$, MIC increases from $16 \mu\text{g mL}^{-1}$ for COE2-3C-C4ethyl ($n = 4$, $m = 2$) to $64 \mu\text{g mL}^{-1}$ for COE2-3C-C2ethyl ($n = 2$, $m = 2$). Upon elongation of the terminal alkyl chains, the MIC is largely insensitive relative to linker length. As seen in Fig. 1a, COEs with N-substituents of butyl or longer ($m \geq 4$) have similar antimicrobial efficacies, with MICs around $4 \mu\text{g mL}^{-1}$. This similarity of the MICs suggests that effects from the terminal alkyl chains dominate over the influence by the linkers. Therefore, for this limited set of compounds, COEs with $m \geq 4$ are desirable for maintaining good antimicrobial efficacy regardless of linkers. In addition to antimicrobial activities against *E. coli* K12, MICs for a subset of COEs against another Gram-negative bacteria (*K. pneumoniae*) and two Gram-positive bacteria (*E. faecium*, and *S. aureus*) were determined (Table S1†) at Emery Pharma (Alameda, CA). In short, similar trends in MICs were observed in those bacterial strains and all these COEs showed relatively low MICs toward tested Gram-positive bacteria.

Mammalian cell cytotoxicity

Antimicrobial compounds need to have low MICs against bacteria but also low cytotoxicity toward mammalian cells. In order to determine cytotoxicity, half maximal inhibitory concentration (IC_{50}) values were determined *in vitro* against human hepatocellular carcinoma cell line (HepG2) by using the MTT assay. As shown in Fig. 1b, COEs with $m \leq 3$ show low cytotoxicity, as their IC_{50} values are higher than $1024 \mu\text{g mL}^{-1}$, the highest concentration tested. Cytotoxicity generally increases in response to an increase in m . For instance, at $n = 4$,

Table 1 Names of the synthesized COEs in this study

m	$n = 2$	$n = 3$	$n = 4$
1	COE2-3C-C2	COE2-3C-C3	COE2-3C-C4
2	COE2-3C-C2ethyl	COE2-3C-C3ethyl	COE2-3C-C4ethyl
3	—	COE2-3C-C3propyl	COE2-3C-C4propyl
4	COE2-3C-C2butyl	COE2-3C-C3butyl	COE2-3C-C4butyl
5	—	—	COE2-3C-C4pentyl
6	COE2-3C-C2hexyl	COE2-3C-C3hexyl	COE2-3C-C4hexyl



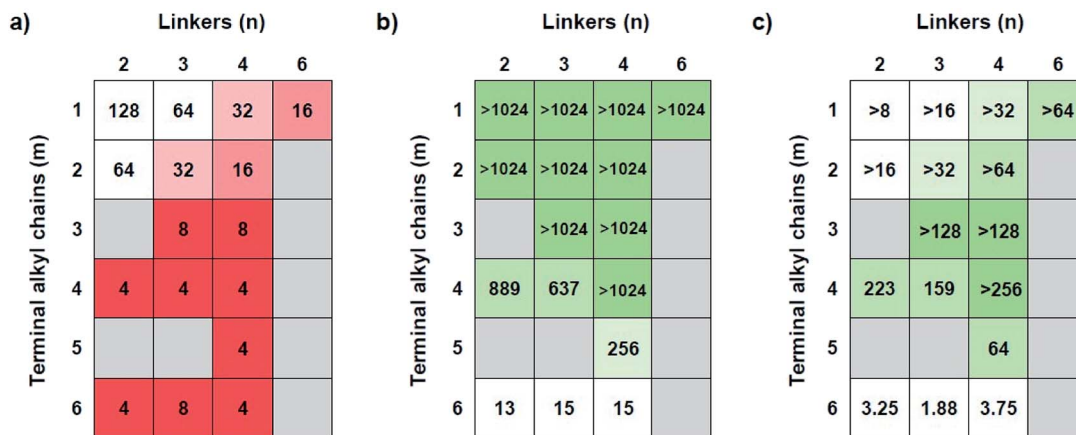


Fig. 1 (a) MICs of COEs against *E. coli* K12; (b) IC_{50} of COEs against HepG2 cell line; (c) heatmap of IC_{50}/MIC values of each COE. The greener box indicates the higher selectivity toward bacterial cells. All MIC and IC_{50} values are reported in $\mu\text{g mL}^{-1}$.

IC_{50} decreases from $>1024 \mu\text{g mL}^{-1}$ ($m = 1-4$) to $256 \mu\text{g mL}^{-1}$ and $15 \mu\text{g mL}^{-1}$ for pentyl ($m = 5$) and hexyl ($m = 6$) groups, respectively. When the terminal groups are butyl ($m = 4$), there is a slight reduction in IC_{50} values for COEs containing linkers with 3 carbon atoms or less ($n \leq 3$; $889 \mu\text{g mL}^{-1}$ and $637 \mu\text{g mL}^{-1}$ for $n = 1$ and 2, respectively). According to the trends of IC_{50} , minimizing cytotoxicity can be realized by having terminal alkyl chains that are butyl groups or shorter ($m \leq 4$). Additionally, at $m = 4$, the four-carbon linker ($n = 4$) is slightly favored due to a small increase in cytotoxicity of COEs with shorter linkers ($n = 2$ and 3).

Bacterial selectivity analysis

An optimal structure of COEs requires balance between antimicrobial properties and mammalian cell cytotoxicity. With the data in Fig. 1a and b, a ratio between IC_{50} and MIC was used to evaluate the bacterial selectivity of COEs. Higher IC_{50}/MIC values are more desirable. Fig. 1c illustrates a heatmap of IC_{50}/MIC values, showing the trend of bacterial selectivity. The heatmap shows that COEs with methyl or ethyl terminal alkyl chains ($m = 1$ and 2) exhibit low selectivity indices due to their low antimicrobial activities ($MIC \geq 16 \mu\text{g mL}^{-1}$) despite their low cytotoxicity. On the other hand, COEs containing longer terminal alkyl chains ($m = 5$ and 6) do not show promising selectivity indices due to cytotoxicity. COEs with hexyl groups ($m = 6$) have extremely low IC_{50}/MIC values at 3.25, 1.88, and 3.75 for $n = 2, 3$ and 4 respectively. **COE2-3C-C4butyl** ($n = 4, m = 4$) is shown to have highest selectivity index with IC_{50}/MIC greater than 256. Thus, it was identified as the optimal structure for this specific set of compounds.

Hemolytic activity

Hemolytic activity is an essential measure that determines suitability for *in vivo* clinical chemistry tests. Thus, the HC_{50} values (defined as a concentration at which 50% of red blood cells were lysed) of COEs were determined toward fresh CD-1 mouse red blood cells, see Table 2. In these experiments, red blood cells were collected from whole blood, washed with

phosphate buffer saline (PBS) and incubated with different concentrations of **COE2-3C-C4** analogs ($n = 4$) at 37°C for 1 hour. After incubation, the red blood cells were removed and the amount of hemoglobin in the supernatant was determined from absorption at 450 nm. Of particular interest from Table 2 is that significant hemolysis is observed only with $m = 6$, with $HC_{50} = 53 \mu\text{g mL}^{-1}$. At the maximum concentration tested ($1024 \mu\text{g mL}^{-1}$), there is no significant hemolysis ($<5\%$) of red blood cells treated with COEs with $m \leq 4$. An abrupt change in hemolytic activity of **COE2-3C-C4pentyl** ($m = 5$) and **COE2-3C-C4hexyl** ($m = 6$) suggests that there is a threshold of the length of alkyl chain termini that triggers significant hemolysis. A similar behavior was reported in membrane-active antimicrobial agents with different alkyl side chains.^{30,31} The observed hemolytic activity of **COE2-3C-C4** series generally agrees with its trend of cytotoxicity (IC_{50} , Fig. 1b, $n = 4, m = 1-6$).

Cell association

Cell association experiments were performed using bacterial and mammalian cells in order to see to what extent COE associating affinity could be correlated to antimicrobial and cytotoxicity profiles. *E. coli* K12 ($OD = 1$) and HepG2 cells (2×10^6 cells per mL) were treated at 37°C for 2 hours in Hank's balanced salt solution (HBSS) without calcium and magnesium ions. Cells were centrifuged and the amount of net unassociated COE in the buffer was determined from the absorbance at

Table 2 HC_{50} and percent hemolysis of CD-1 mouse red blood cells treated with **COE2-3C-C4** series at $1024 \mu\text{g mL}^{-1}$ in PBS

Compound	HC_{50} ($\mu\text{g mL}^{-1}$)	Hemolysis at $1024 \mu\text{g mL}^{-1}$ (%)
COE2-3C-C4	>1024	3.0 ± 0.1
COE2-3C-C4ethyl	>1024	2.3 ± 0.3
COE2-3C-C4propyl	>1024	1.6 ± 0.2
COE2-3C-C4butyl	>1024	4.9 ± 0.6
COE2-3C-C4pentyl	>1024	20.4 ± 2.7
COE2-3C-C4hexyl	53	75.8 ± 1.0



380 nm. The results of these studies are summarized in Fig. 2 and the numbers of COE molecules associated to each *E. coli* K12 and HepG2 cell were calculated (Table S2†). From Fig. 2a, one observes that for $n = 4$ there is an increase in association with *E. coli* K12 as m increases from 1 to 6. COE2-3C-C4hexyl ($n = 4$, $m = 6$) also shows the highest association to HepG2 cells. However, from $m = 1$ to 5, the associations to HepG2 cells are similar within experimental errors (approximately 20% associated). Fig. 2b provides the cell association profile for $m = 1$ and $n = 2$ –6. Highest association to *E. coli* K12 and HepG2 cells is observed for COE2-3C-C6 (*i.e.* $n = 6$, $m = 1$). In this series, association to both cell types decrease for $n \leq 4$. Specifically, association to *E. coli* K12 decreases in a “step function” fashion from $83 \pm 1\%$ when $n = 6$ to less than 10% when $n \leq 4$. A decrease in association was also observed for HepG2 cells from $58 \pm 3\%$ to $14 \pm 11\%$ when n decreases from $n = 6$ to $n = 4$, but is followed by a trend that is difficult to discern given the experimental uncertainties.

By and large, *E. coli* K12 association tracks well with the hydrophobic content in the end groups (Fig. 2a, parameter m). Considering the linker length (Fig. 2b), the most hydrophobic compound ($n = 6$, $m = 1$) in the series also shows the highest cell association. A similar behavior has been observed in certain antimicrobial peptides (AMPs).³² However, the trend of cell association is less clear when $n < 6$ (*i.e.*, 4, 3, and 2). One possible line of thought is that apart from hydrophobicity, the distance between charges also impacts the equilibrium for intercalation within lipid bilayers because of structural mismatch.

It is worth noting that there is a similar cell association for COE2-3C-C4hexyl ($n = 4$, $m = 6$; $101 \pm 2\%$ and $14 \pm 1\%$ to *E. coli* K12 and HepG2 cells, respectively) and COE2-3C-C6 ($n = 6$,

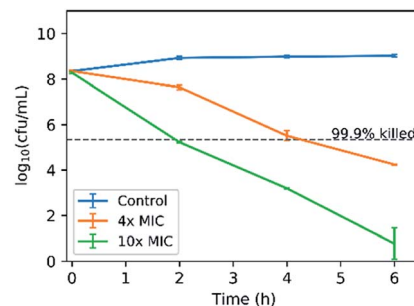


Fig. 3 Time-kill curves of *E. coli* K12 treated with COE2-3C-C4butyl at concentrations of 4× MIC ($16 \mu\text{g mL}^{-1}$) and 10× MIC ($40 \mu\text{g mL}^{-1}$).

$m = 1$; $83 \pm 1\%$ and $12 \pm 1\%$ to *E. coli* K12 and HepG2 cells, respectively). However, their MIC and IC_{50} values are different. Specifically, COE2-3C-C4hexyl has significantly higher antimicrobial activity and cytotoxicity ($\text{MIC} = 4 \mu\text{g mL}^{-1}$ and $\text{IC}_{50} = 15 \mu\text{g mL}^{-1}$) than COE2-3C-C6 ($\text{MIC} = 16 \mu\text{g mL}^{-1}$ and $\text{IC}_{50} > 1024 \mu\text{g mL}^{-1}$; Fig. 1a and b). Another interesting observation arises from a similar percent association to cells for COE2-3C-C4pentyl ($n = 4$, $m = 5$; $80 \pm 1\%$ and $23 \pm 12\%$ to *E. coli* K12 and HepG2 cells, respectively) and COE2-3C-C4butyl ($n = 4$, $m = 4$; $70 \pm 4\%$ and $22 \pm 1\%$ to *E. coli* K12 and HepG2 cells, respectively) (Fig. 2a). Despite their similar associations, they have different $\text{IC}_{50}/\text{MIC}$ values (64 for COE2-3C-C4pentyl and >256 for COE2-3C-C4butyl). These two phenomena imply contribution by a specific, molecular-structure-dependent mechanism that impacts cytotoxicity and antimicrobial efficacy.

Bactericidal activity

As a final note, the bactericidal activity of COE2-3C-C4butyl was investigated by an *in vitro* time-kill kinetic assay based on its optimal $\text{IC}_{50}/\text{MIC}$ ratio. It bears noting that according to a time-dependent cell association assay, COE2-3C-C4butyl reaches its maximal association to *E. coli* K12 within 30 minutes after treatment (Fig. S4†). *E. coli* K12 was challenged with COE2-3C-C4butyl at 4× MIC ($16 \mu\text{g mL}^{-1}$) and 10× MIC ($40 \mu\text{g mL}^{-1}$). As shown in Fig. 3, with an exposure at $40 \mu\text{g mL}^{-1}$ (10× MIC) for 2 hours, a 3 \log_{10} -fold decrease in colony forming units (cfu) was observed. Meanwhile, a 3 \log_{10} -fold decrease in colony forming units was also achieved within 4 hours with exposure at $16 \mu\text{g mL}^{-1}$ (4× MIC). As bactericidal activity is defined as greater than 3 \log_{10} -fold decrease in colony forming units, which is equivalent to 99.9% killing of the inoculum, these results indicate an excellent bactericidal activity.

Conclusions

In conclusion, we synthesized a series of distyrylbenzene-based COEs with different linker and terminal alkyl chain lengths to explore relationships between molecular features and antimicrobial efficacies and mammalian cell cytotoxicity. We found that hydrophobicity exerts influence over the degree of interactions between COEs and the cells. However, their biological activities were also found to be subtle structure dependent, as

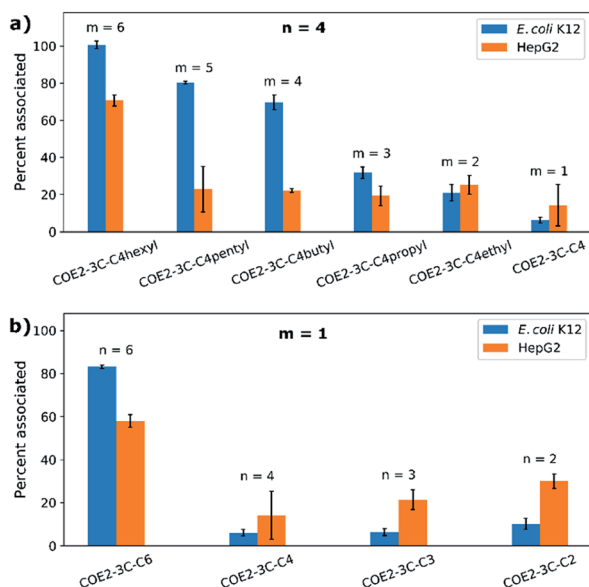


Fig. 2 Cell association of (a) COEs with different terminal alkyl groups (COE2-3C-C4 series, $n = 4$ and $m = 1$ –6) and (b) COEs with different linker length and methyl terminal groups ($n = 2$, 3, 4, and 6 and $m = 1$) at the concentration of $20 \mu\text{M}$ in HBSS with *E. coli* K12 and HepG2 cells.



seen in cases where COEs exhibit different antimicrobial efficacy or cytotoxicity but a similar degree of cell association. Changes in terminal alkyl chains of COEs have more significant impacts on MIC and cytotoxicity compared to changes in linker lengths. Although the mechanism underlying this observation is not clear, the data suggest the presence of poorly understood specific interactions that warrant further investigation. The optimal structure within the series of COEs studied here was found to be **COE2-3C-C4butyl** based on its highest IC_{50}/MIC value and minimal hemolytic activity. It is worth noting that this conclusion is specific for tested cell types. Importantly, **COE2-3C-C4butyl** also shows bactericidal activity against *E. coli* K12. These findings provide indications on how to strategically design antimicrobial COEs with low mammalian cell cytotoxicity based on the distyrylbenzene framework.

Experimental section

Minimum inhibitory concentration experiments

MICs were determined using broth microdilution method.³³ Prior to MIC experiment, bacterial strains were plated out for single colonies on agar plates from frozen glycerol stocks stored at -80°C . *E. coli* K12 (ATCC 47076) from a single colony on agar plate was inoculated in LB medium (10 g L^{-1} bactotryptone, 5 g L^{-1} yeast extract, and 10 g L^{-1} NaCl) and cultured at 37°C with 200 rpm shaking for 5 hours before use. Based on optical density at 600 nm, bacteria suspensions were diluted to the concentration around 1×10^6 cells per mL. Compound solutions were added to sterile 96-well plates and further diluted with 2-fold dilution successively. Later, by adding the bacteria suspensions with the same volume, the final concentrations of compound ranged from 0.5 to $128\text{ }\mu\text{g mL}^{-1}$ and the final concentration of bacteria suspension was 5×10^5 cells per mL. The assay plates were incubated at 37°C with 200 rpm shaking overnight. A group with no compound treatment was set as a negative control and a group with no bacteria was set as background. OD_{600} of the assay plates were measured on a plate reader (Tecan Spark 10M Multimode). The MICs were read as the lowest treatment concentration where less than 10% relative growth (treatment group relative to control) was found. A commercially available antibiotic, colistin, was used as a positive control and the MIC toward *E. coli* K12 was measured to be $1\text{ }\mu\text{g mL}^{-1}$, which is in accordance with what is reported in the literature.³⁴ Experiments were performed with biological replicates.

Mammalian cell viability assay

Cellular cytotoxicity was assessed using an MTT viability assay. HepG2 (ATCC HB-8065) cells were seeded at 1×10^4 cells per well in 96-well plates overnight at 37°C in Dulbecco's Modified Eagle's medium (DMEM, Gibco, GlutaMAX™) supplemented with 10% fetal bovine serum (FBS, Gibco, USDA-approved regions) before use. Compounds were serially diluted with 2-fold dilution in PBS and further diluted in culture media to afford a concentration ranging from 2 to $128\text{ }\mu\text{g mL}^{-1}$ and containing 10% PBS (v/v) each. For tests with higher

concentration of compounds, compound stock solutions were prepared in DMSO and diluted in culture media to obtain a concentration ranging from 128 to $1024\text{ }\mu\text{g mL}^{-1}$ and maintain the final DMSO concentration to be less than 1%. The previous culture media were replaced with 100 μL of treatment solutions per well, the assay plates were incubated at 37°C with 5% CO_2 . After 24 hours of incubation, cells were washed with PBS before adding 100 μL of fresh culture media and 10 μL of a 5 mg mL^{-1} solution of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT, Thermofisher Scientific) to each well. After incubation for 2–4 hours, 100 μL of DMSO as solubilizing solution was added to each well, and absorbance at 570 nm were measured on a plate reader (Tecan Spark 10M Multimode). Percent viability was determined by dividing background-corrected absorbance measurements by background-corrected measurements for untreated cells. The percentage of cell viability was plotted against the concentration of COE (Fig. S2†) and IC_{50} values were determined by curve-fitting (Fig. S1†).

Hemolytic activity

Fresh CD-1 mouse red blood cells (IC05-3054, Innovative Research, Inc.) were washed with PBS for three times with centrifuge at $500 \times g$ for 5 minutes. The pellet was resuspended to yield a 5% (v/v) suspension in PBS. 160 μL of compound solutions were added to a conical bottom 96-well plate as triplicates and dilute with 2-fold dilution sequentially. A 40 μL portion of 5% red blood cell solution was added to each well of the 96-well plate. The final concentration of compounds ranged from 16 to $1024\text{ }\mu\text{g mL}^{-1}$ and the final concentration of the red blood cells was 1%. Red blood cells with PBS alone was used as a negative control and with 1% Triton X-100 (Sigma-Aldrich) treatment as a positive control. After incubation for 1 hour at 37°C , cells were centrifuged at $800 \times g$. A 100 μL portion of the resulting supernatant was transferred to a flat-bottomed 96-well plate and analysed on a microplate reader *via* absorbance measurements at 450 nm. Percent hemolysis was determined by dividing background-corrected absorbance by background-corrected absorbance with 1% Triton X-100 treatment. The percentage of hemolysis was plotted against the concentration of COE (Fig. S3†).

Cell association experiments

Cell association experiments were performed in both Gram-negative bacteria (*E. coli* K12, ATCC 47076) and mammalian cells (HepG2, ATCC HB-8065). In these experiments, the stock solution of COEs were prepared directly with Hank's Balanced Salt Solution with no calcium and magnesium (HBSS, Gibco). Specific protocols for each cell type are described as follows:

Cell association in *E. coli* K12. Prior the experiments, a single colony of *E. coli* K12 from an agar plate was aerobically inoculated in LB medium at 37°C for 5 hours with 200 rpm shaking. Cells were spun down at $6500 \times g$ for 5 minutes and washed twice with HBSS and cell density was adjusted to be twice of 1 OD_{600} . In a 96-well PCR plate, 100 μL of the cell suspension was



mixed with 100 μL of a COE solution in HBSS in each well. The final concentration of COEs was 20 μM . The plate was sealed prior incubation. After the incubation at 37 $^{\circ}\text{C}$ for 2 hours, the cells were spun down at $3000 \times g$ for 30 minutes and 100 μL of supernatant from each well was transferred to another 96-well plate. Amount of the COEs left in the supernatant was determined by measuring absorbance of the supernatant at 380 nm using Tecan Infinite M200 Plate Reader.

Cell association in HepG2 cells. Prior the experiments, HepG2 cells were cultured in DMEM supplemented with 10% FBS. Cells were then harvested from culture and washed three times with HBSS. Then cell density was adjusted to be around $4 \times 10^6 \text{ mL}^{-1}$. In a 96-well PCR plate, 100 μL of the cell suspension was mixed with 100 μL of a COE solution in HBSS in each well. The final concentration of COEs was 20 μM . The plate was sealed prior incubation. After the incubation at 37 $^{\circ}\text{C}$ for 2 hours, the cells were centrifuged at $400 \times g$ for 10 minutes and 100 μL of supernatant from each well was transferred to a flat bottom 96-well plate. Amount of the COEs left in the supernatant was determined by measuring absorbance of the supernatant at 360 nm using Tecan Spark 10M Multimode Plate Reader.

Time-kill kinetics

E. coli K12 (ATCC 47076) from a single colony on agar plate was inoculated in LB medium and cultured at 37 $^{\circ}\text{C}$ and 200 rpm shaking for around 4 hours before use. The bacteria suspension was diluted in LB to afford an OD_{600} value to be around 0.15. The diluted bacteria suspension was then challenged with COE2-3C-C4butyl at $10 \times \text{MIC}$ ($40 \mu\text{g mL}^{-1}$) and $4 \times \text{MIC}$ ($16 \mu\text{g mL}^{-1}$) respectively in culture tubes at 37 $^{\circ}\text{C}$ and 280 rpm shaking. Bacteria suspension without treatment was used as a control. At intervals, 100 μL aliquots from each sample were diluted with 10-fold serial dilution in PBS and were further plated on LB agar plates with 100 μL inoculation. After overnight incubation at 37 $^{\circ}\text{C}$, colonies were counted and cfu per mL was calculated. Experiments were performed in triplicate.

Conflicts of interest

The authors own a related patent filed as PCT/US19/23411. This patent belongs to University of California, Santa Barbara.

Acknowledgements

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References

- 1 J. O'Neill, *Tackling drug-resistant infections globally: final report and recommendations, The review on antimicrobial resistance*, London, 2016.
- 2 M. R. L. Stone, M. S. Butler, W. Phetsang, M. A. Cooper and M. A. T. Blaskovich, *Trends Biotechnol.*, 2018, **36**, 523–536.
- 3 S. C. Davies, T. Fowler, J. Watson, D. M. Livermore and D. Walker, *Lancet*, 2013, **381**, 1606–1609.
- 4 H. Gelband and R. Laxminarayan, *Trends Microbiol.*, 2015, **23**, 524–526.
- 5 I. Torjesen, *BMJ*, 2013, **346**, f1597.
- 6 T. B. Nielsen, E. P. Brass, D. N. Gilbert, J. G. Bartlett and B. Spellberg, *N. Engl. J. Med.*, 2019, **381**, 503–505.
- 7 J. R. Fitchett, *Lancet Infect. Dis.*, 2015, **15**, 1125–1127.
- 8 M. Perros, *Science*, 2015, **347**, 1062–1064.
- 9 T. F. Schäberle and I. M. Hack, *Trends Microbiol.*, 2014, **22**, 165–167.
- 10 H. Cho, T. Uehara and T. G. Bernhardt, *Cell*, 2014, **159**, 1300–1311.
- 11 K. Kimura and T. D. H. Bugg, *Nat. Prod. Rep.*, 2003, **20**, 252–273.
- 12 R. G. Panchal, R. L. Ulrich, D. Lane, M. M. Butler, C. Houseweart, T. Opperman, J. D. Williams, N. P. Peet, D. T. Moir, T. Nguyen, R. Gussio, T. Bowlin and S. Bavari, *Antimicrob. Agents Chemother.*, 2009, **53**, 4283–4291.
- 13 K. Ishiguro, S. Sakiyama, K. Takahashi and T. Arai, *Biochemistry*, 1978, **17**, 2545–2550.
- 14 T. K. Mukhtar and G. D. Wright, *Chem. Rev.*, 2005, **105**, 529–542.
- 15 C. R. Bourne, *Antibiotics*, 2014, **3**, 1–28.
- 16 Y. Wang, S. D. Jett, J. Crum, K. S. Schanze, E. Y. Chi and D. G. Whitten, *Langmuir*, 2013, **29**, 781–792.
- 17 D. Liu, S. Choi, B. Chen, R. J. Doerksen, D. J. Clements, J. D. Winkler, M. L. Klein and W. F. DeGrado, *Angew. Chem., Int. Ed.*, 2004, **43**, 1158–1162.
- 18 S. Choi, A. Issacs, D. Clements, D. Liu, H. Kim, R. W. Scott, J. D. Winkler and W. F. DeGrado, *Proc. Natl. Acad. Sci. U.S.A.*, 2009, **106**, 6968–6973.
- 19 C. Ghosh and J. Haldar, *ChemMedChem*, 2015, **10**, 1606–1624.
- 20 F. Nederberg, Y. Zhang, J. P. K. Tan, K. Xu, H. Wang, C. Yang, S. Gao, X. D. Guo, K. Fukushima, L. Li, J. L. Hedrick and Y.-Y. Yang, *Nat. Chem.*, 2011, **3**, 409–414.
- 21 J. G. Hurdle, A. J. O'Neill, I. Chopra and R. E. Lee, *Nat. Rev. Microbiol.*, 2011, **9**, 62–75.
- 22 H. Yan, C. Catania and G. C. Bazan, *Adv. Mater.*, 2015, **27**, 2958–2973.
- 23 L. E. Garner, J. Park, S. M. Dyar, A. Chworos, J. J. Sumner and G. C. Bazan, *J. Am. Chem. Soc.*, 2010, **132**, 10042–10052.
- 24 B. Wang, S. L. Fronk, Z. D. Rengert, J. Limwongyut and G. C. Bazan, *Chem. Mater.*, 2018, **30**, 5836–5840.
- 25 H. Yan, Z. D. Rengert, A. W. Thomas, C. Rehmann, J. Hinks and G. C. Bazan, *Chem. Sci.*, 2016, **7**, 5714–5722.
- 26 J. Hinks, Y. Wang, W. H. Poh, B. C. Donose, A. W. Thomas, S. Wuertz, S. C. J. Loo, G. C. Bazan, S. Kjelleberg, Y. Mu and T. Seviour, *Langmuir*, 2014, **30**, 2429–2440.



- 27 E. Strandberg, S. Esteban-Martín, A. S. Ulrich and J. Salgado, *Biochim. Biophys. Acta Biomembr.*, 2012, **1818**, 1242–1249.
- 28 C. Zhou, G. W. N. Chia, J. C. S. Ho, T. Seviour, T. Sailov, B. Liedberg, S. Kjelleberg, J. Hinks and G. C. Bazan, *Angew. Chem., Int. Ed.*, 2018, **57**, 8069–8072.
- 29 A. L. Niles, R. A. Moravec and T. L. Riss, *Expert Opin. Drug Discov.*, 2008, **3**, 655–669.
- 30 H. Yamamura, A. Miyagawa, H. Sugiyama, K. Murata, T. Mabuti, R. Mitsunashi, T. Hagiwara, M. Nonaka, K. Tanimoto and H. Tomita, *ChemistrySelect*, 2016, **1**, 469–472.
- 31 J. Hoque, M. M. Konai, S. S. Sequeira, S. Samaddar and J. Haldar, *J. Med. Chem.*, 2016, **59**, 10750–10762.
- 32 T. Wieprecht, M. Dathe, E. Krause, M. Beyermann, W. L. Maloy, D. L. MacDonald and M. Bienert, *FEBS Lett.*, 1997, **147**, 135–140.
- 33 I. Wiegand, K. Hilpert and R. E. W. Hancock, *Nat. Protoc.*, 2008, **3**, 163–175.
- 34 N. S. Sundaramoorthy, P. Suresh, S. S. Ganesan, A. GaneshPrasad and S. Nagarajan, *Sci. Rep.*, 2019, **9**, 19845.

